## **QUALITY ASSURANCE/ QUALITY CONTROL**

This appendix details quality assurance/quality control information for the water quality analyses, sediment geochemistry analyses, tissue chemistry analyses, invertebrate taxonomy, and otter trawl sample collection conducted for the Orange County Sanitation District's (OCSD) 2010-11 ocean monitoring program.

## INTRODUCTION

The Core monitoring program was designed to measure compliance with permit conditions and for temporal and spatial trend analysis. The program includes measurements of:

- Water quality;
- Sediment quality;
- Benthic infaunal community health;
- Fish and macroinvertebrate community health;
- Fish tissue contaminant concentrations (chemical body burden); and
- Fish health (including external parasites and diseases).

The Core monitoring program complies with the Orange County Sanitation District

## WATER QUALITY NARRATIVE

#### Introduction

OCSD's Environmental Laboratory and Ocean Monitoring (ELOM) staff collected 551 discrete ammonia samples during each of the four quarters beginning July 1, 2010 and ending June 30, 2011. All samples were iced upon collection, preserved with 1:1 sulfuric acid upon receipt by the ELOM laboratory staff, and stored at  $4 \pm 2$  °C until

(OCSD) Quality Assurance/Quality Control (QA/QC) Program requirements and applicable federal, state, local, and contract requirements. The objectives of the quality assurance program are as follows:

- Scientific data generated will be of sufficient quality to stand up to scientific and legal scrutiny.
- Data will be gathered or developed in accordance with procedures appropriate for the intended use of the data.
- Data will be of known and acceptable precision, accuracy, representativeness, completeness, and comparability as required by the program.

The various aspects of the program are conducted on a schedule that varies weekly. semi-annually. monthly. quarterly, annually. Table C-1 shows that sampling goals were achieved for >99.6 percent of the required samples. Sampling and data analysis is characterized by quarters 1 through 4, which are representative of summer (July-September), fall (October-December), winter (January-March), and spring (April-June) seasons, respectively. according to ESL analysis Standard Operating Procedures (SOPs), which are found in the Laboratory Operating Procedures Manual (LOPM).

#### Analytical Method - Ammonia

The samples were analyzed for ammonia on a segmented flow analyzer using Standard Method 4500-NH<sub>3</sub> G (Standard Methods 1998). In the analysis, sodium phenolate and sodium hypochlorite react with ammonia to form indophenol blue in a concentration proportional to the ammonia concentration in

Table C-1. Ocean monitoring program sample collection requirements and percent completion, July 2010-June 2011.

Quarter	Program Type	Parameter	Nominal # of Samples	# of Samples Collected	# of QA Duplicates * (≥10%)	%Samples Collected
		CTD Drops	105	105	15	100
	Water Quality	Ammonium	470	551	81	100
		Bacteria	260	259	30	99.6
		Grain size	49	49	5	100
		TOC	49	49	2	100
4	0 - 1 1	Dissolved Sulfides	49	49	5	100
1	Sediment	Metals	49	49	5	100
	Chemistry	PCB/Pesticides	NA	NA	NA	NA
		PAH	49	49	6	100
		LAB	49	49	6	100
	Benthic Infauna	Infauna	49	49	0	100
	Fish Community	Trawls *	23	23	NA	100
	,	CTD Drops	105	105	22	100
	Water Quality	Ammonium	470	551	81	100
		Bacteria	260	260	30	100
		Grain size	10	10	1	100
	Sediment Chemistry	TOC	10	10	0	100
2		Dissolved Sulfides	10	10	1	100
		Metals	10	10	1	100
		PCB/Pesticides	10	10	1	100
		PAH	10	10	1	100
	Benthic Infauna	Infauna	30	30	0	100
	Water Quality	CTD Drops	105	105	21	100
		Ammonium	470	551	81	100
		Bacteria	260	260	30	100
		Grain size	10	10	1	100
		TOC	10	10	0	100
	Sediment	Dissolved Sulfides	10	10	1	100
	Chemistry	Metals	10	10	1	100
3		PCB/Pesticides	10	10	1	100
		PAH	10	10	1	100
	Benthic Infauna	Infauna	30	30	3	100
	Fish Community	Trawls	23	23	NA	100
	,	Hornyhead turbot	20 X 2 *	20 X 2 *	4	100
	Fish Tissue	English sole	20 X 2 *	20 X 2 *	6	100
		Sanddab Guild	12	12	2	100
		CTD Drops	105	105	15	100
	Water Quality	Ammonium	470	551	81	100
	adding	Bacteria	260	260	30	100
		Grain size	10	10	1	100
		TOC	10	10	1	100
4	Sediment	Dissolved Sulfides	10	10	1	100
	Chemistry	Metals	10	10	1	100
	2	PCB/Pesticides	10	10	1	100
		PAH	10	10	1	100
	1	1 / 11 1	1 10	10	1 "	100

NA = not applicable

<sup>\*</sup> Number of QA duplicates indicates number of field duplicates or lab sample splits only. It does not include other QA samples.

\*\* English sole and hornyhead turbot samples were analyzed for both muscle and liver tissue.

the sample. The blue color is intensified with sodium nitroprusside and is measured at 660 nm.

## QA/QC - Ammonia

A typical sample batch includes three blanks, an external reference standard, a spike, and a spike replicate in seawater collected from a control site. One spike and spike replicate is added to the batch every 10 samples. The method detection limit (MDL) for low-level ammonia samples using the segmented flow instrument is 0.02 mg/L. QA/QC summary data are presented in Table C-2. samples were analyzed within the required holding time with 100% of the 78 analyses meeting the QA/QC criteria for blanks. All analyses met the QA/QC criteria for the external reference sample. The 54 first quarter sample matrix spike recoveries, replicate recoveries. and precision measurements were all within target limits.

The 63 second quarter sample matrix spike and replicate recoveries were within target limits, while two of the 63 precision measurements were outside target limits. Three of 89 matrix spike replicate samples, two of 89 matrix spike replicate recoveries, and one of 89 precision measurements for matrix spike and matrix spike replicates were out of control for the third quarter samples. All 69 fourth quarter samples met target limits for matrix spike recoveries, replicates, and precision measurements. In all cases, it was determined that recovery and precision criteria were exceeded due to matrix effects or instrumentation malfunction. Additionally, the set of results following those in question were within the control limits and therefore all results are considered valid.

Table C-2. Water quality ammonium QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	6	6	<2X MDL	N/A
			Matrix Spike	14	10, 4*	80-120	
Summer	NH3WQ100716-1	Ammonium	Matrix Spike Dup	14	11, 3*	80-120	
			Matrix Spike Precision	14	13, 1**		< 11%
			ERA Check Standard	2	2	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	15	14	80-120	
Summer	NH3WQ100804-1	Ammonium	Matrix Spike Dup	15	14, 1*	80-120	
			Matrix Spike Precision	15	12, 3**		< 11%
			ERA Check Standard	1	1	87 - 114	
		Ammonium	Blank	0	0	<2X MDL	N/A
			Matrix Spike	5	5	80-120	
Summer	NH3WQ100804-2		Matrix Spike Dup	5	5	80-120	
			Matrix Spike Precision	5	5		< 11%
			ERA Check Standard	0	0	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	13	12, 1*	80-120	
Summer	NH3WQ100812-1	Ammonium	Matrix Spike Dup	13	13	80-120	
			Matrix Spike Precision	13	13		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	7	7	80-120	
Summer	NH3WQ100813-1	Ammonium	Matrix Spike Dup	7	7	80-120	
			Matrix Spike Precision	7	7		< 11%
			ERA Check Standard	1	1	87 - 114	

Table C-2 Continues.

Table C-2 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	9	9	80-120	
Fall	NH3WQ100924-1	Ammonium	Matrix Spike Dup	9	9	80-120	
			Matrix Spike Precision	9	7, 2**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	6	6	80-120	
Fall	NH3WQ100928-1	Ammonium	Matrix Spike Dup	6	6	80-120	
			Matrix Spike Precision	6	6		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	11	11	80-120	
Fall	NH3WQ101103-1	Ammonium	Matrix Spike Dup	11	11	80-120	
			Matrix Spike Precision	11	8, 3**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	11	10, 1*	80-120	
Fall	NH3WQ101104-1	Ammonium	Matrix Spike Dup	11	10, 1*	80-120	
			Matrix Spike Precision	11	10, 1**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Fall	NH3WQ101105-1	Ammonium	Matrix Spike Dup	10	10	80-120	
			Matrix Spike Precision	10	5, 5**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	6	6	80-120	
Fall	NH3WQ101108-1	Ammonium	Matrix Spike Dup	6	6	80-120	
			Matrix Spike Precision	6	2, 4**		< 11%
			ERA Check Standard	1	1	87 - 114	

Table C-2 Continues.

Table C-2 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Fall	NH3WQ101117-1	Ammonium	Matrix Spike Dup	10	10	80-120	
			Matrix Spike Precision	10	4, 4**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Winter	NH3WQ101216-1	Ammonium	Matrix Spike Dup	10	10	80-120	
			Matrix Spike Precision	10	7, 3**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	5	5	80-120	
Winter	NH3WQ101217-1	Ammonium	Matrix Spike Dup	5	5	80-120	
			Matrix Spike Precision	5	5		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	2	<2X MDL	N/A
			Matrix Spike	7	6	80-120	
Winter	NH3WQ110131-1	Ammonium	Matrix Spike Dup	7	5, 1*	80-120	
			Matrix Spike Precision	7	6, 1**		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	8	7	80-120	
Winter	NH3WQ110201-1	Ammonium	Matrix Spike Dup	8	8	80-120	
			Matrix Spike Precision	8	8		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	8	8	80-120	
Winter	NH3WQ110202-1	Ammonium	Matrix Spike Dup	8	8	80-120	
			Matrix Spike Precision	8	3, 5**		< 11%
			ERA Check Standard	1	1	87 - 114	

Table C-2 Continues.

Table C-2 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	8	8	80-120	
Winter	NH3WQ110208-1	Ammonium	Matrix Spike Dup	8	8	80-120	
			Matrix Spike Precision	8	8		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	7	7	80-120	
Winter	NH3WQ110209-1	Ammonium	Matrix Spike Dup	7	7	80-120	
			Matrix Spike Precision	7	7		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	11	11	80-120	
	NH3WQ110217-1	Ammonium	Matrix Spike Dup	11	11	80-120	
Winter			Matrix Spike Precision	11	11		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	12	11	80-120	
Winter	NH3WQ110223-1	Ammonium	Matrix Spike Dup	12	11	80-120	
			Matrix Spike Precision	12	11		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	13	13	80-120	
Winter	NH3WQ110224-1	Ammonium	Matrix Spike Dup	13	13	80-120	
			Matrix Spike Precision	13	13		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	9	9	80-120	
Spring	NH3WQ110324-1	Ammonium	Matrix Spike Dup	9	9	80-120	
			Matrix Spike Precision	9	9		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	6	6	80-120	
Spring	NH3WQ110328-1	Ammonium	Matrix Spike Dup	6	6	80-120	
-			Matrix Spike Precision	6	6		< 11%
			ERA Check Standard	1	1	87 - 114	

Table C-2 Continues.

Table C-2 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	5	5	80-120	
Spring	NH3WQ110502-1	Ammonium	Matrix Spike Dup	5	5	80-120	
			Matrix Spike Precision	5	5		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	8	8	80-120	
Spring	NH3WQ110503-1	Ammonium	Matrix Spike Dup	8	8	80-120	
			Matrix Spike Precision	8	8		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Spring	NH3WQ110511-1	Ammonium	Matrix Spike Dup	10	10	80-120	
. 0			Matrix Spike Precision	10	10		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
Spring			Matrix Spike	11	11	80-120	
	NH3WQ110512-1	Ammonium	Matrix Spike Dup	11	11	80-120	
. 0			Matrix Spike Precision	11	11		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Spring	NH3WQ110517-1	Ammonium	Matrix Spike Dup	10	10	80-120	
. 0			Matrix Spike Precision	10	10		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	10	10	80-120	
Spring	NH3WQ110518-1	Ammonium	Matrix Spike Dup	10	10	80-120	
			Matrix Spike Precision	10	10		< 11%
			ERA Check Standard	1	1	87 - 114	
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	9	9	80-120	
Summer	NH3WQ110615-1	Ammonium	Matrix Spike Dup	9	9	80-120	
			Matrix Spike Precision	9	9		< 11%
			ERA Check Standard	1	1	87 - 114	<u> </u>
			Blank	3	3	<2X MDL	N/A
			Matrix Spike	6	6	80-120	
Summer	NH3WQ110620-1	Ammonium	Matrix Spike Dup	6	6	80-120	
			Matrix Spike Precision	6	6		< 11%
			ERA Check Standard	1	1	87 - 114	, ,

<sup>\*</sup> Recovery (70% or 130%) was out of control due to rounding.

\*\* Matrix spike precision was out of control due to rounding. The associated method blank and check standard were in control and therefore the data were reported.

## SEDIMENT CHEMISTRY NARRATIVE

## **FIRST QUARTER (JULY 2010)**

### Introduction

OCSD's laboratory received 49 sediment samples from ocean monitoring staff during the month of July 2010. All samples were stored according to the LOPM. All samples were analyzed for polycyclic aromatic hydrocarbons (PAHs), linear alkyl benzenes (LABs), trace metals, mercury, dissolved sulfides (DS), total organic carbon (TOC), First quarter sediment and grain size. samples were not analyzed for organochlorine pesticides or polychlorinated biphenyl congeners (PCBs), with approval from the California Regional Water Quality Control Board and USEPA due to an OCSD sediment-mapping project. In addition, 20 samples typically collected during the first quarter (replicate samples 2 and 3) were not collected, and were redirected toward the sediment mapping study.

## <u>Analytical Methods - PAHs and LABs</u>

The analytical methods used to detect PAHs and LABs in the samples are described in the LOPM. All sediment samples were extracted using an accelerated solvent extractor (ASE) during the months of August through September 2010. Approximately 10 grams (dry weight) of sample were used for each analysis. A separatory funnel extraction was performed using 100 milliliters of sample when field and rinse blanks were included in the batch.

A typical sample batch included 18 field samples with required quality control (QC) samples. Each sample batch analyzed for PAHs and LABs included the following QC samples: one sand blank, one reporting level spike (PAH or LAB), two standard reference materials (SRM), one matrix spike set (PAH or LAB), and two sample extraction duplicates. There were four

batches extracted and analyzed for PAHs. In addition, one batch contained one rinse sample and one field blank. Method detection limits (MDLs) are presented in Table C-3. Acceptance criteria for PAH SRMs are presented in Table C-4.

Sediment PAH and LAB QA/QC summary data are presented in Table C-5. All analyses were performed within holding times and with appropriate quality control measures, as stated in the program's Quality Assurance Project Plan (QAPP). Any variances are noted in the Comments/Notes section of each batch summary.

## <u>Analytical Methods - Organochlorine</u> Pesticides and PCB Congeners

Organochlorine pesticides and PCB congener samples were not analyzed for the summer of 2010. MDLs for PCBs/pesticides for fall through spring samples are presented in Tables C-6 and C-7. Acceptance Criteria for PCB/pesticide SRMs are presented in Table C-8. Sediment PCB/pesticide QA/QC summary data are presented in Table C-9.

