

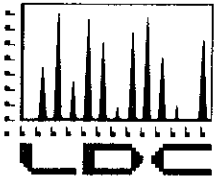
APPENDIX D: DATA VALIDATION REPORTS

ATTACHMENT D-1: SIM ANALYSES

ATTACHMENT D-2: TISSUE CHEMISTRY

ATTACHMENT D-3: SEDIMENT CHEMISTRY

Attachment D-1: SIM Analyses



LABORATORY DATA CONSULTANTS, INC.

7750 El Camino Real, Suite 2L Carlsbad, CA 92009 Phone: 760/634-0437 Fax: 760/634-0439

LDC #13316
April 13, 2005

Windward Environmental, LLC
200 West Mercer Street, Suite 401
Seattle, WA 98119
ATTN: Ms. Susie McGroddy

SUBJECT: Lower Duwamish Waterway Group Sediment Sample Data Validation

Dear Ms. McGroddy,

Enclosed is our EPA Level II data validation of analytical chemistry results generated in support of the Lower Duwamish Waterway Group project. The analyses were performed by Analytical Resources, Inc. Samples were analyzed for GC/MS Semivolatiles by EPA SW 846 Method 8270C using Selected Ion Monitoring (SIM) for Polynuclear Aromatic Hydrocarbons. Samples are referenced under the following Sample Delivery Group: HV45. See the Sample Analysis Table (Attachment 1) for the number of samples reviewed.

Please feel free to contact us if you have any questions.

Sincerely,

Stella S. Cuenco
Project Manager/Senior Chemist

B9a

CHEMICAL DATA QUALITY REVIEW FOR BENTHIC SEDIMENT SAMPLES

Lower Duwamish Waterway Group LDC# 13316

This report details the findings of an EPA Level II data validation review of Analytical chemistry results generated in support of the Lower Duwamish Waterway Group project. The analyses were performed by Analytical Resources, Inc. Samples were analyzed for GC/MS Semivolatiles by EPA SW 846 Method 8270C using Selected Ion Monitoring (SIM) for Polynuclear Aromatic Hydrocarbons. Samples are referenced under the following Sample Delivery Group: HV45. See the Sample Analysis Table (Attachment 1) for the number of samples reviewed and the Sample Validation Table (Attachment 2) for the sample identifications and analyses.

The QC guidelines used for data qualification are those specified in the National Functional Guidelines for Organic Data Review (October 1999). Specific QC criteria used follows the Final Benthic Invertebrate Sampling of the Lower Duwamish Waterway Quality Assurance Project Plan (July 30, 2004). Where specific guidance is not available, the data has been evaluated in a conservative manner consistent with industry standards using professional experience.

The following items were evaluated during the review:

- Holding Times
- Sample Preservation
- Cooler Temperatures
- Instrument Calibration*
- Blanks
- Matrix Spike/Matrix Spike Duplicates
- Internal Standards
- Laboratory Control Samples
- Target Compound Identifications*
- Compound Quantitation and CRQLs*
- System Performance
- Field Duplicates

*Data were not reviewed for Level II.

Attachment 2

SDG#: HV45

VALIDATION SAMPLE TABLE

LDC#: 13316A

Project Name: Lower Duwamish Waterway Group

Parameters/Analytical Method

Project #04-08-06-24

| Client ID # | Lab ID # | Matrix | Date Collected | SVOA (8270C -SIM) | | | | | | | | | | | | | | |
|--------------|----------|----------|----------------|-------------------|--|--|--|--|--|--|--|--|--|--|--|--|--|--|
| LDW-C2-S2 | HV45A | sediment | 08/26/04 | X | | | | | | | | | | | | | | |
| LDW-C3-S1 | HV45B | sediment | 08/29/04 | X | | | | | | | | | | | | | | |
| LDW-C3-S2 | HV45C | sediment | 08/29/04 | X | | | | | | | | | | | | | | |
| LDW-C4-S | HV45D | sediment | 08/27/04 | X | | | | | | | | | | | | | | |
| LDW-C5-S | HV45E | sediment | 08/27/04 | X | | | | | | | | | | | | | | |
| LDW-C6-S | HV45F | sediment | 08/25/04 | X | | | | | | | | | | | | | | |
| LDW-C9-S | HV45G | sediment | 08/15/04 | X | | | | | | | | | | | | | | |
| LDW-B1a-S | HV45H | sediment | 08/12/04 | X | | | | | | | | | | | | | | |
| LDW-2Ba-S | HV45I | sediment | 08/13/04 | X | | | | | | | | | | | | | | |
| LDW-B3a-S | HV45J | sediment | 08/26/04 | X | | | | | | | | | | | | | | |
| LDW-B10b-S | HV45K | sediment | 08/19/04 | X | | | | | | | | | | | | | | |
| LDW-B1b-S | HV45L | sediment | 09/27/04 | X | | | | | | | | | | | | | | |
| LDW-B8b-S | HV45M | sediment | 08/19/04 | X | | | | | | | | | | | | | | |
| LDW-B9b-S | HV45N | sediment | 08/11/04 | X | | | | | | | | | | | | | | |
| LDW-B10a-S | HV45O | sediment | 08/26/04 | X | | | | | | | | | | | | | | |
| LDW-B1a-SMS | HV45HMS | sediment | 08/12/04 | X | | | | | | | | | | | | | | |
| LDW-B1a-SMSD | HV45HMSD | sediment | 08/12/04 | X | | | | | | | | | | | | | | |
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Note: X = Validation was performed.

Only issues which require comment or action are discussed in this report. Data deficiencies are arranged by method. Potential effects of data anomalies have been described where possible.

The following are definitions of the data qualifiers:

- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- J Indicates an estimated value.
- R Quality control indicates the data is not usable.
- N Presumptive evidence of presence of the constituent.
- UJ Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$ Where: ave. RRF = initial calibration average RRF
 RRF = $(A_x)(C_s) / (A_s)(C_x)$ RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

| # | Standard ID | Calibration Date | Compound (Reference Internal Standard) | Average RRF (Initial) | Reported | | Recalculated | |
|---|---------------------|------------------|--|-----------------------|---------------|------|---------------|----|
| | | | | | RRF Conc (CC) | %D | RRF Conc (CC) | %D |
| 1 | PB5C-084 | 7/17/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 52.1 | 52.3 | 4.5 | |
| | PB5C-084 | | 0.83 | 54.0 | 54.0 | 7.8 | | |
| | S-1 | | 0.98 | 100 | 101 | 0.8 | | |
| | | | 0.79 | 52.9 | 53.2 | 6.4 | | |
| 2 | PB5C-084 | 12/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 47.4 | 47.3 | 6.38 | |
| | | | 0.83 | 50.1 | 50.3 | 0.6 | | |
| | | | 0.98 | 97.3 | 97.7 | 2.3 | | |
| | | | 0.79 | 51.5 | 51.8 | 3.60 | | |
| 3 | | | PCB-77 (¹³ C-PCB-77) | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | |

Comments: Refer to Routine Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: B238A3
 SDG #: DPWG 15-17

VALIDATION FINDINGS WORKSHEET

Laboratory Control Sample Results Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: 0

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 * SSC/SA

Where: SSC = Spiked sample concentration
 SA = Spike added

RPD = |LCS - LCSD| * 2 / (LCS + LCSD)

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS ID: OPR

| Compound | Spike Added (NS/M) | | Spiked Sample Concentration (NS/L) | | LCS | | LCSD | | LCS/LCSD | |
|--------------------|--------------------|------|------------------------------------|------|------------------|---------|------------------|---------|----------|--------------|
| | LCS | LCSD | LCS | LCSD | Percent Recovery | | Percent Recovery | | RPD | |
| | | | | | Reported | Recalc. | Reported | Recalc. | Reported | Recalculated |
| PCB-77 | 50 | NA | 47.4 | NA | 94.9 | 94.8 | | | | |
| PCB-81 | ↓ | | 47.2 | | 94.4 | 94.4 | | | | |
| PCB-105 | | | 51.2 | | 102 | 102 | | | | |
| PCB-114 | | | 51.0 | | 102 | 102 | | | | |
| PCB-118 | | | 52.6 | | 105 | 105 | | | | |
| PCB-123 | | | 51.3 | | 103 | 103 | | | | |
| PCB-126 | ↓ | | 50.8 | | 102 | 102 | | | | |
| PCB-156 / PCBIST | 100 | | 97.1 | | 97.1 | 97.1 | | | | |
| PCB-167 | | | | | | | | | | |
| PCB-167 | 50 | | 49.0 | | 97.9 | 98 | | | | |
| PCB-169 | ↓ | | 47.6 | | 95.3 | 95.3 | | | | |
| PCB-170 | | | | | | | | | | |
| PCB-180 | | | | | | | | | | |
| PCB-189 | 50 | ↓ | 51.3 | ↓ | 103 | 103 | | | | |
| | | | | | | | | | | |
| | | | | | | | | | | |

Comments: Refer to Laboratory Control Sample findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

Ions Monitored for HRGC/HRMS Analysis of Polychlorinated Biphenyls

| Descriptor | Accurate mass ^(a) | Ion ID | Analyte | Substance |
|------------|--|--|---|---|
| 1 | 289.9224 291.9194 301.9626 303.9597 325.8804 327.8775 [292.9825] | M M+2 M M+2 M+2 M+4 Lock | C12 H6 35Cl4 C12 H6 35Cl3 37Cl4 13C12 H6 35Cl4 13C12 H6 35Cl3 37Cl C12 H5 35Cl4 37Cl C12 H5 35Cl3 37Cl2 C7 F11 | TCB TCB PeCB PeCB PeCB PeCB PFK |
| 2 | 325.8804 327.8775 337.9207 339.9178 359.8415 361.8385 371.8817 373.8788 393.8025 395.7996 405.8428 407.8398 [354.9892] | M+2 M+4 M+2 M+4 M+2 M+4 M+2 M+4 M+2 M+4 M+2 M+4 Lock | C12 H5 35Cl4 37Cl C12 H5 35Cl3 37Cl2 13C12 H5 35Cl4 37Cl 13C12 H5 35Cl3 37Cl2 C12 H4 35Cl5 37Cl C12 H4 35Cl4 37Cl2 13C12 H4 35Cl5 37Cl 13C12 H4 35Cl4 37Cl2 C12 H3 35Cl6 37Cl C12 H3 35Cl5 37Cl2 13C12 H3 35Cl6 37Cl 13C12 H3 35Cl5 37Cl2 C9F13 | PeCB PeCB PeCB PeCB HxCB HxCB HxCB HxCB HpCB HpCB HpCB HpCB PFK |
| 3 | 509.7229 511.7199 513.7170 [442.9728] | M+4 M+6 M+8 Lock | 13C12 35Cl10 37Cl2 13C12 35Cl9 37Cl3 13C12 35Cl8 37Cl4 C10 F17 | DCB PFK |

S = internal/recovery standard

H = 1.007825
C = 12.000000
¹³C = 13.003355
F = 18.9984

³⁵Cl = 34.968853
³⁷Cl = 36.965903

GC/MS Semivolatiles by EPA SW 846 Method 8270C using Selected Ion Monitoring (SIM)

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. GC/MS Instrument Performance Check

Instrument performance check data were not reviewed for Level II.

III. Initial Calibration

Initial calibration data were not reviewed for Level II.

IV. Continuing Calibration

Continuing calibration data were not reviewed for Level II.

V. Blanks

Method blanks were reviewed for each matrix as applicable. No semivolatile contaminants were found in the method blanks with the following exceptions:

| Method Blank ID | Extraction Date | Compound TIC (RT in minutes) | Concentration | Associated Samples |
|-----------------|-----------------|--|-----------------------|-------------------------|
| MB-031705 | 3/17/05 | Diethylphthalate N-Nitrosodiphenylamine | 15 ug/Kg 6.7 ug/Kg | All samples in SDG HV45 |

Sample concentrations were compared to concentrations detected in the method blanks. The sample concentrations were either not detected or were significantly greater (>10X for common contaminants, >5X for other contaminants) than the concentrations found in the associated method blanks with the following exceptions:

| Associated SDG | Sample | Compound TIC (RT in minutes) | Reported Concentration | Modified Final Concentration |
|----------------|-----------|---------------------------------|------------------------|------------------------------|
| HV45 | LDW-C2-S2 | Diethylphthalate | 6.5 ug/Kg | 6.5U ug/Kg |
| HV45 | LDW-B9b-S | Diethylphthalate | 6.6 ug/Kg | 6.6U ug/Kg |

| Associated SDG | Sample | Compound TIC (RT in minutes) | Reported Concentration | Modified Final Concentration |
|----------------|------------|---------------------------------|------------------------|------------------------------|
| HV45 | LDW-B10a-S | Diethylphthalate | 9.6 ug/Kg | 9.6U ug/Kg |

VI. Surrogate Spikes

Surrogates were added to all samples and blanks as required by the method. All surrogate recoveries (%R) were within QC limits with the following exceptions:

| Associated SDG | Sample | Surrogate | %R (Limits) | Compound | Flag | A or P |
|----------------|-----------|---|--|-------------------|---|--------|
| HV45 | LDW-B9b-S | 2-Fluorobiphenyl 2-Fluorophenol 1,2-Dichlorobenzene-d4 Phenol-d5 2-Chlorophenol-d4 Nitrobenzene-d5 | 37.2 (40-130) 32.5 (40-130) 24.8 (40-130) 33.1 (40-130) 32.5 (40-130) 30.4 (40-130) | All TCL compounds | J (all detects) UJ (all non-detects) | P |

VII. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) samples were reviewed for each matrix as applicable. Percent recoveries (%R) and relative percent differences (RPD) were within QC limits.

VIII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. Percent recoveries (%R) were within QC limits.

Standard reference material was performed at the required frequencies.

IX. Regional Quality Assurance and Quality Control

Not applicable.

X. Internal Standards

Internal standards data were not reviewed for Level II.

XI. Target Compound Identifications

Raw data were not reviewed for this SDG.

XII. Compound Quantitation and CRQLs

Raw data were not reviewed for this SDG.

XIII. Tentatively Identified Compounds (TICs)

Raw data were not reviewed for this SDG.

XIV. System Performance

Raw data were not reviewed for this SDG.

XV. Overall Assessment

Data flags have been summarized at the end of the report.

XVI. Field Duplicates

No field duplicates were identified in this SDG.

XVII. Field Blanks

No field blanks were identified in this SDG.

LDC #: 13316A2b **VALIDATION COMPLETENESS WORKSHEET**

SDG #: HV45 Level II

Laboratory: Analytical Resources, Inc.

