

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NAB-BAY9-SDB6-PRE	100	a	10	100.0	100.0	0.0	107.1	n/a	No
		b	10	100.0					
		c	10	100.0					
NAB-BAY18-SDB6-PRE	100	a	10	100.0	100.0	0.0	107.1	n/a	No
		b	10	100.0					
		c	10	100.0					
NAB-BAY9-SDB6-DUR	100	a	10	100.0	100.0	0.0	107.1	n/a	No
		b	10	100.0					
		c	10	100.0					
NAB-BAY18-SDB6-DUR	100	a	10	100.0	96.7	5.8	103.6	0.343	No
		b	10	100.0					
		c	9	90.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NAB-BAY9-SDB6-PRE	100	a	215	10	95.6	96.4	1.2	98.0	0.0470	No
		b	197	7	96.6					
		c	191	7	96.5					
		d	219	11	95.2					
		e	219	4	98.2					
NAB-BAY18-SDB6-PRE	100	a	164	9	94.8	97.3	1.5	98.9	0.3135	No
		b	218	7	96.9					
		c	213	4	98.2					
		d	243	4	98.4					
		e	216	4	98.2					
NAB-BAY9-SDB6-DUR	100	a	213	4	98.2	97.7	1.3	100.0	0.0670	Yes ^c
		b	208	6	97.2					
		c	200	2	99.0					
		d	207	3	98.6					
		e	221	10	95.7					
NAB-BAY18-SDB6-DUR	100	a	212	13	94.2	95.4	1.5	97.6	0.0384	Yes
		b	188	5	97.4					
		c	187	12	94.0					
		d	245	13	95.0					
		e	209	8	96.3					

QA/QC SAMPLES^a
TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Salt Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Copper Ref. Tox.	50	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	70.0	47.6	70.0	0.148	No
		b	0	0.0					
		c	5	100.0					
		d	4	80.0					
	200	a	1	20.0	10.0	11.5	10.0	0.000	Yes
		b	0	0.0					
		c	0	0.0					
		d	1	20.0					
	400	a	0	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	0.0					
		c	0	0.0					
		d	0	0.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	8	80.0	93.3	11.5	100.0	n/a	n/a
		b	10	100.0					
		c	11	100.0					
Salt Control	n/a	a	10	100.0	100.0	0.0	107.1	0.211	No
		b	10	100.0					
		c	10	100.0					
Copper Ref. Tox.	100	a	10	100.0	100.0	0.0	107.1	0.371	No
		b	10	100.0					
		c	10	100.0					
	200	a	10	100.0	96.7	5.8	103.6	0.500	No
		b	10	100.0					
		c	9	90.0					
	400	a	3	30.0	33.3	5.8	35.7	0.005	Yes
		b	3	30.0					
		c	4	40.0					
	800	a	0	0.0	0.0	0.0	0.0	0.004	Yes
		b	0	0.0					
		c	0	0.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	160	4	97.6	97.7	1.0	100.0	n/a	No
		b	222	4	98.2					
		c	236	6	97.5					
		d	233	9	96.3					
		e	257	3	98.8					
Brine Control 1	n/a	a	204	3	98.6	98.4	0.8	100.7	0.119	No
		b	211	5	97.7					
		c	201	5	97.6					
		d	226	1	99.6					
		e	221	3	98.7					
Brine Control 2	n/a	a	189	3	98.4	97.8	1.1	100.1	0.440	No
		b	231	10	95.9					
		c	210	4	98.1					
		d	190	4	97.9					
		e	210	3	98.6					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Copper Ref. Tox.	2.9	a	231	5	97.9	98.6	0.7	101.0	0.057	No
		b	207	4	98.1					
		c	214	1	99.5					
		d	201	3	98.5					
		e	228	2	99.1					
	4.1	a	214	8	96.4	56.4	39.6	57.7	0.040	Yes
		b	205	21	90.7					
		c	-	-	-					
		d	-	-	-					
		e	-	-	-					
	5.9	a	125	101	55.3	49.3	10.8	50.5	0.000	Yes
		b	125	94	57.1					
		c	132	106	55.5					
		d	114	125	47.7					
		e	64	142	31.1					
	8.4	a	23	187	11.0	10.1	5.2	10.3	0.000	Yes
		b	24	173	12.2					
		c	4	210	1.9					
		d	32	170	15.8					
		e	21	200	9.5					
	12.0	a	0	195	0.0	0.3	0.2	0.3	0.000	Yes
		b	1	246	0.4					
		c	1	221	0.5					
		d	1	218	0.5					
		e	0	219	0.0					
	17.2	a	0	210	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	187	0.0					
		c	0	178	0.0					
		d	0	215	0.0					
		e	0	198	0.0					

SUMMARY RESULTS- QA/QC
COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	100	200	123.5	103.3-147.5
MYSIDS	200	400	352.5	326.3-387.7
MUSSELS	4.1	5.9	6.0	5.9-6.1

Dash indicates no data (replicate was spilled or organisms not added)

^aControls (QA/QC) correspond to all samples from SDB6

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control
n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (*A. affinis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)										
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96					
NAB-OF9-SDB6-FF	12.5%	a	7.6	7.7	7.7	7.6	7.6	7.0	7.2	6.6	6.4	6.3	18.8	18.8	19.6	19.6	20.2	32.9	32.8	32.6	33.3	33.4
	25%	a	7.6	7.7	7.7	7.6	7.6	7.3	6.7	6.5	6.3	6.7	18.9	18.7	19.6	19.4	20.0	33.0	33.0	33.1	33.6	34.0
	50%	a	7.6	7.6	7.7	7.6	7.6	7.3	6.7	6.3	5.7	6.2	19.4	18.8	19.6	19.6	20.2	32.8	32.9	33.0	33.4	33.4
	100%	a	7.8	7.7	7.7	7.6	7.6	7.4	6.0	5.9	5.3	5.3	19.1	18.8	19.6	19.6	20.2	33.3	33.2	33.1	33.6	33.6
NAB-OF9-SDB6-COMP	12.5%	a	7.6	7.7	7.7	7.6	7.6	7.0	7.3	6.8	6.9	6.7	18.7	18.3	19.0	19.6	19.5	33.0	33.2	34.1	32.8	33.5
	25%	a	7.6	7.7	7.6	7.6	7.6	7.0	7.3	6.8	7.1	6.8	18.6	18.3	19.0	19.6	19.5	33.0	33.2	33.1	32.8	33.5
	50%	a	7.7	7.7	7.6	7.6	7.6	7.3	7.1	6.8	6.8	6.9	18.7	18.3	19.2	19.4	19.5	32.6	32.7	32.8	32.9	33.4
	100%	a	7.8	7.7	7.7	7.6	7.6	7.6	6.9	6.5	6.5	6.7	19.0	18.3	19.4	19.3	19.5	32.4	32.8	32.9	32.9	33.1
NAB-OF18-SDB6-FF	12.5%	a	7.6	7.6	7.6	7.5	7.1	7.0	6.5	6.8	6.6	6.6	18.8	18.0	19.4	19.4	19.8	32.9	32.9	32.9	33.1	33.7
	50%	a	7.6	7.6	7.6	7.5	7.1	6.6	6.1	5.7	6.1	6.1	18.8	18.6	19.4	19.6	20.0	32.8	32.8	32.9	33.1	33.2
	100%	a	7.7	7.7	7.7	7.5	7.5	7.9	6.0	5.5	5.3	5.9	18.9	18.6	19.4	19.5	19.9	32.7	32.6	32.9	33.0	33.1
		a	7.6	7.7	7.7	7.6	7.6	6.8	7.4	6.4	7.2	6.7	18.9	18.8	19.6	19.4	20.0	33.0	33.2	33.0	34.0	34.3
NAB-BAY18-SDB6-PRE	100%	a	7.6	7.6	7.7	7.6	7.6	6.9	7.5	6.7	7.4	7.0	18.6	18.2	19.0	19.6	19.4	32.7	32.8	32.9	33.6	34.1
NAB-BAY9-SDB6-DUR	100%	a	7.6	7.6	7.7	7.6	7.6	7.6	7.2	6.7	7.1	7.0	19.9	18.3	19.0	19.5	19.5	31.7	32.0	32.0	31.5	32.0
NAB-BAY18-SDB6-DUR	100%	a	7.6	7.6	7.6	7.6	7.6	7.6	7.2	6.4	7.0	6.6	19.6	18.7	19.5	19.3	19.9	33.1	33.2	33.0	33.2	33.8
Scripps Control	0	a	7.6	7.7	7.7	7.6	7.6	7.8	7.7	7.2	7.4	7.2	18.8	18.2	19.0	19.3	19.1	32.0	32.6	33.5	32.1	33.5
Scripps Cu Ref. Tox	50 µg/l	a	7.6	7.7	7.7	7.6	7.6	6.8	7.3	7.2	7.4	7.1	18.5	18.7	19.3	19.3	19.4	32.2	32.9	33.1	31.7	32.4
	100 µg/l	a	7.6	7.7	7.6	7.6	7.6	6.9	7.7	7.2	6.9	7.1	18.6	18.5	19.0	19.6	19.5	32.5	32.9	33.1	31.9	32.4
	200 µg/l	a	7.6	7.7	7.6	7.7	7.7	6.9	7.8	7.2	7.9	7.0	18.6	18.1	19.3	19.4	19.4	32.1	32.9	33.0	32.5	32.6
	400 µg/l	a	7.6	7.7	7.6	7.6	7.6	7.1	6.9	7.9	7.1	7.1	18.6	18.1	19.1	19.1	19.1	32.1	32.9	33.0	32.1	32.7
Salt Control	n/a	a	7.6	7.7	7.4	7.3	7.3	5.9	7.1	6.9	7.2	6.9	18.8	18.3	19.0	19.4	19.5	31.6	32.7	33.1	32.0	32.7