## Analytical Methods - Trace Metals

Dried sediment samples were analyzed for trace metals in accordance with methods in the LOPM. A typical sample batch for silver, cadmium, chromium, copper, nickel, lead, zinc, selenium, arsenic, and beryllium analyses included three blanks, a blank spike, and one SRM. Additionally. duplicate samples, spiked samples and duplicate spiked samples were analyzed a minimum of once every 10 sediment samples. QC for a typical sample batch for aluminum and iron analyses included three blanks, an SRM, sediment sample with duplicates, spiked samples and duplicate spiked samples analyzed a minimum of once every 10 sediment samples. analysis of the blank spike and SRM provided a measure of the accuracy of the analysis. The analysis of the sample, its

Table C-3. Method detection levels for PAH and LAB compounds in sediments, July 2010–June 2011.

Parameter	Accelerated Solvent Extraction SIM Detection Limit, (ng/g dry weight)	Parameter	Accelerated Solvent Extraction SIM Detection Limit, (ng/g dry weight)
	PAH Cor	npounds	
1,6,7-Trimethylnaphthalene	0.20	Benzo[k]fluoranthene	0.20
1-Methylnaphthalene	0.30	Biphenyl	0.30
1-Methylphenanthrene	0.20	Chrysene	0.20
2,6-Dimethylnaphthalene	0.30	Dibenz[a,h]anthracene	0.10
2-Methylnaphthalene	0.50	Dibenzothiophene	0.20
Acenaphthene	0.40	Fluoranthene	0.30
Acenaphthylene	0.60	Fluorene	0.20
Anthracene	0.70	Indeno[1,2,3-c,d]pyrene	0.20
Benz[a]anthracene	0.20	Naphthalene	0.50
Benzo[a]pyrene	0.10	Perylene	0.20
Benzo[b]fluoranthene	0.30	Phenanthrene	0.40
Benzo[e]pyrene	0.50	Pyrene	0.30
Benzo[g,h,l]perylene	0.30		
	PAH Alkylated	l Homologues	
C1-Chrysenes	2	C1-Fluoranthenes/Pyrenes	2
C2-Chrysenes	2	C1-Naphthalenes	2
C3-Chrysenes	2	C2-Naphthalenes	2
C4-Chrysenes	2	C3-Naphthalenes	2
C1-Dibenzothiophenes	2	C4-Naphthalenes	2
C2-Dibenzothiophenes	2	C1-Phenanthrenes/Anthracenes	2
C3-Dibenzothiophenes	2	C2-Phenanthrenes/Anthracenes	2
C1-Fluorenes	2	C3-Phenanthrenes/Anthracenes	2
C2-Fluorenes	2	C4-Phenanthrenes/Anthracenes	2
C3-Fluorenes	2		
	LAB Cor	npounds	
2-Phenyldecane	0.10	6-Phenyltetradecane	0.40
3-Phenyldecane	0.10	7-Phenyltetradecane	0.10
4-Phenyldecane	0.10	2-Phenylundecane	0.10
5-Phenyldecane	0.10	3-Phenylundecane	0.10
2-Phenyltridecane	0.30	4-Phenylundecane	0.10
3-Phenyltridecane	0.10	5-Phenylundecane	0.10
4-Phenyltridecane	0.20	6-Phenylundecane	0.10
5-Phenyltridecane	0.30	2-Phenyldodecane	0.20
6-Phenyltridecane+7-Phenyltridecane	0.40	3-Phenyldodecane	0.10
2-Phenyltetradecane	0.10	4-Phenyldodecane	0.20
3-Phenyltetradecane	0.10	5-Phenyldodecane	0.20
4-Phenyltetradecane	0.10	6-Phenyldodecane	0.20
5-Phenyltetradecane	0.20		

Table C-4. Acceptance criteria for standard reference materials of PAHs in sediments, July 2010–June 2011.

Compound Name	True Value	Certified Acce	ptance Criteria y/g
, , , , , , , , , , , , , , , , , , ,	hā/ā	Min.	Max.
SRM 1944A	- Organics in Marine Sediment Na	ational Institute of Standards and T	Technology.
Anthracene	1.77	0.44	2.21
Benz[a]anthracene	4.72	1.18	5.90
Benzo[a]pyrene	4.30	1.08	5.38
Benzo[b]fluoranthene	3.87	0.97	4.84
Benzo[e]pyrene	3.28	0.82	4.10
Benzo[g,h,i]perylene	2.84	0.71	3.55
Benzo[k]fluoranthene	2.30	0.58	2.88
Chrysene	4.86	1.22	6.08
Dibenz[a,h]anthracene	0.42	0.11	0.53
Fluoranthene	8.92	2.23	11.15
Indeno(1,2,3-c,d)pyrene	2.78	0.70	3.48
Naphthalene	1.65	0.41	2.06
Perylene	1.17	0.29	1.46
Phenanthrene	5.27	1.32	6.59
Pyrene	9.70	2.43	12.13
SRM 1941E	B - Organics in Marine Sediment N	ational Institute of Standards and	Technology
Anthracene	184	110	258
Benz[a]anthracene	335	201	469
Benzo[a]pyrene	358	215	501
Benzo[b]fluoranthene	453	272	634
Benzo[e]pyrene	325	195	455
Benzo[g,h,i]perylene	307	184	430
Benzo[k]fluoranthene	225	135	315
Chrysene	291	175	407
Dibenz[a,h]anthracene	53	32	74
Fluoranthene	651	391	911
Indeno(1,2,3-c,d)pyrene	341	205	477
Naphthalene	848	509	1,187
Perylene	397	238	556
Phenanthrene	406	244	568
Pyrene	581	349	813

Table C-5. Sediment PAH/LAB QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD	Comments
1		PAH SRM 1944	15	12	25% of the certified or		80% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NIA	93% Pass
		PAH Reporting Level Spike	25	25	60 -120	NA	100% Pass
		LAB Reporting Level Spike	25	20	60 -120		80% Pass
		PAH Matrix Spike					
	Cadaaa Iulio DM	Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
	Sedcore_Jul10_DM	LAB Matrix Spike					
		Based on Mean of MS and MSD	25	23	40 - 120	NA	92% Pass
		PAH Duplicate Analysis - #1	11	9			82% Pass
		PAH Duplicate Analysis - #2	14	14	T NA	< 20% @ 3 x MDL of Sample Mean	100% Pass
		LAB Duplicate Analysis - #1	14	14	- NA		100% Pass
		PAH Duplicate Analysis - #2	11	6			55% Pass
		PAH SRM 1944	15	12	25% of the certified or		80% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NIA	93% Pass
		PAH Reporting Level Spike	25	25	60 -120	NA	100% Pass
		LAB Reporting Level Spike	25	25	60 -120		100% Pass
		PAH Matrix Spike					
1	Cadaaa Iulio DN	Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
1	Sedcore_Jul10_ DN	LAB Matrix Spike					
		Based on Mean of MS and MSD	25	22	40 – 120	NA	88% Pass
		PAH Duplicate Analysis - #1	24	20			83% Pass
		PAH Duplicate Analysis - #2	15	15	] NA	< 20% @ 3 x MDL	100% Pass
		LAB Duplicate Analysis - #1	23	22	- NA	of Sample Mean	96% Pass
		PAH Duplicate Analysis - #2	15	1			6% Pass

Notes: <sup>1</sup> SRM certified values are based on the addition of selected compounds prior to extraction for use as internal standards for quantification purposes. (NIST, Certificate of Analysis, SRM 1941b, SRM 1944a, Organics in Marine Sediment).

OCSD laboratory results are not corrected for surrogate recoveries, causing some analytes with lower molecular weights and boiling points to fail the established criteria for SRM certified values. N/A=not applicable

**Table C-5 Continues.** 

Table C-5 Continued.

Quarter	Sample Set	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD	Comments
		PAH SRM 1944	15	13	25% of the certified or		87% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NA	93% Pass
		PAH Reporting Level Spike	25	22	60 -120	NA	88% Pass
		LAB Reporting Level Spike	25	25	60 -120		100% Pass
		PAH Matrix Spike					
1	Cadasas Iulio DO	Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
1	Sedcore_Jul10_DO	LAB Matrix Spike					
		Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
		PAH Duplicate Analysis - #1	15	10			83% Pass
		PAH Duplicate Analysis - #2	14	14	1	< 20% @ 3 x MDL	78% Pass
		LAB Duplicate Analysis - #1	20	14	- NA	of Sample Mean	82% Pass
		LAB Duplicate Analysis - #2	15	16			100% Pass
		PAH SRM 1944	15	13	25% of the certified or		87% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NA	93% Pass
		PAH Reporting Level Spike	25	24	60 -120	NA	96% Pass
		LAB Reporting Level Spike	NA	NA	60 -120		NA
		PAH Matrix Spike					
2	Cadaras Ort40 DC	Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
2	Sedcore_Oct10_DS	LAB Matrix Spike					
		Based on Mean of MS and MSD	NA	NA	40 - 120	NA	NA
		PAH Duplicate Analysis - #1	17	13			76% Pass
		PAH Duplicate Analysis - #2	NA	NA	NA NA	< 20% @ 3 x MDL	NA
		LAB Duplicate Analysis - #1	NA	NA	] INA	of Sample Mean	NA
		LAB Duplicate Analysis - #2	NA	NA	]		NA

#### Notes:

OCSD laboratory results are not corrected for surrogate recoveries, causing some analytes with lower molecular weights and boiling points to fail the established criteria for SRM certified values. N/A=not applicable.

<sup>&</sup>lt;sup>1</sup> SRM certified values are based on the addition of selected compounds prior to extraction for use as internal standards for quantification purposes. (NIST, Certificate of Analysis, SRM 1941b, SRM 1944, Organics in Marine Sediment).

Table C-5 Continued.

Quarter	Sample Set	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD	Comments
		PAH SRM 1944	15	10	25% of the certified or		*67% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NA	93% Pass
		PAH Reporting Level Spike	25	25	60 -120		100% Pass
3	3 Sedcore_Jan11_DV	PAH Matrix Spike					
		Based on Mean of MS and MSD	25	25	40 - 120	NA	100% Pass
		PAH Duplicate Analysis - #1	17	9	NA	< 20% @ 3 x MDL of Sample Mean	**43% Pass
		PAH Duplicate Analysis - #2	N/A	NA	NA NA		N/A
		PAH SRM 1944	15	14	25% of the certified or		93% Pass
		PAH SRM 1941b	15	14	published acceptance limits <sup>1</sup>	NA	93% Pass
		PAH Reporting Level Spike	25	22	60 -120		88% Pass
4	Sedcore_Apr11_DV	PAH Matrix Spike					
		Based on Mean of MS and MSD	25	21	40 - 120	NA	84% Pass
		PAH Duplicate Analysis - #1	12	8	NA	< 20% @ 3 x MDL	**40% Pass
		PAH Duplicate Analysis - #2	N/A	NA	NA NA	of Sample Mean	N/A

#### Notes:

<sup>1</sup>SRM certified values are based on the addition of selected compounds prior to extraction for use as internal standards for quantification purposes. (NIST, Certificate of Analysis, SRM 1944, SRM 1941b, Organics in Marine Sediment).

OCSD laboratory results are not corrected for surrogate recoveries, thus causing some analytes with lower molecular weights and boiling points to fail the established criteria for SRM certified values.

N/A=not applicable

<sup>\*</sup>PAH results for SRM 1944 were low but the surrogate recoveries were within the limits

<sup>\*\*</sup>Variability within replicate samples are commonly observed, especially for detectable analytes which are near the reporting levels, have caused some analytes to fail the established criteria for precision factors. The surrogate recoveries of these replicates were within the limits, and visual inspection of the replicate samples did not reveal any obvious interferences. No further corrective action is needed for the replicates. The blank and matrix spikes were all within the specifications. Data set integrity was verified and accepted.

Table C-6. Method detection levels for PCB congeners and pesticides in sediments using GC/MS ion trap, July 2010–June 2011.

Parameter	ASE & GC/MS/MS Method Detection Limit (ng/g dry weight)	Parameter	ASE & GC/MS/MS Method Detection Limit (ng/g dry weight)
Aldrin	0.12	PCB 101	0.08
alpha-Chlordane	0.17	PCB 105	0.19
cis-NoNAchlor	0.20	PCB 110	0.16
Dieldrin	0.32	PCB 114	0.22
Endrin	0.53	PCB 118	0.18
gamma-BHC	0.12	PCB 119	0.09
gamma-Chlordane	0.15	PCB 123	0.18
Heptachlor	0.11	PCB 126	0.31
Heptachlor epoxide	0.19	PCB 128	0.22
Hexachlorobenzene	0.21	PCB 138	0.14
Mirex	0.14	PCB 149	0.12
trans-NoNAchlor	0.16	PCB 151	0.11
2,4'-DDD (o,p'-DDD)	0.15	PCB 153	NA
2,4'-DDE (o,p'-DDE)	0.13	PCB 153/168	0.28
2,4'-DDT (o,p'-DDT)	0.16	PCB 156	0.21
4,4'-DDD (p,p'-DDD)	0.17	PCB 157	0.22
4,4'-DDE (p,p'-DDE)	0.15	PCB 158	0.17
4,4'-DDT (p,p'-DDT)	0.18	PCB 167	0.28
4,4'-DDMU	0.50 <sup>1</sup>	PCB 168	NA
PCB 8	0.14	PCB 169	0.30
PCB 18	0.14	PCB 170	0.17
PCB 28	0.09	PCB 177	0.11
PCB 37	0.24	PCB 180	0.16
PCB 44	0.11	PCB 183	0.19
PCB 49	0.09	PCB 187	0.18
PCB 52	0.08	PCB 189	0.22
PCB 66	0.20	PCB 194	0.14
PCB 70	0.20	PCB 195	0.14
PCB 74	0.28	PCB 200	0.21
PCB 77	0.21	PCB 201	0.20
PCB 81	0.24	PCB 206	0.16
PCB 87	0.13	PCB 209	0.10
PCB 99	0.11		

<sup>&</sup>lt;sup>1</sup> Value is the reporting limit (RL).

NA = Not analyzed.