Date: 3/29/05

Page: 1 of 1

Reviewer: [Signature]

2nd Reviewer: [Signature]

METHOD: GC/MS Polynuclear Aromatic Hydrocarbons (EPA SW 846 Method 8270C-SIM)

The samples listed below were reviewed for each of the following validation areas. Validation findings are noted in attached validation findings worksheets.

| | Validation Area | | Comments |
|-------|--|----|--|
| I. | Technical holding times | A | Sampling dates: 8/11 8/11 - 9/27/04 |
| II. | GC/MS Instrument performance check | N | |
| III. | Initial calibration | N | |
| IV. | Continuing calibration | N | |
| V. | Blanks | TW | |
| VI. | Surrogate spikes | TW | |
| VII. | Matrix spike/Matrix spike duplicates | A | |
| VIII. | Laboratory control samples | A | LCS, SRM |
| IX. | Regional Quality Assurance and Quality Control | N | |
| X. | Internal standards | N | |
| XI. | Target compound identification | N | |
| XII. | Compound quantitation/CRQLs | N | |
| XIII. | Tentitatively identified compounds (TICs) | N | |
| XIV. | System performance | N | |
| XV. | Overall assessment of data | A | |
| XVI. | Field duplicates | N | |
| XVII. | Field blanks | N | |

Note: A = Acceptable ND = No compounds detected D = Duplicate
 N = Not provided/applicable R = Rinsate TB = Trip blank
 SW = See worksheet FB = Field blank EB = Equipment blank

Validated Samples

MSLs

| | | | | | | | |
|----|-----------|----|--------------|----|-----------|----|--|
| 1 | LDW-C2-S2 | 11 | LDW-B10b-S | 21 | MB-031705 | 31 | |
| 2 | LDW-C3-S1 | 12 | LDW-B1b-S | 22 | | 32 | |
| 3 | LDW-C3-S2 | 13 | LDW-B8b-S | 23 | | 33 | |
| 4 | LDW-C4-S | 14 | LDW-B9b-S | 24 | | 34 | |
| 5 | LDW-C5-S | 15 | LDW-B10a-S | 25 | | 35 | |
| 6 | LDW-C6-S | 16 | LDW-B1a-SMS | 26 | | 36 | |
| 7 | LDW-C9-S | 17 | LDW-B1a-SMSD | 27 | | 37 | |
| 8 | LDW-B1a-S | 18 | | 28 | | 38 | |
| 9 | LDW-2Ba-S | 19 | | 29 | | 39 | |
| 10 | LDW-B3a-S | 20 | | 30 | | 40 | |

VALIDATION FINDINGS WORKSHEET

METHOD: GC/MS BNA (EPA SW 846 Method 8270)

| | | | | |
|---------------------------------|-------------------------------|----------------------------------|---------------------------------|----------------------------------|
| A. Phenol** | P. Bis(2-chloroethoxy)methane | EE. 2,6-Dinitrotoluene | TT. Pentachlorophenol** | III. Benzo(a)pyrene** |
| B. Bis (2-chloroethyl) ether | Q. 2,4-Dichlorophenol** | FF. 3-Nitroaniline | UU. Phenanthrene | JJJ. Indeno(1,2,3-cd)pyrene |
| C. 2-Chlorophenol | R. 1,2,4-Trichlorobenzene | GG. Acenaphthene** | VV. Anthracene | KKK. Dibenz(a,h)anthracene |
| D. 1,3-Dichlorobenzene | S. Naphthalene | HH. 2,4-Dinitrophenol* | WW. Carbazole | LLL. Benzo(g,h,i)perylene |
| E. 1,4-Dichlorobenzene** | T. 4-Chloroaniline | II. 4-Nitrophenol* | XX. Di-n-butylphthalate | MMM. Bis(2-Chloroisopropyl)ether |
| F. 1,2-Dichlorobenzene | U. Hexachlorobutadiene** | JJ. Dibenzofuran | YY. Fluoranthene** | NNN. Aniline |
| G. 2-Methylphenol | V. 4-Chloro-3-methylphenol** | KK. 2,4-Dinitrotoluene | ZZ. Pyrene | OOO. N-Nitrosodimethylamine |
| H. 2,2'-Oxybis(1-chloropropane) | W. 2-Methylnaphthalene | LL. Diethylphthalate | AAA. Butylbenzylphthalate | PPP. Benzoic Acid |
| I. 4-Methylphenol | X. Hexachlorocyclopentadiene* | MM. 4-Chlorophenyl-phenyl ether | BBB. 3,3'-Dichlorobenzidine | QQQ. Benzyl alcohol |
| J. N-Nitroso-di-n-propylamine* | Y. 2,4,6-Trichlorophenol** | NN. Fluorene | CCC. Benzo(a)anthracene | RRR. Pyridine |
| K. Hexachloroethane | Z. 2,4,5-Trichlorophenol | OO. 4-Nitroaniline | DDD. Chrysene | SSS. Benzidine |
| L. Nitrobenzene | AA. 2-Chloronaphthalene | PP. 4,6-Dinitro-2-methylphenol | EEE. Bis(2-ethylhexyl)phthalate | TTT. |
| M. Isophorone | BB. 2-Nitroaniline | QQ. N-Nitrosodiphenylamine (1)** | FFF. Di-n-octylphthalate** | UUU. |
| N. 2-Nitrophenol** | CC. Dimethylphthalate | RR. 4-Bromophenyl-phenylether | GGG. Benzo(b)fluoranthene | VVV. |
| O. 2,4-Dimethylphenol | DD. Acenaphthylene | SS. Hexachlorobenzene | HHH. Benzo(k)fluoranthene | WWW. |

LDC #: 133160-6
 SDG #: HV45

VALIDATION FINDINGS WORKSHEET

Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: GC/MS BNA (EPA SW 846 Method 8270)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Was a method blank analyzed for each matrix?
- N N/A Was a method blank analyzed for each concentration preparation level?
- N N/A Was a method blank associated with every sample?
- N N/A Was the blank contaminated? If yes, please see qualification below.

Blank extraction date: 3/17/05 Blank analysis date: 3/18/05

Conc. units: µg/kg

Associated Samples: [Signature]

| Compound | Blank ID | Sample Identification | | | | | | | |
|-----------|---------------|-----------------------|--------------|--------------|--|--|--|--|--|
| | | 1 | 14 | 15 | | | | | |
| <u>MB</u> | <u>031705</u> | | | | | | | | |
| <u>2L</u> | <u>15</u> | <u>6.5/U</u> | <u>6.6/U</u> | <u>9.6/U</u> | | | | | |
| <u>QR</u> | <u>6.7</u> | | | | | | | | |
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| | | | | | | | | | |

Blank extraction date: _____ Blank analysis date: _____

Conc. units: _____

Associated Samples: _____

| Compound | Blank ID | Sample Identification | | | | | | | |
|----------|----------|-----------------------|--|--|--|--|--|--|--|
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CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:

Common contaminants such as the phthalates and TICs noted above that were detected in samples within ten times the associated method blank concentration were qualified as not detected, "U". Other contaminants within five times the method blank concentration were also qualified as not detected, "U".

LDC #: 13316026
 SDG #: HV45

VALIDATION FINDINGS WORKSHEET
Surrogate Recovery

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: GC/MS BNA (EPA SW 846 Method 8270)

Please see qualification below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y (N) N/A Were percent recoveries (%R) for surrogates within QC limits?
 Y (N) N/A If 2 or more base neutral or acid surrogates were outside QC limits, was a reanalysis performed to confirm %R?
 Y (N) N/A If any %R was less than 10 percent, was a reanalysis performed to confirm %R?

| # | Date | Sample ID | Surrogate | %R (Limits) | Qualifications |
|---|------|-----------|-----------|---------------|----------------|
| | | 1 | DCB | 38.0 (40-130) | No Anal. |
| | | 8 | ↓ | 39.2 () | ↓ |
| | | 1A | FBP | 37.2 () | ↓ N/A ↓ P |
| | | | 2FP | 32.5 () | |
| | | | DCB | 24.8 () | |
| | | | PHL | 33.1 () | |
| | | | 2CH | 32.5 () | |
| | | | NBZ | 30.4 () | ↓ |
| | | | | () | |
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- * QC limits are advisory
- | | | | |
|------------------------------------|-------------------|---|-------------------|
| QC Limits (Soil) | QC Limits (Water) | QC Limits (Soil) | QC Limits (Water) |
| S1 (NBZ) = Nitrobenzene-d5 23-120 | 35-114 | S5 (2FP) = 2-Fluorophenol 25-121 | 21-100 |
| S2 (FBP) = 2-Fluorobiphenyl 30-115 | 43-116 | S6 (TBP) = 2,4,6-Tribromophenol 19-122 | 10-123 |
| S3 (TPH) = Terphenyl-d14 18-137 | 33-141 | S7 (2CP) = 2-Chlorophenol-d4 20-130* | 33-110* |
| S4 (PHL) = Phenol-d5 24-113 | 10-94 | S8 (DCB) = 1,2-Dichlorobenzene-d4 20-130* | 16-110* |

Attachment D-2: Tissue Chemistry

CHEMICAL DATA QUALITY REVIEW FOR BENTHIC TISSUE SAMPLES**Lower Duwamish Waterway Group
LDC# 13238**

This report details the findings of an EPA Level II and Level IV data validation review of Analytical chemistry results generated in support of the Lower Duwamish Waterway Group project. The analyses were performed by Axys Analytical Services, Ltd. Samples were analyzed for HRGC/HRMS Polychlorinated Biphenyl Congeners by modified EPA Method 1668A. Samples are referenced under the following Sample Delivery Group: DPWG15217. See the Sample Analysis Table (Attachment 1) for the number of samples reviewed and the Sample Validation Table (Attachment 2) for the sample identifications and analyses. Sample IDs ending in "***" underwent Level IV review.

The QC guidelines used for data qualification are those specified in the EPA Region 10 SOP for the Validation of 1668 Toxic, Dioxin-like PCB Data (Revision 1.0, December 8, 1995). Specific QC criteria used follows the Final Benthic Invertebrate Sampling of the Lower Duwamish Waterway Quality Assurance Project Plan (July 30, 2004). Where specific guidance is not available, the data has been evaluated in a conservative manner consistent with industry standards using professional experience.

The following items were evaluated during the review:

- Holding Times
- Sample Preservation
- Cooler Temperatures
- Instrument Calibration*
- Blanks
- Matrix Spike/Matrix Spike Duplicates
- Internal Standards
- Laboratory Control Samples
- Target Compound Identifications*
- Compound Quantitation and CRQLs*
- System Performance
- Field Duplicates

*Data were not reviewed for Level II.

Only issues which require comment or action are discussed in this report. Data deficiencies are arranged by method. Potential effects of data anomalies have been described where possible.

The following are definitions of the data qualifiers:

- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- J Indicates an estimated value.
 - J1 Blank Contamination: Indicates possible high bias and/or false positives.
 - J2 Calibration Range exceeded: Indicates possible low bias.
 - J3 Holding times not met: Indicates low bias for most analytes.
 - J4 Other QC parameters outside control limits: bias not readily determined.
 - J5 Other QC parameters outside control limits. The reported results appear to be biased high. The actual value of target compound in the sample may be lower than the value reported by the laboratory.
 - J6 Other QC parameters outside control limits. The reported results appear to be biased low. The actual value of target compound in the sample may be higher than the value reported by the laboratory.
- R Quality control indicates the data is not usable.
- N Presumptive evidence of presence of the constituent.
- UU Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.

***Overall Data Assessment**

Method blank contamination have warranted the qualification of several compounds as non-detected (U).

*The frequency of matrix spike (MS) and matrix spike duplicate (MSD) was not met as required by the QAPP. MS/MSD analyses are not required for EPA Method 1668A. The laboratory consulted with the client on this discrepancy and was instructed to proceed with the extraction and analysis despite the absence of MS/MSDs.

Field duplicates were not collected for this sampling event.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable. Based upon the data validation all other results are considered valid and usable for all purposes.

Attachment 2

| SDG#: DPWG15217 | | VALIDATION SAMPLE TABLE | | | | | | | | | | LDC#: 13238A | | |
|---|-------------|-------------------------|----------------|------------------------------|--|--|--|--|--|--|----------------------|--------------|--|--|
| Project Name: Lower Duwamish Waterway Group | | | | Parameters/Analytical Method | | | | | | | Project #04-08-06-21 | | | |
| Client ID # | Lab ID # | Matrix | Date Collected | PCB Cong. (1668A) | | | | | | | | | | |
| LDW-B1b-T** | L7510-1 | tissue | 08/10/04 | X | | | | | | | | | | |
| LDW-B2a-T | L7510-2 | tissue | 08/14/04 | X | | | | | | | | | | |
| LDW-B3b-T | L7510-3 | tissue | 08/10/04 | X | | | | | | | | | | |
| LDW-B4b-T | L7510-4 | tissue | 08/17/04 | X | | | | | | | | | | |
| LDW-B5a-T | L7510-5 | tissue | 08/22/04 | X | | | | | | | | | | |
| LDW-B8a-T | L7510-6 | tissue | 08/27/04 | X | | | | | | | | | | |
| LDW-B9b-T** | L7510-7 | tissue | 08/11/04 | X | | | | | | | | | | |
| LDW-B10a-T | L7510-8 | tissue | 08/25/04 | X | | | | | | | | | | |
| LDW-C1-T | L7510-9 | tissue | 08/26/04 | X | | | | | | | | | | |
| LDW-C2-2-T | L7510-10 | tissue | 08/26/04 | X | | | | | | | | | | |
| LDW-C4-T | L7510-11 | tissue | 08/27/04 | X | | | | | | | | | | |
| LDW-C6-T | L7510-12 | tissue | 08/26/04 | X | | | | | | | | | | |
| LDW-C7-T1 | L7510-13 | tissue | 08/26/04 | X | | | | | | | | | | |
| LDW-C8-T | L7510-14 | tissue | 08/26/04 | X | | | | | | | | | | |
| LDW-C9-T | L7510-15 | tissue | 08/25/04 | X | | | | | | | | | | |
| LDW-C10-T1 | L7510-16 | tissue | 08/25/04 | X | | | | | | | | | | |
| LDW-C10-T1DUP | L7510-16DUP | tissue | 08/25/04 | X | | | | | | | | | | |
| | | | | | | | | | | | | | | |
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Note: X = Validation was performed.