MYSIDS (*A. bahia*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)					Dissolved Oxygen (mg/l)					Temperature (°C)					Salinity (‰)				
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96	0	24	48	72	96
NAB-OF9-SDB6-FF	12.5%	a	7.5	7.6	7.5	7.5	7.5	7.0	7.5	5.1	5.5	5.2	18.9	18.5	19.4	19.5	19.4	32.9	32.9	32.6	33.2	33.3
	25%	a	7.6	7.6	7.6	7.5	7.4	7.3	5.1	5.8	5.7	19.0	18.5	19.5	19.4	19.2	32.9	32.9	32.9	33.5	33.5	
	50%	a	7.7	7.6	7.6	7.5	7.4	6.3	5.0	5.3	5.5	19.0	18.6	19.5	19.4	19.3	33.1	32.7	32.6	33.4	33.3	
	100%	a	7.8	7.6	7.6	7.6	7.6	7.7	5.9	4.9	5.0	4.8	19.3	18.6	19.4	19.5	19.3	32.9	33.3	33.3	33.5	33.6
NAB-OF9-SDB6-COMP	12.5%	a	7.7	7.7	7.6	7.5	7.6	7.1	7.0	5.4	6.3	6.1	19.2	18.7	19.5	19.5	19.4	32.4	32.8	32.7	32.9	33.2
	25%	a	7.8	7.7	7.6	7.6	7.6	7.3	6.5	5.2	6.1	6.1	18.8	18.8	19.5	19.5	19.4	32.8	33.0	32.9	32.9	33.3
	50%	a	7.8	7.7	7.6	7.6	7.6	7.4	6.3	5.0	6.0	6.1	19.3	18.8	19.5	19.5	19.3	32.4	32.5	32.6	33.4	33.1
	100%	a	7.9	7.7	7.6	7.6	7.6	7.6	5.3	4.6	5.4	5.5	19.9	18.8	19.5	19.6	19.4	32.3	32.3	32.4	32.5	32.6
NAB-OF18-SDB6-FF	12.5%	a	7.6	7.7	7.5	7.5	7.6	7.3	7.2	5.1	5.0	5.5	18.8	18.6	19.6	19.5	19.3	32.7	33.0	32.9	33.1	33.2
	50%	a	7.7	7.6	7.5	7.4	7.5	7.5	5.6	5.0	4.6	4.6	18.8	18.6	19.6	19.5	19.3	32.6	32.8	32.8	33.1	33.1
	100%	a	7.8	7.6	7.5	7.5	7.5	8.0	5.3	4.5	4.3	4.3	19.1	18.6	19.6	19.5	19.3	32.5	32.8	32.5	33.0	33.0
	100%	a	7.7	7.6	7.5	7.5	7.5	7.3	7.0	5.0	5.9	6.7	19.1	18.6	19.6	19.6	19.4	32.6	33.0	32.6	33.4	34.0
NAB-BAY9-SDB6-PRE	100%	a	7.5	7.7	7.5	7.5	7.5	7.3	7.0	5.0	5.9	6.7	19.1	18.6	19.6	19.6	19.4	31.7	33.3	33.0	32.9	33.4
NAB-BAY18-SDB6-PRE	100%	a	7.7	7.7	7.6	7.6	7.6	6.8	7.2	5.5	6.4	5.6	18.8	18.6	19.7	19.4	19.6	31.7	33.3	33.0	32.9	33.4
NAB-BAY9-SDB6-DUR	100%	a	7.5	7.6	7.5	7.5	ND	7.4	7.1	5.0	6.1	ND	18.9	18.6	19.6	19.5	ND	30.7	30.5	30.7	30.3	ND
NAB-BAY18-SDB6-DUR	100%	a	7.5	7.6	7.5	7.6	7.6	7.8	6.8	5.4	6.8	6.4	20.0	18.6	19.7	19.5	19.4	31.3	31.8	32.7	32.8	32.4
Scripps Control	0	a	7.7	7.7	7.6	7.6	7.6	6.8	7.3	5.4	6.1	6.4	19.3	18.8	19.8	19.8	18.9	32.1	32.1	32.3	31.9	32.1
Scripps Cu Ref. Tox	100 µg/l	a	7.6	7.6	7.5	7.6	7.5	6.9	7.2	5.1	6.1	5.8	19.3	18.8	19.7	19.8	19.6	32.2	32.2	32.6	32.1	32.2
	200 µg/l	a	7.6	7.6	7.6	7.6	7.6	7.0	7.2	5.0	6.5	6.4	19.1	18.7	19.7	19.8	19.5	32.4	32.3	32.6	32.1	32.1
	400 µg/l	a	7.6	7.7	7.6	7.6	7.6	7.0	7.3	5.4	7.0	6.9	19.0	18.8	19.7	19.6	19.4	32.1	32.1	32.6	32.2	32.2
	800 µg/l	a	7.6	7.6	7.6	7.2	7.7	7.1	7.6	6.0	6.2	6.9	19.0	18.8	19.6	19.8	19.4	32.3	32.0	32.3	32.2	32.2
Salt Control	n/a	a	7.6	7.4	7.3	7.7	7.2	5.8	6.6	4.7	7.6	6.4	19.3	18.8	19.8	19.6	19.6	31.8	31.9	32.2	32.2	32.7

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)		Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)			
			0	24	48	0	24	48	0	24	48	0	24	48
NAB-OF9-SDB6-FF	6.2%	f	7.7	7.8	7.8	8.1	7.1	7.1	15.1	15.7	15.3	32.7	32.2	32.6
	12.4%	f	7.8	7.8	7.8	8.0	7.1	7.4	15.2	15.7	15.3	32.4	32.1	32.0
	24.8%	f	7.8	7.8	7.8	8.1	7.2	7.4	15.0	15.7	15.3	31.5	31.8	32.0
	49.5%	f	7.9	7.8	7.8	8.0	7.2	7.3	15.1	15.7	15.3	32.3	31.9	31.5
NAB-OF9-SDB6-COMP	6.2%	f	7.8	7.7	7.8	8.0	7.0	7.0	15.2	15.6	15.6	32.9	32.2	32.2
	12.4%	f	7.8	7.7	7.8	7.4	7.1	7.0	15.2	15.8	15.6	32.6	32.3	32.3
	24.8%	f	7.8	7.7	7.8	7.7	7.1	7.0	15.1	15.8	15.4	32.8	32.4	32.4
	49.5%	f	7.8	7.8	7.8	7.8	7.2	7.0	15.0	15.5	15.4	32.2	32.0	32.0
NAB-OF18-SDB6-FF	6.4%	f	7.8	7.8	7.8	8.1	7.2	7.6	16.2	15.8	15.5	32.2	31.4	32.1
	12.9%	f	7.8	7.8	7.8	8.2	7.1	7.7	16.2	15.9	15.6	32.0	31.1	32.1
	25.7%	f	7.8	7.8	7.8	8.1	7.1	7.6	16.2	15.8	15.5	31.7	31.0	31.6
	51.4%	f	7.9	7.7	7.8	8.1	7.0	7.4	16.0	16.0	15.4	31.1	30.8	30.8
NAB-BAY9-SDB6-PRE	100%	f	7.8	7.7	7.8	7.9	7.1	7.3	15.2	15.6	15.3	32.5	32.2	32.4
NAB-BAY18-SDB6-PRE	100%	f	7.7	7.7	7.8	8.1	7.1	7.0	15.1	15.7	15.5	32.5	32.0	32.1
NAB-BAY9-SDB6-DUR	100%	f	7.8	7.8	7.8	8.1	7.0	7.1	16.3	15.7	15.6	29.8	29.7	30.5
NAB-BAY18-SDB6-DUR	100%	f	7.9	7.8	7.8	7.8	6.8	7.2	16.2	16.0	15.6	32.1	32.0	32.4
Scripps Control	0	f	7.9	7.7	7.7	7.5	6.8	7.6	15.1	15.8	15.3	28.2	28.0	27.8
Scripps Cu Ref. Tox.	2.9 µg/l	f	7.9	7.7	7.8	7.7	6.9	7.7	15.0	15.7	15.4	29.0	28.5	28.4
	4.1 µg/l	f	7.9	7.8	7.8	7.8	6.9	7.6	15.1	15.8	15.3	28.3	28.1	28.0
	5.9 µg/l	f	7.9	7.8	7.8	8.1	6.8	7.8	15.0	15.8	15.3	27.6	27.9	28.4
	8.4 µg/l	f	7.9	7.7	7.7	7.9	6.9	7.8	15.0	15.9	15.4	28.0	28.0	28.2
	12 µg/l	f	7.9	7.8	7.7	8.0	7.0	7.5	15.3	16.0	15.3	28.3	28.1	28.3
	17.2 µg/l	f	7.9	7.8	7.7	8.0	7.2	7.2	15.0	15.8	15.3	28.0	28.1	28.4
Brine Control 1	n/a	f	8.0	7.8	7.8	7.4	7.2	7.3	16.3	16.0	15.8	32.6	32.3	32.1
Brine Control 2	n/a	f	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND

N - water quality not taken due to 100% mortality in treatment

ND - water quality not recorded

OUTFALLS**TOPSMELT (*A. affinis*)**

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NAB-OF9-TIE2-FF	25	95.0
	50	100.0
	100	95.0
NAB-OF18-TIE2-FF	25	95.0
	50	15.0
	100	0.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NAB-OF9-TIE2-FF	25	95.0
	50	90.0
	100	50.0
NAB-OF18-TIE2-FF	25	95.0
	50	20.0
	100	5.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	MEAN NORM DEV (%)
NAB-OF9-TIE2-FF	12.5	57.0
	25	0.0
	50	0.0
	61	0.0
NAB-OF18-TIE2-FF	12.5	81.0
	25	0.0
	57	0.0

Please refer to TIE 2 Report for raw data and water quality

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NAB-BAY9-TIE2-DUR	100	100.0
NAB-BAY18-TIE2-DUR	100	95.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NAB-BAY9-TIE2-DUR	100	100.0
NAB-BAY18-TIE2-DUR	100	100.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	MEAN NORM DEV (%)
NAB-BAY9-TIE2-DUR	100	96.0
NAB-BAY18-TIE2-DUR	100	96.0