Table C-7. Method detection levels for PCB congeners and pesticides in sediments using GC/MS DSQII, July 2010–June 2011

Parameter	ASE & GC/MS/MS Method Detection Limit (ng/g dry weight)	Parameter	ASE & GC/MS/MS Method Detection Limit (ng/g dry weight)
Aldrin	0.06	PCB 101	0.13
alpha-Chlordane	0.13	PCB 105	0.14
cis-NoNAchlor	0.08	PCB 110	0.07
Dieldrin	0.16	PCB 114	0.13
Endrin	0.15	PCB 118	0.07
gamma-BHC	0.06	PCB 119	0.11
gamma-Chlordane	0.05	PCB 123	0.11
Heptachlor	0.06	PCB 126	0.08
Heptachlor epoxide	0.08	PCB 128	0.14
Hexachlorobenzene	0.04	PCB 138	0.13
Mirex	0.14	PCB 149	0.11
trans-NoNAchlor	0.09	PCB 151	0.10
2,4'-DDD (o,p'-DDD)	0.14	PCB 153	NA
2,4'-DDE (o,p'-DDE)	0.11	PCB 153/168	0.25
2,4'-DDT (o,p'-DDT)	0.14	PCB 156	0.07
4,4'-DDD (p,p'-DDD)	0.10	PCB 157	0.09
4,4'-DDE (p,p'-DDE)	0.08	PCB 158	0.12
4,4'-DDT (p,p'-DDT)	0.13	PCB 167	0.11
4,4'-DDMU	0.08	PCB 168	NA
PCB 8	0.06	PCB 169	0.13
PCB 18	0.04	PCB 170	0.08
PCB 28	0.05	PCB 177	0.10
PCB 37	0.15	PCB 180	0.11
PCB 44	0.09	PCB 183	0.13
PCB 49	0.07	PCB 187	0.11
PCB 52	0.05	PCB 189	0.10
PCB 66	0.09	PCB 194	0.17
PCB 70	0.11	PCB 195	0.13
PCB 74	0.11	PCB 200	0.11
PCB 77	0.07	PCB 201	0.17
PCB 81	0.07	PCB 206	0.16
PCB 87	0.06	PCB 209	0.29
PCB 99	0.17		

<sup>&</sup>lt;sup>1</sup> Value is the reporting limit (RL).

NA = Not analyzed.

Table C-8. Acceptance criteria for standard reference materials of pesticides/PCBs in sediments, July 2010–June 2011

Parameter	True Value		nce Range ng/g)	Parameter	True Value		tance Range (ng/g)
	(ng/g)	min.	max.		(ng/g)	min.	max.
	SRM 1944a - Or	ganics in Marin	e Sediment, Na	tional Institute of S	Standards and Ted	chnology,	
		New Y	ork/New Jersey	Waterway Sedime	ent		
alpha-Chlordane	16.51	15.7	17.3	PCB 99	37.5	35.1	39.9
cis-Nonachlor *	3.70	3.00	4.40	PCB 101	73.4	70.9	75.9
gamma-Chlordane *	8.00	6.00	10.0	PCB 105	24.5	23.4	25.6
Hexachlorobenzene	6.0	5.68	6.38	PCB 110	63.5	58.8	68.2
trans-Nonachlor	8.20	7.69	8.71	PCB 118	58.0	53.7	62.3
2,4'-DDD *	38.0	30.0	46.0	PCB 128	8.47	8.19	8.75
2,4'-DDE *	19.0	16.0	22.0	PCB 138	62.1	59.1	65.1
4,4'-DDD *	108	92.0	124	PCB 149	49.7	48.5	50.9
4,4'-DDE *	86.0	74.0	98.0	PCB 151	16.93	16.57	17.3
4,4'-DDT	119	108	130	PCB 153	74.0	71.1	76.9
2,4'-DDD *	38.0	30.0	46.0	PCB 156	6.52	5.86	7.18
PCB 8	22.3	20.0	24.6	PCB 170	22.6	21.2	24.0
PCB 18	51.0	48.4	53.6	PCB 180	44.3	43.1	45.5
PCB 28	80.8	78.1	83.5	PCB 183	12.19	11.6	12.8
PCB 44	60.2	58.2	62.2	PCB 187	25.1	24.1	26.1
PCB 49	53.0	51.3	54.7	PCB 194	11.2	9.80	12.6
PCB 52	79.4	77.4	81.4	PCB 195	3.75	3.36	4.14
PCB 66	71.9	67.6	76.2	PCB 206	9.21	8.70	9.72
PCB 87	29.9	25.6	34.2	1 05 200	0.21	0.10	0.72
	SRM 1941B - Or			ational Institute of S y Waterway Sedime		chnology,	
alpha-Chlordane	0.850	0.740	0.960	PCB 99	2.90	2.54	3.26
cis-Nonachlor	0.378	0.325	0.431	PCB 101	5.11	4.77	5.45
gamma-Chlordane	0.566						
Hexachlorobenzene		0.473	0.659	PCB 105	1.43	1.33	1.53
trans-Nonachlor	5.83	0.473 5.45	0.659 6.21	PCB 105 PCB 110	1.43 4.62	1.33 4.26	
แนกงาเพบเเสบเแบเ	5.83 0.438						1.53
2.4'-DDE *		5.45	6.21	PCB 110	4.62	4.26	1.53 4.98
	0.438	5.45 0.365	6.21 0.511	PCB 110 PCB 118	4.62 4.23	4.26 4.04	1.53 4.98 4.42
2.4'-DDE * 4,4'-DDE	0.438 0.380	5.45 0.365 0.260	6.21 0.511 0.500	PCB 110 PCB 118 PCB 128	4.62 4.23 0.696	4.26 4.04 0.652	1.53 4.98 4.42 0.740
2.4'-DDE * 4,4'-DDE 4,4'-DDD	0.438 0.380 3.22	5.45 0.365 0.260 2.94	6.21 0.511 0.500 3.50	PCB 110 PCB 118 PCB 128 PCB 138	4.62 4.23 0.696 3.60	4.26 4.04 0.652 3.32	1.53 4.98 4.42 0.740 3.88 4.61
2.4'-DDE * 4,4'-DDE 4,4'-DDD 4,4'-DDT *	0.438 0.380 3.22 4.66	5.45 0.365 0.260 2.94 4.20	6.21 0.511 0.500 3.50 5.12 1.54	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149	4.62 4.23 0.696 3.60 4.35	4.26 4.04 0.652 3.32 4.09	1.53 4.98 4.42 0.740 3.88
2.4'-DDE * 4,4'-DDE 4,4'-DDD 4,4'-DDT * PCB 8	0.438 0.380 3.22 4.66 1.12 1.65	5.45 0.365 0.260 2.94 4.20 0.700 1.46	6.21 0.511 0.500 3.50 5.12 1.54 1.84	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156	4.62 4.23 0.696 3.60 4.35 5.47 0.507	4.26 4.04 0.652 3.32 4.09 5.15 0.417	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597
2.4'-DDE * 4,4'-DDE 4,4'-DDD 4,4'-DDT * PCB 8 PCB 18	0.438 0.380 3.22 4.66 1.12 1.65 2.39	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 *	4.62 4.23 0.696 3.60 4.35 5.47 0.507	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800
2.4'-DDE * 4,4'-DDE 4,4'-DDD 4,4'-DDT * PCB 8 PCB 18 PCB 28	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 *	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 156 PCB 170 PCB 180 PCB 183 PCB 187	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39
2.4'-DDE * 4,4'-DDE 4,4'-DDD 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52 PCB 66	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24 4.96	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96 4.43	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52 5.49	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180 PCB 183 PCB 187 PCB 194	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95 0.980	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39 1.10
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52 PCB 66 PCB 70 *	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24 4.96 4.99	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96 4.43 4.70	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52 5.49 5.28	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180 PCB 183 PCB 187 PCB 194 PCB 195	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17 1.04 0.645	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95 0.980 0.585	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39 1.10 0.705
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52 PCB 66 PCB 70 * PCB 74 *	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24 4.96 4.99 2.04	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96 4.43 4.70 1.89	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52 5.49 5.28 2.19	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180 PCB 183 PCB 187 PCB 194 PCB 195 PCB 201	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17 1.04 0.645 0.770	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95 0.980 0.585 0.736	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39 1.10 0.705 0.804
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52 PCB 66 PCB 70 * PCB 74 * PCB 77 *	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24 4.96 4.99 2.04 0.310	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96 4.43 4.70 1.89 0.280	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52 5.49 5.28 2.19 0.340	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180 PCB 183 PCB 187 PCB 194 PCB 195 PCB 201 PCB 206	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17 1.04 0.645 0.770 2.42	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95 0.980 0.585 0.736 2.23	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39 1.10 0.705 0.804 2.61
2.4'-DDE * 4,4'-DDE 4,4'-DDT * PCB 8 PCB 18 PCB 28 PCB 44 PCB 49 PCB 52 PCB 66 PCB 70 * PCB 74 *	0.438 0.380 3.22 4.66 1.12 1.65 2.39 4.52 3.85 4.34 5.24 4.96 4.99 2.04	5.45 0.365 0.260 2.94 4.20 0.700 1.46 2.10 3.95 3.65 4.06 4.96 4.43 4.70 1.89	6.21 0.511 0.500 3.50 5.12 1.54 1.84 2.68 5.09 4.05 4.62 5.52 5.49 5.28 2.19	PCB 110 PCB 118 PCB 128 PCB 138 PCB 149 PCB 153/168 PCB 156 PCB 158 * PCB 170 PCB 180 PCB 183 PCB 187 PCB 194 PCB 195 PCB 201	4.62 4.23 0.696 3.60 4.35 5.47 0.507 0.650 1.35 3.24 0.979 2.17 1.04 0.645 0.770	4.26 4.04 0.652 3.32 4.09 5.15 0.417 0.500 1.26 2.73 0.892 1.95 0.980 0.585 0.736	1.53 4.98 4.42 0.740 3.88 4.61 5.79 0.597 0.800 1.44 3.75 1.07 2.39 1.10 0.705 0.804

<sup>\*</sup> non-certified

Table C-9. Sediment PCB/pesticide QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD	
		PCB	SRM 1944a	27	25	25% of the certified	NA	
		PCB	SRM 1941b	27	26	ranges or published acceptance limits	NA NA	
		PCB	Reporting Level Spike	44	44	60 -120	NA	
		PCB	Matrix Spike	44	37	40 - 120	NA	
		PCB	Matrix Spike Dup	44	37	40 - 120	NA	
		PCB	Matrix Spike Precision	44	44	NA	< 20%	
		Pesticide	SRM 1944a	4	4	25% of the certified	N/A	
		Pesticide	SRM 1941b	7	7	ranges or published acceptance limits	NA	
2	EO	Pesticide	Reporting Level Spike	19	8	60 -120	NA	
		Pesticide	Matrix Spike	19	16	40 - 120	NA	
		Pesticide	Matrix Spike Dup	19	15	40 - 120	NA	
		Pesticide	Matrix Spike Precision	19	18	NA	< 20%	
		PCB	Duplicate 1	3	3	NA	< 20% @ 3 x MDL	
		Pesticides	Duplicate 1	1	1	NA	of Sample Mean.	
		PCBs and Pesticides	Duplicate 1 Sum	1	1	NA	NA	
		PCB	Duplicate 2	NA	NA	NA	< 20% @ 3 x MDL	
		Pesticides	Duplicate 2	NA	NA	NA	of Sample Mean.	
		PCBs and Pesticides	Duplicate 2 Sum	NA	NA	NA	NA	

#### Comments:

Review of calibration check standards injected after sample injections, extraction notes, and instrument conditions did not indicate any atypical circumstances.

NA = Not Applicable

		PCB	SRM 1944a	27	27	25% of the certified	NA
		PCB	SRM 1941b	27	27	ranges or published acceptance limits	INA
		PCB	Reporting Level Spike	44	44	60 -120	NA
		PCB	Matrix Spike	44	44	40 - 120	NA
		PCB	Matrix Spike Dup	44	44	40 - 120	NA
		PCB	Matrix Spike Precision	44	44	NA	< 20%
		Pesticide	SRM 1944a	4	4	25% of the certified	NA
		Pesticide	SRM 1941b	7	7	ranges or published acceptance limits	NA
2.4	FO FB	Pesticide	Reporting Level Spike	19	19	60 -120	NA
3, 4	EQ, ER	Pesticide	Matrix Spike	19	19	40 - 120	NA
		Pesticide	Matrix Spike Dup	19	19	40 - 120	NA
		Pesticide	Matrix Spike Precision	19	19	NA	< 20%
		PCB	Duplicate 1	6	6	NA	< 20% @ 3 x MDL
		Pesticide	Duplicate 1	1	1	NA	of Sample Mean.
		PCBs and Pesticides	Duplicate 1 Sum	11	1	NA	NA
		PCB	Duplicate 2	NA	NA	NA	< 20% @ 3 x MDL
		Pesticide Duplicate 2 NA NA	NA	of Sample Mean.			
		PCBs and Pesticides	Duplicate 2 Sum	NA	NA	NA	NA

#### Comments:

Samples from the  $3^{rd}$  quarter (Set EQ) were re-extracted with samples from the  $4^{th}$  quarter (Set ER) due to contamination during the extraction process.

Review of calibration check standards injected after sample injections, extraction notes, and instrument conditions did not indicate any atypical circumstances.

NA = Not Applicable

duplicate, and the two spiked samples were evaluated for precision. The samples that were spiked with aluminum and iron were not evaluated for spike recoveries because the spike levels were extremely low compared to the concentrations of aluminum and iron in the native samples. The samples were spiked at 20 mg/kg dry weight whereas the native concentrations ranged between 5,000 and 35,000 mg/kg dry weight.

All samples were analyzed within their 6-month holding times. If any analyte exceeded the appropriate calibration curve, and Linear Dynamic Range, the sample was diluted and reanalyzed. MDLs for metals are presented in Table C-10. Acceptance criteria for trace metal SRMs are presented in Table C-11.

The digested samples were analyzed for silver, cadmium, chromium, copper, nickel, lead, zinc, selenium, arsenic, and beryllium by inductively coupled mass spectroscopy (ICPMS). Aluminum and iron were analyzed inductively using coupled emission spectroscopy (ICPES). Sediment trace metal QA/QC summary data are presented in Table C-12. All spike recoveries were between 88% and 106%. The relative percent difference (RPD) between the sample and its duplicate analysis ranged from -27.2% to 9.7%. The RPDs for the spike and spike duplicate analysis ranged from -2.2% to 2.5%.

## Analytical Methods - Mercury

Dried sediment samples were analyzed for mercury in accordance with methods described in the LOPM. QC for a typical batch included a blank, blank spike, and SRM. Sediment samples with duplicates, spiked samples and duplicate spiked samples were run approximately once every 10 sediment samples. All samples were analyzed within their 6-month holding time. When sample mercury concentration exceeded the appropriate calibration curve,

the sample was diluted with the reagent blank and reanalyzed. The samples were analyzed for mercury on a Perkin Elmer FIMS 400 system.

The MDL for sediment mercury is presented in Table C-10. Acceptance criteria for mercury SRM is presented in Table C-11. All QA/QC summary data are presented in Table C-12. The first and second quarter samples were run together in the same analysis batches.

All samples, with some noted exceptions, met the QA/QC criteria guidelines for accuracy and precision. Two duplicate analysis RPDs, two matrix spike recoveries, one matrix spike duplicate recovery, and one matrix spike precision RPD were out of range due to low results and non-homogeneous sample matrices.

## <u>Analytical Methods - Dissolved Sulfides</u>

Dissolved sulfides samples were analyzed in accordance with methods described in the LOPM. The MDL for dissolved sulfides is presented in Table C-13. Sediment dissolved sulfides QA/QC summary data are presented in Table C-14. All samples were analyzed within their required holding times. All analyses met the QA/QC criteria for blanks, blank spikes, matrix spike dups, and matrix spike precisions. One of five matrix spike recoveries was out of control due to matrix interferences.