HRGC/HRMS Polychlorinated Biphenyl Congeners by modified EPA Method 1668A

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. GC/MS Instrument Performance Check

Instrument performance was checked at the required daily frequency.

Retention time windows were established for all congeners. The chromatographic resolution was less than or equal to 40% for congeners PCB-23 and PCB-34 and congeners PCB-182 and PCB-187.

The static resolving power was at least 10,000 (10% valley definition).

Instrument performance check data were not reviewed for Level II.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

The percent relative standard deviations (%RSD) were less than or equal to 20.0% for all compounds.

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound.

Initial calibration data were not reviewed for Level II.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 25.0% for unlabeled compounds and less than or equal to 35.0% for labeled compounds.

Continuing calibration data were not reviewed for Level II.

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated biphenyl contaminants were found in the method blanks with the following exceptions:

| Associated SDG | Method Blank ID | Extraction Date | Compound | Concentration | Associated Samples |
|----------------|-----------------|-----------------|----------|---------------|---------------------------------|
| DPWG15217 | WGK743-101 | 1/27/05 | PCB-1 | 0.215 ng/Kg | All samples in SDG DPWG15217 |
| | | | PCB-2 | 0.245 ng/Kg | |
| | | | PCB-3 | 0.355 ng/Kg | |
| | | | PCB-4 | 0.371 ng/Kg | |
| | | | PCB-6 | 0.180 ng/Kg | |
| | | | PCB-8 | 0.704 ng/Kg | |
| | | | PCB-11 | 1.36 ng/Kg | |
| | | | PCB-15 | 0.294 ng/Kg | |
| | | | PCB-16 | 0.344 ng/Kg | |
| | | | PCB-17 | 0.452 ng/Kg | |
| | | | PCB-18 | 0.812 ng/Kg | |
| | | | PCB-19 | 0.128 ng/Kg | |
| | | | PCB-20 | 0.982 ng/Kg | |
| | | | PCB-21 | 0.450 ng/Kg | |
| | | | PCB-22 | 0.276 ng/Kg | |
| | | | PCB-25 | 0.105 ng/Kg | |
| | | | PCB-26 | 0.251 ng/Kg | |
| | | | PCB-27 | 0.095 ng/Kg | |
| | | | PCB-31 | 0.795 ng/Kg | |
| | | | PCB-32 | 0.264 ng/Kg | |
| | | | PCB-37 | 0.233 ng/Kg | |
| | | | PCB-40 | 0.374 ng/Kg | |
| | | | PCB-42 | 0.166 ng/Kg | |
| | | | PCB-43 | 0.038 ng/Kg | |
| | | | PCB-44 | 0.945 ng/Kg | |
| | | | PCB-45 | 0.170 ng/Kg | |
| | | | PCB-48 | 0.162 ng/Kg | |
| | | | PCB-49 | 0.668 ng/Kg | |
| | | | PCB-50 | 0.166 ng/Kg | |
| | | | PCB-52 | 1.40 ng/Kg | |
| | | | PCB-56 | 0.287 ng/Kg | |
| | | | PCB-59 | 0.097 ng/Kg | |
| | | | PCB-60 | 0.109 ng/Kg | |
| | | | PCB-61 | 1.03 ng/Kg | |
| PCB-64 | 0.309 ng/Kg | | | | |
| PCB-66 | 0.563 ng/Kg | | | | |
| PCB-77 | 0.121 ng/Kg | | | | |
| PCB-83 | 0.698 ng/Kg | | | | |
| PCB-84 | 0.226 ng/Kg | | | | |
| PCB-85 | 0.130 ng/Kg | | | | |
| PCB-86 | 0.695 ng/Kg | | | | |
| PCB-88 | 0.216 ng/Kg | | | | |
| PCB-90 | 1.28 ng/Kg | | | | |
| PCB-92 | 0.154 ng/Kg | | | | |
| PCB-93 | 1.20 ng/Kg | | | | |
| PCB-105 | 0.754 ng/Kg | | | | |
| PCB-110 | 1.29 ng/Kg | | | | |
| PCB-118 | 1.41 ng/Kg | | | | |
| PCB-128 | 0.549 ng/Kg | | | | |
| PCB-129 | 2.43 ng/Kg | | | | |
| PCB-132 | 0.513 ng/Kg | | | | |
| PCB-135 | 0.534 ng/Kg | | | | |
| PCB-136 | 0.167 ng/Kg | | | | |
| PCB-141 | 0.337 ng/Kg | | | | |
| PCB-146 | 0.306 ng/Kg | | | | |
| PCB-147 | 0.997 ng/Kg | | | | |
| PCB-153 | 1.80 ng/Kg | | | | |
| PCB-156 | 0.631 ng/Kg | | | | |
| PCB-158 | 0.250 ng/Kg | | | | |
| PCB-164 | 0.107 ng/Kg | | | | |
| PCB-167 | 0.166 ng/Kg | | | | |
| PCB-170 | 0.676 ng/Kg | | | | |
| PCB-174 | 0.371 ng/Kg | | | | |

*Indicates change as the result of report review.
SDG DPWG15217

| Associated SDG | Method Blank ID | Extraction Date | Compound | Concentration | Associated Samples |
|-----------------------------|-----------------|-----------------|----------------------------|---------------|---------------------------------|
| DPWG15217 | WGK743-101 | 1/27/05 | PCB-177 | 0.271 ng/Kg | All samples in SDG DPWG15217 |
| | | | PCB-179 | 0.133 ng/Kg | |
| | | | PCB-180 | 1.07 ng/Kg | |
| | | | PCB-183 | 0.381 ng/Kg | |
| | | | PCB-187 | 0.393 ng/Kg | |
| | | | PCB-190 | 0.144 ng/Kg | |
| | | | PCB-194 | 0.188 ng/Kg | |
| | | | PCB-195 | 0.111 ng/Kg | |
| | | | PCB-196 | 0.117 ng/Kg | |
| | | | PCB-198 | 0.235 ng/Kg | |
| | | | PCB-201 | 0.023 ng/Kg | |
| | | | PCB-202 | 0.086 ng/Kg | |
| | | | PCB-203 | 0.151 ng/Kg | |
| | | | PCB-204 | 0.012 ng/Kg | |
| | | | PCB-205 | 0.051 ng/Kg | |
| | | | PCB-206 | 0.149 ng/Kg | |
| | | | PCB-208 | 0.133 ng/Kg | |
| | | | PCB-209 | 0.154 ng/Kg | |
| | | | Total Monochloro Biphenyls | 0.460 ng/Kg | |
| | | | Total Dichloro Biphenyls | 1.36 ng/Kg | |
| Total Trichloro Biphenyls | 4.69 ng/Kg | | | | |
| Total Tetrachloro Biphenyls | 5.37 ng/Kg | | | | |
| Total Pentachloro Biphenyls | 6.24 ng/Kg | | | | |
| Total Hexchloro Biphenyls | 7.47 ng/Kg | | | | |
| Total Heptachloro Biphenyls | 2.50 ng/Kg | | | | |
| Total Octachloro Biphenyls | 0.432 ng/Kg | | | | |
| Total Nonachloro Biphenyls | 0.133 ng/Kg | | | | |

Sample concentrations were compared to concentrations detected in the method blanks. The sample concentrations were either not detected or were significantly greater (>5X blank contaminants) than the concentrations found in the associated method blanks with the following exceptions:

| Associated SDG | Sample | Compound | Reported Concentration | Modified Final Concentration |
|----------------|-------------|----------------------------|------------------------|------------------------------|
| DPWG15217 | LDW-B2a-T | PCB-2 | 1.06 ng/Kg | 1.06U ng/Kg |
| | | PCB-3 | 1.60 ng/Kg | 1.60U ng/Kg |
| DPWG15217 | LDW-B8a-T | PCB-2 | 0.895 ng/Kg | 0.895U ng/Kg |
| | | PCB-3 | 1.75 ng/Kg | 1.75U ng/Kg |
| DPWG15217 | LDW-B9b-T** | PCB-2 | 0.259 ng/Kg | 0.259U ng/Kg |
| | | PCB-3 | 0.756 ng/Kg | 0.756U ng/Kg |
| | | PCB-11 | 4.45 ng/Kg | 4.45U ng/Kg |
| | | Total Monochloro Biphenyls | 1.56 ng/Kg | 1.56U ng/Kg |
| DPWG15217 | LDW-B10a-T | PCB-2 | 0.526 ng/Kg | 0.526U ng/Kg |
| | | PCB-3 | 0.442 ng/Kg | 0.442U ng/Kg |
| | | Total Monochloro Biphenyls | 1.35 ng/Kg | 1.35U ng/Kg |
| | | PCB-1 | 0.382 ng/Kg | 0.382U ng/Kg |

*Indicates change as the result of report review.
SDG DPWG15217

| Associated SDG | Sample | Compound | Reported Concentration | Modified Final Concentration |
|----------------|------------|--|--|---|
| DPWG15217 | LDW-C1-T | PCB-2 PCB-3 | 0.311 ng/Kg 0.819 ng/Kg | 0.311U ng/Kg 0.819U ng/Kg |
| DPWG15217 | LDW-C2-2-T | PCB-2 PCB-3 Total Monochloro Biphenyls PCB-1 PCB-204 | 0.293 ng/Kg 0.715 ng/Kg 2.01 ng/Kg 1.00 ng/Kg 0.021 ng/Kg | 0.293U ng/Kg 0.715U ng/Kg 2.01U ng/Kg 1.00U ng/Kg 0.021U ng/Kg |
| DPWG15217 | LDW-C4-T | PCB-2 PCB-3 Total Monochloro Biphenyls PCB-1 PCB-204 | 0.316 ng/Kg 0.711 ng/Kg 2.00 ng/Kg 0.971 ng/Kg 0.049 ng/Kg | 0.316U ng/Kg 0.711U ng/Kg 2.00U ng/Kg 0.971U ng/Kg 0.049U ng/Kg |
| DPWG15217 | LDW-C6-T | PCB-2 PCB-3 | 0.847 ng/Kg 1.23 ng/Kg | 0.847U ng/Kg 1.23U ng/Kg |
| DPWG15217 | LDW-C7-T1 | PCB-2 PCB-3 | 0.665 ng/Kg 1.24 ng/Kg | 0.665U ng/Kg 1.24U ng/Kg |
| DPWG15217 | LDW-C9-T | PCB-2 PCB-3 | 0.310 ng/Kg 0.781 ng/Kg | 0.310U ng/Kg 0.781U ng/Kg |
| DPWG15217 | LDW-C10-T1 | PCB-2 | 0.853 ng/Kg | 0.853 U ng/Kg |

No field blanks were identified in this SDG.

*VI. Matrix Spike/Matrix Spike Duplicates

*Matrix spike (MS) and matrix spike duplicate (MSD) analyses were not required by the method.

Duplicate (DUP) sample analyses were reviewed for each matrix as applicable. Results were within QC limits.

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. The percent recoveries (%R) were within the QC limits.

Standard reference material was performed at the required frequencies.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits.

X. Target Compound Identifications

All target compound identifications were within validation criteria.

Target compound identifications data were not reviewed for Level II.

XI. Compound Quantitation and CRQLs

All compound quantitation and CRQLs were within validation criteria.

Compound quantitation and CRQLs data were not reviewed for Level II.

XII. System Performance

The system performance was acceptable.

System performance data were not reviewed for Level II.

XIII. Overall Assessment of Data

The overall assessment of data was acceptable.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

METHOD: HRGC/HRMS Polychlorinated Biphenyl Congeners (EPA Method 1668A)-Med

The samples listed below were reviewed for each of the following validation areas. Validation findings are noted in attached validation findings worksheets.

| | Validation Area | | Comments |
|-------|--|------------------|---|
| I. | Technical holding times | A | Sampling dates: 8/10 - 27/02 |
| II. | GC/MS Instrument performance check | A | REVIEW Not reviewed for level II |
| III. | Initial calibration | A | 70 RSD ≤ 20. |
| IV. | Routine calibration | A | 70 RSD ≤ 25 / 35 (native / Label) |
| V. | Blanks | SW | |
| VI. | Matrix spike/Matrix spike duplicates / DIAP | N/A | Not req'd |
| VII. | Laboratory control samples | A | OPR. SRU |
| VIII. | Regional quality assurance and quality control | N | |
| IX. | Internal standards | A | Not reviewed for level II |
| X. | Target compound identifications | A | ↓ |
| XI. | Compound quantitation and CRQLs | A | |
| XII. | System performance | A | |
| XIII. | Overall assessment of data | N/A A | |
| XIV. | Field duplicates | N | |
| XV. | Field blanks | N | |

Note: A = Acceptable ND = No compounds detected D = Duplicate
 N = Not provided/applicable R = Rinsate TB = Trip blank
 SW = See worksheet FB = Field blank EB = Equipment blank

Validated Samples: ** level IV
 M Tissues

| | | | | | | | |
|----|--------------|----|---------------|----|-------------|----|--|
| 1 | LDW-B1b-T ** | 11 | LDW-C4-T | 21 | WF14743-101 | 31 | |
| 2 | LDW-B2a-T | 12 | LDW-C6-T | 22 | | 32 | |
| 3 | LDW-B3b-T | 13 | LDW-C7-T1 | 23 | | 33 | |
| 4 | LDW-B4b-T | 14 | LDW-C8-T | 24 | | 34 | |
| 5 | LDW-B5a-T | 15 | LDW-C9-T | 25 | | 35 | |
| 6 | LDW-B8a-T | 16 | LDW-C10-T1 | 26 | | 36 | |
| 7 | LDW-B9b-T ** | 17 | LDW-C10-T1DUP | 27 | | 37 | |
| 8 | LDW-B10a-T | 18 | | 28 | | 38 | |
| 9 | LDW-C1-T | 19 | | 29 | | 39 | |
| 10 | LDW-C2-2-T | 20 | | 30 | | 40 | |

LDC #: 13238A3
 SDG #: DPWG/15217

VALIDATION FINDINGS CHECKLIST

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 Reviewer: 9
 2nd Reviewer: 8

