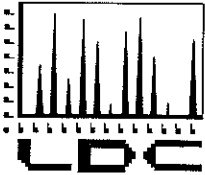


APPENDIX C DATA VALIDATION REPORT



LABORATORY DATA CONSULTANTS, INC.

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

LDC #15449

September 26, 2006

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

**SUBJECT: Lower Duwamish Waterway Group Fish Tissue Sample Data
Validation**

Dear Ms. Mitchell,

Enclosed is our final EPA 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: DPWG19975/WG19626. 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

Attachment 1

PDF **LDC #15449 (Windward Environmental, LLC - Seattle WA / Lower Duwamish Waterway Group)**

LDC	SDG#	DATE REC'D	(3) DATE DUE	PCB Cong. (1668)																												
					W	T	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S
Matrix: Water/Tissue																																
A	DPWG19975/WG19626	09/05/06	09/26/06	0	6																											
Total	B/SC			0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6	

CHEMICAL DATA QUALITY REVIEW FOR FISH TISSUE SAMPLES

Lower Duwamish Waterway Group LDC# 15449

This report details the findings of an EPA 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: DPWG19975/WG19626. 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 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 Fish and Crab Collection and Chemical Analyses Quality Assurance Project Plan (August 27, 2004). Where specific guidance is not available, the data has been evaluated in a conservative manner 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 Replicates

Attachment 1

PDF LDC #15449 (Windward Environmental, LLC - Seattle WA / Lower Duwamish Waterway Group)

LDC	SDG#	DATE REC'D	(3) DATE DUE	PCB Cong. (1668)																												
					W	T	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S
Matrix: Water/Tissue					W	T	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S	W	S		
A	DPWG19975/WG19626	09/05/06	09/26/06		0	6																										
Total	B/SC				0	6	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	6		

Attachment 2

SDG#: DPWG19975/WG19626

VALIDATION SAMPLE TABLE

LDC#: 15449A

Project Name: Lower Duwamish Waterway Group

Parameters/Analytical Method

Project #04-08-06-22

Client ID #	Lab ID #	Matrix	Date Collected	PCBs Cong. (1668A)													
LDW-05-T2-B-SS-WB-COMP1	L9071-1	tissue	09/01/05	X													
LDW-05-T3-D-SS-WB-COMP1	L9071-2	tissue	08/31/06	X													
LDW-05-T1-M-ES-WB-COMP3	L9071-3	tissue	08/29/06	X													
LDW-05-T2-M-ES-WB-COMP3	L9071-4	tissue	09/01/06	X													
LDW-05-T3-M-ES-WB-COMP2	L9071-5	tissue	09/01/06	X													
LDW-05-T1-B-SS-WB-COMP1	L9071-6	tissue	08/31/06	X													
LDW-05-T3-M-ES-WB-COMP2DUP	L9071-5DUP	tissue	09/01/06	X													

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.
 - 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

I. Method Compliance

The ion abundance ratio for PCB-169 in the lowest standard of the initial calibration did not meet the method QC limits.

II. Usability

The quality control criteria reviewed were met and are considered acceptable. Based upon the data validation all results are considered valid and usable for all purposes.

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 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).

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 ion abundance ratios for all PCBs were within validation criteria with the following exceptions:

SDG	Date	Compound	Ion Abundance Ratio (Limits)	Associated Samples	Flag	A or P
DPWG19975/ WG19626	5/5/06 (CS0 Standard)	PCB-169	0.99 (1.05-1.43)	LDW-05-T2-B-SS-WB-COMP1 LDW-05-T3-D-SS-WB-COMP1 LDW-05-T1-M-ES-WB-COMP3 LDW-05-T2-M-ES-WB-COMP3	NA	-

N/A = Not applicable

For the result above flagged "Not applicable", since the ion abundance ratio in the continuing calibration was within the method QC limits, this finding did not warrant the qualification of the data.

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.

IV. Routine Calibration (Continuing)

Routine calibration was performed at the required frequency.

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.

The ion abundance ratios for all PCBs were within validation criteria.

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:

SDG	Method Blank ID	Extraction Date	Compound	Concentration	Associated Samples
DPWG19975/ WG19626	WG19626-101	7/14/06	PCB-18	0.802 ng/Kg	LDW-05-T2-B-SS-WB-COMP1 LDW-05-T3-D-SS-WB-COMP1 LDW-05-T1-M-ES-WB-COMP3 LDW-05-T2-M-ES-WB-COMP3 LDW-05-T3-M-ES-WB-COMP2 LDW-05-T1-B-SS-WB-COMP1 LDW-05-T3-M-ES-WB-COMP2DUP
			PCB-20	1.70 ng/Kg	
			PCB-40	0.638 ng/Kg	
			PCB-44	2.67 ng/Kg	
			PCB-45	0.870 ng/Kg	
			PCB-49	1.66 ng/Kg	
			PCB-50	0.381 ng/Kg	
			PCB-52	2.54 ng/Kg	
			PCB-59	0.362 ng/Kg	
			PCB-60	0.608 ng/Kg	
			PCB-61	2.81 ng/Kg	
			PCB-64	0.414 ng/Kg	
			PCB-66	1.60 ng/Kg	
			PCB-82	0.782 ng/Kg	
			PCB-83	1.81 ng/Kg	
			PCB-86	0.766 ng/Kg	
			PCB-88	0.472 ng/Kg	
			PCB-90	4.46 ng/Kg	
			PCB-92	0.826 ng/Kg	
			PCB-93	4.20 ng/Kg	
			PCB-105	1.07 ng/Kg	
			PCB-109	0.577 ng/Kg	
			PCB-110	2.05 ng/Kg	
			PCB-118	2.28 ng/Kg	
			PCB-128	0.498 ng/Kg	
			PCB-129	2.46 ng/Kg	
			PCB-131	0.440 ng/Kg	
			PCB-132	0.704 ng/Kg	
			PCB-135	1.61 ng/Kg	
			PCB-137	0.424 ng/Kg	
			PCB-141	0.745 ng/Kg	
			PCB-146	1.02 ng/Kg	
			PCB-147	1.72 ng/Kg	
			PCB-153	2.72 ng/Kg	
PCB-156	1.13 ng/Kg				
PCB-167	0.392 ng/Kg				
PCB-170	0.526 ng/Kg				
PCB-180	1.47 ng/Kg				
PCB-183	0.412 ng/Kg				
PCB-187	0.662 ng/Kg				
PCB-190	0.735 ng/Kg				
PCB-194	0.757 ng/Kg				
PCB-198	0.575 ng/Kg				
PCB-81	0.115 ng/Kg				
PCB-169	0.079 ng/Kg				
Total tetrachlorobiphenyls	5.05 ng/Kg				
Total pentachlorobiphenyls	2.28 ng/Kg				
Total hexachlorobiphenyls	4.44 ng/Kg				
Total octachlorobiphenyls	0.757 ng/Kg				
Total PCBs	12.5 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.

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.

XI. Compound Quantitation and CRQLs

All compound quantitation and CRQLs were within validation criteria with the following exceptions:

SDG	Sample	Compound	Finding	Criteria	Flag	A or P
DPWG19975/ WG19626	LDW-05-T2-M-ES-WB-COMP3	PCB-118	Sample result exceeded calibration range.	Reported result should be within calibration range.	J2 (all detects)	P

XII. System Performance

The system performance was acceptable.

XIII. Overall Assessment of Data

Data flags are summarized at the end of this report if data has been qualified.

XIV. Field Replicates

No field replicates were identified in this SDG.

**Lower Duwamish Waterway Group
Polychlorinated Biphenyls - Data Qualification Summary - SDG DPWG19975/WG19626**

SDG	Sample	Compound	Flag	A or P	Reason
DPWG19975/ WG19626	LDW-05-T2-M-ES-WB-COMP3	PCB-118	J2 (all detects)	P	Compound quantitation and CRQLs

**Lower Duwamish Waterway Group
Polychlorinated Biphenyls - Laboratory Blank Data Qualification Summary - SDG
DPWG19975/WG19626**

No Sample Data Qualified in this SDG

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		Comments
I.	Technical holding times	A	Sampling dates: 8/29 - 9/1/05
II.	GC/MS Instrument performance check	A	
III.	Initial calibration	SPW	70 RSD ≤ 20
IV.	Routine calibration	A	70 D ≤ 25/35 (unline/labeled)
V.	Blanks	SW	
VI.	Matrix spike/Matrix spike duplicates	DUP	N/A
VII.	Laboratory control samples	A	LES. CRM
VIII.	Regional quality assurance and quality control	N	
IX.	Internal standards	SPW	
X.	Target compound identifications	A	
XI.	Compound quantitation and CRQLs	SW	
XII.	System performance	A	
XIII.	Overall assessment of data	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:
 All TISSUES

1	LDW-05-T2-B-SS-WB-COMP1	11	WG19626-10	21		31	
2	LDW-05-T3-D-SS-WB-COMP1	12		22		32	
3	LDW-05-T1-M-ES-WB-COMP3	13		23		33	
4	LDW-05-T2-M-ES-WB-COMP3	14		24		34	
5	LDW-05-T3-M-ES-WB-COMP2	15		25		35	
6	LDW-05-T1-B-SS-WB-COMP1	16		26		36	
7	LDW-05-T3-M-ES-WB-COMP2DUP	17		27		37	
8		18		28		38	
9		19		29		39	
10		20		30		40	

LDC #: 16449A3
 SDG #: DPWG19975

VALIDATION FINDINGS CHECKLIST

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

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 $\leq 30\%$ for labeled standards?	/			
Did all calibration standards meet the Ion Abundance Ratio criteria?		/		
Was the signal to noise ratio for each target compound ≥ 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 #: 15419A3
 SDG #: DPW#19975

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.			/	

LDC #: 15449A3
 SDG #: DPW199TS

VALIDATION FINDINGS WORKSHEET

Initial Calibration

Page: 1 of 1
 Reviewer: Q
 2nd Reviewer: a

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 Was the initial calibration performed at 5 concentration levels?
- N N/A Were all percent relative standard deviations (%RSD) $\leq 65\%$ for unlabeled standards and labeled standards?
- N N/A Did all calibration standards meet the Ion Abundance Ratio criteria?
- N N/A Was the signal to noise ratio for each target compound ≥ 2.5 and for each recovery and internal standard ≥ 10 ?