<u>Analytical Methods - Total Organic Carbon</u> Total Organic Carbon (TOC) samples were contract laboratory. analvzed by а Columbia Analytical Services, Kelso, WA. The MDL for TOC is presented in Table C-13. Sediment TOC QA/QC summary data are presented in Table C-15. The samples were analyzed within their required holding times. Three samples were analyzed in duplicate. The samples and their duplicate analyses had a RPD of less than 10%. The recoveries for matrix spike were within the 80-120% range.

Table C-10. Method detection limits for trace metals in sediments, July 2010–June 2011.

Parameter	Detection Limits (mg/kg dry weight)			
Aluminum	50			
Arsenic	0.15			
Beryllium	0.01			
Cadmium	0.01			
Chromium	0.15			
Copper	0.10			
Iron	50			
Lead	0.10			
Nickel	0.10			
Mercury	0.00011			
Selenium	0.15			
Silver	0.02			
Zinc	0.15			

Table C-11. Acceptance criteria for standard reference materials of metals in sediments, July 2010–June 2011

Parameter	True Value (mg/kg)		ptance Criteria //kg)						
	(9/9)	Min.	Max.						
Priority Pollutn	Environmental Resource Associates D056-540 Priority PollutnT™/CLP Inorganic Soils – Microwave Digestion Environmental Resource Associates								
Aluminum	10400	6370	14400						
Arsenic	280	226	333						
Beryllium	51	42.4	59.6						
Cadmium	182	149	215						
Chromium	142	115	170						
Copper	132	110	155						
Iron	16600	9490	23700						
Lead	72.2	59.1	85.4						
Nickel	155	128	182						
Selenium	165	128	203						
Silver	126	83.7	169						
Zinc	346	273	418						
		corporation CRM016-050 ference Material Lot BE016							
Mercury	0.158	0.00	0.357						

Table C-12. Sediment metals QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
		Arsenic.	Blank	10	10	<3X MDL	N/A
		Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Summer	HMSED100928-1	Copper,	Matrix Spike Dup	10	10	70-130	
Cummon	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Lead, Nickel,	Matrix Spike Precision	10	10		< 20%
		Selenium,	Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
		Silver, Zinc	CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
		Arsenic,	Blank	10	10	<3X MDL	N/A
		Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Summer	HMSED101007-1	Copper,	Matrix Spike Dup	10	10	70-130	
		Lead, Nickel,	Matrix Spike Precision	10	10		< 20%
		Selenium,	Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
		Silver, Zinc	CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
		Arsenic,	Blank	10	10	<3X MDL	N/A
		Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Summer	HMSED101026-1	Copper,	Matrix Spike Dup	10	10	70-130	
		Lead, Nickel.	Matrix Spike Precision	10	10		< 20%
		Selenium,	Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
		Silver, Zinc	CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
			Blank	2	2	<3X MDL	N/A
Summer	r   ALEESED101004-1	Aluminum, Iron	Duplicate Analysis	2	2	NA	@ ≥ 10 X MDL < 20%
		IIOII	CRM Analysis	2	2	80-120% or certified value, whichever is greater	
			Blank	2	2	<3X MDL	N/A
Summer	ALFESED101012-1	Aluminum, Iron	Duplicate Analysis	2	2	NA	@ <u>&gt;</u> 10 X MDL < 20%
		II OII	CRM Analysis	2	2	80-120% or certified value, whichever is greater.	

Table C-12 Continues.

Table C-12 Continued.

Quarter	Sample Set	Parameter	Description	Description  Number of Compounds Compounds Passed  Number of Compounds Passed  Target Accuracy % Recovery		Target Precision % RPD	
			Blank	2	2	<3X MDL	N/A
Summer	ALFESED101104-1	Aluminum,	Duplicate Analysis	2	2	NA	@ <u>&gt;</u> 10 X MDL < 20%
Summer	ALI ESEDIOTIO4-1	Iron	CRM Analysis	2	2	80-120% or certified value, whichever is greater.	
			Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	90-110	N/A
			Matrix Spike	2	**1	70-130	
Summer	HGSED100928-1	Mercury	Matrix Spike Dup	2	**1	70-130	
Summer	1103LD100920-1	Wiercury	Matrix Spike Precision	2	**1		< 25%
			Duplicate Analysis	2	**1	NA	@ <u>&gt;</u> 10 X MDL < 30%
			CRM Analysis	1	1	80-120% or certified value, whichever is greater.	
			Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	90-110	N/A
			Matrix Spike	3	**2	70-130	
Summer	HGSED101006-1	Mercury	Matrix Spike Dup	3	3	70-130	
Summer	110320101000-1	Wiercury	Matrix Spike Precision	3	**2		< 25%
			Duplicate Analysis	3	**2	NA	@ <u>&gt;</u> 10 X MDL < 30%
			CRM Analysis	1	1	80-120% or certified value, whichever is greater.	
			Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	90-110	N/A
			Matrix Spike	2	2	70-130	
Summer/	Summer/ HGSED101102-1	Mercury	Matrix Spike Dup	2	2	70-130	
Fall	110000101102-1	ivicicuiy	Matrix Spike Precision	2	2		< 25%
			Duplicate Analysis	2	2	NA	@ <u>&gt;</u> 10 X MDL < 30%
			CRM Analysis	1	1	80-120% or certified value, whichever is greater.	

<sup>\*\*</sup> Two duplicate analysis RPDs were out of range due to low results and non-homogeneous samples. Two matrix spike recoveries, one matrix spike duplicate recovery, and one matrix spike precision RPD failed due to non-homogeneous sample matrices.

Table C-12 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
		Arsenic.	Blank	10	10	<3X MDL	N/A
		Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Fall	HMSED101103-1	Copper,	Matrix Spike Dup	10	10	70-130	
ı alı	TIMOLD TO TIOS-T	Lead, Nickel,	Matrix Spike Precision	10	10		< 20%
		Selenium,	Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
		Silver, Zinc	CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
			Blank	2	2	<3X MDL	N/A
Fall	ALFESED101104-2	Aluminum,	Duplicate Analysis	2	2	NA	@ ≥ 10 X MDL < 20%
		Iron	CRM Analysis	2	2	80-120% or certified value, whichever is greater	
		Arsenic,	Blank	10	10	<3X MDL	N/A
		Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Winter	HMSED110217-1	Copper,	Matrix Spike Dup	10	10	70-130	
VVIIILEI	LIMPED LIDZ I I-1	Lead, Nickel,	Matrix Spike Precision	10	10		< 20%
		Selenium,	Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
	Silver, Zinc		CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
			Blank	2	2	<3X MDL	N/A
Winter	ALFESED110218-1	Aluminum,	Duplicate Analysis	2	2	NA	@ <u>&gt;</u> 10 X MDL < 20%
vviillei	ALITEGED   10210-1	Iron	CRM Analysis	2	2	80-120% or certified value, whichever is greater.	

Table C-12 Continues.

Table C-12 Continued.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
			Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	90-110	N/A
			Matrix Spike	1	**0	70-130	
Winter	HGSED110222-1	Mercury	Matrix Spike Dup	1	**0	70-130	
VVIIICI	110000110222-1	Wichcury	Matrix Spike Precision	1	**0		< 25%
			Duplicate Analysis	1	1	NA	@ <u>&gt;</u> 10 X MDL < 30%
			CRM Analysis	1	1	80-120% or certified value, whichever is greater.	_
		Arsenic,	Blank	10	10	<3X MDL	N/A
	Beryllium, Cadmium,	Beryllium,	Blank Spike	10	10	90-110	N/A
		Cadmium, Chromium,	Matrix Spike	10	10	70-130	
Spring	UMSED110526 1	Copper, Lead, Nickel, Selenium,	Matrix Spike Dup	10	10	70-130	
Opinig	11W3LD110320-1		Matrix Spike Precision	10	10		< 20%
			Duplicate Analysis	10	10	NA	@ <u>&gt;</u> 10 X MDL < 20%
		Silver, Zinc	CRM Analysis	10	10	80-120% or certified value, whichever is greater.	
			Blank	2	2	<3X MDL	N/A
Spring	ALFESED110602-1	Aluminum,	Duplicate Analysis	2	2	NA	@ <u>&gt;</u> 10 X MDL < 20%
Opinig	NEI EGEBT 10002 1	Iron	CRM Analysis	2	2	80-120% or certified value, whichever is greater.	
			Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	90-110	N/A
			Matrix Spike	1	1	70-130	
Carina	UCCED110521 1	Morouni	Matrix Spike Dup	1	1	70-130	
opring	Spring HGSED110531-1	Mercury	Matrix Spike Precision	1	1		< 25%
			Duplicate Analysis	1	1	NA	@ <u>&gt;</u> 10 X MDL < 30%
			CRM Analysis	1	1	80-120% or certified value, whichever is greater.	

<sup>\*\*</sup> One Matrix spike recovery, one matrix spike duplicate recovery, and one matrix spike precision RPD were out of range due to non-homogeneous sample matrices

NA = Not applicable.

Table C-13. Method detection limits for dissolved sulfides, total organic carbon, and grain size in sediments, July 2010–June 2011.

Orange County Sanitation District, California.

Parameter	Detection Limits
Dissolved Sulfides (OCSD)	1.03 mg/kg dry weight
Total Organic Carbon (Columbia Analytical Services)	0.05%
Grain Size (Weston Solutions, Inc.)	0.001 %

Table C-14. Sediment dissolved sulfides QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
	SULFIDE100720-1		Method Blank	5	5	<2X MDL	N/A
	SULFIDE100728-1		Blank Spike	5	5	80 -120	N/A
Summer	SULFIDE100819-1 SULFIDE100824-1	Dissolved Sulfides	Matrix Spike	5	4*	70 - 130	
	SULFIDE100826-1		Matrix Spike Dup	5	5	70 - 130	
			Matrix Spike Precision	5	5		<30%
			Method Blank	1	1	<2X MDL	N/A
		DE101021-1 Dissolved Sulfides	Blank Spike	1	1	80 -120	N/A
Fall	SULFIDE101021-1		Matrix Spike	1	1	70 - 130	
			Matrix Spike Dup	1	1	70 - 130	
			Matrix Spike Precision	1	1		<30%
			Method Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	80 -120	N/A
Winter	SULFIDE110111-1	Dissolved Sulfides	Matrix Spike	1	1	70 - 130	
			Matrix Spike Dup	1	1	70 - 130	
			Matrix Spike Precision	1	1		<30%
			Method Blank	1	1	<2X MDL	N/A
			Blank Spike	1	1	80 -120	N/A
Spring	SULFIDE110412-1	412-1 Dissolved Sulfides	Matrix Spike	1	1	70 - 130	
			Matrix Spike Dup	1	1	70 - 130	
			Matrix Spike Precision	1	1		<30%

<sup>\*,</sup> Matrix spike recovery (67%) was out of control due to matrix interferences.

Table C-15. Sediment total organic carbon QA/QC summary, July 2010–June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
Summer	TOC-100928-1	Total Organic Carbon	Duplicate and matrix spike	3	3	80-120 <sup>1</sup>	10% <sup>1</sup>
Fall	TOC-101123-1	Total Organic Carbon	Duplicate and Matrix Spike	1	1*	80-120 <sup>1</sup>	10% <sup>1</sup>
Winter	TOC-110201-1	Total Organic Carbon	Duplicate and Matrix Spike	1	1	80-120 <sup>1</sup>	10% <sup>1</sup>
Spring	TOC-110512-1	Total Organic Carbon	Duplicate and Matrix Spike	2	2	80-120 <sup>1</sup>	10% <sup>1</sup>

<sup>&</sup>lt;sup>1</sup> TOC Target Precision/Accuracy of QC Criteria is not described in the Core Monitoring Quality Assurance Project Plan. \* One sample dup precision (16.7%) was out of range due to matrix interference.

## Table C-16. Sediment Grain Size QA/QC Summary, July 2010 – June 2011.

Quarter	Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
Summer	PSIZE101012-1	2-1 Grain Size	Reference Standard	15	15	NA	Mean ± 3 σ of the reference standard for median phi, skewness, dispersion, % gravel, % sand, % clay, and % silt
			Duplicate	11	11		≤10%
Fall	PSIZ-110107-1	-1 Grain Size	Reference Standard	1	1	NA	Mean ± 3 σ of the reference standard for median phi, skewness, dispersion, % gravel, % sand, % clay, and % silt
			Duplicate	1	1		≤10%
Winter	PSIZ-110317-1	Z-110317-1 Grain Size	Reference Standard	2	2	NA	Mean ± 3 σ of the reference standard for median phi, skewness, dispersion, % gravel, % sand, % clay, and % silt
			Duplicate	1	1		≤10%
Spring	PSIZE110531-1	PSIZE110531-1 Grain Size	Reference Standard	2	2	NA	Mean ± 3 σ of the reference standard for median phi, skewness, dispersion, % gravel, % sand, % clay, and % silt
			Duplicate	1	1		≤10%

## Analytical Methods - Grain Size

Grain size samples were analyzed by a contract laboratory, Weston Solutions, Carlsbad, CA. The MDL for sediment grain size is presented in Table C-13. Sediment grain size QA/QC summary data are presented in Table C-16. Fifteen standard reference material (SRM) samples were analyzed. All analyses were within three standard deviations of SRM for the statistical parameters (median phi. dispersion, and skewness), percent gravel, percent sand, percent clay, and percent silt. samples and their duplicate Eleven analyses had a RPD ≤10%.

## **SECOND QUARTER (OCTOBER 2010)**

OCSD's laboratory received 10 sediment samples from the ocean monitoring staff during the month of October 2010. All samples were stored according to methods described in the LOPM. All samples were analyzed for organochlorine pesticides, PCB congeners, PAHs, trace metals, mercury, dissolved sulfides, grain size, and TOC.

All sediment samples that were analyzed for organochlorine pesticides and PCB congeners were extracted on November 11, 2010. All sediment samples that were analyzed for PAHs were extracted on November 17, 2010. Any variances that occurred during sample processing or analysis are noted in the Comments/Notes section of each batch summary. All sediment samples were extracted using an ASE.

## <u>Analytical Methods - Organochlorine</u> <u>Pesticides and PCB Congeners</u>

The analytical methods used to process the organochlorine pesticides and PCB congeners samples are described in the LOPM. All sediment extracts were analyzed by GC/MS. Approximately 10 grams (dry weight) of sample were used for each analysis. If a field blank and rinse

were included in the batch, a separatory funnel extraction was performed using 100 milliliters of sample.

A typical sample batch consisted of 10 field samples with required QC samples, which included one sand blank, two SRM, one PCB/pesticide reporting level spike, one PCB/pesticide matrix spike set, and one duplicate sample extraction. There were two batches extracted. In addition, one batch contained a rinse sample and a field MDLs for PCBs/pesticides are Table C-6 presented in and C-7. Acceptance Criteria for PCB/pesticide SRMs are presented in Table C-8.

Sediment PCB/pesticide QA/QC summary data are presented in Table C-9. analyses were performed within QAPP stated holding times and with appropriate quality control measures. When constituent concentrations exceeded the calibration range of the instrument, dilutions were performed and the samples reanalyzed. Anv variances are noted in the Comments/Notes section of each batch summary.

