

SGS

AXYS

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SGS AXYS Client No.: 4972

Client Address: AECOM
1111 Third Avenue, Suite 1600
Seattle, WA, US, 98101

The SGS AXYS contact for these data is Sean Campbell.

BATCH SUMMARY

| | |
|--|---|
| Batch ID: WG66481 | Date: 04-Feb-2019 |
| Analysis Type: PCB Congener | Matrix Type: Filter |
| BATCH MAKEUP | |
| Contract: 4972 Samples: L30523-1 PDI-WS-T05-1811 L30523-2 PDI-WS-T01-1811 L30523-3 PDI-WS-T03-1811 L30523-4 PDI-WS-T07-1811 L30523-5 PDI-WS-T02-1811 L30523-6 PDI-WS-T04-1812 L30523-7 PDI-WS-T06-1811 L30523-8 PDI-RB-XF-181129 | Blank: WG66481-101 Reference or Spike: WG66481-102 |
| <p>Resubmission 06-Feb-19: The data are being resubmitted to include the correct 'sample' file for the EDD. No other changes have been made to the data.</p> <p>Comments:</p> <ol style="list-style-type: none"> 1. Data are considered final. 2. Data are not blank corrected. Blank data should be taken into consideration when evaluating sample data. 3. Blank data should be evaluated against specifications using the same blank sample size as the size of the client samples. 4. An interference known to originate during extraction from the high boiling point of the toluene was observed in the OPR, WG66481-102, the Lab Blank, WG66481-101, and sample 'PDI-RB-XF-181129' (AXYS ID: L30523-8) near the mono- and/or di-substituted PCBs. The affected compounds have been flagged 'NQ' – not quantifiable. 5. The relative retention times (RRT) of several PCB congeners were observed to be slightly outside of the nominal RRT acceptance windows in the Lab Blank, WG66481-101, and sample 'PDI-RB-XF-181129' (AXYS ID: L30523-8). The congeners were determined to be present based on a detailed inspection of sample and calibration chromatography. 6. The concentration of clean-up standard ¹³C-labeled PCB-28 in closing calibration filename PB9C_028 S:9 was observed to be slightly below the method control limit. Given that clean-up standards are not used for target analyte quantification, sample data are not affected by this variance. | |

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February 2017

FQA-006 Rev. 4. 20-Sep-2013

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T05-1811
Sample Collection:
27-Nov-2018 15:44

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-1 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 30-Jan-2019 Time: 15:03:24 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_027 S: 6 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_027 S: 1 |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | G | 22.7 | 1.55 (S) | 2.76 | 1.001 |
| 3-MoCB | 2 | | | 15.0 | 1.93 (S) | 3.52 | 0.989 |
| 4-MoCB | 3 | | | 17.0 | 1.34 (S) | 3.35 | 1.001 |
| 2,2'-DiCB | 4 | | | 60.0 | 5.81 (S) | 1.50 | 1.001 |
| 2,3-DiCB | 5 | | U | | 5.21 (S) | | |
| 2,3'-DiCB | 6 | | | 31.1 | 4.72 (S) | 1.47 | 1.176 |
| 2,4-DiCB | 7 | | K | 7.05 | 4.84 (S) | 1.97 | 1.160 |
| 2,4'-DiCB | 8 | | | 140 | 4.32 (S) | 1.50 | 1.208 |
| 2,5-DiCB | 9 | | K | 8.68 | 4.56 (S) | 2.27 | 1.146 |
| 2,6-DiCB | 10 | | U | | 4.66 (S) | | |
| 3,3'-DiCB | 11 | | | 433 | 5.25 (S) | 1.47 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C | 17.2 | 4.99 (S) | 1.74 | 0.985 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 4.84 (S) | | |
| 4,4'-DiCB | 15 | | | 112 | 4.04 (S) | 1.42 | 1.001 |
| 2,2',3-TriCB | 16 | | | 145 | 0.868 (Q) | 1.04 | 1.166 |
| 2,2',4-TriCB | 17 | | | 202 | 0.868 (Q) | 1.13 | 1.139 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 342 | 0.868 (Q) | 1.06 | 1.114 |
| 2,2',6-TriCB | 19 | | | 61.1 | 0.868 (Q) | 1.03 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 612 | 1.87 (S) | 0.98 | 0.848 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 249 | 1.81 (S) | 0.98 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 189 | 2.11 (S) | 0.93 | 0.872 |
| 2,3,5-TriCB | 23 | | U | | 2.06 (S) | | |
| 2,3,6-TriCB | 24 | | J | 4.29 | 0.868 (Q) | 1.09 | 1.161 |
| 2,3',4-TriCB | 25 | | | 100 | 1.60 (S) | 1.00 | 0.825 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 94.6 | 1.90 (S) | 0.90 | 1.304 |
| 2,3',6-TriCB | 27 | | | 31.7 | 0.868 (Q) | 1.07 | 1.153 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 465 | 1.77 (S) | 0.94 | 0.837 |
| 2,4',6-TriCB | 32 | | | 162 | 1.84 (S) | 0.95 | 1.199 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | J | 2.86 | 1.99 (S) | 1.01 | 1.276 |
| 3,3',4-TriCB | 35 | | | 14.2 | 2.02 (S) | 0.90 | 0.985 |
| 3,3',5-TriCB | 36 | | K J | 5.46 | 1.93 (S) | 1.21 | 0.933 |
| 3,4,4'-TriCB | 37 | | | 142 | 1.54 (S) | 0.90 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 1.82 (S) | | |
| 3,4',5-TriCB | 39 | | | 8.77 | 1.87 (S) | 1.07 | 0.947 |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 378 | 0.868 (Q) | 0.76 | 1.337 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 177 | 0.868 (Q) | 0.77 | 1.312 |
| 2,2',3,5'-TeCB | 43 | | | 25.9 | 0.868 (Q) | 0.67 | 1.248 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 1470 | 0.868 (Q) | 0.77 | 1.287 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 414 | 0.868 (Q) | 0.74 | 1.148 |
| 2,2',3,6'-TeCB | 46 | | | 45.9 | 0.868 (Q) | 0.77 | 1.161 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 145 | 0.868 (Q) | 0.79 | 1.275 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 419 | 0.868 (Q) | 0.78 | 1.260 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 120 | 0.868 (Q) | 0.79 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 720 | 0.868 (Q) | 0.76 | 1.234 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | | 7.68 | 0.868 (Q) | 0.83 | 1.001 |
| 2,3,3',4'-TeCB | 55 | | | 10.8 | 6.06 (S) | 0.70 | 0.890 |
| 2,3,3',4'-TeCB | 56 | | | 374 | 5.73 (S) | 0.73 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 5.40 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 5.60 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 58.0 | 0.868 (Q) | 0.80 | 1.302 |
| 2,3,4,4'-TeCB | 60 | | | 132 | 5.47 (S) | 0.71 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 1180 | 5.31 (S) | 0.73 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | | 27.3 | 5.35 (S) | 0.76 | 0.864 |
| 2,3,4',6'-TeCB | 64 | | | 284 | 0.868 (Q) | 0.76 | 1.349 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | | 706 | 5.43 (S) | 0.77 | 0.884 |
| 2,3',4,5'-TeCB | 67 | | K | 17.9 | 4.56 (S) | 0.56 | 0.857 |
| 2,3',4,5'-TeCB | 68 | | | 214 | 5.10 (S) | 0.74 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | K J | 5.63 | 5.31 (S) | 0.98 | 0.823 |
| 2,3',5',6'-TeCB | 73 | | U | | 0.868 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 66.4 | 4.40 (S) | 0.75 | 1.001 |
| 3,3',4,5'-TeCB | 78 | | U | | 5.15 (S) | | |
| 3,3',4,5'-TeCB | 79 | | | 11.7 | 4.23 (S) | 0.81 | 0.970 |
| 3,3',5,5'-TeCB | 80 | | U | | 4.86 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 3.98 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 123 | 0.868 (Q) | 1.60 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 550 | 0.868 (Q) | 1.52 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 253 | 0.868 (Q) | 1.45 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 180 | 0.868 (Q) | 1.52 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 639 | 0.868 (Q) | 1.56 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 150 | 0.868 (Q) | 1.60 | 1.154 |
| 2,2',3,4,6'-PeCB | 89 | | | 13.0 | 0.868 (Q) | 1.62 | 1.182 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 887 | 0.868 (Q) | 1.56 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 166 | 0.868 (Q) | 1.72 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 742 | 0.868 (Q) | 1.59 | 1.121 |
| 2,2',3,5,6'-PeCB | 94 | | | 8.07 | 0.868 (Q) | 1.48 | 1.102 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | | 9.20 | 0.868 (Q) | 1.57 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | | 15.6 | 0.868 (Q) | 1.45 | 1.094 |
| 2,2',4,6,6'-PeCB | 104 | | K J | 1.26 | 0.868 (Q) | 0.32 | 1.003 |
| 2,3,3',4,4'-PeCB | 105 | | | 300 | 2.36 (S) | 1.53 | 1.000 |
| 2,3,3',4,5-PeCB | 106 | | U | | 2.76 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C | 34.5 | 2.81 (S) | 1.45 | 0.991 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 58.7 | 2.57 (S) | 1.34 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 1070 | 0.868 (Q) | 1.59 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.868 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.868 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | | 14.9 | 2.32 (S) | 1.58 | 1.000 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 681 | 2.45 (S) | 1.50 | 1.001 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 4.79 | 0.868 (Q) | 1.27 | 0.958 |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.868 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | K | 14.6 | 3.31 (S) | 1.10 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K | 14.8 | 2.54 (S) | 1.33 | 1.000 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 6.06 | 2.56 (S) | 2.18 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 2.42 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 161 | 1.63 (S) | 1.11 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 1340 | 1.83 (S) | 1.25 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 72.9 | 2.17 (S) | 1.27 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | | 14.6 | 2.40 (S) | 1.37 | 1.159 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 475 | 2.60 (S) | 1.20 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 21.0 | 2.37 (S) | 1.42 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 64.5 | 2.34 (S) | 1.29 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 500 | 0.868 (Q) | 1.24 | 1.103 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 159 | 0.868 (Q) | 1.34 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | | 41.4 | 2.05 (S) | 1.09 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C K | 20.0 | 2.11 (S) | 0.99 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 230 | 2.04 (S) | 1.10 | 0.904 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 2.50 (S) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 58.4 | 0.868 (Q) | 1.22 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.868 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 223 | 1.90 (S) | 1.23 | 0.885 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 1010 | 2.12 (S) | 1.27 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | K J | 3.42 | 0.868 (Q) | 2.07 | 1.083 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | K J | 3.77 | 0.868 (Q) | 2.54 | 1.012 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | J | 1.59 | 0.868 (Q) | 1.22 | 1.006 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 1090 | 1.62 (S) | 1.29 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | J | 1.76 | 0.868 (Q) | 1.12 | 1.001 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 143 | 1.79 (S) | 1.22 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 117 | 1.37 (S) | 1.29 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | K | 18.7 | 1.36 (S) | 1.00 | 0.981 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

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|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 1.46 (S) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | K J | 3.03 | 1.42 (S) | 1.48 | 0.989 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 95.8 | 1.43 (S) | 1.24 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 1.81 (S) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 55.5 | 1.46 (S) | 1.14 | 1.000 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 1.46 (S) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 398 | 0.868 (Q) | 0.99 | 1.000 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 104 | 0.868 (Q) | 1.01 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 67.7 | 0.868 (Q) | 1.06 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 321 | 0.868 (Q) | 1.03 | 1.133 |
| 2,2',3,3',4,5',6-HpCB | 175 | | | 15.1 | 0.868 (Q) | 1.01 | 1.102 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 46.6 | 0.868 (Q) | 1.05 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 238 | 0.868 (Q) | 1.03 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 84.1 | 0.868 (Q) | 0.94 | 1.085 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 162 | 0.868 (Q) | 1.06 | 1.009 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 1080 | 0.868 (Q) | 1.07 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | K J | 4.04 | 0.868 (Q) | 0.67 | 1.156 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | J | 2.35 | 0.868 (Q) | 0.96 | 1.116 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 241 | 0.868 (Q) | 0.98 | 1.127 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | J | 2.10 | 0.868 (Q) | 0.98 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.868 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 443 | 0.868 (Q) | 1.06 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | K J | 1.35 | 0.868 (Q) | 1.26 | 1.001 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 14.0 | 0.868 (Q) | 1.14 | 1.001 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 76.3 | 0.868 (Q) | 0.99 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | K | 15.8 | 0.868 (Q) | 1.41 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.868 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | | 245 | 0.868 (Q) | 0.95 | 0.991 |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | | 97.9 | 0.868 (Q) | 0.88 | 0.946 |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | | 140 | 0.868 (Q) | 0.95 | 0.916 |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C | 45.0 | 0.868 (Q) | 0.78 | 1.046 |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C | 317 | 0.868 (Q) | 0.92 | 1.115 |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | | 38.0 | 0.868 (Q) | 0.85 | 1.023 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | K | 75.8 | 0.868 (Q) | 1.05 | 1.000 |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | | 213 | 0.868 (Q) | 0.89 | 0.920 |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.868 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | | 13.4 | 0.868 (Q) | 0.80 | 1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 167 | 3.57 (S) | 0.79 | 1.000 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | K | 20.7 | 2.69 (S) | 0.94 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 60.4 | 3.00 (S) | 0.76 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 129 | 0.868 (Q) | 1.04 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T05-1811
Sample Collection:
27-Nov-2018 15:44

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 15:03:24
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-1
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_027 S: 6
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_027 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | G | 4000 | 917 | 22.9 | 3.19 | 0.717 |
| 13C12-4-MoCB | 3L | | | 4000 | 1150 | 28.7 | 3.12 | 0.856 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1250 | 31.3 | 1.55 | 0.873 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1470 | 36.8 | 1.53 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2300 | 57.5 | 1.04 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1290 | 32.3 | 0.99 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1800 | 45.1 | 0.79 | 0.810 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1750 | 43.7 | 0.71 | 1.395 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1870 | 46.9 | 0.67 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2030 | 50.8 | 1.61 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1890 | 47.2 | 1.54 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1730 | 43.2 | 1.58 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1850 | 46.2 | 1.53 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1830 | 45.8 | 1.55 | 1.151 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1790 | 44.7 | 1.51 | 1.300 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2950 | 73.8 | 1.26 | 0.786 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 5410 | 67.6 | 1.27 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2760 | 69.0 | 1.27 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2830 | 70.8 | 1.26 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 3350 | 83.8 | 1.08 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 3180 | 79.4 | 1.04 | 0.872 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 3340 | 83.6 | 1.08 | 0.712 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1940 | 48.5 | 0.95 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2900 | 72.4 | 0.91 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 3140 | 78.5 | 0.80 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3710 | 92.7 | 0.74 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3630 | 90.7 | 0.76 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 4110 | 103 | 1.18 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1420 | 35.6 | 1.02 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2620 | 65.5 | 1.60 | 1.088 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3260 | 81.4 | 1.10 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T01-1811
Sample Collection:
28-Nov-2018 14:26

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-2 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 30-Jan-2019 Time: 16:07:37 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_027 S: 7 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_027 S: 1 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | | 16.9 | 1.09 (S) | 3.16 | 1.001 |
| 3-MoCB | 2 | | | 15.5 | 1.32 (S) | 3.04 | 0.987 |
| 4-MoCB | 3 | | | 14.3 | 0.897 (S) | 3.22 | 1.000 |
| 2,2'-DiCB | 4 | | | 31.9 | 4.92 (S) | 1.68 | 1.001 |
| 2,3-DiCB | 5 | | U | | 4.84 (S) | | |
| 2,3'-DiCB | 6 | | | 20.6 | 4.39 (S) | 1.38 | 1.177 |
| 2,4-DiCB | 7 | | K J | 4.69 | 4.49 (S) | 2.28 | 1.159 |
| 2,4'-DiCB | 8 | | | 79.3 | 4.02 (S) | 1.43 | 1.208 |
| 2,5-DiCB | 9 | | K J | 6.75 | 4.24 (S) | 1.96 | 1.147 |
| 2,6-DiCB | 10 | | U | | 4.33 (S) | | |
| 3,3'-DiCB | 11 | | | 300 | 4.88 (S) | 1.54 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C K | 11.0 | 4.64 (S) | 2.89 | 0.984 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 4.50 (S) | | |
| 4,4'-DiCB | 15 | | | 58.7 | 4.00 (S) | 1.55 | 1.001 |
| 2,2',3-TriCB | 16 | | | 33.1 | 0.862 (Q) | 1.18 | 1.166 |
| 2,2',4-TriCB | 17 | | | 68.3 | 0.862 (Q) | 1.02 | 1.139 |
| 2,2',5-TriCB | 18 | 18 + 30 | C G | 87.5 | 0.862 (Q) | 1.08 | 1.114 |
| 2,2',6-TriCB | 19 | | K | 26.1 | 0.862 (Q) | 1.26 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 187 | 0.862 (Q) | 1.00 | 0.848 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 83.8 | 0.862 (Q) | 0.97 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 55.5 | 0.885 (S) | 1.02 | 0.872 |
| 2,3,5-TriCB | 23 | | K J | 1.31 | 0.866 (S) | 0.77 | 1.284 |
| 2,3,6-TriCB | 24 | | K J | 1.72 | 0.862 (Q) | 1.83 | 1.159 |
| 2,3',4-TriCB | 25 | | | 27.5 | 0.862 (Q) | 1.02 | 0.825 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 30.2 | 0.862 (Q) | 0.93 | 1.302 |
| 2,3',6-TriCB | 27 | | | 10.5 | 0.862 (Q) | 0.99 | 1.152 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 126 | 0.862 (Q) | 0.92 | 0.837 |
| 2,4',6-TriCB | 32 | | G | 42.9 | 0.862 (Q) | 1.02 | 1.198 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | K J | 2.80 | 0.862 (Q) | 1.76 | 1.275 |
| 3,3',4-TriCB | 35 | | J | 5.28 | 0.862 (Q) | 0.99 | 0.985 |
| 3,3',5-TriCB | 36 | | K J | 2.48 | 0.862 (Q) | 2.03 | 0.932 |
| 3,4,4'-TriCB | 37 | | | 52.2 | 0.862 (Q) | 0.93 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.862 (Q) | | |
| 3,4',5-TriCB | 39 | | J | 3.14 | 0.862 (Q) | 0.98 | 0.947 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 103 | 0.862 (Q) | 0.74 | 1.338 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 55.3 | 0.862 (Q) | 0.78 | 1.312 |
| 2,2',3,5'-TeCB | 43 | | K J | 3.07 | 0.862 (Q) | 1.32 | 1.245 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 455 | 0.862 (Q) | 0.77 | 1.289 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 222 | 0.862 (Q) | 0.75 | 1.149 |
| 2,2',3,6'-TeCB | 46 | | K | 13.5 | 0.862 (Q) | 0.92 | 1.161 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 31.3 | 0.862 (Q) | 0.76 | 1.275 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 168 | 0.862 (Q) | 0.74 | 1.261 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 41.7 | 0.862 (Q) | 0.74 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 293 | 0.862 (Q) | 0.77 | 1.236 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | J | 6.57 | 0.862 (Q) | 0.78 | 1.001 |
| 2,3,3',4'-TeCB | 55 | | U | | 2.71 (S) | | |
| 2,3,3',4'-TeCB | 56 | | | 96.0 | 2.56 (S) | 0.73 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 2.42 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 2.51 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 17.9 | 0.862 (Q) | 0.74 | 1.303 |
| 2,3,4,4'-TeCB | 60 | | | 36.3 | 2.45 (S) | 0.70 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 393 | 2.38 (S) | 0.76 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | | 10.5 | 2.40 (S) | 0.78 | 0.865 |
| 2,3,4',6'-TeCB | 64 | | | 89.1 | 0.862 (Q) | 0.82 | 1.349 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | | 273 | 2.43 (S) | 0.68 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | J | 5.98 | 2.04 (S) | 0.83 | 0.857 |
| 2,3',4,5'-TeCB | 68 | | | 64.2 | 2.28 (S) | 0.74 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | K J | 5.81 | 2.38 (S) | 1.15 | 0.824 |
| 2,3',5',6'-TeCB | 73 | | U | | 0.862 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 26.5 | 1.95 (S) | 0.68 | 1.000 |
| 3,3',4,5'-TeCB | 78 | | U | | 2.30 (S) | | |
| 3,3',4,5'-TeCB | 79 | | | 8.83 | 1.89 (S) | 0.83 | 0.971 |
| 3,3',5,5'-TeCB | 80 | | U | | 2.17 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 1.85 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 61.9 | 1.01 (S) | 1.39 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 357 | 1.01 (S) | 1.56 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 131 | 1.16 (S) | 1.41 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 88.6 | 0.862 (Q) | 1.56 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 356 | 0.862 (Q) | 1.55 | 0.902 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 90.7 | 1.01 (S) | 1.67 | 1.155 |
| 2,2',3,4,6'-PeCB | 89 | | J | 5.76 | 1.06 (S) | 1.50 | 1.182 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 578 | 0.862 (Q) | 1.57 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 121 | 0.980 (S) | 1.46 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 425 | 0.983 (S) | 1.53 | 1.121 |
| 2,2',3,5,6'-PeCB | 94 | | J | 5.29 | 1.11 (S) | 1.53 | 1.102 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | J | 5.49 | 0.862 (Q) | 1.52 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | K | 10.2 | 0.909 (S) | 1.89 | 1.094 |
| 2,2',4,6,6'-PeCB | 104 | | K J | 2.21 | 0.862 (Q) | 1.92 | 1.001 |
| 2,3,3',4,4'-PeCB | 105 | | K | 158 | 1.85 (S) | 1.32 | 1.000 |
| 2,3,3',4,5-PeCB | 106 | | U | | 2.21 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C K | 18.3 | 2.25 (S) | 1.78 | 0.991 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 37.2 | 2.05 (S) | 1.58 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 633 | 0.862 (Q) | 1.54 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | K J | 0.889 | 0.862 (Q) | 0.71 | 0.946 |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.862 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | | 9.59 | 1.88 (S) | 1.61 | 1.001 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 426 | 2.00 (S) | 1.52 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 3.59 | 0.862 (Q) | 1.88 | 0.959 |
| 2,3',4,5',6-PeCB | 121 | | K J | 1.05 | 0.862 (Q) | 0.43 | 1.201 |
| 2',3,3',4,5-PeCB | 122 | | K J | 6.35 | 2.65 (S) | 0.95 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K J | 6.72 | 2.00 (S) | 1.14 | 1.001 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 2.66 | 2.01 (S) | 0.74 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 1.93 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 95.8 | 0.862 (Q) | 1.38 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 723 | 0.862 (Q) | 1.26 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 44.1 | 0.862 (Q) | 1.27 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | | 9.34 | 0.862 (Q) | 1.14 | 1.158 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 253 | 0.862 (Q) | 1.26 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 13.9 | 0.862 (Q) | 1.11 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 38.4 | 0.862 (Q) | 1.30 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 285 | 0.862 (Q) | 1.25 | 1.102 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 93.3 | 0.862 (Q) | 1.15 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | | 20.4 | 0.862 (Q) | 1.39 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C K | 14.4 | 0.862 (Q) | 1.48 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 116 | 0.862 (Q) | 1.28 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 0.862 (Q) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 29.0 | 0.862 (Q) | 1.27 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.862 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 139 | 0.862 (Q) | 1.34 | 0.884 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 585 | 0.862 (Q) | 1.22 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | K J | 3.06 | 0.862 (Q) | 0.68 | 1.083 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | J | 4.56 | 0.862 (Q) | 1.06 | 1.011 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | K J | 1.31 | 0.862 (Q) | 2.91 | 1.006 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 609 | 0.862 (Q) | 1.24 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | K J | 1.46 | 0.862 (Q) | 0.54 | 1.001 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 82.0 | 0.862 (Q) | 1.13 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 63.0 | 0.862 (Q) | 1.33 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | U | | 0.862 (Q) | | |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 0.862 (Q) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | K J | 0.933 | 0.862 (Q) | 0.55 | 0.988 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 50.3 | 0.862 (Q) | 1.36 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 0.862 (Q) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 30.1 | 0.862 (Q) | 1.15 | 1.000 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 0.862 (Q) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 184 | 0.862 (Q) | 1.01 | 1.001 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 52.8 | 0.862 (Q) | 1.11 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | K | 35.8 | 0.862 (Q) | 1.22 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 158 | 0.862 (Q) | 1.08 | 1.132 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K J | 6.22 | 0.862 (Q) | 0.64 | 1.103 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 23.1 | 0.862 (Q) | 1.05 | 1.033 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 123 | 0.862 (Q) | 1.03 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 40.0 | 0.862 (Q) | 0.99 | 1.085 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 77.0 | 0.862 (Q) | 1.06 | 1.009 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 527 | 0.862 (Q) | 1.03 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | K J | 3.15 | 0.862 (Q) | 0.60 | 1.156 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | K J | 2.34 | 0.862 (Q) | 1.68 | 1.115 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 109 | 0.862 (Q) | 1.00 | 1.127 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | K J | 1.25 | 0.862 (Q) | 1.56 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.862 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 215 | 0.862 (Q) | 1.06 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | J | 0.931 | 0.862 (Q) | 1.00 | 1.000 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | K | 7.30 | 0.862 (Q) | 1.79 | 1.000 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 36.5 | 0.862 (Q) | 1.15 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | | 8.44 | 0.862 (Q) | 1.10 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.862 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | | 111 | 0.862 (Q) | 0.84 | 0.991 |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | | 46.7 | 0.862 (Q) | 0.97 | 0.945 |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | | 63.4 | 0.862 (Q) | 1.02 | 0.916 |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C K | 23.1 | 0.862 (Q) | 0.73 | 1.047 |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C | 136 | 0.862 (Q) | 0.85 | 1.115 |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | | 16.4 | 0.862 (Q) | 0.92 | 1.023 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | K | 35.5 | 0.862 (Q) | 0.75 | 1.001 |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | | 90.4 | 0.862 (Q) | 0.91 | 0.920 |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.862 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | K J | 6.00 | 0.862 (Q) | 1.11 | 1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 101 | 3.43 (S) | 0.76 | 1.001 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | K | 10.9 | 2.59 (S) | 0.96 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 35.0 | 2.88 (S) | 0.74 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 104 | 0.862 (Q) | 0.99 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T01-1811
Sample Collection:
28-Nov-2018 14:26

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 16:07:37
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-2
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_027 S: 7
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_027 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|---------------------------|-------------|--------------------------|----------------|----------------|-------------------|------------------------|-------|
| 13C12-2-MoCB | 1L | | | 4000 | 1190 | 29.8 | 3.15 | 0.718 |
| 13C12-4-MoCB | 3L | | | 4000 | 1490 | 37.2 | 3.07 | 0.857 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1520 | 37.9 | 1.58 | 0.873 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1640 | 41.0 | 1.54 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2550 | 63.8 | 1.02 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1390 | 34.7 | 1.03 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1960 | 49.0 | 0.79 | 0.810 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1850 | 46.3 | 0.70 | 1.396 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1870 | 46.8 | 0.68 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 1940 | 48.6 | 1.59 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1820 | 45.5 | 1.56 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1630 | 40.8 | 1.60 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1700 | 42.6 | 1.52 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1740 | 43.4 | 1.52 | 1.150 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1740 | 43.4 | 1.50 | 1.299 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2620 | 65.6 | 1.28 | 0.787 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 5580 | 69.7 | 1.24 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2740 | 68.6 | 1.23 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2870 | 71.8 | 1.26 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 2930 | 73.2 | 1.06 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 2660 | 66.5 | 1.02 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2720 | 68.0 | 1.06 | 0.712 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1950 | 48.7 | 0.93 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2520 | 62.9 | 0.90 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2940 | 73.5 | 0.83 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3410 | 85.3 | 0.79 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3340 | 83.5 | 0.77 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3640 | 90.9 | 1.15 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1510 | 37.8 | 0.98 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2480 | 62.1 | 1.61 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3250 | 81.2 | 1.03 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T03-1811
Sample Collection:
27-Nov-2018 16:22

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-3 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 30-Jan-2019 Time: 17:11:47 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_027 S: 8 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_027 S: 1 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | G | 17.4 | 1.06 (S) | 3.14 | 1.001 |
| 3-MoCB | 2 | | | 13.5 | 1.24 (S) | 3.07 | 0.988 |
| 4-MoCB | 3 | | K | 14.1 | 0.842 (Q) | 3.63 | 1.001 |
| 2,2'-DiCB | 4 | | | 35.4 | 4.62 (S) | 1.52 | 1.000 |
| 2,3-DiCB | 5 | | U | | 3.96 (S) | | |
| 2,3'-DiCB | 6 | | | 19.3 | 3.59 (S) | 1.39 | 1.175 |
| 2,4-DiCB | 7 | | J | 4.13 | 3.68 (S) | 1.73 | 1.157 |
| 2,4'-DiCB | 8 | | | 74.1 | 3.29 (S) | 1.47 | 1.207 |
| 2,5-DiCB | 9 | | K J | 5.79 | 3.46 (S) | 2.77 | 1.145 |
| 2,6-DiCB | 10 | | U | | 3.54 (S) | | |
| 3,3'-DiCB | 11 | | G | 282 | 3.99 (S) | 1.53 | 0.968 |
| 3,4-DiCB | 12 | 12 + 13 | C | 13.6 | 3.80 (S) | 1.78 | 0.983 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 3.68 (S) | | |
| 4,4'-DiCB | 15 | | | 50.1 | 3.02 (S) | 1.46 | 1.001 |
| 2,2',3-TriCB | 16 | | | 36.0 | 0.842 (Q) | 1.06 | 1.166 |
| 2,2',4-TriCB | 17 | | | 93.2 | 0.842 (Q) | 1.02 | 1.139 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 84.7 | 0.842 (Q) | 1.10 | 1.115 |
| 2,2',6-TriCB | 19 | | | 39.4 | 0.842 (Q) | 0.90 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 183 | 0.842 (Q) | 0.96 | 0.848 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 91.0 | 0.842 (Q) | 0.96 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 62.0 | 0.842 (Q) | 0.93 | 0.872 |
| 2,3,5-TriCB | 23 | | U | | 0.842 (Q) | | |
| 2,3,6-TriCB | 24 | | J | 1.05 | 0.842 (Q) | 1.12 | 1.159 |
| 2,3',4-TriCB | 25 | | | 30.4 | 0.842 (Q) | 0.98 | 0.825 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 29.0 | 0.842 (Q) | 1.00 | 1.302 |
| 2,3',6-TriCB | 27 | | | 9.72 | 0.842 (Q) | 1.10 | 1.152 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 131 | 0.842 (Q) | 0.97 | 0.837 |
| 2,4',6-TriCB | 32 | | G | 36.9 | 0.842 (Q) | 0.92 | 1.198 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | K J | 1.09 | 0.842 (Q) | 0.87 | 1.275 |
| 3,3',4-TriCB | 35 | | | 7.79 | 0.842 (Q) | 1.01 | 0.985 |
| 3,3',5-TriCB | 36 | | J | 2.89 | 0.842 (Q) | 1.19 | 0.932 |
| 3,4,4'-TriCB | 37 | | | 53.7 | 0.842 (Q) | 0.93 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.842 (Q) | | |
| 3,4',5-TriCB | 39 | | U | | 0.842 (Q) | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 98.1 | 0.842 (Q) | 0.74 | 1.337 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 51.4 | 0.842 (Q) | 0.78 | 1.312 |
| 2,2',3,5'-TeCB | 43 | | K J | 6.25 | 0.842 (Q) | 0.89 | 1.247 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 433 | 0.842 (Q) | 0.80 | 1.287 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 259 | 0.842 (Q) | 0.78 | 1.149 |
| 2,2',3,6'-TeCB | 46 | | | 12.0 | 0.842 (Q) | 0.82 | 1.161 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 31.7 | 0.842 (Q) | 0.85 | 1.274 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 163 | 0.842 (Q) | 0.76 | 1.260 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 36.9 | 0.842 (Q) | 0.78 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 271 | 0.842 (Q) | 0.78 | 1.235 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | | 9.36 | 0.842 (Q) | 0.77 | 1.001 |
| 2,3,3',4'-TeCB | 55 | | K J | 2.88 | 2.69 (S) | 0.60 | 0.889 |
| 2,3,3',4'-TeCB | 56 | | | 95.8 | 2.54 (S) | 0.79 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 2.39 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 2.48 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C K | 16.0 | 0.842 (Q) | 0.64 | 1.302 |
| 2,3,4,4'-TeCB | 60 | | | 34.7 | 2.43 (S) | 0.77 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 375 | 2.35 (S) | 0.76 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | | 8.61 | 2.37 (S) | 0.79 | 0.865 |
| 2,3,4',6'-TeCB | 64 | | | 78.1 | 0.842 (Q) | 0.74 | 1.349 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | G | 255 | 2.40 (S) | 0.75 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | J | 4.57 | 2.02 (S) | 0.83 | 0.857 |
| 2,3',4,5'-TeCB | 68 | | | 77.9 | 2.26 (S) | 0.74 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | K | 8.01 | 2.35 (S) | 0.94 | 0.823 |
| 2,3',5',6'-TeCB | 73 | | K J | 2.43 | 0.842 (Q) | 0.58 | 1.242 |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 28.5 | 1.79 (S) | 0.73 | 1.000 |
| 3,3',4,5'-TeCB | 78 | | U | | 2.28 (S) | | |
| 3,3',4,5'-TeCB | 79 | | K J | 6.25 | 1.87 (S) | 0.97 | 0.970 |
| 3,3',5,5'-TeCB | 80 | | U | | 2.15 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 1.80 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 53.6 | 0.957 (S) | 1.57 | 0.933 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 349 | 0.962 (S) | 1.62 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 112 | 1.10 (S) | 1.50 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 83.0 | 0.842 (Q) | 1.68 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 331 | 0.842 (Q) | 1.53 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 84.3 | 0.956 (S) | 1.59 | 1.155 |
| 2,2',3,4,6'-PeCB | 89 | | K J | 3.31 | 1.00 (S) | 1.05 | 1.182 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 522 | 0.842 (Q) | 1.56 | 0.869 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 111 | 0.929 (S) | 1.40 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 389 | 0.932 (S) | 1.56 | 1.121 |
| 2,2',3,5,6'-PeCB | 94 | | K J | 5.49 | 1.05 (S) | 1.13 | 1.102 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | K J | 5.33 | 0.842 (Q) | 1.23 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | | 11.1 | 0.862 (S) | 1.41 | 1.094 |
| 2,2',4,6,6'-PeCB | 104 | | K J | 1.51 | 0.842 (Q) | 1.21 | 1.001 |
| 2,3,3',4,4'-PeCB | 105 | | | 145 | 1.78 (S) | 1.37 | 1.000 |
| 2,3,3',4,5-PeCB | 106 | | U | | 2.14 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C | 17.2 | 2.18 (S) | 1.51 | 0.990 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 41.7 | 1.99 (S) | 1.44 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 592 | 0.842 (Q) | 1.56 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.842 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.842 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | J | 6.30 | 1.78 (S) | 1.43 | 1.000 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 404 | 1.95 (S) | 1.45 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 4.51 | 0.842 (Q) | 1.90 | 0.959 |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.842 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | K J | 5.08 | 2.57 (S) | 1.92 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K | 7.55 | 1.84 (S) | 1.95 | 1.001 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 2.50 | 1.94 (S) | 2.34 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 1.87 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 93.3 | 1.30 (S) | 1.33 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 753 | 1.46 (S) | 1.25 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | K | 41.2 | 1.73 (S) | 1.43 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | K | 9.27 | 1.92 (S) | 0.98 | 1.158 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 266 | 2.08 (S) | 1.18 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | K | 14.6 | 1.89 (S) | 1.70 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C K | 39.1 | 1.87 (S) | 1.03 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 311 | 0.842 (Q) | 1.25 | 1.103 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 96.2 | 0.842 (Q) | 1.29 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | | 24.8 | 1.64 (S) | 1.20 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C | 16.6 | 1.69 (S) | 1.35 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 115 | 1.63 (S) | 1.24 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 2.00 (S) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 31.4 | 0.842 (Q) | 1.41 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.842 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 148 | 1.52 (S) | 1.26 | 0.884 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 677 | 1.70 (S) | 1.24 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | J | 3.29 | 0.842 (Q) | 1.33 | 1.083 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | J | 4.13 | 0.842 (Q) | 1.25 | 1.012 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | K J | 1.41 | 0.842 (Q) | 3.65 | 1.006 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 624 | 1.30 (S) | 1.28 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | J | 1.14 | 0.842 (Q) | 1.10 | 1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 82.7 | 1.38 (S) | 1.33 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 62.8 | 1.09 (S) | 1.11 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | U | | 1.08 (S) | | |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 1.16 (S) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | K J | 1.49 | 1.14 (S) | 0.99 | 0.989 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 48.7 | 1.14 (S) | 1.26 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 1.45 (S) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 28.9 | 1.14 (S) | 1.33 | 1.000 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 1.06 (S) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 195 | 0.842 (Q) | 1.05 | 1.000 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 51.3 | 0.842 (Q) | 1.10 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 35.6 | 0.842 (Q) | 1.00 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 153 | 0.842 (Q) | 1.00 | 1.133 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K | 7.06 | 0.842 (Q) | 1.25 | 1.102 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 24.9 | 0.842 (Q) | 0.97 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 124 | 0.842 (Q) | 1.01 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 46.5 | 0.842 (Q) | 0.94 | 1.085 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 85.6 | 0.842 (Q) | 0.92 | 1.009 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 517 | 0.842 (Q) | 1.04 | 1.001 |
| 2,2',3,4,4',5,6-HpCB | 181 | | J | 2.82 | 0.842 (Q) | 1.03 | 1.155 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | K J | 1.92 | 0.842 (Q) | 1.49 | 1.115 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 112 | 0.842 (Q) | 1.04 | 1.127 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | J | 1.97 | 0.842 (Q) | 1.02 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.842 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 221 | 0.842 (Q) | 1.02 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | K J | 1.03 | 0.842 (Q) | 0.33 | 1.000 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | K | 7.71 | 0.842 (Q) | 1.49 | 1.001 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 41.6 | 0.842 (Q) | 1.08 | 0.948 |
| 2,3,3',4,4',5',6-HpCB | 191 | | | 9.08 | 0.842 (Q) | 1.14 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.842 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OcCB | 194 | | | 125 | 0.842 (Q) | 0.87 | 0.991 |
| 2,2',3,3',4,4',5,6-OcCB | 195 | | | 53.7 | 0.842 (Q) | 0.89 | 0.945 |
| 2,2',3,3',4,4',5,6'-OcCB | 196 | | | 72.1 | 0.842 (Q) | 0.91 | 0.916 |
| 2,2',3,3',4,4',6,6'-OcCB | 197 | 197 + 200 | C | 22.1 | 0.842 (Q) | 0.92 | 1.047 |
| 2,2',3,3',4,5,5',6-OcCB | 198 | 198 + 199 | C | 147 | 0.842 (Q) | 0.88 | 1.115 |
| 2,2',3,3',4,5,5',6'-OcCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OcCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OcCB | 201 | | | 19.4 | 0.842 (Q) | 0.87 | 1.023 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 39.5 | 0.842 (Q) | 0.86 | 1.001 |
| 2,2',3,4,4',5,5',6-OcCB | 203 | | | 94.6 | 0.842 (Q) | 0.94 | 0.920 |
| 2,2',3,4,4',5,6,6'-OcCB | 204 | | U | | 0.842 (Q) | | |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | J | 5.72 | 0.842 (Q) | 0.96 | 1.000 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 85.3 | 2.76 (S) | 0.77 | 1.001 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | | 11.6 | 2.11 (S) | 0.76 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 31.0 | 2.38 (S) | 0.81 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 75.3 | 0.842 (Q) | 1.18 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T03-1811
Sample Collection:
27-Nov-2018 16:22