#	Date	Standard ID	Compound	Finding %RSD	Finding Ion Abundance Ratio	Associated Samples	Qualifications
	5/5/06	DT63-123B (CSO)	PCB169		0.99 (1.05-1.43)	1-4, Bk	NA For No label (ccv in)

Halogen	Selected ions (m/z)	Ion Abundance Ratio	Halogen	Selected ions (m/z)	Ion Abundance Ratio
1 Cl	M/M+2	2.66-3.60	7 Cl	M/M+2	0.37-0.51
2 Cl	M/M+2	1.33-1.81	7 Cl	M+2/M+4	0.88-1.20
3 Cl	M/M+2	0.88-1.20	8 Cl	M+2/M+4	0.76-1.02
4 Cl	M/M+2	0.65-0.89	9 Cl	M/M+2	1.14-1.54
5 Cl	M+2/M+4	1.32-1.78	9 Cl	M/M-2	0.66-0.90
6 Cl	M/M+2	0.43-0.59	10 Cl	M/M+2	0.99-1.35
6 Cl	M+2/M+4	1.05-1.43			

LDC #: 10119A3
SDG #: DPIN&199TS

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: 7/14/06 Blank analysis date: 7/24/06

Conc. units: ng/kg Associated samples: ML

Compound	Blank ID	Sample Identification									
<i>see attachment</i>	<u>9626-10</u>	<u>All</u>									
		<u>75X</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".

AXYS METHOD MLA-010 Rev 08

Form 1A
PCB CONGENER ANALYSIS REPORT

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

AXYS ANALYTICAL SERVICES

P.O. Box 2219, 2045 MILLS RD. WEST, SIDNEY, B.C., CANADA
V8L 3S8 TEL (250) 655-5800 FAX (250) 655-5811
Contract No.: 4033

Project No. N/A
Lab Sample I.D.: WG19626-101
Sample Size: 5.00 g
Initial Calibration Date: 05-Jun-2006
Instrument ID: HR GC/MS
GC Column ID: SPB OCTYL
Sample Data Filename: PB6C_327 S: 5
Blank Data Filename: PB6C_327 S: 5
Cal. Ver. Data Filename: PB6C_327 S: 1

Matrix: CORN OIL
Sample Receipt Date: N/A
Extraction Date: 14-Jul-2006
Analysis Date: 24-Jul-2006 Time: 11:52:21
Extract Volume (uL): 400
Injection Volume (uL): 1.0
Dilution Factor: 20
Concentration Units: ng/kg

COMPOUND	IUPAC NO.	CO-ELUTIONS	LAB FLAG ¹	CONC. FOUND	REPORTING LIMIT	ION ABUND. RATIO	RRT
2-MoCB	1		UD		0.472		
3-MoCB	2		UD		0.546		
4-MoCB	3		UD		0.664		
2,2'-DiCB	4		UD		2.97		
2,3-DiCB	5		UD		2.46		
2,3'-DiCB	6		UD		2.26		
2,4-DiCB	7		UD		2.32		
2,4'-DiCB	8		UD		2.14		
2,5-DiCB	9		UD		2.24		
2,6-DiCB	10		UD		2.31		
3,3'-DiCB	11		UD		2.37		
3,4-DiCB	12	12 + 13	CUD		2.47		
3,4'-DiCB	13	12 + 13	C12				
3,5-DiCB	14		UD		2.31		
4,4'-DiCB	15		UD		3.15		
2,2',3-TrICB	16		UD		1.09		
2,2',4-TrICB	17		UD		0.922		
2,2',5-TrICB	18	18 + 30	4.01 CKDJ	0.802	0.771	1.51	1.112
2,2',6-TrICB	19		UD		1.07		
2,3,3'-TriCB	20	20 + 28	8.50 CKDJ	1.70	0.982	0.83	0.848
2,3,4-TriCB	21	21 + 33	CUD		0.955		
2,3,4'-TriCB	22		UD		1.07		
2,3,5-TriCB	23		UD		1.02		
2,3,6-TriCB	24		UD		0.693		
2,3',4-TriCB	25		UD		0.879		
2,3',5-TriCB	26	26 + 29	CUD		0.980		
2,3',6-TriCB	27		UD		0.634		
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		UD		0.934		
2,4',6-TriCB	32		UD		0.962		
2',3,4-TriCB	33	21 + 33	C21				
2',3,5-TriCB	34		UD		0.999		
3,3',4-TriCB	35		UD		1.10		
3,3',5-TriCB	36		UD		0.987		
3,4,4'-TriCB	37		UD		1.18		
3,4,5-TriCB	38		UD		1.03		