All samples were analyzed for metals within their holding times. All of the metals analyses met the QA criteria guidelines. Sediment metals QA/QC summary data are presented in Table C-12. All spike recoveries were between 86.9% and 103.1%. The RPDs of the sample and its duplicate ranged from -16.1% to 2.1%. The RPDs of the spike and spike duplicate ranged from -3.3% and 3.1%.

Sediment Mercury QA/QC summary data are presented in Table C-12. Fall samples were batched with summer samples and all samples met the QA criteria guidelines.

The analyses for dissolved sulfides, TOC, and grain size met criteria guidelines as specified in the project QAPP. MDL, SRM, and QA/QC summary data are presented in

Tables C-13 through C-16. One TOC sediment duplicate analysis had a precision greater than 10% due to matrix interference.

## **THIRD QUARTER (JANUARY 2011)**

OCSD's laboratory received 10 sediment samples from the ocean monitoring staff during the month of January 2011. All samples were stored according to methods described in the LOPM. All samples were analyzed for organochlorine pesticides, PCB congeners, PAHs, trace metals, mercury, dissolved sulfides, grain size, and TOC.

All sediment samples that were analyzed for organochlorine pesticides and PCB congeners were first extracted on February 24, 2011. Due to contamination during the extraction process, all samples were reextracted on April 20, 2011. All sediment samples that were analyzed for PAHs were extracted on January 20, 2011. variances are noted in the Comments/Notes section of each batch All sediment samples were extracted using an ASE. All sediment extracts for PCB congeners and pesticides were analyzed by GC/MS.

All samples were analyzed for metals within their holding times. Sediment metals QA/QC summary data are presented in Table C-12. All spike recoveries were between 89.5% and 113.0%. The RPDs of the sample and its duplicate ranged from -17.0% and 3.1%. The RPDs of the spike and spike duplicate ranged from -2.9% and -1.1%.

Sediment mercury QA/QC summary data are presented in Table C-12. All samples met the QA criteria guidelines except for one matrix spike recovery, one matrix spike duplicate recovery, and one matrix spike precision RPD due to low results and non-homogeneous sample matrices.

The analyses for dissolved sulfide, TOC, and grain size met the QA criteria guidelines specified in the QAPP. MDL, SRM, and QA/QC summary data are presented in Tables C-13 through C-16.

## **FOURTH QUARTER (APRIL 2011)**

OCSD's laboratory received 10 sediment samples from the ocean monitoring staff during the month of April 2011. All samples were stored according to methods described in the LOPM. All samples were analyzed for organochlorine pesticides, PCB congeners, PAHs, trace metals, mercury, dissolved sulfides, grain size, and TOC.

All sediment samples being analyzed for organochlorine pesticides and PCB congeners were extracted on April 20. All sediment samples being analyzed for PAHs were extracted on May 15, 2011. Any variances, which may have occurred during sample processing or analysis, are noted in the Comments/Notes section of each batch summary. sediment samples were extracted using an All sediment extracts for PCB ASE. congeners and pesticides were analyzed by Ion Trap GC/MS.

All samples were analyzed for metals within their holding times. All metal analyses met the QA objectives. Sediment metals QA/QC summary data are presented in Table C-12. All spike recoveries were between 89.5% and 105.8%. The RPDs of the sample and its duplicate ranged from -7.6% and 12.8%. The RPDs of the spike and spike duplicate ranged from 0.9% and 2.2%. Sediment mercury QA/QC summary data are presented in Table C-12. samples met the QA criteria guidelines. The analyses for dissolved sulfide, TOC, and grain size met the QA criteria guidelines specified in the QAPP. MDL, SRM, and QA/QC summary data are presented in Tables C-13 through C-16.

# FISH TISSUE CHEMISTRY NARRATIVE

## **THIRD QUARTER (JANUARY 2011)**

### Introduction

OCSD's laboratory received 40 individual fish samples and 12 composite samples (containing six fish per bag), from the ocean monitoring staff during the month of The individual samples January 2011. were stored, dissected, and homogenized according to methods described in the LOPM. A 1:1 muscle to water ratio was used. No water was used during liver homogenization. After the individual samples were homogenized, equal aliquots of muscle and liver from each sample were frozen and distributed to the inorganic and organic chemistry sections of the laboratory for analyses. Each of the 12 composites were weighed and homogenized using a 1:1 whole body fish to water ratio, according to methods described in the LOPM. After the composites were homogenized, equal aliquots were frozen and distributed to the inorganic and organic chemistry sections of the laboratory for analyses.

The Organic Chemistry Section extracted 40 fish muscle samples, 40 fish liver samples, and 12 whole body composite samples and analyzed them for PCB congeners and organochlorine pesticides. Percent lipid content was also determined for each sample. A typical organic tissue sample batch included 15 field samples with required QC samples. The QC samples included one hydromatrix blank, two duplicate sample extractions, one matrix spike, one matrix duplicate spike, two SRMs, and one reporting level spike (matrix of store bought, farm-raised, Tilapia).

For mercury analysis, one sample batch consisted of 15–20 fish tissue samples and the required QC samples, which included a

blank, blank spike, SRM, sample duplicates, matrix spikes, and matrix spike duplicates.

## <u>Analytical Methods - Organochlorine</u> Pesticides and PCB Congeners

The analytical methods used for organochlorine pesticides and PCB congeners were according to methods described in the LOPM. All fish tissue was extracted using an ASE 200 and analyzed by GC/MS.

The MDLs for pesticides and PCBs in fish tissue are presented in Tables C-17 and C18. Acceptance criteria for PCB SRMs in fish tissue are presented in Tables C-19 and C-20. Fish tissue pesticide and PCB QA/QC summary data are presented in Table C-21. All analyses were performed within the required holding times and with appropriate quality control measures. cases where constituent concentrations exceeded the calibration range of the instrument, the samples were diluted and reanalyzed. Any variances that occurred during sample preparation or analyses are noted in the Comments/Notes section of each batch summary.

## <u>Analytical Methods – Lipid Content</u>

Percent lipid content was determined for each sample using methods described in the LOPM. Lipids were extracted by dichloromethane from approximately 1 to 2 g of sample and concentrated to 2 mL. A 100 uL aliquot of the extract was placed in a tarred aluminum weighing boat and the solvent allowed to evaporate to dryness. The remaining residue was weighed, and the percent lipid content calculated. Lipid content QA/QC summary data presented in Table C-22. All analyses were performed within the required holding times appropriate quality control and with Any variances that occurred measures. during sample preparation or analyses are noted in the Comments/Notes section of the Fish Tissue Percent QA/QC Summary.

Table C-17. Method detection levels for pesticides and PCB congeners in fish tissue using GC/MS ion trap, July 2010–June 2011.

Parameters	Method Detection Limit ng/g wet weight	Parameters	Method Detection Limit ng/g wet weight						
Pesticides									
o,p'-DDD	0.90	Dieldrin	1.0						
o,p'-DDE	0.80	Endrin	1.4						
o,p'-DDT	0.68	gamma-BHC	0.72						
p,p'-DDD	1.2	gamma-Chlordane	0.78						
p,p'-DDE	0.92	Heptachlor	0.71						
p,p'-DDT	0.85	Heptachlor epoxide	0.72						
p,p'-DDMU	0.50	Hexachlorobenzene	0.83						
Aldrin	0.67	Mirex	0.63						
alpha-Chlordane	0.75	trans-Nonachlor	0.83						
cis-Nonachlor	0.70								
	PCB Con	geners							
PCB 8	0.86	PCB 128	0.65						
PCB 18	0.54	PCB 138	0.86						
PCB 28	0.70	PCB 149	1.1						
PCB 37	0.66	PCB 151	0.61						
PCB 44	0.68	PCB 156	1.0						
PCB 49	0.87	PCB 157	1.2						
PCB 52	0.73	PCB 158	1.2						
PCB 66	0.65	PCB 167	1.3						
PCB 70	1.2	PCB 168/153	2.6						
PCB 74	1.1	PCB 169	1.5						
PCB 77	1.3	PCB 170	1.3						
PCB 81	0.83	PCB 177	1.2						
PCB 87	0.87	PCB 180	0.64						
PCB 99	0.90	PCB 183	0.88						
PCB 101	0.84	PCB 187	1.1						
PCB 105	1.1	PCB 189	1.3						
PCB 110	0.84	PCB 194	0.97						
PCB 114	0.59	PCB 195	0.77						
PCB 118	1.1	PCB 200	1.2						
PCB 119	0.84	PCB 201	0.91						
PCB 123	1.1	PCB 206	1.1						
PCB 126	1.1	PCB 209	1.2						

Table C-18. Method detection levels for pesticides and PCB congeners in fish tissue using GC/MS DSQII, July 2010–June 2011.

Parameters	Method Detection Limit ng/g wet weight	Parameters	Method Detection Limit ng/g wet weight						
Pesticides									
o,p'-DDD	0.33	Dieldrin	0.31						
o,p'-DDE	0.23	Endrin	0.64						
o,p'-DDT	0.33	gamma-BHC	0.21						
p,p'-DDD	0.16	gamma-Chlordane	0.25						
p,p'-DDE	0.31	Heptachlor	0.23						
p,p'-DDT	0.24	Heptachlor epoxide	0.37						
p,p'-DDMU	0.43	Hexachlorobenzene	0.32						
Aldrin	0.30	Mirex	0.29						
alpha-Chlordane	0.33	trans-Nonachlor	0.21						
cis-Nonachlor	0.19								
	PCB Cor	ngeners							
PCB 8	0.24	PCB 128	0.08						
PCB 18	0.24	PCB 138	0.16						
PCB 28	0.21	PCB 149	0.33						
PCB 37	0.27	PCB 151	0.22						
PCB 44	0.36	PCB 156	0.10						
PCB 49	0.17	PCB 157	0.10						
PCB 52	0.17	PCB 158	0.18						
PCB 66	0.26	PCB 167	0.09						
PCB 70	0.23	PCB 168/153	0.23						
PCB 74	0.24	PCB 169	0.15						
PCB 77	0.21	PCB 170	0.18						
PCB 81	0.19	PCB 177	0.09						
PCB 87	0.17	PCB 180	0.18						
PCB 99	0.44	PCB 183	0.13						
PCB 101	0.14	PCB 187	0.06						
PCB 105	0.13	PCB 189	0.12						
PCB 110	0.19	PCB 194	0.17						
PCB 114	0.10	PCB 195	0.13						
PCB 118	0.22	PCB 200	0.08						
PCB 119	0.14	PCB 201	0.20						
PCB 123	0.21	PCB 206	0.11						
PCB 126	0.11	PCB 209	0.29						

Table C-19. Acceptance criteria for standard reference materials of PCB congeners in fish tissue, CARP-2, July 2010–June 2011.

Parameter	True Value (ng/g)	Acceptance Range (ng/g)			
	(ng/g)	Minimum	Maximum		
PCB 18	27.3	23.3	31.3		
PCB 28	34.0	26.8	41.2		
PCB 52	138	95.0	181		
PCB 44	86.6	60.7	112		
PCB 118	148	115	181		
PCB 153	105	83.0	127		
PCB 128	20.4	16.0	24.8		
PCB 180	53.3	40.3	66.3		
PCB 194	10.9	7.80	14.0		
PCB 206	4.40	3.30	5.50		

CARP-2, Ground Whole Carp Reference Material for Organochlorine Compounds, National Research Council Canada.

Table C-20. Acceptance criteria for standard reference materials of pesticides and PCB congeners in fish tissue, SRM-1946, July 2010–June 2011.

Orange County Sanitation District, California.

Parameter	True Value (ng/g)	Acceptance Range (ng/g)		Parameter	True Value (ng/g)	Acceptance Range (ng/g)	
	(119/9)	Minimum	Maximum		(119/9)	Minimum	Maximum
gamma-BHC	1.14	0.96	1.32	PCB 99	25.6	23.3	27.9
Dieldrin	32.5	29.0	36.0	PCB 101	34.6	32.0	37.2
Heptachlor epoxide	5.50	5.27	5.73	PCB 105	19.9	19.0	20.8
Hexachlorobenzene	7.25	6.42	8.08	PCB 110	22.8	20.8	24.8
alpha-Chlordane	32.5	30.7	34.3	PCB 118	52.1	51.1	53.1
gamma-Chlordane	8.36	7.45	9.27	PCB 126	0.380	0.363	0.397
cis-Nonachlor	59.1	55.5	62.7	PCB 128	22.8	20.9	24.7
trans-Nonachlor	99.6	92.0	107	PCB 138	115	102	128
Mirex	6.47	5.70	7.24	PCB 149	26.3	25.0	27.6
o,p'-DDD	2.20	1.95	2.45	PCB 153/168	170	161	179
p,p'-DDD	17.7	14.9	20.5	PCB 156	9.52	9.01	10.0
p,p'-DDE	373	325	421	PCB 169	0.106	0.092	0.120
p,p'-DDT	37.2	33.7	40.7	PCB 170	25.2	23.0	27.4
PCB 44	4.66	3.80	5.52	PCB 180	74.4	70.4	78.4
PCB 49	3.80	3.41	4.19	PCB 183	21.9	19.4	24.4
PCB 52	8.1	7.10	9.10	PCB 187	55.2	53.1	57.3
PCB 66	10.8	8.90	12.7	PCB 194	13.0	11.7	14.3
PCB 70	14.9	14.3	15.5	PCB 195	5.30	4.85	5.75
PCB 74	4.83	4.32	5.34	PCB 206	5.40	4.97	5.83
PCB 77	0.327	0.302	0.352	PCB 209	1.30	1.09	1.51
PCB 87	9.4	8.00	10.8				

SRM 1946, Organics in Lake Superior Fish Tissue, National Institute of Standards and Technology.

Table C-21. Fish tissue PCB/pesticide QA/QC summary, July 2010–June 2011.

Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD				
Sample Set – MB (15 Muscle Tissue Samples)								
NRCC CARP-2	10	10	according to published acceptance criteria	NA				
SRM 1946	40	30	according to published acceptance criteria	NA				
PCB Reporting Level Spike	44	44	75 -125	NA				
PCB Matrix Spike:	44	44	70 - 130	NA				
PCB Matrix Spike Dup	44	43	70 - 130	INA				
Precision	44	44	NA	< 25%				
Pesticide Reporting Level Spike	19	18	75 -125	NA				
Pesticide Matrix Spike	19	19	70-130	NA				
Pesticide Matrix Spike Dup	19	18	70-130	INA				
Precision	19	19	NA	< 25%				
PCB/Pesticide Duplicate Analysis								
Duplicate 1 PCB	1	0		0-0/ 0 0 1/0/ 6				
Duplicate 1 Pesticides	3	3	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 1 Sum of Pesticides and PCBs	1	1		Sample Mean.				
Duplicate 2 PCB	8	8		0-0/ 0 0 1/0/ 6				
Duplicate 2 Pesticides	4	4	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 2 Sum of Pesticides and PCBs	1	1		Sample Mean.				
Sa	mple Set – MA2	(15 Muscle T	issue Samples)					
NRCC CARP-2	10	10	according to published acceptance criteria	NA				
SRM 1946	40	38	according to published acceptance criteria	NA				
PCB Reporting Level Spike	44	43	75 -125	NA				
PCB Matrix Spike:	44	43	70 - 130	NA				
PCB Matrix Spike Dup	44	43	70 - 130	INA				
Precision	44	44	NA	< 25%				
Pesticide Reporting Level Spike	19	19	75 -125	NA				
Pesticide Matrix Spike	19	19	70 120	NIA				
Pesticide Matrix Spike Dup	19	19	70-130	NA				
Precision	19	19	NA	< 25%				
PCB/Pesticide Duplicate Analysis								
Duplicate 1 PCB	0	0		070/ 0 0 110/ 5				
Duplicate 1 Pesticides	1	1	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 1 Sum of Pesticides and PCBs	cate 1 Sum of Pesticides and PCBs 1 1			Gample Mean.				
Duplicate 2 PCB	0	0						
Duplicate 2 Pesticides	1	1	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 2 Sum of Pesticides and PCBs	1	1		Gampie Mean.				