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 17:11:47
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-3
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_027 S: 8
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_027 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | G | 4000 | 887 | 22.2 | 3.23 | 0.718 |
| 13C12-4-MoCB | 3L | | | 4000 | 1220 | 30.4 | 3.10 | 0.857 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1210 | 30.2 | 1.57 | 0.874 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1560 | 39.0 | 1.56 | 1.253 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2100 | 52.5 | 1.02 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1440 | 35.9 | 1.01 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1780 | 44.6 | 0.80 | 0.810 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1950 | 48.8 | 0.72 | 1.395 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1940 | 48.4 | 0.72 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 1970 | 49.3 | 1.61 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1960 | 49.1 | 1.51 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1790 | 44.7 | 1.57 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1840 | 45.9 | 1.49 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1980 | 49.5 | 1.53 | 1.151 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1860 | 46.5 | 1.53 | 1.299 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2370 | 59.3 | 1.26 | 0.787 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 5430 | 67.9 | 1.25 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2700 | 67.4 | 1.23 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2870 | 71.7 | 1.24 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 2960 | 74.1 | 1.02 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 2810 | 70.3 | 1.08 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2740 | 68.5 | 1.04 | 0.713 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1980 | 49.5 | 0.95 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2390 | 59.8 | 0.88 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2960 | 74.1 | 0.83 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3410 | 85.3 | 0.77 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3290 | 82.3 | 0.76 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3430 | 85.6 | 1.18 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1480 | 36.9 | 1.02 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2580 | 64.4 | 1.64 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3110 | 77.7 | 1.04 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T07-1811
Sample Collection:
28-Nov-2018 13:18

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-4 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 30-Jan-2019 Time: 22:48:48 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_028 S: 4 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_028 S: 1 |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | G | 13.5 | 1.06 (S) | 2.73 | 1.001 |
| 3-MoCB | 2 | | | 9.75 | 1.30 (S) | 3.36 | 0.989 |
| 4-MoCB | 3 | | K | 9.59 | 0.855 (S) | 3.62 | 1.001 |
| 2,2'-DiCB | 4 | | K | 26.6 | 4.68 (S) | 1.09 | 1.001 |
| 2,3-DiCB | 5 | | U | | 4.26 (S) | | |
| 2,3'-DiCB | 6 | | | 13.4 | 3.88 (S) | 1.57 | 1.175 |
| 2,4-DiCB | 7 | | U | | 3.87 (S) | | |
| 2,4'-DiCB | 8 | | | 51.1 | 3.45 (S) | 1.53 | 1.206 |
| 2,5-DiCB | 9 | | J | 5.29 | 3.68 (S) | 1.54 | 1.146 |
| 2,6-DiCB | 10 | | U | | 3.78 (S) | | |
| 3,3'-DiCB | 11 | | | 273 | 4.34 (S) | 1.42 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C U | | 4.10 (S) | | |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 3.97 (S) | | |
| 4,4'-DiCB | 15 | | | 26.3 | 3.18 (S) | 1.40 | 1.001 |
| 2,2',3-TriCB | 16 | | | 20.3 | 0.831 (Q) | 0.89 | 1.165 |
| 2,2',4-TriCB | 17 | | | 41.2 | 0.831 (Q) | 0.98 | 1.138 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 48.8 | 0.831 (Q) | 1.02 | 1.113 |
| 2,2',6-TriCB | 19 | | K | 7.89 | 0.831 (Q) | 1.36 | 1.000 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 83.8 | 0.831 (Q) | 0.98 | 0.849 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 48.2 | 0.831 (Q) | 0.92 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 30.7 | 0.831 (Q) | 1.00 | 0.873 |
| 2,3,5-TriCB | 23 | | U | | 0.831 (Q) | | |
| 2,3,6-TriCB | 24 | | K J | 1.05 | 0.831 (Q) | 1.30 | 1.157 |
| 2,3',4-TriCB | 25 | | | 13.8 | 0.831 (Q) | 1.00 | 0.826 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 17.1 | 0.831 (Q) | 0.95 | 1.302 |
| 2,3',6-TriCB | 27 | | K J | 4.22 | 0.831 (Q) | 1.23 | 1.151 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 66.4 | 0.831 (Q) | 0.95 | 0.837 |
| 2,4',6-TriCB | 32 | | | 19.5 | 0.831 (Q) | 0.92 | 1.197 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | K J | 0.993 | 0.831 (Q) | 1.51 | 1.274 |
| 3,3',4-TriCB | 35 | | J | 5.58 | 0.831 (Q) | 1.15 | 0.985 |
| 3,3',5-TriCB | 36 | | J | 3.05 | 0.831 (Q) | 0.98 | 0.932 |
| 3,4,4'-TriCB | 37 | | K | 20.9 | 0.831 (Q) | 0.88 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.831 (Q) | | |
| 3,4',5-TriCB | 39 | | U | | 0.831 (Q) | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 43.9 | 0.831 (Q) | 0.78 | 1.338 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 20.4 | 0.831 (Q) | 0.67 | 1.313 |
| 2,2',3,5'-TeCB | 43 | | K J | 3.45 | 0.915 (S) | 1.05 | 1.248 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 749 | 0.831 (Q) | 0.78 | 1.289 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 244 | 0.831 (Q) | 0.79 | 1.150 |
| 2,2',3,6'-TeCB | 46 | | U | | 0.831 (Q) | | |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | K | 13.6 | 0.831 (Q) | 1.07 | 1.275 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 60.7 | 0.831 (Q) | 0.79 | 1.260 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 12.7 | 0.831 (Q) | 0.71 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 129 | 0.831 (Q) | 0.78 | 1.235 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | U | | 0.831 (Q) | | |
| 2,3,3',4'-TeCB | 55 | | U | | 3.79 (S) | | |
| 2,3,3',4'-TeCB | 56 | | | 44.2 | 3.64 (S) | 0.76 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 3.36 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 3.63 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C K | 8.69 | 0.831 (Q) | 0.91 | 1.303 |
| 2,3,4,4'-TeCB | 60 | | K | 18.5 | 3.49 (S) | 0.60 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 162 | 3.33 (S) | 0.81 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | U | | 3.41 (S) | | |
| 2,3,4',6'-TeCB | 64 | | | 36.9 | 0.831 (Q) | 0.75 | 1.349 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | G | 98.4 | 3.46 (S) | 0.71 | 0.884 |
| 2,3',4,5'-TeCB | 67 | | U | | 2.83 (S) | | |
| 2,3',4,5'-TeCB | 68 | | | 148 | 3.21 (S) | 0.74 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | U | | 3.28 (S) | | |
| 2,3',5',6'-TeCB | 73 | | U | | 0.831 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 14.4 | 2.69 (S) | 0.79 | 1.000 |
| 3,3',4,5'-TeCB | 78 | | U | | 3.46 (S) | | |
| 3,3',4,5'-TeCB | 79 | | J | 3.32 | 2.65 (S) | 0.88 | 0.970 |
| 3,3',5,5'-TeCB | 80 | | U | | 3.12 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 2.53 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | K | 23.6 | 0.831 (Q) | 1.91 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 129 | 0.831 (Q) | 1.47 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 43.7 | 0.831 (Q) | 1.73 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 41.8 | 0.831 (Q) | 1.51 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 141 | 0.831 (Q) | 1.53 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 25.9 | 0.831 (Q) | 1.52 | 1.154 |
| 2,2',3,4,6'-PeCB | 89 | | U | | 0.831 (Q) | | |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 203 | 0.831 (Q) | 1.60 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 42.3 | 0.831 (Q) | 1.50 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 141 | 0.831 (Q) | 1.70 | 1.120 |
| 2,2',3,5,6'-PeCB | 94 | | U | | 0.831 (Q) | | |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | K J | 0.837 | 0.831 (Q) | 1.23 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | K J | 1.75 | 0.831 (Q) | 1.05 | 1.093 |
| 2,2',4,6,6'-PeCB | 104 | | U | | 0.831 (Q) | | |
| 2,3,3',4,4'-PeCB | 105 | | | 85.1 | 0.917 (S) | 1.41 | 1.000 |
| 2,3,3',4,5-PeCB | 106 | | U | | 1.12 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C K | 11.2 | 1.20 (S) | 1.15 | 0.990 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 18.5 | 1.10 (S) | 1.54 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 246 | 0.831 (Q) | 1.48 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.831 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.831 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | K J | 5.37 | 0.923 (S) | 1.84 | 1.000 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 189 | 1.01 (S) | 1.54 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 1.73 | 0.831 (Q) | 0.88 | 0.959 |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.831 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | K J | 2.86 | 1.31 (S) | 1.14 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K J | 3.69 | 0.976 (S) | 1.08 | 1.001 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 2.51 | 1.01 (S) | 2.18 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 1.23 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 58.2 | 1.14 (S) | 1.16 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 406 | 1.16 (S) | 1.21 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 22.8 | 1.52 (S) | 1.22 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | J | 2.78 | 1.43 (S) | 1.17 | 1.159 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 111 | 1.50 (S) | 1.23 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | K J | 4.82 | 1.37 (S) | 1.58 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 18.5 | 1.42 (S) | 1.19 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 107 | 0.831 (Q) | 1.18 | 1.103 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 30.1 | 0.831 (Q) | 1.09 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | K | 14.5 | 1.49 (S) | 0.89 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C K J | 3.32 | 1.30 (S) | 2.16 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 72.1 | 1.34 (S) | 1.18 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 1.48 (S) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | K | 12.2 | 0.831 (Q) | 0.89 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.831 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 58.5 | 1.10 (S) | 1.16 | 0.885 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 273 | 1.27 (S) | 1.17 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | U | | 0.831 (Q) | | |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | U | | 0.831 (Q) | | |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | U | | 0.831 (Q) | | |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 326 | 1.04 (S) | 1.30 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | J | 1.37 | 0.831 (Q) | 1.32 | 1.001 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 42.9 | 1.01 (S) | 1.23 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 37.8 | 0.906 (S) | 1.39 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | K | 9.09 | 0.952 (S) | 1.58 | 0.981 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 0.974 (S) | | |
| 2,3,3',4',5,5',6-HxCB | 162 | | U | | 0.932 (S) | | |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | K | 27.1 | 0.971 (S) | 1.51 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 1.14 (S) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5',6-HxCB | 167 | | | 15.0 | 0.846 (S) | 1.21 | 1.001 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5',6-HxCB | 169 | | U | | 0.831 (Q) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 123 | 0.831 (Q) | 1.06 | 1.000 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 35.4 | 0.831 (Q) | 0.98 | 1.162 |
| 2,2',3,3',4,5,5',6-HpCB | 172 | | | 26.0 | 0.831 (Q) | 1.13 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6',6-HpCB | 174 | | | 135 | 0.831 (Q) | 1.07 | 1.132 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K J | 4.35 | 0.831 (Q) | 1.32 | 1.102 |
| 2,2',3,3',4,6,6',6-HpCB | 176 | | K | 14.6 | 0.831 (Q) | 1.25 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 71.4 | 0.831 (Q) | 0.99 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 30.5 | 0.831 (Q) | 1.01 | 1.085 |
| 2,2',3,3',5,6,6',6-HpCB | 179 | | | 50.5 | 0.831 (Q) | 0.92 | 1.009 |
| 2,2',3,4,4',5,5',6-HpCB | 180 | 180 + 193 | C | 359 | 0.831 (Q) | 1.01 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | K J | 1.05 | 0.831 (Q) | 0.75 | 1.155 |
| 2,2',3,4,4',5,6',6-HpCB | 182 | | K J | 0.934 | 0.831 (Q) | 2.96 | 1.115 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 86.4 | 0.831 (Q) | 1.00 | 1.127 |
| 2,2',3,4,4',6,6',6-HpCB | 184 | | K J | 3.07 | 0.831 (Q) | 0.83 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6',6-HpCB | 186 | | U | | 0.831 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 193 | 0.831 (Q) | 1.12 | 1.110 |
| 2,2',3,4',5,6,6',6-HpCB | 188 | | U | | 0.831 (Q) | | |
| 2,3,3',4,4',5,5',6-HpCB | 189 | | K J | 3.88 | 0.831 (Q) | 1.54 | 1.001 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 32.1 | 0.831 (Q) | 1.04 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | J | 6.64 | 0.831 (Q) | 1.03 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.831 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5',6-OcCB | 194 | | | 92.0 | 0.831 (Q) | 0.91 | 0.991 |
| 2,2',3,3',4,4',5,6-OcCB | 195 | | | 34.5 | 0.831 (Q) | 0.98 | 0.946 |
| 2,2',3,3',4,4',5,6',6-OcCB | 196 | | | 48.5 | 0.831 (Q) | 0.79 | 0.916 |
| 2,2',3,3',4,4',6,6',6-OcCB | 197 | 197 + 200 | C | 14.8 | 0.831 (Q) | 0.85 | 1.046 |
| 2,2',3,3',4,5,5',6-OcCB | 198 | 198 + 199 | C | 127 | 0.831 (Q) | 0.89 | 1.115 |
| 2,2',3,3',4,5,5',6',6-OcCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6',6-OcCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6',6-OcCB | 201 | | | 10.9 | 0.831 (Q) | 0.96 | 1.023 |
| 2,2',3,3',5,5',6,6',6-OcCB | 202 | | | 23.2 | 0.831 (Q) | 0.82 | 1.000 |
| 2,2',3,4,4',5,5',6-OcCB | 203 | | | 80.9 | 0.831 (Q) | 0.90 | 0.920 |
| 2,2',3,4,4',5,6,6',6-OcCB | 204 | | U | | 0.831 (Q) | | |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | J | 5.53 | 0.831 (Q) | 0.88 | 1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 65.8 | 2.75 (S) | 0.76 | 1.000 |
| 2,2',3,3',4,4',5,6,6',6-NoCB | 207 | | K | 7.67 | 2.11 (S) | 1.03 | 1.020 |
| 2,2',3,3',4,5,5',6,6',6-NoCB | 208 | | | 18.2 | 2.48 (S) | 0.88 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6',6-DeCB | 209 | | | 43.5 | 0.831 (Q) | 1.07 | 1.001 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T07-1811
Sample Collection:
28-Nov-2018 13:18

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 22:48:48
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-4
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_028 S: 4
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_028 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | G | 4000 | 1010 | 25.2 | 3.27 | 0.718 |
| 13C12-4-MoCB | 3L | | | 4000 | 1330 | 33.2 | 3.01 | 0.857 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1290 | 32.3 | 1.62 | 0.874 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1660 | 41.5 | 1.52 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2350 | 58.7 | 1.05 | 1.073 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1500 | 37.4 | 1.01 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1960 | 48.9 | 0.79 | 0.811 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1890 | 47.3 | 0.69 | 1.396 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1910 | 47.7 | 0.69 | 1.373 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2090 | 52.3 | 1.56 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1890 | 47.3 | 1.56 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1700 | 42.6 | 1.60 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1760 | 44.0 | 1.54 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1840 | 46.1 | 1.57 | 1.151 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1790 | 44.8 | 1.53 | 1.300 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2770 | 69.2 | 1.26 | 0.786 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 5440 | 68.1 | 1.23 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2740 | 68.6 | 1.25 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2830 | 70.8 | 1.19 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 3310 | 82.8 | 1.07 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 3340 | 83.6 | 1.09 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2900 | 72.6 | 1.07 | 0.713 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1870 | 46.7 | 0.96 | 0.959 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2610 | 65.2 | 0.89 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2960 | 74.1 | 0.84 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3560 | 88.9 | 0.74 | 1.044 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3200 | 79.9 | 0.78 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3770 | 94.4 | 1.19 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1530 | 38.1 | 1.01 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2600 | 65.0 | 1.63 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3180 | 79.5 | 1.02 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T02-1811
Sample Collection:
30-Nov-2018 15:06

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-5 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 30-Jan-2019 Time: 23:53:04 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_028 S: 5 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_028 S: 1 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | G | 19.3 | 1.25 (S) | 2.75 | 1.001 |
| 3-MoCB | 2 | | | 19.3 | 1.52 (S) | 2.85 | 0.988 |
| 4-MoCB | 3 | | | 15.1 | 0.990 (S) | 3.47 | 1.000 |
| 2,2'-DiCB | 4 | | | 35.9 | 4.80 (S) | 1.68 | 1.000 |
| 2,3-DiCB | 5 | | U | | 4.42 (S) | | |
| 2,3'-DiCB | 6 | | | 23.2 | 4.02 (S) | 1.62 | 1.176 |
| 2,4-DiCB | 7 | | K J | 4.81 | 4.01 (S) | 2.48 | 1.159 |
| 2,4'-DiCB | 8 | | | 87.5 | 3.58 (S) | 1.46 | 1.208 |
| 2,5-DiCB | 9 | | J | 6.03 | 3.81 (S) | 1.75 | 1.147 |
| 2,6-DiCB | 10 | | U | | 3.91 (S) | | |
| 3,3'-DiCB | 11 | | | 353 | 4.49 (S) | 1.45 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C K | 15.1 | 4.25 (S) | 2.74 | 0.984 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 4.11 (S) | | |
| 4,4'-DiCB | 15 | | | 63.4 | 3.30 (S) | 1.48 | 1.001 |
| 2,2',3-TriCB | 16 | | | 36.1 | 0.884 (Q) | 1.10 | 1.166 |
| 2,2',4-TriCB | 17 | | | 61.4 | 0.884 (Q) | 1.01 | 1.139 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 82.3 | 0.884 (Q) | 0.96 | 1.114 |
| 2,2',6-TriCB | 19 | | | 33.5 | 0.884 (Q) | 1.11 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 213 | 0.884 (Q) | 1.01 | 0.848 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 90.5 | 0.884 (Q) | 0.93 | 0.857 |
| 2,3,4'-TriCB | 22 | | K | 68.0 | 0.888 (S) | 0.88 | 0.872 |
| 2,3,5-TriCB | 23 | | U | | 0.884 (Q) | | |
| 2,3,6-TriCB | 24 | | J | 1.35 | 0.884 (Q) | 1.03 | 1.161 |
| 2,3',4-TriCB | 25 | | | 41.0 | 0.884 (Q) | 1.00 | 0.826 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 37.2 | 0.884 (Q) | 1.19 | 1.304 |
| 2,3',6-TriCB | 27 | | | 10.4 | 0.884 (Q) | 1.04 | 1.152 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 160 | 0.884 (Q) | 0.97 | 0.837 |
| 2,4',6-TriCB | 32 | | | 44.6 | 0.884 (Q) | 1.03 | 1.198 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | K J | 2.02 | 0.884 (Q) | 1.48 | 1.275 |
| 3,3',4-TriCB | 35 | | | 8.77 | 0.884 (Q) | 1.01 | 0.985 |
| 3,3',5-TriCB | 36 | | K J | 3.54 | 0.884 (Q) | 1.47 | 0.932 |
| 3,4,4'-TriCB | 37 | | | 56.4 | 0.884 (Q) | 0.92 | 1.001 |
| 3,4,5-TriCB | 38 | | K J | 1.00 | 0.884 (Q) | 0.71 | 0.968 |
| 3,4',5-TriCB | 39 | | J | 1.21 | 0.884 (Q) | 1.13 | 0.946 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 111 | 0.884 (Q) | 0.77 | 1.338 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 58.7 | 0.884 (Q) | 0.77 | 1.314 |
| 2,2',3,5'-TeCB | 43 | | J | 6.65 | 0.884 (Q) | 0.84 | 1.249 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 384 | 0.884 (Q) | 0.78 | 1.288 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 129 | 0.884 (Q) | 0.80 | 1.149 |
| 2,2',3,6'-TeCB | 46 | | | 12.4 | 0.884 (Q) | 0.77 | 1.162 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 39.2 | 0.884 (Q) | 0.76 | 1.276 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 209 | 0.884 (Q) | 0.73 | 1.261 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 48.5 | 0.884 (Q) | 0.83 | 1.112 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 346 | 0.884 (Q) | 0.77 | 1.235 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | | 7.30 | 0.884 (Q) | 0.82 | 1.001 |
| 2,3,3',4'-TeCB | 55 | | U | | 4.18 (S) | | |
| 2,3,3',4'-TeCB | 56 | | | 130 | 4.02 (S) | 0.75 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 3.71 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 4.00 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 20.5 | 0.884 (Q) | 0.83 | 1.304 |
| 2,3,4,4'-TeCB | 60 | | | 40.9 | 3.85 (S) | 0.72 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 522 | 3.67 (S) | 0.74 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | | 12.0 | 3.76 (S) | 0.77 | 0.865 |
| 2,3,4',6'-TeCB | 64 | | | 96.2 | 0.884 (Q) | 0.78 | 1.350 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | G | 348 | 3.82 (S) | 0.76 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | K J | 6.52 | 3.13 (S) | 0.24 | 0.857 |
| 2,3',4,5'-TeCB | 68 | | | 50.7 | 3.54 (S) | 0.77 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | K | 7.85 | 3.62 (S) | 0.57 | 0.824 |
| 2,3',5',6'-TeCB | 73 | | U | | 0.884 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 32.9 | 2.93 (S) | 0.71 | 1.000 |
| 3,3',4,5'-TeCB | 78 | | U | | 3.82 (S) | | |
| 3,3',4,5'-TeCB | 79 | | K | 11.2 | 2.93 (S) | 1.05 | 0.971 |
| 3,3',5,5'-TeCB | 80 | | U | | 3.45 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 2.74 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 74.4 | 0.884 (Q) | 1.61 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 454 | 0.884 (Q) | 1.51 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 150 | 0.884 (Q) | 1.48 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 111 | 0.884 (Q) | 1.64 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 439 | 0.884 (Q) | 1.57 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 113 | 0.884 (Q) | 1.75 | 1.153 |
| 2,2',3,4,6'-PeCB | 89 | | K J | 5.62 | 0.884 (Q) | 2.00 | 1.181 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 685 | 0.884 (Q) | 1.58 | 0.869 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 150 | 0.884 (Q) | 1.58 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 505 | 0.884 (Q) | 1.57 | 1.120 |
| 2,2',3,5,6'-PeCB | 94 | | K J | 5.27 | 0.884 (Q) | 1.18 | 1.101 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | J | 4.91 | 0.884 (Q) | 1.73 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | | 16.2 | 0.884 (Q) | 1.78 | 1.093 |
| 2,2',4,6,6'-PeCB | 104 | | K J | 1.32 | 0.884 (Q) | 4.50 | 1.001 |
| 2,3,3',4,4'-PeCB | 105 | | | 221 | 1.52 (S) | 1.46 | 1.001 |
| 2,3,3',4,5-PeCB | 106 | | U | | 1.84 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C K | 27.8 | 1.98 (S) | 1.21 | 0.990 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 58.3 | 1.81 (S) | 1.38 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 801 | 0.884 (Q) | 1.58 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | K J | 1.62 | 0.884 (Q) | 2.65 | 0.946 |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.884 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | K | 10.9 | 1.52 (S) | 2.47 | 1.001 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 576 | 1.62 (S) | 1.46 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 6.17 | 0.884 (Q) | 1.91 | 0.958 |
| 2,3',4,5',6-PeCB | 121 | | J | 1.86 | 0.884 (Q) | 1.55 | 1.200 |
| 2',3,3',4,5-PeCB | 122 | | K | 9.42 | 2.16 (S) | 2.68 | 1.009 |
| 2',3,4,4',5-PeCB | 123 | | K | 8.51 | 1.61 (S) | 1.77 | 1.001 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 5.30 | 1.65 (S) | 3.13 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 2.03 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 163 | 1.65 (S) | 1.36 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 1130 | 1.68 (S) | 1.23 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 71.0 | 2.20 (S) | 1.14 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | | 9.85 | 2.07 (S) | 1.11 | 1.158 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 373 | 2.17 (S) | 1.25 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 20.5 | 1.98 (S) | 1.08 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C K | 51.8 | 2.05 (S) | 0.99 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 388 | 0.884 (Q) | 1.25 | 1.104 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 122 | 0.884 (Q) | 1.19 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | | 48.4 | 2.15 (S) | 1.15 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C K | 21.8 | 1.87 (S) | 0.98 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 190 | 1.93 (S) | 1.13 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 2.14 (S) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 38.5 | 0.884 (Q) | 1.27 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.884 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 205 | 1.58 (S) | 1.27 | 0.884 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 824 | 1.82 (S) | 1.20 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | K J | 5.13 | 0.884 (Q) | 0.76 | 1.083 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | K J | 3.00 | 0.884 (Q) | 0.65 | 1.012 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | K J | 1.58 | 0.884 (Q) | 0.83 | 1.006 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 970 | 1.50 (S) | 1.25 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | K J | 1.72 | 0.884 (Q) | 2.34 | 1.001 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 115 | 1.46 (S) | 1.19 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 103 | 1.31 (S) | 1.11 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | K | 17.6 | 1.37 (S) | 0.83 | 0.981 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 1.40 (S) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | K J | 2.00 | 1.34 (S) | 1.61 | 0.989 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 72.9 | 1.40 (S) | 1.34 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 1.65 (S) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 37.0 | 1.23 (S) | 1.36 | 1.001 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 1.31 (S) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 263 | 0.884 (Q) | 1.00 | 1.001 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 87.3 | 0.884 (Q) | 0.96 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 52.4 | 0.884 (Q) | 1.04 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 304 | 0.884 (Q) | 0.98 | 1.132 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K | 11.2 | 0.884 (Q) | 0.84 | 1.102 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 39.8 | 0.884 (Q) | 0.99 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 165 | 0.884 (Q) | 1.07 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 79.5 | 0.884 (Q) | 1.05 | 1.084 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 131 | 0.884 (Q) | 1.05 | 1.009 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 729 | 0.884 (Q) | 1.07 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | K J | 3.21 | 0.884 (Q) | 1.37 | 1.156 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | J | 4.24 | 0.884 (Q) | 1.06 | 1.115 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 204 | 0.884 (Q) | 1.08 | 1.126 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | K J | 2.57 | 0.884 (Q) | 2.12 | 1.024 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.884 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 411 | 0.884 (Q) | 0.96 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | K J | 1.39 | 0.884 (Q) | 2.28 | 1.000 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 10.7 | 0.884 (Q) | 1.08 | 1.000 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 60.1 | 0.884 (Q) | 1.03 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | | 13.4 | 0.884 (Q) | 0.99 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.884 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | | 150 | 0.884 (Q) | 0.92 | 0.991 |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | | 66.8 | 0.884 (Q) | 0.94 | 0.946 |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | | 77.1 | 0.884 (Q) | 0.87 | 0.916 |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C K | 32.1 | 0.884 (Q) | 1.04 | 1.046 |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C | 211 | 0.884 (Q) | 0.91 | 1.115 |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | | 23.0 | 0.884 (Q) | 0.98 | 1.023 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | | 49.7 | 0.884 (Q) | 0.85 | 1.000 |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | | 126 | 0.884 (Q) | 0.96 | 0.920 |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.884 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | K | 8.72 | 0.884 (Q) | 1.15 | 1.000 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 159 | 2.89 (S) | 0.76 | 1.000 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | | 16.0 | 2.26 (S) | 0.69 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 53.6 | 2.69 (S) | 0.68 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 208 | 0.884 (Q) | 1.10 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T02-1811
Sample Collection:
30-Nov-2018 15:06

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 23:53:04
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-5
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_028 S: 5
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_028 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | G | 4000 | 892 | 22.3 | 3.23 | 0.717 |
| 13C12-4-MoCB | 3L | | | 4000 | 1230 | 30.7 | 3.12 | 0.857 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1250 | 31.2 | 1.53 | 0.873 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1530 | 38.3 | 1.53 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2350 | 58.8 | 1.07 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1460 | 36.6 | 1.01 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1920 | 47.9 | 0.79 | 0.810 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1950 | 48.9 | 0.71 | 1.396 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 2000 | 50.1 | 0.70 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2120 | 52.9 | 1.59 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1960 | 49.0 | 1.54 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1760 | 43.9 | 1.53 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1840 | 46.0 | 1.56 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1890 | 47.3 | 1.55 | 1.151 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1850 | 46.4 | 1.58 | 1.300 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2960 | 73.9 | 1.27 | 0.786 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 5440 | 68.0 | 1.24 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2720 | 68.1 | 1.23 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2800 | 70.1 | 1.26 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 3630 | 90.7 | 1.04 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 3550 | 88.7 | 1.07 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 3050 | 76.3 | 1.04 | 0.713 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1970 | 49.4 | 0.91 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2590 | 64.7 | 0.89 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 3180 | 79.5 | 0.86 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3800 | 95.1 | 0.75 | 1.044 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3430 | 85.6 | 0.75 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 4060 | 102 | 1.17 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1560 | 39.0 | 1.00 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2660 | 66.6 | 1.62 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3110 | 77.8 | 1.03 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T04-1812
Sample Collection:
01-Dec-2018 13:10