COMPOUND	IUPAC NO.	CO-ELUTIONS	5X	LAB FLAG ¹	CONC. FOUND	REPORTING LIMIT	ION ABUND. RATIO	RRT
3,4',5'-TriCB	39			UD		1.02		
2,2',3,3'-TeCB	40	40 + 41 + 71	3.19	CDJ	0.638	0.269	0.70	1.334
2,2',3,4'-TeCB	41	40 + 41 + 71		C40				
2,2',3,4'-TeCB	42			UD		0.282		
2,2',3,5'-TeCB	43			UD		0.328		
2,2',3,5'-TeCB	44	44 + 47 + 65	13.35	CKDJ	2.67	0.245	0.63	1.285
2,2',3,6'-TeCB	45	45 + 51	4.35	CKDJ	0.870	0.274	1.08	1.145
2,2',3,6'-TeCB	46			UD		0.316		
2,2',4,4'-TeCB	47	44 + 47 + 65		C44				
2,2',4,5'-TeCB	48			UD		0.274		
2,2',4,5'-TeCB	49	49 + 69	8.3	CKDJ	1.66	0.229	0.58	1.258
2,2',4,6'-TeCB	50	50 + 53	1.905	CKDJ	0.381	0.266	3.90	1.110
2,2',4,6'-TeCB	51	45 + 51		C45				
2,2',5,5'-TeCB	52		12.7	KDJ	2.54	0.258	1.24	1.233
2,2',5,6'-TeCB	53	50 + 53		C50				
2,2',6,6'-TeCB	54			UD		0.244		
2,3,3',4'-TeCB	55			UD		0.533		
2,3,3',4'-TeCB	56			UD		0.518		
2,3,3',5'-TeCB	57			UD		0.494		
2,3,3',5'-TeCB	58			UD		0.507		
2,3,3',6'-TeCB	59	59 + 62 + 75	1.81	CKDJ	0.362	0.204	0.96	1.300
2,3,4,4'-TeCB	60		3.04	KDJ	0.608	0.535	1.24	0.912
2,3,4,5'-TeCB	61	61 + 70 + 74 + 76	14.05	CDJ	2.81	0.492	0.73	0.875
2,3,4,6'-TeCB	62	59 + 62 + 75		C59				
2,3,4',5'-TeCB	63			UD		0.476		
2,3,4',6'-TeCB	64		2.07	KDJ	0.414	0.196	3.38	1.347
2,3,5,6'-TeCB	65	44 + 47 + 65		C44				
2,3',4,4'-TeCB	66		8.0	DJ	1.60	0.499	0.82	0.885
2,3',4,5'-TeCB	67			UD		0.431		
2,3',4,5'-TeCB	68			UD		0.477		
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			UD		0.469		
2,3',5,6'-TeCB	73			UD		0.199		
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			UD		0.534		
3,3',4,5'-TeCB	78			UD		0.532		
3,3',4,5'-TeCB	79			UD		0.437		
3,3',5,5'-TeCB	80			UD		0.463		
3,4,4',5'-TeCB	81			X				
2,2',3,3',4'-PeCB	82		3.91	KDJ	0.782	0.464	0.62	0.934
2,2',3,3',5'-PeCB	83	83 + 99	9.05	CKDJ	1.81	0.421	0.73	0.885
2,2',3,3',6'-PeCB	84			UD		0.471		
2,2',3,4,4'-PeCB	85	85 + 116 + 117		CUD		0.345		
2,2',3,4,5'-PeCB	86	86 + 87 + 97 + 108 + 119 + 125	383	CKDJ	0.766	0.358	1.87	0.899
2,2',3,4,5'-PeCB	87	86 + 87 + 97 + 108 + 119 + 125		C86				
2,2',3,4,6'-PeCB	88	88 + 91	2.36	CKDJ	0.472	0.417	0.38	1.151
2,2',3,4,6'-PeCB	89			UD		0.436		
2,2',3,4',5'-PeCB	90	90 + 101 + 113	22.3	CKDJ	4.46	0.363	1.28	0.869
2,2',3,4',6'-PeCB	91	88 + 91		C88				
2,2',3,5,5'-PeCB	92		4.13	KDJ	0.826	0.421	2.34	0.853
2,2',3,5,6'-PeCB	93	93 + 95 + 98 + 100 + 102	21	CKDJ	4.20	0.404	1.82	1.123
2,2',3,5,6'-PeCB	94			UD		0.455		
2,2',3,5,6'-PeCB	95	93 + 95 + 98 + 100 + 102		C93				
2,2',3,6,6'-PeCB	96			UD		0.247		
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				