Table C-21 Continues.

Table C-21 Continued.

Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD	
Sample Set – I	MC2 (10 Muscle	Tissue and 5 W	hole Body Tissue Samples)		
NRCC CARP-2	10	10	according to published acceptance criteria	NA	
SRM 1946	40	38	according to published acceptance criteria	NA	
PCB Reporting Level Spike	44	44	75 -125	NA	
PCB Matrix Spike:	44	42	70 - 130	NIA	
PCB Matrix Spike Dup	44	42	70 - 130	NA	
Precision	44	44	NA	< 25%	
Pesticide Reporting Level Spike	19	19	75 -125	NA	
Pesticide Matrix Spike	19	19	70-130	NA	
Pesticide Matrix Spike Dup	19	19	70-130	INA	
Precision	19	19	NA	< 25%	
PCB/Pesticide Duplicate Analysis					
Duplicate 1 PCB	0	0		4 0 E 0 / A 0 / M D 1 - 4	
Duplicate 1 Pesticides	2	1	NA	< 25% @ 3 x MDL of Sample Mean.	
Duplicate 1 Sum of Pesticides and PCBs	1	1		Campio Moan.	
Duplicate 2 PCB	0	0		1050/ @ 0 MDL - f	
Duplicate 2 Pesticides	2	2	NA	< 25% @ 3 x MDL of Sample Mean.	
Duplicate 2 Sum of Pesticides and PCBs	1	1		oup.oou	
Sar	nple Set – MD2	(7 Whole Body	Tissue Samples)		
NRCC CARP-2	10	10	according to published acceptance criteria	NA	
SRM 1946	40	40	according to published acceptance criteria	NA	
PCB Reporting Level Spike	44	43	75 -125	NA	
PCB Matrix Spike:	44	43	70 420	NIA	
PCB Matrix Spike Dup	44	43	70 - 130	NA	
Precision	44	44	NA	< 25%	
Pesticide Reporting Level Spike	19	19	75 -125	NA	
Pesticide Matrix Spike	19	19	70-130	NIA	
Pesticide Matrix Spike Dup	19	19	70-130	NA	
Precision	19	19	NA	< 25%	
PCB/Pesticide Duplicate Analysis					
Duplicate 1 PCB	0	0		4 0 E 0 / A 0 / A D 1 - 5	
Duplicate 1 Pesticides	3	2	NA	< 25% @ 3 x MDL of Sample Mean.	
Duplicate 1 Sum of Pesticides and PCBs	11	1		Campic Mcan.	
Duplicate 2 PCBs	0	0		4.0E0/ @ 0 MDL = 5	
Duplicate 2 Pesticides	1	1	NA	< 25% @ 3 x MDL of Sample Mean.	
Duplicate 2 Sum of Pesticides and PCBs	1	1		Sample Modifi	

Table C-21 Continues.

Table C-21 Continued.

Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD				
Sample Set – LA (15 Liver Tissue Samples)								
NRCC CARP-2	10	10	according to published acceptance criteria	NA				
SRM 1946 *	40	37	according to published acceptance criteria					
PCB Reporting Level Spike	44	43	75 -125	NA				
PCB Matrix Spike:	44	43	70 120	NIA				
PCB Matrix Spike Dup	44	43	70 - 130	NA				
Precision	44	44	NA	< 25%				
Pesticide Reporting Level Spike	19	19	75 -125	NA				
Pesticide Matrix Spike	19	18	70 120	NIA				
Pesticide Matrix Spike Dup	19	18	70-130	NA				
Precision	19	19	NA	< 25%				
PCB/Pesticide Duplicate Analysis								
Duplicate 1 PCB	1	1		0-0/ 0 0 1/0/ 6				
Duplicate 1 Pesticides	3	2	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 1 Sum of Pesticides and PCBs	1	1		Campic Mean.				
Duplicate 2 PCB	1	1		. 050/ O 0 MDL 1				
Duplicate 2 Pesticides	3	2	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 2 Sum of Pesticides and PCBs	1	1		Campio Woan.				
	Sample Set – L	B (15 Liver Tiss	ue Samples)					
NRCC CARP-2	10	7	according to published acceptance criteria	NA				
SRM 1946	40	36	according to published acceptance criteria					
PCB Reporting Level Spike	44	44	75 -125	NA				
PCB Matrix Spike:	44	44	70 - 130	NA				
PCB Matrix Spike Dup	44	44	70 - 130	INA				
Precision	44	44	NA	< 25%				
Pesticide Reporting Level Spike	19	19	75 -125	NA				
Pesticide Matrix Spike	19	17	70-130	NA				
Pesticide Matrix Spike Dup	19	16	70-130	INA				
Precision	19	17	NA	< 25%				
PCB/Pesticide Duplicate Analysis								
Duplicate 1 PCB	2	1		20E0/ @ 2 :: MDL =f				
Duplicate 1 Pesticides	3	1	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 1 Sum of Pesticides and PCBs	1	1		Jampie Mean.				
Duplicate 2 PCB	3	0		4.050/ @ 2 MDL -5				
Duplicate 2 Pesticides	1	0	NA	< 25% @ 3 x MDL of Sample Mean.				
Duplicate 2 Sum of Pesticides and PCBs	1	0		25				

Table C-21 Continues.

Table C-21 Continued.

Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD						
Sample Set – LC (10 Liver Tissue Samples)										
NRCC CARP-2	10	9	according to published acceptance criteria	NA						
SRM 1946	40	338	according to published acceptance criteria							
PCB Reporting Level Spike	44	142	75 -125	NA						
PCB Matrix Spike:	44	843	70 - 130	NA						
PCB Matrix Spike Dup	44	44	70 - 130	INA						
Precision	44	44	NA	< 25%						
Pesticide Reporting Level Spike	19	19	75 -125	NA						
Pesticide Matrix Spike	19	17	70-130	NA						
Pesticide Matrix Spike Dup	19	18	70-130	INA						
Precision	19	19	NA	< 25%						
PCB/Pesticide Duplicate Analysis										
Duplicate 1 PCB	1	3		0.50/ 0.0 1/5/ 6						
Duplicate 1 Pesticides	4	3	NA	< 25% @ 3 x MDL of Sample Mean.						
Duplicate 1 Sum of Pesticides and PCBs	1	1		Campic Mean.						
Duplicate 2 PCB	9	8		. 050/ O 0 MDI (						
Duplicate 2 Pesticides	4	4	NA	< 25% @ 3 x MDL of Sample Mean.						
Duplicate 2 Sum of Pesticides and PCBs	1	1		Gampie Mean.						

CARP-2: National Research Council Canada; SRM 1946: National Institute of Standards & Technology

Table C-22. Fish tissue percent lipid QA/QC summary, July 2010–June 2011.

Sample Set	Tissue Type	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Precision % RPD
MB	Muscle	Percent Lipid	Duplicate Samples	2	2	<25%
MA2	Muscle	Percent Lipid	Duplicate Samples	2	2	<25%
MC2	Muscle	Percent Lipid	Duplicate Samples	2	2	<25%
MD2	Whole Body	Percent Lipid	Duplicate Samples	2	2	<25%
LA	Liver	Percent Lipid	Duplicate Samples	2	2	<25%
LB	Liver	Percent Lipid	Duplicate Samples	2	2	<25%
LC	Liver	Percent Lipid	Duplicate Samples	2	2	<25%

Table C-23. Method detection levels for mercury in fish tissue, July 2010–June 2011.

Orange County Sanitation District, California.

Parameter	Method Detection Limit (ng/g wet weight)
Mercury	0.002

Table C-24. Acceptance criteria for standard reference materials of mercury in fish tissue, July 2010–June 2011.

Orange County Sanitation District, California.

Mercury	True Value	Acceptance Range (ng/g)		
	(ng/g)	Minimum	Maximum	
DORM-2	4.64	4.38	4.90	
DORM-3	0.382	0.322	0.442	

Dogfish Muscle and Liver Reference Material for Mercury, National Research Council Canada.

#### Analytical Methods - Mercury

Fish tissue samples were analyzed for mercury in accordance with ELOM SOP 245.1A. Typical QC analyses for a tissue sample batch included a blank, a blank spike, and SRMs (liver and muscle). In the same batch, additional QC samples included duplicate analyses of the sample, spiked samples, and duplicate spiked samples, which were run approximately once every 10 samples.

The MDL for fish mercury is presented in Table C-23. Acceptance criteria for the mercury SRMs are presented in Table C-24. Fish tissue mercury QA/QC summary data are presented in Table C-25. All samples were analyzed within their 6-month holding times and met the QA criteria guidelines.

Pretreated (resected and 1:1 muscle:water homogenized) fish samples were analyzed

for mercury in accordance with methods described in the LOPM. QC for a typical batch included a blank, a blank spike, and an SRM (whole fish). Fish samples with duplicates, spiked samples, and duplicate spiked samples were run approximately once every 10 fish samples. When sample mercury concentration exceeded the appropriate calibration curve, the sample was diluted with the reagent blank and reanalyzed. The samples were analyzed for mercury on a Perkin Elmer FIMS 400 system.

All samples met the QA criteria guidelines for accuracy and precision with a few exceptions. Two sample duplicate RPDs were out of acceptable range due to low results and a non-homogeneous sample matrix. One sample duplicate spike recovery was out of its acceptable range due to a non-homogeneous sample matrix.

Table C-25. Fish tissue mercury QA/QC summary, July 2010–June 2011.

Sample Set	Parameter	Description	Number of Compounds Tested	Number of Compounds Passed	Target Accuracy % Recovery	Target Precision % RPD
		Blank	1	1	<2X MDL	NA
		Blank Spike	1	1	90-110	NA
		Matrix Spike	2	2	70-130	
HGFISH110228-1	Mercury	Matrix Spike Dup	2	2	70-130	
11011011110220 1	Wichdary	Matrix Spike Precision	2	2		< 25%
		Duplicate Analysis	2	**1	NA	@ <u>&gt;</u> 10 X MDL < 30%
		CRM Analysis	1	1	80-120% or certified value, whichever is greater.	
		Blank	1	1	<2X MDL	NA
		Blank Spike	1	1	90-110	NA
		Matrix Spike	3	3	70-130	
HGFISH110303-1	Mercury	Matrix Spike Dup	3	3	70-130	
	Wichouty	Matrix Spike Precision	3	3		< 25%
		Duplicate Analysis	3	3	NA	@ <u>&gt;</u> 10 X MDL < 30%
		CRM Analysis	1	1	80-120% or certified value, whichever is greater.	
		Blank	1	1	<2X MDL	NA
		Blank Spike	1	1	90-110	NA
		Matrix Spike	4	4	70-130	
HGFISH110307-1	Mercury	Matrix Spike Dup	4	**3	70-130	
1.51 15111 15507-1	Wichouty	Matrix Spike Precision	4	4		< 25%
		Duplicate Analysis	4	**3	NA	@ <u>&gt;</u> 10 X MDL < 30%
		CRM Analysis	1	1	80-120% or certified value, whichever is greater.	

<sup>\*\*</sup> Two duplicate analysis RPDs and one matrix spike duplicate recovery failed due to low results and non-homogeneous sample matrices. NA = Not applicable.

#### **BENTHIC INFAUNA NARRATIVE**

#### **SORTING AND TAXONOMY QA/QC**

The OCSD is in the final stages of insourcing the infaunal taxonomy program element. This work was done exclusively (Weston Solutions, contractors Inc., Carlsbad, CA) since about 1985. As part of the taxonomic training of OCSD staff, a series of re-identifications with contractors were conducted. New metrics were used to evaluate the results as part of the in-house taxonomy QA/QC development. These methods differ from the 2010-11 QAPP, which will be modified next year to reflect the new protocols. However, for the sorting QA/QC, the 2010-11 QAPP procedures were followed. The following sections describe QA/QC protocols used under the program and the status of samples that have received sorting and taxonomic QA/QC. Sorting QA/QC procedures have been completed for three surveys: summer (July 2010, Cruise # OC-2010-025), fall (October 2010, Cruise # OC-2009-040), and winter (January 2011, Cruise # OC-2011-001) surveys. Taxonomic identifications were conducted for the winter survey (January 2011).

# Sorting QA/QC Procedures

OCSD's NPDES permit designates 10 quarterly (summer, fall, winter, and spring) benthic-sampling stations and 39 annual (summer) benthic-sampling stations. Sorting procedures were performed on one replicate infaunal sample collected from each of three randomly selected quarterly stations in the summer, fall, and winter quarters and an additional seven samples (at least one from each of the four major depth contour intervals) for the annual survey; no QA samples were processed for the spring The sorting procedure involved removal by Weston Solutions, Inc. (Weston) personnel of all biological organisms and fragments from benthic samples. Organisms were further sorted by taxa, transferred to

separate vials, and total counts per station replicate were made. When all samples from a cruise passed Weston's in-house sorting efficiency criteria, they were shipped along with any remaining particulates (RPs), including sediments and shell and kelp fragments, to OCSD for reanalysis. OCSD re-sorted the sample RPs and collected any organisms or fragments that had been missed by Weston. The sample passed the QA procedure if the total number of animals collected by OCSD from the RPs was less than or equal to 5% of the total number of individuals collected by Weston for that sample.

# 2010-11 Sorting QA/QC Status

Sorting results for all 2010-11 QA samples were well within the 5% QC limit (99.5 % accuracy).

