

U.S. Department of the Interior
NATIONAL BIOLOGICAL SERVICE
Midwest Science Center
Organic, Biological and Physiological Chemistry
4200 New Haven Road, Columbia, Missouri 65201

Final Laboratory Report FY-96-30-30

GC/HRMS ANALYSES of NON-*ortho*-CHLORO-SUBSTITUTED POLYCHLORINATED
BIPHENYLS in DUCKS and TREE SWALLOWS, TREE SWALLOW EGGS, and INSECTS
from the Hudson River Area

WORK UNIT #30096

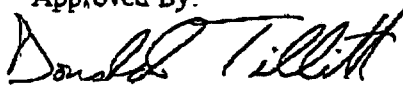
September 20, 1996

Submitted by:



Paul H. Peterman
Chemist, GC/MS

Approved By:



Donald E. Tillitt
Chief, Organic, Biological &
Physiological Chemistry Branch

To: Ms. Anne Secord
US Fish and Wildlife Service
3817 Luker Road
Cortland, NY 13045

Subject: Measured concentrations of non-*ortho*-chloro-substituted polychlorinated biphenyls (PCBs) in ducks and tree swallows, tree swallow eggs, and insects from the Hudson River Area. 11 1995.

Sample History: These samples are part of the study to track the movement of PCDDs, PCDFs, and dioxin-like PCBs. Birds, bird eggs, and other food-chain samples were collected from four primary sampling sites and another control site and were received from Anne Secord. Eggs had been weighed within a day of collection and were used as egg fresh weights. The samples were assigned individual data-base identification numbers and the descriptions were entered into a sample log. Labels on the sample containers were checked and reconciled with the packing list. Samples were stored frozen at -20°C in their original containers until sample processing began.

Analytical Preparation Methods Summary: The eggs were prepared according to MSC SOPC 5.143 (1). The samples were blended, and an aliquant (typically 10 g) was homogenized with three-to-four times its weight of anhydrous sodium sulfate.

Additional 18 quality control samples were also prepared and spiked: three samples in triplicate, three procedural blanks, one chicken egg (matrix) blank, two bluegill (matrix) blanks, one chicken egg spike, two bluegill spikes, and three positive control carp samples (6806, Saginaw Bay, Michigan). Approximately 10-g aliquants of the samples were homogenized with three times their weight of

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Table 1. Non-o-Chloro-Substituted PCBs (pg/g) in Birds, Bird Eggs, and Insects from the Hudson River

August 8, 1998 Work Unit 30098		GC/MS Sets: N30PCB and N31PCB Dates: 7/23/96 and 8/15/96		Non-o-Polychlorinated Biphenyls					
NFCR Number:	Submitter Number:	Sample Description:	GC/MS Run No.	Tetra: 3,4,4',5-TCB (81)	3,3',4,4'-TCB (77)	Penta: 3,3',4,4',5-PeCB (126)	Hexa: 3,3',4,4',5,5'-HxCB (169)		
12402-A	HUDWDA-4	2 Wood Duck Hen Composite, Replicate A, Near Special Area 13/Saratoga N.H. Park, 25 g	30-29	240	7,200	151	6.3		
12402-B	HUDWDA-4	2 Wood Duck Hen Composite, Replicate B, Near Special Area 13/Saratoga N.H. Park, 25 g	30-30	238	7,310	160	6.6		
12402-C	HUDWDA-4	2 Wood Duck Hen Composite, Replicate C, Near Special Area 13/Saratoga N.H. Park, 25 g	30-31	239	7,360	157	6.6		
12403	HUDMLE-4	Mallard Egg Composite, Saratoga N.H. Park, 10 g	30-32	1,100	15,400	476	8.4		
12404	HUDMLA-3	2 Mallard Hen Composite, Near Saratoga N.H. Park, 25 g	30-34 31-3	6,400	46,000	2,020	15		
12405	TS-E100	Tree Swallow Eggs, Chicks, and Adults: 3 Nest Egg (Tree Swallow) Composite, Lock 8, Reference Site, 10 g	30-35	2,900	50,200	2,340	34		
12406	TS-C100	3 Nest Tree Swallow Chick Composite, Lock 8, Reference Site, 10 g	30-36	700	12,400	376	8.4		
12407	TS-A100	Tree Swallow Adult Female, Lock 8 Reference Site, 10 g	30-37 31-4	16,000	220,000	5,900	120		
12408	TS-E204	1 Nest Tree Swallow Egg Composite, Remnant Site 4, Nest 204, 2.51 g	30-39 31-14	17,000	220,000	4,820	32		
12409	TS-C204-5	Tree Swallow 5-day old Chick Composite, Remnant Site 4, Nest 204, 7.32 g	30-40 31-6	6,900	120,000	1,740	8.4		
12410	TS-C204-10	Tree Swallow 10-day old Chick Composite, Remnant Site 4, Nest 204, 10 g	30-41 31-8	21,000	230,000	3,440	9.1 NO		
12411	TS-C204-15	Tree Swallow 15-day old Chick Composite, Remnant Site 4, Nest 204, 10 g	30-42 31-9	36,000	480,000	5,950	16 NO		

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Table 1. Non-o-Chloro-Substituted PCBs (pg/g) in Birds, Bird Eggs, and Insects from the Hudson River

August 8, 1986 Work Unit 30098			GC/MS Sets: N30PCB and N31PCB Dates: 7/23/96 and 8/15/96						Non-o-Polychlorinated Biphenyls			
NFCR Number:	Submitter Number:	Sample Description:	GC/MS Run No.	Tetra: (81)	3,4,4',5'-TCB	3,3',4,4'-TCB (77)	3,3',4,4',5'-PeCB (126)	Hexa: (169)				
12396	HUDBUG 2	Insects: Odonates, Remnant Site 4, 9.44 g	30-19	1,470	25,600	650	10					
12397	HUDBUG 3	Odonates only, Special Area 13, 10 g	30-20	230	10,800	376	6.8					
12398	HUDBUG 4	Odonates only, Saratoga Natl Historic Park, 10 g	30-21	53	3,600	96	4.8					
12436	HUDBUG-1M	Insect Mouth Sample, Lock 8, Ref. Site, 0.17 g	30-74 31-34	550	7,200	430 ND	100 ND					
12437	HUDBUG-2M	Insect Mouth Sample, Remnant Site 4, 2.08 g	30-75	6,180	92,800	1,250	8 ND					
12438	HUDBUG-3M	Insect Mouth Sample, Special Area 13, 2.9 g	30-76	1,760	35,200	430	6 ND					
12439	HUDBUG-4M	Insect Mouth Sample, Saratoga N.H. Park, 0.51 g	30-77 31-35	200	1,900	160	40 ND					
12399	HUDWDE-1	Duck and Duck eggs: Wood duck egg composite, A Remote Area, 10 g	30-22	10	160	21	3.7					
12400-A	HUDWDE-2	Wood duck egg composite, Replicate A, by Remnant Site 4 and Special Area 13, 10 g	30-24	1,400	20,300	430	6.8					
12400-B	HUDWDE-2	Wood duck egg composite, Replicate B, by Remnant Site 4 and Special Area 13, 10 g	30-25	1,390	20,700	428	7.5					
12400-C	HUDWDE-2	Wood duck egg composite, Replicate C, by Remnant Site 4 and Special Area 13, 10 g	30-26	1,380	20,350	426	6.4					
12401	HUDWDA-3	Adult Wood Duck Hen (eggs are HUDWDE-2), Near Remnant Site 4/Special Area 13, 25 g	30-27	430	3,450	207	2.2					

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Midwest Science Center
BRD, US Geological Survey
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Final Laboratory Report FY-97-30-01

**CONGENER-SPECIFIC ANALYSIS OF POLYCHLORINATED BIPHENYL RESIDUES IN TREE
SWALLOW CHICKS, EGGS AND OTHER BIOTA FROM THE HUDSON RIVER, W.U. 30096**

November 25, 1996

Submitted By:

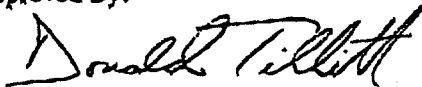


K.R. Echols
Chromatography



J.C. Meadows
Analytical Separations

Approved By:



D.E. Tillitt
Leader
Organic/Biochemistry Section

To: Anne Secord
US Fish and Wildlife Service
3817 Luker Rd.
Corland, NY 13045

Subject: Results of the congener-specific PCB analysis of tree swallow chicks, eggs and other biota from the Hudson River.