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN SURVIVAL (%)
Scripps Control	n/a	100.0
Salt Control	n/a	100.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN SURVIVAL (%)
Natural Seawater	n/a	100.0
Salt Control	n/a	95.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN NORM DEV (%)
Natural Seawater	n/a	96.0
Brine Control	n/a	95.0

SUMMARY RESULTS- QA/QC

COPPER REFERENCE TOXICANT TEST

SPECIES	DATE	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	4/6/2005	50	100	101.8	86.1-120.5
MYSIDS	5/19/2005	214.4	326	271.5	236.1-305.75
MUSSELS	3/19/2005	10	20.0	13.04	12.8-13.3

Reference Toxicant tests are within two standard deviations of Nautilus' control chart mean
Please refer to TIE II Report for raw data and water quality

SDB7- 04/27/2005

OUTFALLS

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NAB-OF9-SDB7-FF	12.5	a	4	80.0	90.0	11.5	94.7	0.500	No
		b	4	80.0					
		c	5	100.0					
		d	6	100.0					
	50	a	6	100.0	95.0	10.0	100.0	0.312	No
		b	5	100.0					
		c	4	80.0					
		d	5	100.0					
	100	a	4	80.0	85.0	10.0	89.5	0.104	No
		b	5	100.0					
		c	4	80.0					
		d	4	80.0					
NAB-OF9-SDB7-COMP	12.5	a	5	100.0	90.8	10.7	95.6	0.500	No
		b	4	80.0					
		c	5	100.0					
		d	5	83.3					
	50	a	4	66.7	81.7	13.7	86.0	0.104	No
		b	4	80.0					
		c	5	100.0					
		d	4	80.0					
	100	a	3	60.0	60.0	16.3	63.2	0.007	Yes
		b	4	80.0					
		c	3	60.0					
		d	2	40.0					
NAB-OF18-SDB7-FF	12.5	a	4	80.0	85.0	10.0	85.0	0.029	Yes
		b	5	100.0					
		c	4	80.0					
		d	4	80.0					
	50	a	5	100.0	90.0	11.5	90.0	0.091	No
		b	5	100.0					
		c	4	80.0					
		d	4	80.0					
	100	a	5	100.0	90.0	11.5	90.0	0.091	No
		b	4	80.0					
		c	5	100.0					
		d	4	80.0					
NAB-OF18-SDB7-COMP	12.5	a	5	100.0	95.0	10.0	95.0	0.196	No
		b	5	100.0					
		c	4	80.0					
		d	5	100.0					
	50	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	90.0	11.5	90.0	0.091	No
		b	5	100.0					
		c	4	80.0					
		d	4	80.0					

^aControls (QA/QC) correspond to all samples from SDB7

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NAB-BAY9-SDB7-PRE	100	a	4	80.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NAB-BAY18-SDB7-PRE	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NAB-BAY9-SDB7-DUR	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NAB-BAY18-SDB7-DUR	100	a	4	80.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NAB-BAY9-SDB7-PRE	100	a	137	8	94.5	94.6	1.0	102.7	0.064	No
		b	115	7	94.3					
		c	133	9	93.7					
		d	134	5	96.4					
		e	135	8	94.4					
NAB-BAY18-SDB7-PRE	100	a	133	16	89.3	91.6	3.6	99.4	0.395	No
		b	116	15	88.5					
		c	138	7	95.2					
		d	138	6	95.8					
		e	129	16	89.0					
NAB-BAY9-SDB7-DUR	100	a	136	9	93.8	93.2	2.1	101.2	0.248	No
		b	125	13	90.6					
		c	128	11	92.1					
		d	122	5	96.1					
		e	117	8	93.6					
NAB-BAY18-SDB7-DUR	100	a	112	18	86.2	92.3	3.6	100.2	0.470	No
		b	136	9	93.8					
		c	139	10	93.3					
		d	128	10	92.8					
		e	124	6	95.4					

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	95.0	10.0	n/a	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
Salt Control	n/a	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Copper Ref. Tox.	50	a	3	60.0	85.0	19.1	89.5	0.201	No
		b	4	80.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	75.0	19.1	78.9	0.065	No
		b	3	60.0					
		c	4	80.0					
		d	3	60.0					
	200	a	2	40.0	70.0	34.6	73.7	0.124	No
		b	5	100.0					
		c	5	100.0					
		d	2	40.0					
	400	a	0	0.0	25.0	25.2	26.3	0.004	Yes
		b	3	60.0					
		c	1	20.0					
		d	1	20.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	153	12	92.7	92.1	2.4	100.0	n/a	n/a
		b	154	20	88.5					
		c	137	9	93.8					
		d	99	7	93.4					
Copper Ref. Tox.	2.9	a	120	18	87.0	82.7	8.5	89.8	0.034	Yes
		b	131	17	88.5					
		c	95	37	72.0					
		d	130	13	90.9					
	4.1	e	101	33	75.4					
		a	95	81	54.0	54.7	10.4	59.3	0.000	Yes
		b	87	50	63.5					
		c	-	-	-					
		d	-	-	-					
		e	-	-	-					
	5.9	a	0	131	0.0	2.3	3.2	2.5	0.000	Yes
		b	4	165	2.4					
		c	9	106	7.8					
		d	2	149	1.3					
	8.4	e	0	147	0.0					
		a	0	131	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	135	0.0					
		c	0	151	0.0					
		d	0	154	0.0					
		e	0	137	0.0					

SUMMARY RESULTS- QA/QC

COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	200.0	400.0	268.2	160.3-506.5
MUSSELS	2.9	4.1	4.3	3.78-4.69

Dash indicates no data (replicate was spilled or organisms not added)

^aControls (QA/QC) correspond to all samples from SDB7

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control
n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (*A. affinis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)											
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96						
NAB-OF9-SDB7-FF	12.5%	a	7.9	7.8	7.7	7.8	7.7	7.4	6.8	6.6	7.2	7.1	18.9	18.6	19.3	19.0	19.1	33.1	33.3	33.3	33.6	33.6	
	25%	a	8.0	7.8	7.7	7.8	7.7	7.9	6.5	6.4	7.0	7.0	18.8	18.6	19.3	19.0	19.1	32.9	32.9	32.9	33.2	33.1	
	100%	a	8.1	7.9	7.7	7.8	7.7	7.9	6.2	6.5	6.4	6.4	18.6	18.5	19.3	18.9	19.0	32.1	32.2	32.2	33.2	33.4	
NAB-OF9-SDB7-COMP	12.5%	a	7.8	7.8	7.7	7.8	7.6	7.6	6.3	6.1	6.6	6.1	19.2	18.3	19.4	18.6	19.3	33.4	33.3	33.3	33.2	33.7	
	25%	a	7.9	7.8	7.6	7.7	7.7	7.8	5.5	5.8	7.0	6.6	18.6	18.3	19.4	18.6	19.2	33.1	33.0	33.1	33.2	33.3	
	100%	a	8.1	7.9	7.7	7.8	7.8	8.2	5.6	6.0	6.3	6.6	18.2	18.3	19.4	18.3	19.2	32.5	32.6	32.8	32.9	33.0	
NAB-OF18-SDB7-FF	12.5%	a	7.8	7.8	7.6	7.8	7.7	7.5	6.2	6.1	6.8	6.8	19.0	18.5	19.5	18.7	19.0	33.1	33.2	33.2	33.3	33.4	
	50%	a	8.0	7.8	7.6	7.8	7.7	7.7	6.3	6.3	6.8	6.5	18.8	18.5	19.4	18.6	18.9	32.8	32.9	32.7	33.1	33.1	
	100%	a	8.1	7.9	7.7	7.6	7.6	7.6	7.8	5.7	5.7	5.7	6.2	18.6	18.4	19.3	18.6	18.9	32.2	32.2	32.1	32.2	32.3
NAB-OF18-SDB7-COMP	12.5%	a	7.8	7.7	7.7	7.8	7.7	7.8	6.6	6.3	6.9	7.2	18.9	18.0	19.3	18.0	18.8	33.4	33.5	33.3	33.8	34.0	
	50%	a	7.8	7.7	7.6	7.7	7.7	7.8	6.7	6.4	7.2	7.1	18.6	18.0	19.3	18.0	18.8	32.8	32.9	32.9	33.1	33.3	
	100%	a	7.9	7.7	7.5	7.7	7.7	8.0	5.6	5.8	6.1	6.6	18.6	18.1	19.3	18.2	18.8	32.2	32.2	32.2	33.2	32.3	
NAB-BAY9-SDB7-PRE	100%	a	7.8	7.8	7.7	7.8	7.8	7.4	6.9	6.3	7.3	6.8	19.0	18.6	19.3	18.8	19.3	33.6	33.7	33.9	33.9	33.8	
	100%	a	7.8	7.8	7.7	7.8	7.7	7.3	6.6	6.4	7.3	6.3	19.1	18.4	19.4	18.8	19.3	33.5	33.7	33.9	33.7	33.7	
	100%	a	7.7	7.6	7.7	7.7	7.7	7.3	7.0	6.4	7.3	6.2	18.6	18.3	19.3	18.6	19.1	32.2	32.2	32.2	33.2	32.2	
NAB-BAY18-SDB7-DUR	100%	a	7.7	7.7	7.7	7.8	7.8	7.5	7.0	6.7	7.5	7.1	18.4	18.3	19.4	18.6	19.0	32.9	33.0	33.0	33.1	33.0	
	Scripps Control	0	a	7.8	7.8	7.6	7.8	7.7	7.7	6.9	6.6	7.5	7.2	18.6	18.3	19.3	18.3	19.1	31.6	31.5	31.7	31.7	31.9
	Scripps Cu Ref. Tox.	50 µg/l	a	7.8	7.8	7.6	7.8	7.8	7.0	6.7	7.7	7.4	7.4	18.8	18.3	19.3	18.3	19.0	31.5	31.5	31.6	31.6	31.6
	100 µg/l	a	7.8	7.8	7.6	7.8	7.8	7.7	7.0	6.6	7.7	7.6	18.8	18.3	19.3	18.4	19.1	31.5	31.5	31.6	31.5	31.6	
	200 µg/l	a	7.8	7.8	7.6	7.8	7.7	7.7	7.1	6.6	7.9	7.4	18.8	18.2	19.1	18.4	19.0	31.5	31.6	31.8	31.9	31.6	
	400 µg/l	a	7.9	7.8	7.7	7.8	7.7	7.7	7.2	7.1	7.9	7.3	18.8	18.3	19.1	18.6	19.0	31.5	31.6	31.8	31.8	31.6	
Salt Control	n/a	a	7.9	7.7	7.5	7.6	7.5	7.4	6.1	6.4	7.0	6.6	19.4	18.3	19.2	18.6	19.0	31.4	31.5	31.6	31.4	31.5	