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-6 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 31-Jan-2019 Time: 00:57:17 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_028 S: 6 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_028 S: 1 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | | 11.8 | 0.867 (Q) | 2.88 | 1.001 |
| 3-MoCB | 2 | | | 9.77 | 0.957 (S) | 2.66 | 0.989 |
| 4-MoCB | 3 | | | 11.5 | 0.867 (Q) | 3.30 | 1.001 |
| 2,2'-DiCB | 4 | | | 30.8 | 3.35 (S) | 1.62 | 1.001 |
| 2,3-DiCB | 5 | | U | | 3.16 (S) | | |
| 2,3'-DiCB | 6 | | | 15.8 | 2.87 (S) | 1.49 | 1.175 |
| 2,4-DiCB | 7 | | K J | 3.99 | 2.86 (S) | 1.20 | 1.159 |
| 2,4'-DiCB | 8 | | | 56.7 | 2.56 (S) | 1.36 | 1.208 |
| 2,5-DiCB | 9 | | K J | 6.27 | 2.73 (S) | 1.28 | 1.146 |
| 2,6-DiCB | 10 | | U | | 2.80 (S) | | |
| 3,3'-DiCB | 11 | | | 213 | 3.21 (S) | 1.49 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C K | 9.45 | 3.04 (S) | 2.11 | 0.984 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 2.94 (S) | | |
| 4,4'-DiCB | 15 | | | 34.1 | 2.40 (S) | 1.44 | 1.000 |
| 2,2',3-TriCB | 16 | | K | 28.9 | 0.867 (Q) | 0.87 | 1.165 |
| 2,2',4-TriCB | 17 | | | 53.5 | 0.867 (Q) | 0.96 | 1.138 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 69.9 | 0.867 (Q) | 1.10 | 1.113 |
| 2,2',6-TriCB | 19 | | | 21.7 | 0.867 (Q) | 1.16 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 136 | 0.867 (Q) | 0.99 | 0.849 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 68.4 | 0.867 (Q) | 0.95 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 45.2 | 0.867 (Q) | 0.95 | 0.873 |
| 2,3,5-TriCB | 23 | | U | | 0.867 (Q) | | |
| 2,3,6-TriCB | 24 | | J | 1.09 | 0.867 (Q) | 0.98 | 1.158 |
| 2,3',4-TriCB | 25 | | | 14.7 | 0.867 (Q) | 0.99 | 0.826 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 24.2 | 0.867 (Q) | 0.96 | 1.302 |
| 2,3',6-TriCB | 27 | | J | 6.64 | 0.867 (Q) | 0.98 | 1.152 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 106 | 0.867 (Q) | 0.99 | 0.837 |
| 2,4',6-TriCB | 32 | | K | 35.3 | 0.867 (Q) | 0.84 | 1.198 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | U | | 0.867 (Q) | | |
| 3,3',4-TriCB | 35 | | J | 5.68 | 0.867 (Q) | 1.15 | 0.986 |
| 3,3',5-TriCB | 36 | | K J | 2.00 | 0.867 (Q) | 0.80 | 0.932 |
| 3,4,4'-TriCB | 37 | | | 37.9 | 0.867 (Q) | 0.99 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.867 (Q) | | |
| 3,4',5-TriCB | 39 | | U | | 0.867 (Q) | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 73.1 | 0.867 (Q) | 0.75 | 1.338 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 37.4 | 0.867 (Q) | 0.83 | 1.313 |
| 2,2',3,5'-TeCB | 43 | | K J | 6.68 | 0.867 (Q) | 0.92 | 1.247 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 685 | 0.867 (Q) | 0.74 | 1.289 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 176 | 0.867 (Q) | 0.78 | 1.150 |
| 2,2',3,6'-TeCB | 46 | | K | 8.06 | 0.867 (Q) | 0.64 | 1.161 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 24.2 | 0.867 (Q) | 0.73 | 1.275 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 118 | 0.867 (Q) | 0.73 | 1.261 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 29.0 | 0.867 (Q) | 0.77 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 216 | 0.867 (Q) | 0.81 | 1.235 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | K J | 3.50 | 0.867 (Q) | 0.91 | 1.002 |
| 2,3,3',4'-TeCB | 55 | | U | | 2.35 (S) | | |
| 2,3,3',4'-TeCB | 56 | | | 78.1 | 2.26 (S) | 0.79 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 2.09 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 2.25 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 13.4 | 0.867 (Q) | 0.84 | 1.302 |
| 2,3,4,4'-TeCB | 60 | | | 29.2 | 2.17 (S) | 0.76 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 297 | 2.07 (S) | 0.75 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | K J | 6.22 | 2.11 (S) | 1.07 | 0.864 |
| 2,3,4',6'-TeCB | 64 | | | 60.4 | 0.867 (Q) | 0.82 | 1.350 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | G | 203 | 2.15 (S) | 0.76 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | K J | 3.18 | 1.76 (S) | 0.38 | 0.857 |
| 2,3',4,5'-TeCB | 68 | | | 115 | 1.99 (S) | 0.74 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | U | | 2.04 (S) | | |
| 2,3',5',6'-TeCB | 73 | | U | | 0.867 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 22.4 | 1.71 (S) | 0.76 | 1.000 |
| 3,3',4,5'-TeCB | 78 | | U | | 2.15 (S) | | |
| 3,3',4,5'-TeCB | 79 | | J | 6.35 | 1.65 (S) | 0.70 | 0.970 |
| 3,3',5,5'-TeCB | 80 | | U | | 1.94 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 1.66 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 46.5 | 1.01 (S) | 1.34 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 235 | 0.964 (S) | 1.55 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 89.0 | 1.04 (S) | 1.59 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 64.5 | 0.867 (Q) | 1.50 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 253 | 0.867 (Q) | 1.62 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 55.5 | 0.912 (S) | 1.57 | 1.155 |
| 2,2',3,4,6'-PeCB | 89 | | K J | 4.70 | 0.980 (S) | 1.22 | 1.182 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 371 | 0.867 (Q) | 1.58 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 80.7 | 0.930 (S) | 1.64 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 293 | 0.890 (S) | 1.69 | 1.120 |
| 2,2',3,5,6'-PeCB | 94 | | K J | 2.53 | 0.994 (S) | 1.31 | 1.102 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | J | 3.02 | 0.867 (Q) | 1.76 | 1.015 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | K | 7.53 | 0.867 (Q) | 1.09 | 1.094 |
| 2,2',4,6,6'-PeCB | 104 | | U | | 0.867 (Q) | | |
| 2,3,3',4,4'-PeCB | 105 | | | 134 | 1.49 (S) | 1.52 | 1.001 |
| 2,3,3',4,5-PeCB | 106 | | U | | 1.79 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C | 16.3 | 1.92 (S) | 1.71 | 0.990 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | K | 30.3 | 1.76 (S) | 1.27 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 459 | 0.867 (Q) | 1.55 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.867 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.867 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | K J | 6.87 | 1.49 (S) | 1.25 | 1.000 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 331 | 1.60 (S) | 1.46 | 1.001 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | K J | 2.80 | 0.867 (Q) | 2.08 | 0.959 |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.867 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | J | 4.92 | 2.10 (S) | 1.73 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K J | 5.41 | 1.59 (S) | 1.79 | 1.000 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 2.28 | 1.65 (S) | 1.39 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 1.97 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 87.1 | 1.47 (S) | 1.26 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 622 | 1.50 (S) | 1.28 | 0.928 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 39.7 | 1.96 (S) | 1.20 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | K | 7.83 | 1.85 (S) | 0.79 | 1.159 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 216 | 1.94 (S) | 1.33 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 11.0 | 1.77 (S) | 1.41 | 1.190 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 30.0 | 1.83 (S) | 1.38 | 1.138 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 213 | 0.867 (Q) | 1.20 | 1.103 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 68.5 | 0.867 (Q) | 1.38 | 1.022 |
| 2,2',3,4,4',5-HxCB | 137 | | K | 26.2 | 1.92 (S) | 1.03 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C | 10.7 | 1.67 (S) | 1.12 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 106 | 1.72 (S) | 1.18 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 1.91 (S) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 24.4 | 0.867 (Q) | 1.28 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.867 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 111 | 1.41 (S) | 1.37 | 0.884 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 447 | 1.63 (S) | 1.27 | 1.132 |
| 2,2',3,4',5,6'-HxCB | 148 | | K J | 1.64 | 0.867 (Q) | 2.81 | 1.083 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | K J | 1.75 | 0.867 (Q) | 1.47 | 1.012 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | U | | 0.867 (Q) | | |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 509 | 1.34 (S) | 1.25 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | K J | 0.990 | 0.867 (Q) | 2.13 | 1.001 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 63.7 | 1.34 (S) | 1.14 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 55.2 | 1.17 (S) | 1.22 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | | 9.41 | 1.23 (S) | 1.12 | 0.981 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 1.25 (S) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | J | 1.39 | 1.20 (S) | 1.17 | 0.989 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 42.5 | 1.25 (S) | 1.41 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 1.47 (S) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 21.2 | 1.09 (S) | 1.24 | 1.000 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 1.04 (S) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 141 | 0.867 (Q) | 0.99 | 1.001 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 45.8 | 0.867 (Q) | 1.13 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 23.9 | 0.867 (Q) | 0.99 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 166 | 0.867 (Q) | 1.09 | 1.133 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K J | 6.90 | 0.867 (Q) | 0.73 | 1.102 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 19.2 | 0.867 (Q) | 1.08 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 89.5 | 0.867 (Q) | 1.00 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 35.9 | 0.867 (Q) | 1.07 | 1.085 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 65.6 | 0.867 (Q) | 1.15 | 1.010 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 354 | 0.867 (Q) | 1.07 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | K J | 1.90 | 0.867 (Q) | 2.90 | 1.156 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | U | | 0.867 (Q) | | |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 107 | 0.867 (Q) | 1.14 | 1.127 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | K J | 1.42 | 0.867 (Q) | 2.49 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.867 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 215 | 0.867 (Q) | 1.07 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | U | | 0.867 (Q) | | |
| 2,3,3',4,4',5,5'-HpCB | 189 | | K J | 4.61 | 0.867 (Q) | 1.21 | 1.001 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 36.4 | 0.867 (Q) | 0.90 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | K J | 5.52 | 0.867 (Q) | 1.21 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.867 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | | 73.3 | 0.867 (Q) | 0.94 | 0.991 |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | | 29.3 | 0.867 (Q) | 0.97 | 0.946 |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | K | 39.7 | 0.867 (Q) | 1.23 | 0.916 |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C | 20.8 | 0.867 (Q) | 0.82 | 1.047 |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C | 113 | 0.867 (Q) | 0.82 | 1.115 |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | K | 12.1 | 0.867 (Q) | 1.10 | 1.023 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | | 25.8 | 0.867 (Q) | 0.81 | 1.001 |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | | 62.0 | 0.867 (Q) | 0.98 | 0.920 |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.867 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | K J | 2.79 | 0.867 (Q) | 1.24 | 1.000 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 61.6 | 3.31 (S) | 0.77 | 1.000 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | K | 7.18 | 2.49 (S) | 0.90 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 21.6 | 2.88 (S) | 0.79 | 1.000 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 47.9 | 0.867 (Q) | 1.05 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T04-1812
Sample Collection:
01-Dec-2018 13:10

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 31-Jan-2019 Time: 00:57:17
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-6
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_028 S: 6
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_028 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | | 4000 | 1280 | 32.1 | 3.15 | 0.718 |
| 13C12-4-MoCB | 3L | | | 4000 | 1540 | 38.5 | 3.06 | 0.856 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1490 | 37.2 | 1.55 | 0.873 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1800 | 44.9 | 1.54 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2640 | 66.1 | 1.04 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1460 | 36.5 | 1.02 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 2110 | 52.8 | 0.79 | 0.810 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1780 | 44.6 | 0.68 | 1.395 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1840 | 46.1 | 0.71 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2090 | 52.2 | 1.60 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1830 | 45.7 | 1.54 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1610 | 40.2 | 1.56 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1680 | 42.0 | 1.52 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1750 | 43.8 | 1.56 | 1.150 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1660 | 41.5 | 1.52 | 1.299 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2740 | 68.6 | 1.28 | 0.787 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 4940 | 61.8 | 1.27 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2490 | 62.2 | 1.27 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2570 | 64.3 | 1.25 | 1.190 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 3170 | 79.2 | 1.05 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 3110 | 77.8 | 1.05 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2740 | 68.5 | 1.03 | 0.712 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1750 | 43.8 | 0.97 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2310 | 57.8 | 0.96 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2790 | 69.7 | 0.86 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3280 | 82.0 | 0.76 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3100 | 77.5 | 0.74 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3520 | 87.9 | 1.17 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1550 | 38.8 | 1.03 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2410 | 60.2 | 1.61 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 2920 | 72.9 | 1.05 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T06-1811
Sample Collection:
30-Nov-2018 16:26

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-7 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 31-Jan-2019 Time: 02:01:34 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_028 S: 7 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_028 S: 1 |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | G | 11.0 | 2.00 (S) | 3.22 | 1.003 |
| 3-MoCB | 2 | | | 17.1 | 1.56 (S) | 3.25 | 0.989 |
| 4-MoCB | 3 | | | 12.0 | 0.846 (Q) | 3.35 | 1.001 |
| 2,2'-DiCB | 4 | | | 20.3 | 2.99 (S) | 1.33 | 1.001 |
| 2,3-DiCB | 5 | | U | | 2.81 (S) | | |
| 2,3'-DiCB | 6 | | | 13.6 | 2.56 (S) | 1.48 | 1.175 |
| 2,4-DiCB | 7 | | J | 2.58 | 2.55 (S) | 1.63 | 1.159 |
| 2,4'-DiCB | 8 | | | 41.4 | 2.28 (S) | 1.55 | 1.206 |
| 2,5-DiCB | 9 | | K J | 5.30 | 2.43 (S) | 0.88 | 1.145 |
| 2,6-DiCB | 10 | | U | | 2.49 (S) | | |
| 3,3'-DiCB | 11 | | | 189 | 2.86 (S) | 1.43 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C K | 9.04 | 2.70 (S) | 0.88 | 0.985 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 2.62 (S) | | |
| 4,4'-DiCB | 15 | | | 22.4 | 2.14 (S) | 1.47 | 1.001 |
| 2,2',3-TriCB | 16 | | K | 15.9 | 0.846 (Q) | 1.22 | 1.165 |
| 2,2',4-TriCB | 17 | | | 28.6 | 0.846 (Q) | 1.16 | 1.138 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 47.0 | 0.846 (Q) | 1.10 | 1.113 |
| 2,2',6-TriCB | 19 | | K | 7.51 | 0.846 (Q) | 0.76 | 1.000 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 75.8 | 0.846 (Q) | 0.99 | 0.849 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 38.0 | 0.846 (Q) | 1.09 | 0.857 |
| 2,3,4'-TriCB | 22 | | | 23.4 | 0.976 (S) | 0.92 | 0.873 |
| 2,3,5-TriCB | 23 | | U | | 0.929 (S) | | |
| 2,3,6-TriCB | 24 | | U | | 0.846 (Q) | | |
| 2,3',4-TriCB | 25 | | | 9.93 | 0.846 (Q) | 1.00 | 0.826 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 13.0 | 0.874 (S) | 0.95 | 1.302 |
| 2,3',6-TriCB | 27 | | J | 3.71 | 0.846 (Q) | 1.11 | 1.152 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 58.7 | 0.846 (Q) | 0.98 | 0.837 |
| 2,4',6-TriCB | 32 | | | 17.3 | 0.846 (Q) | 1.05 | 1.198 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | K J | 1.08 | 0.942 (S) | 1.78 | 1.276 |
| 3,3',4-TriCB | 35 | | K J | 4.09 | 0.936 (S) | 0.77 | 0.985 |
| 3,3',5-TriCB | 36 | | K J | 1.65 | 0.878 (S) | 0.66 | 0.932 |
| 3,4,4'-TriCB | 37 | | | 17.8 | 0.846 (Q) | 1.03 | 1.001 |
| 3,4,5-TriCB | 38 | | K J | 2.37 | 0.846 (Q) | 1.98 | 0.968 |
| 3,4',5-TriCB | 39 | | U | | 0.873 (S) | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 38.0 | 0.846 (Q) | 0.76 | 1.337 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | | 18.8 | 0.846 (Q) | 0.75 | 1.312 |
| 2,2',3,5'-TeCB | 43 | | J | 3.98 | 0.846 (Q) | 0.79 | 1.248 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 203 | 0.846 (Q) | 0.76 | 1.287 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 103 | 0.846 (Q) | 0.75 | 1.149 |
| 2,2',3,6'-TeCB | 46 | | K J | 4.48 | 0.846 (Q) | 1.02 | 1.161 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | | 16.5 | 0.846 (Q) | 0.84 | 1.275 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 65.3 | 0.846 (Q) | 0.82 | 1.260 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C | 12.6 | 0.846 (Q) | 0.80 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 191 | 0.846 (Q) | 0.78 | 1.234 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | U | | 0.846 (Q) | | |
| 2,3,3',4'-TeCB | 55 | | U | | 3.32 (S) | | |
| 2,3,3',4'-TeCB | 56 | | | 54.0 | 3.19 (S) | 0.74 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 2.94 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 3.17 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 7.20 | 0.846 (Q) | 0.75 | 1.302 |
| 2,3,4,4'-TeCB | 60 | | K | 20.1 | 3.05 (S) | 0.61 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 234 | 2.91 (S) | 0.77 | 0.875 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | U | | 2.98 (S) | | |
| 2,3,4',6'-TeCB | 64 | | | 40.7 | 0.846 (Q) | 0.76 | 1.349 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | G | 117 | 3.03 (S) | 0.78 | 0.884 |
| 2,3',4,5'-TeCB | 67 | | U | | 2.48 (S) | | |
| 2,3',4,5'-TeCB | 68 | | | 38.9 | 2.81 (S) | 0.80 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | U | | 2.87 (S) | | |
| 2,3',5',6'-TeCB | 73 | | U | | 0.846 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | | 15.7 | 2.36 (S) | 0.87 | 1.001 |
| 3,3',4,5'-TeCB | 78 | | U | | 3.03 (S) | | |
| 3,3',4,5'-TeCB | 79 | | | 7.78 | 2.32 (S) | 0.75 | 0.971 |
| 3,3',5,5'-TeCB | 80 | | U | | 2.73 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 2.27 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | | 42.9 | 0.846 (Q) | 1.50 | 0.934 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 222 | 0.846 (Q) | 1.58 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | | 90.1 | 0.846 (Q) | 1.50 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 70.2 | 0.846 (Q) | 1.60 | 0.920 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C G | 263 | 0.846 (Q) | 1.53 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 51.0 | 0.846 (Q) | 1.54 | 1.154 |
| 2,2',3,4,6'-PeCB | 89 | | J | 3.48 | 0.846 (Q) | 1.39 | 1.181 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 369 | 0.846 (Q) | 1.58 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 73.2 | 0.846 (Q) | 1.57 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 273 | 0.846 (Q) | 1.66 | 1.120 |
| 2,2',3,5,6'-PeCB | 94 | | K J | 1.22 | 0.846 (Q) | 0.38 | 1.104 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | K J | 1.93 | 0.846 (Q) | 1.06 | 1.014 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | J | 3.74 | 0.846 (Q) | 1.59 | 1.093 |
| 2,2',4,6,6'-PeCB | 104 | | U | | 0.846 (Q) | | |
| 2,3,3',4,4'-PeCB | 105 | | | 147 | 0.931 (S) | 1.48 | 1.000 |
| 2,3,3',4,5-PeCB | 106 | | U | | 1.10 (S) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C | 21.3 | 1.19 (S) | 1.54 | 0.991 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | | 25.5 | 1.08 (S) | 1.45 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 425 | 0.846 (Q) | 1.60 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.846 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.846 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | K J | 6.59 | 0.936 (S) | 1.17 | 1.000 |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | | 318 | 0.993 (S) | 1.49 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | U | | 0.846 (Q) | | |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.846 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | K J | 4.06 | 1.30 (S) | 1.04 | 1.010 |
| 2',3,4,4',5-PeCB | 123 | | K J | 5.24 | 0.983 (S) | 1.74 | 1.001 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | K J | 2.12 | 0.999 (S) | 1.45 | 1.000 |
| 3,3',4,5,5'-PeCB | 127 | | U | | 1.22 (S) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 95.9 | 0.846 (Q) | 1.32 | 0.959 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 572 | 0.846 (Q) | 1.26 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 38.5 | 0.846 (Q) | 1.13 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | K | 8.60 | 0.846 (Q) | 0.79 | 1.159 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 191 | 0.846 (Q) | 1.29 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | K | 10.2 | 0.846 (Q) | 1.55 | 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C K | 30.2 | 0.846 (Q) | 1.04 | 1.139 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 150 | 0.846 (Q) | 1.19 | 1.103 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 49.9 | 0.846 (Q) | 1.42 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | | 31.6 | 0.846 (Q) | 1.41 | 0.918 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C K | 12.2 | 0.846 (Q) | 1.49 | 1.152 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 95.9 | 0.846 (Q) | 1.12 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 0.846 (Q) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 19.0 | 0.846 (Q) | 1.18 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.846 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | | 80.9 | 0.846 (Q) | 1.20 | 0.885 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 384 | 0.846 (Q) | 1.25 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | U | | 0.846 (Q) | | |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | K J | 1.13 | 0.846 (Q) | 0.47 | 1.012 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | U | | 0.846 (Q) | | |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 442 | 0.846 (Q) | 1.26 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | K J | 1.27 | 0.846 (Q) | 0.28 | 1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 69.8 | 0.846 (Q) | 1.17 | 1.000 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 60.2 | 0.846 (Q) | 1.24 | 0.938 |
| 2,3,3',4,5,5'-HxCB | 159 | | K | 6.98 | 0.846 (Q) | 1.03 | 0.981 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

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| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 0.846 (Q) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | U | | 0.846 (Q) | | |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 37.9 | 0.846 (Q) | 1.28 | 0.921 |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 0.846 (Q) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 23.1 | 0.846 (Q) | 1.20 | 1.000 |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 0.846 (Q) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 110 | 0.846 (Q) | 0.99 | 1.000 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C K | 38.4 | 0.846 (Q) | 1.46 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 23.3 | 0.846 (Q) | 0.94 | 0.897 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 113 | 0.846 (Q) | 1.02 | 1.133 |
| 2,2',3,3',4,5',6-HpCB | 175 | | K J | 4.76 | 0.846 (Q) | 1.38 | 1.102 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 14.3 | 0.846 (Q) | 1.13 | 1.034 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 65.8 | 0.846 (Q) | 1.10 | 1.145 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 30.0 | 0.846 (Q) | 1.11 | 1.085 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 45.8 | 0.846 (Q) | 1.13 | 1.010 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 280 | 0.846 (Q) | 1.01 | 1.001 |
| 2,2',3,4,4',5,6-HpCB | 181 | | U | | 0.846 (Q) | | |
| 2,2',3,4,4',5,6'-HpCB | 182 | | U | | 0.846 (Q) | | |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 70.5 | 0.846 (Q) | 1.06 | 1.127 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | J | 1.85 | 0.846 (Q) | 1.04 | 1.025 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.846 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 155 | 0.846 (Q) | 1.01 | 1.110 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | U | | 0.846 (Q) | | |
| 2,3,3',4,4',5,5'-HpCB | 189 | | K J | 2.75 | 0.846 (Q) | 1.71 | 1.000 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 28.0 | 0.846 (Q) | 1.05 | 0.947 |
| 2,3,3',4,4',5',6-HpCB | 191 | | K J | 4.59 | 0.846 (Q) | 1.41 | 0.918 |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.846 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OcCB | 194 | | | 65.5 | 0.846 (Q) | 0.91 | 0.991 |
| 2,2',3,3',4,4',5,6-OcCB | 195 | | | 27.5 | 0.846 (Q) | 0.86 | 0.946 |
| 2,2',3,3',4,4',5,6'-OcCB | 196 | | | 30.9 | 0.846 (Q) | 0.90 | 0.916 |
| 2,2',3,3',4,4',6,6'-OcCB | 197 | 197 + 200 | C | 11.3 | 0.846 (Q) | 0.80 | 1.046 |
| 2,2',3,3',4,5,5',6-OcCB | 198 | 198 + 199 | C | 86.2 | 0.846 (Q) | 0.88 | 1.114 |
| 2,2',3,3',4,5,5',6'-OcCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OcCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OcCB | 201 | | | 10.7 | 0.846 (Q) | 0.80 | 1.023 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 23.6 | 0.846 (Q) | 1.01 | 1.000 |
| 2,2',3,4,4',5,5',6-OcCB | 203 | | | 49.6 | 0.846 (Q) | 0.77 | 0.920 |
| 2,2',3,4,4',5,6,6'-OcCB | 204 | | U | | 0.846 (Q) | | |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | J | 3.44 | 0.846 (Q) | 0.88 | 1.000 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 53.4 | 2.97 (S) | 0.74 | 1.000 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | K | 7.37 | 2.28 (S) | 0.49 | 1.020 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 21.0 | 2.67 (S) | 0.82 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 49.3 | 0.846 (Q) | 1.22 | 1.001 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-WS-T06-1811
Sample Collection:
30-Nov-2018 16:26

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 31-Jan-2019 Time: 02:01:34
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-7
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_028 S: 7
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_028 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG 2 | SPIKE CONC. | CONC. FOUND | R(%) 3 | ION ABUND. RATIO | RRT |
|-------------------------------------|----------------|-------------|---------------|----------------|----------------|--------|------------------------|-------|
| 13C12-2-MoCB | 1L | | G | 4000 | 750 | 18.7 | 3.53 | 0.719 |
| 13C12-4-MoCB | 3L | | | 4000 | 1580 | 39.6 | 3.10 | 0.857 |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 1560 | 39.1 | 1.57 | 0.874 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 1920 | 47.9 | 1.53 | 1.252 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2650 | 66.3 | 1.08 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1450 | 36.1 | 0.99 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1980 | 49.4 | 0.77 | 0.811 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1800 | 45.1 | 0.69 | 1.396 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1780 | 44.4 | 0.72 | 1.373 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2020 | 50.6 | 1.58 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1700 | 42.5 | 1.57 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1520 | 37.9 | 1.55 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1610 | 40.3 | 1.56 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1660 | 41.4 | 1.56 | 1.151 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1600 | 40.0 | 1.55 | 1.300 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2980 | 74.5 | 1.29 | 0.786 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 4980 | 62.3 | 1.25 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2530 | 63.2 | 1.25 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2580 | 64.5 | 1.28 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 3150 | 78.7 | 1.00 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 3240 | 81.0 | 1.03 | 0.872 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2890 | 72.2 | 1.08 | 0.712 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1800 | 44.9 | 0.96 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2510 | 62.8 | 0.89 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2860 | 71.4 | 0.87 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3370 | 84.2 | 0.78 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3050 | 76.3 | 0.76 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3610 | 90.3 | 1.19 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1590 | 39.7 | 0.98 | 0.924 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2330 | 58.2 | 1.62 | 1.088 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 2930 | 73.2 | 1.02 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Kristen Bowes _____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-RB-XF-181129
Sample Collection:
29-Nov-2018 09:50

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---|
| Contract No.: | 4972 | Project No. | PORTLAND HARBOR PDI AND BASELINE WATER |
| Matrix: | FILTER | Lab Sample I.D.: | L30523-8 |
| Sample Receipt Date: | 04-Dec-2018 | Sample Size: | 1 sample |
| Extraction Date: | 14-Jan-2019 | Initial Calibration Date: | 15-Jan-2019 |
| Analysis Date: | 31-Jan-2019 Time: 03:05:49 | Instrument ID: | HR GC/MS |
| Extract Volume (uL): | 20 | GC Column ID: | SPB OCTYL |
| Injection Volume (uL): | 1.0 | Sample Data Filename: | PB9C_028 S: 8 |
| Dilution Factor: | N/A | Blank Data Filename: | PB9C_027 S: 5 |
| Concentration Units: | pg/sample | Cal. Ver. Data Filename: | PB9C_028 S: 1 |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | NQ | | | | |
| 3-MoCB | 2 | | NQ | | | | |
| 4-MoCB | 3 | | NQ | | | | |
| 2,2'-DiCB | 4 | | K G | 16.9 | 5.35 (S) | 1.87 | 1.001 |
| 2,3-DiCB | 5 | | NQ | | | | |
| 2,3'-DiCB | 6 | | NQ | | | | |
| 2,4-DiCB | 7 | | NQ | | | | |
| 2,4'-DiCB | 8 | | NQ | | | | |
| 2,5-DiCB | 9 | | NQ | | | | |
| 2,6-DiCB | 10 | | NQ | | | | |
| 3,3'-DiCB | 11 | | NQ | | | | |
| 3,4-DiCB | 12 | 12 + 13 | C NQ | | | | |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | NQ | | | | |
| 4,4'-DiCB | 15 | | NQ | | | | |
| 2,2',3-TriCB | 16 | | K | 16.1 | 0.841 (Q) | 1.67 | 1.159 |
| 2,2',4-TriCB | 17 | | | 57.2 | 0.841 (Q) | 1.04 | 1.133 |
| 2,2',5-TriCB | 18 | 18 + 30 | C K | 41.4 | 0.841 (Q) | 1.31 | 1.109 |
| 2,2',6-TriCB | 19 | | K J | 6.37 | 0.841 (Q) | 1.29 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 43.2 | 0.843 (S) | 0.95 | 0.850 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 26.6 | 0.847 (S) | 0.88 | 0.858 |
| 2,3,4'-TriCB | 22 | | | 14.3 | 0.983 (S) | 0.92 | 0.874 |
| 2,3,5-TriCB | 23 | | U | | 0.936 (S) | | |
| 2,3,6-TriCB | 24 | | K J | 0.890 | 0.841 (Q) | 1.96 | 1.154 |
| 2,3',4-TriCB | 25 | | K | 8.82 | 0.841 (Q) | 0.65 | 0.827 |
| 2,3',5-TriCB | 26 | 26 + 29 | C K | 8.39 | 0.880 (S) | 0.82 | 1.293 |
| 2,3',6-TriCB | 27 | | J | 2.42 | 0.841 (Q) | 1.05 | 1.145 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | K | 32.6 | 0.841 (Q) | 0.84 | 0.839 |
| 2,4',6-TriCB | 32 | | | 14.0 | 0.852 (S) | 1.02 | 1.191 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | U | | 0.949 (S) | | |
| 3,3',4-TriCB | 35 | | K J | 1.39 | 0.943 (S) | 0.23 | 0.985 |
| 3,3',5-TriCB | 36 | | U | | 0.885 (S) | | |
| 3,4,4'-TriCB | 37 | | K J | 5.15 | 0.841 (Q) | 1.60 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.841 (Q) | | |
| 3,4',5-TriCB | 39 | | K J | 1.76 | 0.879 (S) | 0.65 | 0.947 |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C K | 9.35 | 0.841 (Q) | 0.65 | 1.330 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | K | 6.88 | 0.841 (Q) | 0.99 | 1.307 |
| 2,2',3,5'-TeCB | 43 | | U | | 0.841 (Q) | | |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 103 | 0.841 (Q) | 0.83 | 1.282 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C | 98.8 | 0.841 (Q) | 0.66 | 1.146 |
| 2,2',3,6'-TeCB | 46 | | K J | 1.73 | 0.841 (Q) | 0.45 | 1.157 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | J | 4.72 | 0.841 (Q) | 0.76 | 1.269 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 15.6 | 0.841 (Q) | 0.71 | 1.255 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C J | 3.74 | 0.841 (Q) | 0.72 | 1.108 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 28.0 | 0.841 (Q) | 0.82 | 1.230 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | U | | 0.841 (Q) | | |
| 2,3,3',4'-TeCB | 55 | | U | | 1.75 (S) | | |
| 2,3,3',4'-TeCB | 56 | | K J | 4.20 | 1.68 (S) | 0.56 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 1.55 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 1.68 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C K J | 1.65 | 0.841 (Q) | 0.63 | 1.295 |
| 2,3,4,4'-TeCB | 60 | | J | 2.47 | 1.61 (S) | 0.77 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 18.1 | 1.54 (S) | 0.68 | 0.876 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | U | | 1.57 (S) | | |
| 2,3,4',6'-TeCB | 64 | | J | 6.40 | 0.841 (Q) | 0.80 | 1.343 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | K | 7.78 | 1.60 (S) | 0.97 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | U | | 1.31 (S) | | |
| 2,3',4,5'-TeCB | 68 | | | 54.0 | 1.48 (S) | 0.72 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | U | | 1.52 (S) | | |
| 2,3',5',6'-TeCB | 73 | | U | | 0.841 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | U | | 1.16 (S) | | |
| 3,3',4,5'-TeCB | 78 | | U | | 1.60 (S) | | |
| 3,3',4,5'-TeCB | 79 | | U | | 1.23 (S) | | |
| 3,3',5,5'-TeCB | 80 | | U | | 1.44 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 1.20 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | U | | 0.841 (Q) | | |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C J | 4.32 | 0.841 (Q) | 1.68 | 0.887 |
| 2,2',3,3',6'-PeCB | 84 | | K J | 3.98 | 0.841 (Q) | 2.28 | 1.161 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C K J | 1.37 | 0.841 (Q) | 2.98 | 0.922 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C J G | 5.52 | 0.841 (Q) | 1.57 | 0.902 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C K J | 1.94 | 0.841 (Q) | 1.84 | 1.153 |
| 2,2',3,4,6'-PeCB | 89 | | U | | 0.841 (Q) | | |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C K | 8.89 | 0.841 (Q) | 1.17 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | J | 1.31 | 0.841 (Q) | 1.62 | 0.854 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C G | 7.92 | 0.841 (Q) | 1.63 | 1.118 |
| 2,2',3,5,6'-PeCB | 94 | | U | | 0.841 (Q) | | |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | U | | 0.841 (Q) | | |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | U | | 0.841 (Q) | | |
| 2,2',4,6,6'-PeCB | 104 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4'-PeCB | 105 | | K J | 2.42 | 0.841 (Q) | 3.16 | 1.001 |
| 2,3,3',4,5-PeCB | 106 | | U | | 0.841 (Q) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C K J | 1.07 | 0.841 (Q) | 4.73 | 0.991 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | J | 1.05 | 0.841 (Q) | 1.64 | 0.997 |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C | 8.86 | 0.841 (Q) | 1.53 | 0.926 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.841 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.841 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | U | | 0.841 (Q) | | |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | K | 7.17 | 0.841 (Q) | 2.15 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | U | | 0.841 (Q) | | |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.841 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | U | | 0.841 (Q) | | |
| 2',3,4,4',5-PeCB | 123 | | U | | 0.841 (Q) | | |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | U | | 0.841 (Q) | | |
| 3,3',4,5,5'-PeCB | 127 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C K J | 1.92 | 0.841 (Q) | 0.79 | 0.957 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 8.54 | 0.841 (Q) | 1.32 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | K J | 1.39 | 0.841 (Q) | 2.42 | 0.913 |
| 2,2',3,3',4,6-HxCB | 131 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,6'-HxCB | 132 | | K J | 4.04 | 0.841 (Q) | 0.80 | 1.172 |
| 2,2',3,3',5,5'-HxCB | 133 | | U | | 0.841 (Q) | | |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C U | | 0.841 (Q) | | |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C J | 3.62 | 0.841 (Q) | 1.26 | 1.101 |
| 2,2',3,3',6,6'-HxCB | 136 | | K J | 1.56 | 0.841 (Q) | 0.76 | 1.023 |
| 2,2',3,4,4',5-HxCB | 137 | | U | | 0.841 (Q) | | |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C U | | 0.841 (Q) | | |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | K J | 2.44 | 0.841 (Q) | 2.74 | 0.903 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 0.841 (Q) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | K J | 1.16 | 0.841 (Q) | 0.77 | 1.121 |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.841 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | K J | 1.58 | 0.841 (Q) | 1.63 | 0.884 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C K | 9.76 | 0.841 (Q) | 1.02 | 1.132 |
| 2,2',3,4',5,6'-HxCB | 148 | | U | | 0.841 (Q) | | |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | U | | 0.841 (Q) | | |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | U | | 0.841 (Q) | | |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C K | 7.18 | 0.841 (Q) | 1.03 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C U | | 0.841 (Q) | | |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | K J | 1.01 | 0.841 (Q) | 0.69 | 0.939 |
| 2,3,3',4,5,5'-HxCB | 159 | | U | | 0.841 (Q) | | |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 0.841 (Q) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | U | | 0.841 (Q) | | |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | U | | 0.841 (Q) | | |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 0.841 (Q) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | U | | 0.841 (Q) | | |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | K J | 1.29 | 0.841 (Q) | 0.71 | 1.000 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C J | 1.13 | 0.841 (Q) | 1.06 | 1.162 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | K J | 1.25 | 0.841 (Q) | 3.03 | 1.132 |
| 2,2',3,3',4,5',6-HpCB | 175 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,6,6'-HpCB | 176 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4',5,6-HpCB | 177 | | K J | 0.888 | 0.841 (Q) | 0.62 | 1.144 |
| 2,2',3,3',5,5',6-HpCB | 178 | | U | | 0.841 (Q) | | |
| 2,2',3,3',5,6,6'-HpCB | 179 | | K J | 1.63 | 0.841 (Q) | 0.26 | 1.009 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C K J | 4.28 | 0.841 (Q) | 0.61 | 1.000 |
| 2,2',3,4,4',5,6-HpCB | 181 | | U | | 0.841 (Q) | | |
| 2,2',3,4,4',5,6'-HpCB | 182 | | U | | 0.841 (Q) | | |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C K J | 1.68 | 0.841 (Q) | 0.85 | 1.126 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | U | | 0.841 (Q) | | |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.841 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | K J | 2.63 | 0.841 (Q) | 0.87 | 1.109 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4',5,5'-HpCB | 189 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4',5,6-HpCB | 190 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4',5',6-HpCB | 191 | | U | | 0.841 (Q) | | |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.841 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C U | | 0.841 (Q) | | |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C K J | 1.26 | 0.841 (Q) | 0.35 | 1.115 |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | U | | 0.841 (Q) | | |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | U | | 0.841 (Q) | | |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | K J | 1.13 | 0.841 (Q) | 0.53 | 0.920 |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.841 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | U | | 0.841 (Q) | | |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | U | | 5.89 (S) | | |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | U | | 4.14 (S) | | |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | U | | 4.51 (S) | | |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | K J | 2.57 | 0.841 (Q) | 0.59 | 1.001 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener; NQ = data not quantifiable.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