Method: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

| Validation Area | Yes | No | NA | Findings/Comments |
|--|-----|----|----|-------------------|
| I. Technical holding times | | | | |
| All technical holding times were met. | / | | | |
| Cooler temperature criteria was met. | / | | | |
| II. GC/MS instrument performance check | | | | |
| Was PFK exact mass 336.9700 verified? | / | | | |
| Were the retention time windows established for all homologues? | / | | | |
| Is the static resolving power at least 10,000 (10% valley definition)? | / | | | |
| Was the mass resolution adequately check with PFK? | / | | | |
| III. Initial calibration | | | | |
| Was the initial calibration performed at 5 concentration levels? | / | | | |
| Were all percent relative standard deviations (%RSD) \leq 25% for unlabeled standards and $<$ 30% for labeled standards? | / | | | |
| Did all calibration standards meet the Ion Abundance Ratio criteria? | / | | | |
| Was the signal to noise ratio for each target compound \geq 2.5 and for each recovery and internal standard $>$ 10? | / | | | |
| IV. Continuing calibration | | | | |
| Was a routine calibration performed at the beginning of each 12 hour period? | / | | | |
| Were all percent differences (%D) $<$ 40% for unlabeled and labeled standards? | / | | | |
| Did all routine calibration standards meet the Ion Abundance Ratio criteria? | / | | | |
| V. Blanks | | | | |
| Was a method blank associated with every sample in this SDG? | / | | | |
| Was a method blank performed for each matrix and concentration? | / | | | |
| Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. | / | | | |
| VI. Matrix spike/Matrix spike duplicates | | | | |
| Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. | | / | | DWP |
| Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? | | / | | |
| VII. Laboratory control samples | | | | |
| Was an LCS analyzed for this SDG? | / | | | |
| Was an LCS analyzed per extraction batch? | / | | | |
| Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits? | / | | | |

LDC #: 13238A3
 SDG #: DPWG/5217

VALIDATION FINDINGS CHECKLIST

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 2nd Reviewer: [Signature]

| Validation Area | Yes | No | NA | Findings/Comments |
|--|-----|----|----|-------------------|
| VIII. Regional Quality Assurance and Quality Control | | | | |
| Were performance evaluation (PE) samples performed? | | / | | |
| Were the performance evaluation (PE) samples within the acceptance limits? | | | / | |
| IX. Internal standards | | | | |
| Were internal standard recoveries within the 25-150% criteria? | / | | | |
| Was the minimum S/N ratio of all internal standard peaks > 10? | / | | | |
| X. Target compound identification | | | | |
| For polychlorinated biphenyl congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard? | / | | | |
| For polychlorinated biphenyl congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration? | / | | | |
| For other polychlorinated biphenyl congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution? | / | | | |
| Did compound spectra contain all characteristic ions listed in the table attached? | / | | | |
| Was the Ion Abundance Ratio for the two quantitation ions within criteria? | / | | | |
| Was the signal to noise ratio for each target compound and labeled standard > 2.5? | / | | | |
| Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)? | / | | | |
| Was an acceptable lock mass recorded and monitored? | / | | | |
| XI. Compound quantitation/CRQLs | | | | |
| Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound? | / | | | |
| Were compound quantitation and CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation? | / | | | |
| XII. System performance | | | | |
| System performance was found to be acceptable. | / | | | |
| XIII. Overall assessment of data | | | | |
| Overall assessment of data was found to be acceptable. | / | | | |
| XIV. Field duplicates | | | | |
| Field duplicate pairs were identified in this SDG. | | / | | |
| Target compounds were detected in the field duplicates. | | | / | |
| XV. Field blanks | | | | |
| Field blanks were identified in this SDG. | | / | | |
| Target compounds were detected in the field blanks. | | | / | |

LDC #: 13238A3
 SDG #: DPWGIS217

VALIDATION FINDINGS WORKSHEET
Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- N N/A Were all samples associated with a method blank?
- N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?
- N N/A Was method blank contamination less < CRQL for all target compounds?

Blank extraction date: 12/15/05 Blank analysis date: 3/2/05
 Conc. units: ng/g

Associated samples: M

| Compound | Blank ID | Sample Identification | | | | | | | | | | | | |
|----------------------------|-----------------------|-----------------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|--|--|--|--|
| | | 2 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | | | | |
| | <u>WSP43-101</u> | | | | | | | | | | | | | |
| PCB 2 | <u>0.249</u> | <u>1.06/U</u> | <u>0.895/U</u> | <u>0.259/U</u> | <u>0.526/U</u> | <u>0.311/U</u> | <u>0.293/U</u> | <u>0.316/U</u> | <u>0.847/U</u> | <u>0.665/U</u> | | | | |
| PCB 3 | <u>0.355</u> | <u>1.60/U</u> | <u>1.75/U</u> | <u>0.756/U</u> | <u>0.442/U</u> | <u>0.819/U</u> | <u>0.715/U</u> | <u>0.711/U</u> | <u>1.23/U</u> | <u>1.24/U</u> | | | | |
| PCB 11 | <u>1.36</u> | | | <u>4.45/U</u> | | | | | | | | | | |
| Total Monochloro Biphenyls | <u>0.460</u> | | | <u>1.56/U</u> | <u>1.25/U</u> | | | <u>2.01/U</u> | <u>2.00/U</u> | | | | | |
| PCB 1 | <u>0.215</u> | | | | <u>0.382/U</u> | | | <u>1.00/U</u> | <u>0.971/U</u> | | | | | |
| PCB 204 | <u>0.102</u> | | | | <u>0.054/U</u> | | | <u>0.021/U</u> | <u>0.049/U</u> | | | | | |
| | <u>see attachment</u> | | | | | | | | | | | | | |
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CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 All contaminants within five times the method blank concentration were qualified as not detected, "U".

LDC #: B238A3
SDG #: DPWGIS217

VALIDATION FINDINGS WORKSHEET
Blanks

Page: 2 of 2
Reviewer: Q
2nd Reviewer: AL

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

- Y N N/A Were all samples associated with a method blank?
- Y N N/A Was a method blank performed for each matrix and whenever a sample extraction was performed?
- Y N N/A Was method blank contamination less < CRQL for all target compounds?

Blank extraction date: ~~2/25/05~~ / 2/25/05 Blank analysis date: 3/12/05
Conc. units: ng/kg

Associated samples: 111

| Compound | Blank ID | Sample Identification | |
|-----------------------------------|-----------------------|-----------------------|----------------|
| | <u>43-10</u> | <u>15</u> | <u>16</u> |
| <u>PCB 2</u> | <u>0.244</u> | <u>0.310/U</u> | <u>0.853/U</u> |
| <u>PCB 3</u> | <u>0.255</u> | <u>0.781/U</u> | |
| <u>PCB 11</u> | <u>1.76</u> | | |
| <u>Total Monochloro Biphenyls</u> | <u>0.460</u> | | |
| <u>PCB 1</u> | <u>0.215</u> | | |
| <u>PCB 204</u> | <u>0.912</u> | | |
| | <u>see attachment</u> | | |

CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
All contaminants within five times the method blank concentration were qualified as not detected, "U".

C:\WPDOCS\WRK\PCB\BLANKS.166

Lab Name: AXYS ANALYTICAL SERVICES

Contract No.: 4033

Matrix: CORN OIL

Sample Receipt Date: N/A

Extraction Date: 27-Jan-2005

Analysis Date: 12-Feb-2005

Extract Volume (µL): 20

Injection Volume (µL): 1.0

Dilution Factor: N/A

Concentration Units: ng/kg

Time: 2:26:42

Sample Collection: N/A

Project Number: N/A

Lab Sample ID: WG14743-101

Sample Size: 5.00 g

Initial Calibration Date: 04-Feb-2005

Instrument ID: HR GC/MS

GC Column ID: SPB-OCTYL

Sample Data Filename: PB5C_074 S:7

Blank Data Filename: PB5C_074 S:7

Cal. Ver. Data Filename: PB5C_074 S:1

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RR1 |
|---------------------------|---------------|--------------------|-----------------------|------------------|-------------------|------------------|------------------|
| 2 - MoCB | 1 | 5X | JB | 0.215 | 0.0348 | 3.45 | 1.001 |
| 3 - MoCB | 2 | 1.075 | JB | 0.245 | 0.0439 | 3.42 | 0.988 |
| 4 - MoCB | 3 | 1.225 | KJB | 0.355 | 0.0488 | 3.71 | 1.000 |
| 2,2 - DiCB | 4 | 1.775 | KJB | 0.371 | 0.166 | 1.05 | 1.001 |
| 2,3 - DiCB | 5 | 1.855 | U | U | 0.150 | U | U |
| 2,3' - DiCB | 6 | 0.9 | KJB | 0.180 | 0.141 | 1.61 | 1.176 |
| 2,4 - DiCB | 7 | U | U | U | 0.139 | U | U |
| 2,4' - DiCB | 8 | 3.52 | KJB | 0.704 | 0.133 | 1.26 | 1.207 |
| 2,5 - DiCB | 9 | U | U | U | 0.138 | U | U |
| 2,6 - DiCB | 10 | U | U | U | 0.137 | U | U |
| 3,3' - DiCB | 11 | 6.8 | JB | 1.36 | 0.160 | 1.61 | 0.989 |
| 3,4 - DiCB | 12 | 12 + 13 | C U | U | 0.159 | U | U |
| 3,4' - DiCB | 13 | 12 + 13 | C12 | U | U | U | U |
| 3,5 - DiCB | 14 | U | U | U | 0.152 | U | U |
| 4,4' - DiCB | 15 | 1.47 | KJB | 0.294 | 0.188 | 1.14 | 1.000 |
| 2,2'3 - TriCB | 16 | 1.72 | JB | 0.344 | 0.0391 | 1.07 | 1.166 |
| 2,2'4 - TriCB | 17 | 2.26 | JB | 0.452 | 0.0355 | 1.05 | 1.140 |
| 2,2'5 - TriCB | 18 | 4.06 | C JB | 0.812 | 0.0289 | 1.14 | 1.113 |
| 2,2'6 - TriCB | 19 | 0.64 | JB | 0.128 | 0.0320 | 1.04 | 1.002 |
| 2,2'3' - TriCB | 20 | 4.91 | C JB | 0.982 | 0.0359 | 1.02 | 0.848 |

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Approved by: Kearney QA/QC Chemist

27-02-2005
 dd-mm-yyyy

0342

WG14743-101

CLIENT ID: LAB BLANK
 Project Number: N/A
 Sample Data Filename: PB5C_074 S:7

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RRT | | | |
|------------------|----------------|-------------|-----------------------|-------------|-----------------|------------------|--------|--------|-------|-------|
| PCB | 2,3,4 - TriCB | 21 | 5X | 0.25 | 21 + 33 | C JB | 0.450 | 0.0354 | 0.95 | 0.857 |
| | 2,3,4' - TriCB | 22 | 1.38 | | | JB | 0.276 | 0.0395 | 1.04 | 0.871 |
| | 2,3,5 - TriCB | 23 | | | | U | | 0.0367 | | |
| | 2,3,6 - TriCB | 24 | | | | U | | 0.0252 | | |
| | 2,3,4 - TriCB | 25 | 0.525 | | | JB | 0.105 | 0.0324 | 1.00 | 0.825 |
| | 2,3,5 - TriCB | 26 | 1.255 | 26 + 29 | | C JB | 0.251 | 0.0371 | 0.96 | 1.302 |
| | 2,3,6 - TriCB | 27 | 0.475 | | | JB | 0.095 | 0.0246 | 1.10 | 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 | 3.975 | | | JB | 0.795 | 0.0362 | 0.99 | 0.837 | |
| 2,4,6 - TriCB | 32 | 1.32 | | | KJB | 0.264 | 0.0338 | 1.21 | 1.198 | |
| 2,3,4 - TriCB | 33 | | 21 + 33 | | C21 | | | | | |
| 2,3,5 - TriCB | 34 | | | | U | | 0.0369 | | | |
| 3,3,4 - TriCB | 35 | | | | U | | 0.0436 | | | |
| 3,3,4 - TriCB | 36 | | | | U | | 0.0361 | | | |
| 3,4,4' - TriCB | 37 | 1.165 | | | KJB | 0.233 | 0.0451 | 1.26 | 1.001 | |
| 3,4,5 - TriCB | 38 | | | | U | | 0.0374 | | | |
| 3,4,5 - TriCB | 39 | | | | U | | 0.0373 | | | |
| 2,2',3,3' - TeCB | 40 | 1.87 | 40 + 41 + 71 | | C KJB | 0.374 | 0.0261 | 0.55 | 1.338 | |
| 2,2',3,4' - TeCB | 41 | | 40 + 41 + 71 | | C40 | | | | | |
| 2,2',3,4' - TeCB | 42 | 0.83 | | | JB | 0.166 | 0.0277 | 0.72 | 1.313 | |
| 2,2',3,5' - TeCB | 43 | 0.19 | | | KJB | 0.038 | 0.0308 | 0.29 | 1.247 | |
| 2,2',3,5' - TeCB | 44 | 4.725 | 44 + 47 + 65 | | C JB | 0.945 | 0.0243 | 0.78 | 1.287 | |
| 2,2',3,6' - TeCB | 45 | 0.85 | 45 + 51 | | C JB | 0.170 | 0.0256 | 0.80 | 1.147 | |
| 2,2',3,6' - TeCB | 46 | | | | U | | 0.0302 | | | |
| 2,2',4,4' - TeCB | 47 | | 44 + 47 + 65 | | C44 | | | | | |
| 2,2',4,5' - TeCB | 48 | 0.81 | | | KJB | 0.162 | 0.0260 | 0.61 | 1.275 | |
| 2,2',4,5' - TeCB | 49 | 3.34 | 49 + 69 | | C JB | 0.668 | 0.0231 | 0.83 | 1.260 | |
| 2,2',4,6' - TeCB | 50 | 0.83 | 50 + 53 | | C KJB | 0.166 | 0.0247 | 0.98 | 1.112 | |
| 2,2',4,6' - TeCB | 51 | | 45 + 51 | | C45 | | | | | |
| 2,2',5,5' - TeCB | 52 | 7.0 | | | JB | 1.40 | 0.0251 | 0.81 | 1.235 | |
| 2,2',5,6' - TeCB | 53 | | 50 + 53 | | C50 | | | | | |
| 2,2',5,6' - TeCB | 54 | | | | U | | 0.0157 | | | |
| 2,3,3',4' - TeCB | 55 | | | | U | | 0.0650 | | | |
| 2,3,3',4' - TeCB | 56 | 1.435 | | | KJB | 0.287 | 0.0664 | 1.16 | 0.904 | |
| 2,3,3',5' - TeCB | 57 | | | | U | | 0.0634 | | | |
| 2,3,3',5' - TeCB | 58 | | | | U | | 0.0622 | | | |
| 2,3,3',6' - TeCB | 59 | 0.485 | 59 + 62 + 75 | | C KJB | 0.097 | 0.0200 | 0.48 | 1.303 | |
| 2,3,4,4' - TeCB | 60 | 0.545 | | | KJB | 0.109 | 0.0640 | 1.52 | 0.911 | |
| 2,3,4,5' - TeCB | 61 | 5.15 | 61 + 70 + 74 + 76 | | C JB | 1.03 | 0.0597 | 0.66 | 0.875 | |
| 2,3,4,6' - TeCB | 62 | | 59 + 62 + 75 | | C59 | | | | | |
| 2,3,4',5' - TeCB | 63 | | | | U | | 0.0611 | | | |