Taxonomic Identification QA/QC Procedures

Benthic infauna samples undergo comparative taxonomic analysis by two independent groups of taxonomists. selected infauna samples were first identified by OCSD taxonomists and re-identified by Weston then taxonomists from Weston. compared the two sets of data and reported discrepancies. Taxonomic discrepancies were reviewed and resolved by OCSD taxonomists. Following their review, any necessary corrections to taxon names or abundances were made and the project database modified to reflect these changes. The results were tallied by station and percent errors were calculated using the equations below:

Equation 1.  $%Error_{\#Taxa} = [(\#Taxa_{Resolved} - \#Taxa_{Original}) \div \#Taxa_{Resolved}] \cdot 100$ 

Equation 2. %Error  $_{\# Individuals} = (\# Individuals)$   $_{Resolved} - \# Individuals$   $_{Original}) \div \#$   $_{Individuals} = (\# Individuals)$  $_{Resolved} = (\# Individuals)$ 

Equation 3. %Error  $\#ID Taxa = (\# Taxa MissID \div \# Taxa Resolved) \cdot 100$ 

Equation 4. %Error  $\#_{ID\ Individuals} = (\#_{Individuals} = 100)$   $\#_{ID\ Individuals} = (\#_{Individuals} = 100)$ 

These equations were adapted from the Macrobenthic (Infaunal) Sample Analysis Laboratory Manual (SCCWRP 2008). In each equation the taxa or individuals "resolved" represents the final taxonomic determination or count following resolution by OCSD staff; taxa or individuals "original" represents the originating taxonomist's taxonomic determination or count; and taxa or individuals "mis-ID" represents the number of taxa or individuals that the originating taxonomist identified incorrectly.

When applied to individual taxonomists these equations are a measure of taxonomic accuracy (i.e., QA). The first three equations considered gauges of errors in accounting (e.g., recording on wrong line, miscounting, etc.), which by their random nature are hard to predict. Sample accuracy (i.e., QC) is calculated by station using the fourth equation reported herein. Equation 4 is the preferred measure of identification accuracy. It is weighted by abundance and has a more rigorous set of consequences (corrective actions) when errors are greater than 10%. Corrective actions include a reanalysis of additional samples for the effected taxa and additional, targeted, training. Equation 3, while included herein, technically was an assessment

identification accuracy but is considered too sensitive a measure for samples with low diversities, which are commonly found in samples of Echinodermata and Minor Phyla.

#### 2010-11 Taxonomic QA/QC Results

The QA/QC results are presented in Table C-26 and C-27. All stations met their QC objectives for percent error of number of identified individuals (Eq. 4) with a mean of 4.8%.

All samples were under the actionable threshold for QA measures. However, the relative high value at Station C of 9.2% error of identified taxa was due to the somewhat high number of Polychaete taxa misidentified (12.1% error of identified taxa) for all three stations. This, in turn, was caused by a few select problematic taxa present in these samples. In one example: the SvIlid Polychaete Syllides mikeli was misidentified as Syllides reishi. All effected records were corrected. In another case, the Lumbrinerid polychaete *Lumbrineris cruzensis* was, upon reanalysis, determined to be Lumbrineris latreilli. It was decided that due to the subjective nature of the diagnostic character used to separate the species, no further action was warranted and better in-house conventions are being developed.

Table C-26. Re-identification results for January 2011 QA samples.

Station	Rep	Description	Original Count	Mis-identified	Final Count
1	1 1	No. of Individuals	631	12	634
'		No. of Taxa	142	10	142
12	40 4	No. of Individuals	206	14	203
12	1	No. of Taxa	96	6	98
	1	No. of Individuals	422	24	430
С	I	No. of Taxa	140	13	141

Table C-27. Percent error rates calculated for January 2011 QA samples.

Error Typo	Station (rep)						
Error Type	1(1)	12(1)	C(1)	Mean			
%Error # Taxa	0.0	2.0	0.7	0.9			
%Error # Individuals	0.5	-1.5	1.9	0.3			
%Error # ID Taxa	7.0	6.1	9.2	7.5			
%Error # ID Individuals	1.9	6.9	5.6	4.8			

#### **OTTER TRAWL NARRATIVE**

The OCSD trawl sampling protocols are based upon regionally developed sampling methods (Mearns and Stubs 1974; Mearns and Allen 1978) and US Environmental Protection Agency 301(h) guidance documents (Tetra Tech 1986). These include a maximum distance from the nominal trawl station co-ordinates, sampling depth, vessel speed, and distance (trawl track) covered. Table C-28 lists the trawl quality assurance objectives (QAO).

Established regional survey methods for southern California requires that a portion of the trawl track must pass within a 100-m circle that originates from the nominal sample station position and be within 10% of the station's nominal depth. The speed of the trawl should range from 0.77 to 1.0 m/s or 1.5 to 2.0 kts. Since 1985, the District has trawled a set distance of 450 meters (the distance that the net is actually on the bottom collecting fish and invertebrates); regional surveys trawls are based on time on the bottom, not distance.

# Summer 2010

For summer 2010, trawl distances ranged from 408 to 472 m with the average trawl length being 455.4 m and the average trawl

speed being 1.9 kts for all trawls combined (Table C-29). All of the trawls passed through the designated 100-meter circle (Figure C-1). Trawl depths and time on the bottom were determined using an attached pressure sensor that showed excellent trawl repeatability in both depth (Table C-30) and distance traveled (Figure C-2). Station T3, which is located on the edge of the Newport submarine canyon where depth changes rapidly, was the only anomalous station (Figure 6-1). A perfectly flat trawl along an isobath is difficult to maintain at this station. While Station T3 appears not to follow the bottom depth contour, OCSD staff believe that the net is trawling properly along an irregular bottom.

#### Winter 2011

For winter 2011, all trawl lengths ranged from 449 to 547 m with the average trawl length being 464.4 m and the average trawl speed being 2.1 kts for all trawls combined (Table C-31). All the trawls passed through the designated 100-meter circle (Figure C-3). Trawl depths and time on the bottom were determined using an attached pressure sensor that showed excellent trawl repeatability in both depth (Table C-32) and distance traveled (Figure C-4). Station T3 was again the only anomalous station.

Table C-28. Districts quality assurance objectives for trawl sampling, July 2010–June 2011.

Measure	Quality Assurance Objective (QAO)
Trawl Track Depth	±10% of nominal station depth (at any point during the trawl)
Trawl Track Length	450 m
Distance from nominal	100 m
Vessel Speed	1.5–2.0 knots

Table C-29. Trawl sample dates, track distances, percent difference from target track distance, elapsed time, and vessel speed, August 2010.

Date	Station	Haul	Distance Trawled (meters)	Percent Difference from Target Distance *	Elapsed Time (seconds)	Trawl speed (knots)**
August 18, 2010	T1	1	470.2	4.5	538	1.7
August 18, 2010	T1	2	448.0	-0.4	542	1.6
August 18, 2010	T1	3	450.1	0.0	551	1.6
August 17, 2010	T2	1	454.7	1.0	532	1.7
August 17, 2010	T2	2	461.5	2.6	556	1.6
August 17, 2010	T3	1	408.0	-9.3	382	2.1
August 18, 2010	Т3	2	452.8	0.6	465	1.9
August 18, 2010	Т3	3	455.8	1.3	471	1.9
August 17, 2010	T6	1	451.0	0.2	642	1.4
August 24, 2010	T6	2	455.5	1.2	408	2.2
August 17, 2010	T10	1	472.0	4.9	360	2.5
August 17, 2010	T10	2	455.3	1.2	435	2.0
August 24, 2010	T11	1	460.0	2.2	472	1.9
August 25, 2010	T11	2	463.6	3.0	497	1.8
August 25, 2010	T11	3	457.5	1.7	475	1.9
August 23, 2010	T12	1	463.6	3.0	401	2.2
August 23, 2010	T12	2	458.2	1.8	432	2.1
August 23, 2010	T12	3	458.2	1.8	397	2.2
August 23, 2010	T13	1	453.8	0.8	433	2.0
August 24, 2010	T13	2	453.1	0.7	418	2.1
August 24, 2010	T13	3	453.8	0.8	448	2.0
August 23, 2010	T14	1	453.5	0.8	470	1.9
August 25, 2010	T14	2	463.9	3.1	524	1.7
	Mea	an value	455.4	1.2	472	1.9

Hauls with speeds less than 1.5 knots or greater than 2 knots are denoted in bold.

<sup>\*</sup> Target Distance – 450 meters

<sup>\*\*</sup> Target Speed - 1.5 - 2.0 knot

Table C-30. Ten percent trawl depth QA, August 2010.

Date	Station	Haul	Nominal Depth (m)	QA Range (m)	Data Source	Average Bottom Depth (m)	10% Y/N
August 18, 2010	T1	1			SBE data	55.8	Υ
August 16, 2010	11	ı			SOD data	54.5	Υ
August 18, 2010	T1	2	55	49.5–60.5	SBE data	56.6	Υ
August 10, 2010	1 1		33	49.5-00.5	SOD data	54.0	Υ
August 18, 2010	T1	3			SBE data	56.8	Υ
August 10, 2010	11	3			SOD data	54.5	Υ
August 17, 2010	T2	1			SBE data	36.5	Υ
August 17, 2010	12	1	35	31.5–38.5	SOD data	35.0	Υ
August 17, 2010	T2	2	33	31.5–36.5	SBE data	36.5	Υ
August 17, 2010	12				SOD data	35.0	Υ
August 17, 2010	T3	1			SBE data	67.7	N
August 17, 2010	13	1			SOD data	59.0	Υ
August 18, 2010	Т3	2	55	49.5–60.5	SBE data	74.7	N
August 16, 2010	13		33		SOD data	67.0	N
August 19, 2010	Т3	3			SBE data	69.8	N
August 18, 2010	13	3			SOD data	64.0	N
August 17, 2010	Т6	1			SBE data	37.8	Υ
August 17, 2010	10	1	36	32.4–39.6	SOD data	36.5	Υ
August 24, 2010	Т6	2	30	32.4-39.0	SBE data	37.4	Υ
August 24, 2010	10				SOD data	36.5	Υ
August 17, 2010	T10	1			SBE data	136.8	Υ
August 17, 2010	110	!	137	123.3–150.7	SOD data	133.0	Υ
August 17, 2010	T10	2	137	123.3-150.7	SBE data	143.6	Υ
August 17, 2010	110				SOD data	133.5	Υ
August 24, 2010	T11	1			SBE data	60.2	Υ
August 24, 2010	111	!			SOD data	61.5	Υ
August 25, 2010	T11	2	60	54.0–66.0	SBE data	54.6	Υ
August 25, 2010	111	2	00	54.0-00.0	SOD data	59.0	Υ
August 25, 2010	T11	3			SBE data	62.6	Υ
August 25, 2010	111	J			SOD data	62.0	Υ
August 23, 2010	T12	1			SBE data	58.1	Υ
August 23, 2010	114	'			SOD data	57.5	Υ
August 23, 2010	August 23, 2010 T12	T12 2	60	54.0–66.0	SBE data	59.5	Υ
August 23, 2010	114	2	00	J <del>4</del> .0—00.0	SOD data	56.5	Υ
August 23, 2010	T12	3			SBE data	58.2	Υ
August 25, 2010	1 12	,			SOD data	56.5	Υ

Table C-30 Continues.

Table C-30 Continued.

Date	Station	Haul	Nominal Depth (m)	QA Range (m)	Data Source	Average Bottom Depth (m)	10% Y/N		
August 23, 2010	T13	1			SBE data	62.9	Υ		
August 23, 2010	113	I			SOD data	59.5	Υ		
August 24, 2010	T13	2	60	54.0–66.0	SBE data	62.7	Υ		
August 24, 2010	113				SOD data	61.0	Υ		
A	T13	3			SBE data	65.2	Υ		
August 24, 2010		3			SOD data	60.0	Υ		
August 22, 2010	T14	T4.4	T4.4	1			SBE data	139.4	Y
August 23, 2010		'	137	123.3–150.7	SOD data	140.0	Υ		
A	T14	2			SBE data	144.3	Υ		
August 25, 2010	T14				SOD data	138.0	Υ		

# Notes:

Station T3 depth varies widely. 10% QA may not be applicable.

SBE = Seabird Electronics

SOD = Station occupation data

Y = Yes (Pass)

N = No (Fail)

N/A = Not analyzed

Table C-31. Trawl sample dates, track distances, percent difference from target track distance, elapsed time, and vessel speed, January 2011.

Date	Station	Haul	Distance Trawled (meters)	Percent Difference from Target Distance *	Elapsed Time (seconds)	Trawl speed (knots)**
January 18, 2011	T0	1	463.0	2.9	427	2.1
January 19, 2011	T1	1	464.8	3.3	448	2.0
January 19, 2011	T1	2	467.6	3.9	438	2.1
January 19, 2011	T1	3	546.9	21.5	505	2.1
January 11, 2011	T2	1	461.1	2.5	459	2.0
January 12, 2011	T2	2	459.4	2.1	381	2.3
January 11, 2011	T3	1	454.8	1.1	403	2.2
January 11, 2011	Т3	2	462.7	2.8	430	2.1
January 12, 2011	Т3	3	460.1	2.2	418	2.1
January 12, 2011	T6	1	455.8	1.3	426	2.1
January 12, 2011	T6	2	453.2	0.7	442	2.0
January 11, 2011	T10	1	472.6	5.0	447	2.1
January 11, 2011	T10	2	459.0	2.0	436	2.0
January 18, 2011	T11	1	458.1	1.8	440	2.0
January 19, 2011	T11	2	461.8	2.6	431	2.1
January 19, 2011	T11	3	458.3	1.8	443	2.0
January 18, 2011	T12	1	463.7	3.0	419	2.2
January 18, 2011	T12	2	463.5	3.0	434	2.1
January 18, 2011	T12	3	456.5	1.5	442	2.0
January 12, 2011	T13	1	449.1	-0.2	462	1.9
January 18, 2011	T13	2	461.1	2.5	435	2.1
January 18, 2011	T13	3	471.3	4.7	444	2.1
January 12, 2011	T14	1	467.4	3.9	431	2.1
January 12, 2011	T14	2	453.7	0.8	435	2.0
	Mea	n value	464.4	3.2	437	2.1

Hauls with speeds less than 1.5 knots or greater than 2 knots are denoted in bold.

<sup>\*</sup> Target Distance – 450 meters

<sup>\*\*</sup> Target Speed – 1.5 – 2.0 knots

Table C-32. Ten percent trawl depth QA, January 2011.

Date	Station	Haul	Nominal Depth (m)	QA Range (m)	Data Source	Average Bottom Depth (m)	10% Y/N
January 18, 2011	T0	1	18	16.2–19.8	SBE data	NA	NA
					SOD data	NA	NA
January 19, 2011 January 19, 2011	T1	1	55	49.5–60.5	SBE data	57.6	Υ
					SOD data	55.5	Υ
	T1	2			SBE data	57.3	Υ
					SOD data	55.5	Y
January 19, 2011	T1	3			SBE data	56.8	Υ
		Ŭ			SOD data	55.0	Υ
January 11, 2011	T2	1	35	31.5–38.5	SBE data	35.7	Υ
					SOD data	34.5	Υ
January 11, 2011	T2	2			SBE data	35.5	Υ
	, _				SOD data	34.5	Υ
January 11, 2011	Т3	1		49.5–60.5	SBE data	60.3	Υ
	10	•			SOD data	56.5	Υ
January 11, 2011	Т3	2	55		SBE data	61.2	N
					SOD data	58.0	Y
January 12, 2011	Т3	3			SBE data	67.4	N
					SOD data	63.0	N
January 12, 2011	Т6	1	36	32.4–39.6	SBE data	37.1	Υ
					SOD data	36.0	Υ
January 12, 2011	Т6	3			SBE data	37.3	Υ
					SOD data	36.0	Υ
January 11, 2011	T10	1	137	123.3–150.7	SBE data	146.2	Y
					SOD data	133.0	Υ
January 11, 2011	T10	2			SBE data	137.0	Υ
					SOD data	136.5	Υ
January 18, 2011	T11	1	60	54.0–66.0	SBE data	65.5	Υ
					SOD data	58.5	Υ
January 19, 2011	T11	2			SBE data	61.2	Υ
					SOD data	53.5	N
January 19, 2011	T11	3			SBE data	59.9	Y
					SOD data	62.0	Υ
January 18, 2011	T12	1	60	54.0–66.0	SBE data	58.9	Υ
					SOD data	57.5	Y
January 18, 2011	T12	2			SBE data	59.0	Υ
					SOD data	57.0	Υ
January 18, 2011	T12	3			SBE data	59.0	Υ
					SOD data	55.5	Υ

Table C-32 Continues.