COMPOUND	IUPAC NO.	CO-ELUTIONS	5X LAB FLAG ¹	CONC. FOUND	REPORTING LIMIT	ION ABUND. RATIO	RRT
2,2',4,5',6-PeCB	103		UD		0.378		
2,2',4,6,6'-PeCB	104		UD		0.277		
2,3,3',4,4'-PeCB	105		5.35 KDJ	1.07	0.458	2.47	1.000
2,3,3',4,5-PeCB	106		UD		0.415		
2,3,3',4',5-PeCB	107	107 + 124	CUD		0.428		
2,3,3',4,5'-PeCB	108	86 + 87 + 97 + 108 + 119 + 125	C86				
2,3,3',4,6-PeCB	109		2.885 KDJ	0.577	0.384	3.41	0.997
2,3,3',4',6-PeCB	110	110 + 115	10.25 CKDJ	2.05	0.303	2.24	0.925
2,3,3',5,5'-PeCB	111		UD		0.312		
2,3,3',5,6-PeCB	112		UD		0.303		
2,3,3',5',6-PeCB	113	90 + 101 + 113	C90				
2,3,4,4',5-PeCB	114		UD		0.414		
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		11.4 DJ	2.28	0.428	1.60	1.000
2,3',4,4',6-PeCB	119	86 + 87 + 97 + 108 + 119 + 125	C86				
2,3',4,5,5'-PeCB	120		UD		0.295		
2,3',4,5',6-PeCB	121		UD		0.317		
2',3,3',4,5-PeCB	122		UD		0.451		
2',3,4,4',5-PeCB	123		UD		0.561		
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		X				
3,3',4,5,5'-PeCB	127		UD		0.440		
2,2',3,3',4,4'-HxCB	128	128 + 166	2.49 CKDJ	0.498	0.253	5.15	0.959
2,2',3,3',4,5-HxCB	129	129 + 138 + 160 + 163	12.3 CKDJ	2.46	0.254	0.93	0.929
2,2',3,3',4,5'-HxCB	130		UD		0.317		
2,2',3,3',4,6-HxCB	131		2.2 KDJ	0.440	0.301	1.52	1.160
2,2',3,3',4,6'-HxCB	132		3.52 KDJ	0.704	0.315	0.51	1.174
2,2',3,3',5,5'-HxCB	133		UD		0.289		
2,2',3,3',5,6-HxCB	134	134 + 143	CUD		0.297		
2,2',3,3',5,6'-HxCB	135	135 + 151 + 154	8.05 CKDJ	1.61	0.338	0.96	1.102
2,2',3,3',6,6'-HxCB	136		UD		0.264		
2,2',3,4,4',5-HxCB	137		2.12 KDJ	0.424	0.291	0.21	0.918
2,2',3,4,4',5'-HxCB	138	129 + 138 + 160 + 163	C129				
2,2',3,4,4',6-HxCB	139	139 + 140	CUD		0.265		
2,2',3,4,4',6'-HxCB	140	139 + 140	C139				
2,2',3,4,5,5'-HxCB	141		3.725 KDJ	0.745	0.273	0.88	0.904
2,2',3,4,5,6-HxCB	142		UD		0.307		
2,2',3,4,5,6'-HxCB	143	134 + 143	C134				
2,2',3,4,5',6-HxCB	144		UD		0.347		
2,2',3,4,6,6'-HxCB	145		UD		0.277		
2,2',3,4',5,5'-HxCB	146		5.1 KDJ	1.02	0.249	1.03	0.885
2,2',3,4',5,6-HxCB	147	147 + 149	8.6 CDJ	1.72	0.262	1.20	1.133
2,2',3,4',5,6'-HxCB	148		UD		0.356		
2,2',3,4',5',6-HxCB	149	147 + 149	C147				
2,2',3,4',6,6'-HxCB	150		UD		0.265		
2,2',3,5,5',6-HxCB	151	135 + 151 + 154	C135				
2,2',3,5,6,6'-HxCB	152		UD		0.257		
2,2',4,4',5,5'-HxCB	153	153 + 168	13.6 CDJ	2.72	0.222	1.21	0.899
2,2',4,4',5,6'-HxCB	154	135 + 151 + 154	C135				
2,2',4,4',6,6'-HxCB	155		UD		0.248		
2,3,3',4,4',5-HxCB	156	156 + 157	5.65 CKDJ	1.13	0.294	0.88	1.000
2,3,3',4,4',5'-HxCB	157	156 + 157	C156				
2,3,3',4,4',6-HxCB	158		UD		0.198		
2,3,3',4,5,5'-HxCB	159		UD		0.218		
2,3,3',4,5,6-HxCB	160	129 + 138 + 160 + 163	C129				
2,3,3',4,5',6-HxCB	161		UD		0.209		
2,3,3',4',5,5'-HxCB	162		UD		0.222		
2,3,3',4',5,6-HxCB	163	129 + 138 + 160 + 163	C129				
2,3,3',4',5',6-HxCB	164		UD		0.212		
2,3,3',5,5',6-HxCB	165		UD		0.233		
2,3,4,4',5,6-HxCB	166	128 + 166	C128				