Form 2
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
PDI-RB-XF-181129
Sample Collection:
29-Nov-2018 09:50

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: 04-Dec-2018
Extraction Date: 14-Jan-2019
Analysis Date: 31-Jan-2019 Time: 03:05:49
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absolute

Project No. PORTLAND HARBOR PDI AND
BASELINE WATER
Lab Sample I.D.: L30523-8
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_028 S: 8
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_028 S: 1

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | NQ | | | | | |
| 13C12-4-MoCB | 3L | | NQ | | | | | |
| 13C12-2,2'-DiCB | 4L | | G | 4000 | 1040 | 25.9 | 1.59 | 0.889 |
| 13C12-4,4'-DiCB | 15L | | NQ | | | | | |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 2140 | 53.4 | 1.04 | 1.070 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1120 | 28.0 | 1.02 | 1.089 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 1570 | 39.3 | 0.78 | 0.814 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 1590 | 39.8 | 0.71 | 1.393 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 4000 | 1480 | 37.0 | 0.72 | 1.370 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 1580 | 39.6 | 1.57 | 0.810 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1240 | 31.1 | 1.57 | 1.199 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 4000 | 1250 | 31.3 | 1.61 | 1.178 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 4000 | 1240 | 31.1 | 1.53 | 1.161 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 4000 | 1270 | 31.7 | 1.54 | 1.150 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 4000 | 1070 | 26.7 | 1.53 | 1.299 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2420 | 60.4 | 1.27 | 0.787 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 8000 | 3550 | 44.4 | 1.27 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 1830 | 45.7 | 1.22 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 1760 | 44.0 | 1.27 | 1.191 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 4000 | 2510 | 62.7 | 1.07 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 2570 | 64.1 | 1.07 | 0.873 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 2740 | 68.6 | 1.07 | 0.713 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1330 | 33.3 | 0.98 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 1990 | 49.7 | 0.89 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 1720 | 43.1 | 0.85 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 2100 | 52.6 | 0.74 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 2230 | 55.8 | 0.76 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 2260 | 56.5 | 1.18 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1760 | 44.0 | 1.01 | 0.925 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2520 | 63.0 | 1.59 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3040 | 76.0 | 1.03 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener; NQ = data not quantifiable.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 1A
PCB CONGENER ANALYSIS REPORT

CLIENT SAMPLE NO.
Lab Blank
Sample Collection:
N/A

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: N/A
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 13:59:01
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg/sample

Project No. N/A
Lab Sample I.D.: WG66481-101
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_027 S: 5
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_027 S: 1

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|--------------|-----------|-------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2-MoCB | 1 | | NQ | | | | |
| 3-MoCB | 2 | | NQ | | | | |
| 4-MoCB | 3 | | NQ | | | | |
| 2,2'-DiCB | 4 | | | 13.6 | 3.71 (S) | 1.65 | 1.000 |
| 2,3-DiCB | 5 | | U | | 3.70 (S) | | |
| 2,3'-DiCB | 6 | | | 9.64 | 3.35 (S) | 1.59 | 1.168 |
| 2,4-DiCB | 7 | | U | | 3.43 (S) | | |
| 2,4'-DiCB | 8 | | | 38.2 | 3.07 (S) | 1.44 | 1.198 |
| 2,5-DiCB | 9 | | K J | 3.78 | 3.23 (S) | 1.99 | 1.139 |
| 2,6-DiCB | 10 | | U | | 3.31 (S) | | |
| 3,3'-DiCB | 11 | | | 32.7 | 3.73 (S) | 1.62 | 0.969 |
| 3,4-DiCB | 12 | 12 + 13 | C U | | 3.54 (S) | | |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | |
| 3,5-DiCB | 14 | | U | | 3.44 (S) | | |
| 4,4'-DiCB | 15 | | | 10.5 | 3.08 (S) | 1.76 | 1.001 |
| 2,2',3-TriCB | 16 | | | 14.8 | 0.864 (Q) | 1.00 | 1.164 |
| 2,2',4-TriCB | 17 | | | 14.7 | 0.864 (Q) | 0.99 | 1.136 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 33.4 | 0.864 (Q) | 1.01 | 1.112 |
| 2,2',6-TriCB | 19 | | J | 3.78 | 0.864 (Q) | 1.12 | 1.001 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C K | 28.5 | 0.864 (Q) | 0.86 | 0.849 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 15.4 | 0.864 (Q) | 1.01 | 0.858 |
| 2,3,4'-TriCB | 22 | | | 10.4 | 0.864 (Q) | 1.01 | 0.873 |
| 2,3,5-TriCB | 23 | | U | | 0.864 (Q) | | |
| 2,3,6-TriCB | 24 | | J | 0.894 | 0.864 (Q) | 1.10 | 1.158 |
| 2,3',4-TriCB | 25 | | J | 2.22 | 0.864 (Q) | 0.90 | 0.826 |
| 2,3',5-TriCB | 26 | 26 + 29 | C K J | 5.58 | 0.864 (Q) | 0.80 | 1.299 |
| 2,3',6-TriCB | 27 | | K J | 2.18 | 0.864 (Q) | 0.77 | 1.149 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | |
| 2,4',5-TriCB | 31 | | | 23.7 | 0.864 (Q) | 0.97 | 0.838 |
| 2,4',6-TriCB | 32 | | K G | 8.93 | 0.864 (Q) | 1.23 | 1.196 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | |
| 2',3,5-TriCB | 34 | | U | | 0.864 (Q) | | |
| 3,3',4-TriCB | 35 | | K J | 1.10 | 0.864 (Q) | 1.73 | 0.985 |
| 3,3',5-TriCB | 36 | | U | | 0.864 (Q) | | |
| 3,4,4'-TriCB | 37 | | J | 4.13 | 0.864 (Q) | 1.19 | 1.001 |
| 3,4,5-TriCB | 38 | | U | | 0.864 (Q) | | |
| 3,4',5-TriCB | 39 | | U | | 0.864 (Q) | | |

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This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 7.79 | 0.864 (Q) | 0.77 | 1.335 |
| 2,2',3,4'-TeCB | 41 | 40 + 41 + 71 | C40 | | | | |
| 2,2',3,4'-TeCB | 42 | | K J | 3.77 | 0.864 (Q) | 1.11 | 1.311 |
| 2,2',3,5'-TeCB | 43 | | U | | 0.864 (Q) | | |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 17.2 | 0.864 (Q) | 0.73 | 1.286 |
| 2,2',3,6'-TeCB | 45 | 45 + 51 | C J | 4.97 | 0.864 (Q) | 0.78 | 1.146 |
| 2,2',3,6'-TeCB | 46 | | J | 2.12 | 0.864 (Q) | 0.74 | 1.160 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | |
| 2,2',4,5'-TeCB | 48 | | K J | 4.21 | 0.864 (Q) | 1.06 | 1.273 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 7.34 | 0.864 (Q) | 0.73 | 1.259 |
| 2,2',4,6'-TeCB | 50 | 50 + 53 | C K J | 2.30 | 0.864 (Q) | 1.05 | 1.111 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | |
| 2,2',5,5'-TeCB | 52 | | | 18.5 | 0.864 (Q) | 0.88 | 1.233 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | |
| 2,2',6,6'-TeCB | 54 | | U | | 0.864 (Q) | | |
| 2,3,3',4'-TeCB | 55 | | U | | 1.45 (S) | | |
| 2,3,3',4'-TeCB | 56 | | J | 1.84 | 1.37 (S) | 0.70 | 0.905 |
| 2,3,3',5'-TeCB | 57 | | U | | 1.29 (S) | | |
| 2,3,3',5'-TeCB | 58 | | U | | 1.34 (S) | | |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C K J | 1.44 | 0.864 (Q) | 0.61 | 1.301 |
| 2,3,4,4'-TeCB | 60 | | K J | 1.60 | 1.31 (S) | 1.95 | 0.911 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 11.0 | 1.27 (S) | 0.76 | 0.876 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | |
| 2,3,4',5'-TeCB | 63 | | U | | 1.28 (S) | | |
| 2,3,4',6'-TeCB | 64 | | K J | 5.83 | 0.864 (Q) | 0.65 | 1.348 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | |
| 2,3',4,4'-TeCB | 66 | | J G | 5.23 | 1.30 (S) | 0.67 | 0.885 |
| 2,3',4,5'-TeCB | 67 | | U | | 1.09 (S) | | |
| 2,3',4,5'-TeCB | 68 | | K J | 1.68 | 1.22 (S) | 0.62 | 0.832 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | |
| 2,3',5,5'-TeCB | 72 | | U | | 1.27 (S) | | |
| 2,3',5',6'-TeCB | 73 | | U | | 0.864 (Q) | | |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | |
| 3,3',4,4'-TeCB | 77 | | U | | 1.01 (S) | | |
| 3,3',4,5'-TeCB | 78 | | U | | 1.23 (S) | | |
| 3,3',4,5'-TeCB | 79 | | U | | 1.01 (S) | | |
| 3,3',5,5'-TeCB | 80 | | U | | 1.16 (S) | | |
| 3,4,4',5'-TeCB | 81 | | U | | 1.01 (S) | | |
| 2,2',3,3',4'-PeCB | 82 | | K J | 1.26 | 0.864 (Q) | 0.93 | 0.933 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C J | 4.08 | 0.864 (Q) | 1.38 | 0.886 |
| 2,2',3,3',6'-PeCB | 84 | | J | 2.10 | 0.864 (Q) | 1.45 | 1.162 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C K J | 1.06 | 0.864 (Q) | 0.98 | 0.919 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C K J G | 6.03 | 0.864 (Q) | 0.74 | 0.901 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C K J | 0.870 | 0.864 (Q) | 0.41 | 1.153 |
| 2,2',3,4,6'-PeCB | 89 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C K | 6.99 | 0.864 (Q) | 1.88 | 0.870 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | |
| 2,2',3,5,5'-PeCB | 92 | | J | 1.12 | 0.864 (Q) | 1.54 | 0.853 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C K J | 6.08 | 0.864 (Q) | 1.93 | 1.119 |
| 2,2',3,5,6'-PeCB | 94 | | U | | 0.864 (Q) | | |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',3,6,6'-PeCB | 96 | | U | | 0.864 (Q) | | |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|---------------------|-----------|--------------------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,2',4,4',6-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | |
| 2,2',4,5',6-PeCB | 103 | | U | | 0.864 (Q) | | |
| 2,2',4,6,6'-PeCB | 104 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4'-PeCB | 105 | | J | 2.69 | 0.864 (Q) | 1.73 | 1.001 |
| 2,3,3',4,5-PeCB | 106 | | U | | 0.864 (Q) | | |
| 2,3,3',4',5-PeCB | 107 | 107 + 124 | C U | | 0.864 (Q) | | |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3,3',4,6-PeCB | 109 | | U | | 0.864 (Q) | | |
| 2,3,3',4',6-PeCB | 110 | 110 + 115 | C K | 6.94 | 0.864 (Q) | 1.82 | 0.925 |
| 2,3,3',5,5'-PeCB | 111 | | U | | 0.864 (Q) | | |
| 2,3,3',5,6-PeCB | 112 | | U | | 0.864 (Q) | | |
| 2,3,3',5',6-PeCB | 113 | 90 + 101 + 113 | C90 | | | | |
| 2,3,4,4',5-PeCB | 114 | | U | | 0.864 (Q) | | |
| 2,3,4,4',6-PeCB | 115 | 110 + 115 | C110 | | | | |
| 2,3,4,5,6-PeCB | 116 | 85 + 116 + 117 | C85 | | | | |
| 2,3,4',5,6-PeCB | 117 | 85 + 116 + 117 | C85 | | | | |
| 2,3',4,4',5-PeCB | 118 | | K J | 5.73 | 0.864 (Q) | 1.17 | 1.000 |
| 2,3',4,4',6-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 2,3',4,5,5'-PeCB | 120 | | U | | 0.864 (Q) | | |
| 2,3',4,5',6-PeCB | 121 | | U | | 0.864 (Q) | | |
| 2',3,3',4,5-PeCB | 122 | | U | | 0.864 (Q) | | |
| 2',3,4,4',5-PeCB | 123 | | U | | 0.864 (Q) | | |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | |
| 3,3',4,4',5-PeCB | 126 | | U | | 0.864 (Q) | | |
| 3,3',4,5,5'-PeCB | 127 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C J | 1.10 | 0.864 (Q) | 1.07 | 0.958 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C K J | 5.07 | 0.864 (Q) | 0.89 | 0.929 |
| 2,2',3,3',4,5'-HxCB | 130 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,6-HxCB | 131 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,6'-HxCB | 132 | | K J | 3.60 | 0.864 (Q) | 1.56 | 1.173 |
| 2,2',3,3',5,5'-HxCB | 133 | | U | | 0.864 (Q) | | |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C U | | 0.864 (Q) | | |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C K J | 1.97 | 0.864 (Q) | 2.70 | 1.102 |
| 2,2',3,3',6,6'-HxCB | 136 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5-HxCB | 137 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C U | | 0.864 (Q) | | |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | K J | 1.07 | 0.864 (Q) | 1.44 | 0.904 |
| 2,2',3,4,5,6-HxCB | 142 | | U | | 0.864 (Q) | | |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | U | | 0.864 (Q) | | |
| 2,2',3,4,6,6'-HxCB | 145 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5,5'-HxCB | 146 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C J | 3.20 | 0.864 (Q) | 1.20 | 1.133 |
| 2,2',3,4',5,6'-HxCB | 148 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | U | | 0.864 (Q) | | |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | U | | 0.864 (Q) | | |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C K J | 4.01 | 0.864 (Q) | 1.50 | 0.899 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6'-HxCB | 155 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C U | | 0.864 (Q) | | |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | U | | 0.864 (Q) | | |
| 2,3,3',4,5,5'-HxCB | 159 | | U | | 0.864 (Q) | | |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |

This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | REPORTING LIMIT (RL) ² | ION ABUND. RATIO | RRT |
|-------------------------------|-----------|-----------------------|-----------------------|-------------|-----------------------------------|------------------|-------|
| 2,3,3',4,5',6-HxCB | 161 | | U | | 0.864 (Q) | | |
| 2,3,3',4',5,5'-HxCB | 162 | | U | | 0.864 (Q) | | |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | U | | 0.864 (Q) | | |
| 2,3,3',5,5',6-HxCB | 165 | | U | | 0.864 (Q) | | |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | U | | 0.864 (Q) | | |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | |
| 3,3',4,4',5,5'-HxCB | 169 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C U | | 0.864 (Q) | | |
| 2,2',3,3',4,5,5'-HpCB | 172 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,5',6-HpCB | 175 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,6,6'-HpCB | 176 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4',5,6-HpCB | 177 | | U | | 0.864 (Q) | | |
| 2,2',3,3',5,5',6-HpCB | 178 | | U | | 0.864 (Q) | | |
| 2,2',3,3',5,6,6'-HpCB | 179 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C U | | 0.864 (Q) | | |
| 2,2',3,4,4',5,6-HpCB | 181 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5,6'-HpCB | 182 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C U | | 0.864 (Q) | | |
| 2,2',3,4,4',6,6'-HpCB | 184 | | U | | 0.864 (Q) | | |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5,5',6-HpCB | 187 | | U | | 0.864 (Q) | | |
| 2,2',3,4',5,6,6'-HpCB | 188 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4',5,5'-HpCB | 189 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4',5,6-HpCB | 190 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4',5',6-HpCB | 191 | | U | | 0.864 (Q) | | |
| 2,3,3',4,5,5',6-HpCB | 192 | | U | | 0.864 (Q) | | |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5'-OxCB | 194 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',5,6-OxCB | 195 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',5,6'-OxCB | 196 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',6,6'-OxCB | 197 | 197 + 200 | C U | | 0.864 (Q) | | |
| 2,2',3,3',4,5,5',6-OxCB | 198 | 198 + 199 | C U | | 0.864 (Q) | | |
| 2,2',3,3',4,5,5',6'-OxCB | 199 | 198 + 199 | C198 | | | | |
| 2,2',3,3',4,5,6,6'-OxCB | 200 | 197 + 200 | C197 | | | | |
| 2,2',3,3',4,5',6,6'-OxCB | 201 | | U | | 0.864 (Q) | | |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5,5',6-OxCB | 203 | | U | | 0.864 (Q) | | |
| 2,2',3,4,4',5,6,6'-OxCB | 204 | | U | | 0.864 (Q) | | |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | U | | 0.864 (Q) | | |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | U | | 3.35 (S) | | |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | U | | 2.54 (S) | | |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | U | | 2.84 (S) | | |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | K J | 3.15 | 0.864 (Q) | 0.87 | 1.000 |

(1) Where applicable, custom lab flags have been used on this report; U = not detected at RL; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than lowest calibration equivalent; G = lock mass interference present; C = co-eluting congener; NQ = data not quantifiable.

(2) Reporting Limit (Code): S = sample detection limit; M = method detection limit; L = lowest calibration level equivalent; Q = minimum reporting level.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 2
PCB CONGENER ANALYSIS REPORTCLIENT SAMPLE NO.
Lab Blank
Sample Collection:
N/A

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811Contract No.: 4972
Matrix: FILTER
Sample Receipt Date: N/A
Extraction Date: 14-Jan-2019
Analysis Date: 30-Jan-2019 Time: 13:59:01
Extract Volume (uL): 20
Injection Volume (uL): 1.0
Dilution Factor: N/A
Concentration Units: pg absoluteProject No. N/A
Lab Sample I.D.: WG66481-101
Sample Size: 1 sample
Initial Calibration Date: 15-Jan-2019
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB9C_027 S: 5
Blank Data Filename: PB9C_027 S: 5
Cal. Ver. Data Filename: PB9C_027 S: 1This page is part of a total report that contains information necessary for accreditation compliance.
This test is not NELAP accredited. Sample results relate only to the sample tested.

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | SPIKE CONC. | CONC. FOUND | R(%) ³ | ION ABUND. RATIO | RRT |
|-------------------------------------|------------------------|-------------|-----------------------|-------------|-------------|-------------------|------------------|-------|
| 13C12-2-MoCB | 1L | | NQ | | | | | |
| 13C12-4-MoCB | 3L | | NQ | | | | | |
| 13C12-2,2'-DiCB | 4L | | | 4000 | 2020 | 50.5 | 1.56 | 0.879 |
| 13C12-4,4'-DiCB | 15L | | | 4000 | 2180 | 54.6 | 1.53 | 1.249 |
| 13C12-2,2',6-TriCB | 19L | | | 4000 | 3070 | 76.6 | 1.03 | 1.072 |
| 13C12-3,4,4'-TriCB | 37L | | | 4000 | 1650 | 41.3 | 1.01 | 1.090 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 4000 | 2320 | 58.0 | 0.78 | 0.811 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 4000 | 2170 | 54.2 | 0.70 | 1.395 |
| 13C12-3,4,4',5-TeCB | 81L | | | 4000 | 2140 | 53.6 | 0.70 | 1.372 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 4000 | 2220 | 55.5 | 1.57 | 0.808 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 4000 | 1970 | 49.2 | 1.53 | 1.199 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 4000 | 1800 | 45.0 | 1.55 | 1.178 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 4000 | 1820 | 45.6 | 1.50 | 1.161 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 4000 | 1970 | 49.4 | 1.54 | 1.150 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 4000 | 1890 | 47.2 | 1.53 | 1.299 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 4000 | 2670 | 66.8 | 1.27 | 0.787 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 8000 | 4920 | 61.5 | 1.26 | 1.107 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 4000 | 2460 | 61.5 | 1.24 | 1.078 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 4000 | 2530 | 63.2 | 1.29 | 1.191 |
| 13C12-2,2',3,3',4,4',5-HpCB | 170L | | | 4000 | 2970 | 74.3 | 1.06 | 0.897 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 4000 | 2840 | 71.0 | 1.07 | 0.872 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 4000 | 3420 | 85.5 | 1.06 | 0.712 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 4000 | 1950 | 48.6 | 0.93 | 0.958 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 4000 | 2560 | 64.1 | 0.86 | 0.818 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 4000 | 2900 | 72.5 | 0.84 | 1.009 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 4000 | 3300 | 82.6 | 0.77 | 1.043 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 4000 | 3250 | 81.3 | 0.78 | 0.949 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 4000 | 3570 | 89.3 | 1.15 | 1.075 |
| CLEANUP STANDARD | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 4000 | 1750 | 43.6 | 1.01 | 0.925 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 4000 | 2700 | 67.5 | 1.63 | 1.087 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 4000 | 3130 | 78.3 | 1.05 | 1.012 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener; NQ = data not quantifiable.

(3) R% = percent recovery of labeled compounds.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____Kristen Bowes_____

SGS AXYS METHOD MLA-010 Rev 12

Form 8A

PCB CONGENER ONGOING PRECISION AND RECOVERY (OPR)

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|-------------------------------|----------------------------|----------------------------------|---------------|
| Contract No.: | 4972 | Lab Sample I.D.: | WG66481-102 |
| Matrix: | FILTER | Initial Calibration Date: | 15-Jan-2019 |
| Extraction Date: | 16-Jan-2019 | Instrument ID: | HR GC/MS |
| Analysis Date: | 30-Jan-2019 Time: 10:46:16 | GC Column ID: | SPB OCTYL |
| Extract Volume (uL): | 20 | OPR Data Filename: | PB9C_027 S: 2 |
| Injection Volume (uL): | 1.0 | Blank Data Filename: | PB9C_027 S: 5 |
| Dilution Factor: | N/A | Cal. Ver. Data Filename: | PB9C_027 S: 1 |

CONCENTRATIONS REPORTED ARE CONCENTRATIONS IN EXTRACT, BASED ON A 20 uL EXTRACT VOLUME.

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | ION ABUND. RATIO | SPIKE CONC. (ng/mL) | CONC. FOUND (ng/mL) | OPR CONC. LIMITS (ng/mL) | % RECOVERY |
|-------------------------------|-----------|-------------|-----------------------|------------------|---------------------|---------------------|--------------------------|------------|
| 2-MoCB | 1 | | NQ | | | | | |
| 4-MoCB | 3 | | NQ | | | | | |
| 2,2'-DiCB | 4 | | G | 1.48 | 100 | 80.9 | 50.0 - 150 | 80.9 |
| 4,4'-DiCB | 15 | | | 1.47 | 100 | 72.3 | 50.0 - 150 | 72.3 |
| 2,2',6-TriCB | 19 | | | 1.06 | 100 | 93.2 | 50.0 - 150 | 93.2 |
| 3,4,4'-TriCB | 37 | | | 0.98 | 100 | 77.2 | 50.0 - 150 | 77.2 |
| 2,2',6,6'-TeCB | 54 | | | 0.77 | 100 | 91.3 | 50.0 - 150 | 91.3 |
| 3,3',4,4'-TeCB | 77 | | | 0.76 | 100 | 81.6 | 50.0 - 150 | 81.6 |
| 3,4,4',5-TeCB | 81 | | | 0.76 | 100 | 82.2 | 50.0 - 150 | 82.2 |
| 2,2',4,6,6'-PeCB | 104 | | | 1.56 | 100 | 102 | 50.0 - 150 | 102 |
| 2,3,3',4,4'-PeCB | 105 | | | 1.50 | 100 | 82.1 | 50.0 - 150 | 82.1 |
| 2,3,4,4',5-PeCB | 114 | | | 1.51 | 100 | 79.6 | 50.0 - 150 | 79.6 |
| 2,3',4,4',5-PeCB | 118 | | | 1.49 | 100 | 82.9 | 50.0 - 150 | 82.9 |
| 2',3,4,4',5-PeCB | 123 | | | 1.49 | 100 | 80.4 | 50.0 - 150 | 80.4 |
| 3,3',4,4',5-PeCB | 126 | | | 1.46 | 100 | 79.4 | 50.0 - 150 | 79.4 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 1.29 | 100 | 104 | 50.0 - 150 | 104 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 1.25 | 200 | 182 | 100 - 300 | 91.0 |
| 2,3,3',4,4',5',5'-HxCB | 157 | 156 + 157 | C156 | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 1.24 | 100 | 91.6 | 50.0 - 150 | 91.6 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 1.24 | 100 | 91.4 | 50.0 - 150 | 91.4 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 1.05 | 100 | 98.6 | 50.0 - 150 | 98.6 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 1.00 | 100 | 82.9 | 50.0 - 150 | 82.9 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 0.89 | 100 | 104 | 50.0 - 150 | 104 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | 0.89 | 100 | 94.0 | 50.0 - 150 | 94.0 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 0.80 | 100 | 98.7 | 50.0 - 150 | 98.7 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 0.77 | 100 | 101 | 50.0 - 150 | 101 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 1.19 | 100 | 100 | 50.0 - 150 | 100 |

(1) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener; NQ = data not quantifiable.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Kristen Bowes _____

These pages are part of a larger report that may contain information necessary for full data evaluation. Results reported relate only to the sample tested.

For Axy Internal Use Only [XSL Template: Form16688A.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5; Report Filename: 1668_PCB1668_PCBTF_WG66481-102_Form8A_SJ2506285.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 8B

PCB CONGENER ONGOING PRECISION AND RECOVERY (OPR)

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

| | | | |
|------------------------|----------------------------|---------------------------|---------------|
| Contract No.: | 4972 | Lab Sample I.D.: | WG66481-102 |
| Matrix: | FILTER | Initial Calibration Date: | 15-Jan-2019 |
| Extraction Date: | 16-Jan-2019 | Instrument ID: | HR GC/MS |
| Analysis Date: | 30-Jan-2019 Time: 10:46:16 | GC Column ID: | SPB OCTYL |
| Extract Volume (uL): | 20 | OPR Data Filename: | PB9C_027 S: 2 |
| Injection Volume (uL): | 1.0 | Blank Data Filename: | PB9C_027 S: 5 |
| Dilution Factor: | N/A | Cal. Ver. Data Filename: | PB9C_027 S: 1 |

CONCENTRATIONS REPORTED ARE CONCENTRATIONS IN EXTRACT, BASED ON A 20 uL EXTRACT VOLUME.

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG ² | ION ABUND. RATIO | SPIKE CONC. (ng/mL) | CONC. FOUND (ng/mL) | OPR CONC. LIMITS (ng/mL) | % RECOVERY |
|-------------------------------------|-------------|-------------|-----------------------|------------------|---------------------|---------------------|--------------------------|------------|
| 13C12-2-MoCB | 1L | | NQ | | | | | |
| 13C12-4-MoCB | 3L | | NQ | | | | | |
| 13C12-2,2'-DiCB | 4L | | G | 1.54 | 200 | 101 | 60.0 - 280 | 50.3 |
| 13C12-4,4'-DiCB | 15L | | | 1.53 | 200 | 113 | 60.0 - 280 | 56.4 |
| 13C12-2,2',6-TriCB | 19L | | | 1.05 | 200 | 157 | 60.0 - 280 | 78.3 |
| 13C12-3,4,4'-TriCB | 37L | | | 0.99 | 200 | 83.3 | 60.0 - 280 | 41.7 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 0.78 | 200 | 117 | 60.0 - 280 | 58.3 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 0.71 | 200 | 103 | 60.0 - 280 | 51.3 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 0.71 | 200 | 103 | 60.0 - 280 | 51.4 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 1.58 | 200 | 109 | 60.0 - 280 | 54.7 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 1.58 | 200 | 90.4 | 60.0 - 280 | 45.2 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 1.57 | 200 | 89.6 | 60.0 - 280 | 44.8 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 1.54 | 200 | 87.2 | 60.0 - 280 | 43.6 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 1.54 | 200 | 88.0 | 60.0 - 280 | 44.0 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 1.49 | 200 | 86.7 | 60.0 - 280 | 43.4 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 1.26 | 200 | 140 | 60.0 - 280 | 70.2 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 1.23 | 400 | 259 | 120 - 560 | 64.7 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 1.27 | 200 | 130 | 60.0 - 280 | 64.8 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 1.25 | 200 | 131 | 60.0 - 280 | 65.5 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 1.06 | 200 | 182 | 60.0 - 280 | 91.0 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 0.92 | 200 | 90.1 | 60.0 - 280 | 45.1 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 0.88 | 200 | 139 | 60.0 - 280 | 69.3 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 0.82 | 200 | 139 | 60.0 - 280 | 69.3 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 0.75 | 200 | 162 | 60.0 - 280 | 80.9 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 0.78 | 200 | 154 | 60.0 - 280 | 76.9 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 1.21 | 200 | 166 | 60.0 - 280 | 83.1 |

CLEANUP STANDARD

| | | | | | | | | |
|-----------------------------|------|--|--|------|-----|------|------------|------|
| 13C12-2,4,4'-TriCB | 28L | | | 1.00 | 200 | 89.5 | 80.0 - 250 | 44.7 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 1.62 | 200 | 126 | 80.0 - 250 | 63.1 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 1.05 | 200 | 150 | 80.0 - 250 | 75.0 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; G = lock mass interference present; C = co-eluting congener; NQ = data not quantifiable.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Kristen Bowes _____

These pages are part of a larger report that may contain information necessary for full data evaluation. Results reported relate only to the sample tested.