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NAB-BAY9-SDB7-PRE	100%	f	7.8	7.8	7.8	7.5	8.4	8.2	16.0	15.1	14.9	33.8	33.8	33.8
NAB-BAY18-SDB7-PRE	100%	f	7.8	7.8	7.8	7.9	8.5	8.4	16.0	15.3	15.1	29.7	30.0	30.2
NAB-BAY9-SDB7-DUR	100%	f	7.8	7.8	7.8	7.9	8.4	8.3	16.0	15.0	14.8	29.7	29.9	30.2
NAB-BAY18-SDB7-DUR	100%	f	7.9	7.8	7.8	7.8	8.5	8.6	16.0	15.1	15.0	29.6	29.6	29.7
Scripps Control	0	f	8.1	8.0	7.7	7.5	8.2	8.2	16.0	15.5	15.2	29.4	29.3	29.0
Scripps Cu Ref. Tox.	2.9 µg/l	f	7.9	7.9	7.9	7.2	8.3	8.5	15.8	15.2	15.0	29.5	29.6	29.7
	4.1 µg/l	f	7.8	7.8	7.7	7.9	8.4	8.4	15.8	15.2	15.3	29.5	29.6	29.7
	5.9 µg/l	f	7.8	7.8	7.7	7.8	8.3	8.4	15.9	15.2	15.4	29.5	29.6	29.6
	8.4 µg/l	f	7.8	7.8	7.7	7.9	8.6	8.2	15.9	15.2	15.2	29.6	29.5	29.6
	12 µg/l	f	7.8	7.8	7.7	7.9	8.5	8.3	15.8	15.0	15.1	29.6	29.5	29.6
	17.2 µg/l	f	7.8	7.8	7.8	7.7	8.2	8.5	15.8	15.0	15.1	29.6	29.6	29.7

Appendix C4

N/

SDB4- 10/17/2004

SDB5- 01/10/2005

SDB6- 2/10/2005

TIE2- 3/19/2005

SDB7- 4/17/2005

SDB4 – 10/17/2004

OUTFALLS

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB4-FF	12.5	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	25	a	5	100.0	85.0	19.1	85.0	0.108	No
		b	4	80.0					
		c	3	60.0					
		d	5	100.0					
	50	a	5	100.0	90.0	11.5	90.0	0.091	No
		b	4	80.0					
		c	5	100.0					
		d	4	80.0					
	100	a	3	60.0	80.0	16.3	80.0	0.046	Yes
		b	4	80.0					
		c	4	80.0					
		d	5	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB4-FF	12.5	a	9	90.0	96.7	5.8	96.7	0.211	No
		b	10	100.0					
		c	10	100.0					
	25	a	10	100.0	96.7	5.8	96.7	0.211	No
		b	9	90.0					
		c	10	100.0					
	50	a	8	80.0	76.7	17.3	76.7	0.048	Yes
		b	9	90.0					
		c	6	60.0					
	100	a	6	60.0	56.7	5.8	56.7	0.003	Yes
		b	5	50.0					
		c	6	60.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB4-FF	6.25	a	164	7	95.9	92.6	3.6	95.0	0.018	Yes
		b	164	13	92.7					
		c	168	6	96.6					
		d	137	17	89.0					
		e	145	18	89.0					
	12.5	a	121	41	74.7	83.2	7.3	85.3	0.006	Yes
		b	169	20	89.4					
		c	163	32	83.6					
		d	175	17	91.1					
		e	157	47	77.0					
	25	a	5	161	3.0	6.7	5.6	6.9	0.000	Yes
		b	1	170	0.6					
		c	10	146	6.4					
		d	13	140	8.5					
		e	27	152	15.1					
	50	a	1	200	0.5	0.3	0.5	0.3	0.000	Yes
		b	0	126	0.0					
		c	2	187	1.1					
		d	0	149	0.0					
		e	0	145	0.0					
	61.4	a	0	196	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	196	0.0					
		c	0	196	0.0					
		d	0	196	0.0					
		e	0	196	0.0					

^aControls (QA/QC) correspond to all samples from SDB4

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB4-DUR	100	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB4-DUR	100	a	10	100.0	100.0	0.0	100.0	n/a	No
		b	10	100.0					
		c	10	100.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB4-DUR	100	a	167	5	97.1	97.6	0.5	100.1	0.3966	No
		b	162	3	98.2					
		c	187	4	97.9					
		d	167	5	97.1					
		e	188	4	97.9					

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Salt Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Copper Ref. Tox.	50	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	4	80.0	90.0	11.5	90.0	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
	200	a	0	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	0.0					
		c	0	0.0					
		d	0	0.0					
	400	a	0	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	0.0					
		c	0	0.0					
		d	0	0.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	9	90.0	93.3	5.8	100.0	n/a	n/a
		b	9	90.0					
		c	10	100.0					
Salt Control	n/a	a	10	100.0	100.0	0.0	107.1	0.092	No
		b	10	100.0					
		c	10	100.0					
Copper Ref. Tox.	25	a	10	100.0	100.0	0.0	107.1	0.092	No
		b	10	100.0					
		c	10	100.0					
	50	a	10	100.0	100.0	0.0	107.1	0.092	No
		b	10	100.0					
		c	10	100.0					
	100	a	10	100.0	100.0	0.0	107.1	0.092	No
		b	10	100.0					
		c	10	100.0					
	200	a	9	90.0	83.3	5.8	89.3	0.051	No
		b	8	80.0					
		c	8	80.0					
	400	a	2	20.0	6.7	11.5	7.1	0.001	Yes
		b	0	0.0					
		c	0	0.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	148	25	85.5	94.5	5.4	100.0	n/a	No
		b	175	5	97.2					
		c	139	10	93.3					
		d	193	4	98.0					
		e	174	3	98.3					
Brine Control	n/a	a	177	6	96.7	95.7	1.1	98.1	0.323	No
		b	170	10	94.4					
		c	186	6	96.9					
		d	171	8	95.5					
		e	164	9	94.8					
Copper Ref. Tox.	2.9	a	167	9	94.9	95.1	0.7	100.7	0.397	No
		b	200	11	94.8					
		c	168	10	94.4					
		d	176	8	95.7					
		e	168	7	96.0					
	4.1	a	166	3	98.2	90.3	10.0	95.6	0.221	No
		b	202	7	96.7					
		c	164	17	90.6					
		d	118	43	73.3					
		e	141	11	92.8					
	5.9	a	178	9	95.2	79.0	14.3	83.7	0.036	Yes
		b	169	20	89.4					
		c	157	36	81.3					
		d	128	60	68.1					
		e	124	79	61.1					
	8.4	a	69	106	39.4	23.7	13.7	25.1	0.000	Yes
		b	56	141	28.4					
		c	58	126	31.5					
		d	12	177	6.3					
		e	24	162	12.9					
	12.0	a	1	177	0.6	1.3	1.3	1.3	0.000	Yes
		b	5	172	2.8					
		c	5	203	2.4					
		d	1	207	0.5					
		e	0	171	0.0					
	17.2	a	3	177	1.7	0.5	0.7	0.5	0.000	Yes
		b	1	167	0.6					
		c	0	191	0.0					
		d	0	175	0.0					
		e	0	199	0.0					

SUMMARY RESULTS- QA/QC
COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	50	100	132.0	120.2-144.8
MYSIDS	200	400	265.3	232.5-302.4
MUSSELS	5.9	8.4	7.29	6.1-8.3

^aControls (QA/QC) correspond to all samples from SDB4

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control
n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (*A. affinis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																			
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																														
NI-OF23A-SDB4-FF	12.5%	a	7.9	7.7	7.6	7.7	7.6	6.8	5.2	5.1	5.2	6.0	19.9	18.3	19.3	19.3	19.6	33.9	34.2	34.3	34.3	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	34.2	