COMPOUND	IUPAC NO.	CO-ELUTIONS	LAB FLAG ¹	CONC. FOUND	REPORTING LIMIT	ION ABUND. RATIO	RRT
2,3',4,4',5,5'-HxCB	167						
2,3',4,4',5',6-HxCB	168	153 + 168	1.96 KDJ	0.392	0.210	0.54	1.001
3,3',4,4',5,5'-HxCB	169		C153				
2,2',3,3',4,4',5-HpCB	170		X				
2,2',3,3',4,4',6-HpCB	171	171 + 173	2.63 KDJ	0.526	0.407	2.94	0.936
2,2',3,3',4,5,5'-HpCB	172		CUD		0.389		
2,2',3,3',4,5,6-HpCB	173	171 + 173	UD		0.393		
2,2',3,3',4,5,6'-HpCB	174		C171				
2,2',3,3',4,5',6-HpCB	175		UD		0.356		
2,2',3,3',4,6,6'-HpCB	176		UD		0.343		
2,2',3,3',4,5,6-HpCB	177		UD		0.262		
2,2',3,3',5,5',6-HpCB	178		UD		0.370		
2,2',3,3',5,6,6'-HpCB	179		UD		0.365		
2,2',3,4,4',5,5'-HpCB	180	180 + 193	7.35 CKDJ	1.47	0.309	2.04	0.910
2,2',3,4,4',5,6-HpCB	181		UD		0.367		
2,2',3,4,4',5,6'-HpCB	182		UD		0.336		
2,2',3,4,4',5',6-HpCB	183	183 + 185	2.06 CKDJ	0.412	0.349	0.65	1.127
2,2',3,4,4',6,6'-HpCB	184		UD		0.254		
2,2',3,4,5,5',6-HpCB	185	183 + 185	C183				
2,2',3,4,5,6,6'-HpCB	186		UD		0.271		
2,2',3,4',5,5',6-HpCB	187		3.315 KDJ	0.663	0.326	0.43	1.109
2,2',3,4',5,6,6'-HpCB	188		UD		0.267		
2,3,3',4,4',5,5'-HpCB	189		UD		0.529		
2,3,3',4,4',5,6-HpCB	190		3.675 KDJ	0.735	0.302	1.52	0.947
2,3,3',4,4',5',6-HpCB	191		UD		0.291		
2,3,3',4,5,5',6-HpCB	192		UD		0.316		
2,3,3',4',5,5',6-HpCB	193	180 + 193	C180				
2,2',3,3',4,4',5,5'-OoCB	194		3.785 DJ	0.757	0.572	0.92	0.992
2,2',3,3',4,4',5,6-OoCB	195		UD		0.595		
2,2',3,3',4,4',5,6'-OoCB	196		UD		0.445		
2,2',3,3',4,4',6,6'-OoCB	197	197 + 200	UD		0.317		
2,2',3,3',4,5,5',6-OoCB	198	198 + 199	2.875 CKDJ	0.575	0.461	2.04	1.115
2,2',3,3',4,5,5',6'-OoCB	199	198 + 199	C198				
2,2',3,3',4,5,6,6'-OoCB	200	197 + 200	C197				
2,2',3,3',4,5',6,6'-OoCB	201		UD		0.313		
2,2',3,3',5,5',6,6'-OoCB	202		UD		0.403		
2,2',3,4,4',5,5',6-OoCB	203		UD		0.430		
2,2',3,4,4',5,6,6'-OoCB	204		UD		0.315		
2,3,3',4,4',5,5',6-OoCB	205		UD		0.472		
2,2',3,3',4,4',5,5',6-NoCB	206		UD		2.12		
2,2',3,3',4,4',5,6,6'-NoCB	207		UD		1.48		
2,2',3,3',4,5,5',6,6'-NoCB	208		UD		1.69		
2,2',3,3',4,4',5,5',6,6'-DeCB	209		UD		0.705		

(1) Where applicable, custom lab flags have been used on this report; U = not detected; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; D = dilution data; J = concentration less than LMCL; C = co-eluting congener; X = result reported separately.