SGS AXYS METHOD MLA-010 Rev 12

Form 3A

PCB CONGENERS INITIAL CALIBRATION RELATIVE RESPONSES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

Instrument ID: HR GC/MS

GC Column ID: SPB OCTYL

CS0 Data Filename: PB9C_009E S: 3

CS1 Data Filename: PB9C_009E S: 5

CS2 Data Filename: PB9C_009F S: 1

CS3 Data Filename: PB9C_009E S: 8

CS4 Data Filename: PB9C_009E S: 7

CS5 Data Filename: PB9C_009F S: 3

CS6 Data Filename: PB9C_009F S: 4

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RELATIVE RESPONSE (RR) | | | | | | MEAN RR | CV ² (%RSD) |
|-------------------------------|-----------|-------------|-----------------------|------------------------|------|------|------|------|------|---------|------------------------|
| | | | | CS0 | CS1 | CS2 | CS3 | CS4 | CS5 | | |
| 2-MoCB | 1 | | | 1.09 | 1.09 | 1.14 | 1.13 | 1.14 | 1.03 | 1.10 | 3.74 |
| 4-MoCB | 3 | | | 1.21 | 1.11 | 1.11 | 1.12 | 1.14 | 1.09 | 1.13 | 3.73 |
| 2,2'-DiCB | 4 | | | 0.99 | 0.94 | 1.00 | 1.01 | 1.04 | 1.06 | 1.01 | 4.15 |
| 4,4'-DiCB | 15 | | | 0.91 | 0.86 | 0.90 | 0.93 | 0.94 | 0.96 | 0.92 | 3.85 |
| 2,2',6-TriCB | 19 | | | 1.16 | 1.07 | 1.10 | 1.10 | 1.14 | 1.15 | 1.11 | 2.75 |
| 3,4,4'-TriCB | 37 | | | 0.99 | 0.96 | 1.02 | 1.01 | 1.05 | 1.07 | 0.97 | 3.77 |
| 2,2',6,6'-TeCB | 54 | | | 1.08 | 1.02 | 1.07 | 1.10 | 1.12 | 1.15 | 1.07 | 3.77 |
| 3,3',4,4'-TeCB | 77 | | | 1.11 | 1.02 | 1.05 | 1.03 | 1.05 | 1.07 | 1.04 | 2.97 |
| 3,4,4',5-TeCB | 81 | | | 1.04 | 1.01 | 1.01 | 1.05 | 1.08 | 1.10 | 1.05 | 3.16 |
| 2,2',4,6,6'-PeCB | 104 | | | 1.14 | 1.08 | 1.07 | 1.09 | 1.13 | 1.17 | 1.13 | 3.08 |
| 2,3,3',4,4'-PeCB | 105 | | | 0.98 | 1.02 | 1.01 | 1.02 | 1.05 | 1.09 | 1.05 | 3.46 |
| 2,3,4,4',5-PeCB | 114 | | | 1.06 | 1.06 | 1.06 | 1.07 | 1.11 | 1.15 | 1.07 | 3.10 |
| 2,3',4,4',5-PeCB | 118 | | | 0.97 | 0.95 | 1.00 | 1.01 | 1.05 | 1.11 | 1.07 | 5.35 |
| 2',3,4,4',5-PeCB | 123 | | | 0.88 | 0.95 | 0.95 | 0.96 | 1.00 | 1.04 | 1.01 | 5.19 |
| 3,3',4,4',5-PeCB | 126 | | | 1.03 | 0.99 | 1.00 | 1.04 | 1.06 | 1.10 | 1.07 | 3.53 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 1.01 | 1.02 | 1.02 | 1.07 | 1.09 | 1.09 | 1.05 | 3.69 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 1.11 | 1.07 | 1.15 | 1.14 | 1.16 | 1.19 | 1.10 | 3.68 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 1.10 | 1.08 | 1.14 | 1.16 | 1.18 | 1.22 | 1.18 | 4.24 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 1.08 | 1.02 | 1.08 | 1.09 | 1.11 | 1.14 | 1.09 | 3.21 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 0.99 | 0.97 | 0.97 | 0.98 | 1.01 | 1.03 | 0.99 | 2.33 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 0.97 | 1.00 | 1.04 | 1.03 | 1.04 | 1.09 | 1.04 | 3.90 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 0.88 | 0.84 | 0.83 | 0.86 | 0.88 | 0.94 | 0.92 | 4.32 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | 1.01 | 0.94 | 1.01 | 1.00 | 1.02 | 1.05 | 1.01 | 3.39 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 1.12 | 0.91 | 1.00 | 0.99 | 1.01 | 1.03 | 0.98 | 6.25 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 0.89 | 0.95 | 0.96 | 0.97 | 1.00 | 1.02 | 0.97 | 4.34 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 1.35 | 1.06 | 1.01 | 1.03 | 1.04 | 1.07 | 1.02 | 11.2 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) For contract CV specifications, see Section 10.4.4, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Jason MacKenzie _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 3B

PCB CONGENERS INITIAL CALIBRATION RELATIVE RESPONSES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

Instrument ID: HR GC/MS

GC Column ID: SPB OCTYL

CS0 Data Filename: PB9C_009E S: 3

CS1 Data Filename: PB9C_009E S: 5

CS2 Data Filename: PB9C_009F S: 1

CS3 Data Filename: PB9C_009E S: 8

CS4 Data Filename: PB9C_009E S: 7

CS5 Data Filename: PB9C_009F S: 3

CS6 Data Filename: PB9C_009F S: 4

| COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RELATIVE RESPONSE (RR) | | | | | | MEAN RR | CV ³ (%RSD) | |
|-------------------------------------|------------------------|-------------|-----------------------|------------------------|------|------|------|------|------|---------|------------------------|------|
| | | | | CS0 | CS1 | CS2 | CS3 | CS4 | CS5 | | | CS6 |
| 13C12-2-MoCB | 1L | | | 1.12 | 1.12 | 1.16 | 1.14 | 1.15 | 1.18 | 1.21 | 1.16 | 2.74 |
| 13C12-4-MoCB | 3L | | | 1.06 | 1.08 | 1.10 | 1.08 | 1.13 | 1.20 | 1.27 | 1.13 | 6.80 |
| 13C12-2,2'-DiCB | 4L | | | 0.65 | 0.65 | 0.66 | 0.66 | 0.68 | 0.71 | 0.77 | 0.68 | 6.23 |
| 13C12-4,4'-DiCB | 15L | | | 0.99 | 1.01 | 0.99 | 0.99 | 1.09 | 1.20 | 1.32 | 1.08 | 12.0 |
| 13C12-2,2',6-TriCB | 19L | | | 0.46 | 0.45 | 0.45 | 0.48 | 0.49 | 0.54 | 0.60 | 0.50 | 11.3 |
| 13C12-3,4,4'-TriCB | 37L | | | 1.84 | 1.86 | 1.88 | 1.85 | 1.96 | 2.30 | 2.69 | 2.05 | 15.7 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 1.47 | 1.47 | 1.52 | 1.52 | 1.60 | 1.75 | 2.03 | 1.62 | 12.5 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 1.43 | 1.46 | 1.47 | 1.42 | 1.52 | 1.79 | 2.13 | 1.60 | 16.5 |
| 13C12-3,4,4',5-TeCB | 81L | | | 1.45 | 1.46 | 1.49 | 1.43 | 1.49 | 1.76 | 2.13 | 1.60 | 16.2 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 1.24 | 1.22 | 1.25 | 1.31 | 1.39 | 1.62 | 1.93 | 1.42 | 18.4 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 1.51 | 1.49 | 1.49 | 1.46 | 1.56 | 1.79 | 2.10 | 1.63 | 14.6 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 1.54 | 1.52 | 1.53 | 1.51 | 1.67 | 1.92 | 2.10 | 1.69 | 14.0 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 1.48 | 1.45 | 1.44 | 1.43 | 1.54 | 1.74 | 2.09 | 1.60 | 15.2 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 1.47 | 1.47 | 1.45 | 1.43 | 1.56 | 1.78 | 2.14 | 1.61 | 16.2 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 1.42 | 1.45 | 1.46 | 1.41 | 1.53 | 1.72 | 1.96 | 1.56 | 13.2 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 1.27 | 1.22 | 1.27 | 1.30 | 1.45 | 1.75 | | 1.37 | 14.5 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 1.29 | 1.31 | 1.29 | 1.31 | 1.46 | 1.72 | 1.95 | 1.47 | 17.8 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | | | | | |
| 13C12-2,3,3',4,4',5,5'-HxCB | 167L | | | 1.26 | 1.29 | 1.25 | 1.26 | 1.38 | 1.57 | 1.80 | 1.40 | 14.9 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 1.40 | 1.42 | 1.36 | 1.39 | 1.50 | 1.69 | 1.89 | 1.52 | 13.0 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 1.34 | 1.28 | 1.36 | 1.43 | 1.60 | 2.02 | | 1.51 | 18.2 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 1.51 | 1.47 | 1.46 | 1.46 | 1.57 | 1.68 | 1.72 | 1.55 | 6.94 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 1.34 | 1.33 | 1.39 | 1.42 | 1.42 | 1.64 | 1.81 | 1.48 | 12.1 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 1.39 | 1.36 | 1.36 | 1.37 | 1.43 | 1.57 | 1.60 | 1.44 | 6.98 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 0.93 | 0.90 | 0.92 | 0.92 | 0.96 | 1.03 | 1.08 | 0.96 | 6.90 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 1.13 | 1.11 | 1.10 | 1.15 | 1.20 | 1.32 | 1.46 | 1.21 | 11.1 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 1.02 | 0.97 | 1.02 | 1.01 | 1.05 | 1.17 | 1.20 | 1.06 | 8.12 |
| CLEAN-UP STANDARD | | | | | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | 1.92 | 1.92 | 1.90 | 1.90 | 1.87 | 1.87 | 1.85 | 1.89 | 1.31 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 1.30 | 1.30 | 1.31 | 1.30 | 1.38 | 1.47 | 1.69 | 1.39 | 10.5 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 0.85 | 0.84 | 0.82 | 0.84 | 0.85 | 0.85 | 0.87 | 0.85 | 1.81 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) For contract CV specifications, see Section 10.4.4, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Jason MacKenzie _____

SGS AXYS METHOD MLA-010 Rev 12

Form 3C
PCB CONGENER INITIAL CALIBRATION ION ABUNDANCE RATIOS

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

Instrument ID: HR GC/MS

GC Column ID: SPB OCTYL

CS0 Data Filename: PB9C_009E S: 3

CS1 Data Filename: PB9C_009E S: 5

CS2 Data Filename: PB9C_009F S: 1

CS3 Data Filename: PB9C_009E S: 8

CS4 Data Filename: PB9C_009E S: 7

CS5 Data Filename: PB9C_009F S: 3

CS6 Data Filename: PB9C_009F S: 4

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | M/Z's FORMING RATIO ² | ION ABUNDANCE RATIO | | | | | | QC LIMITS ² | |
|-------------------------------|-----------|-------------|-----------------------|----------------------------------|---------------------|------|------|------|------|------|------------------------|-----------|
| | | | | | CS0 | CS1 | CS2 | CS3 | CS4 | CS5 | | CS6 |
| 2-MoCB | 1 | | | M/M+2 | 3.13 | 3.21 | 3.21 | 3.12 | 3.12 | 2.67 | | 2.66-3.60 |
| 4-MoCB | 3 | | | M/M+2 | 3.57 | 3.29 | 3.14 | 3.12 | 3.12 | 2.91 | | 2.66-3.60 |
| 2,2'-DiCB | 4 | | | M/M+2 | 1.69 | 1.58 | 1.53 | 1.55 | 1.55 | 1.54 | | 1.33-1.79 |
| 4,4'-DiCB | 15 | | | M/M+2 | 1.46 | 1.55 | 1.56 | 1.55 | 1.54 | 1.56 | | 1.33-1.79 |
| 2,2',6-TriCB | 19 | | | M/M+2 | 1.03 | 0.97 | 1.02 | 1.08 | 1.07 | 1.04 | 1.05 | 0.88-1.20 |
| 3,4,4'-TriCB | 37 | | | M/M+2 | 1.08 | 1.04 | 1.01 | 1.02 | 1.04 | 1.04 | 1.02 | 0.88-1.20 |
| 2,2',6,6'-TeCB | 54 | | | M/M+2 | 0.73 | 0.80 | 0.78 | 0.79 | 0.79 | 0.79 | 0.82 | 0.65-0.89 |
| 3,3',4,4'-TeCB | 77 | | | M/M+2 | 0.80 | 0.81 | 0.76 | 0.78 | 0.77 | 0.78 | 0.77 | 0.65-0.89 |
| 3,4,4',5-TeCB | 81 | | | M/M+2 | 0.72 | 0.74 | 0.76 | 0.78 | 0.78 | 0.77 | 0.78 | 0.65-0.89 |
| 2,2',4,6,6'-PeCB | 104 | | | M+2/M+4 | 1.53 | 1.48 | 1.47 | 1.57 | 1.56 | 1.56 | 1.55 | 1.32-1.78 |
| 2,3,3',4,4'-PeCB | 105 | | | M+2/M+4 | 1.55 | 1.51 | 1.58 | 1.57 | 1.57 | 1.54 | 1.55 | 1.32-1.78 |
| 2,3,4,4',5-PeCB | 114 | | | M+2/M+4 | 1.50 | 1.72 | 1.61 | 1.59 | 1.60 | 1.61 | 1.61 | 1.32-1.78 |
| 2,3',4,4',5-PeCB | 118 | | | M+2/M+4 | 1.46 | 1.55 | 1.62 | 1.56 | 1.55 | 1.55 | 1.56 | 1.32-1.78 |
| 2',3,4,4',5-PeCB | 123 | | | M+2/M+4 | 1.73 | 1.53 | 1.58 | 1.57 | 1.54 | 1.54 | 1.55 | 1.32-1.78 |
| 3,3',4,4',5-PeCB | 126 | | | M+2/M+4 | 1.39 | 1.57 | 1.54 | 1.57 | 1.55 | 1.55 | 1.55 | 1.32-1.78 |
| 2,2',4,4',6,6'-HxCB | 155 | | | M+2/M+4 | 1.42 | 1.22 | 1.27 | 1.28 | 1.25 | 1.25 | | 1.05-1.43 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | M+2/M+4 | 1.16 | 1.22 | 1.27 | 1.26 | 1.25 | 1.26 | 1.21 | 1.05-1.43 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | M+2/M+4 | 1.42 | 1.25 | 1.28 | 1.27 | 1.25 | 1.26 | 1.26 | 1.05-1.43 |
| 3,3',4,4',5,5'-HxCB | 169 | | | M+2/M+4 | 1.01 | 1.23 | 1.21 | 1.25 | 1.27 | 1.27 | 1.27 | 1.05-1.43 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | M+2/M+4 | 1.00 | 1.19 | 1.03 | 1.05 | 1.05 | 1.04 | | 0.89-1.21 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | M+2/M+4 | 1.11 | 0.94 | 1.03 | 1.04 | 1.05 | 1.05 | 1.05 | 0.89-1.21 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | M+2/M+4 | 0.81 | 0.92 | 0.94 | 0.89 | 0.89 | 0.90 | 0.90 | 0.76-1.02 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | M+2/M+4 | 0.85 | 0.86 | 0.89 | 0.89 | 0.89 | 0.90 | 0.89 | 0.76-1.02 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | M+2/M+4 | 0.81 | 0.78 | 0.79 | 0.78 | 0.78 | 0.79 | 0.79 | 0.65-0.89 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | M+2/M+4 | 0.88 | 0.76 | 0.75 | 0.79 | 0.79 | 0.79 | 0.78 | 0.65-0.89 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | M+4/M+6 | 1.22 | 1.05 | 1.28 | 1.20 | 1.18 | 1.17 | 1.17 | 0.99-1.33 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) See Table 8 Method 1668A for m/z specifications and ion abundance ratio control limits.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Jason MacKenzie _____

For Axys Internal Use Only [XSL Template: Form16683C.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5; Report Filename: 1668_PCB1668_15-Jan-2019_PB9C_Form3C_GS80037.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 3D
PCB CONGENER INITIAL CALIBRATION ION ABUNDANCE RATIOS

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

Instrument ID: HR GC/MS

GC Column ID: SPB OCTYL

CS0 Data Filename: PB9C_009E S: 3

CS1 Data Filename: PB9C_009E S: 5

CS2 Data Filename: PB9C_009F S: 1

CS3 Data Filename: PB9C_009E S: 8

CS4 Data Filename: PB9C_009E S: 7

CS5 Data Filename: PB9C_009F S: 3

CS6 Data Filename: PB9C_009F S: 4

| LABELED COMPOUND | IUPAC NO. ¹ | CO- ELUTIONS | LAB FLAG ² | M/Z's FORMING RATIO ³ | ION ABUNDANCE RATIO | | | | | | QC LIMITS ³ | |
|-------------------------------------|---------------------------|-----------------|--------------------------|--|---------------------|------|------|------|------|------|---------------------------|-----------|
| | | | | | CS0 | CS1 | CS2 | CS3 | CS4 | CS5 | | CS6 |
| 13C12-2-MoCB | 1L | | | M/M+2 | 3.24 | 3.21 | 3.15 | 3.18 | 3.21 | 3.19 | 3.19 | 2.66-3.60 |
| 13C12-4-MoCB | 3L | | | M/M+2 | 3.17 | 3.18 | 3.20 | 3.18 | 3.14 | 3.11 | 3.13 | 2.66-3.60 |
| 13C12-2,2'-DiCB | 4L | | | M/M+2 | 1.59 | 1.60 | 1.59 | 1.59 | 1.61 | 1.59 | 1.58 | 1.33-1.79 |
| 13C12-4,4'-DiCB | 15L | | | M/M+2 | 1.60 | 1.59 | 1.60 | 1.57 | 1.59 | 1.58 | 1.57 | 1.33-1.79 |
| 13C12-2,2',6-TriCB | 19L | | | M/M+2 | 1.07 | 1.07 | 1.07 | 1.06 | 1.07 | 1.08 | 1.08 | 0.88-1.20 |
| 13C12-3,4,4'-TriCB | 37L | | | M/M+2 | 1.04 | 1.05 | 1.05 | 1.05 | 1.05 | 1.04 | 1.04 | 0.88-1.20 |
| 13C12-2,2',6,6'-TeCB | 54L | | | M/M+2 | 0.79 | 0.79 | 0.79 | 0.79 | 0.78 | 0.79 | 0.79 | 0.65-0.89 |
| 13C12-3,3',4,4'-TeCB | 77L | | | M/M+2 | 0.72 | 0.74 | 0.76 | 0.74 | 0.74 | 0.76 | 0.76 | 0.65-0.89 |
| 13C12-3,4,4',5-TeCB | 81L | | | M/M+2 | 0.72 | 0.72 | 0.75 | 0.71 | 0.73 | 0.72 | 0.75 | 0.65-0.89 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | M+2/M+4 | 1.60 | 1.60 | 1.58 | 1.58 | 1.59 | 1.57 | 1.59 | 1.32-1.78 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | M+2/M+4 | 1.58 | 1.56 | 1.57 | 1.59 | 1.56 | 1.56 | 1.57 | 1.32-1.78 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | M+2/M+4 | 1.62 | 1.61 | 1.65 | 1.64 | 1.62 | 1.65 | 1.64 | 1.32-1.78 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | M+2/M+4 | 1.57 | 1.59 | 1.57 | 1.58 | 1.55 | 1.56 | 1.54 | 1.32-1.78 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | M+2/M+4 | 1.57 | 1.58 | 1.58 | 1.56 | 1.56 | 1.54 | 1.57 | 1.32-1.78 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | M+2/M+4 | 1.55 | 1.56 | 1.57 | 1.58 | 1.56 | 1.58 | 1.56 | 1.32-1.78 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | M+2/M+4 | 1.25 | 1.26 | 1.25 | 1.24 | 1.26 | 1.27 | 1.27 | 1.05-1.43 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | M+2/M+4 | 1.27 | 1.27 | 1.25 | 1.26 | 1.28 | 1.25 | 1.25 | 1.05-1.43 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | M+2/M+4 | 1.26 | 1.27 | 1.26 | 1.25 | 1.28 | 1.24 | 1.24 | 1.05-1.43 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | M+2/M+4 | 1.26 | 1.27 | 1.23 | 1.26 | 1.27 | 1.26 | 1.23 | 1.05-1.43 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | M+2/M+4 | 1.05 | 1.06 | 1.06 | 1.06 | 1.05 | 1.07 | | 0.89-1.21 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | M+2/M+4 | 1.01 | 1.02 | 0.99 | 1.00 | 1.01 | 1.02 | 1.04 | 0.89-1.21 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | M+2/M+4 | 0.91 | 0.90 | 0.89 | 0.92 | 0.93 | 0.91 | 0.90 | 0.76-1.02 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | M+2/M+4 | 0.86 | 0.87 | 0.87 | 0.87 | 0.87 | 0.86 | 0.87 | 0.76-1.02 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | M+2/M+4 | 0.76 | 0.77 | 0.77 | 0.77 | 0.78 | 0.77 | 0.78 | 0.65-0.89 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | M+2/M+4 | 0.77 | 0.76 | 0.74 | 0.77 | 0.77 | 0.77 | 0.77 | 0.65-0.89 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | M+4/M+6 | 1.17 | 1.19 | 1.19 | 1.17 | 1.18 | 1.18 | 1.14 | 0.99-1.33 |
| CLEAN-UP STANDARD | | | | | | | | | | | | |
| 13C12-2,4,4'-TriCB | 28L | | | M/M+2 | 1.05 | 1.05 | 1.06 | 1.04 | 1.05 | 1.06 | 1.05 | 0.88-1.20 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | M+2/M+4 | 1.59 | 1.60 | 1.61 | 1.58 | 1.61 | 1.62 | 1.59 | 1.32-1.78 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | M+2/M+4 | 1.05 | 1.06 | 1.04 | 1.04 | 1.04 | 1.06 | 1.04 | 0.89-1.21 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) See Table 8 Method 1668A for m/z specifications and ion abundance ratio control limits.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Jason MacKenzie _____

SGS AXYS METHOD MLA-010 Rev 12

Form 4A
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_027 S: 1
 Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
 GC Column ID: SPB OCTYL Analysis Time: 09:39:33

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | MZ's FORMING RATIO ² | ION ABUND. RATIO | QC LIMITS ³ | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------|-----------|-------------|-----------------------|---------------------------------|------------------|------------------------|---------------------|---------------------|
| 2-MoCB | 1 | | | M/M+2 | 2.98 | 2.66-3.60 | 19.0 | 17.5 - 32.5 |
| 4-MoCB | 3 | | | M/M+2 | 2.98 | 2.66-3.60 | 19.4 | 17.5 - 32.5 |
| 2,2'-DiCB | 4 | | | M/M+2 | 1.51 | 1.33-1.79 | 19.0 | 17.5 - 32.5 |
| 4,4'-DiCB | 15 | | | M/M+2 | 1.50 | 1.33-1.79 | 22.2 | 19.6 - 36.4 |
| 2,2',6-TriCB | 19 | | | M/M+2 | 1.03 | 0.88-1.20 | 24.9 | 17.5 - 32.5 |
| 3,4,4'-TriCB | 37 | | | M/M+2 | 0.98 | 0.88-1.20 | 19.2 | 17.5 - 32.5 |
| 2,2',6,6'-TeCB | 54 | | | M/M+2 | 0.78 | 0.65-0.89 | 46.7 | 35.0 - 65.0 |
| 3,3',4,4'-TeCB | 77 | | | M/M+2 | 0.77 | 0.65-0.89 | 40.1 | 35.0 - 65.0 |
| 3,4,4',5-TeCB | 81 | | | M/M+2 | 0.77 | 0.65-0.89 | 42.2 | 35.0 - 65.0 |
| 2,2',4,6,6'-PeCB | 104 | | | M+2/M+4 | 1.54 | 1.32-1.78 | 51.5 | 35.0 - 65.0 |
| 2,3,3',4,4'-PeCB | 105 | | | M+2/M+4 | 1.53 | 1.32-1.78 | 42.5 | 35.0 - 65.0 |
| 2,3,4,4',5-PeCB | 114 | | | M+2/M+4 | 1.55 | 1.32-1.78 | 41.9 | 35.0 - 65.0 |
| 2,3',4,4',5-PeCB | 118 | | | M+2/M+4 | 1.53 | 1.32-1.78 | 40.8 | 35.0 - 65.0 |
| 2',3,4,4',5-PeCB | 123 | | | M+2/M+4 | 1.53 | 1.32-1.78 | 41.3 | 35.0 - 65.0 |
| 3,3',4,4',5-PeCB | 126 | | | M+2/M+4 | 1.55 | 1.32-1.78 | 43.5 | 39.0 - 72.4 |
| 2,2',4,4',6,6'-HxCB | 155 | | | M+2/M+4 | 1.29 | 1.05-1.43 | 53.3 | 35.0 - 65.0 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | M+2/M+4 | 1.24 | 1.05-1.43 | 92.3 | 70.0 - 130 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | M+2/M+4 | 1.23 | 1.05-1.43 | 51.1 | 35.0 - 65.0 |
| 3,3',4,4',5,5'-HxCB | 169 | | | M+2/M+4 | 1.24 | 1.05-1.43 | 49.7 | 35.0 - 65.0 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | M+2/M+4 | 1.05 | 0.89-1.21 | 48.8 | 35.0 - 65.0 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | M+2/M+4 | 1.01 | 0.89-1.21 | 42.6 | 35.0 - 65.0 |
| 2,2',3,3',5,5',6,6'-OoCB | 202 | | | M+2/M+4 | 0.92 | 0.76-1.02 | 85.8 | 58.9 - 110 |
| 2,3,3',4,4',5,5',6-OoCB | 205 | | | M+2/M+4 | 0.87 | 0.76-1.02 | 70.0 | 52.5 - 97.5 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | M+2/M+4 | 0.78 | 0.65-0.89 | 73.5 | 52.5 - 97.5 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | M+2/M+4 | 0.78 | 0.65-0.89 | 80.8 | 58.7 - 109 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | M+4/M+6 | 1.18 | 0.99-1.33 | 71.5 | 52.5 - 97.5 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

For Axy Internal Use Only [XSL Template: Form16684A.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5;
 Report Filename: 1668_PCB1668_PB9C_027S1__Form4A_SJ2506283.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 4B
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_027 S: 1
Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
GC Column ID: SPB OCTYL Analysis Time: 09:39:33

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG 2 | MZ's FORMING RATIO 3 | ION ABUND. RATIO | QC LIMITS 4 | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------------|-------------|-------------|------------|----------------------|------------------|-------------|---------------------|---------------------|
| 13C12-2-MoCB | 1L | | | M/M+2 | 3.14 | 2.66-3.60 | 107 | 50.0 - 150 |
| 13C12-4-MoCB | 3L | | | M/M+2 | 3.06 | 2.66-3.60 | 98.0 | 50.0 - 150 |
| 13C12-2,2'-DiCB | 4L | | | M/M+2 | 1.54 | 1.33-1.79 | 93.1 | 50.0 - 150 |
| 13C12-4,4'-DiCB | 15L | | | M/M+2 | 1.53 | 1.33-1.79 | 87.3 | 50.0 - 150 |
| 13C12-2,2',6-TriCB | 19L | | | M/M+2 | 1.03 | 0.88-1.20 | 129 | 50.0 - 150 |
| 13C12-3,4,4'-TriCB | 37L | | | M/M+2 | 1.02 | 0.88-1.20 | 57.0 | 50.0 - 150 |
| 13C12-2,2',6,6'-TeCB | 54L | | | M/M+2 | 0.80 | 0.65-0.89 | 86.9 | 50.0 - 150 |
| 13C12-3,3',4,4'-TeCB | 77L | | | M/M+2 | 0.70 | 0.65-0.89 | 69.4 | 50.0 - 150 |
| 13C12-3,4,4',5-TeCB | 81L | | | M/M+2 | 0.65 | 0.65-0.89 | 70.8 | 50.0 - 150 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 77.8 | 50.0 - 150 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | M+2/M+4 | 1.50 | 1.32-1.78 | 68.1 | 50.0 - 150 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | M+2/M+4 | 1.58 | 1.32-1.78 | 60.7 | 50.0 - 150 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | M+2/M+4 | 1.52 | 1.32-1.78 | 63.7 | 50.0 - 150 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | M+2/M+4 | 1.56 | 1.32-1.78 | 66.4 | 50.0 - 150 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | M+2/M+4 | 1.55 | 1.32-1.78 | 78.3 | 50.0 - 150 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | M+2/M+4 | 1.28 | 1.05-1.43 | 72.0 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | M+2/M+4 | 1.26 | 1.05-1.43 | 161 | 100 - 300 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | M+2/M+4 | 1.24 | 1.05-1.43 | 85.7 | 50.0 - 150 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | M+2/M+4 | 1.20 | 1.05-1.43 | 79.2 | 50.0 - 150 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | M+2/M+4 | 1.06 | 0.89-1.21 | 101 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | M+2/M+4 | 0.92 | 0.89-1.21 | 59.4 | 50.0 - 150 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | M+2/M+4 | 0.87 | 0.76-1.02 | 95.8 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | M+2/M+4 | 0.85 | 0.76-1.02 | 96.8 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | M+2/M+4 | 0.77 | 0.65-0.89 | 111 | 50.0 - 150 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | M+2/M+4 | 0.76 | 0.65-0.89 | 113 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | M+4/M+6 | 1.17 | 0.99-1.33 | 121 | 50.0 - 150 |

CLEAN-UP STANDARD

| | | | | | | | | |
|-----------------------------|------|--|--|---------|------|-----------|------|------------|
| 13C12-2,4,4'-TriCB | 28L | | | M/M+2 | 1.01 | 0.88-1.20 | 65.6 | 60.0 - 130 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | M+2/M+4 | 1.61 | 1.32-1.78 | 93.7 | 60.0 - 130 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | M+2/M+4 | 1.05 | 0.89-1.21 | 105 | 60.0 - 130 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

SGS AXYS METHOD MLA-010 Rev 12

Form 6A
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

VER Data Filename: PB9C_027 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 09:39:33

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RETENTION TIME REFERENCE | IUPAC NO. ² | RRT | RRT QC LIMITS |
|-------------------------------|-----------|-------------|-----------------------|---|------------------------|-------|---------------|
| 2-MoCB | 1 | | | 13C12-2-MoCB | 1L | 1.001 | 0.999-1.004 |
| 4-MoCB | 3 | | | 13C12-4-MoCB | 3L | 1.001 | 0.999-1.004 |
| 2,2'-DiCB | 4 | | | 13C12-2,2'-DiCB | 4L | 1.000 | 0.999-1.004 |
| 4,4'-DiCB | 15 | | | 13C12-4,4'-DiCB | 15L | 1.001 | 0.999-1.002 |
| 2,2',6-TriCB | 19 | | | 13C12-2,2',6-TriCB | 19L | 1.001 | 0.999-1.003 |
| 3,4,4'-TriCB | 37 | | | 13C12-3,4,4'-TriCB | 37L | 1.001 | 0.999-1.002 |
| 2,2',6,6'-TeCB | 54 | | | 13C12-2,2',6,6'-TeCB | 54L | 1.002 | 0.999-1.002 |
| 3,3',4,4'-TeCB | 77 | | | 13C12-3,3',4,4'-TeCB | 77L | 1.000 | 1.000-1.001 |
| 3,4,4',5-TeCB | 81 | | | 13C12-3,4,4',5-TeCB | 81L | 1.000 | 1.000-1.001 |
| 2,2',4,6,6'-PeCB | 104 | | | 13C12-2,2',4,6,6'-PeCB | 104L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4'-PeCB | 105 | | | 13C12-2,3,3',4,4'-PeCB | 105L | 1.000 | 1.000-1.001 |
| 2,3,4,4',5-PeCB | 114 | | | 13C12-2,3,4,4',5-PeCB | 114L | 1.000 | 1.000-1.001 |
| 2,3',4,4',5-PeCB | 118 | | | 13C12-2,3',4,4',5-PeCB | 118L | 1.000 | 1.000-1.001 |
| 2',3,4,4',5-PeCB | 123 | | | 13C12-2',3,4,4',5-PeCB | 123L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5-PeCB | 126 | | | 13C12-3,3',4,4',5-PeCB | 126L | 1.000 | 1.000-1.001 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 13C12-2,2',4,4',6,6'-HxCB | 155L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 13C12-2,3,3',4,4',5-HxCB and 13C12-2,3,3',4,4',5'-HxCB | 156L/157L | 1.001 | 0.999-1.003 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 13C12-2,3',4,4',5,5'-HxCB | 167L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 13C12-3,3',4,4',5,5'-HxCB | 169L | 1.000 | 1.000-1.001 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 13C12-2,2',3,4',5,6,6'-HpCB | 188L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 13C12-2,3,3',4,4',5,5'-HpCB | 189L | 1.000 | 1.000-1.001 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | 1.000 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | 1.000 | 1.000-1.001 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) Suffix "L" indicates labeled compound

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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Report Filename: 1668_PCB1668_PB9C_027S1__Form6A_SJ2506283.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 6B
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_027 S: 1

Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL Analysis Time: 09:39:33

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RETENTION TIME REFERENCE | IUPAC NO. ¹ | RRT | RRT QC LIMITS |
|-------------------------------------|------------------------|-------------|-----------------------|--------------------------------|------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 13C12-2,5-DiCB | 9L | 0.719 | 0.688-0.750 |
| 13C12-4-MoCB | 3L | | | 13C12-2,5-DiCB | 9L | 0.857 | 0.826-0.889 |
| 13C12-2,2'-DiCB | 4L | | | 13C12-2,5-DiCB | 9L | 0.874 | 0.843-0.905 |
| 13C12-4,4'-DiCB | 15L | | | 13C12-2,5-DiCB | 9L | 1.253 | 1.222-1.284 |
| 13C12-2,2',6-TriCB | 19L | | | 13C12-2,5-DiCB | 9L | 1.073 | 1.042-1.104 |
| 13C12-3,4,4'-TriCB | 37L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.090 | 1.070-1.110 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.810 | 0.797-0.824 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.395 | 1.382-1.408 |
| 13C12-3,4,4',5-TeCB | 81L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.371 | 1.358-1.385 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 0.808 | 0.798-0.818 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.199 | 1.188-1.209 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.178 | 1.168-1.188 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.161 | 1.151-1.171 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.150 | 1.140-1.160 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.299 | 1.288-1.309 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 0.787 | 0.779-0.795 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.107 | 1.099-1.115 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.078 | 1.070-1.086 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.191 | 1.182-1.199 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.713 | 0.707-0.719 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.959 | 0.953-0.965 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.818 | 0.811-0.824 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.009 | 1.000-1.019 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.043 | 1.034-1.053 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.949 | 0.943-0.955 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.075 | 1.066-1.085 |

CLEANUP STANDARD

| | | | | | | | |
|-----------------------------|------|--|--|---------------------------|------|-------|-------------|
| 13C12-2,4,4'-TriCB | 28L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.924 | 0.911-0.938 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.087 | 1.077-1.097 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.012 | 1.004-1.020 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

PCB CONGENER INITIAL CALIBRATION RELATIVE RESPONSES,
ION ABUNDANCE RATIOS, AND RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

CAL Data Filename: PB9C_027 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 09:39:33