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Approved by:  QA/QC Chemist

27-02-2005
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0343

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RRT | | |
|---------------------|-------------------|--------------------------------|-----------------------|-------------|-----------------|------------------|--------|-------|-------|
| PCB | 2,3,4',6 - TeCB | 64 | 5X | 1.545 | JB | 0.309 | 0.0194 | 0.76 | 1.350 |
| | 2,3,5,6 - TeCB | 65 | | | C44 | | | | |
| | 2,3',4,4' - TeCB | 66 | 44 + 47 + 65 | 2.815 | JB | 0.563 | 0.0617 | 0.71 | 0.884 |
| | 2,3',4,5' - TeCB | 67 | | | U | | 0.0578 | | |
| | 2,3',4,5' - TeCB | 68 | | | U | | 0.0615 | | |
| | 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 | | 0.0646 | | |
| | 2,3',5',6' - TeCB | 73 | | | U | | 0.0192 | | |
| 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 | 0.605 | | JB | 0.121 | 0.0727 | 0.85 | 1.001 | |
| 3,3',4,5' - TeCB | 78 | | | U | | 0.0683 | | | |
| 3,3',4,5' - TeCB | 79 | | | U | | 0.0548 | | | |
| 3,3',5,5' - TeCB | 80 | | | U | | 0.0616 | | | |
| 3,4,4',5' - TeCB | 81 | | | U | | 0.0671 | | | |
| 2,2',3,3',4' - PeCB | 82 | | | U | | 0.0816 | | | |
| 2,2',3,3',5' - PeCB | 83 | 83 + 99 | 3.49 | C JB | 0.698 | 0.0726 | 1.74 | 0.886 | |
| 2,2',3,3',6' - PeCB | 84 | | 1.13 | KJB | 0.226 | 0.0805 | 2.37 | 1.163 | |
| 2,2',3,4,4' - PeCB | 85 | 85 + 116 + 117 | 0.65 | C KJB | 0.130 | 0.0625 | 2.31 | 0.920 | |
| 2,2',3,4,5' - PeCB | 86 | 86 + 87 + 97 + 108 + 119 + 125 | 3.475 | C KJB | 0.695 | 0.0618 | 1.28 | 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 | 1.08 | C JB | 0.216 | 0.0694 | 1.32 | 1.155 | |
| 2,2',3,4,6' - PeCB | 89 | | | U | | 0.0749 | | | |
| 2,2',3,4,5' - PeCB | 90 | 90 + 101 + 113 | 6.4 | C JB | 1.28 | 0.0637 | 1.39 | 0.869 | |
| 2,2',3,4',6' - PeCB | 91 | 88 + 91 | | C88 | | | | | |
| 2,2',3,3,5' - PeCB | 92 | | 0.77 | JB | 0.154 | 0.0743 | 1.49 | 0.853 | |
| 2,2',3,5,6' - PeCB | 93 | 93 + 95 + 98 + 100 + 102 | 6.0 | C JB | 1.20 | 0.0686 | 1.42 | 1.120 | |
| 2,2',3,5,6' - PeCB | 94 | | | U | | 0.0742 | | | |
| 2,2',3,5',6' - PeCB | 95 | 93 + 95 + 98 + 100 + 102 | | C93 | | | | | |
| 2,2',3,6,6' - PeCB | 96 | | | U | | 0.142 | | | |
| 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 | | | U | | 0.0652 | | | |
| 2,2',4,6,6' - PeCB | 104 | | | U | | 0.112 | | | |
| 2,3,3',4,4' - PeCB | 105 | | 3.77 | KJB | 0.754 | 0.0604 | 2.80 | 1.001 | |
| 2,3,3',4,5' - PeCB | 106 | | | U | | 0.0831 | | | |

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RRT |
|-----------------------|---------------------|-----------------------|--------------------------------|-------------|-----------------|------------------|------|
| PCB | 1,3,3',4',5 - PeCB | 107 | 107 + 124 | | | | |
| | 2,3,3',4,5' - PeCB | 108 | 86 + 87 + 97 + 108 + 119 + 125 | | | | |
| | 2,3,3',4,6 - PeCB | 109 | | | | | |
| | 2,3,3',4',6 - PeCB | 110 | 110 + 115 | C JB | 1.29 | 0.0546 | 1.49 |
| | 2,3,3',5,5' - PeCB | 111 | | U | | 0.0548 | |
| | 2,3,3',5,6 - PeCB | 112 | | U | | 0.0560 | |
| | 2,3,3',5',6 - PeCB | 113 | 90 + 101 + 113 | C90 | | | |
| | 2,3,4,4',5 - PeCB | 114 | | U | | 0.0866 | |
| | 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 | | JB | 1.41 | 0.0822 | 1.49 |
| | 2,3',4,4',6 - PeCB | 119 | 86 + 87 + 97 + 108 + 119 + 125 | C86 | | | |
| | 2,3',4,5,5' - PeCB | 120 | | U | | 0.0554 | |
| | 2,3',4,5',6 - PeCB | 121 | | U | | 0.0545 | |
| | 2',3,3',4,5' - PeCB | 122 | | U | | 0.0913 | |
| | 2',3,4,4',5 - PeCB | 123 | | U | | 0.0835 | |
| | 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 | | 0.102 | | |
| 3,3',4,5,5' - PeCB | 127 | | U | | 0.0905 | | |
| 2,2',3,3',4,4' - HxCB | 128 | 128 + 166 | C JB | 0.549 | 0.0889 | 1.38 | |
| 2,2',3,3',4,5' - HxCB | 129 | 129 + 138 + 160 + 163 | C JB | 2.43 | 0.0843 | 1.10 | |
| 2,2',3,3',4,5' - HxCB | 130 | | U | | 0.107 | | |
| 2,2',3,3',4,6 - HxCB | 131 | | U | | 0.0975 | | |
| 2,2',3,3',4,6' - HxCB | 132 | | KJB | 0.513 | 0.0959 | 1.50 | |
| 2,2',3,3',5,5' - HxCB | 133 | | U | | 0.0926 | | |
| 2,2',3,3',5,6 - HxCB | 134 | 134 + 143 | C U | | 0.0937 | | |
| 2,2',3,3',5,6' - HxCB | 135 | 135 + 151 + 154 | C JB | 0.534 | 0.106 | 1.16 | |
| 2,2',3,3',6,6' - HxCB | 136 | | JB | 0.167 | 0.0798 | 1.09 | |
| 2,2',3,4,4',5 - HxCB | 137 | | U | | 0.0971 | | |
| 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 | | 0.0877 | | |
| 2,2',3,4,4',6' - HxCB | 140 | 139 + 140 | C130 | | | | |
| 2,2',3,4,5,5' - HxCB | 141 | | KJB | 0.337 | 0.0931 | 0.73 | |
| 2,2',3,4,5,6 - HxCB | 142 | | U | | 0.0979 | | |
| 2,2',3,4,5,6' - HxCB | 143 | 134 + 143 | C134 | | | | |
| 2,2',3,4,5',6 - HxCB | 144 | | U | | 0.110 | | |
| 2,2',3,4,6,6' - HxCB | 145 | | U | | 0.0796 | | |
| 2,2',3,4',5,5' - HxCB | 146 | | KJB | 0.306 | 0.0881 | 1.66 | |
| 2,2',3,4',5,6 - HxCB | 147 | 147 + 149 | C JB | 0.997 | 0.0866 | 1.22 | |
| 2,2',3,4',5,6' - HxCB | 148 | | U | | 0.111 | | |
| 2,2',3,4',5',6 - HxCB | 149 | 147 + 149 | C147 | | | | |

0345

| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RRT |
|----------------------------|-----------|-----------------------|-----------------------|-------------|-----------------|------------------|-------|
| 2,2',3,4',6,6' - HxCB | 150 | | U | | 0.0787 | | |
| 2,2',3,5,5',6 - HxCB | 151 | 135 + 151 + 154 | C135 | | | | |
| 2,2',3,5,6,6' - HxCB | 152 | | U | | 0.0762 | | |
| 2,2',4,4',5,5' - HxCB | 153 | 153 + 168 | C JB | 1.80 | 0.0766 | 1.28 | 0.899 |
| 2,2',4,4',5,6' - HxCB | 154 | 135 + 151 + 154 | C135 | | | | |
| 2,2',4,4',6,6' - HxCB | 155 | | U | | 0.0579 | | |
| 2,3,3',4,4',5 - HxCB | 156 | 156 + 157 | C JB | 0.631 | 0.0930 | 1.29 | 1.000 |
| 2,3,3',4,4',5' - HxCB | 157 | 156 + 157 | C156 | | | | |
| 2,3,3',4,4',6 - HxCB | 158 | | JB | 0.250 | 0.0720 | 1.12 | 0.938 |
| 2,3,3',4,5,5' - HxCB | 159 | | U | | 0.0751 | | |
| 2,3,3',4,5,6 - HxCB | 160 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4,5',6 - HxCB | 161 | | U | | 0.0690 | | |
| 2,3,3',4',5,5' - HxCB | 162 | | U | | 0.0730 | | |
| 2,3,3',4',5,6 - HxCB | 163 | 129 + 138 + 160 + 163 | C129 | | | | |
| 2,3,3',4',5',6 - HxCB | 164 | | JB | 0.107 | 0.0740 | 1.12 | 0.921 |
| 2,3,3',5,5',6 - HxCB | 165 | | U | | 0.0771 | | |
| 2,3,4,4',5,6 - HxCB | 166 | 128 + 166 | C128 | | | | |
| 2,3',4,4',5,5' - HxCB | 167 | | KJB | 0.166 | 0.0664 | 1.47 | 1.001 |
| 2,3',4,4',5',6 - HxCB | 168 | 153 + 166 | C153 | | | | |
| 3,3',4,4',5,5' - HxCB | 169 | | U | | 0.0786 | | |
| 2,2',3,3',4,4',5 - HpCB | 170 | | JB | 0.676 | 0.169 | 1.00 | 0.936 |
| 2,2',3,3',4,4',6 - HpCB | 171 | 171 + 173 | C U | | 0.156 | | |
| 2,2',3,3',4,4',5,5' - HpCB | 172 | | U | | 0.159 | | |
| 2,2',3,3',4,5,6 - HpCB | 173 | 171 + 173 | C171 | | | | |
| 2,2',3,3',4,5,6' - HpCB | 174 | | JB | 0.371 | 0.140 | 1.02 | 1.133 |
| 2,2',3,3',4,5',6 - HpCB | 175 | | U | | 0.135 | | |
| 2,2',3,3',4,6,6' - HpCB | 176 | | U | | 0.101 | | |
| 2,2',3,3',4',5,6 - HpCB | 177 | | KJB | 0.271 | 0.151 | 1.60 | 1.145 |
| 2,2',3,3',5,5',6 - HpCB | 178 | | U | | 0.138 | | |
| 2,2',3,3',5,6,6' - HpCB | 179 | | KJB | 0.133 | 0.0963 | 2.25 | 1.010 |
| 2,2',3,4,4',5,5' - HpCB | 180 | 180 + 193 | C JB | 1.07 | 0.130 | 1.13 | 0.910 |
| 2,2',3,4,4',5,6 - HpCB | 181 | | U | | 0.142 | | |
| 2,2',3,4,4',5,6' - HpCB | 182 | | U | | 0.140 | | |
| 2,2',3,4,4',5',6 - HpCB | 183 | 183 + 185 | C JB | 0.381 | 0.136 | 1.15 | 1.127 |
| 2,2',3,4,4',6,6' - HpCB | 184 | | U | | 0.0927 | | |
| 2,2',3,4,5,5',6 - HpCB | 185 | 183 + 185 | C183 | | | | |
| 2,2',3,4,5,6,6' - HpCB | 186 | | U | | 0.101 | | |
| 2,2',3,4',5,5',6 - HpCB | 187 | | KJB | 0.383 | 0.126 | 1.39 | 1.110 |
| 2,2',3,4',5,6,6' - HpCB | 188 | | U | | 0.0770 | | |
| 2,3,3',4,4',5,5' - HpCB | 189 | | U | | 0.108 | | |
| 2,3,3',4,4',5,6 - HpCB | 190 | | KJB | 0.144 | 0.129 | 0.68 | 0.947 |
| 2,3,3',4,4',5',6 - HpCB | 191 | | U | | 0.125 | | |
| 2,3,3',4,5,5',6 - HpCB | 192 | | U | | 0.127 | | |



| COMPOUND | IUPAC NO. | CO-ELUTIONS | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT | ION ABUND. RATIO | RRT |
|--|----------------|----------------------|-----------------------|------------------|-------------------|------------------|------------------|
| 2,3,3',4',5,5',6 - HpCB | 193 | 180 + 193 | C180 | | | | |
| 2,2',3,3',4,4',5,5' - OcCB | 194 | 0.94 | KJB | 0.188 | 0.0067 | 1.11 | 0.991 |
| 2,2',3,3',4,4',5,6 - OcCB | 195 | 0.555 | JB | 0.111 | 0.0075 | 0.84 | 0.945 |
| 2,2',3,3',4,4',5,6' - OcCB | 196 | 0.585 | KJB | 0.117 | 0.0079 | 1.35 | 0.916 |
| 2,2',3,3',4,4',6,6' - OcCB | 197 | 197 + 200 | C U | | 0.0052 | | |
| 2,2',3,3',4,5,5',6 - OcCB | 198 | 1.175 | C JB | 0.235 | 0.0079 | 0.84 | 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 | 0.115 | KJB | 0.023 | 0.0053 | 5.00 | 1.023 |
| 2,2',3,3',5,5',6,6' - OcCB | 202 | 0.43 | JB | 0.086 | 0.0051 | 0.79 | 1.000 |
| 2,2',3,4,4',5,5',6 - OcCB | 203 | 0.755 | KJB | 0.151 | 0.0072 | 1.33 | 0.920 |
| 2,2',3,4,4',5,6,6' - OcCB | 204 | 0.06 | KJB | 0.012 | 0.0054 | 45.99 | 1.040 |
| 2,3,3',4,4',5,5',6 - OcCB | 205 | 0.255 | KJB | 0.051 | 0.0061 | 3.40 | 1.000 |
| 2,2',3,3',4,4',5,5',6 - NoCB | 206 | 0.745 | KJB | 0.149 | 0.111 | 0.96 | 1.001 |
| 2,2',3,3',4,4',5,6,6' - NoCB | 207 | | U | | 0.0915 | | |
| 2,2',3,3',4,5,5',6,6' - NoCB | 208 | 0.665 | JB | 0.133 | 0.0922 | 0.82 | 1.001 |
| 2,2',3,3',4,4',5,5',6,6' - DeCB | 209 | 0.77 | KJB | 0.154 | 0.0076 | 0.96 | 1.001 |

(1) C = co-eluting congener; U = not detected; K = peak detected, but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; E = exceeds calibrated linear range, see dilution data; D = dilution data; Z = compound not requested; J = concentration less than LMCL; B = analyte found in sample and the associated blank; X = results reported separately

These pages are part of a larger report that may contain information necessary for full data evaluation.