Approved by: Teresa Rawsthorne QA/QC Chemist

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



AXYS METHOD MLA-010 Rev 08

Form 1A
PCB CONGENER ANALYSIS REPORT

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

AXYS ANALYTICAL SERVICES
P.O. Box 2219, 2045 MILLS RD. WEST, SIDNEY, B.C., CANADA
V8L 3S8 TEL (250) 655-5800 FAX (250) 655-5811

Contract No.: 4033

Project No. N/A
Lab Sample I.D.: WG19626-101

Matrix: CORN OIL

Sample Size: 5.00 g

Sample Receipt Date: N/A

Initial Calibration Date: 08-May-2006

Extraction Date: 14-Jul-2006

Instrument ID: HR GC/MS

Analysis Date: 01-Aug-2006 Time: 01:12:21

GC Column ID: DB1

Extract Volume (uL): 20

Sample Data Filename: DT63_189B S: 4

Injection Volume (uL): 2.0

Blank Data Filename: DT63_189B S: 4

Dilution Factor: N/A

Cal. Ver. Data Filename: DT63_189B S: 1

Concentration Units: ng/kg

COMPOUND	IUPAC NO.	CO-ELUTIONS	LAB FLAG ¹	CONC. FOUND	REPORTING LIMIT	ION ABUND. RATIO	RRT
3,3',4,4'-TeCB	77		X				
3,4,4',5'-TeCB	81	5X	KJ	0.115	0.0634	23.96	1.000
2,3,3',4,4'-PeCB	105		X				
2,3,4,4',5'-PeCB	114		X				
2,3',4,4',5'-PeCB	118		X				
2',3,4,4',5'-PeCB	123		X				
3,3',4,4',5'-PeCB	126		U		0.157		
2,3,3',4,4',5'-HxCB	156		X				
2,3,3',4,4',5'-HxCB	157		X				
2,3',4,4',5,5'-HxCB	167		X				
3,3',4,4',5,5'-HxCB	169	0.393	KJ	0.079	0.0427	0.92	1.000
2,2',3,3',4,4',5'-HpCB	170		X				
2,2',3,4,4',5,5'-HpCB	180		X				
2,3,3',4,4',5,5'-HpCB	189		X				

(1) Where applicable, custom lab flags have been used on this report; U = not detected; K = peak detected but did not meet quantification criteria, result reported represents the estimated maximum possible concentration; J = concentration less than LMCL; X = result reported separately.

Approved by: Teresa Rawsthorne QA/QC Chemist

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



AXYS METHOD MLA-010 Rev 08

Form 1A
HOMOLOGUE TOTAL PCB ANALYSIS REPORT

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

AXYS ANALYTICAL SERVICES

P.O. Box 2219, 2045 MILLS RD. WEST, SIDNEY, B.C., CANADA
VBL 3S8 TEL (250) 655-5800 FAX (250) 655-5811
Contract No.: 4033

Project No. N/A
Lab Sample I.D.: WG19626-101
Sample Size: 5.00 g
Initial Calibration Date: 05-Jun-2006
Instrument ID: HR GC/MS
GC Column ID: DB1, SPB OCTYL
Sample Data Filename(s): DT63_189B S: 4, PB6C_327 S: 5
Blank Data Filename: PB6C_327 S: 5
Cal. Ver. Data Filename: PB6C_327 S: 1

Matrix: CORN OIL
Sample Receipt Date: N/A
Extraction Date: 14-Jul-2006
Analysis Date: 24-Jul-2006 Time: 11:52:21
Extract Volume (uL): 400
Injection Volume (uL): 1.0
Dilution Factor: 20
Concentration Units: ng/kg

PCB HOMOLOGUE GROUP	LAB FLAG ¹	CONC. FOUND	
Total Monochloro Biphenyls	U		<u>5X</u>
Total Dichloro Biphenyls	U		
Total Trichloro Biphenyls	U		
Total Tetrachloro Biphenyls		5.05	25.25
Total Pentachloro Biphenyls		2.28	11.4
Total Hexachloro Biphenyls		4.44	22.2
Total Heptachloro Biphenyls	U		
Total Octachloro Biphenyls		0.757	3.785
Total Nonachloro Biphenyls	U		
Decachloro Biphenyl	U		
TOTAL PCBs		12.5	62.5

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

Approved by: Teresa Rawsthorne QA/QC Chemist

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



LDC #: LS449A2
SDG #: DNV/G19975/WG19625

VALIDATION FINDINGS WORKSHEET

Compound Quantitation and Reported CRQLs

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

METHOD: HRGC/HRMS PCB Congeners (EPA Method 1668)

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

- N N/A Were the correct internal standard (IS), quantitation ions and relative response factors (RRF) used to quantitate the compound?
 N N/A Compound quantitation and CRQLs were adjusted to reflect all sample dilutions and dry weight factors (if necessary).

#	Date	Sample ID	Finding	Associated Samples	Qualifications
		4	PCB 118 > calib range		J2 detg/P

Comments: See sample calculation verification worksheet for recalculations

LDC #: 15419A3
 SDG #: OPNS 19975

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 (CS3 std)	RRF (CS3 std)	%RSD	%RSD
1	ICAL	5/30/06	PCB-77 (¹³ C-PCB-77)	1.04	1.04	1.11	1.10	3.72	3.82
			PCB-105 (¹³ C-PCB-105)	0.94	0.94	1.00	1.00	5.08	4.91
			PCB-156 (¹³ C-PCB-156)	0.96	0.96	1.04	1.03	4.53	4.81
			PCB-180 (¹³ C-PCB-180)	0.93	0.93	0.99	0.98	3.81	3.93
2	ICAL	5/30/06	PCB-77 (¹³ C-PCB-77)	0.88	0.88	0.94	0.94	4.01	3.96
			PCB-105 (¹³ C-PCB-105) (26)	0.91	0.91	1.00	1.00	7.64	7.42
			PCB-156 (¹³ C-PCB-156) (69)	0.86	0.86	0.92	0.92	5.00	4.84
			PCB-180 (¹³ C-PCB-180)						
3	ICAL	5/30/06	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 #: 15449A3
 SDG #: DPW/19975