Table C-32 Continued.

Date	Station	Haul	Nominal Depth (m)	QA Range (m)	Data Source	Average Bottom Depth (m)	10% Y/N
January 12, 2011	T13	1	60	54.0–66.0	SBE data	61.9	Υ
					SOD data	57.5	Υ
January 18, 2011	T13	2			SBE data	61.6	Υ
					SOD data	55.0	Υ
January 18, 2011	T13	3			SBE data	62.8	Υ
					SOD data	58.5	Y
January 12, 2011	T14	1	137	123.3–150.7	SBE data	141.6	Υ
					SOD data	140.5	Υ
January 12, 2011	T14	2			SBE data	142.3	Υ
					SOD data	136.0	Υ

# Notes:

Station T3 depth varies widely. 10% QA may not be applicable.

SBE = Seabird Electronics

SOD = Station occupation data

Y = Yes (Pass)

N = No (Fail)

NA = Not analyzed

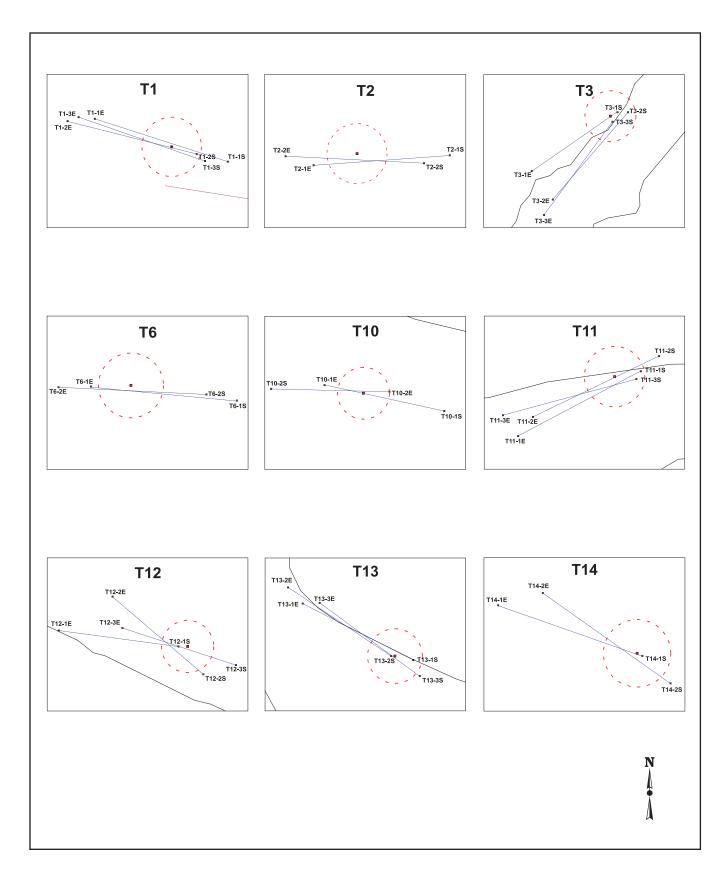


Figure C-1. Quality assurance plots of distance to station for otter trawl hauls, August 2010.

Red circle represents 100 meter distance from nominal trawl station center point. Blue lines represent trawl path while net is on the bottom. Trawl endpoints are labeled by station name, haul number, start (S) and end (E).

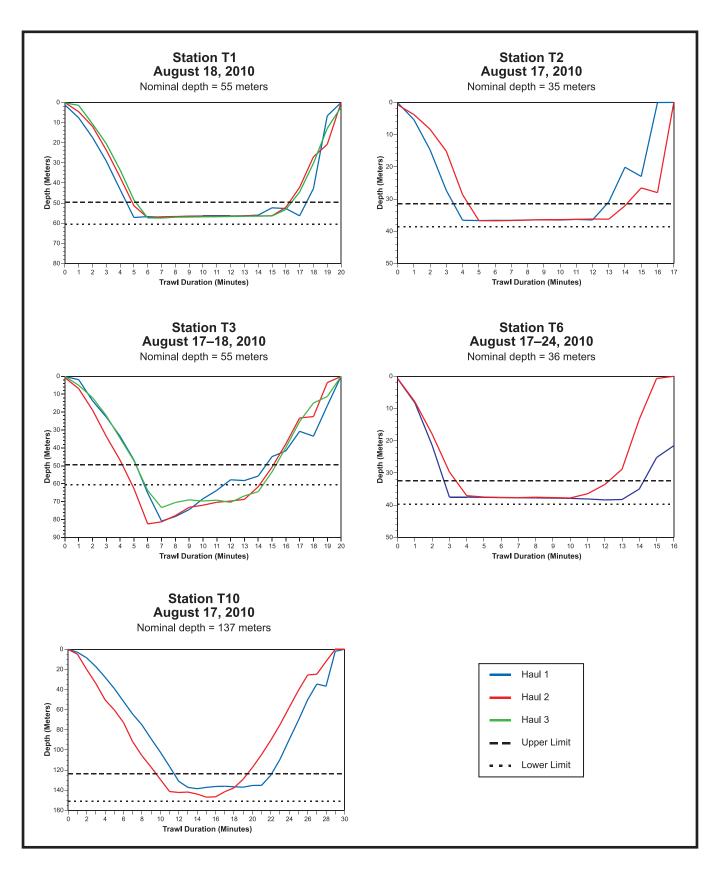


Figure C-2. Quality assurance plots of trawl depth and trawl duration per haul for otter trawl stations, August 2010.

Upper and lower limit lines are ± 10% of nominal trawl depth.

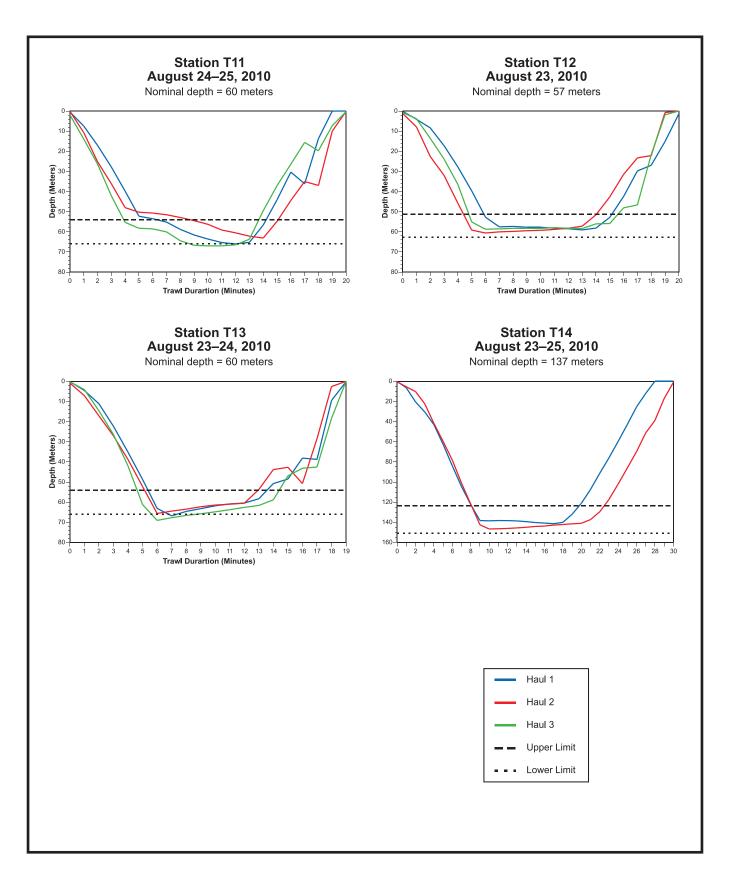


Figure C-2 Continued.

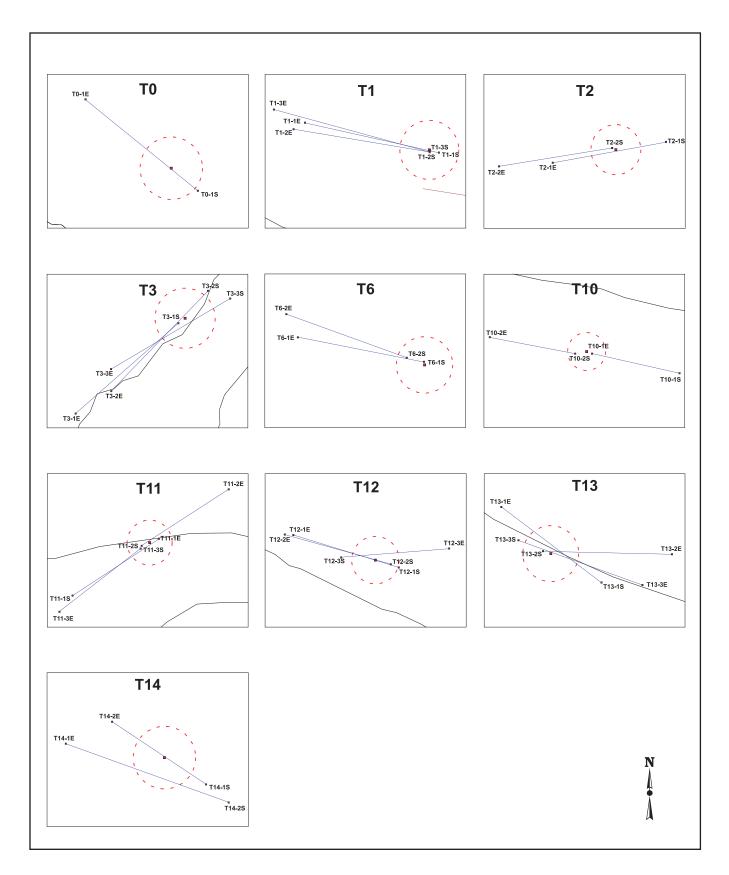


Figure C-3. Quality assurance plots of distance to station for otter trawl hauls, January 2011.

Red circle represents 100 meter distance from nominal trawl station center point. Blue lines represent trawl path while net is on the bottom. Trawl endpoints are labeled by station name, haul number, start (S) and end (E).

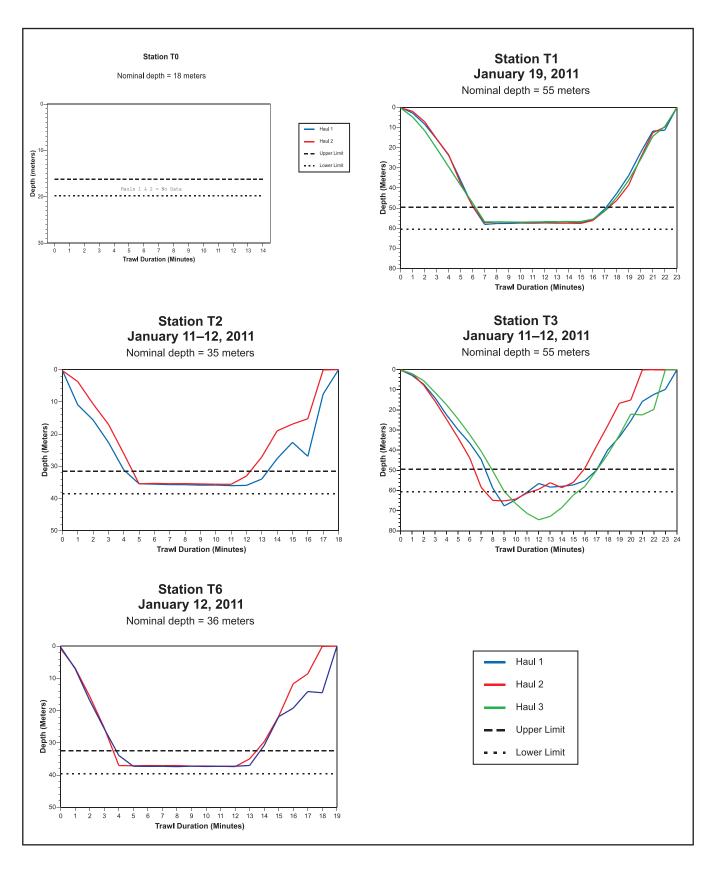


Figure C-4. Quality assurance plots of trawl depth and trawl duration per haul for otter trawl stations, January 2011.

Upper and lower limit lines are ± 10% of nominal trawl depth.

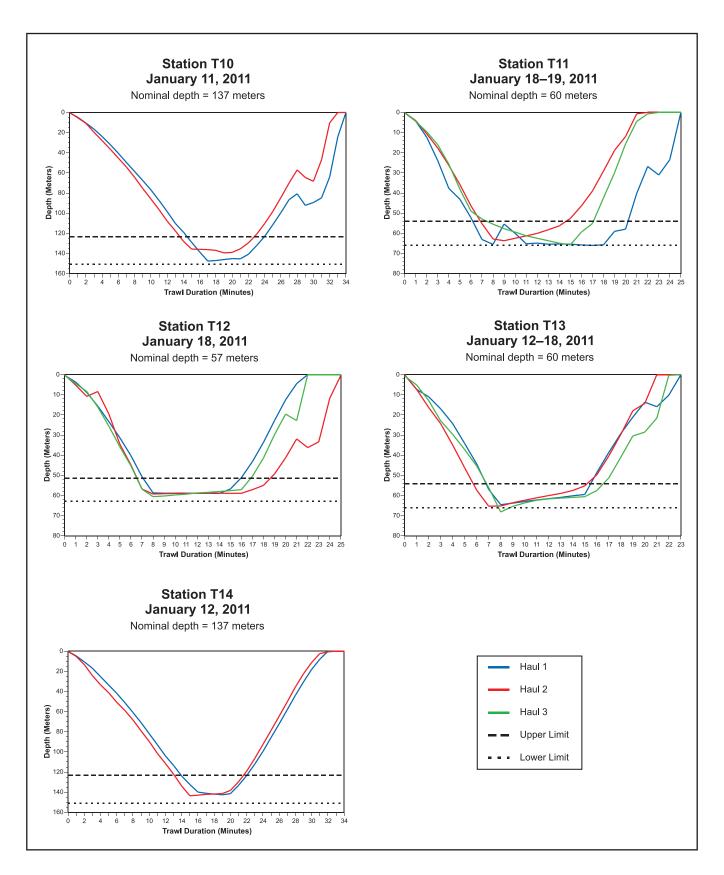


Figure C-4 Continued.

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