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|----------------|-----------|--------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 3-MoCB | 2 | | | 0.83 | M/M+2 | 2.98 | 2.66-3.60 | 0.988 | 0.984 - 0.992 |
| 2,3-DiCB | 5 | | | 0.86 | M/M+2 | 1.51 | 1.33-1.79 | 1.198 | 1.194 - 1.201 |
| 2,3'-DiCB | 6 | | | 0.94 | M/M+2 | 1.49 | 1.33-1.79 | 1.175 | 1.171 - 1.179 |
| 2,4-DiCB | 7 | | | 0.92 | M/M+2 | 1.49 | 1.33-1.79 | 1.158 | 1.155 - 1.162 |
| 2,4'-DiCB | 8 | | | 1.03 | M/M+2 | 1.51 | 1.33-1.79 | 1.206 | 1.202 - 1.210 |
| 2,5-DiCB | 9 | | | 0.98 | M/M+2 | 1.50 | 1.33-1.79 | 1.145 | 1.142 - 1.149 |
| 2,6-DiCB | 10 | | | 0.96 | M/M+2 | 1.46 | 1.33-1.79 | 1.013 | 1.010 - 1.017 |
| 3,3'-DiCB | 11 | | | 0.85 | M/M+2 | 1.51 | 1.33-1.79 | 0.969 | 0.967 - 0.972 |
| 3,4-DiCB | 12 | 12 + 13 | C | 0.89 | M/M+2 | 1.47 | 1.33-1.79 | 0.985 | 0.983 - 0.988 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | | | |
| 3,5-DiCB | 14 | | | 0.92 | M/M+2 | 1.46 | 1.33-1.79 | 0.926 | 0.924 - 0.929 |
| 2,2',3-TriCB | 16 | | | 0.89 | M/M+2 | 1.05 | 0.88-1.20 | 1.166 | 1.163 - 1.169 |
| 2,2',4-TriCB | 17 | | | 1.10 | M/M+2 | 1.05 | 0.88-1.20 | 1.139 | 1.136 - 1.142 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 1.32 | M/M+2 | 1.07 | 0.88-1.20 | 1.113 | 1.110 - 1.115 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 0.86 | M/M+2 | 0.97 | 0.88-1.20 | 0.849 | 0.846 - 0.852 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 0.88 | M/M+2 | 0.98 | 0.88-1.20 | 0.856 | 0.853 - 0.859 |
| 2,3,4'-TriCB | 22 | | | 0.76 | M/M+2 | 0.98 | 0.88-1.20 | 0.872 | 0.870 - 0.873 |
| 2,3,5-TriCB | 23 | | | 0.78 | M/M+2 | 0.98 | 0.88-1.20 | 1.284 | 1.281 - 1.287 |
| 2,3,6-TriCB | 24 | | | 1.54 | M/M+2 | 1.05 | 0.88-1.20 | 1.159 | 1.156 - 1.162 |
| 2,3',4-TriCB | 25 | | | 1.00 | M/M+2 | 0.99 | 0.88-1.20 | 0.826 | 0.824 - 0.828 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 0.84 | M/M+2 | 1.00 | 0.88-1.20 | 1.303 | 1.298 - 1.307 |
| 2,3',6-TriCB | 27 | | | 1.53 | M/M+2 | 1.04 | 0.88-1.20 | 1.151 | 1.148 - 1.154 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | | | |
| 2,4',5-TriCB | 31 | | | 0.91 | M/M+2 | 1.00 | 0.88-1.20 | 0.837 | 0.835 - 0.839 |
| 2,4',6-TriCB | 32 | | | 0.87 | M/M+2 | 0.95 | 0.88-1.20 | 1.197 | 1.194 - 1.200 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | | | |
| 2',3,5-TriCB | 34 | | | 0.80 | M/M+2 | 0.97 | 0.88-1.20 | 1.274 | 1.272 - 1.277 |
| 3,3',4-TriCB | 35 | | | 0.79 | M/M+2 | 1.01 | 0.88-1.20 | 0.985 | 0.984 - 0.987 |
| 3,3',5-TriCB | 36 | | | 0.83 | M/M+2 | 0.96 | 0.88-1.20 | 0.932 | 0.931 - 0.934 |
| 3,4,5-TriCB | 38 | | | 0.88 | M/M+2 | 0.98 | 0.88-1.20 | 0.968 | 0.966 - 0.970 |
| 3,4',5-TriCB | 39 | | | 0.86 | M/M+2 | 0.98 | 0.88-1.20 | 0.946 | 0.944 - 0.948 |
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 0.87 | M/M+2 | 0.78 | 0.65-0.89 | 1.336 | 1.332 - 1.340 |
| 2,2',3,4-TeCB | 41 | 40 + 41 + 71 | C40 | | | | | | |
| 2,2',3,4'-TeCB | 42 | | | 0.80 | M/M+2 | 0.77 | 0.65-0.89 | 1.313 | 1.310 - 1.315 |
| 2,2',3,5-TeCB | 43 | | | 0.73 | M/M+2 | 0.78 | 0.65-0.89 | 1.247 | 1.245 - 1.250 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 0.94 | M/M+2 | 0.78 | 0.65-0.89 | 1.287 | 1.283 - 1.291 |
| 2,2',3,6-TeCB | 45 | 45 + 51 | C | 0.84 | M/M+2 | 0.77 | 0.65-0.89 | 1.148 | 1.144 - 1.152 |
| 2,2',3,6'-TeCB | 46 | | | 0.75 | M/M+2 | 0.76 | 0.65-0.89 | 1.161 | 1.159 - 1.164 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | | | |
| 2,2',4,5-TeCB | 48 | | | 0.82 | M/M+2 | 0.77 | 0.65-0.89 | 1.275 | 1.273 - 1.277 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 0.98 | M/M+2 | 0.78 | 0.65-0.89 | 1.259 | 1.255 - 1.263 |
| 2,2',4,6-TeCB | 50 | 50 + 53 | C | 0.88 | M/M+2 | 0.78 | 0.65-0.89 | 1.113 | 1.109 - 1.117 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | | | |
| 2,2',5,5'-TeCB | 52 | | | 0.90 | M/M+2 | 0.79 | 0.65-0.89 | 1.235 | 1.232 - 1.237 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | | | |
| 2,3,3',4-TeCB | 55 | | | 0.67 | M/M+2 | 0.80 | 0.65-0.89 | 0.890 | 0.888 - 0.891 |
| 2,3,3',4'-TeCB | 56 | | | 0.71 | M/M+2 | 0.77 | 0.65-0.89 | 0.905 | 0.903 - 0.906 |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|-------------------|-----------|--------------------------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 2,3,3',5'-TeCB | 57 | | | 0.75 | M/M+2 | 0.76 | 0.65-0.89 | 0.845 | 0.843 - 0.846 |
| 2,3,3',5'-TeCB | 58 | | | 0.72 | M/M+2 | 0.74 | 0.65-0.89 | 0.852 | 0.850 - 0.853 |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 1.17 | M/M+2 | 0.78 | 0.65-0.89 | 1.303 | 1.299 - 1.307 |
| 2,3,4,4'-TeCB | 60 | | | 0.74 | M/M+2 | 0.78 | 0.65-0.89 | 0.912 | 0.910 - 0.913 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 0.76 | M/M+2 | 0.76 | 0.65-0.89 | 0.875 | 0.872 - 0.878 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | | | |
| 2,3,4',5'-TeCB | 63 | | | 0.76 | M/M+2 | 0.81 | 0.65-0.89 | 0.865 | 0.863 - 0.866 |
| 2,3,4',6'-TeCB | 64 | | | 1.19 | M/M+2 | 0.77 | 0.65-0.89 | 1.349 | 1.347 - 1.352 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | | | |
| 2,3',4,4'-TeCB | 66 | | | 0.75 | M/M+2 | 0.80 | 0.65-0.89 | 0.884 | 0.883 - 0.886 |
| 2,3',4,5'-TeCB | 67 | | | 0.89 | M/M+2 | 0.75 | 0.65-0.89 | 0.857 | 0.855 - 0.858 |
| 2,3',4,5'-TeCB | 68 | | | 0.79 | M/M+2 | 0.78 | 0.65-0.89 | 0.832 | 0.831 - 0.834 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | | | |
| 2,3',5,5'-TeCB | 72 | | | 0.76 | M/M+2 | 0.76 | 0.65-0.89 | 0.823 | 0.822 - 0.824 |
| 2,3',5,6'-TeCB | 73 | | | 1.07 | M/M+2 | 0.75 | 0.65-0.89 | 1.242 | 1.240 - 1.245 |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 3,3',4,5'-TeCB | 78 | | | 0.79 | M/M+2 | 0.74 | 0.65-0.89 | 0.987 | 0.986 - 0.989 |
| 3,3',4,5'-TeCB | 79 | | | 0.96 | M/M+2 | 0.75 | 0.65-0.89 | 0.971 | 0.969 - 0.972 |
| 3,3',5,5'-TeCB | 80 | | | 0.83 | M/M+2 | 0.75 | 0.65-0.89 | 0.925 | 0.923 - 0.926 |
| 2,2',3,3',4'-PeCB | 82 | | | 0.89 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.934 | 0.932 - 0.935 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 0.89 | M+2/M+4 | 1.59 | 1.32-1.78 | 0.885 | 0.882 - 0.887 |
| 2,2',3,3',6'-PeCB | 84 | | | 0.78 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.162 | 1.160 - 1.164 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 1.19 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.919 | 0.917 - 0.922 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C | 1.08 | M+2/M+4 | 1.58 | 1.32-1.78 | 0.900 | 0.897 - 0.904 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 0.89 | M+2/M+4 | 1.61 | 1.32-1.78 | 1.153 | 1.149 - 1.156 |
| 2,2',3,4,6'-PeCB | 89 | | | 0.85 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.182 | 1.180 - 1.184 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 1.07 | M+2/M+4 | 1.56 | 1.32-1.78 | 0.869 | 0.867 - 0.871 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 0.92 | M+2/M+4 | 1.56 | 1.32-1.78 | 0.853 | 0.852 - 0.855 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 0.91 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.130 | 1.119 - 1.140 |
| 2,2',3,5,6'-PeCB | 94 | | | 0.81 | M+2/M+4 | 1.59 | 1.32-1.78 | 1.102 | 1.100 - 1.104 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',3,6,6'-PeCB | 96 | | | 1.21 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.015 | 1.012 - 1.019 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | | | |
| 2,2',4,4',6'-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,5',6'-PeCB | 103 | | | 0.99 | M+2/M+4 | 1.59 | 1.32-1.78 | 1.094 | 1.092 - 1.096 |
| 2,3,3',4,5'-PeCB | 106 | | | 0.87 | M+2/M+4 | 1.51 | 1.32-1.78 | 1.004 | 1.002 - 1.005 |
| 2,3,3',4',5'-PeCB | 107 | 107 + 124 | C | 0.86 | M+2/M+4 | 1.47 | 1.32-1.78 | 0.991 | 0.988 - 0.993 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,3,3',4,6'-PeCB | 109 | | | 0.94 | M+2/M+4 | 1.48 | 1.32-1.78 | 0.997 | 0.996 - 0.999 |
| 2,3,3',4',6'-PeCB | 110 | 110 + 115 | C | 1.35 | M+2/M+4 | 1.56 | 1.32-1.78 | 0.926 | 0.924 - 0.929 |
| 2,3,3',5,5'-PeCB | 111 | | | 1.33 | M+2/M+4 | 1.58 | 1.32-1.78 | 0.946 | 0.944 - 0.947 |
| 2,3,3',5,6'-PeCB | 112 | | | 1.28 | M+2/M+4 | 1.55 | 1.32-1.78 | 0.889 | 0.888 - 0.891 |
| 2,3,3',5',6'-PeCB | 113 | 90 + 101 + 113 | C90 | | | | | | |
| 2,3,4,4',6'-PeCB | 115 | 110 + 115 | C110 | | | | | | |
| 2,3,4,5,6'-PeCB | 116 | 85 + 116 + 117 | C85 | | | | | | |
| 2,3,4',5,6'-PeCB | 117 | 85 + 116 + 117 | C85 | | | | | | |
| 2,3',4,4',6'-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,3',4,5,5'-PeCB | 120 | | | 1.44 | M+2/M+4 | 1.55 | 1.32-1.78 | 0.958 | 0.957 - 0.960 |
| 2,3',4,5',6'-PeCB | 121 | | | 1.21 | M+2/M+4 | 1.56 | 1.32-1.78 | 1.200 | 1.198 - 1.202 |
| 2',3,3',4,5'-PeCB | 122 | | | 0.73 | M+2/M+4 | 1.48 | 1.32-1.78 | 1.010 | 1.008 - 1.011 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|--------------------------|-----------|-----------------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 3,3',4,5,5'-PeCB | 127 | | | 1.00 | M+2/M+4 | 1.49 | 1.32-1.78 | 1.040 | 1.039 - 1.042 |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 1.04 | M+2/M+4 | 1.24 | 1.05-1.43 | 0.958 | 0.956 - 0.960 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 0.93 | M+2/M+4 | 1.22 | 1.05-1.43 | 0.930 | 0.927 - 0.932 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 0.78 | M+2/M+4 | 1.23 | 1.05-1.43 | 0.913 | 0.912 - 0.914 |
| 2,2',3,3',4,6-HxCB | 131 | | | 0.71 | M+2/M+4 | 1.28 | 1.05-1.43 | 1.158 | 1.156 - 1.159 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 0.65 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.173 | 1.170 - 1.176 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 0.71 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.190 | 1.188 - 1.191 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 0.72 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.140 | 1.137 - 1.142 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 0.77 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.105 | 1.100 - 1.111 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 0.92 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.023 | 1.021 - 1.024 |
| 2,2',3,4,4',5-HxCB | 137 | | | 0.83 | M+2/M+4 | 1.22 | 1.05-1.43 | 0.918 | 0.917 - 0.920 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C | 0.80 | M+2/M+4 | 1.25 | 1.05-1.43 | 1.152 | 1.149 - 1.154 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 0.83 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.904 | 0.903 - 0.905 |
| 2,2',3,4,5,6-HxCB | 142 | | | 0.68 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.163 | 1.161 - 1.164 |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 0.77 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.121 | 1.119 - 1.122 |
| 2,2',3,4,6,6'-HxCB | 145 | | | 0.84 | M+2/M+4 | 1.29 | 1.05-1.43 | 1.033 | 1.031 - 1.034 |
| 2,2',3,4',5,5'-HxCB | 146 | | | 0.89 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.884 | 0.883 - 0.885 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 0.80 | M+2/M+4 | 1.25 | 1.05-1.43 | 1.132 | 1.130 - 1.135 |
| 2,2',3,4',5,6'-HxCB | 148 | | | 0.74 | M+2/M+4 | 1.28 | 1.05-1.43 | 1.083 | 1.082 - 1.085 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | | 0.89 | M+2/M+4 | 1.22 | 1.05-1.43 | 1.011 | 1.010 - 1.013 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | | 0.99 | M+2/M+4 | 1.27 | 1.05-1.43 | 1.006 | 1.005 - 1.008 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 1.04 | M+2/M+4 | 1.26 | 1.05-1.43 | 0.900 | 0.898 - 0.902 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 1.24 | M+2/M+4 | 1.23 | 1.05-1.43 | 0.938 | 0.937 - 0.939 |
| 2,3,3',4,5,5'-HxCB | 159 | | | 1.25 | M+2/M+4 | 1.26 | 1.05-1.43 | 0.982 | 0.981 - 0.983 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,3,3',4,5',6-HxCB | 161 | | | 1.16 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.888 | 0.887 - 0.889 |
| 2,3,3',4',5,5'-HxCB | 162 | | | 1.19 | M+2/M+4 | 1.23 | 1.05-1.43 | 0.989 | 0.988 - 0.991 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 1.19 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.921 | 0.920 - 0.922 |
| 2,3,3',5,5',6-HxCB | 165 | | | 0.93 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.878 | 0.877 - 0.879 |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | | | |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 1.13 | M+2/M+4 | 1.06 | 0.89-1.21 | 1.000 | 0.999 - 1.001 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 1.00 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.161 | 1.159 - 1.163 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 0.93 | M+2/M+4 | 1.04 | 0.89-1.21 | 0.897 | 0.896 - 0.898 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 1.20 | M+2/M+4 | 1.07 | 0.89-1.21 | 1.132 | 1.131 - 1.134 |
| 2,2',3,3',4,5',6-HpCB | 175 | | | 1.20 | M+2/M+4 | 1.01 | 0.89-1.21 | 1.102 | 1.100 - 1.103 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 1.34 | M+2/M+4 | 1.06 | 0.89-1.21 | 1.033 | 1.032 - 1.035 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 1.10 | M+2/M+4 | 1.03 | 0.89-1.21 | 1.144 | 1.143 - 1.146 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 1.04 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.085 | 1.083 - 1.086 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 1.27 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.009 | 1.008 - 1.011 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 1.00 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.000 | 0.999 - 1.001 |
| 2,2',3,4,4',5,6-HpCB | 181 | | | 1.05 | M+2/M+4 | 1.01 | 0.89-1.21 | 1.155 | 1.154 - 1.157 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | | 1.20 | M+2/M+4 | 1.06 | 0.89-1.21 | 1.115 | 1.114 - 1.116 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 1.18 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.127 | 1.126 - 1.129 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | | 1.40 | M+2/M+4 | 1.08 | 0.89-1.21 | 1.025 | 1.023 - 1.026 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | | 1.31 | M+2/M+4 | 1.06 | 0.89-1.21 | 1.046 | 1.045 - 1.047 |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 1.23 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.109 | 1.108 - 1.110 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 1.32 | M+2/M+4 | 1.02 | 0.89-1.21 | 0.947 | 0.946 - 0.948 |
| 2,3,3',4,4',5',6-HpCB | 191 | | | 1.28 | M+2/M+4 | 1.06 | 0.89-1.21 | 0.918 | 0.917 - 0.919 |
| 2,3,3',4,5,5',6-HpCB | 192 | | | 1.16 | M+2/M+4 | 1.03 | 0.89-1.21 | 0.903 | 0.902 - 0.904 |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | | | |
| 2,2',3,3',4,4',5,5'-OcCB | 194 | | | 0.80 | M+2/M+4 | 0.88 | 0.76-1.02 | 0.991 | 0.990 - 0.992 |
| 2,2',3,3',4,4',5,6-OcCB | 195 | | | 0.71 | M+2/M+4 | 0.87 | 0.76-1.02 | 0.945 | 0.944 - 0.946 |
| 2,2',3,3',4,4',5,6'-OcCB | 196 | | | 0.74 | M+2/M+4 | 0.91 | 0.76-1.02 | 0.916 | 0.915 - 0.917 |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|-----------------------------------|-----------|-------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 2,2',3,3',4,4',6,6'-OcCB | 197 | 197 + 200 | C | 0.98 | M+2/M+4 | 0.90 | 0.76-1.02 | 1.046 | 1.043 - 1.048 |
| 2,2',3,3',4,5,5',6-OcCB | 198 | 198 + 199 | C | 0.75 | M+2/M+4 | 0.90 | 0.76-1.02 | 1.114 | 1.112 - 1.116 |
| 2,2',3,3',4,5,5',6'-OcCB | 199 | 198 + 199 | C198 | | | | | | |
| 2,2',3,3',4,5,6,6'-OcCB | 200 | 197 + 200 | C197 | | | | | | |
| 2,2',3,3',4,5',6,6'-OcCB | 201 | | | 1.01 | M+2/M+4 | 0.88 | 0.76-1.02 | 1.023 | 1.021 - 1.025 |
| 2,2',3,4,4',5,5',6-OcCB | 203 | | | 0.80 | M+2/M+4 | 0.91 | 0.76-1.02 | 0.920 | 0.919 - 0.921 |
| 2,2',3,4,4',5,6,6'-OcCB | 204 | | | 0.96 | M+2/M+4 | 0.92 | 0.76-1.02 | 1.039 | 1.038 - 1.040 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | | 1.19 | M+2/M+4 | 0.78 | 0.65-0.89 | 1.020 | 1.019 - 1.021 |

(1) Where applicable, custom lab flags have been used on this report.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 3B

PCB CONGENER INITIAL CALIBRATION RELATIVE RESPONSES,
ION ABUNDANCE RATIOS, AND RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

CAL Data Filename: PB9C_027 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 09:39:33

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RRF | MZ's FORMING RATIO ³ | ION ABUND. RATIO | RATIO QC LIMITS ⁴ | RRT | RRT QC LIMITS |
|----------------------------------|------------------------|-------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 1.24 | M/M+2 | 3.14 | 2.66-3.60 | 0.719 | 0.703 - 0.735 |
| 13C12-4-MoCB | 3L | | | 1.11 | M/M+2 | 3.06 | 2.66-3.60 | 0.857 | 0.842 - 0.873 |
| 13C12-2,2'-DiCB | 4L | | | 0.64 | M/M+2 | 1.54 | 1.33-1.79 | 0.874 | 0.858 - 0.890 |
| 13C12-4,4'-DiCB | 15L | | | 0.95 | M/M+2 | 1.53 | 1.33-1.79 | 1.253 | 1.237 - 1.268 |
| 13C12-2,2',6-TriCB | 19L | | | 0.64 | M/M+2 | 1.03 | 0.88-1.20 | 1.073 | 1.057 - 1.088 |
| 13C12-3,4,4'-TriCB | 37L | | | 1.17 | M/M+2 | 1.02 | 0.88-1.20 | 1.090 | 1.080 - 1.100 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 1.41 | M/M+2 | 0.80 | 0.65-0.89 | 0.810 | 0.804 - 0.817 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 1.11 | M/M+2 | 0.70 | 0.65-0.89 | 1.395 | 1.389 - 1.402 |
| 13C12-3,4,4',5-TeCB | 81L | | | 1.13 | M/M+2 | 0.65 | 0.65-0.89 | 1.371 | 1.365 - 1.378 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 1.11 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.808 | 0.803 - 0.813 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 1.11 | M+2/M+4 | 1.50 | 1.32-1.78 | 1.199 | 1.193 - 1.204 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 1.02 | M+2/M+4 | 1.58 | 1.32-1.78 | 1.178 | 1.173 - 1.183 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 1.02 | M+2/M+4 | 1.52 | 1.32-1.78 | 1.161 | 1.156 - 1.166 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 1.07 | M+2/M+4 | 1.56 | 1.32-1.78 | 1.150 | 1.145 - 1.155 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 1.22 | M+2/M+4 | 1.55 | 1.32-1.78 | 1.299 | 1.293 - 1.304 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 0.99 | M+2/M+4 | 1.28 | 1.05-1.43 | 0.787 | 0.783 - 0.791 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 1.19 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.107 | 1.103 - 1.111 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 1.20 | M+2/M+4 | 1.24 | 1.05-1.43 | 1.078 | 1.074 - 1.082 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 1.20 | M+2/M+4 | 1.20 | 1.05-1.43 | 1.191 | 1.186 - 1.195 |
| 13C12-2,2',3,3',4,4',5-HpCB | 170L | | | 1.01 | M+2/M+4 | 1.06 | 0.89-1.21 | 0.897 | 0.894 - 0.900 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 1.23 | M+2/M+4 | 1.06 | 0.89-1.21 | 0.873 | 0.870 - 0.876 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 1.52 | M+2/M+4 | 1.06 | 0.89-1.21 | 0.713 | 0.710 - 0.716 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 0.92 | M+2/M+4 | 0.92 | 0.89-1.21 | 0.959 | 0.956 - 0.962 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 1.42 | M+2/M+4 | 0.87 | 0.76-1.02 | 0.818 | 0.815 - 0.821 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 1.39 | M+2/M+4 | 0.85 | 0.76-1.02 | 1.009 | 1.005 - 1.014 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 1.07 | M+2/M+4 | 0.77 | 0.65-0.89 | 1.043 | 1.039 - 1.048 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 1.36 | M+2/M+4 | 0.76 | 0.65-0.89 | 0.949 | 0.946 - 0.952 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 4A
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_027 S: 9
Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
GC Column ID: SPB OCTYL Analysis Time: 18:16:03

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | MZ's FORMING RATIO ² | ION ABUND. RATIO | QC LIMITS ³ | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------|-----------|-------------|-----------------------|---------------------------------|------------------|------------------------|---------------------|---------------------|
| 2-MoCB | 1 | | | M/M+2 | 2.98 | 2.66-3.60 | 19.0 | 17.5 - 32.5 |
| 4-MoCB | 3 | | | M/M+2 | 2.99 | 2.66-3.60 | 19.2 | 17.5 - 32.5 |
| 2,2'-DiCB | 4 | | | M/M+2 | 1.49 | 1.33-1.79 | 18.7 | 17.5 - 32.5 |
| 4,4'-DiCB | 15 | | | M/M+2 | 1.50 | 1.33-1.79 | 21.7 | 19.6 - 36.4 |
| 2,2',6-TriCB | 19 | | | M/M+2 | 1.05 | 0.88-1.20 | 24.3 | 17.5 - 32.5 |
| 3,4,4'-TriCB | 37 | | | M/M+2 | 0.98 | 0.88-1.20 | 18.8 | 17.5 - 32.5 |
| 2,2',6,6'-TeCB | 54 | | | M/M+2 | 0.78 | 0.65-0.89 | 46.4 | 35.0 - 65.0 |
| 3,3',4,4'-TeCB | 77 | | | M/M+2 | 0.76 | 0.65-0.89 | 39.8 | 35.0 - 65.0 |
| 3,4,4',5-TeCB | 81 | | | M/M+2 | 0.79 | 0.65-0.89 | 41.5 | 35.0 - 65.0 |
| 2,2',4,6,6'-PeCB | 104 | | | M+2/M+4 | 1.54 | 1.32-1.78 | 51.9 | 35.0 - 65.0 |
| 2,3,3',4,4'-PeCB | 105 | | | M+2/M+4 | 1.47 | 1.32-1.78 | 41.0 | 35.0 - 65.0 |
| 2,3,4,4',5-PeCB | 114 | | | M+2/M+4 | 1.51 | 1.32-1.78 | 40.4 | 35.0 - 65.0 |
| 2,3',4,4',5-PeCB | 118 | | | M+2/M+4 | 1.49 | 1.32-1.78 | 40.5 | 35.0 - 65.0 |
| 2',3,4,4',5-PeCB | 123 | | | M+2/M+4 | 1.52 | 1.32-1.78 | 45.0 | 35.0 - 65.0 |
| 3,3',4,4',5-PeCB | 126 | | | M+2/M+4 | 1.47 | 1.32-1.78 | 42.2 | 39.0 - 72.4 |
| 2,2',4,4',6,6'-HxCB | 155 | | | M+2/M+4 | 1.27 | 1.05-1.43 | 52.0 | 35.0 - 65.0 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | M+2/M+4 | 1.25 | 1.05-1.43 | 90.8 | 70.0 - 130 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | M+2/M+4 | 1.26 | 1.05-1.43 | 50.2 | 35.0 - 65.0 |
| 3,3',4,4',5,5'-HxCB | 169 | | | M+2/M+4 | 1.32 | 1.05-1.43 | 48.1 | 35.0 - 65.0 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | M+2/M+4 | 1.06 | 0.89-1.21 | 49.1 | 35.0 - 65.0 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | M+2/M+4 | 0.98 | 0.89-1.21 | 42.1 | 35.0 - 65.0 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | | M+2/M+4 | 0.92 | 0.76-1.02 | 81.1 | 58.9 - 110 |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | | M+2/M+4 | 0.88 | 0.76-1.02 | 70.0 | 52.5 - 97.5 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | M+2/M+4 | 0.79 | 0.65-0.89 | 74.3 | 52.5 - 97.5 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | M+2/M+4 | 0.79 | 0.65-0.89 | 81.7 | 58.7 - 109 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | M+4/M+6 | 1.18 | 0.99-1.33 | 72.2 | 52.5 - 97.5 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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Report Filename: 1668_PCB1668_PB9C_027S9_Form4A_SJ2506299.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 4B
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_027 S: 9
Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
GC Column ID: SPB OCTYL Analysis Time: 18:16:03

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG 2 | MZ's FORMING RATIO 3 | ION ABUND. RATIO | QC LIMITS 4 | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------------|-------------|-------------|------------|----------------------|------------------|-------------|---------------------|---------------------|
| 13C12-2-MoCB | 1L | | | M/M+2 | 3.06 | 2.66-3.60 | 105 | 50.0 - 150 |
| 13C12-4-MoCB | 3L | | | M/M+2 | 3.03 | 2.66-3.60 | 98.2 | 50.0 - 150 |
| 13C12-2,2'-DiCB | 4L | | | M/M+2 | 1.53 | 1.33-1.79 | 91.3 | 50.0 - 150 |
| 13C12-4,4'-DiCB | 15L | | | M/M+2 | 1.52 | 1.33-1.79 | 87.6 | 50.0 - 150 |
| 13C12-2,2',6-TriCB | 19L | | | M/M+2 | 1.04 | 0.88-1.20 | 130 | 50.0 - 150 |
| 13C12-3,4,4'-TriCB | 37L | | | M/M+2 | 1.01 | 0.88-1.20 | 57.2 | 50.0 - 150 |
| 13C12-2,2',6,6'-TeCB | 54L | | | M/M+2 | 0.80 | 0.65-0.89 | 88.6 | 50.0 - 150 |
| 13C12-3,3',4,4'-TeCB | 77L | | | M/M+2 | 0.71 | 0.65-0.89 | 62.5 | 50.0 - 150 |
| 13C12-3,4,4',5-TeCB | 81L | | | M/M+2 | 0.71 | 0.65-0.89 | 66.1 | 50.0 - 150 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | M+2/M+4 | 1.51 | 1.32-1.78 | 82.4 | 50.0 - 150 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | M+2/M+4 | 1.54 | 1.32-1.78 | 57.4 | 50.0 - 150 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | M+2/M+4 | 1.55 | 1.32-1.78 | 57.0 | 50.0 - 150 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | M+2/M+4 | 1.51 | 1.32-1.78 | 56.5 | 50.0 - 150 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 58.4 | 50.0 - 150 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | M+2/M+4 | 1.50 | 1.32-1.78 | 57.2 | 50.0 - 150 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | M+2/M+4 | 1.27 | 1.05-1.43 | 93.5 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | M+2/M+4 | 1.27 | 1.05-1.43 | 176 | 100 - 300 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | M+2/M+4 | 1.27 | 1.05-1.43 | 88.6 | 50.0 - 150 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | M+2/M+4 | 1.23 | 1.05-1.43 | 93.3 | 50.0 - 150 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | M+2/M+4 | 1.04 | 0.89-1.21 | 95.1 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | M+2/M+4 | 0.95 | 0.89-1.21 | 61.0 | 50.0 - 150 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | M+2/M+4 | 0.91 | 0.76-1.02 | 87.8 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | M+2/M+4 | 0.83 | 0.76-1.02 | 97.0 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | M+2/M+4 | 0.76 | 0.65-0.89 | 115 | 50.0 - 150 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | M+2/M+4 | 0.78 | 0.65-0.89 | 108 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | M+4/M+6 | 1.18 | 0.99-1.33 | 129 | 50.0 - 150 |

CLEAN-UP STANDARD

| | | | | | | | | |
|-----------------------------|------|--|--|---------|------|-----------|------|------------|
| 13C12-2,4,4'-TriCB | 28L | | | M/M+2 | 1.01 | 0.88-1.20 | 64.6 | 60.0 - 130 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | M+2/M+4 | 1.60 | 1.32-1.78 | 87.2 | 60.0 - 130 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | M+2/M+4 | 1.02 | 0.89-1.21 | 104 | 60.0 - 130 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

SGS AXYS METHOD MLA-010 Rev 12

Form 6A
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

VER Data Filename: PB9C_027 S: 9

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 18:16:03

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RETENTION TIME REFERENCE | IUPAC NO. ² | RRT | RRT QC LIMITS |
|-------------------------------|-----------|-------------|-----------------------|---|------------------------|-------|---------------|
| 2-MoCB | 1 | | | 13C12-2-MoCB | 1L | 1.003 | 0.999-1.004 |
| 4-MoCB | 3 | | | 13C12-4-MoCB | 3L | 1.002 | 0.999-1.004 |
| 2,2'-DiCB | 4 | | | 13C12-2,2'-DiCB | 4L | 1.001 | 0.999-1.004 |
| 4,4'-DiCB | 15 | | | 13C12-4,4'-DiCB | 15L | 1.000 | 0.999-1.002 |
| 2,2',6-TriCB | 19 | | | 13C12-2,2',6-TriCB | 19L | 1.001 | 0.999-1.003 |
| 3,4,4'-TriCB | 37 | | | 13C12-3,4,4'-TriCB | 37L | 1.001 | 0.999-1.002 |
| 2,2',6,6'-TeCB | 54 | | | 13C12-2,2',6,6'-TeCB | 54L | 1.001 | 0.999-1.002 |
| 3,3',4,4'-TeCB | 77 | | | 13C12-3,3',4,4'-TeCB | 77L | 1.000 | 1.000-1.001 |
| 3,4,4',5-TeCB | 81 | | | 13C12-3,4,4',5-TeCB | 81L | 1.000 | 1.000-1.001 |
| 2,2',4,6,6'-PeCB | 104 | | | 13C12-2,2',4,6,6'-PeCB | 104L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4'-PeCB | 105 | | | 13C12-2,3,3',4,4'-PeCB | 105L | 1.000 | 1.000-1.001 |
| 2,3,4,4',5-PeCB | 114 | | | 13C12-2,3,4,4',5-PeCB | 114L | 1.001 | 1.000-1.001 |
| 2,3',4,4',5-PeCB | 118 | | | 13C12-2,3',4,4',5-PeCB | 118L | 1.001 | 1.000-1.001 |
| 2',3,4,4',5-PeCB | 123 | | | 13C12-2',3,4,4',5-PeCB | 123L | 1.001 | 1.000-1.001 |
| 3,3',4,4',5-PeCB | 126 | | | 13C12-3,3',4,4',5-PeCB | 126L | 1.000 | 1.000-1.001 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 13C12-2,2',4,4',6,6'-HxCB | 155L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 13C12-2,3,3',4,4',5-HxCB and 13C12-2,3,3',4,4',5'-HxCB | 156L/157L | 1.000 | 0.998-1.003 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 13C12-2,3',4,4',5,5'-HxCB | 167L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 13C12-3,3',4,4',5,5'-HxCB | 169L | 1.000 | 1.000-1.001 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 13C12-2,2',3,4',5,6,6'-HpCB | 188L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 13C12-2,3,3',4,4',5,5'-HpCB | 189L | 1.000 | 1.000-1.001 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | | 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | | 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | 1.000 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | 1.000 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | 1.001 | 1.000-1.001 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) Suffix "L" indicates labeled compound

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 6B
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

VER Data Filename: PB9C_027 S: 9

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 18:16:03

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RETENTION TIME REFERENCE | IUPAC NO. ¹ | RRT | RRT QC LIMITS |
|-------------------------------------|------------------------|-------------|-----------------------|--------------------------------|------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 13C12-2,5-DiCB | 9L | 0.717 | 0.686-0.749 |
| 13C12-4-MoCB | 3L | | | 13C12-2,5-DiCB | 9L | 0.856 | 0.825-0.887 |
| 13C12-2,2'-DiCB | 4L | | | 13C12-2,5-DiCB | 9L | 0.873 | 0.842-0.904 |
| 13C12-4,4'-DiCB | 15L | | | 13C12-2,5-DiCB | 9L | 1.253 | 1.222-1.285 |
| 13C12-2,2',6-TriCB | 19L | | | 13C12-2,5-DiCB | 9L | 1.072 | 1.041-1.103 |
| 13C12-3,4,4'-TriCB | 37L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.090 | 1.070-1.110 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.809 | 0.796-0.823 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.395 | 1.382-1.408 |
| 13C12-3,4,4',5-TeCB | 81L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.372 | 1.359-1.385 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 0.808 | 0.798-0.818 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.199 | 1.189-1.209 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.178 | 1.168-1.189 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.161 | 1.151-1.171 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.150 | 1.140-1.161 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.300 | 1.289-1.310 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 0.787 | 0.779-0.795 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.107 | 1.099-1.115 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.078 | 1.070-1.086 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.191 | 1.182-1.199 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.713 | 0.706-0.719 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.959 | 0.953-0.965 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.818 | 0.811-0.824 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.009 | 1.000-1.019 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.043 | 1.034-1.053 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-NoCB | 208L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.949 | 0.943-0.956 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.075 | 1.066-1.085 |

CLEANUP STANDARD

| | | | | | | | |
|-----------------------------|------|--|--|---------------------------|------|-------|-------------|
| 13C12-2,4,4'-TriCB | 28L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.924 | 0.910-0.937 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.087 | 1.077-1.098 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.012 | 1.004-1.020 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

SGS AXYS METHOD MLA-010 Rev 12

Form 4A
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 1
 Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
 GC Column ID: SPB OCTYL Analysis Time: 19:36:07