0347

Form 1A
 HOMOLOGUE TOTAL POLYCHLORINATED BIPHENYLS (PCB) ANALYSIS REPORT

| | | |
|------------------------------------|---------------------------|--------------|
| Lab Name: AXYS ANALYTICAL SERVICES | Sample Collection: | N/A |
| Contract No.: 4033 | Project Number: | N/A |
| Matrix: CORN OIL | Lab Sample ID: | WG14743-101 |
| Sample Receipt Date: N/A | Sample Size: | 5.00 g |
| Extraction Date: 27-Jan-2005 | Initial Calibration Date: | 04-Feb-2005 |
| Analysis Date: 12-Feb-2005 | Instrument ID: | HR GC/MS |
| Time: 2:26:42 | GC Column ID: | SPB-OCTYL |
| Extract Volume (µL): 20 | Blank Data Filename: | PB5C_074 S:7 |
| Injection Volume (µL): 1.0 | Cal. Ver. Data Filename: | PB5C_074 S:1 |
| Dilution Factor: N/A | Sample Datafile(s): | PB5C_074 S:7 |
| Concentration Units : ng/kg | | |

| PCB HOMOLOGUE GROUP | LAB FLAG ¹ | CONC. FOUND | DETECTION LIMIT |
|---|-----------------------|-----------------|-----------------|
| Total Monochloro Biphenyls <i>5X</i> 2.3 | | 0.460 | 0.0488 |
| Total Dichloro Biphenyls 6.8 | | 1.36 | 0.188 |
| Total Trichloro Biphenyls 23.45 | | 4.69 | 0.0451 |
| Total Tetrachloro Biphenyls 26.85 | | 5.37 | 0.0727 |
| Total Pentachloro Biphenyls 31.2 | | 6.24 | 0.142 |
| Total Hexachloro Biphenyls 37.35 | | 7.47 | 0.111 |
| Total Heptachloro Biphenyls 12.5 | | 2.50 | 0.169 |
| Total Octachloro Biphenyls 2.16 | | 0.432 | 0.0079 |
| Total Nonachloro Biphenyls 0.665 | | 0.133 | 0.111 |
| Decachloro Biphenyl | U | | 0.0076 |
| TOTAL PCBs | | 25.0 | |

(1) U = Not detected
 (2) All header information pertains to the initial instrumental analysis of the sample extract.
 Additional sample datafiles listed refer to secondary analysis of the sample extract.

These pages are part of a larger report that may contain information necessary for full data evaluation.

LDC #: 13238A3
 SDG #: D-2/15217

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: X

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

$RRF = (A_x)(C_s)/(A_s)(C_x)$
 average RRF = sum of the RRFs/number of standards
 $\%RSD = 100 * (S/\bar{X})$

A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard
 S = Standard deviation of the RRFs, \bar{X} = Mean of the RRFs

| # | Standard ID | Calibration Date | Compound (Reference Internal Standard) | Reported | Recalculated | Reported | Recalculated | Reported | Recalculated |
|---|-------------|------------------|--|-----------------------|-----------------------|--------------|--------------|----------|--------------|
| | | | | Average RRF (Initial) | Average RRF (Initial) | RRF (std) | RRF (std) | %RSD | %RSD |
| 1 | 1CAZ | 2/2/05 | PCE-77 (¹³ C-PCB-77) | 0.91 | 0.91 | 0.87 | 0.86 | 3.54 | 3.20 |
| | | | PCE-105 (¹³ C-PCB-105) | 0.83 | 0.83 | 0.82 | 0.82 | 3.00 | 3.19 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 0.98 | 0.96 | 0.96 | 1.52 | 1.61 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 0.79 | 0.79 | 0.79 | 2.01 | 1.65 |
| 2 | | | PCB-77 (¹³ C-PCB-77) | | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | | |
| 3 | | | PCB-77 (¹³ C-PCB-77) | | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | | |

Comments: Refer to Initial Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 1323843
 SDG #: DPW 15217

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = 100 * (ave. RRF - RRF)/ave. RRF
 RRF = (A_s)(C_s)/(A_c)(C)

Where: ave. RRF = initial calibration average RRF
 RRF = continuing calibration RRF
 A_s = Area of compound, A_s = Area of associated internal standard
 C_s = Concentration of compound, C_s = Concentration of internal standard

| # | Standard ID | Calibration Date | Compound (Reference Internal Standard) | Average RRF (Initial) | Reported | Recalculated | Reported | Recalculated |
|---|-----------------|------------------|--|-----------------------|---------------------------|---------------------------|----------|--------------|
| | | | | | RRF C _{inc} (CC) | RRF C _{inc} (CC) | %D | %D |
| 1 | DBSC-084 S=1 | 2/17/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 52.1 | 52.3 | reported | 4.5 |
| | | | PCB-105 (¹³ C-PCB-105) | 0.83 | 54.0 | 54.0 | | 7.8 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 100 | 101 | | 0.8 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 52.9 | 53.2 | | 6.4 |
| 2 | DBSC-074 S=1 | 1/2/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 47.4 | 47.3 | reported | 5.38 |
| | | | PCB-105 (¹³ C-PCB-105) | 0.83 | 50.4 | 50.3 | | 0.6 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 97.3 | 97.7 | | 2.3 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 51.5 | 51.8 | | 3.60 |
| 3 | | | PCB-77 (¹³ C-PCB-77) | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | |

Comments: Refer to Routine Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 3238A3
 SDG #: DPWGLS-17
 METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

VALIDATION FINDINGS WORKSHEET
Laboratory Control Sample Results Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: 0

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

% Recovery = 100 * SSC/SA

Where: SSC = Spiked sample concentration
 SA = Spike added

RPD = |LCS - LCSD| * 2 / (LCS + LCSD)

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS ID: OTR

| Compound | Spike Added (ng/ml) | | Spiked Sample Concentration (ng/L) | | LCS | | LCSD | | LCS/LCSD | |
|--------------------|---------------------|------|------------------------------------|------|------------------|---------|------------------|---------|----------|--------------|
| | LCS | LCSD | LCS | LCSD | Percent Recovery | | Percent Recovery | | RPD | |
| | | | | | Reported | Recalc. | Reported | Recalc. | Reported | Recalculated |
| PCB-77 | 50 | NA | 47.4 | NA | 94.9 | 94.8 | | | | |
| PCB-81 | ↓ | | 47.2 | | 94.4 | 94.4 | | | | |
| PCB-105 | ↓ | | 51.2 | | 102 | 102 | | | | |
| PCB-114 | ↓ | | 51.0 | | 102 | 102 | | | | |
| PCB-118 | ↓ | | 52.6 | | 105 | 105 | | | | |
| PCB-123 | ↓ | | 51.3 | | 103 | 103 | | | | |
| PCB-126 | ↓ | | 50.8 | | 102 | 102 | | | | |
| PCB-156 / PCB157 | 100 | | 97.1 | | 97.1 | 97.1 | | | | |
| PCB-167 | 50 | | 49.0 | | 97.9 | 98 | | | | |
| PCB-169 | ↓ | | 47.6 | | 95.3 | 95.3 | | | | |
| PCB-170 | | | | | | | | | | |
| PCB-180 | | | | | | | | | | |
| PCB-189 | 50 | ↓ | 51.3 | ↓ | 103 | 103 | | | | |
| | | | | | | | | | | |
| | | | | | | | | | | |

Comments: Refer to Laboratory Control Sample findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

Ions Monitored for HRGC/HRMS Analysis of Polychlorinated Biphenyls

| Descriptor | Accurate mass ⁽ⁿ⁾ | Ion ID | Analyte | Substance | |
|------------|------------------------------|----------|----------------------|--------------------|-----|
| 1 | 289.9224 | M | C12 H6 35Cl4 | TCB | |
| | 291.9194 | M+2 | C12 H6 35Cl3 37Cl4 | TCB | |
| | 301.9626 | M | 13C12 H6 35Cl4 | PeCB | |
| | 303.9597 | M+2 | 13C12 H6 35Cl3 37Cl | PeCB | |
| | 325.8804 | M+2 | C12 H5 35Cl4 37Cl | PeCB | |
| | 327.8775 | M+4 | C12 H5 35Cl3 37Cl2 | PeCB | |
| | [292.9825] | Lock | C7 F11 | FFK | |
| 2 | 325.8804 | M+2 | C12 H5 35Cl4 37Cl | PeCB | |
| | 327.8775 | M+4 | C12 H5 35Cl3 37Cl2 | PeCB | |
| | 337.9207 | M+2 | 13C12 H5 35Cl4 37Cl | PeCB | |
| | 339.9178 | M+4 | 13C12 H5 35Cl3 37Cl2 | PeCB | |
| | 359.8415 | M+2 | C12 H4 35Cl5 37Cl | HxCB | |
| | 361.8385 | M+4 | C12 H4 35Cl4 37Cl2 | HxCB | |
| | 371.8817 | M+2 | 13C12 H4 35Cl5 37Cl | HxCB | |
| | 373.8788 | M+4 | 13C12 H4 35Cl4 37Cl2 | HxCB | |
| | 393.8025 | M+2 | C12 H3 35Cl6 37Cl | HpCB | |
| | 395.7996 | M+4 | C12 H3 35Cl5 37Cl2 | HpCB | |
| | 405.8428 | M+2 | 13C12 H3 35Cl6 37Cl | HpCB | |
| | 407.8398 | M+4 | 13C12 H3 35Cl5 37Cl2 | HpCB | |
| | [354.9892] | Lock | C9F13 | FFK | |
| | 3 | 509.7229 | M+4 | 13C12 35Cl10 37Cl2 | DCB |
| | | 511.7199 | M+6 | 13C12 35Cl9 37Cl3 | |
| 513.7170 | | M+8 | 13C12 35Cl8 37Cl4 | | |
| [442.9728] | | Lock | C10 F17 | FFK | |
| | | | | | |

S = internal/recovery standard

H = 1.007825
 C = 12.000000
¹³C = 13.003355
 F = 18.9984

³⁵Cl = 34.968853
³⁷Cl = 36.965903

LDC #: 13238A3
 SDG #: OPWGLS=IT

VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)
Y N N/A Were all reported results recalculated and verified for all level IV samples?
Y N N/A Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

Concentration = $\frac{A_x(I_s)(DF)}{A_s(RRF)(V_s)(\%S)}$

A_x = Area of the characteristic ion (EICP) for the compound to be measured

A_s = Area of the characteristic ion (EICP) for the specific internal standard

I_s = Amount of internal standard added in nanograms (ng)

RRF = Relative response factor of the calibration standard.

V_s = Volume or weight of sample pruged in milliliters (ml) or grams (g).

Df = Dilution factor.

%S = Percent solids, applicable to soils and solid matrices only.

Example:
 Sample I.D. 1, PCBT:

Conc. = $\frac{(1.99e+07)(2000)(1)}{(9.82e+07)(0.91)(1.44)(1)}$
 = 309.29 ng/kg

| # | Sample ID | Compound | Reported Concentration () | Calculated Concentration () | Qualification |
|---|-----------|----------|-------------------------------|---------------------------------|---------------|
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Attachment D-3: Sediment Chemistry

CHEMICAL DATA QUALITY REVIEW FOR BENTHIC SEDIMENT SAMPLES**Lower Duwamish Waterway Group
LDC# 13247**

This report details the findings of an EPA Level II and Level IV data validation review of Analytical chemistry results generated in support of the Lower Duwamish Waterway Group project. The analyses were performed by Axys Analytical Services, Ltd. Samples were analyzed for HRGC/HRMS Polychlorinated Biphenyl Congeners by modified EPA Method 1668A. Samples are referenced under the following Sample Delivery Group: DPWG15252. See the Sample Analysis Table (Attachment 1) for the number of samples reviewed and the Sample Validation Table (Attachment 2) for the sample identifications and analyses. Sample IDs ending in "***" underwent Level IV review.

The QC guidelines used for data qualification are those specified in the EPA Region 10 SOP for the Validation of 1668 Toxic, Dioxin-like PCB Data (Revision 1.0, December 8, 1995). Specific QC criteria used follows the Final Benthic Invertebrate Sampling of the Lower Duwamish Waterway Quality Assurance Project Plan (July 30, 2004). Where specific guidance is not available, the data has been evaluated in a conservative manner consistent with industry standards using professional experience.

The following items were evaluated during the review:

- Holding Times
- Sample Preservation
- Cooler Temperatures
- Instrument Calibration*
- Blanks
- Matrix Spike/Matrix Spike Duplicates
- Internal Standards
- Laboratory Control Samples
- Target Compound Identifications*
- Compound Quantitation and CRQLs*
- System Performance
- Field Duplicates

*Data were not reviewed for Level II.

An asterisk (*) will be placed in the margin to the left of any revised section in the text.

Only issues which require comment or action are discussed in this report. Data deficiencies are arranged by method. Potential effects of data anomalies have been described where possible.