Sample History:

The purpose of this study is to determine whether migratory birds can accumulate detrimental levels of PCBs from various areas along the Hudson River. The indicator species chosen was the Tree Swallow (*Tachycineta bicolor*). Eggs, pre-fledgling chicks and adult tree swallows have been assessed for contaminant concentrations. Additionally, other biota were analyzed—insects which are the food of the tree swallows and two species of ducks. Higher burdens of PCBs in the chicks than those in the eggs represent local accumulation. Studies have indicated that pre-fledgling chicks can accumulate contaminants from sediment via consumption of adult midges, an emergent insect that comprises a large proportion of their diet. Since these midges spend their larval stages in sediment and emerge as breeding adults that do not feed, the contaminants that they contain must be derived from the sediment.

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	Sample Matrix	Grams Extracted	% Lipid	004,010	007,009	008	009,008	010	011	017,015	024,027	016,032	020	026
12396	HUDBUG-2	Insect	9.44	3.6	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	3.48	<0.85	<0.85	<0.85	<0.85
12397	HUDBUG-3	Insect	10.00	5.6	4.63	<0.85	<0.85	<0.85	6.40	2.63	2.74	<2.33	2.97	<0.85	<2.33
12398	HUDBUG-4	Insect	10.00	4.0	11.42	<0.85	<0.85	<2.33	<2.33	<2.33	3.49	<0.85	<2.33	<0.85	<2.33
12399	HUDWDE-1	Wood Duck Egg Composite	10.00	14.9	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12400 A	HUDWDE-2	Wood Duck Egg Composite	10.00	15.0	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	19.07	<0.85	<0.85	<0.85	<2.33
12400 B	HUDWDE-2	Wood Duck Egg Composite	10.00	15.2	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	18.33	<0.85	<0.85	<0.85	<2.33
12400 C	HUDWDE-2	Wood Duck Egg Composite	10.00	14.8	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	16.88	<0.85	<0.85	<0.85	<2.33
12401	HUDWQA-3	Adult Wood Duck Hen	25.00	1.4	4.68	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33
12402 A	HUDWQA-4	Two Wood Dk. Hen Composite	25.00	3.4	3.20	<0.85	<0.85	<2.33	<0.85	<2.33	7.19	<0.85	<2.33	<0.85	<2.33
12402 B	HUDWQA-4	Two Wood Dk. Hen Composite	25.00	3.4	2.75	<0.85	<0.85	<0.85	<0.85	<0.85	6.49	<0.85	<2.33	<0.85	<2.33
12402 C	HUDWQA-4	Two Wood Dk. Hen Composite	25.00	3.4	2.82	<0.85	<0.85	<2.33	<0.85	<0.85	7.38	<0.85	<2.33	<0.85	<2.33
12403	HUDMLE-4	Mallard Egg Composite	10.00	13.7	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	4.02	<0.85	<0.85	<0.85	<2.33
12404	HUDMLE-3	Two Mallard Hen Composite	25.00	4.1	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<2.33	<0.85	<2.33
12405	TS-E100	Tree Swallow Egg Composite	10.00	8.9	3.68	<0.85	<0.85	<0.85	<2.33	<2.33	4.87	<2.33	2.36	<0.85	2.78
12406	TS-C100	Tree Swallow Chick Composite	10.00	5.1	5.93	<0.85	<0.85	<0.85	3.64	3.17	5.68	2.36	6.02	<0.85	4.30
12407(a)	TS-A100	Tree Swallow Adult Female	10.00	8.3	<2.33	<0.85	<0.85	4.13	6.06	41.22	49.59	3.84	36.91	3.21	22.01
12408(a)	TS-E204	Tree Swallow Egg Composite	2.51	7.2	11.08	2.33	<2.33	14.28	16.98	154.42	151.01	13.24	111.43	10.47	110.39
12409	TS-C204-5	Tree Swallow Chick Composite	7.32	2.9	19.11	8.34	2.37	21.58	25.28	282.73	146.54	15.15	216.89	10.86	84.32
12410	TS-C204-10	Tree Swallow Chick Composite	10.00	7.1	115.15	30.46	33.19	240.43	89.33	1495.74	1090.05	51.72	1072.33	45.01	373.55
12411	TS-C204-15	Tree Swallow Chick Composite	10.00	9.5	75.33	13.78	7.89	128.49	115.79	895.07	1145.48	128.81	1381.52	71.39	712.68
12412	TS-E227	Tree Swallow Egg Composite	4.14	7.7	11.75	4.19	3.20	31.89	16.81	207.43	156.12	14.23	855.57	12.37	134.21
12413	TS-C227-5	Tree Swallow Chick Composite	10.00	4.1	17.52	<2.33	<2.33	6.84	31.79	298.81	163.48	13.83	124.80	6.80	73.91
12414	TS-C227-10	Tree Swallow Chick Composite	10.00	7.6	47.78	8.17	10.31	68.83	70.60	758.03	603.80	32.34	437.51	27.38	272.03
12415	TS-C227-15	Tree Swallow Chick Composite	10.00	10.5	119.82	29.13	27.25	251.51	129.33	1237.56	1464.82	79.22	1533.33	56.14	606.68
12416	TS-E313	Tree Swallow Egg Composite	3.48	5.7	271.36	4.77	5.35	50.02	27.63	91.59	75.38	10.14	85.80	6.04	75.54
12417	TS-C313-5	Tree Swallow Chick Composite	10.00	3.6	46.99	<2.33	<2.33	9.52	27.74	138.18	103.46	13.38	82.81	3.19	38.33
12418	TS-C313-15	Tree Swallow Chick Composite	10.00	7.6	115.98	4.49	4.77	35.30	57.94	194.29	169.16	28.24	230.54	8.39	114.75
12419	TS-E340	Tree Swallow Egg Composite	3.50	4.9	15.68	<2.33	<0.85	4.09	3.78	12.17	11.54	4.15	18.47	<2.33	21.42
12420	TS-C340-15	Tree Swallow Chick Composite	10.00	8.7	35.87	<2.33	3.39	13.20	25.28	125.13	159.29	19.50	167.33	6.24	107.94
12421	TS-E337	Tree Swallow Egg Composite	3.83	7.0	25.47	<2.33	<2.33	10.92	7.42	27.08	44.85	7.18	87.80	<2.33	31.64
12422	TS-C337-15	Tree Swallow Chick Composite	10.00	9.5	102.44	7.34	9.67	66.43	89.55	376.94	440.82	47.35	337.72	16.80	163.53
12423	TS-C339-5	Tree Swallow Chick Composite	10.00	5.2	44.55	3.12	3.12	13.35	7.39	154.45	101.96	12.14	137.81	3.46	63.27
12424(a)	TS-C339-10	Tree Swallow Chick Composite	10.00	6.7	138.08	5.49	7.66	38.75	64.86	232.61	188.52	22.22	218.95	10.89	108.28
12425	TS-C339-15	Tree Swallow Chick Composite	10.00	6.2	310.87	16.39	20.68	119.48	131.49	374.42	453.53	51.63	528.78	32.77	260.81
12426	TS-S315	Tree Swallow (salaged)	7.97	4.3	22.37	<0.85	<0.85	<2.33	31.99	104.89	88.74	19.59	81.98	4.57	61.85
12427	TS-A322	Tree Swallow Adult Female	10.00	7.1	24.69	<0.85	<0.85	<2.33	56.27	21.39	34.33	33.02	83.28	19.82	47.88
12428	TS-E411	Tree Swallow Egg Composite	4.11	7.5	82.48	<0.85	<2.33	6.21	47.30	12.23	20.62	9.19	18.94	<2.33	21.93
12429	TS-C411-15	Tree Swallow Chick Composite	10.00	7.3	115.37	<2.33	<2.33	15.09	62.45	32.48	40.79	24.95	65.78	2.36	51.17
12430	TS-E421	Tree Swallow Egg Composite	4.52	8.0	30.81	<0.85	<0.85	<2.33	7.80	3.90	9.40	5.35	8.07	<0.85	7.58
12431	TS-C421-15	Tree Swallow Chick Composite	10.00	6.2	29.85	<0.85	<0.85	2.35	21.81	7.01	11.08	14.87	24.80	<2.33	33.95
12432	TS-E428	Tree Swallow Egg Composite	3.15	8.3	55.77	<2.33	<2.33	3.05	12.94	6.57	7.64	7.47	13.37	<2.33	12.67
12433	TS-C428-15	Tree Swallow Chick Composite	10.00	7.1	34.58	<0.85	<2.33	4.28	23.87	11.17	18.50	13.19	33.59	<2.33	36.27
12434	TS-E500	Tree Swallow Egg Composite	10.00	6.8	37.83	<0.85	<0.85	2.68	10.26	4.79	10.70	8.85	15.23	<2.33	12.04
12435	TS-C500	Tree Swallow Chick Composite	10.00	8.8	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	<2.33	<0.85	<2.33
12436(a)	HUDBUG-A	Insect Mouth Sample & Insect	0.17	—	<0.85	20.21	9.02	6.17	<0.85	4.28	11.25	<0.85	<0.85	<0.85	<0.85
12437	HUDBUG-B	Insect Mouth Sample	2.88	3.8	30.35	8.88	5.35	66.73	22.88	293.00	253.04	18.50	313.84	20.71	124.86
12438	HUDBUG-C	Insect Mouth Sample	2.80	3.1	95.15	2.85	3.37	15.96	24.68	105.99	95.96	9.99	92.44	2.89	32.98
12439(a)	HUDBUG-D	Insect Mouth Sample	0.51	—	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12440	TS-E1TH	Tree Swallow Egg Composite	6.07	7.1	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12511 A	TS-C1TH	Tree Swallow Chick Composite	10.00	7.2	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12511 B	TS-C1TH	Tree Swallow Chick Composite	10.00	7.5	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12511 C	TS-C1TH	Tree Swallow Chick Composite	10.00	7.9	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85

(a) GC Replicate Average

(b) A contaminant has biased this congeners concentration.