MYSIDS (*A. bahia*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)										
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96					
NI-OF23A-SDB4-FF	12.5%	a	7.9	7.7	7.8	7.8	7.9	6.7	3.9	6.9	6.8	6.9	19.3	18.3	19.5	19.1	19.3	34.3	34.5	34.5	34.6	34.6
	25%	a	8.1	7.6	7.7	7.9	7.9	6.8	6.2	6.5	7.0	7.1	19.2	18.8	19.6	19.3	19.4	34.4	34.5	34.5	34.6	34.6
	50%	a	8.3	7.6	7.8	7.7	7.9	7.0	6.6	6.9	6.0	6.7	18.8	18.8	19.3	19.1	19.3	34.1	34.2	34.4	34.1	34.2
	100%	a	8.7	7.7	7.8	7.8	7.9	7.1	6.3	6.7	6.5	6.6	18.3	18.0	19.5	19.1	19.4	33.6	34.2	34.4	34.0	34.5
NI-BAY23A-SDB4-DUR	100%	a	7.8	7.7	7.6	7.7	7.7	7.2	5.6	5.3	5.9	6.4	18.6	18.3	19.7	19.5	19.7	34.3	34.5	34.5	34.4	34.6
Scripps Control	0	a	7.9	7.7	7.7	7.7	7.6	6.9	4.7	5.3	5.0	4.8	18.8	18.3	19.4	19.3	19.5	34.4	34.6	34.6	34.2	34.3
Scripps Cu Ref. Tox.	25 µg/l	a	7.9	7.7	7.7	7.8	7.6	6.9	5.4	5.7	5.4	5.2	18.8	18.6	19.4	19.3	19.6	34.3	34.4	34.6	34.3	34.3
	50 µg/l	a	7.9	7.7	7.6	7.7	7.6	7.1	5.3	5.3	5.0	4.5	18.7	18.6	19.5	19.3	19.6	34.5	34.5	34.5	34.2	34.2
	100 µg/l	a	7.9	7.7	7.6	7.7	7.7	7.1	5.7	4.9	5.6	5.1	18.6	18.5	19.4	19.3	19.6	34.2	34.4	34.5	34.3	34.3
	200 µg/l	a	7.9	7.8	7.8	7.8	7.7	7.0	5.9	6.3	6.1	5.7	18.7	18.6	19.1	19.3	19.6	34.4	34.6	34.2	34.3	34.4
	400 µg/l	a	7.9	7.8	7.6	7.9	7.8	7.0	6.1	5.5	6.4	6.1	18.7	18.6	19.4	19.3	19.6	34.3	34.5	34.6	34.2	34.2
Salt Control	n/a	a	7.9	7.8	7.6	7.9	7.8	7.0	6.1	5.5	6.4	6.1	18.7	18.6	19.4	19.3	19.6	34.3	34.5	34.6	34.2	34.2

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)			D.O. (mg/l)	Temp (°C)	Salinity (‰)		
			0	48	72			0	48	72
NI-OF23A-SDB4-FF	6.25%	f	7.7	7.6	7.0	6.6	15.7	33.6	33.8	33.8
	25%	f	7.7	7.6	7.0	5.8	15.5	34.0	33.9	33.9
	61.4%	f	7.7	7.3	6.9	3.3	15.6	33.6	34.0	34.0
NI-BAY23A-SDB4-DUR	100%	f	7.7	7.8	7.0	6.8	15.1	34.0	34.2	34.2
Scripps Control	0	f	7.8	7.6	6.9	6.8	15.6	34.0	34.2	34.2
Scripps Cu Ref. Tox.	2.9 µg/l	f	7.8	7.8	7.0	6.8	15.8	33.9	34.1	34.1
	8.4 µg/l	f	7.8	7.7	6.9	6.8	15.7	33.9	34.1	34.1
	24 µg/l	f	7.8	7.8	6.9	7.1	15.8	34.1	34.1	34.1
Brine Control	0	f	7.9	7.9	7.0	7.0	15.5	33.7	34.2	34.2

N - water quality not taken due to 100% mortality in treatment

SDB5 - 01/10/2005

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-DOWNTOWN PIER-SDB5-AFT	100	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-DOWNTOWN PIER-SDB5-AFT	100	a	10	100.0	93.3	5.8		n/a	No
		b	9	90.0					
		c	9	90.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB5-AFT	100	a	184	14	92.9	93.9	1.1	104.2	0.007	Yes
		b	144	7	95.4					
		c	116	9	92.8					
		d	178	11	94.2					
		e	164	10	94.3					
NI-DOWNTOWN PIER-SDB5-AFT	100	a	150	8	94.9	93.6	1.6	103.9	0.012	Yes
		b	170	11	93.9					
		c	139	7	95.2					
		d	144	11	92.9					
		e	155	15	91.2					

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Copper Ref. Tox.	25	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	4	80.0	95.0	10.0	95.0	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	4	80.0	90.0	11.5	94.7	0.091	No
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
	200	a	1	20.0	15.0	10.0	16.7	0.000	Yes
		b	1	20.0					
		c	1	20.0					
		d	0	0.0					
	400	a	0	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	0.0					
		c	0	0.0					
		d	0	0.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	9	90.0	93.3	5.8	n/a	n/a	n/a
		b	9	90.0					
		c	10	100.0					
Copper Ref. Tox.	25	a	9	90.0	93.3	5.8	100.0	0.500	No
		b	9	90.0					
		c	10	100.0					
	50	a	10	100.0	103.3	5.8	110.7	0.051	No
		b	10	100.0					
		c	11	110.0					
	100	a	10	100.0	100.0	0.0	96.8	0.092	No
		b	10	100.0					
		c	10	100.0					
	200	a	8	80.0	90.0	10.0	90.0	0.325	No
		b	9	90.0					
		c	10	100.0					
	400	a	2	20.0	26.7	11.5	29.6	0.002	Yes
		b	2	20.0					
		c	4	40.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	160	4	97.6	97.7	1.0	100.0	n/a	No
		b	222	4	98.2					
		c	236	6	97.5					
		d	233	9	96.3					
		e	257	3	98.8					
Brine Control 1	n/a	a	204	3	98.6	98.4	0.8	100.7	0.119	No
		b	211	5	97.7					
		c	201	5	97.6					
		d	226	1	99.6					
		e	221	3	98.7					
Brine Control 2	n/a	a	189	3	98.4	97.8	1.1	100.1	0.440	No
		b	231	10	95.9					
		c	210	4	98.1					
		d	190	4	97.9					
		e	210	3	98.6					
Copper Ref. Tox.	2.9	a	231	5	97.9	98.6	0.7	101.0	0.057	No
		b	207	4	98.1					
		c	214	1	99.5					
		d	201	3	98.5					
		e	228	2	99.1					
	4.1	a	214	8	96.4	56.4	39.6	57.7	0.040	Yes
		b	205	21	90.7					
		c	-	-	-					
		d	-	-	-					
		e	-	-	-					
	5.9	a	125	101	55.3	49.3	10.8	50.5	0.000	Yes
		b	125	94	57.1					
		c	132	106	55.5					
		d	114	125	47.7					
		e	64	142	31.1					
	8.4	a	23	187	11.0	10.1	5.2	10.3	0.000	Yes
		b	24	173	12.2					
		c	4	210	1.9					
		d	32	170	15.8					
		e	21	200	9.5					
	12.0	a	0	195	0.0	0.3	0.2	0.3	0.000	Yes
		b	1	246	0.4					
		c	1	221	0.5					
		d	1	218	0.5					
		e	0	219	0.0					
	17.2	a	0	210	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	187	0.0					
		c	0	178	0.0					
		d	0	215	0.0					
		e	0	198	0.0					

SUMMARY RESULTS- QA/QC

COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	100	200	138.5	114.4-167.8
MYSIDS	200	400	324.9	276.2-379.8
MUSSELS ^d	4.1	5.9	6.0	5.9-6.1

Dash indicates no data (replicate was spilled or organisms not added)

^aControls (QA/QC) correspond to all samples from SDB5

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

^dCopper reference toxicant test performed on 02/10/2005

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (*A. affinis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NI-DOWNTOWN PIER-SDB5-AFT	100%	a	7.9	7.8	7.5	7.7	7.6	7.2	6.7	6.5	6.5	19.3	19.9	19.8
Scripps Control	n/a	a	7.9	7.8	7.6	7.8	7.7	7.5	6.2	6.4	6.1	19.0	18.8	19.7
												18.2	32.7	30.9
												30.9	30.7	30.9

MYSIDS (*A. bahia*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NI-DOWNTOWN PIER-SDB5-AFT	100%	a	8.0	7.8	7.6	7.8	7.7	7.6	6.1	5.0	5.8	18.9	18.8	19.6
Scripps Control	n/a	a	7.9	7.8	7.5	7.7	7.5	7.3	5.9	5.4	6.2	18.9	19.1	19.8
												18.2	31.8	30.7
												30.8	30.8	31.0
												31.1	30.8	30.9

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)			D.O. (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NI-OF23A-SDB5-AFT	100%	f	7.8	7.8	7.9	7.5	7.8	7.5	17.0	15.8	16.3	30.2	30.1	29.9
NI-DOWNTOWN PIER-SDB5-AFT	100%	f	7.8	7.8	7.7	7.4	7.7	7.4	17.0	16.3	16.3	27.0	26.7	26.7
Scripps Control	n/a	f	7.8	7.8	7.8	6.9	7.9	7.5	17.0	16.0	16.3	28.4	28.3	28.4