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

Page: 1 of
 Reviewer:
 2nd Reviewer:

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

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	DB6C-327 S=1 6/5	7/24/06	PCB-77 (¹³ C-PCB-77)	1.04	47.3	47.2		
			PCB-105 (¹³ C-PCB-105)	0.94	51.5	51.5		
			PCB-156 (¹³ C-PCB-156)	0.96	104	104		
			PCB-180 (¹³ C-PCB-180)	0.93	47.7	47.9		
2	DB63-189B S=1 5/8	8/	PCB-77 (¹³ C-PCB-77-81)	0.88	49.6	49.8		
			PCB-105 (¹³ C-PCB-105-126)	0.91	42.1	42.6		
			PCB-156 (¹³ C-PCB-156-169)	0.86	49.3	49.6		
			PCB-180 (¹³ C-PCB-180)					
3	DB6C-330 S=1 5/30	7/26/06	PCB-77 (¹³ C-PCB-77)	1.04	47.4	47.3		
			PCB-105 (¹³ C-PCB-105)	0.94	52.8	53.0		
			PCB-156 (¹³ C-PCB-156)	0.96	107	107		
			PCB-180 (¹³ C-PCB-180)	0.93	49.3	49.6		

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 #: 15117A3
 SDG #: DPN19975

VALIDATION FINDINGS WORKSHEET
Routine Calibration Results Verification

Page: 2 of 2
 Reviewer: 9
 2nd Reviewer: AL

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 = $100 * (\text{ave. RRF} - \text{RRF}) / \text{ave. RRF}$
 $\text{RRF} = (A_x)(C_u) / (A_u)(C_x)$

Where: ave. RRF = initial calibration average RRF
 RRF = continuing calibration RRF
 A_x = Area of compound, A_u = Area of associated internal standard
 C_x = Concentration of compound, C_u = 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	FB6C-328A S=2 6/5	7/24/06	PCB-77 (¹² C-PCB-77)	1.04	47.7			
			PCB-105 (¹³ C-PCB-105)	0.94	53.9			
			PCB-156 (¹³ C-PCB-156)	0.96	10.7			
			PCB-180 (¹³ C-PCB-180)	0.93	48.5			
2	FB6C-33C S=1	7/27/06	PCB-77 (¹² C-PCB-77)	1.04	46.9	46.9		
			PCB-105 (¹³ C-PCB-105)	0.94	52.6	52.7		
			PCB-156 (¹³ C-PCB-156)	0.96	106	106		
			PCB-180 (¹³ C-PCB-180)	0.93	48.2	48.4		
3	FB6C-338A S=2	7/24/06	PCB-77 (¹² C-PCB-77)	1.04	46.6	46.7		
			PCB-105 (¹³ C-PCB-105)	0.94	53.9	54.0		
			PCB-156 (¹³ C-PCB-156)	0.96	106	106		
			PCB-180 (¹³ C-PCB-180)	0.93	47.8	48.0		

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 #: 15A9A3
 SDG #: DPW 19975

VALIDATION FINDINGS WORKSHEET
Laboratory Control Sample Results Verification

Page: 1 of 1

Reviewer: [Signature]

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

2nd 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:

% 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: W19026-103

Compound	Spike Added (ns/ml)		Spiked Sample Concentration (ns/ml)		LCS		LCSD		LCS/LCSD	
	LCS	LCSD	LCS	LCSD	Percent Recovery		Percent Recovery		RPD	
					Reported	Recalc.	Reported	Recalc.	Reported	Recalculated
PCB-77	100	100	97.0		97.0	97.0				
PCB-81			101		101	101				
PCB-105			99.9		99.9	99.9				
PCB-114			104		104	104				
PCB-118			104		104	104				
PCB-123			104		104	104				
PCB-126			108		108	108				
PCB-156	200		218		109	109				
PCB-157										
PCB-167	100		105		105	105				
PCB-169	↓		101		101	101				
PCB-170										
PCB-180										
PCB-189	100		97.5		97.5	97.5				

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

LDC #: 15449A3
SDG #: DPNS 19975

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?

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

- 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 = 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, ~~JOBT7~~

$$\begin{aligned} \text{Conc.} &= \frac{(6.862 + 0.6)(4000)}{(6.62 + 0.0)(1.04)(3.20)(1)} \\ &= 1245.5 \text{ ng/g} \end{aligned}$$

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