(c) Not analyzed for lipid.

(d) ND for whole sample.

MDL = 0.85 ng/g

MDL = 2.33 ng/g

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Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	025	031	028	028,033	033	051	022	045	046	052	043	049	047	048	044
12396	HUDBUG-2	4.20	3.84	76.38	< 0.85	< 0.85	< 0.85	7.54	< 0.85	< 0.85	15.46	< 0.85	38.22	< 0.85	104.30	7.59
12397	HUDBUG-3	< 2.33	3.23	31.75	< 2.33	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	14.11	< 0.85	16.98	24.41	3.32	5.08
12398	HUDBUG-4	< 2.33	< 2.33	19.98	< 0.85	< 2.33	< 0.85	< 2.33	< 0.85	< 0.85	12.64	< 0.85	16.47	16.94	< 2.33	2.35
12399	HUDWDE-1	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12400 A	HUDWDE-2	< 0.85	11.95	219.78	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	24.04	< 0.85	27.77	159.37	5.84	< 0.85
12400 B	HUDWDE-2	< 0.85	12.23	197.39	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	23.12	< 0.85	26.81	155.29	4.02	< 0.85
12400 C	HUDWDE-2	< 0.85	10.79	200.32	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	21.91	< 0.85	25.46	148.45	2.85	< 0.85
12401	HUDWDA-3	< 0.85	< 2.33	10.41	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	2.88	< 0.85	< 2.33	2.38	8.16	< 0.85
12402 A	HUDWDA-4	< 0.85	< 2.33	38.83	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	4.92	< 0.85	5.14	24.47	< 2.33	< 0.85
12402 B	HUDWDA-4	< 0.85	2.44	38.05	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	5.08	< 0.85	5.23	25.08	< 2.33	< 0.85
12402 C	HUDWDA-4	< 0.85	< 2.33	40.35	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	5.27	< 0.85	5.32	25.16	2.75	< 0.85
12403	HUDWLE-4	< 0.85	8.88	145.12	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	15.66	< 0.85	13.18	69.89	< 2.33	< 2.33
12404	HUDWLA-3	< 2.33	12.39	289.91	2.89	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	19.32	< 0.85	8.41	180.72	22.04	2.71
12405	TS-E100	5.00	87.09	318.49	< 2.33	10.13	< 2.33	< 2.33	< 0.85	< 0.85	440.82	< 0.85	284.74	153.80	50.71	11.17
12406	TS-C100	3.48	23.83	89.76	< 2.33	11.80	2.83	5.35	< 0.85	< 2.33	125.43	< 0.85	117.24	64.37	11.72	43.29
12407(a)	TS-A100	30.68	428.71	1438.01	23.32	60.49	11.79	18.51	4.81	4.83	2428.69	4.58	2084.19	1111.19	170.57	118.66
12408(a)	TS-E204	91.66	565.50	1832.90	63.78	161.40	47.83	71.89	20.00	31.55	2949.35	17.57	2230.95	1289.65	208.84	538.12
12409	TS-C204-5	60.85	414.39	1040.49	129.23	96.17	42.42	141.84	61.75	29.33	1114.79	24.85	1048.72	811.44	303.65	586.71
12410	TS-C204-10	325.83	1467.89	5807.26	851.29	256.86	116.84	898.91	173.23	78.13	4125.12	31.81	4571.99	1809.58	2157.88	1869.68
12411	TS-C204-15	779.47	5871.02	10862.47	448.59	957.00	328.48	992.95	117.75	119.09	8044.78	32.63	9602.52	4884.44	1857.37	3720.55
12412	TS-E227	85.93	1187.58	2686.70	137.57	108.91	34.85	122.20	20.95	36.13	2816.52	22.34	2200.18	1384.91	197.34	530.40
12413	TS-C227-5	42.89	888.22	334.48	62.72	108.55	38.93	87.19	85.15	37.64	1130.04	14.98	962.40	824.70	240.50	527.89
12414	TS-C227-40	231.33	1435.94	4082.68	410.92	317.40	75.98	410.79	108.29	71.59	3184.14	85.95	3640.87	1889.70	805.24	1856.11
12415	TS-C227-15	604.24	3241.78	9052.58	13.93	1388.59	185.24	1088.54	146.08	119.28	7837.98	32.38	8323.76	3324.00	3381.15	3788.02
12416	TS-E313	51.06	412.09	879.96	54.91	51.55	10.94	60.84	5.99	4.14	925.81	< 0.85	650.16	754.23	156.88	122.37
12417	TS-C313-5	38.42	126.92	377.43	45.16	68.80	25.01	80.06	38.09	16.47	574.15	12.82	881.98	388.41	204.42	286.05
12418	TS-C313-16	104.82	825.62	2284.77	88.16	144.65	52.98	303.82	33.13	30.79	1820.07	10.93	2486.31	1571.83	311.15	1028.20
12419	TS-E340	17.61	710.10	1695.53	34.47	63.74	14.01	14.27	< 2.33	< 2.33	1918.67	4.34	1218.41	981.50	133.83	81.63
12420	TS-C340-15	84.02	451.15	1667.39	76.74	131.63	45.43	131.98	19.40	28.51	2058.71	1.33	2153.43	1287.05	200.88	618.85
12421	TS-E337	21.92	361.89	1015.11	22.44	58.98	9.69	21.81	2.49	3.00	1039.33	< 2.33	678.69	441.36	29.15	76.39
12422	TS-C337-18	190.30	1171.35	3811.81	161.24	267.54	82.06	416.28	82.78	73.54	2827.94	12.78	3583.58	2160.12	711.30	1805.89
12423	TS-C339-6	60.34	185.44	647.28	42.95	45.03	20.68	113.63	35.46	17.87	685.02	6.21	823.82	516.04	166.35	322.58
12424(a)	TS-C339-10	88.40	612.49	1811.06	157.53	92.44	41.24	248.84	43.73	22.66	1305.34	13.28	1608.54	980.87	320.02	640.62
12425	TS-C338-15	203.10	994.99	3469.67	642.68	132.49	70.27	56.48	71.95	39.20	2741.17	21.44	3244.85	1689.27	1148.38	1112.23
12426	TS-E315	66.34	513.40	1656.61	15.92	108.13	40.37	87.34	30.87	28.73	1812.42	7.44	1914.25	1154.73	226.02	717.07
12427	TS-A322	52.12	2280.15	9042.74	69.26	136.40	42.43	19.51	4.38	6.70	10587.12	5.78	11348.72	7828.97	124.62	197.10
12428	TS-E411	11.22	135.02	345.80	2.58	47.13	5.61	7.16	< 2.33	< 2.33	750.32	4.38	558.70	421.98	10.97	42.72
12429	TS-C411-15	15.90	81.27	95.59	3.80	48.46	15.56	9.38	4.32	3.01	282.34	< 2.33	248.84	287.52	25.45	61.80
12430	TS-E421	4.79	54.18	138.98	< 2.33	22.03	2.88	< 2.33	< 0.85	< 0.85	281.94	< 0.85	189.68	180.06	7.38	14.27
12431	TS-C421-15	11.86	67.28	68.42	< 0.85	37.55	11.32	3.89	< 2.33	< 2.33	282.14	2.34	233.80	122.99	30.03	53.98
12432	TS-E426	9.59	129.67	380.94	< 0.85	24.63	3.54	4.50	< 0.85	< 2.33	1018.53	< 0.85	634.34	402.84	474.67	21.98
12433	TS-C426-15	10.59	59.86	93.82	2.53	45.39	13.53	5.54	< 2.33	< 2.33	243.58	< 2.33	208.86	193.99	15.23	55.80
12434	TS-E508	7.21	38.67	105.27	< 2.33	20.97	4.82	2.96	< 0.85	< 0.85	208.79	< 0.85	152.10	96.80	5.72	19.79
12435	TS-C500	< 0.85	3.87	8.52	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	25.27	< 0.85	22.47	15.28	3.94	2.45
12436(c)	HUDBUG-A	24.81	< 2.33	19.40	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	18.08	< 0.85	11.85	27.68	< 0.85	6.18
12437	HUDBUG-B	122.98	446.21	1182.50	375.62	141.43	43.51	316.47	65.17	18.87	1298.62	12.89	1411.52	758.06	484.88	633.66
12438	HUDBUG-C	24.81	215.11	835.56	36.59	43.31	17.35	99.44	30.46	9.50	433.70	5.73	558.28	302.17	131.95	285.98
12439(c)(d)	HUDBUG-D	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12440	TS-E11H	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	2.97	< 0.85	< 2.33	< 2.33	< 0.85	< 0.85
12511 A	TS-C11H	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12511 B	TS-C11H	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12511 C	TS-C11H	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85

(a) GC Replicate Average

(b) A contaminant has
biased this congener
concentration.