SDB6 – 02/10/2005

OUTFALLS

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB6-FF	12.5	a	5	100.0	95.0	10.0	95.0	0.196	No
		b	5	100.0					
		c	4	80.0					
		d	5	100.0					
	25	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	90.0	20.0	90.0	0.196	No
		b	5	100.0					
		c	3	60.0					
		d	5	100.0					
NI-OF26-SDB6-FF	12.5	a	4	80.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	25	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	5	100.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
	100	a	5	100.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
NI-OF26-SDB6-COMP	12.5	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	4	80.0	95.0	10.0	100.0	0.500	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB6-FF	12.5	a	10	100.0	93.3	5.8	93.3	0.092	No
		b	9	90.0					
		c	9	90.0					
	25	a	10	100.0	100.0	0.0	100.0	n/a	No
		b	10	100.0					
		c	10	100.0					
	50	a	10	100.0	100.0	0.0	100.0	n/a	No
		b	10	100.0					
		c	10	100.0					
	100	a	9	90.0	96.7	5.8	96.7	0.211	No
		b	10	100.0					
		c	10	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF26-SDB6-FF	12.5	a	10	100.0	100.0	0.0	100.0	0.211	No
		b	10	100.0					
		c	10	100.0					
	25	a	10	100.0	93.3	5.8	93.3	0.051	No
		b	9	90.0					
		c	9	90.0					
	50	a	10	100.0	96.7	5.8	96.7	0.115	No
		b	9	90.0					
		c	10	100.0					
	100	a	8	80.0	73.3	11.5	73.3	0.014	Yes
		b	8	80.0					
		c	6	60.0					
NI-OF26-SDB6-COMP	12.5	a	10	100.0	93.3	11.5	93.3	0.137	No
		b	8	80.0					
		c	10	100.0					
	50	a	10	100.0	100.0	0.0	100.0	0.211	No
		b	10	100.0					
		c	10	100.0					
	100	a	10	100.0	100.0	0.0	100.0	0.211	No
		b	10	100.0					
		c	10	100.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB6-FF	6.2	a	219	5	97.8	98.3	1.5	100.3	0.346	No
		b	244	1	99.6					
		c	215	9	96.0					
		d	230	2	99.1					
		e	229	2	99.1					
	12.4	a	210	6	97.2	96.7	1.0	98.7	0.034	Yes
		b	212	10	95.5					
		c	201	6	97.1					
		d	198	4	98.0					
		e	208	9	95.9					
	24.8	a	28	198	12.4	10.3	2.9	10.5	0.000	Yes
		b	22	208	9.6					
		c	15	199	7.0					
		d	34	206	14.2					
		e	21	226	8.5					
	49.5	a	0	221	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	203	0.0					
		c	0	219	0.0					
		d	0	219	0.0					
		e	0	235	0.0					
NI-OF26-SDB6-FF	6.2	a	207	4	98.1	97.7	0.7	100.3	0.358	No
		b	209	6	97.2					
		c	190	4	97.9					
		d	238	8	96.7					
		e	201	3	98.5					
	12.4	a	185	6	96.9	96.5	1.4	99.1	0.163	No
		b	214	13	94.3					
		c	229	8	96.6					
		d	212	7	96.8					
		e	215	4	98.2					
	24.8	a	172	22	88.7	92.6	2.5	95.0	0.004	Yes
		b	185	10	94.9					
		c	200	16	92.6					
		d	201	17	92.2					
		e	196	11	94.7					
	49.5	a	1	144	0.7	0.1	0.3	0.1	0.000	Yes
		b	0	218	0.0					
		c	0	231	0.0					
		d	0	222	0.0					
		e	0	215	0.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF26-SDB6-COMP	7.0	a	238	8	96.7	97.7	1.2	100.2	0.398	No
		b	216	7	96.9					
		c	203	2	99.0					
		d	191	2	99.0					
		e	207	7	96.7					
	13.9	a	215	9	96.0	97.0	1.4	99.6	0.319	No
		b	213	5	97.7					
		c	228	2	99.1					
		d	196	8	96.1					
		e	207	8	96.3					
	27.9	a	237	13	94.8	95.8	2.1	98.3	0.095	No
		b	202	6	97.1					
		c	218	14	94.0					
		d	181	2	98.9					
		e	218	13	94.4					
	55.7	a	198	14	93.4	95.5	1.8	97.9	0.042	Yes
		b	212	8	96.4					
		c	201	8	96.2					
		d	201	5	97.6					
		e	210	14	93.8					

^aControls (QA/QC) correspond to all samples from SDB6

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB6-PRE	100	a	5	100	100.0	0.0	100.0	n/a	No
		b	5	100					
		c	5	100					
		d	5	100					
NI-BAY26-SDB6-PRE	100	a	4	80	95.0	10.0	95.0	0.196	No
		b	5	100					
		c	5	100					
		d	5	100					
NI-OF23A-SDB6-DUR	100	a	5	100	100.0	0.0	100.0	n/a	No
		b	5	100					
		c	5	100					
		d	5	100					
NI-BAY26-SDB6-DUR	100	a	5	100	100.0	0.0	100.0	n/a	No
		b	5	100					
		c	5	100					
		d	5	100					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB6-PRE	100	a	10	100	100.0	0.0	107.1	n/a	No
		b	10	100					
		c	10	100					
NI-BAY26-SDB6-PRE	100	a	10	100	100.0	0.0	107.1	n/a	No
		b	11	100					
		c	10	100					
NI-OF23A-SDB6-DUR	100	a	10	100	100.0	0.0	107.1	n/a	No
		b	10	100					
		c	10	100					
NI-BAY26-SDB6-DUR	100	a	10	100	100.0	0.0	107.1	n/a	No
		b	10	100					
		c	10	100					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP.	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB6-PRE	100.0	a	221	4	98.2	98.0	0.8	99.6	0.294	No
		b	203	5	97.6					
		c	214	4	98.2					
		d	188	6	96.9					
		e	234	2	99.2					
NI-BAY26-SDB6-PRE	100	a	213	3	98.6	97.5	1.3	99.0	0.377	No
		b	234	6	97.5					
		c	211	3	98.6					
		d	203	6	97.1					
		e	208	10	95.4					
NI-OF23A-SDB6-DUR	100	a	207	4	98.1	97.1	1.5	99.4	0.130	No
		b	154	3	98.1					
		c	193	6	97.0					
		d	205	5	97.6					
		e	193	11	94.6					
NI-BAY26-SDB6-DUR	100	a	218	9	96.0	96.4	0.8	98.6	0.079	No
		b	219	8	96.5					
		c	233	8	96.7					
		d	221	11	95.3					
		e	183	5	97.3					

QA/QC SAMPLES^a
TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Salt Control	n/a	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
Copper Ref. Tox.	50	a	5	100	100.0	0.0	100.0	n/a	No
		b	5	100					
		c	5	100					
		d	5	100					
	100	a	5	100	70.0	47.6	70.0	0.148	No
		b	0	0					
		c	5	100					
		d	4	80					
	200	a	1	20	10.0	11.5	10.0	0.000	Yes
		b	0	0					
		c	0	0					
		d	1	20					
	400	a	0	0	0.0	0.0	0.0	0.000	Yes
		b	0	0					
		c	0	0					
		d	0	0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	8	80.0	93.3	11.5	100.0	n/a	n/a
		b	10	100.0					
		c	11	100.0					
Salt Control	n/a	a	10	100.0	100.0	0.0	107.1	0.211	No
		b	10	100.0					
		c	10	100.0					

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Copper Ref. Tox.	100	a	10	100	100.0	0.0	107.1	0.371	No
		b	10	100					
		c	10	100					
	200	a	10	100	96.7	5.8	103.6	0.500	No
		b	10	100					
		c	9	90					
	400	a	3	30	33.3	5.8	35.7	0.005	Yes
		b	3	30					
		c	4	40					
	800	a	0	0	0.0	0.0	0.0	0.004	Yes
		b	0	0					
		c	0	0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	ABNORMA	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	160	4	97.6	97.7	1.0	100.0	n/a	n/a
		b	222	4	98.2					
		c	236	6	97.5					
		d	233	9	96.3					
		e	257	3	98.8					
Brine Control 1	n/a	a	204	3	98.6	98.4	0.8	100.7	0.119	No
		b	211	5	97.7					
		c	201	5	97.6					
		d	226	1	99.6					
		e	221	3	98.7					
Brine Control 2	n/a	a	189	3	98.4	97.8	1.1	100.1	0.440	No
		b	231	10	95.9					
		c	210	4	98.1					
		d	190	4	97.9					
		e	210	3	98.6					
Copper Ref. Tox.	2.9	a	231	5	97.9	98.6	0.7	101.0	0.057	No
		b	207	4	98.1					
		c	214	1	99.5					
		d	201	3	98.5					
		e	228	2	99.1					
	4.1	a	214	8	96.4	56.4	39.6	57.7	0.040	Yes
		b	205	21	90.7					
		c	-	-	-					
		d	-	-	-					
		e	-	-	-					
	5.9	a	125	101	55.3	49.3	10.8	50.5	0.000	Yes
		b	125	94	57.1					
		c	132	106	55.5					
		d	114	125	47.7					
		e	64	142	31.1					
	8.4	a	23	187	11.0	10.1	5.2	10.3	0.000	Yes
		b	24	173	12.2					
		c	4	210	1.9					
		d	32	170	15.8					
		e	21	200	9.5					
	12.0	a	0	195	0.0	0.3	0.2	0.3	0.000	Yes
		b	1	246	0.4					
		c	1	221	0.5					
		d	1	218	0.5					
		e	0	219	0.0					
	17.2	a	0	210	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	187	0.0					
		c	0	178	0.0					
		d	0	215	0.0					
		e	0	198	0.0					

SUMMARY RESULTS- QA/QC

COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	100.0	200.0	123.5	103.3-147.5
MYSIDS	200.0	400.0	352.5	326.3-387.7
MUSSELS	4.1	5.9	6.0	5.9-6.1

Dash indicates no data (replicate was spilled or organisms not added)