The following are definitions of the data qualifiers:

- U Indicates the compound or analyte was analyzed for but not detected at or above the stated limit.
- J Indicates an estimated value.
 - J1 Blank Contamination: Indicates possible high bias and/or false positives.
 - J2 Calibration Range exceeded: Indicates possible low bias.
 - J3 Holding times not met: Indicates low bias for most analytes.
 - J4 Other QC parameters outside control limits: bias not readily determined.
 - J5 Other QC parameters outside control limits. The reported results appear to be biased high. The actual value of target compound in the sample may be lower than the value reported by the laboratory.
 - J6 Other QC parameters outside control limits. The reported results appear to be biased low. The actual value of target compound in the sample may be higher than the value reported by the laboratory.
- R Quality control indicates the data is not usable.
- N Presumptive evidence of presence of the constituent.
- UJ Indicates the compound or analyte was analyzed for but not detected. The sample detection limit is an estimated value.
- A Indicates the finding is based upon technical validation criteria.
- P Indicates the finding is related to a protocol/contractual deviation.

***Overall Data Assessment**

Laboratory duplicate precision exceedances have warranted the qualification of detected results as estimated (J).

*The frequency of matrix spike (MS) and matrix spike duplicate (MSD) was not met as required by the QAPP. MS/MSD analyses are not required for EPA Method 1668A. The laboratory consulted with the client on this discrepancy and was instructed to proceed with the extraction and analysis despite the absence of MS/MSDs.

Field duplicates were not collected for this sampling event.

The quality control criteria reviewed, other than those discussed above, were met and are considered acceptable. Sample results that were found to be estimated (J) are usable for limited purposes only. Based upon the data validation all other results are considered valid and usable for all purposes.

Attachment 2

SDG#: DPWG15252

VALIDATION SAMPLE TABLE

LDC#: 13247A

Project Name: Lower Duwamish Waterway Group

Parameters/Analytical Method

Project #04-08-06-21

| Client ID # | Lab ID # | Matrix | Date Collected | PCB Cong. (1668A) | | | | | | | | | | | | | | | |
|--------------|------------|----------|----------------|-------------------|--|--|--|--|--|--|--|--|--|--|--|--|--|--|--|
| LDW-B1b-S | L7505-1 | sediment | 08/10/04 | X | | | | | | | | | | | | | | | |
| LDW-B2a-S | L7505-2 | sediment | 08/14/04 | X | | | | | | | | | | | | | | | |
| LDW-B3b-S | L7505-3 | sediment | 08/10/04 | X | | | | | | | | | | | | | | | |
| LDW-B4b-S | L7505-4 | sediment | 08/17/04 | X | | | | | | | | | | | | | | | |
| LDW-B5a-S | L7505-5 | sediment | 08/22/04 | X | | | | | | | | | | | | | | | |
| LDW-B8a-S | L7505-6 | sediment | 08/27/04 | X | | | | | | | | | | | | | | | |
| LDW-B9b-S | L7505-7 | sediment | 08/11/04 | X | | | | | | | | | | | | | | | |
| LDW-B10a-S** | L7505-8 | sediment | 08/25/04 | X | | | | | | | | | | | | | | | |
| LDW-C1-S** | L7505-9 | sediment | 08/26/04 | X | | | | | | | | | | | | | | | |
| LDW-C2-S2 | L7505-10 | sediment | 08/26/04 | X | | | | | | | | | | | | | | | |
| LDW-C4-S | L7505-11 | sediment | 08/27/04 | X | | | | | | | | | | | | | | | |
| LDW-C6-S | L7505-12 | sediment | 08/26/04 | X | | | | | | | | | | | | | | | |
| LDW-C7-S-1 | L7505-13 | sediment | 08/26/04 | X | | | | | | | | | | | | | | | |
| LDW-C8-S | L7505-14 | sediment | 08/26/04 | X | | | | | | | | | | | | | | | |
| LDW-C9-S | L7505-15 | sediment | 08/25/04 | X | | | | | | | | | | | | | | | |
| LDW-C10-S-1 | L7505-16 | sediment | 08/25/04 | X | | | | | | | | | | | | | | | |
| LDW-B1b-SDUP | L7505-1DUP | sediment | 08/10/04 | X | | | | | | | | | | | | | | | |
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Note: X = Validation was performed.

HRGC/HRMS Polychlorinated Biphenyl Congeners by modified EPA Method 1668A

I. Technical Holding Times

All technical holding time requirements were met.

The chain-of-custodies were reviewed for documentation of cooler temperatures. All cooler temperatures met validation criteria.

II. GC/MS Instrument Performance Check

Instrument performance was checked at the required daily frequency.

Retention time windows were established for all congeners. The chromatographic resolution was less than or equal to 40% for congeners PCB-23 and PCB-34 and congeners PCB-182 and PCB-187.

The static resolving power was at least 10,000 (10% valley definition).

Instrument performance check data were not reviewed for Level II.

III. Initial Calibration

A five point initial calibration was performed as required by the method.

The percent relative standard deviations (%RSD) were less than or equal to 20.0% for compounds.

The minimum S/N ratio for each target compound was greater than or equal to 2.5 and greater than or equal to 10 for each recovery and internal standard compound.

Initial calibration data were not reviewed for Level II.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequencies.

All of the routine calibration percent differences (%D) between the initial calibration RRF and the routine calibration RRF were less than or equal to 25.0% for unlabeled compounds and less than or equal to 35.0% for labeled compounds.

Continuing calibration data were not reviewed for Level II.

V. Blanks

Method blanks were reviewed for each matrix as applicable. No polychlorinated biphenyl contaminants were found in the method blanks with the following exceptions:

| Associated SDG | Method Blank ID | Extraction Date | Compound | Concentration | Associated Samples |
|----------------|-----------------|-----------------|---------------------|---------------|---------------------------------|
| DPWG15252 | WG14745-101 | 1/25/05 | PCB-66 | 0.108 ng/Kg | All samples in SDG DPWG15252 |
| | | | PCB-77 | 0.042 ng/Kg | |
| | | | PCB-99+101+113 | 0.236 ng/Kg | |
| | | | PCB-105 | 0.078 ng/Kg | |
| | | | PCB-110+115 | 0.289 ng/Kg | |
| | | | PCB-114 | 0.044 ng/Kg | |
| | | | PCB-118 | 0.195 ng/Kg | |
| | | | PCB-123 | 0.027 ng/Kg | |
| | | | PCB-126 | 0.041 ng/Kg | |
| | | | PCB-129+138+160+163 | 0.310 ng/Kg | |
| | | | PCB-153+168 | 0.257 ng/Kg | |
| | | | PCB-156+157 | 0.088 ng/Kg | |
| | | | PCB-180+193 | 0.160 ng/Kg | |

Sample concentrations were compared to concentrations detected in the method blanks. The sample concentrations were either not detected or were significantly greater (>5X blank contaminants) than the concentrations found in the associated method blanks.

No field blanks were identified in this SDG.

*VI. Matrix Spike/Matrix Spike Duplicates

Matrix spike (MS) and matrix spike duplicate (MSD) analyses were not required by the method.

Duplicate (DUP) sample analyses were reviewed for each matrix as applicable. Results were within QC limits with the following exceptions:

*Indicates change as the result of report review.
SDG DPWG15252

| Associated SDG | DUP ID (Associated Samples) | Compounds | RPD (Limits) | Flag | A or P |
|----------------|-----------------------------|--|--|--|--------|
| DPWG15252 | LDW-B1b-SDUP (LDW-B1b-S) | PCB-99+101+113 PCB-129+138+160+163 PCB-153+168 PCB-156+157 PCB-167 PCB-180+193 PCB-189 | 53.7 (≤ 50) 75.6 (≤ 50) 89.5 (≤ 50) 60.8 (≤ 50) 59.9 (≤ 50) 112 (≤ 50) 91.6 (< 50) | J4 (all detects) J4 (all detects) J4 (all detects) J4 (all detects) J4 (all detects) J4 (all detects) J4 (all detects) | A |

VII. Laboratory Control Samples (LCS)

Laboratory control samples were reviewed for each matrix as applicable. The percent recoveries (%R) were within the QC limits.

Standard reference material was performed at the required frequencies.

VIII. Regional Quality Assurance and Quality Control

Not applicable.

IX. Internal Standards

All internal standard recoveries were within QC limits.

X. Target Compound Identifications

All target compound identifications were within validation criteria.

Target compound identifications data were not reviewed for Level II.

XI. Compound Quantitation and CRQLs

All compound quantitation and CRQLs were within validation criteria.

Compound quantitation and CRQLs data were not reviewed for Level II.

XII. System Performance

The system performance was acceptable.

System performance data were not reviewed for Level II.

*Indicates change as the result of report review.
SDG DPWG15252

XIII. Overall Assessment of Data

The overall assessment of data was acceptable.

XIV. Field Duplicates

No field duplicates were identified in this SDG.

*Indicates change as the result of report review.
SDG DPWG15252

METHOD: HRGC/HRMS Polychlorinated Biphenyl Congeners (EPA Method 1668A)

The samples listed below were reviewed for each of the following validation areas. Validation findings are noted in attached validation findings worksheets.

| Validation Area | | Findings | Comments |
|-----------------|--|------------|------------------------------|
| I. | Technical holding times | A | Sampling dates: 8/10 - 27/04 |
| II. | GC/MS Instrument performance check | A | NR for level 1/1 |
| III. | Initial calibration | A | 70 PSD ≤ 20 |
| IV. | Routine calibration | A | 70 D ≤ 25/35. |
| V. | Blanks | SW | |
| VI. | Matrix spike/Matrix spike duplicates | DUP N / SW | not req'd |
| VII. | Laboratory control samples | A | OPR, SEM |
| VIII. | Regional quality assurance and quality control | N | |
| IX. | Internal standards | A | NR for level 1/1 |
| X. | Target compound identifications | A | |
| XI. | Compound quantitation and CRQLs | A | |
| XII. | System performance | A | |
| XIII. | Overall assessment of data | SW | |
| XIV. | Field duplicates | N | |
| XV. | Field blanks | N | |

Note: A = Acceptable ND = No compounds detected D = Duplicate
 N = Not provided/applicable R = Rinsate TB = Trip blank
 SW = See worksheet FB = Field blank EB = Equipment blank

Validated Samples:

M seeds

| | | | | | | |
|----|---------------|----|--------------|----|------------|----|
| 1 | LDW-B1b-S | 11 | LDW-C4-S | 21 | NS1245-101 | 31 |
| 2 | LDW-B2a-S | 12 | LDW-C6-S | 22 | | 32 |
| 3 | LDW-B3b-S | 13 | LDW-C7-S-1 | 23 | | 33 |
| 4 | LDW-B4b-S | 14 | LDW-C8-S | 24 | | 34 |
| 5 | LDW-B5a-S | 15 | LDW-C9-S | 25 | | 35 |
| 6 | LDW-B8a-S | 16 | LDW-C10-S-1 | 26 | | 36 |
| 7 | LDW-B9b-S | 17 | LDW-B1b-SDUP | 27 | | 37 |
| 8 | LDW-B10a-S ** | 18 | | 28 | | 38 |
| 9 | LDW-C1-S ** | 19 | | 29 | | 39 |
| 10 | LDW-C2-S2 | 20 | | 30 | | 40 |

LDC #: 13247A3
 SDG #: DPW/15252

VALIDATION FINDINGS CHECKLIST

Page: 1 of 2
 Reviewer: 9
 2nd Reviewer: X

Method: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

| Validation Area | Yes | No | NA | Findings/Comments |
|--|-----|----|----|-------------------|
| I. Technical holding times | | | | |
| All technical holding times were met. | / | | | |
| Cooler temperature criteria was met. | / | | | |
| II. GC/MS instrument performance check | | | | |
| Was PFK exact mass 380.9760 verified? | / | | | |
| Were the retention time windows established for all homologues? | / | | | |
| Is the static resolving power at least 10,000 (10% valley definition)? | / | | | |
| Was the mass resolution adequately check with PFK? | / | | | |
| III. Initial calibration | | | | |
| Was the initial calibration performed at 5 concentration levels? | / | | | |
| Were all percent relative standard deviations (%RSD) \leq 25% for unlabeled standards and < 30% for labeled standards? | / | | | |
| Did all calibration standards meet the Ion Abundance Ratio criteria? | / | | | |
| Was the signal to noise ratio for each target compound \geq 2.5 and for each recovery and internal standard > 10? | / | | | |
| IV. Continuing calibration | | | | |
| Was a routine calibration performed at the beginning of each 12 hour period? | / | | | |
| Were all percent differences (%D) < 40% for unlabeled and labeled standards? | / | | | |
| Did all routine calibration standards meet the Ion Abundance Ratio criteria? | / | | | |
| V. Blanks | | | | |
| Was a method blank associated with every sample in this SDG? | / | | | |
| Was a method blank performed for each matrix and concentration? | / | | | |
| Was there contamination in the method blanks? If yes, please see the Blanks validation completeness worksheet. | / | | | |
| VI. Matrix spike/Matrix spike duplicates | | | | |
| Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water. | | / | | dup |
| Were the MS/MSD percent recoveries (%R) and the relative percent differences (RPD) within the QC limits? | | | / | |
| VII. Laboratory control samples | | | | |
| Was an LCS analyzed for this SDG? | / | | | |
| Was an LCS analyzed per extraction batch? | / | | | |
| Were the LCS percent recoveries (%R) and relative percent difference (RPD) within the QC limits? | / | | | |

LDC #: 1324783
 SDG #: DPNG15252

VALIDATION FINDINGS CHECKLIST

Page: 2 of 2
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

| Validation Area | Yes | No | NA | Findings/Comments |
|--|-----|----|----|-------------------|
| VIII. Regional Quality Assurance and Quality Control | | | | |
| Were performance evaluation (PE) samples performed? | | / | | |
| Were the performance evaluation (PE) samples within the acceptance limits? | | | / | |
| IX. Internal standards | | | | |
| Were internal standard recoveries within the 25-150% criteria? | / | | | |
| Was the minimum S/N ratio of all internal standard peaks > 10? | / | | | |
| X. Target compound identification | | | | |
| For polychlorinated biphenyl congeners with associated labeled standards, were the retention times of the two quantitation peaks within -1 to 3 sec. of the RT of the labeled standard? | / | | | |
| For polychlorinated biphenyl congeners without associated labeled standards, were the relative retention times of the two quantitation peaks within 0.005 time units of the RRT measured in the routine calibration? | / | | | |
| For other polychlorinated biphenyl congeners, were the retention times of the two quantitation peaks within RT established in the performance check solution? | / | | | |
| Did compound spectra contain all characteristic ions listed in the table attached? | / | | | |
| Was the Ion Abundance Ratio for the two quantitation ions within criteria? | / | | | |
| Was the signal to noise ratio for each target compound and labeled standard > 2.5? | / | | | |
| Does the maximum intensity of each specified characteristic ion coincide within ± 2 seconds (includes labeled standards)? | / | | | |
| Was an acceptable lock mass recorded and monitored? | / | | | |
| XI. Compound quantitation/CRQLs | | | | |
| Were the correct internal standard (IS), quantitation ion and relative response factor (RRF) used to quantitate the compound? | / | | | |
| Were compound quantitation and CRQLs adjusted to reflect all sample dilutions and dry weight factors applicable to level IV validation? | / | | | |
| XII. System performance | | | | |
| System performance was found to be acceptable. | / | | | |
| XIII. Overall assessment of data | | | | |
| Overall assessment of data was found to be acceptable. | / | | | |
| XIV. Field duplicates | | | | |
| Field duplicate pairs were identified in this SDG. | | / | | |
| Target compounds were detected in the field duplicates. | | | / | |
| XV. Field blanks | | | | |
| Field blanks were identified in this SDG. | | / | | |
| Target compounds were detected in the field blanks. | | | / | |