(c) Not analysed for lipid.

(d) ND for whole sample.

NDL = 0.85 ng/g

MQL = 2.33 ng/g

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	042	041	054	040	047	053	074	075,076	066	095	091	056,060	092	064	001
12396	HUDBUG-2	77.98	< 2.33	6.09	2.71	< 2.33	22.43	213.83	9.06	50.97	10.86	2.74	35.84	3.81	< 2.33	24.81
12397	HUDBUG-3	7.59	2.70	3.87	< 2.33	< 2.33	9.56	96.80	9.44	38.33	5.63	< 2.33	17.17	2.83	< 2.33	12.46
12398	HUDBUG-4	7.94	< 0.85	2.34	< 0.85	< 0.85	4.85	37.85	< 2.33	< 0.85	4.19	< 2.33	5.42	2.57	< 0.85	8.07
12399	HUDWDE-1	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12400 A	HUDWDE-2	4.44	< 0.85	< 2.33	< 0.85	< 2.33	36.18	209.98	6.68	7.83	7.59	< 2.33	36.51	30.51	< 0.85	23.51
12400 B	HUDWDE-2	4.53	< 0.85	< 2.33	< 0.85	< 2.33	35.21	188.93	8.92	7.30	8.73	< 2.33	35.30	28.83	< 0.85	22.63
12400 C	HUDWDE-2	4.02	< 0.85	< 2.33	< 0.85	< 2.33	33.71	193.11	8.24	7.16	7.10	< 2.33	33.63	28.45	< 0.85	21.51
12401	HUDWQA-3	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	7.42	39.86	< 2.33	3.08	8.60	< 0.85	< 2.33	< 2.33	< 0.85	< 2.33
12402 A	HUDWQA-4	11.34	< 0.85	< 0.85	< 0.85	< 0.85	7.87	36.50	< 2.33	24.65	4.12	< 0.85	4.44	5.61	< 0.85	4.51
12402 B	HUDWQA-4	12.17	< 0.85	< 0.85	< 0.85	< 0.85	8.04	37.48	< 2.33	26.68	3.96	< 0.85	4.40	5.82	< 0.85	4.51
12402 C	HUDWQA-4	13.68	< 0.85	< 0.85	< 0.85	< 0.85	8.43	39.28	< 2.33	27.80	4.28	< 0.85	4.80	6.19	< 0.85	4.74
12403	HUDWQA-4	< 2.33	< 0.85	< 2.33	< 0.85	< 0.85	30.13	148.18	10.06	5.57	2.45	< 2.33	27.16	9.78	2.55	13.05
12404	HUDWQA-3	15.55	< 2.33	2.36	< 0.85	< 0.85	115.33	656.38	17.88	34.32	6.92	3.89	88.08	28.11	< 2.33	42.30
12405	TS-E100	69.64	7.48	58.07	3.89	6.13	45.92	198.82	198.36	346.67	65.89	35.03	95.12	53.66	24.34	212.52
12406	TS-C100	65.11	11.84	28.59	5.99	4.92	13.00	79.87	63.34	5.95	13.45	12.22	38.05	14.89	5.77	73.85
12407(a)	TS-A100	112.37	80.70	12.49	22.47	58.16	203.89	1344.68	1681.80	1088.52	1176.43	297.34	412.03	254.56	16.80	1128.12
12408(a)	TS-E204	378.08	295.00	414.63	80.89	75.89	259.60	1538.80	1289.15	983.71	543.22	255.54	370.82	291.52	84.70	778.98
12409	TS-C204-6	682.22	274.64	361.06	107.40	58.12	122.55	799.79	649.94	1417.38	235.46	116.41	401.84	98.13	65.77	508.59
12410	TS-C204-10	2416.12	1381.77	748.89	373.41	160.03	281.37	2430.34	3412.50	4178.54	1731.91	393.29	1334.77	393.53	268.19	1412.40
12411	TS-C204-15	6273.49	2619.93	2755.57	377.87	540.94	591.48	5459.39	46.97	468.02	507.83	627.77	2563.63	436.08	274.04	2793.04
12412	TS-E227	1019.41	233.79	458.24	111.33	82.89	240.77	1439.25	1560.06	128.11	238.82	218.04	897.74	204.56	69.96	829.21
12413	TS-C227-5	603.94	173.24	271.93	95.85	41.99	68.14	595.88	539.72	409.60	512.58	90.72	178.98	88.57	65.01	339.55
12414	TS-C227-10	2816.22	787.03	1032.51	230.47	190.41	256.45	2235.88	2407.47	152.00	277.18	237.11	1009.22	189.83	149.87	1231.20
12415	TS-C227-15	3947.64	2183.52	2497.96	520.28	427.51	529.79	4505.71	6325.41	5778.59	4084.79	472.15	2189.85	712.05	370.55	2579.06
12416	TS-E315	377.48	64.67	215.01	28.85	34.20	233.83	825.88	689.60	825.80	304.96	138.84	220.37	127.86	11.10	357.83
12417	TS-C315-5	423.50	92.81	188.39	77.93	33.24	76.68	402.77	279.06	241.98	144.77	75.18	116.75	81.01	50.80	271.16
12418	TS-C315-15	1648.65	936.37	831.88	133.58	60.70	211.79	1660.19	1351.80	101.94	171.06	154.34	690.09	186.74	88.48	1040.54
12419	TS-E340	334.52	101.80	407.55	18.02	28.28	185.64	956.47	1142.78	61.18	305.17	141.35	332.44	178.55	6.19	530.44
12420	TS-C340-15	851.15	202.93	584.88	90.48	91.69	196.52	1455.51	1239.25	1531.11	1185.78	181.86	483.82	272.38	87.20	1024.62
12421	TS-E397	272.22	45.31	210.64	10.18	23.80	78.81	582.32	632.04	44.28	56.88	68.32	201.12	87.89	18.36	378.59
12422	TS-C337-85	2505.08	530.81	1155.75	285.36	161.71	290.00	2344.82	1792.82	142.48	321.28	233.98	985.35	234.93	119.92	1355.21
12423	TS-C338-5	939.57	132.01	275.51	72.28	38.79	73.74	523.04	452.89	319.82	413.77	73.57	178.43	77.80	48.12	358.93
12424(a)	TS-C338-10	1107.80	274.18	506.82	114.89	71.40	111.94	1109.94	1145.41	76.97	145.20	114.59	548.05	112.12	69.56	668.47
12425	TS-C338-15	1859.25	678.54	885.93	225.72	150.95	278.20	2075.55	2372.45	2297.47	1979.37	334.54	994.20	408.07	174.72	1192.21
12426	TS-S315	1118.09	209.32	599.17	114.57	77.45	194.16	1334.87	8072.45	92.18	161.99	184.13	595.03	175.80	98.93	892.42
12427	TS-A322	1451.84	184.18	2402.94	68.39	138.38	1267.98	9979.78	5447.82	687.27	872.89	1218.98	2986.40	653.58	53.78	6444.07
12428	TS-E411	88.01	26.38	84.88	3.58	15.48	65.40	260.19	238.78	21.74	91.01	73.75	88.67	107.70	12.19	345.85
12429	TS-C411-15	77.17	38.47	48.83	12.01	11.45	27.63	100.02	108.62	165.87	44.28	36.39	38.83	39.55	18.43	120.71
12430	TS-E421	34.68	10.65	27.12	< 2.33	5.29	23.70	99.72	79.53	7.99	20.87	22.63	28.85	30.75	10.44	100.57
12431	TS-C421-15	55.37	35.04	47.87	7.13	11.09	24.28	95.73	93.40	72.47	72.28	40.59	35.28	46.73	16.29	129.56
12432	TS-E426	82.98	11.88	104.15	8.82	15.88	98.93	336.61	308.04	265.28	67.87	98.37	142.19	187.34	4.13	350.88
12433	TS-C426-15	67.00	33.10	41.23	6.47	10.13	22.56	96.06	98.10	7.32	35.14	80.88	39.05	36.26	13.92	115.98
12434	TS-E500	35.51	15.36	28.69	4.04	5.42	16.08	63.21	68.58	92.55	29.81	23.48	23.03	27.39	2.89	85.90
12435	TS-C500	3.57	< 2.33	3.42	< 0.85	< 0.85	2.37	9.51	9.33	10.34	10.13	< 2.33	3.54	2.94	< 2.33	11.98
12436(c)	HUDBUG-A	< 0.85	< 2.33	3.24	< 0.85	< 0.85	< 0.85	8.49	12.44	< 0.85	22.02	< 0.85	2.78	4.89	7.08	86.20
12437	HUDBUG-B	941.09	350.98	421.05	120.32	102.48	123.78	813.09	1281.64	710.08	378.72	140.82	417.83	105.25	67.98	595.58
12438	HUDBUG-C	406.04	94.17	195.46	57.25	27.32	45.59	377.87	352.99	25.16	44.36	44.25	185.02	42.48	23.29	237.12
12439(c)(d)	HUDBUG-D	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12440	TS-C1TH	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	< 2.33	< 0.85	< 2.33	< 2.33	< 0.85	< 2.33	< 2.33	7.27
12511 A	TS-C1TH	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33
12511 B	TS-C1TH	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33
12511 C	TS-C1TH	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33