^aControls (QA/QC) correspond to all samples from SDB6

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (A. affinis)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)										
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96					
NI-OF23A-SDB6-FF	12.5%	a	7.6	7.7	7.7	7.7	7.4	7.0	7.3	6.5	6.1	5.5	19.5	18.6	19.6	19.6	20.0	33.4	33.4	33.4	33.6	33.8
	25%	a	7.6	7.7	7.7	7.6	7.5	7.0	7.4	6.5	6.3	6.3	19.2	18.6	19.6	19.7	20.0	33.4	33.4	33.4	34.0	34.1
	50%	a	7.7	7.7	7.7	7.6	7.5	7.2	7.0	6.4	6.2	6.1	18.9	18.7	19.6	19.6	20.1	33.4	33.4	33.5	33.9	34.0
	100%	a	7.7	7.7	7.7	7.6	7.6	7.5	6.9	6.5	6.3	6.5	18.8	18.7	19.5	19.5	19.9	33.0	33.2	33.2	34.1	34.4
NI-OF26-SDB6-FF	12.5%	a	7.6	7.7	7.7	7.6	7.5	7.1	7.4	6.5	6.7	6.5	18.9	18.8	19.6	19.5	19.9	33.3	33.3	33.3	33.6	33.7
	25%	a	7.6	7.7	7.7	7.6	7.6	7.3	7.4	6.5	6.7	6.7	18.9	18.4	19.5	19.4	19.9	33.2	33.3	33.3	33.6	33.8
	50%	a	7.6	7.6	7.7	7.6	7.6	7.2	7.0	6.5	6.4	6.9	19.0	18.5	19.6	19.3	19.8	32.9	32.9	32.9	33.5	33.8
	100%	a	7.8	7.7	7.6	7.6	7.6	7.5	6.9	6.4	6.0	6.4	19.1	18.4	19.5	19.4	19.8	32.8	32.9	33.1	33.3	33.3
NI-OF26-SDB6-COMP	12.5%	a	7.7	7.7	7.7	7.6	7.6	7.3	7.3	6.5	6.3	6.6	18.8	18.4	19.6	19.5	19.7	32.9	32.9	33.0	33.7	33.8
	50%	a	7.6	7.7	7.7	7.6	7.6	7.3	7.3	6.5	6.4	6.7	18.8	18.4	19.6	19.5	19.8	33.1	32.9	33.1	33.4	33.5
	100%	a	7.8	7.7	7.6	7.6	7.6	7.8	7.3	6.7	6.7	6.8	18.9	18.3	19.5	19.0	19.6	33.2	33.1	33.6	34.1	34.8
	100%	a	7.6	7.7	7.7	7.6	7.6	7.5	6.8	7.4	6.6	7.3	5.8	19.1	18.7	19.6	19.4	20.0	33.5	33.5	33.3	33.5
NI-BAY26-SDB6-PRE	100%	a	7.6	7.7	7.7	7.6	7.6	7.2	7.2	6.6	7.3	6.4	18.9	18.8	19.6	19.3	19.9	33.0	33.3	33.3	33.3	33.8
	100%	a	7.6	7.6	7.7	7.6	7.5	7.3	7.6	6.7	7.2	6.3	19.7	18.8	19.5	19.4	19.9	32.9	32.9	33.0	33.2	33.7
	100%	a	7.6	7.7	7.7	7.5	7.5	7.3	7.2	6.5	7.0	6.6	18.8	18.2	19.7	19.6	20.3	30.7	30.8	30.4	31.2	31.2
	0	a	7.6	7.7	7.7	7.6	7.6	7.8	7.7	7.2	7.4	7.2	18.8	18.8	19.0	19.3	19.1	32.0	32.6	33.5	32.1	33.5
Scripps Control	50 µg/l	a	7.6	7.7	7.7	7.6	7.6	6.8	7.3	7.2	7.4	7.1	18.5	18.7	19.3	19.3	19.4	32.2	32.9	33.1	31.7	32.4
	100 µg/l	a	7.6	7.7	7.6	7.6	7.6	6.9	7.7	7.2	6.9	7.1	18.6	18.5	19.0	19.6	19.5	32.5	32.9	33.1	31.9	32.4
	200 µg/l	a	7.6	7.7	7.6	7.7	7.7	6.9	7.8	7.2	7.9	7.0	18.6	18.1	19.3	19.4	19.4	32.1	32.9	33.0	32.5	32.6
	400 µg/l	a	7.6	7.7	7.6	N	N	6.9	7.9	7.1	N	N	N	18.6	18.1	19.1	N	N	32.1	32.9	33.0	N
Salt Control	n/a	a	7.6	7.7	7.4	7.3	7.3	5.9	7.1	6.9	7.2	6.9	18.8	18.3	19.0	19.4	19.5	31.6	32.7	33.1	32.0	32.7

MYSIDS (A. bahia)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)					Dissolved Oxygen (mg/l)					Temperature (°C)					Salinity (‰)				
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96	0	24	48	72	96
NI-OF23A-SDB6-FF	12.5%	a	7.7	7.6	7.4	7.5	7.5	7.0	6.7	5.0	6.0	5.2	20.3	18.6	19.5	19.4	19.3	33.2	33.4	33.1	33.5	33.7
	25%	a	7.7	7.5	7.4	7.5	7.5	6.9	5.6	4.4	5.3	5.0	19.9	18.8	19.5	19.4	19.4	33.1	33.4	33.3	33.7	33.8
	50%	a	7.8	7.6	7.6	7.5	7.5	6.7	5.0	5.5	5.3	5.1	19.1	18.8	19.5	19.5	19.4	33.0	33.4	33.3	33.7	33.7
	100%	a	7.9	7.7	7.5	7.6	7.6	7.7	6.3	4.7	4.8	5.2	18.6	18.8	19.5	19.6	19.3	33.0	33.4	33.3	33.6	33.6
NI-OF26-SDB6-FF	12.5%	a	7.7	7.6	7.5	7.5	7.5	7.0	6.9	5.0	5.5	5.7	19.2	18.6	19.4	19.6	19.3	32.3	33.1	33.0	33.4	33.6
	25%	a	7.7	7.6	7.5	7.5	7.6	7.4	6.5	4.9	5.4	5.8	19.4	18.6	19.4	19.6	19.4	32.6	32.9	32.9	33.4	33.5
	50%	a	7.8	7.7	7.5	7.5	7.5	7.6	6.2	4.8	4.8	5.2	19.5	18.7	19.4	19.7	19.4	32.6	32.7	32.7	33.1	33.2
	100%	a	7.9	7.7	7.5	7.6	7.6	7.2	5.8	4.9	5.2	5.2	19.8	18.7	19.4	19.4	19.3	31.7	32.7	32.6	32.8	33.0
NI-OF26-SDB6-COMP	12.5%	a	7.7	7.7	7.6	7.5	7.5	7.3	6.9	5.0	5.9	5.8	18.9	18.7	19.7	19.6	19.4	32.9	33.0	33.0	33.8	33.5
	50%	a	7.7	7.7	7.6	7.5	7.6	7.4	6.4	4.8	5.5	5.2	18.8	18.4	19.7	19.5	19.3	32.9	33.0	33.1	33.3	33.4
	100%	a	7.8	7.7	7.6	7.6	7.5	7.8	6.2	4.4	4.9	5.1	19.2	18.6	19.7	19.6	19.4	33.0	33.2	33.1	33.1	33.4
	100%	a	7.7	7.7	7.6	7.5	7.5	6.7	7.3	5.5	5.9	4.4	20.2	18.7	19.6	19.4	19.6	33.1	33.7	33.4	33.8	34.2
NI-BAY26-SDB6-PRE	100%	a	7.7	7.6	7.5	7.5	7.6	7.1	6.8	5.3	6.2	6.0	19.3	18.6	19.6	19.6	19.4	33.1	33.4	33.3	33.8	33.8
NI-OF23A-SDB6-DUR	100%	a	7.5	7.7	7.5	7.5	7.8	7.4	7.0	5.7	6.3	5.4	19.8	18.8	19.5	19.5	19.3	32.6	33.0	33.0	33.8	33.5
NI-BAY26-SDB6-DUR	100%	a	7.6	7.7	7.6	7.6	7.6	7.9	6.9	4.7	5.5	5.5	19.9	18.6	19.7	19.7	19.4	32.6	32.9	32.9	33.3	33.4
Scripps Control	0	a	7.7	7.7	7.6	7.6	7.6	6.8	7.3	5.4	6.1	6.4	19.3	18.8	19.8	19.8	18.9	32.1	32.1	32.3	31.9	32.1
Scripps Cu Ref. Tox	100 µg/l	a	7.6	7.6	7.5	7.6	7.5	6.9	7.2	5.1	6.1	5.8	19.3	18.8	19.7	19.8	19.6	32.2	32.2	32.6	32.1	32.2
	200 µg/l	a	7.6	7.6	7.6	7.6	7.6	7.0	7.2	5.0	6.5	6.4	19.1	18.7	19.7	19.8	19.5	32.4	32.3	32.6	32.1	32.1
	400 µg/l	a	7.6	7.7	7.6	7.6	7.6	7.0	7.3	5.4	7.0	6.9	19.0	18.8	19.7	19.6	19.4	32.1	32.1	32.6	32.2	32.2
	800 µg/l	a	7.6	7.6	7.6	7.2	7.7	7.1	7.6	6.0	6.2	6.9	19.0	18.8	19.6	19.8	19.4	32.3	32.0	32.6	32.2	32.2
Salt Control	n/a	a	7.6	7.4	7.3	7.7	7.2	5.8	6.6	4.7	7.6	6.4	19.3	18.8	19.8	19.6	19.6	31.8	31.9	32.2	32.2	32.7

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)			D.O. (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NI-OF23A-SDB6-FF	6.2%	f	7.8	7.8	7.8	8.0	6.9	7.5	15.2	15.8	15.5	32.5	32.6	32.5
	12.4%	f	7.8	7.8	7.8	8.0	7.0	7.7	16.2	15.8	15.4	32.9	32.0	32.6
	24.8%	f	7.8	7.8	7.8	7.8	7.0	7.6	16.2	15.7	15.4	31.6	31.7	32.9
	49.5%	f	7.8	7.8	7.8	8.0	7.0	7.7	16.3	15.8	15.5	31.5	31.5	32.0
NI-OF26-SDB6-FF	6.2%	f	7.8	7.8	7.8	7.8	7.2	7.5	15.0	15.8	15.4	32.7	32.6	32.4
	12.4%	f	7.8	7.8	7.8	7.8	7.2	7.7	15.0	15.9	15.4	31.5	32.5	32.3
	24.8%	f	7.8	7.8	7.8	8.0	7.1	7.6	15.1	15.8	15.4	32.5	32.1	31.9
	49.5%	f	7.8	7.8	7.8	7.9	7.2	7.7	15.0	15.7	15.5	32.4	31.9	31.9
NI-OF26-SDB6-COMP	7.0%	f	7.8	7.8	7.8	8.2	7.1	7.5	16.3	15.9	15.7	32.4	32.6	33.0
	13.9%	f	7.9	7.8	7.8	8.2	7.1	7.6	16.2	15.9	15.7	32.7	32.5	32.5
	27.9%	f	7.9	7.7	7.8	8.2	7.1	7.3	16.2	16.0	15.6	32.2	31.9	32.3
	55.7%	f	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND
NI-OF23A-SDB6-PRE	100%	f	7.8	7.8	7.8	7.8	6.9	7.5	15.3	15.8	15.5	33.2	32.9	32.5
NI-BAY26-SDB6-PRE	100%	f	7.8	7.8	7.8	7.7	7.1	7.6	15.3	15.8	15.4	32.4	32.9	32.5
NI-OF23A-SDB6-DUR	100%	f	7.8	7.8	7.8	8.3	7.2	7.4	16.3	15.8	15.5	31.6	31.5	31.7
NI-BAY26-SDB6-DUR	100%	f	7.8	7.8	7.8	8.1	6.8	7.6	16.3	15.7	15.6	31.2	31.2	31.2
Scripps Control	0	f	7.9	7.7	7.7	7.5	6.8	7.6	15.1	15.8	15.3	28.2	28.0	27.8
Scripps Cu Ref. Tox.	2.9 µg/l	f	7.9	7.7	7.8	7.7	6.9	7.7	15.0	15.7	15.4	29.0	28.5	28.4
	4.1 µg/l	f	7.9	7.8	7.8	7.8	6.9	7.6	15.1	15.8	15.3	28.3	28.1	28.0
	5.9 µg/l	f	7.9	7.8	7.8	8.1	6.8	7.8	15.0	15.8	15.3	27.6	27.9	28.4
	8.4 µg/l	f	7.9	7.7	7.7	7.9	6.9	7.8	15.0	15.9	15.4	28.0	28.0	28.2
	12 µg/l	f	7.9	7.8	7.7	8.0	7.0	7.5	15.3	16.0	15.3	28.3	28.1	28.3
	17.2 µg/l	f	7.9	7.8	7.7	8.0	7.2	7.2	15.0	15.8	15.3	28.0	28.1	28.4
Brine Control 1	n/a	f	8.0	7.8	7.8	7.4	7.2	7.3	16.3	16.0	15.8	32.6	32.3	32.1
Brine Control 2	n/a	f	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND	ND