Ions Monitored for HRGC/HRMS Analysis of Polychlorinated Biphenyls

| Descriptor | Accurate mass ^(a) | Ion ID | Analyte | Substance | |
|------------|------------------------------|----------|----------------------|--------------------|-----|
| 1 | 289.9224 | M | C12 H6 35Cl4 | TCB | |
| | 291.9194 | M+2 | C12 H6 35Cl3 37Cl4 | TCB | |
| | 301.9626 | M | 13C12 H6 35Cl4 | PeCB | |
| | 303.9597 | M+2 | 13C12 H6 35Cl3 37Cl | PeCB | |
| | 325.8804 | M+2 | C12 H5 35Cl4 37Cl | PeCB | |
| | 327.8775 | M+4 | C12 H5 35Cl3 37Cl2 | PeCB | |
| | [292.9825] | Lock | C7 F11 | PFK | |
| 2 | 325.8804 | M+2 | C12 H5 35Cl4 37Cl | PeCB | |
| | 327.8775 | M+4 | C12 H5 35Cl3 37Cl2 | PeCB | |
| | 337.9207 | M+2 | 13C12 H5 35Cl4 37Cl | PeCB | |
| | 339.9178 | M+4 | 13C12 H5 35Cl3 37Cl2 | PeCB | |
| | 359.8415 | M+2 | C12 H4 35Cl5 37Cl | HxCB | |
| | 361.8385 | M+4 | C12 H4 35Cl4 37Cl2 | HxCB | |
| | 371.8817 | M+2 | 13C12 H4 35Cl5 37Cl | HxCB | |
| | 373.8788 | M+4 | 13C12 H4 35Cl4 37Cl2 | HxCB | |
| | 393.8025 | M+2 | C12 H3 35Cl6 37Cl | HpCB | |
| | 395.7996 | M+4 | C12 H3 35Cl5 37Cl2 | HpCB | |
| | 405.8428 | M+2 | 13C12 H3 35Cl6 37Cl | HpCB | |
| | 407.8398 | M+4 | 13C12 H3 35Cl5 37Cl2 | HpCB | |
| | [354.9892] | Lock | C9F13 | PFK | |
| | 3 | 509.7229 | M+4 | 13C12 35Cl10 37Cl2 | DCB |
| | | 511.7199 | M+6 | 13C12 35Cl9 37Cl3 | |
| 513.7170 | | M+8 | 13C12 35Cl8 37Cl4 | | |
| [442.9728] | | Lock | C10 F17 | PFK | |
| | | | | | |

S = internal/recovery standard

H = 1.007825
 C = 12.000000
¹³C = 13.003355
 F = 18.9984

³⁵Cl = 34.968853
³⁷Cl = 36.965903

LDC #: 1327A3
 SDG #: DPWF15252

VALIDATION FINDINGS WORKSHEET
Blanks

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".
 N N/A Were all samples associated with a method blank?
 N N/A Was a method blank performed for each matrix and wherever a sample extraction was performed?
 N N/A Was method blank contamination less < CRQL for all target compounds?

Blank extraction date: 1/25/05 Blank analysis date: 2/3/05
 Conc. units: ng/kg

Associated samples: M

| Compound | Blank ID | Sample Identification | | | | | | | | | |
|-----------------|---------------------|-----------------------|--|--|--|--|--|--|--|--|--|
| | 1745-101 | All | | | | | | | | | |
| PCB 66 | 0.108 | | | | | | | | | | |
| 77 | 0.042 | | | | | | | | | | |
| 90+101+113 | 0.336 | | | | | | | | | | |
| 105 | 0.078 | | | | | | | | | | |
| 110+115 | 0.076 ²⁸ | | | | | | | | | | |
| 114 | 0.044 | | | | | | | | | | |
| 118 | 0.195 | | | | | | | | | | |
| 123 | 0.027 | | | | | | | | | | |
| 126 | 0.041 | | | | | | | | | | |
| 129+138+160+163 | 0.310 | | | | | | | | | | |
| 153+168 | 0.257 | | | | | | | | | | |
| 156+157 | 0.088 | | | | | | | | | | |
| 180+193 | 0.160 | | | | | | | | | | |
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CIRCLED RESULTS WERE NOT QUALIFIED. ALL RESULTS NOT CIRCLED WERE QUALIFIED BY THE FOLLOWING STATEMENT:
 All contaminants within five times the method blank concentration were qualified as not detected, "U".

LDC #: 13247A3
 SDG #: DPW 415752

VALIDATION FINDINGS WORKSHEET
Matrix Spike/Matrix Spike Duplicates

Page: 1 of 1
 Reviewer: C
 2nd Reviewer: R

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Please see qualifications below for all questions answered "N". Not applicable questions are identified as "N/A".

Y N N/A Were a matrix spike (MS) and matrix spike duplicate (MSD) analyzed for each matrix in this SDG? If no, indicate which matrix does not have an associated MS/MSD. Soil / Water.

Y N N/A Was a MS/MSD analyzed every 20 samples of each matrix?

Y N N/A Were the MS/MSD concentrations and the relative percent differences (RPD) within the QC limits stated below?

| # | Date | MS/MSD ID | Compound | MS %R (Limits) | MSD %R (Limits) | RPD (Limits) | Associated Samples | Qualifications |
|---|------|-----------|--------------------|----------------|-----------------|--------------|--------------------|----------------|
| | 1/17 | | PCB 90+10+113 | () | () | 53.7 (≤50) | 1 | Jf adets / A |
| | | | PCB 129+138+160+63 | () | () | 75.6 () | | |
| | | | PCB 153+168 | () | () | 89.5 () | | |
| | | | PCB 156+157 | () | () | 60.8 () | | |
| | | | PCB 167 | () | () | 59.9 () | | |
| | | | PCB 180+193 | () | () | 112 () | | |
| | | | PCB 9#189 | () | () | 91.6 () | | |
| | | | | () | () | () | | |
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LDC #: 1324703
 SDG #: DPW/EIS/252

VALIDATION FINDINGS WORKSHEET
Initial Calibration Calculation Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The Relative Response Factor (RRF), average RRF, and percent relative standard deviation (%RSD) were recalculated for the compounds identified below using the following calculations:

$$RRF = (A_x)(C_s)/(A_s)(C_x)$$

average RRF = sum of the RRFs/number of standards

$$\%RSD = 100 * (S/X)$$

A_x = Area of compound,

C_x = Concentration of compound,

S = Standard deviation of the RRFs,

A_s = Area of associated internal standard

C_s = Concentration of internal standard

X = Mean of the RRFs

| # | Standard ID | Calibration Date | Compound (Reference Internal Standard) | Reported | Recalculated | Reported | Recalculated | Reported | Recalculated |
|---|-------------|------------------|--|-----------------------|-----------------------|-----------|--------------|----------|--------------|
| | | | | Average RRF (Initial) | Average RRF (Initial) | RRF (std) | RRF (std) | %RSD | %RSD |
| 1 | 1CAZ | 2/4/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 0.91 | 0.87 | 0.86 | 3.54 | 3.20 |
| | | | PCB-105 (¹³ C-PCB-105) | 0.83 | 0.83 | 0.82 | 0.82 | 3.00 | 3.19 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 0.98 | 0.96 | 0.96 | 1.52 | 1.61 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 0.79 | 0.79 | 0.79 | 2.01 | 1.65 |
| 2 | | | PCB-77 (¹³ C-PCB-77) | | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | | |
| 3 | | | PCB-77 (¹³ C-PCB-77) | | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | | |

Comments: Refer to Initial Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 1327A3
 SDG #: DPW 15252

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

Page: 1 of 1
 Reviewer: [Signature]
 2nd Reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

The percent difference (%D) of the initial calibration average Relative Response Factors (RRFs) and the continuing calibration RRFs were recalculated for the compounds identified below using the following calculation:

% Difference = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$
 $\text{RRF} = (A_x / C_x) / (A_s / C_s)$

Where: ave. RRF = initial calibration average RRF
 RRF = continuing calibration RRF
 A_x = Area of compound, A_s = Area of associated internal standard
 C_x = Concentration of compound, C_s = Concentration of internal standard

| # | Standard ID | Calibration Date | Compound (Reference Internal Standard) | Average RRF (Initial) | Reported | Recalculated | Reported | Recalculated |
|---|-------------|------------------|--|-----------------------|--------------|--------------|--------------|--------------|
| | | | | | RRF Amt (CC) | RRF Amt (CC) | %D | %D |
| 1 | PB5C-067 | 2/8/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 48.3 | 48.2 | Not reported | 3.6 |
| | | | PCB-105 (¹³ C-PCB-105) | 0.83 | 49.3 | 49.2 | | 1.6 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 98.6 | 98.5 | | 1.5 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 49.8 | 50.1 | | 0.2 |
| 2 | PB5C-069 | 2/9/05 | PCB-77 (¹³ C-PCB-77) | 0.91 | 48.8 | 48.7 | Not reported | 2.6 |
| | | | PCB-105 (¹³ C-PCB-105) | 0.83 | 50.5 | 50.2 | | 0.4 |
| | | | PCB-156 (¹³ C-PCB-156) | 0.98 | 100 | 100 | | 0 |
| | | | PCB-180 (¹³ C-PCB-180) | 0.79 | 49.2 | 49.6 | | 0.7 |
| 3 | | | PCB-77 (¹³ C-PCB-77) | | | | | |
| | | | PCB-105 (¹³ C-PCB-105) | | | | | |
| | | | PCB-156 (¹³ C-PCB-156) | | | | | |
| | | | PCB-180 (¹³ C-PCB-180) | | | | | |

Comments: Refer to Routine Calibration findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 13247A3
 SDG #: DPW/IS252

VALIDATION FINDINGS WORKSHEET
Laboratory Control Sample Results Verification

Page: 1 of 1

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Reviewer: [Signature]

The percent recoveries (%R) and Relative Percent Difference (RPD) of the laboratory control sample and laboratory control sample duplicate (if applicable) were recalculated for the compounds identified below using the following calculation:

2nd Reviewer: [Signature]

% Recovery = 100 * SSC/SA

Where: SSC = Spiked sample concentration
 SA = Spike added

RPD = |LCS - LCSD| * 2 / (LCS + LCSD)

LCS = Laboratory control sample percent recovery

LCSD = Laboratory control sample duplicate percent recovery

LCS ID: OPR

| Compound | Spike Added (ng/ml) | | Spiked Sample Concentration (ng/ml) | | LCS | | LCSD | | LCS/LCSD | |
|----------------|---------------------|------|-------------------------------------|------|------------------|---------|------------------|---------|----------|--------------|
| | LCS | LCSD | LCS | LCSD | Percent Recovery | | Percent Recovery | | RPD | |
| | | | | | Reported | Recalc. | Reported | Recalc. | Reported | Recalculated |
| PCB-77 | 50.0 | NA | 48.8 | NA | 97.6 | 97.6 | | | | |
| PCB-81 | ↓ | | 48.5 | | 97.0 | 97.0 | | | | |
| PCB-105 | ↓ | | 49.4 | | 98.9 | 98.8 | | | | |
| PCB-114 | ↓ | | 49.1 | | 98.1 | 98.2 | | | | |
| PCB-113 | ↓ | | 50.0 | | 100 | 100 | | | | |
| PCB-123 | ↓ | | 50.1 | | 100 | 100 | | | | |
| PCB-125 | ↓ | | 48.6 | | 97.2 | 97.2 | | | | |
| PCB-156 + 165T | 100 | | 97.9 | | 97.9 | 97.9 | | | | |
| PCB-157 | | | | | | | | | | |
| PCB-167 | 50.0 | | 49.6 | | 99.1 | 99.2 | | | | |
| PCB-169 | ↓ | | 49.3 | | 98.6 | 98.6 | | | | |
| PCB-170 | | | | | | | | | | |
| PCB-180 | | | | | | | | | | |
| PCB-189 | 50.0 | ↓ | 49.7 | ↓ | 99.5 | 99.4 | | | | |
| | | | | | | | | | | |
| | | | | | | | | | | |

Comments: Refer to Laboratory Control Sample findings worksheet for list of qualifications and associated samples when reported results do not agree within 10.0% of the recalculated results.

LDC #: 13247A3
SDG #: OPNG-15252

VALIDATION FINDINGS WORKSHEET Sample Calculation Verification

Page: 1 of 1
Reviewer: [Signature]
2nd reviewer: [Signature]

METHOD: HRGC/HRMS Polychlorinated Biphenyls (EPA Method 1668)

Y N N/A Were all reported results recalculated and verified for all level IV samples?
Y N N/A Were all recalculated results for detected target compounds agree within 10.0% of the reported results?

$$\text{Concentration} = \frac{(A_s)(I_s)(DF)}{(A_i)(RRF)(V_s)(\%S)}$$

- A_s = Area of the characteristic ion (EICP) for the compound to be measured
- A_i = Area of the characteristic ion (EICP) for the specific internal standard
- I_s = Amount of internal standard added in nanograms (ng)
- RRF = Relative response factor of the calibration standard.
- V_s = Volume or weight of sample pruged in milliliters (ml) or grams (g).
- DF = Dilution factor.
- %S = Percent solids, applicable to soils and solid matrices only.

Example:

Sample I.D. 8, PCBT7:

$$\begin{aligned} \text{Conc.} &= \frac{(3.31 \times 10^7) (2000) (1)}{(1.36 \times 10^8) (0.91) (10.8) ()} \\ &= 49.53 \text{ ng/kg} \end{aligned}$$

| # | Sample ID | Compound | Reported Concentration () | Calculated Concentration () | Qualification |
|---|-----------|----------|----------------------------|------------------------------|---------------|
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