(a) GC Replicate Average
 (b) A contaminant has
 biased this congeners
 concentration.
 (c) Not analyzed for lipid.
 (d) ND for whole sample.

MDL = 0.85 ng/g
 MQL = 2.33 ng/g

08/30/99 MON 15:05 FAX 518 458 6372
08/25/98 11:31 6607 753 9899

BUREAU OF TOXIC SUBSTANCE
FWS NYFO

009
009/012

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	089	119	083	097	087	136	110	082	191	135,144,124	147	107	123,149	119	134
12396	HUDBUG-2	88.41	<2.33	<2.33	21.45	11.86	<0.85	14.09	4.23	<2.33	<2.33	<0.85	3.68	11.56	56.81	<0.85
12397	HUDBUG-3	29.12	<0.85	<0.85	4.48	5.71	<0.85	3.65	<2.33	<2.33	<2.33	<0.85	3.26	3.84	38.08	<0.85
12398	HUDBUG-4	13.57	<0.85	<0.85	2.52	<2.33	<0.85	3.90	<0.85	<0.85	<0.85	<2.33	<2.33	3.16	12.04	<0.85
12399	HUDWDE-1	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85
12400 A	HUDWDE-2	114.47	4.08	<0.85	<0.85	5.88	<0.85	11.17	<0.85	6.70	4.88	4.82	9.51	15.86	112.26	<0.85
12400 B	HUDWDE-2	111.22	3.80	<0.85	<0.85	6.15	<0.85	10.70	<0.85	6.46	4.84	3.96	8.82	15.43	108.75	<0.85
12400 C	HUDWDE-2	105.78	3.49	<0.85	<0.85	5.88	<0.85	10.26	<0.85	6.21	3.63	4.24	8.59	14.84	104.58	<0.85
12401	HUDWDA-3	18.75	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	30.65	<0.85
12402 A	HUDWDA-4	23.06	<0.85	<0.85	<0.85	<2.33	<0.85	2.73	<0.85	<2.33	<0.85	<2.33	<2.33	2.53	21.18	<0.85
12402 B	HUDWDA-4	23.20	<0.85	<0.85	<0.85	<2.33	<0.85	2.40	<0.85	<0.85	<0.85	<2.33	<2.33	2.45	21.10	<0.85
12402 C	HUDWDA-4	24.28	<0.85	<0.85	<0.85	<2.33	<0.85	3.27	<0.85	<2.33	<0.85	<2.33	<2.33	2.71	22.62	<0.85
12403	HUDMLE-4	120.30	<2.33	<0.85	<2.33	7.42	<0.85	8.82	<0.85	<2.33	<2.33	2.68	9.58	10.14	89.16	<0.85
12404	HUDMLA-3	389.95	5.68	<0.85	<2.33	26.82	<2.33	24.11	<0.85	7.51	4.02	24.01	33.87	27.32	448.45	<0.85
12405	TS-E100	142.11	7.51	<0.85	16.36	61.13	3.43	117.28	<2.33	10.75	15.53	6.48	27.05	74.44	253.58	<0.85
12406	TS-C100	54.48	2.56	2.62	26.29	30.16	<0.85	43.36	7.02	3.59	4.24	<2.33	7.97	15.84	70.01	<0.85
12407(a)	TS-A100	745.66	48.01	12.80	85.97	413.22	21.60	577.04	16.03	54.89	62.56	23.94	84.22	283.15	828.89	<2.33
12408(a)	TS-E204	632.00	40.47	30.23	205.77	279.00	16.02	438.88	52.45	54.03	55.63	26.08	110.83	316.97	557.00	<2.33
12409	TS-C204-5	340.79	20.28	33.51	210.89	262.47	9.77	378.81	84.12	21.40	35.08	4.15	41.84	121.87	388.01	4.02
12400	TS-C204-10	828.34	38.25	74.63	532.51	643.40	21.91	882.17	230.89	43.04	87.50	12.33	87.99	201.34	752.98	16.16
12411	TS-C204-15	2147.27	82.14	149.29	1172.68	1356.14	40.28	1939.25	343.75	87.73	175.16	24.09	203.82	451.42	1688.03	18.13
12412	TS-E227	716.57	40.45	24.46	169.17	345.77	15.40	556.00	56.98	49.37	76.77	20.48	88.40	212.71	755.15	4.48
12413	TS-C227-4	244.31	13.39	26.63	128.15	148.06	6.99	188.53	60.49	16.07	28.31	5.13	38.07	87.51	174.87	4.38
12414	TS-C227-10	887.00	34.01	66.06	517.69	586.02	20.24	822.36	174.21	42.24	75.08	11.50	89.17	183.80	630.14	14.07
12415	TS-C227-18	1754.21	63.15	114.67	1089.48	1216.32	35.79	1485.04	367.84	77.70	127.35	15.32	68.10	372.64	1542.66	16.01
12416	TS-E313	335.37	25.51	9.91	66.00	238.07	8.56	179.42	22.47	25.34	48.83	22.83	117.45	187.05	463.03	6.38
12417	TS-C313-5	219.75	11.84	24.17	88.25	116.40	4.30	157.24	52.28	14.61	23.13	5.35	39.15	61.13	150.82	4.11
12418	TS-C313-15	785.09	30.39	53.48	424.59	452.34	13.48	684.88	138.31	45.18	64.30	13.31	75.21	168.68	751.15	6.81
12419	TS-E340	442.29	28.38	5.10	128.02	208.26	16.57	407.82	11.23	33.90	50.41	12.39	80.88	179.28	452.78	<2.33
12420	TS-C340-15	713.27	33.35	53.72	300.88	405.39	14.43	581.83	78.17	55.79	78.89	13.17	68.02	148.41	578.24	7.19
12421	TS-E337	302.28	66.64	3.36	84.70	141.87	2.96	264.82	12.36	17.79	25.18	8.20	38.65	85.23	299.45	<0.85
12422	TS-C337-15	1065.11	51.09	80.62	572.93	593.33	23.18	881.22	189.93	75.05	94.06	12.95	<2.33	196.78	937.58	13.72
12423	TS-C339-5	255.52	11.48	21.93	138.83	153.75	4.41	208.00	47.71	17.24	24.85	6.56	33.98	82.78	252.26	5.88
12424(a)	TS-C339-10	527.08	22.83	41.87	273.13	303.14	6.83	450.00	94.34	30.87	52.71	9.24	66.67	153.07	518.55	4.12
12425	TS-C339-15	880.68	43.33	73.28	458.52	546.94	22.59	743.11	182.43	62.75	66.03	18.32	85.19	208.29	792.39	16.80
12426	TS-E315	732.56	31.07	58.90	353.67	418.48	13.22	623.83	101.83	49.19	68.40	13.67	97.89	228.34	800.04	11.26
12427	TS-A322	5912.37	238.00	43.41	344.12	2286.91	54.45	3743.11	42.88	221.79	341.24	93.40	788.29	1315.11	6773.42	3.34
12428	TS-E411	270.58	17.56	4.47	94.38	84.44	7.17	166.63	5.73	31.24	29.36	12.09	30.39	128.33	238.88	<0.85
12429	TS-C411-15	95.83	5.86	6.11	30.28	39.22	4.24	68.39	7.86	12.32	11.10	4.83	11.07	38.00	77.89	3.81
12430	TS-E421	94.37	6.21	<2.33	8.92	22.99	<2.33	46.85	<2.33	8.07	8.07	5.28	11.04	35.39	85.67	<0.85
12431	TS-C421-15	99.79	6.22	7.02	35.27	42.71	4.39	68.14	7.00	18.18	13.19	5.48	10.89	47.92	40.56	<2.33
12432	TS-E426	325.88	20.27	<2.33	21.32	82.60	6.10	174.40	2.89	24.73	28.54	15.95	38.87	132.38	170.27	<0.85
12433	TS-C428-15	95.88	6.19	5.86	30.58	36.76	4.11	67.02	7.38	12.28	11.01	5.28	12.54	40.87	85.11	<2.33
12434	TS-E500	62.09	3.66	<2.33	14.63	23.16	2.80	44.19	3.24	6.20	7.85	2.98	7.58	28.84	83.37	<0.85
12435	TS-C500	7.34	<0.85	<0.85	<2.33	3.30	<0.85	4.95	<0.85	<2.33	<2.33	<0.85	<2.33	3.84	6.10	<0.85
12438(c)	HUDBUG-A	21.80	12.02	<0.85	8.61	20.05	<0.85	32.79	2.71	3.85	2.80	<0.85	<2.33	20.08	24.33	<0.85
12437	HUDBUG-B	408.21	16.09	38.47	246.23	288.93	8.96	321.30	108.74	20.33	38.62	5.13	44.30	104.84	199.35	6.42
12436	HUDBUG-C	205.13	8.73	17.34	106.02	108.68	2.43	159.03	38.69	8.11	12.54	3.82	20.59	37.93	187.83	2.67
12439(c)(d)	HUDBUG-D	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12440	TS-ETH	3.78	<0.85	<0.85	<0.85	<2.33	<0.85	2.93	<0.85	<2.33	<0.85	<0.85	<0.85	5.08	6.88	<0.85
12411 A	TS-CITH	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85
12411 B	TS-CITH	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85
12411 C	TS-CITH	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85