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | MZ's FORMING RATIO ² | ION ABUND. RATIO | QC LIMITS ³ | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------|-----------|-------------|-----------------------|---------------------------------|------------------|------------------------|---------------------|---------------------|
| 2-MoCB | 1 | | | M/M+2 | 2.99 | 2.66-3.60 | 18.8 | 17.5 - 32.5 |
| 4-MoCB | 3 | | | M/M+2 | 2.98 | 2.66-3.60 | 19.1 | 17.5 - 32.5 |
| 2,2'-DiCB | 4 | | | M/M+2 | 1.49 | 1.33-1.79 | 18.8 | 17.5 - 32.5 |
| 4,4'-DiCB | 15 | | | M/M+2 | 1.45 | 1.33-1.79 | 21.3 | 19.6 - 36.4 |
| 2,2',6-TriCB | 19 | | | M/M+2 | 1.04 | 0.88-1.20 | 24.3 | 17.5 - 32.5 |
| 3,4,4'-TriCB | 37 | | | M/M+2 | 0.98 | 0.88-1.20 | 18.8 | 17.5 - 32.5 |
| 2,2',6,6'-TeCB | 54 | | | M/M+2 | 0.78 | 0.65-0.89 | 46.1 | 35.0 - 65.0 |
| 3,3',4,4'-TeCB | 77 | | | M/M+2 | 0.77 | 0.65-0.89 | 38.6 | 35.0 - 65.0 |
| 3,4,4',5-TeCB | 81 | | | M/M+2 | 0.80 | 0.65-0.89 | 41.3 | 35.0 - 65.0 |
| 2,2',4,6,6'-PeCB | 104 | | | M+2/M+4 | 1.59 | 1.32-1.78 | 52.1 | 35.0 - 65.0 |
| 2,3,3',4,4'-PeCB | 105 | | | M+2/M+4 | 1.43 | 1.32-1.78 | 42.5 | 35.0 - 65.0 |
| 2,3,4,4',5-PeCB | 114 | | | M+2/M+4 | 1.53 | 1.32-1.78 | 42.2 | 35.0 - 65.0 |
| 2,3',4,4',5-PeCB | 118 | | | M+2/M+4 | 1.45 | 1.32-1.78 | 41.6 | 35.0 - 65.0 |
| 2',3,4,4',5-PeCB | 123 | | | M+2/M+4 | 1.48 | 1.32-1.78 | 40.9 | 35.0 - 65.0 |
| 3,3',4,4',5-PeCB | 126 | | | M+2/M+4 | 1.54 | 1.32-1.78 | 41.0 | 39.0 - 72.4 |
| 2,2',4,4',6,6'-HxCB | 155 | | | M+2/M+4 | 1.28 | 1.05-1.43 | 52.3 | 35.0 - 65.0 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | M+2/M+4 | 1.24 | 1.05-1.43 | 92.8 | 70.0 - 130 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | M+2/M+4 | 1.22 | 1.05-1.43 | 51.1 | 35.0 - 65.0 |
| 3,3',4,4',5,5'-HxCB | 169 | | | M+2/M+4 | 1.26 | 1.05-1.43 | 48.4 | 35.0 - 65.0 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | M+2/M+4 | 1.02 | 0.89-1.21 | 48.3 | 35.0 - 65.0 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | M+2/M+4 | 1.02 | 0.89-1.21 | 40.8 | 35.0 - 65.0 |
| 2,2',3,3',5,5',6,6'-OxCB | 202 | | | M+2/M+4 | 0.91 | 0.76-1.02 | 82.8 | 58.9 - 110 |
| 2,3,3',4,4',5,5',6-OxCB | 205 | | | M+2/M+4 | 0.89 | 0.76-1.02 | 69.9 | 52.5 - 97.5 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | M+2/M+4 | 0.77 | 0.65-0.89 | 73.2 | 52.5 - 97.5 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | M+2/M+4 | 0.78 | 0.65-0.89 | 83.1 | 58.7 - 109 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | M+4/M+6 | 1.19 | 0.99-1.33 | 70.6 | 52.5 - 97.5 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

For Axy Internal Use Only [XSL Template: Form16684A.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5;
 Report Filename: 1668_PCB1668_PB9C_028S1__Form4A_SJ2506476.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 4B
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 1
 Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
 GC Column ID: SPB OCTYL Analysis Time: 19:36:07

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG 2 | MZ's FORMING RATIO 3 | ION ABUND. RATIO | QC LIMITS 4 | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------------|-------------|-------------|------------|----------------------|------------------|-------------|---------------------|---------------------|
| 13C12-2-MoCB | 1L | | | M/M+2 | 3.10 | 2.66-3.60 | 108 | 50.0 - 150 |
| 13C12-4-MoCB | 3L | | | M/M+2 | 3.07 | 2.66-3.60 | 97.8 | 50.0 - 150 |
| 13C12-2,2'-DiCB | 4L | | | M/M+2 | 1.55 | 1.33-1.79 | 91.8 | 50.0 - 150 |
| 13C12-4,4'-DiCB | 15L | | | M/M+2 | 1.53 | 1.33-1.79 | 86.4 | 50.0 - 150 |
| 13C12-2,2',6-TriCB | 19L | | | M/M+2 | 1.03 | 0.88-1.20 | 132 | 50.0 - 150 |
| 13C12-3,4,4'-TriCB | 37L | | | M/M+2 | 1.01 | 0.88-1.20 | 55.3 | 50.0 - 150 |
| 13C12-2,2',6,6'-TeCB | 54L | | | M/M+2 | 0.79 | 0.65-0.89 | 88.8 | 50.0 - 150 |
| 13C12-3,3',4,4'-TeCB | 77L | | | M/M+2 | 0.70 | 0.65-0.89 | 61.8 | 50.0 - 150 |
| 13C12-3,4,4',5-TeCB | 81L | | | M/M+2 | 0.69 | 0.65-0.89 | 65.3 | 50.0 - 150 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | M+2/M+4 | 1.60 | 1.32-1.78 | 81.5 | 50.0 - 150 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 56.9 | 50.0 - 150 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | M+2/M+4 | 1.58 | 1.32-1.78 | 55.6 | 50.0 - 150 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | M+2/M+4 | 1.52 | 1.32-1.78 | 54.9 | 50.0 - 150 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 56.1 | 50.0 - 150 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | M+2/M+4 | 1.52 | 1.32-1.78 | 56.0 | 50.0 - 150 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | M+2/M+4 | 1.29 | 1.05-1.43 | 95.2 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | M+2/M+4 | 1.24 | 1.05-1.43 | 177 | 100 - 300 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | M+2/M+4 | 1.25 | 1.05-1.43 | 88.8 | 50.0 - 150 |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | M+2/M+4 | 1.28 | 1.05-1.43 | 91.9 | 50.0 - 150 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | M+2/M+4 | 1.28 | 1.05-1.43 | 91.9 | 50.0 - 150 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | M+2/M+4 | 1.03 | 0.89-1.21 | 94.7 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | M+2/M+4 | 0.92 | 0.89-1.21 | 58.7 | 50.0 - 150 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | M+2/M+4 | 0.91 | 0.76-1.02 | 82.2 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | M+2/M+4 | 0.85 | 0.76-1.02 | 96.9 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | M+2/M+4 | 0.78 | 0.65-0.89 | 118 | 50.0 - 150 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | M+2/M+4 | 0.75 | 0.65-0.89 | 105 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | M+4/M+6 | 1.18 | 0.99-1.33 | 132 | 50.0 - 150 |

CLEAN-UP STANDARD

| | | | | | | | | |
|-----------------------------|------|--|--|---------|------|-----------|------|------------|
| 13C12-2,4,4'-TriCB | 28L | | | M/M+2 | 0.98 | 0.88-1.20 | 63.7 | 60.0 - 130 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | M+2/M+4 | 1.62 | 1.32-1.78 | 86.0 | 60.0 - 130 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | M+2/M+4 | 1.02 | 0.89-1.21 | 96.6 | 60.0 - 130 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

SGS AXYS METHOD MLA-010 Rev 12

Form 6A
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

VER Data Filename: PB9C_028 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 19:36:07

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RETENTION TIME REFERENCE | IUPAC NO. ² | RRT | RRT QC LIMITS |
|-------------------------------|-----------|-------------|-----------------------|---|------------------------|-------|---------------|
| 2-MoCB | 1 | | | 13C12-2-MoCB | 1L | 1.001 | 0.999-1.004 |
| 4-MoCB | 3 | | | 13C12-4-MoCB | 3L | 1.001 | 0.999-1.004 |
| 2,2'-DiCB | 4 | | | 13C12-2,2'-DiCB | 4L | 1.001 | 0.999-1.004 |
| 4,4'-DiCB | 15 | | | 13C12-4,4'-DiCB | 15L | 1.001 | 0.999-1.002 |
| 2,2',6-TriCB | 19 | | | 13C12-2,2',6-TriCB | 19L | 1.001 | 0.999-1.003 |
| 3,4,4'-TriCB | 37 | | | 13C12-3,4,4'-TriCB | 37L | 1.001 | 0.999-1.002 |
| 2,2',6,6'-TeCB | 54 | | | 13C12-2,2',6,6'-TeCB | 54L | 1.001 | 0.999-1.002 |
| 3,3',4,4'-TeCB | 77 | | | 13C12-3,3',4,4'-TeCB | 77L | 1.000 | 1.000-1.001 |
| 3,4,4',5-TeCB | 81 | | | 13C12-3,4,4',5-TeCB | 81L | 1.000 | 1.000-1.001 |
| 2,2',4,6,6'-PeCB | 104 | | | 13C12-2,2',4,6,6'-PeCB | 104L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4'-PeCB | 105 | | | 13C12-2,3,3',4,4'-PeCB | 105L | 1.001 | 1.000-1.001 |
| 2,3,4,4',5-PeCB | 114 | | | 13C12-2,3,4,4',5-PeCB | 114L | 1.000 | 1.000-1.001 |
| 2,3',4,4',5-PeCB | 118 | | | 13C12-2,3',4,4',5-PeCB | 118L | 1.000 | 1.000-1.001 |
| 2',3,4,4',5-PeCB | 123 | | | 13C12-2',3,4,4',5-PeCB | 123L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5-PeCB | 126 | | | 13C12-3,3',4,4',5-PeCB | 126L | 1.000 | 1.000-1.001 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 13C12-2,2',4,4',6,6'-HxCB | 155L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 13C12-2,3,3',4,4',5-HxCB and 13C12-2,3,3',4,4',5'-HxCB | 156L/157L | 1.000 | 0.998-1.003 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 13C12-2,3',4,4',5,5'-HxCB | 167L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 13C12-3,3',4,4',5,5'-HxCB | 169L | 1.000 | 1.000-1.001 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 13C12-2,2',3,4',5,6,6'-HpCB | 188L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 13C12-2,3,3',4,4',5,5'-HpCB | 189L | 1.001 | 1.000-1.001 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | 1.001 | 1.000-1.001 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | 1.000 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | 1.001 | 1.000-1.001 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) Suffix "L" indicates labeled compound

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

For Axy Internal Use Only [XSL Template: Form16686A.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5;
Report Filename: 1668_PCB1668_PB9C_028S1__Form6A_SJ2506476.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 6B
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 1
 Instrument ID: HR GC/MS Analysis Date: 30-Jan-2019
 GC Column ID: SPB OCTYL Analysis Time: 19:36:07

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RETENTION TIME REFERENCE | IUPAC NO. ¹ | RRT | RRT QC LIMITS |
|-------------------------------------|------------------------|-------------|-----------------------|--------------------------------|------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 13C12-2,5-DiCB | 9L | 0.718 | 0.687-0.749 |
| 13C12-4-MoCB | 3L | | | 13C12-2,5-DiCB | 9L | 0.856 | 0.825-0.888 |
| 13C12-2,2'-DiCB | 4L | | | 13C12-2,5-DiCB | 9L | 0.873 | 0.842-0.904 |
| 13C12-4,4'-DiCB | 15L | | | 13C12-2,5-DiCB | 9L | 1.253 | 1.222-1.284 |
| 13C12-2,2',6-TriCB | 19L | | | 13C12-2,5-DiCB | 9L | 1.072 | 1.041-1.103 |
| 13C12-3,4,4'-TriCB | 37L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.090 | 1.070-1.110 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.810 | 0.797-0.824 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.395 | 1.381-1.408 |
| 13C12-3,4,4',5-TeCB | 81L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.371 | 1.358-1.385 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 0.809 | 0.798-0.819 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.199 | 1.188-1.209 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.178 | 1.168-1.188 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.161 | 1.151-1.171 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.151 | 1.140-1.161 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.299 | 1.289-1.310 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 0.787 | 0.779-0.795 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.108 | 1.099-1.116 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.078 | 1.070-1.086 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.191 | 1.182-1.199 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.713 | 0.707-0.719 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.959 | 0.953-0.965 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.818 | 0.812-0.824 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.009 | 1.000-1.019 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.044 | 1.034-1.053 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-NoCB | 208L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.949 | 0.943-0.956 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.075 | 1.066-1.085 |

CLEANUP STANDARD

| | | | | | | | |
|-----------------------------|------|--|--|---------------------------|------|-------|-------------|
| 13C12-2,4,4'-TriCB | 28L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.924 | 0.910-0.937 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.087 | 1.077-1.098 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.012 | 1.004-1.020 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

PCB CONGENER INITIAL CALIBRATION RELATIVE RESPONSES,
ION ABUNDANCE RATIOS, AND RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

CAL Data Filename: PB9C_028 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 19:36:07

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|----------------|-----------|--------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 3-MoCB | 2 | | | 0.81 | M/M+2 | 2.97 | 2.66-3.60 | 0.988 | 0.984 - 0.991 |
| 2,3-DiCB | 5 | | | 0.82 | M/M+2 | 1.44 | 1.33-1.79 | 1.198 | 1.194 - 1.201 |
| 2,3'-DiCB | 6 | | | 0.91 | M/M+2 | 1.49 | 1.33-1.79 | 1.176 | 1.173 - 1.180 |
| 2,4-DiCB | 7 | | | 0.91 | M/M+2 | 1.48 | 1.33-1.79 | 1.159 | 1.155 - 1.162 |
| 2,4'-DiCB | 8 | | | 1.02 | M/M+2 | 1.48 | 1.33-1.79 | 1.209 | 1.205 - 1.212 |
| 2,5-DiCB | 9 | | | 0.95 | M/M+2 | 1.45 | 1.33-1.79 | 1.147 | 1.143 - 1.150 |
| 2,6-DiCB | 10 | | | 0.93 | M/M+2 | 1.47 | 1.33-1.79 | 1.014 | 1.011 - 1.018 |
| 3,3'-DiCB | 11 | | | 0.81 | M/M+2 | 1.48 | 1.33-1.79 | 0.968 | 0.966 - 0.971 |
| 3,4-DiCB | 12 | 12 + 13 | C | 0.86 | M/M+2 | 1.45 | 1.33-1.79 | 0.984 | 0.982 - 0.987 |
| 3,4'-DiCB | 13 | 12 + 13 | C12 | | | | | | |
| 3,5-DiCB | 14 | | | 0.88 | M/M+2 | 1.50 | 1.33-1.79 | 0.925 | 0.923 - 0.928 |
| 2,2',3-TriCB | 16 | | | 0.95 | M/M+2 | 1.02 | 0.88-1.20 | 1.167 | 1.164 - 1.170 |
| 2,2',4-TriCB | 17 | | | 1.09 | M/M+2 | 1.01 | 0.88-1.20 | 1.140 | 1.137 - 1.143 |
| 2,2',5-TriCB | 18 | 18 + 30 | C | 1.32 | M/M+2 | 1.06 | 0.88-1.20 | 1.114 | 1.111 - 1.117 |
| 2,3,3'-TriCB | 20 | 20 + 28 | C | 0.83 | M/M+2 | 0.97 | 0.88-1.20 | 0.849 | 0.845 - 0.852 |
| 2,3,4-TriCB | 21 | 21 + 33 | C | 0.83 | M/M+2 | 0.98 | 0.88-1.20 | 0.856 | 0.853 - 0.859 |
| 2,3,4'-TriCB | 22 | | | 0.71 | M/M+2 | 0.99 | 0.88-1.20 | 0.872 | 0.870 - 0.874 |
| 2,3,5-TriCB | 23 | | | 0.75 | M/M+2 | 0.98 | 0.88-1.20 | 1.284 | 1.282 - 1.287 |
| 2,3,6-TriCB | 24 | | | 1.47 | M/M+2 | 1.05 | 0.88-1.20 | 1.160 | 1.157 - 1.163 |
| 2,3',4-TriCB | 25 | | | 0.96 | M/M+2 | 0.97 | 0.88-1.20 | 0.825 | 0.824 - 0.827 |
| 2,3',5-TriCB | 26 | 26 + 29 | C | 0.80 | M/M+2 | 0.98 | 0.88-1.20 | 1.304 | 1.299 - 1.309 |
| 2,3',6-TriCB | 27 | | | 1.62 | M/M+2 | 1.06 | 0.88-1.20 | 1.152 | 1.150 - 1.155 |
| 2,4,4'-TriCB | 28 | 20 + 28 | C20 | | | | | | |
| 2,4,5-TriCB | 29 | 26 + 29 | C26 | | | | | | |
| 2,4,6-TriCB | 30 | 18 + 30 | C18 | | | | | | |
| 2,4',5-TriCB | 31 | | | 0.85 | M/M+2 | 0.95 | 0.88-1.20 | 0.837 | 0.835 - 0.839 |
| 2,4',6-TriCB | 32 | | | 0.82 | M/M+2 | 0.96 | 0.88-1.20 | 1.199 | 1.196 - 1.202 |
| 2',3,4-TriCB | 33 | 21 + 33 | C21 | | | | | | |
| 2',3,5-TriCB | 34 | | | 0.74 | M/M+2 | 0.98 | 0.88-1.20 | 1.276 | 1.273 - 1.279 |
| 3,3',4-TriCB | 35 | | | 0.74 | M/M+2 | 0.98 | 0.88-1.20 | 0.985 | 0.983 - 0.987 |
| 3,3',5-TriCB | 36 | | | 0.79 | M/M+2 | 1.01 | 0.88-1.20 | 0.932 | 0.930 - 0.934 |
| 3,4,5-TriCB | 38 | | | 0.84 | M/M+2 | 0.99 | 0.88-1.20 | 0.967 | 0.965 - 0.969 |
| 3,4',5-TriCB | 39 | | | 0.80 | M/M+2 | 0.99 | 0.88-1.20 | 0.945 | 0.943 - 0.947 |
| 2,2',3,3'-TeCB | 40 | 40 + 41 + 71 | C | 0.88 | M/M+2 | 0.79 | 0.65-0.89 | 1.336 | 1.331 - 1.340 |
| 2,2',3,4-TeCB | 41 | 40 + 41 + 71 | C40 | | | | | | |
| 2,2',3,4'-TeCB | 42 | | | 0.84 | M/M+2 | 0.78 | 0.65-0.89 | 1.312 | 1.309 - 1.314 |
| 2,2',3,5-TeCB | 43 | | | 0.67 | M/M+2 | 0.78 | 0.65-0.89 | 1.247 | 1.245 - 1.250 |
| 2,2',3,5'-TeCB | 44 | 44 + 47 + 65 | C | 0.97 | M/M+2 | 0.78 | 0.65-0.89 | 1.286 | 1.282 - 1.291 |
| 2,2',3,6-TeCB | 45 | 45 + 51 | C | 0.89 | M/M+2 | 0.77 | 0.65-0.89 | 1.147 | 1.143 - 1.151 |
| 2,2',3,6'-TeCB | 46 | | | 0.77 | M/M+2 | 0.79 | 0.65-0.89 | 1.160 | 1.158 - 1.163 |
| 2,2',4,4'-TeCB | 47 | 44 + 47 + 65 | C44 | | | | | | |
| 2,2',4,5-TeCB | 48 | | | 0.83 | M/M+2 | 0.77 | 0.65-0.89 | 1.274 | 1.272 - 1.277 |
| 2,2',4,5'-TeCB | 49 | 49 + 69 | C | 1.02 | M/M+2 | 0.79 | 0.65-0.89 | 1.259 | 1.255 - 1.263 |
| 2,2',4,6-TeCB | 50 | 50 + 53 | C | 0.93 | M/M+2 | 0.77 | 0.65-0.89 | 1.112 | 1.108 - 1.116 |
| 2,2',4,6'-TeCB | 51 | 45 + 51 | C45 | | | | | | |
| 2,2',5,5'-TeCB | 52 | | | 0.94 | M/M+2 | 0.79 | 0.65-0.89 | 1.234 | 1.232 - 1.236 |
| 2,2',5,6'-TeCB | 53 | 50 + 53 | C50 | | | | | | |
| 2,3,3',4-TeCB | 55 | | | 0.66 | M/M+2 | 0.78 | 0.65-0.89 | 0.889 | 0.888 - 0.891 |
| 2,3,3',4'-TeCB | 56 | | | 0.69 | M/M+2 | 0.77 | 0.65-0.89 | 0.905 | 0.903 - 0.906 |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|-------------------|-----------|--------------------------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 2,3,3',5'-TeCB | 57 | | | 0.75 | M/M+2 | 0.75 | 0.65-0.89 | 0.845 | 0.843 - 0.846 |
| 2,3,3',5'-TeCB | 58 | | | 0.69 | M/M+2 | 0.76 | 0.65-0.89 | 0.852 | 0.850 - 0.853 |
| 2,3,3',6'-TeCB | 59 | 59 + 62 + 75 | C | 1.20 | M/M+2 | 0.77 | 0.65-0.89 | 1.302 | 1.298 - 1.306 |
| 2,3,4,4'-TeCB | 60 | | | 0.72 | M/M+2 | 0.76 | 0.65-0.89 | 0.911 | 0.910 - 0.912 |
| 2,3,4,5'-TeCB | 61 | 61 + 70 + 74 + 76 | C | 0.75 | M/M+2 | 0.75 | 0.65-0.89 | 0.874 | 0.871 - 0.877 |
| 2,3,4,6'-TeCB | 62 | 59 + 62 + 75 | C59 | | | | | | |
| 2,3,4',5'-TeCB | 63 | | | 0.74 | M/M+2 | 0.82 | 0.65-0.89 | 0.865 | 0.863 - 0.866 |
| 2,3,4',6'-TeCB | 64 | | | 1.24 | M/M+2 | 0.77 | 0.65-0.89 | 1.349 | 1.346 - 1.351 |
| 2,3,5,6'-TeCB | 65 | 44 + 47 + 65 | C44 | | | | | | |
| 2,3',4,4'-TeCB | 66 | | | 0.72 | M/M+2 | 0.78 | 0.65-0.89 | 0.885 | 0.883 - 0.886 |
| 2,3',4,5'-TeCB | 67 | | | 0.88 | M/M+2 | 0.75 | 0.65-0.89 | 0.856 | 0.855 - 0.858 |
| 2,3',4,5'-TeCB | 68 | | | 0.78 | M/M+2 | 0.80 | 0.65-0.89 | 0.832 | 0.830 - 0.833 |
| 2,3',4,6'-TeCB | 69 | 49 + 69 | C49 | | | | | | |
| 2,3',4',5'-TeCB | 70 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 2,3',4',6'-TeCB | 71 | 40 + 41 + 71 | C40 | | | | | | |
| 2,3',5,5'-TeCB | 72 | | | 0.76 | M/M+2 | 0.75 | 0.65-0.89 | 0.823 | 0.822 - 0.824 |
| 2,3',5,6'-TeCB | 73 | | | 1.19 | M/M+2 | 0.77 | 0.65-0.89 | 1.242 | 1.240 - 1.245 |
| 2,4,4',5'-TeCB | 74 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 2,4,4',6'-TeCB | 75 | 59 + 62 + 75 | C59 | | | | | | |
| 2',3,4,5'-TeCB | 76 | 61 + 70 + 74 + 76 | C61 | | | | | | |
| 3,3',4,5'-TeCB | 78 | | | 0.72 | M/M+2 | 0.80 | 0.65-0.89 | 0.987 | 0.986 - 0.989 |
| 3,3',4,5'-TeCB | 79 | | | 0.94 | M/M+2 | 0.75 | 0.65-0.89 | 0.971 | 0.969 - 0.972 |
| 3,3',5,5'-TeCB | 80 | | | 0.80 | M/M+2 | 0.76 | 0.65-0.89 | 0.925 | 0.923 - 0.926 |
| 2,2',3,3',4'-PeCB | 82 | | | 0.89 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.933 | 0.932 - 0.935 |
| 2,2',3,3',5'-PeCB | 83 | 83 + 99 | C | 0.93 | M+2/M+4 | 1.58 | 1.32-1.78 | 0.884 | 0.882 - 0.887 |
| 2,2',3,3',6'-PeCB | 84 | | | 0.87 | M+2/M+4 | 1.55 | 1.32-1.78 | 1.161 | 1.160 - 1.163 |
| 2,2',3,4,4'-PeCB | 85 | 85 + 116 + 117 | C | 1.23 | M+2/M+4 | 1.56 | 1.32-1.78 | 0.919 | 0.917 - 0.922 |
| 2,2',3,4,5'-PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | C | 1.15 | M+2/M+4 | 1.55 | 1.32-1.78 | 0.900 | 0.897 - 0.904 |
| 2,2',3,4,5'-PeCB | 87 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,2',3,4,6'-PeCB | 88 | 88 + 91 | C | 0.99 | M+2/M+4 | 1.59 | 1.32-1.78 | 1.152 | 1.148 - 1.156 |
| 2,2',3,4,6'-PeCB | 89 | | | 0.92 | M+2/M+4 | 1.55 | 1.32-1.78 | 1.181 | 1.179 - 1.183 |
| 2,2',3,4',5'-PeCB | 90 | 90 + 101 + 113 | C | 1.17 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.869 | 0.866 - 0.871 |
| 2,2',3,4',6'-PeCB | 91 | 88 + 91 | C88 | | | | | | |
| 2,2',3,5,5'-PeCB | 92 | | | 0.97 | M+2/M+4 | 1.57 | 1.32-1.78 | 0.853 | 0.852 - 0.854 |
| 2,2',3,5,6'-PeCB | 93 | 93 + 95 + 98 + 100 + 102 | C | 1.01 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.129 | 1.118 - 1.140 |
| 2,2',3,5,6'-PeCB | 94 | | | 0.91 | M+2/M+4 | 1.54 | 1.32-1.78 | 1.101 | 1.100 - 1.103 |
| 2,2',3,5',6'-PeCB | 95 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',3,6,6'-PeCB | 96 | | | 1.42 | M+2/M+4 | 1.61 | 1.32-1.78 | 1.015 | 1.011 - 1.018 |
| 2,2',3',4,5'-PeCB | 97 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,2',3',4,6'-PeCB | 98 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,4',5'-PeCB | 99 | 83 + 99 | C83 | | | | | | |
| 2,2',4,4',6'-PeCB | 100 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,5,5'-PeCB | 101 | 90 + 101 + 113 | C90 | | | | | | |
| 2,2',4,5,6'-PeCB | 102 | 93 + 95 + 98 + 100 + 102 | C93 | | | | | | |
| 2,2',4,5',6'-PeCB | 103 | | | 1.10 | M+2/M+4 | 1.54 | 1.32-1.78 | 1.093 | 1.091 - 1.095 |
| 2,3,3',4,5'-PeCB | 106 | | | 0.85 | M+2/M+4 | 1.48 | 1.32-1.78 | 1.004 | 1.002 - 1.005 |
| 2,3,3',4',5'-PeCB | 107 | 107 + 124 | C | 0.79 | M+2/M+4 | 1.49 | 1.32-1.78 | 0.990 | 0.988 - 0.992 |
| 2,3,3',4,5'-PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,3,3',4,6'-PeCB | 109 | | | 0.87 | M+2/M+4 | 1.44 | 1.32-1.78 | 0.997 | 0.996 - 0.998 |
| 2,3,3',4',6'-PeCB | 110 | 110 + 115 | C | 1.41 | M+2/M+4 | 1.55 | 1.32-1.78 | 0.926 | 0.924 - 0.928 |
| 2,3,3',5,5'-PeCB | 111 | | | 1.36 | M+2/M+4 | 1.58 | 1.32-1.78 | 0.945 | 0.944 - 0.947 |
| 2,3,3',5,6'-PeCB | 112 | | | 1.40 | M+2/M+4 | 1.61 | 1.32-1.78 | 0.889 | 0.887 - 0.890 |
| 2,3,3',5',6'-PeCB | 113 | 90 + 101 + 113 | C90 | | | | | | |
| 2,3,4,4',6'-PeCB | 115 | 110 + 115 | C110 | | | | | | |
| 2,3,4,5,6'-PeCB | 116 | 85 + 116 + 117 | C85 | | | | | | |
| 2,3,4',5,6'-PeCB | 117 | 85 + 116 + 117 | C85 | | | | | | |
| 2,3',4,4',6'-PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |
| 2,3',4,5,5'-PeCB | 120 | | | 1.45 | M+2/M+4 | 1.54 | 1.32-1.78 | 0.959 | 0.957 - 0.960 |
| 2,3',4,5',6'-PeCB | 121 | | | 1.30 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.200 | 1.198 - 1.202 |
| 2',3,3',4,5'-PeCB | 122 | | | 0.72 | M+2/M+4 | 1.48 | 1.32-1.78 | 1.010 | 1.009 - 1.012 |
| 2',3,4,5,5'-PeCB | 124 | 107 + 124 | C107 | | | | | | |
| 2',3,4,5,6'-PeCB | 125 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | | | | |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|--------------------------|-----------|-----------------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 3,3',4,5,5'-PeCB | 127 | | | 0.77 | M+2/M+4 | 1.38 | 1.32-1.78 | 1.041 | 1.040 - 1.042 |
| 2,2',3,3',4,4'-HxCB | 128 | 128 + 166 | C | 0.85 | M+2/M+4 | 1.26 | 1.05-1.43 | 0.958 | 0.956 - 0.960 |
| 2,2',3,3',4,5-HxCB | 129 | 129 + 138 + 160 + 163 | C | 0.83 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.930 | 0.928 - 0.933 |
| 2,2',3,3',4,5'-HxCB | 130 | | | 0.64 | M+2/M+4 | 1.27 | 1.05-1.43 | 0.913 | 0.912 - 0.914 |
| 2,2',3,3',4,6-HxCB | 131 | | | 0.68 | M+2/M+4 | 1.25 | 1.05-1.43 | 1.159 | 1.157 - 1.160 |
| 2,2',3,3',4,6'-HxCB | 132 | | | 0.65 | M+2/M+4 | 1.24 | 1.05-1.43 | 1.173 | 1.170 - 1.176 |
| 2,2',3,3',5,5'-HxCB | 133 | | | 0.71 | M+2/M+4 | 1.22 | 1.05-1.43 | 1.190 | 1.189 - 1.192 |
| 2,2',3,3',5,6-HxCB | 134 | 134 + 143 | C | 0.68 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.141 | 1.138 - 1.143 |
| 2,2',3,3',5,6'-HxCB | 135 | 135 + 151 + 154 | C | 0.82 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.106 | 1.100 - 1.112 |
| 2,2',3,3',6,6'-HxCB | 136 | | | 1.06 | M+2/M+4 | 1.28 | 1.05-1.43 | 1.023 | 1.021 - 1.024 |
| 2,2',3,4,4',5-HxCB | 137 | | | 0.65 | M+2/M+4 | 1.22 | 1.05-1.43 | 0.919 | 0.918 - 0.920 |
| 2,2',3,4,4',5'-HxCB | 138 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,2',3,4,4',6-HxCB | 139 | 139 + 140 | C | 0.75 | M+2/M+4 | 1.26 | 1.05-1.43 | 1.152 | 1.150 - 1.155 |
| 2,2',3,4,4',6'-HxCB | 140 | 139 + 140 | C139 | | | | | | |
| 2,2',3,4,5,5'-HxCB | 141 | | | 0.73 | M+2/M+4 | 1.27 | 1.05-1.43 | 0.904 | 0.902 - 0.905 |
| 2,2',3,4,5,6-HxCB | 142 | | | 0.65 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.164 | 1.162 - 1.165 |
| 2,2',3,4,5,6'-HxCB | 143 | 134 + 143 | C134 | | | | | | |
| 2,2',3,4,5',6-HxCB | 144 | | | 0.79 | M+2/M+4 | 1.28 | 1.05-1.43 | 1.121 | 1.119 - 1.122 |
| 2,2',3,4,6,6'-HxCB | 145 | | | 0.94 | M+2/M+4 | 1.27 | 1.05-1.43 | 1.033 | 1.032 - 1.035 |
| 2,2',3,4',5,5'-HxCB | 146 | | | 0.88 | M+2/M+4 | 1.24 | 1.05-1.43 | 0.885 | 0.883 - 0.886 |
| 2,2',3,4',5,6-HxCB | 147 | 147 + 149 | C | 0.77 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.133 | 1.130 - 1.135 |
| 2,2',3,4',5,6'-HxCB | 148 | | | 0.78 | M+2/M+4 | 1.23 | 1.05-1.43 | 1.084 | 1.082 - 1.085 |
| 2,2',3,4',5',6-HxCB | 149 | 147 + 149 | C147 | | | | | | |
| 2,2',3,4',6,6'-HxCB | 150 | | | 1.03 | M+2/M+4 | 1.27 | 1.05-1.43 | 1.012 | 1.010 - 1.014 |
| 2,2',3,5,5',6-HxCB | 151 | 135 + 151 + 154 | C135 | | | | | | |
| 2,2',3,5,6,6'-HxCB | 152 | | | 1.15 | M+2/M+4 | 1.29 | 1.05-1.43 | 1.006 | 1.005 - 1.008 |
| 2,2',4,4',5,5'-HxCB | 153 | 153 + 168 | C | 0.93 | M+2/M+4 | 1.23 | 1.05-1.43 | 0.900 | 0.898 - 0.902 |
| 2,2',4,4',5,6'-HxCB | 154 | 135 + 151 + 154 | C135 | | | | | | |
| 2,3,3',4,4',6-HxCB | 158 | | | 1.07 | M+2/M+4 | 1.22 | 1.05-1.43 | 0.938 | 0.937 - 0.939 |
| 2,3,3',4,5,5'-HxCB | 159 | | | 1.02 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.983 | 0.981 - 0.984 |
| 2,3,3',4,5,6-HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,3,3',4,5',6-HxCB | 161 | | | 1.00 | M+2/M+4 | 1.24 | 1.05-1.43 | 0.888 | 0.887 - 0.889 |
| 2,3,3',4',5,5'-HxCB | 162 | | | 1.04 | M+2/M+4 | 1.20 | 1.05-1.43 | 0.989 | 0.988 - 0.991 |
| 2,3,3',4',5,6-HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | | | |
| 2,3,3',4',5',6-HxCB | 164 | | | 1.00 | M+2/M+4 | 1.23 | 1.05-1.43 | 0.921 | 0.920 - 0.922 |
| 2,3,3',5,5',6-HxCB | 165 | | | 0.85 | M+2/M+4 | 1.25 | 1.05-1.43 | 0.879 | 0.878 - 0.880 |
| 2,3,4,4',5,6-HxCB | 166 | 128 + 166 | C128 | | | | | | |
| 2,3',4,4',5',6-HxCB | 168 | 153 + 168 | C153 | | | | | | |
| 2,2',3,3',4,4',5-HpCB | 170 | | | 1.14 | M+2/M+4 | 1.03 | 0.89-1.21 | 1.000 | 0.999 - 1.001 |
| 2,2',3,3',4,4',6-HpCB | 171 | 171 + 173 | C | 0.81 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.162 | 1.159 - 1.164 |
| 2,2',3,3',4,5,5'-HpCB | 172 | | | 0.79 | M+2/M+4 | 1.05 | 0.89-1.21 | 0.897 | 0.896 - 0.898 |
| 2,2',3,3',4,5,6-HpCB | 173 | 171 + 173 | C171 | | | | | | |
| 2,2',3,3',4,5,6'-HpCB | 174 | | | 0.84 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.133 | 1.131 - 1.134 |
| 2,2',3,3',4,5',6-HpCB | 175 | | | 0.89 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.102 | 1.101 - 1.103 |
| 2,2',3,3',4,6,6'-HpCB | 176 | | | 1.15 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.034 | 1.032 - 1.035 |
| 2,2',3,3',4',5,6-HpCB | 177 | | | 1.10 | M+2/M+4 | 1.06 | 0.89-1.21 | 1.145 | 1.143 - 1.146 |
| 2,2',3,3',5,5',6-HpCB | 178 | | | 0.79 | M+2/M+4 | 1.03 | 0.89-1.21 | 1.085 | 1.083 - 1.086 |
| 2,2',3,3',5,6,6'-HpCB | 179 | | | 1.14 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.009 | 1.008 - 1.011 |
| 2,2',3,4,4',5,5'-HpCB | 180 | 180 + 193 | C | 1.00 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.000 | 0.999 - 1.001 |
| 2,2',3,4,4',5,6-HpCB | 181 | | | 0.87 | M+2/M+4 | 1.03 | 0.89-1.21 | 1.156 | 1.155 - 1.157 |
| 2,2',3,4,4',5,6'-HpCB | 182 | | | 0.89 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.115 | 1.114 - 1.117 |
| 2,2',3,4,4',5',6-HpCB | 183 | 183 + 185 | C | 0.90 | M+2/M+4 | 1.04 | 0.89-1.21 | 1.127 | 1.126 - 1.129 |
| 2,2',3,4,4',6,6'-HpCB | 184 | | | 1.19 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.025 | 1.023 - 1.026 |
| 2,2',3,4,5,5',6-HpCB | 185 | 183 + 185 | C183 | | | | | | |
| 2,2',3,4,5,6,6'-HpCB | 186 | | | 1.04 | M+2/M+4 | 1.03 | 0.89-1.21 | 1.046 | 1.045 - 1.048 |
| 2,2',3,4',5,5',6-HpCB | 187 | | | 0.87 | M+2/M+4 | 1.05 | 0.89-1.21 | 1.110 | 1.109 - 1.111 |
| 2,3,3',4,4',5,6-HpCB | 190 | | | 1.15 | M+2/M+4 | 1.04 | 0.89-1.21 | 0.947 | 0.946 - 0.948 |
| 2,3,3',4,4',5',6-HpCB | 191 | | | 1.17 | M+2/M+4 | 1.04 | 0.89-1.21 | 0.918 | 0.917 - 0.919 |
| 2,3,3',4,5,5',6-HpCB | 192 | | | 1.00 | M+2/M+4 | 1.02 | 0.89-1.21 | 0.903 | 0.902 - 0.904 |
| 2,3,3',4',5,5',6-HpCB | 193 | 180 + 193 | C180 | | | | | | |
| 2,2',3,3',4,4',5,5'-OcCB | 194 | | | 0.84 | M+2/M+4 | 0.87 | 0.76-1.02 | 0.991 | 0.990 - 0.992 |
| 2,2',3,3',4,4',5,6-OcCB | 195 | | | 0.71 | M+2/M+4 | 0.87 | 0.76-1.02 | 0.946 | 0.945 - 0.946 |
| 2,2',3,3',4,4',5,6'-OcCB | 196 | | | 0.75 | M+2/M+4 | 0.87 | 0.76-1.02 | 0.916 | 0.915 - 0.917 |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RRF | MZ's FORMING RATIO ² | ION ABUND. RATIO | RATIO QC LIMITS ³ | RRT | RRT QC LIMITS |
|-----------------------------------|-----------|-------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 2,2',3,3',4,4',6,6'-OcCB | 197 | 197 + 200 | C | 0.92 | M+2/M+4 | 0.89 | 0.76-1.02 | 1.045 | 1.043 - 1.048 |
| 2,2',3,3',4,5,5',6-OcCB | 198 | 198 + 199 | C | 0.69 | M+2/M+4 | 0.88 | 0.76-1.02 | 1.114 | 1.112 - 1.116 |
| 2,2',3,3',4,5,5',6'-OcCB | 199 | 198 + 199 | C198 | | | | | | |
| 2,2',3,3',4,5,6,6'-OcCB | 200 | 197 + 200 | C197 | | | | | | |
| 2,2',3,3',4,5',6,6'-OcCB | 201 | | | 0.88 | M+2/M+4 | 0.89 | 0.76-1.02 | 1.023 | 1.021 - 1.024 |
| 2,2',3,4,4',5,5',6-OcCB | 203 | | | 0.77 | M+2/M+4 | 0.92 | 0.76-1.02 | 0.920 | 0.919 - 0.921 |
| 2,2',3,4,4',5,6,6'-OcCB | 204 | | | 0.88 | M+2/M+4 | 0.90 | 0.76-1.02 | 1.039 | 1.038 - 1.040 |
| 2,2',3,3',4,4',5,6,6'-NoCB | 207 | | | 1.22 | M+2/M+4 | 0.78 | 0.65-0.89 | 1.020 | 1.019 - 1.021 |