(a) GC Replicate Average

(b) A contaminant has
biased this congeners
concentration.

(c) Not analyzed for lipid.

(d) MD for whole sample.

MDL = 0.85 ng/g

MQL = 2.33 ng/g

WU 30096, USDO, USGS, BRD, WSC, 11/25/96/F76 = FSECFC3M.WK4

318038

08/30/99 MON 15:06 FAX 518 458 6372
08/25/99 11:32 607 753 9899

BUREAU OF TOXIC SUBSTANCE
FWS NYFO

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	114	121,122	146	153	132	105	141	179	137	176	130	138	158	129	178
12396	HUDBUG-2	7.98	< 0.85	12.06	33.08	< 2.33	45.72	< 2.33	< 0.85	2.77	< 0.85	3.69	26.24	9.61	< 2.33	< 0.85
12397	HUDBUG-3	3.57	< 0.85	6.50	19.81	< 2.33	18.96	< 2.33	< 0.85	< 0.85	< 0.85	< 2.33	7.53	< 2.33	< 0.85	< 0.85
12398	HUDBUG-4	< 0.85	< 2.33	5.78	9.68	< 0.85	6.55	< 0.85	< 0.85	< 0.85	< 0.85	< 2.33	11.97	< 0.85	< 0.85	< 2.33
12399	HUDWDE-1	< 0.85	< 0.85	< 0.85	2.53	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	2.39	< 0.85	< 0.85	< 0.85
12400 A	HUDWDE-2	7.91	2.91	18.94	42.83	< 2.33	53.00	< 2.33	< 0.85	4.26	< 0.85	4.21	60.46	4.63	< 0.85	5.03
12400 B	HUDWDE-2	7.89	2.58	18.18	41.64	< 2.33	52.16	< 2.33	< 0.85	4.00	< 0.85	3.73	69.22	4.85	< 0.85	4.56
12400 C	HUDWDE-2	7.37	2.41	17.39	40.04	< 2.33	49.81	< 2.33	< 0.85	3.83	< 0.85	3.52	56.20	4.07	< 0.85	4.46
12401	HUDWDA-3	2.75	< 0.85	6.68	22.18	< 0.85	19.09	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	20.90	< 2.33	< 0.85	< 0.85
12402 A	HUDWDA-4	< 2.33	< 0.85	5.14	11.34	< 0.85	8.54	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	13.26	< 2.33	< 0.85	< 2.33
12402 B	HUDWDA-4	< 2.33	< 0.85	5.10	11.37	< 0.85	8.15	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	13.18	< 2.33	< 0.85	< 2.33
12402 C	HUDWDA-4	< 2.33	< 0.85	5.40	12.05	< 0.85	8.77	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	14.08	< 2.33	< 0.85	< 2.33
12403	HUDWDA-4	8.02	< 2.33	19.05	64.22	< 0.85	71.89	< 0.85	< 0.85	3.05	< 0.85	< 2.33	62.10	5.83	< 0.85	8.44
12404	HUDWDA-3	25.35	4.79	64.69	189.18	5.20	184.16	2.64	< 0.85	18.33	< 0.85	13.38	288.81	21.20	< 0.85	17.27
12405	TS-EK00	19.23	3.89	45.51	112.75	18.75	159.13	10.54	2.64	21.88	< 0.85	18.77	192.82	22.01	< 2.33	80.07
12406	TS-C100	5.49	< 2.33	9.04	24.08	5.94	27.40	4.05	< 0.85	3.73	< 0.85	3.03	36.31	3.63	< 2.33	< 2.33
12407(a)	TS-A100	48.05	11.42	110.70	358.33	139.89	342.91	62.76	18.47	58.03	5.84	57.05	577.44	64.07	13.68	24.57
12408(a)	TS-E204	56.03	8.14	86.38	183.52	129.68	172.71	51.38	16.60	48.52	2.81	45.78	317.55	55.92	15.78	19.38
12408	TS-C204-5	22.68	9.14	28.24	102.20	72.78	240.63	22.96	16.57	18.60	2.38	18.51	188.13	20.87	14.50	4.48
12410	TS-C204-10	88.35	24.51	51.29	157.83	171.95	124.89	65.59	4.73	35.88	< 2.33	36.56	311.18	40.70	31.61	7.21
12411	TS-C204-15	117.95	39.20	80.30	320.83	240.54	1143.66	117.53	62.20	73.53	4.98	65.39	617.43	75.16	16.59	2.94
12412	TS-E227	51.10	10.20	78.82	177.18	119.89	487.10	62.22	6.06	38.87	3.78	38.87	233.19	41.46	13.16	16.04
12413	TS-C227-5	10.46	5.65	22.55	53.40	44.25	60.00	20.88	< 0.85	14.51	< 0.85	12.77	93.16	16.42	9.51	2.33
12414	TS-C227-10	55.20	18.94	47.27	151.81	122.08	468.18	50.44	20.57	31.95	3.14	30.51	287.77	36.95	8.27	< 2.33
12415	TS-C227-15	68.24	< 0.85	85.22	301.71	198.80	765.74	64.92	7.38	58.99	5.68	58.08	564.08	68.48	51.89	11.50
12416	TS-E913	94.88	13.49	138.43	243.08	48.33	241.93	31.29	6.40	51.78	3.38	87.20	435.28	69.79	6.55	34.76
12417	TS-C313-5	19.01	3.48	30.85	65.71	34.72	57.75	15.21	4.96	15.53	< 0.85	16.29	114.83	20.27	9.43	6.89
12418	TS-C313-15	40.31	19.87	56.80	169.74	87.81	414.29	48.36	4.24	38.49	< 2.33	33.14	316.84	57.09	23.48	10.63
12419	TS-E348	33.56	6.05	62.88	156.75	76.46	307.52	29.80	8.12	25.13	3.69	28.35	218.46	30.31	12.04	10.16
12420	TS-C340-15	33.89	11.79	66.80	170.63	99.38	54.39	36.40	8.24	37.69	< 2.33	37.19	294.14	41.48	28.27	11.89
12421	TS-E337	25.01	3.00	41.30	104.99	22.89	196.21	16.93	< 2.33	16.03	< 0.85	18.13	179.69	18.59	6.57	8.76
12422	TS-C337-10	55.37	24.64	80.40	199.98	135.82	485.45	52.02	7.60	50.49	< 0.85	47.74	366.78	39.61	45.24	9.50
12423	TS-C339-5	15.29	4.91	25.79	63.32	31.81	89.57	15.65	4.20	12.80	< 0.85	13.42	117.47	14.75	8.51	4.13
12424(a)	TS-C339-10	37.90	15.64	40.71	103.53	67.67	290.74	58.10	3.13	22.24	< 2.33	24.91	185.28	28.15	17.64	7.34
12425	TS-C339-15	33.06	22.22	72.91	209.22	160.70	237.15	53.76	11.04	42.46	2.38	43.58	382.05	39.61	34.97	9.15
12426	TS-S315	48.97	11.71	71.10	188.05	78.36	441.22	52.29	5.22	38.94	2.57	38.68	242.95	46.50	28.51	13.00
12427	TS-A322	487.12	24.40	515.58	1844.65	563.40	3803.32	304.77	90.53	288.84	12.54	312.74	2863.80	313.15	59.78	120.83
12428	TS-E411	17.20	7.47	49.90	128.23	43.95	118.05	19.08	3.58	16.76	< 2.33	19.56	188.42	16.00	3.76	19.57
12429	TS-C411-15	5.27	< 2.33	14.59	38.88	15.64	36.55	5.48	2.73	4.66	< 0.85	5.38	59.21	4.07	< 2.33	4.07
12430	TS-E421	5.73	3.59	21.77	47.80	11.28	41.89	8.53	< 0.85	8.25	< 0.85	8.50	70.95	5.99	< 2.33	4.13
12431	TS-C421-15	5.57	< 0.85	16.06	37.78	19.51	36.81	6.30	3.01	5.10	< 0.85	5.63	55.77	4.55	2.51	4.38
12432	TS-E426	22.08	< 2.33	65.42	137.08	37.59	162.35	17.07	< 2.33	21.82	< 2.33	22.80	209.90	21.70	< 2.33	18.57
12433	TS-C426-15	6.35	4.45	23.24	41.25	16.65	41.71	6.22	< 2.33	5.11	< 0.85	6.83	64.18	4.22	2.81	4.84
12434	TS-E500	3.86	< 2.33	13.34	34.03	14.72	25.07	4.74	< 2.33	4.58	< 0.85	4.98	53.00	8.06	< 2.33	3.67
12435	TS-C500	< 0.85	< 0.85	< 2.33	6.28	< 2.33	2.89	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	8.08	< 0.85	< 0.85	< 0.85
12436(c)	HUDBUG-A	< 0.85	< 0.85	4.24	26.37	17.21	10.80	7.33	< 0.85	< 0.85	< 0.85	< 0.85	45.71	4.78	< 0.85	< 0.85
12437	HUDBUG-B	18.16	11.74	21.38	72.59	74.54	222.52	23.98	7.62	18.40	< 2.33	13.73	136.20	14.84	14.06	2.77
12438	HUDBUG-C	12.56	3.21	11.25	38.81	19.63	96.89	8.81	< 0.85	8.04	< 0.85	5.73	70.29	7.67	5.78	< 2.33
12439(c)(d)	HUDBUG-D	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85
12440	TS-ENH	< 0.85	< 0.85	4.55	12.04	< 2.33	2.81	< 2.33	< 0.85	< 0.85	< 0.85	< 0.85	14.05	< 2.33	< 0.85	< 2.33
12511 A	TS-CITH	< 0.85	< 0.85	< 0.85	3.14	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	3.25	< 0.85	< 0.85	< 0.85
12511 B	TS-CITH	< 0.85	< 0.85	< 0.85	3.00	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	3.13	< 0.85	< 0.85	< 0.85
12511 C	TS-CITH	< 0.85	< 0.85	< 0.85	3.02	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	< 0.85	3.24	< 0.85	< 0.85	< 0.85

(a) GC Replicate Average
(b) A contaminant has
biased this congeners
concentration.
(c) Not analysed for lipid.
(d) ND for whole sample.

MDL = 0.85ng/g
MQL = 2.33 ng/g

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200
12396	HUDBUG-2	2.74	4.53	2.72	<0.85	<0.85	<2.33	<0.85	12.23	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12397	HUDBUG-3	7.05	4.80	<2.33	<0.85	<0.85	<2.33	<0.85	8.71	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12398	HUDBUG-4	8.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	2.91	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12399	HUDWDE-1	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	8.16	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12400 A	HUDWDE-2	16.74	16.50	3.34	<0.85	<0.85	3.89	<0.85	8.14	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12400 B	HUDWDE-3	4.66	17.29	3.09	<0.85	<0.85	3.70	<0.85	7.71	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12400 C	HUDWDE-4	15.05	16.60	2.98	<0.85	<0.85	3.56	<0.85	4.43	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12401	HUDWDA-1	15.12	4.22	<0.85	<0.85	<0.85	<2.33	<0.85	12.66	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12401	HUDWDA-2	7.77	6.69	<0.85	<0.85	<0.85	<2.33	<0.85	4.37	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12402 A	HUDWDA-3	5.85	4.02	<0.85	<0.85	<0.85	<2.33	<0.85	2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12402 B	HUDWDA-4	5.85	3.99	<0.85	<0.85	<0.85	<2.33	<0.85	2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12402 C	HUDWDA-5	5.85	4.21	<0.85	<0.85	<0.85	<2.33	<0.85	2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12403	HUDWDA-6	5.85	25.19	4.33	<0.85	<0.85	<2.33	<0.85	12.66	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12404	HUDWDA-7	18.02	5.90	<0.85	<0.85	<0.85	<2.33	<0.85	4.37	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12404	HUDWDA-8	152.78	31.57	17.20	<0.85	<0.85	<2.33	<0.85	33.24	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12405	TS-E200	55.39	18.73	15.18	<0.85	<0.85	<2.33	<0.85	6.80	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12406	TS-E200	6.92	3.02	2.84	<0.85	<0.85	<2.33	<0.85	6.80	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12407 (a)	TS-E200	121.62	58.01	38.59	5.12	38.67	46.25	23.67	60.79	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12408 (a)	TS-E200	89.05	34.40	25.99	2.94	28.49	27.79	16.32	62.86	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12409	TS-E200	28.82	11.91	8.28	<2.33	23.02	10.10	10.10	51.09	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12410	TS-E200	40.51	22.44	16.93	2.87	30.66	16.26	17.92	94.71	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12411	TS-E200	84.55	188.07	31.50	7.73	50.66	33.99	17.92	56.75	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12412	TS-E200	81.80	34.72	20.02	3.18	24.22	26.87	14.42	58.15	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12413	TS-E200	18.28	6.45	7.83	<2.33	7.92	6.65	3.41	48.12	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12414	TS-E200	38.49	20.29	16.53	3.18	19.29	15.17	7.69	48.12	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12415	TS-E200	65.36	37.83	27.97	5.13	37.85	26.73	11.40	11.42	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12416	TS-E200	177.38	67.41	44.73	<2.33	13.02	62.24	33.97	11.42	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12417	TS-E200	29.38	11.63	9.83	<0.85	5.71	10.82	5.31	23.80	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12418	TS-E200	50.04	23.42	16.53	<2.33	14.39	17.08	9.00	45.74	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12419	TS-E200	50.06	20.18	13.49	<2.33	14.33	17.08	9.00	45.74	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12420	TS-E200	56.47	24.83	19.67	2.84	19.92	20.40	9.76	52.73	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12421	TS-E200	66.39	37.36	24.78	4.08	25.07	11.42	5.38	51.70	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12422	TS-E200	24.12	16.02	11.80	<0.85	6.31	7.34	2.70	14.25	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12423	TS-E200	24.55	15.02	11.80	<0.85	6.31	7.34	2.70	14.25	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12424 (a)	TS-E200	67.76	30.40	21.79	3.08	28.61	28.17	13.34	30.94	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12425	TS-E200	64.18	31.71	22.99	2.84	17.36	23.32	11.21	47.72	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12426	TS-E200	523.20	203.90	123.95	18.55	127.77	205.99	89.58	437.52	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33	<2.33
12427	TS-E200	52.64	20.00	10.20	<2.33	12.81	18.20	7.74	22.52	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12428	TS-E200	15.75	3.80	2.86	<0.85	8.72	4.17	<2.33	5.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12429	TS-E200	23.16	7.35	4.11	<0.85	4.73	5.55	3.02	8.46	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12430	TS-E200	20.95	3.47	2.92	<0.85	4.40	4.41	<2.33	5.98	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12431	TS-E200	55.75	16.84	12.44	<2.33	12.42	16.97	6.14	25.39	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12432	TS-E200	19.59	4.11	3.44	<0.85	4.00	4.97	<2.33	7.53	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12433	TS-E200	20.78	5.51	3.27	<0.85	4.87	4.33	2.80	5.91	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12434	TS-E200	8.45	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12435	TS-E200	13.58	4.63	<0.85	<0.85	10.57	6.88	<2.33	2.46	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12436 (a)	HUDBUG-A	17.28	8.54	6.54	<2.33	10.57	6.88	8.55	10.46	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12437	HUDBUG-B	9.65	3.72	3.03	<0.85	2.61	<0.85	<2.33	9.26	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12438	HUDBUG-C	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12439	HUDBUG-D	9.60	2.89	<2.33	<0.85	<0.85	<2.33	<0.85	2.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12440	TS-E200	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	2.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12441 A	TS-E200	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	2.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12441 B	TS-E200	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	2.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85
12441 C	TS-E200	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	2.84	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85

(a) GC Replicate Average
(b) A contaminant has
biased this congeners
concentration.
(c) Not analyzed for lipid.
(d) ND for whole sample.