(1) Where applicable, custom lab flags have been used on this report.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 3B

PCB CONGENER INITIAL CALIBRATION RELATIVE RESPONSES,
ION ABUNDANCE RATIOS, AND RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

CAL Data Filename: PB9C_028 S: 1

Instrument ID: HR GC/MS

Analysis Date: 30-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 19:36:07

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RRF | MZ's FORMING RATIO ³ | ION ABUND. RATIO | RATIO QC LIMITS ⁴ | RRT | RRT QC LIMITS |
|-------------------------------------|------------------------|-------------|-----------------------|------|---------------------------------|------------------|------------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 1.24 | M/M+2 | 3.10 | 2.66-3.60 | 0.718 | 0.702 - 0.734 |
| 13C12-4-MoCB | 3L | | | 1.11 | M/M+2 | 3.07 | 2.66-3.60 | 0.856 | 0.841 - 0.872 |
| 13C12-2,2'-DiCB | 4L | | | 0.63 | M/M+2 | 1.55 | 1.33-1.79 | 0.873 | 0.857 - 0.889 |
| 13C12-4,4'-DiCB | 15L | | | 0.94 | M/M+2 | 1.53 | 1.33-1.79 | 1.253 | 1.237 - 1.268 |
| 13C12-2,2',6'-TriCB | 19L | | | 0.65 | M/M+2 | 1.03 | 0.88-1.20 | 1.072 | 1.056 - 1.087 |
| 13C12-3,4,4'-TriCB | 37L | | | 1.14 | M/M+2 | 1.01 | 0.88-1.20 | 1.090 | 1.080 - 1.100 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 1.44 | M/M+2 | 0.79 | 0.65-0.89 | 0.810 | 0.804 - 0.817 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 0.99 | M/M+2 | 0.70 | 0.65-0.89 | 1.395 | 1.388 - 1.401 |
| 13C12-3,4,4',5'-TeCB | 81L | | | 1.05 | M/M+2 | 0.69 | 0.65-0.89 | 1.371 | 1.365 - 1.378 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 1.16 | M+2/M+4 | 1.60 | 1.32-1.78 | 0.809 | 0.803 - 0.814 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 0.93 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.199 | 1.193 - 1.204 |
| 13C12-2,3,4,4',5'-PeCB | 114L | | | 0.94 | M+2/M+4 | 1.58 | 1.32-1.78 | 1.178 | 1.173 - 1.183 |
| 13C12-2,3',4,4',5'-PeCB | 118L | | | 0.88 | M+2/M+4 | 1.52 | 1.32-1.78 | 1.161 | 1.156 - 1.166 |
| 13C12-2',3,4,4',5'-PeCB | 123L | | | 0.90 | M+2/M+4 | 1.57 | 1.32-1.78 | 1.151 | 1.146 - 1.156 |
| 13C12-3,3',4,4',5'-PeCB | 126L | | | 0.88 | M+2/M+4 | 1.52 | 1.32-1.78 | 1.299 | 1.294 - 1.304 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 1.31 | M+2/M+4 | 1.29 | 1.05-1.43 | 0.787 | 0.783 - 0.791 |
| 13C12-2,3,3',4,4',5'-HxCB | 156L | 156L + 157L | C | 1.30 | M+2/M+4 | 1.24 | 1.05-1.43 | 1.108 | 1.103 - 1.112 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 1.24 | M+2/M+4 | 1.25 | 1.05-1.43 | 1.078 | 1.074 - 1.082 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 1.40 | M+2/M+4 | 1.28 | 1.05-1.43 | 1.191 | 1.187 - 1.195 |
| 13C12-2,2',3,3',4,4',5'-HpCB | 170L | | | 0.88 | M+2/M+4 | 1.05 | 0.89-1.21 | 0.897 | 0.894 - 0.900 |
| 13C12-2,2',3,4,4',5,5'-HpCB | 180L | | | 1.01 | M+2/M+4 | 1.07 | 0.89-1.21 | 0.873 | 0.870 - 0.876 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 1.43 | M+2/M+4 | 1.03 | 0.89-1.21 | 0.713 | 0.710 - 0.716 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 0.91 | M+2/M+4 | 0.92 | 0.89-1.21 | 0.959 | 0.956 - 0.962 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | 1.22 | M+2/M+4 | 0.91 | 0.76-1.02 | 0.818 | 0.815 - 0.821 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | 1.40 | M+2/M+4 | 0.85 | 0.76-1.02 | 1.009 | 1.005 - 1.014 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 1.13 | M+2/M+4 | 0.78 | 0.65-0.89 | 1.044 | 1.039 - 1.048 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-NoCB | 208L | | | 1.27 | M+2/M+4 | 0.75 | 0.65-0.89 | 0.949 | 0.946 - 0.953 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 4A
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 9
 Instrument ID: HR GC/MS Analysis Date: 31-Jan-2019
 GC Column ID: SPB OCTYL Analysis Time: 04:10:02

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | MZ's FORMING RATIO ² | ION ABUND. RATIO | QC LIMITS ³ | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------|-----------|-------------|-----------------------|---------------------------------|------------------|------------------------|---------------------|---------------------|
| 2-MoCB | 1 | | | M/M+2 | 2.95 | 2.66-3.60 | 18.1 | 17.5 - 32.5 |
| 4-MoCB | 3 | | | M/M+2 | 2.95 | 2.66-3.60 | 18.2 | 17.5 - 32.5 |
| 2,2'-DiCB | 4 | | | M/M+2 | 1.46 | 1.33-1.79 | 18.0 | 17.5 - 32.5 |
| 4,4'-DiCB | 15 | | | M/M+2 | 1.46 | 1.33-1.79 | 20.7 | 19.6 - 36.4 |
| 2,2',6-TriCB | 19 | | | M/M+2 | 1.04 | 0.88-1.20 | 24.0 | 17.5 - 32.5 |
| 3,4,4'-TriCB | 37 | | | M/M+2 | 0.97 | 0.88-1.20 | 18.6 | 17.5 - 32.5 |
| 2,2',6,6'-TeCB | 54 | | | M/M+2 | 0.77 | 0.65-0.89 | 45.2 | 35.0 - 65.0 |
| 3,3',4,4'-TeCB | 77 | | | M/M+2 | 0.74 | 0.65-0.89 | 39.2 | 35.0 - 65.0 |
| 3,4,4',5-TeCB | 81 | | | M/M+2 | 0.75 | 0.65-0.89 | 41.4 | 35.0 - 65.0 |
| 2,2',4,6,6'-PeCB | 104 | | | M+2/M+4 | 1.54 | 1.32-1.78 | 52.0 | 35.0 - 65.0 |
| 2,3,3',4,4'-PeCB | 105 | | | M+2/M+4 | 1.41 | 1.32-1.78 | 39.2 | 35.0 - 65.0 |
| 2,3,4,4',5-PeCB | 114 | | | M+2/M+4 | 1.37 | 1.32-1.78 | 38.7 | 35.0 - 65.0 |
| 2,3',4,4',5-PeCB | 118 | | | M+2/M+4 | 1.36 | 1.32-1.78 | 38.9 | 35.0 - 65.0 |
| 2',3,4,4',5-PeCB | 123 | | | M+2/M+4 | 1.44 | 1.32-1.78 | 39.8 | 35.0 - 65.0 |
| 3,3',4,4',5-PeCB | 126 | | | M+2/M+4 | 1.52 | 1.32-1.78 | 41.4 | 39.0 - 72.4 |
| 2,2',4,4',6,6'-HxCB | 155 | | | M+2/M+4 | 1.29 | 1.05-1.43 | 51.4 | 35.0 - 65.0 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | M+2/M+4 | 1.18 | 1.05-1.43 | 91.0 | 70.0 - 130 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | M+2/M+4 | 1.21 | 1.05-1.43 | 49.5 | 35.0 - 65.0 |
| 3,3',4,4',5,5'-HxCB | 169 | | | M+2/M+4 | 1.29 | 1.05-1.43 | 49.4 | 35.0 - 65.0 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | M+2/M+4 | 1.09 | 0.89-1.21 | 47.2 | 35.0 - 65.0 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | M+2/M+4 | 1.01 | 0.89-1.21 | 41.2 | 35.0 - 65.0 |
| 2,2',3,3',5,5',6,6'-OoCB | 202 | | | M+2/M+4 | 0.92 | 0.76-1.02 | 77.5 | 58.9 - 110 |
| 2,3,3',4,4',5,5',6-OoCB | 205 | | | M+2/M+4 | 0.89 | 0.76-1.02 | 70.2 | 52.5 - 97.5 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | M+2/M+4 | 0.78 | 0.65-0.89 | 74.3 | 52.5 - 97.5 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | M+2/M+4 | 0.78 | 0.65-0.89 | 83.1 | 58.7 - 109 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | M+4/M+6 | 1.19 | 0.99-1.33 | 71.6 | 52.5 - 97.5 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) See Table 8, Method 1668A, for m/z specifications.

(3) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

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SGS AXYS METHOD MLA-010 Rev 12

Form 4B
PCB CONGENER CALIBRATION VERIFICATION

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 9
Instrument ID: HR GC/MS Analysis Date: 31-Jan-2019
GC Column ID: SPB OCTYL Analysis Time: 04:10:02

| LABELLED COMPOUND | IUPAC NO. 1 | CO-ELUTIONS | LAB FLAG 2 | MZ's FORMING RATIO 3 | ION ABUND. RATIO | QC LIMITS 4 | CONC. FOUND (ng/mL) | CONC. RANGE (ng/mL) |
|-------------------------------------|-------------|-------------|------------|----------------------|------------------|-------------|---------------------|---------------------|
| 13C12-2-MoCB | 1L | | | M/M+2 | 3.10 | 2.66-3.60 | 109 | 50.0 - 150 |
| 13C12-4-MoCB | 3L | | | M/M+2 | 3.06 | 2.66-3.60 | 102 | 50.0 - 150 |
| 13C12-2,2'-DiCB | 4L | | | M/M+2 | 1.50 | 1.33-1.79 | 91.4 | 50.0 - 150 |
| 13C12-4,4'-DiCB | 15L | | | M/M+2 | 1.51 | 1.33-1.79 | 88.0 | 50.0 - 150 |
| 13C12-2,2',6-TriCB | 19L | | | M/M+2 | 1.06 | 0.88-1.20 | 142 | 50.0 - 150 |
| 13C12-3,4,4'-TriCB | 37L | | | M/M+2 | 0.99 | 0.88-1.20 | 51.9 | 50.0 - 150 |
| 13C12-2,2',6,6'-TeCB | 54L | | | M/M+2 | 0.79 | 0.65-0.89 | 88.0 | 50.0 - 150 |
| 13C12-3,3',4,4'-TeCB | 77L | | | M/M+2 | 0.71 | 0.65-0.89 | 58.8 | 50.0 - 150 |
| 13C12-3,4,4',5-TeCB | 81L | | | M/M+2 | 0.68 | 0.65-0.89 | 62.7 | 50.0 - 150 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | M+2/M+4 | 1.60 | 1.32-1.78 | 85.7 | 50.0 - 150 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | M+2/M+4 | 1.52 | 1.32-1.78 | 54.8 | 50.0 - 150 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | M+2/M+4 | 1.59 | 1.32-1.78 | 54.0 | 50.0 - 150 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | M+2/M+4 | 1.47 | 1.32-1.78 | 52.3 | 50.0 - 150 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 54.9 | 50.0 - 150 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | M+2/M+4 | 1.55 | 1.32-1.78 | 53.4 | 50.0 - 150 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | M+2/M+4 | 1.27 | 1.05-1.43 | 106 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | M+2/M+4 | 1.26 | 1.05-1.43 | 178 | 100 - 300 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | M+2/M+4 | 1.23 | 1.05-1.43 | 90.1 | 50.0 - 150 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | M+2/M+4 | 1.23 | 1.05-1.43 | 91.8 | 50.0 - 150 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | M+2/M+4 | 1.03 | 0.89-1.21 | 109 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | M+2/M+4 | 0.95 | 0.89-1.21 | 63.9 | 50.0 - 150 |
| 13C12-2,2',3,3',5,5',6,6'-OxCB | 202L | | | M+2/M+4 | 0.89 | 0.76-1.02 | 99.3 | 50.0 - 150 |
| 13C12-2,3,3',4,4',5,5',6-OxCB | 205L | | | M+2/M+4 | 0.84 | 0.76-1.02 | 99.0 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | M+2/M+4 | 0.76 | 0.65-0.89 | 119 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-NoCB | 208L | | | M+2/M+4 | 0.75 | 0.65-0.89 | 109 | 50.0 - 150 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | M+4/M+6 | 1.20 | 0.99-1.33 | 140 | 50.0 - 150 |

CLEAN-UP STANDARD

| | | | | | | | | |
|-----------------------------|------|--|--|---------|------|-----------|------|------------|
| 13C12-2,4,4'-TriCB | 28L | | | M/M+2 | 1.00 | 0.88-1.20 | 59.6 | 60.0 - 130 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | M+2/M+4 | 1.57 | 1.32-1.78 | 86.7 | 60.0 - 130 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | M+2/M+4 | 1.04 | 0.89-1.21 | 105 | 60.0 - 130 |

(1) Suffix "L" indicates labeled compound.

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(3) See Table 8, Method 1668A, for m/z specifications.

(4) Ion Abundance Ratio Control Limits as specified in Table 8, Method 1668A.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

SGS AXYS METHOD MLA-010 Rev 12

Form 6A
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019

VER Data Filename: PB9C_028 S: 9

Instrument ID: HR GC/MS

Analysis Date: 31-Jan-2019

GC Column ID: SPB OCTYL

Analysis Time: 04:10:02

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | RETENTION TIME REFERENCE | IUPAC NO. ² | RRT | RRT QC LIMITS |
|-------------------------------|-----------|-------------|-----------------------|---|------------------------|-------|---------------|
| 2-MoCB | 1 | | | 13C12-2-MoCB | 1L | 1.001 | 0.999-1.004 |
| 4-MoCB | 3 | | | 13C12-4-MoCB | 3L | 1.001 | 0.999-1.004 |
| 2,2'-DiCB | 4 | | | 13C12-2,2'-DiCB | 4L | 1.001 | 0.999-1.004 |
| 4,4'-DiCB | 15 | | | 13C12-4,4'-DiCB | 15L | 1.001 | 0.999-1.002 |
| 2,2',6-TriCB | 19 | | | 13C12-2,2',6-TriCB | 19L | 1.001 | 0.999-1.003 |
| 3,4,4'-TriCB | 37 | | | 13C12-3,4,4'-TriCB | 37L | 1.001 | 0.999-1.002 |
| 2,2',6,6'-TeCB | 54 | | | 13C12-2,2',6,6'-TeCB | 54L | 1.001 | 0.999-1.002 |
| 3,3',4,4'-TeCB | 77 | | | 13C12-3,3',4,4'-TeCB | 77L | 1.000 | 1.000-1.001 |
| 3,4,4',5-TeCB | 81 | | | 13C12-3,4,4',5-TeCB | 81L | 1.000 | 1.000-1.001 |
| 2,2',4,6,6'-PeCB | 104 | | | 13C12-2,2',4,6,6'-PeCB | 104L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4'-PeCB | 105 | | | 13C12-2,3,3',4,4'-PeCB | 105L | 1.000 | 1.000-1.001 |
| 2,3,4,4',5-PeCB | 114 | | | 13C12-2,3,4,4',5-PeCB | 114L | 1.000 | 1.000-1.001 |
| 2,3',4,4',5-PeCB | 118 | | | 13C12-2,3',4,4',5-PeCB | 118L | 1.000 | 1.000-1.001 |
| 2',3,4,4',5-PeCB | 123 | | | 13C12-2',3,4,4',5-PeCB | 123L | 1.000 | 1.000-1.001 |
| 3,3',4,4',5-PeCB | 126 | | | 13C12-3,3',4,4',5-PeCB | 126L | 1.000 | 1.000-1.001 |
| 2,2',4,4',6,6'-HxCB | 155 | | | 13C12-2,2',4,4',6,6'-HxCB | 155L | 1.001 | 0.999-1.002 |
| 2,3,3',4,4',5-HxCB | 156 | 156 + 157 | C | 13C12-2,3,3',4,4',5-HxCB and 13C12-2,3,3',4,4',5'-HxCB | 156L/157L | 1.000 | 0.998-1.003 |
| 2,3,3',4,4',5'-HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3',4,4',5,5'-HxCB | 167 | | | 13C12-2,3',4,4',5,5'-HxCB | 167L | 1.001 | 1.000-1.001 |
| 3,3',4,4',5,5'-HxCB | 169 | | | 13C12-3,3',4,4',5,5'-HxCB | 169L | 1.000 | 1.000-1.001 |
| 2,2',3,4',5,6,6'-HpCB | 188 | | | 13C12-2,2',3,4',5,6,6'-HpCB | 188L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5'-HpCB | 189 | | | 13C12-2,3,3',4,4',5,5'-HpCB | 189L | 1.001 | 1.000-1.001 |
| 2,2',3,3',5,5',6,6'-OcCB | 202 | | | 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | 1.000 | 1.000-1.001 |
| 2,3,3',4,4',5,5',6-OcCB | 205 | | | 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6-NoCB | 206 | | | 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,5,5',6,6'-NoCB | 208 | | | 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | 1.001 | 1.000-1.001 |
| 2,2',3,3',4,4',5,5',6,6'-DeCB | 209 | | | 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | 1.000 | 1.000-1.001 |

(1) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

(2) Suffix "L" indicates labeled compound

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

For Axy Internal Use Only [XSL Template: Form16686A.xsl; Created: 04-Feb-2019 14:14:50; Application: XMLTransformer-1.17.5;
Report Filename: 1668_PCB1668_PB9C_028S9_Form6A_SJ2506492.html; Workgroup: WG66481; Design ID: 3360]

SGS AXYS METHOD MLA-010 Rev 12

Form 6B
PCB CONGENER RELATIVE RETENTION TIMES

SGS AXYS ANALYTICAL SERVICES

2045 MILLS RD., SIDNEY, B.C., CANADA
V8L 5X2 TEL (250) 655-5800 FAX (250) 655-5811

Initial Calibration Date: 15-Jan-2019 VER Data Filename: PB9C_028 S: 9

Instrument ID: HR GC/MS Analysis Date: 31-Jan-2019

GC Column ID: SPB OCTYL Analysis Time: 04:10:02

| LABELLED COMPOUND | IUPAC NO. ¹ | CO-ELUTIONS | LAB FLAG ² | RETENTION TIME REFERENCE | IUPAC NO. ¹ | RRT | RRT QC LIMITS |
|-------------------------------------|------------------------|-------------|-----------------------|--------------------------------|------------------------|-------|---------------|
| 13C12-2-MoCB | 1L | | | 13C12-2,5-DiCB | 9L | 0.718 | 0.686-0.749 |
| 13C12-4-MoCB | 3L | | | 13C12-2,5-DiCB | 9L | 0.856 | 0.825-0.888 |
| 13C12-2,2'-DiCB | 4L | | | 13C12-2,5-DiCB | 9L | 0.872 | 0.841-0.903 |
| 13C12-4,4'-DiCB | 15L | | | 13C12-2,5-DiCB | 9L | 1.252 | 1.221-1.283 |
| 13C12-2,2',6-TriCB | 19L | | | 13C12-2,5-DiCB | 9L | 1.072 | 1.041-1.103 |
| 13C12-3,4,4'-TriCB | 37L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.090 | 1.070-1.110 |
| 13C12-2,2',6,6'-TeCB | 54L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.810 | 0.797-0.823 |
| 13C12-3,3',4,4'-TeCB | 77L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.395 | 1.382-1.408 |
| 13C12-3,4,4',5-TeCB | 81L | | | 13C12-2,2',5,5'-TeCB | 52L | 1.372 | 1.359-1.385 |
| 13C12-2,2',4,6,6'-PeCB | 104L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 0.808 | 0.798-0.819 |
| 13C12-2,3,3',4,4'-PeCB | 105L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.199 | 1.189-1.210 |
| 13C12-2,3,4,4',5-PeCB | 114L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.179 | 1.168-1.189 |
| 13C12-2,3',4,4',5-PeCB | 118L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.162 | 1.151-1.172 |
| 13C12-2',3,4,4',5-PeCB | 123L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.151 | 1.140-1.161 |
| 13C12-3,3',4,4',5-PeCB | 126L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.300 | 1.289-1.310 |
| 13C12-2,2',4,4',6,6'-HxCB | 155L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 0.787 | 0.778-0.795 |
| 13C12-2,3,3',4,4',5-HxCB | 156L | 156L + 157L | C | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.107 | 1.099-1.115 |
| 13C12-2,3,3',4,4',5'-HxCB | 157L | 156L + 157L | C156L | | | | |
| 13C12-2,3',4,4',5,5'-HxCB | 167L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.077 | 1.069-1.085 |
| 13C12-3,3',4,4',5,5'-HxCB | 169L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.190 | 1.182-1.198 |
| 13C12-2,2',3,4',5,6,6'-HpCB | 188L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.712 | 0.706-0.719 |
| 13C12-2,3,3',4,4',5,5'-HpCB | 189L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.958 | 0.952-0.965 |
| 13C12-2,2',3,3',5,5',6,6'-OcCB | 202L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.818 | 0.811-0.824 |
| 13C12-2,3,3',4,4',5,5',6-OcCB | 205L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.009 | 1.000-1.018 |
| 13C12-2,2',3,3',4,4',5,5',6-NoCB | 206L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.043 | 1.034-1.053 |
| 13C12-2,2',3,3',4,5,5',6,6'-NoCB | 208L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 0.949 | 0.943-0.955 |
| 13C12-2,2',3,3',4,4',5,5',6,6'-DeCB | 209L | | | 13C12-2,2',3,3',4,4',5,5'-OcCB | 194L | 1.075 | 1.066-1.085 |

CLEANUP STANDARD

| | | | | | | | |
|-----------------------------|------|--|--|---------------------------|------|-------|-------------|
| 13C12-2,4,4'-TriCB | 28L | | | 13C12-2,2',5,5'-TeCB | 52L | 0.924 | 0.910-0.937 |
| 13C12-2,3,3',5,5'-PeCB | 111L | | | 13C12-2,2',4,5,5'-PeCB | 101L | 1.088 | 1.077-1.098 |
| 13C12-2,2',3,3',5,5',6-HpCB | 178L | | | 13C12-2,2',3,4,4',5'-HxCB | 138L | 1.012 | 1.004-1.020 |

(1) Suffix "L" indicates labeled compound

(2) Where applicable, custom lab flags have been used on this report; C = co-eluting congener.

These data are validated and reported as accurate and in accord with SGS AXYS Analytical Services Ltd. ISO17025 compliant quality assurance processes.

Signed: _____ Eleanor Andaya _____

Accreditation Scope

SGS AXYS Analytical Services Ltd.
file ref.: ACC-101 Rev. 41

| Compound Class | Compound | Accredited Method ID | SGS AXYS Method ID | Serum | | | | | | | | | | Tissue | Urine | | Water | | Water, Non-Potable | | | | | | | | |
|----------------|---|----------------------|--------------------|-------|----------------|-------------|---------------|----------------|--------------|--------------|---------------|-----------|----------------|--------|-------------|------|----------------|-------------|--------------------|---------------|----------------|--------------|--------------|-----------------|-----------|------------------|----------------|
| | | | | CALA | California DPH | Florida DOH | Minnesota DOH | New Jersey DEP | New York DOH | Virginia DGS | Washington DE | Maine DOH | ANAB ISO 17025 | | ANAB DoD ** | CALA | California DPH | Florida DOH | | Minnesota DOH | New Jersey DEP | New York DOH | Virginia DGS | Washington DE * | Maine DOH | Pennsylvania DEP | ANAB ISO 17025 |
| | Decenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | | | | | | | | | | | | |
| | deoxycholic acid | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | Y | | | | | | | | | | | | |
| | docosaheptaenoic acid (DHA) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | docosatetraenoic acid (adrenic acid) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | Dodecanediolcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Dodecanoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Dodecenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Dopamine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | eicosapentaenoic acid (EPA) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | Eicosatetraenoic acid (arachidonic acid) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | eicosatrienoic acid (dihomo-γ-linolenic acid) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | Glutaconylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Glutamate | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Glutamine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Glutaryl carnitine (Hydroxyhexanoylcarnitine) | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Glycine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | glycochenodeoxycholic acid | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | glycocholic acid | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | glycodeoxycholic acid | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hexadecadienylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | hexadecanoic acid (palmitic acid) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | Hexadecanoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | hexadecenoic acid (palmitoleic acid) | SGS AXYS MLM-001 | MLM-001 | | | | | | | | | | | Y | | | | | | | | | | | | | |
| | Hexadecenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hexanoylcarnitine (Fumaryl carnitine) | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hexenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hexose (sum isomers) | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Histamine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Histidine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyhexadecadienylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyhexadecanoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyhexadecenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxybutyrylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyoctadecenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyproline | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxypropionylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxy sphingomyeline C14:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxy sphingomyeline C16:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxy sphingomyeline C22:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxy sphingomyeline C22:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxy sphingomyeline C24:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxytetradecadienylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxytetradecenoylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Hydroxyvaleryl carnitine (Methylmalonylcarnitine) | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Isoleucine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Kynurenine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Leucine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lithocholic acid | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | Lysine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C14:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C16:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C16:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C17:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C18:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C18:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C18:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C20:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |
| | lysoPhosphatidylcholine acyl C20:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | Y | | Y | | | | | | | | | | | |

Accreditation Scope

SGS AXYS Analytical Services Ltd.
file ref.: ACC-101 Rev. 41

| Compound Class | Compound | Accredited Method ID | SGS AXYS Method ID | Serum | | Solids | | Tissue | | Urine | | Water | | Water, Non-Potable | |
|----------------|----------------------------------|----------------------|--------------------|-------|---|--------|---|--------|---|-------|---|-------|---|--------------------|---|
| | | | | CALA | Y | CALA | Y | CALA | Y | CALA | Y | CALA | Y | CALA | Y |
| | Phosphatidylcholine diacyl C28:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C30:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C30:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C32:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C32:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C32:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C32:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C34:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C34:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C34:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C34:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:5 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C36:6 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:5 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C38:6 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:5 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C40:6 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:4 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:5 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Phosphatidylcholine diacyl C42:6 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Pimelylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Proline | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Propionylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Propionylcarnitine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Putrescine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sarcosine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Serine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Serotonin | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Spermidine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Spermine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C16:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C16:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C18:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C18:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C20:2 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C22:3 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C24:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C24:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C26:0 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Sphingomyeline C26:1 | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Symmetric dimethylarginine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |
| | Taurine | SGS AXYS MLM-001 | MLM-001 | Y | | | | | | | | | | | |

Legend

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|------------------|--|
| Y | Accreditation scope |
| BFR | Brominated flame retardants (non-PBDPE) |
| BPA and mPE | Bisphenol A and mono-Phthalate Esters |
| HBCDD | Hexabromocyclododecane |
| OC Pesticides | Organochlorine Pesticides |
| PAH | Polycyclic Aromatic Hydrocarbons |
| PBDPE | Polybrominated diphenylethers |
| PCB | Polychlorinated Biphenyls |
| PCDDF | Polychlorinated dibenzodioxins/furans |
| PFAS | Per- and Polyfluoroalkyl Substances |
| PPCP | Pharmaceutical and Personal Care Products |
| TBBPA | Tetrabromobisphenol A |
| TOP | Total Oxidizable Precursors |
| California DPH | California Department of Public Health, Lab ID 2911 |
| Florida DOH | Florida Department of Health, Lab ID E871007, (NELAC Standard) |
| Pennsylvania DEP | Pennsylvania Department of Environmental Protection |
| Minnesota DOH | Minnesota Department of Health, Lab ID 232-999-430, (NELAC Standard) |
| New Jersey DEP | New Jersey Department of Environmental Protection, Lab ID CANA005, (NELAC Standard) |
| New York DOH | New York Department of Health, Lab ID 11674, (NELAC Standard) |
| Washington DE | Washington Department of Ecology, Lab ID C404 |
| Virginia DGS | Virginia Department of General Services, Division of Consolidated Laboratory Services, Lab ID 460224, (NELAC Standard) |
| Maine DOH | Maine Center for Disease Control and Prevention, Department of Health and Human Services, Lab ID CN00003 |

ANAB DoD ANSI-ASQ National Accreditation Board, certificate ADE-1861, (US DoD QSM 5.1 Standard)



CALA Canadian Association for Laboratory Accreditation Inc., Lab ID A2637, (ISO/IEC 17025:2005 Standard)



ANAB ISO 17025 ANSI-ASQ National Accreditation Board, certificate ADE-1861.01, (ISO/IEC 17025:2005 Standard)