MDL = 0.35 ng/g
MQL = 2.33 ng/g

WU 30096,

Table 1. Congener Specific PCBs (ng/g) - Hudson River Samples (WU30096)

Sample Name	Field ID	191	200	170,190	198 (b)	189	196,203	189	205,195	207	194	205	206	209	Total cPCBs
12388	HUDBUG-2	<0.85	<0.85	4.84	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	1085
12397	HUDBUG-3	<0.85	<0.85	4.38	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	509
12399	HUDBUG-4	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	240
12399	HUDWDE-1	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	5
12400 A	HUDWDE-2	<0.85	<0.85	7.07	<0.85	3.63	3.20	<0.85	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	1410
12400 B	HUDWDE-2	<0.85	<0.85	6.80	<0.85	3.76	3.97	<0.85	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	1341
12400 C	HUDWDE-2	<0.85	<0.85	6.51	<0.85	3.71	2.94	<0.85	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	1300
12401	HUDWDA-3	<0.85	<0.85	4.64	<0.85	2.81	2.49	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	250
12402 A	HUDWDA-4	<0.85	<0.85	<2.33	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	283
12402 B	HUDWDA-4	<0.85	<0.85	<2.33	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	286
12402 C	HUDWDA-4	<0.85	<0.85	<2.33	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<2.33	<0.85	306
12403	HUDWDA-4	<0.85	<0.85	8.37	<0.85	3.92	4.24	<0.85	<2.33	<0.85	2.38	<0.85	<0.85	<0.85	1118
12404	HUDWDA-3	3.43	<0.85	39.57	<2.33	29.87	19.03	<2.33	8.92	<2.33	16.62	<2.33	11.60	2.62	3966
12405	TS-E100	2.85	<0.85	31.75	<2.33	14.95	13.86	<2.33	5.21	<0.85	9.75	<2.33	10.13	3.39	4561
12406	TS-C100	<0.85	<0.85	4.32	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	1411
12407(a)	TS-A100	4.35	<2.33	70.83	2.59	39.61	37.58	3.81	12.15	2.45	28.27	2.62	16.85	4.81	22225
12408(a)	TS-E204	8.49	<2.33	59.27	<2.33	23.99	21.04	<2.33	7.80	<2.33	16.51	<2.33	12.81	3.32	22820
12409	TS-C204-5	<2.33	<0.85	20.32	<2.33	0.38	4.62	<0.85	2.54	<0.85	6.16	<0.85	4.56	<0.85	15832
12410	TS-C204-10	<2.33	<2.33	32.41	<0.85	10.07	10.28	<2.33	2.50	<0.85	7.17	<0.85	<0.85	<0.85	55628
12411	TS-C204-15	6.07	<2.33	51.60	<0.85	19.22	20.42	2.45	5.95	<2.33	15.33	<2.33	10.32	<2.33	94409
12412	TS-E227	4.51	<2.33	49.25	2.87	22.18	21.14	<2.33	6.76	<2.33	14.58	<2.33	7.90	2.52	24890
12413	TS-C227-4	<0.85	<0.85	13.43	<0.85	4.85	4.76	<0.85	<2.33	<0.85	3.44	<0.85	<2.33	<0.85	11488
12414	TS-C227-10	2.76	<0.85	29.10	<0.85	6.88	9.71	<2.33	2.75	<0.85	7.31	<0.85	4.07	<0.85	39660
12415	TS-C227-15	3.72	<2.33	53.28	<0.85	14.87	15.89	<2.33	4.27	<0.85	11.13	<0.85	<0.85	<0.85	98280
12416	TS-E313	9.59	<0.85	100.30	<2.33	80.88	49.32	3.95	17.73	3.68	43.20	2.86	37.10	9.08	141339
12417	TS-C313-5	<2.33	<0.85	19.30	<0.85	7.84	6.76	<0.85	<2.33	<0.85	5.22	<0.85	3.12	<0.85	7822
12418	TS-C313-15	<2.33	<0.85	35.00	<0.85	11.87	11.40	<2.33	3.18	<0.85	8.70	<0.85	<0.85	<0.85	26116
12419	TS-E340	2.56	<0.85	31.07	2.50	15.18	12.82	<2.33	4.29	<0.85	9.34	<0.85	7.82	<2.33	15174
12420	TS-C340-15	2.58	<0.85	40.58	<2.33	13.91	13.81	<2.33	3.68	<0.85	10.56	<0.85	4.41	<0.85	23519
12421	TS-E337	<2.33	<0.85	19.83	<2.33	11.37	8.30	<0.85	3.19	<0.85	8.99	<0.85	<0.85	<2.33	8917
12422	TS-C337-15	3.72	<0.85	32.73	<2.33	14.25	14.68	<2.33	4.08	<0.85	11.67	<0.85	6.10	<0.85	37956
12423	TS-C339-5	<0.85	<0.85	13.84	<0.85	6.02	5.21	<0.85	<2.33	<0.85	3.78	<0.85	<2.33	<0.85	5659
12424(a)	TS-C339-10	<2.33	<0.85	23.60	<0.85	8.79	7.86	<2.33	<2.33	<0.85	5.82	<0.85	<0.85	<0.85	18180
12425	TS-C339-15	3.59	<2.33	34.58	<2.33	17.80	15.91	<2.33	4.96	<0.85	12.20	<0.85	8.51	<2.33	39646
12426	TS-E315	2.90	<0.85	44.14	<0.85	16.53	16.45	<2.33	4.92	<0.85	12.78	<2.33	4.87	<2.33	21289
12427	TS-A322	26.57	8.19	322.07	7.99	188.38	122.53	14.27	39.86	8.11	95.87	9.10	<2.33	10.76	113822
12428	TS-E411	<2.33	<0.85	24.17	<0.85	17.21	14.04	<2.33	4.63	<0.85	8.47	<0.85	<0.85	2.35	8150
12429	TS-C411-15	<0.85	<0.85	5.70	<0.85	3.42	2.59	<0.85	<2.33	<0.85	<2.33	<0.85	<2.33	<0.85	3068
12430	TS-E421	<0.85	<0.85	10.38	<0.85	8.10	5.46	<0.85	<2.33	<0.85	3.45	<0.85	<0.85	<2.33	2135
12431	TS-C421-15	<0.85	<0.85	5.78	<0.85	8.82	2.71	<0.85	<2.33	<0.85	<2.33	<0.85	<2.33	<0.85	2512
12432	TS-E426	<2.33	<0.85	26.49	<2.33	14.47	12.92	<2.33	4.52	<0.85	7.85	<2.33	7.50	2.69	7379
12433	TS-C426-10	<0.85	<0.85	6.39	<0.85	4.05	3.11	<0.85	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	2456
12434	TS-E600	<0.85	<0.85	7.58	<0.85	5.78	5.02	<0.85	<2.33	<0.85	3.13	<2.33	4.04	3.15	1405
12435	TS-C500	<0.85	<2.33	<2.33	<0.85	<2.33	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	197
12436(c)	HUDBUG-A	<0.85	<0.85	2.71	10.28	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	7.68	5.62	573
12437	HUDBUG-B	<0.85	<0.85	12.30	2.93	4.82	4.35	<0.85	<2.33	<0.85	3.91	<0.85	7.45	<0.85	17743
12438	HUDBUG-C	<0.85	<0.85	6.87	<0.85	<2.33	<2.33	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<0.85	6928
12439(c)(d)	HUDBUG-D	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	0
12440	TS-E17H	<0.85	<0.85	3.88	<0.85	3.48	2.83	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	<2.33	103
12511 A	TS-C17H	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	6
12511 B	TS-C17H	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	6
12511 C	TS-C17H	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<0.85	<2.33	<0.85	<0.85	6

(a) GC Replicate Average

(b) A contaminant has biased this congeners concentration.

(c) Not analysed for lipid.

(d) ND for whole sample.

MQL = 0.85 ng/g

MQL = 2.33 ng/g