N - water quality not taken due to 100% mortality in treatment

ND - water quality not recorded

TIE2 – 03/19/2005

OUTFALLS

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NI-OF23A-TIE2-FF	25	100.0
	50	95.0
	100	65.0
NI-OF26-TIE2-FF	25	95.0
	50	100.0
	100	100.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NI-OF23A-TIE2-FF	25	100.0
	50	100.0
	100	65.0
NI-OF26-TIE2-FF	25	95.0
	50	100.0
	100	95.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	MEAN NORM DEV (%)
NI-OF23A-TIE2-FF	12.50	95.0
	25	24.0
	50	0.0
	57	0.0
NI-OF26-TIE2-FF	12.50	96.0
	25	94.0
	50	94.0
	69	89.0

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NI-BAY23A-TIE2-DUR	100	95.0
NI-BAY26-TIE2-DUR	100	100.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (%)	MEAN SURVIVAL (%)
NI-BAY23A-TIE2-DUR	100	100.0
NI-BAY26-TIE2-DUR	100	95.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	MEAN NORM DEV (%)
NI-BAY23A-TIE2-DUR	100	96.0
NI-BAY26-TIE2-DUR	100	95.0

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN SURVIVAL (%)
Scripps Control	n/a	100.0
Salt Control	n/a	100.0

MYSIDS (*A. bahia*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN SURVIVAL (%)
Scripps Control	n/a	100.0
Salt Control	n/a	95.0

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	MEAN NORM DEV (%)
Natural Seawater	n/a	96.0
Brine Control	n/a	95.0

Please refer to TIE II Report for raw data and water quality

^aControls (QA/QC) correspond to all samples from TIE2

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

SUMMARY RESULTS- QA/QC

COPPER REFERENCE TOXICANT TEST

SPECIES	DATE	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	4/6/2005	50	100	101.8	86.1-120.5
MYSIDS	5/19/2005	214.4	326	271.5	236.1-305.75
MUSSELS	3/19/2005	10	20.0	13.04	12.8-13.3

Reference Toxicant tests are within two standard deviations of Nautilus' control chart mean

SDB7 – 04/27/2005

OUTFALLS

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ¹	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB7-FF	12.5	a	5	100.0	100.0	0.0	100.0	n/a	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	5	100.0	95.0	10.0	95.0	0.196	No
		b	5	100.0					
		c	4	80.0					
		d	5	100.0					
	100	a	4	80.0	95.0	10.0	95.0	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NI-OF26-SDB7-FF	12.5	a	5	100.0	100.0	0.0	111.1	0.091	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	5	100.0	100.0	0.0	111.1	0.091	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	100	a	4	80.0	80.0	16.3	88.9	0.180	No
		b	5	100.0					
		c	3	60.0					
		d	4	80.0					
NI-OF26-SDB7-COMP	12.5	a	5	100.0	100.0	0.0	111.1	0.091	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
	50	a	5	100.0	90.0	20.0	100.0	0.500	No
		b	5	100.0					
		c	3	60.0					
		d	5	100.0					
	100	a	5	100.0	100.0	0.0	111.1	0.091	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

^aControls (QA/QC) correspond to all samples from SDB7

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

BAY SAMPLES

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (%)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-BAY23A-SDB7-PRE	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NI-BAY26-SDB7-PRE	100	a	4	80.0	90.0	11.5	94.7	0.269	No
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
NI-OF23A-SDB7-DUR	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					
NI-BAY26-SDB7-DUR	100	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (%)	REP	# NORMAL	# ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
NI-OF23A-SDB7-PRE	100.0	a	128	16	88.9	90.0	2.8	97.7	0.135	No
		b	106	17	86.2					
		c	148	17	89.7					
		d	122	11	91.7					
		e	116	8	93.5					
NI-BAY26-SDB7-PRE	100	a	133	2	98.5	96.8	2.1	105.0	0.012	Yes ^c
		b	124	2	98.4					
		c	128	4	97.0					
		d	112	4	96.6					
		e	126	9	93.3					
NI-OF23A-SDB7-DUR	100	a	117	17	87.3	91.8	3.2	99.7	0.443	No
		b	130	13	90.9					
		c	136	12	91.9					
		d	150	11	93.2					
		e	141	6	95.9					
NI-BAY26-SDB7-DUR	100	a	130	8	94.2	95.7	2.7	103.9	0.0393	Yes ^c
		b	139	3	97.9					
		c	142	1	99.3					
		d	98	7	93.3					
		e	104	7	93.7					

QA/QC SAMPLES^a

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	5	100.0	95.0	10.0	n/a	n/a	n/a
		b	5	100.0					
		c	5	100.0					
		d	4	80.0					
Salt Control	n/a	a	5	100.0	100.0	0.0	105.3	0.196	No
		b	5	100.0					
		c	5	100.0					
		d	5	100.0					

TOPSMELT (*A. affinis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP	SURVIVAL (#)	SURVIVAL (%)	MEAN SURVIVAL (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Copper Ref. Tox.	50	a	3	60.0	85.0	19.1	89.5	0.201	No
		b	4	80.0					
		c	5	100.0					
		d	5	100.0					
	100	a	5	100.0	75.0	19.1	78.9	0.065	No
		b	3	60.0					
		c	4	80.0					
		d	3	60.0					
	200	a	2	40.0	70.0	34.6	73.7	0.124	No
		b	5	100.0					
		c	5	100.0					
		d	2	40.0					
	400	a	0	0.0	25.0	25.2	26.3	0.004	Yes
		b	3	60.0					
		c	1	20.0					
		d	1	20.0					

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	CONC (% or µg/l Cu)	REP.	# NORMAL	ABNORMAL	NORM DEVEL (%)	MEAN NORM DEV (%)	STD DEV	% of CONTROL ²	P-VALUE ^b	SIG DIFF FROM CONTROL?
Scripps Control	n/a	a	153	12	92.7	92.1	2.4	100.0	n/a	n/a
		b	154	20	88.5					
		c	137	9	93.8					
		d	99	7	93.4					
Copper Ref. Tox.	2.9	a	120	18	87.0	82.7	8.5	89.8	0.034	Yes
		b	131	17	88.5					
		c	95	37	72.0					
		d	130	13	90.9					
		e	101	33	75.4					
	4.1	a	95	81	54.0	54.7	10.4	59.3	0.000	Yes
		b	87	50	63.5					
		c	-	-	-					
		d	-	-	-					
		e	-	-	-					
	5.9	a	0	131	0.0	2.3	3.2	2.5	0.000	Yes
		b	4	165	2.4					
		c	9	106	7.8					
		d	2	149	1.3					
		e	0	147	0.0					
	8.4	a	0	131	0.0	0.0	0.0	0.0	0.000	Yes
		b	0	135	0.0					
		c	0	151	0.0					
		d	0	154	0.0					
		e	0	137	0.0					

SUMMARY RESULTS- QA/QC
COPPER REFERENCE TOXICANT TEST

SPECIES	NOEC (µg/l)	LOEC (µg/l)	EC50 (µg/l)	95% C.L. (µg/l)
TOPSMELT	200	400.0	268.18	160.3-506.5
MUSSELS	2.9	4.1	4.30	3.78-4.69

Dash indicates no data (replicate was spilled or organisms not added)

^aControls (QA/QC) correspond to all samples from SDB7

^bStudent's t-test with a one tailed distribution and two sample unequal variance

^cp-value is significant because treatment had a significantly greater proportion normal compared to the control

n/a - t-test not used since control and treatment have same percentage survival

¹Controls were the Bay water samples taken prior to storm (PRE) with comparable sample ID

²Controls were Scripps filtered seawater

WATER QUALITY

TOPSMELT (*A. affinis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep	pH (SU)			Dissolved Oxygen (mg/l)			Temperature (°C)			Salinity (‰)										
			0	24	48	72	96	0	24	48	72	96	0	24	48	72	96					
NI-OF23A-SDB7-FF	12.5%	a	7.7	7.7	7.6	7.7	7.7	7.5	6.8	6.4	7.3	6.8	18.8	18.6	19.3	18.6	19.1	33.6	33.8	33.9	34.1	34.2
	50%	a	7.7	7.7	7.6	7.7	7.6	7.7	6.4	6.5	7.4	6.4	18.6	18.6	19.4	18.8	19.2	34.0	34.2	34.1	34.3	34.3
	100%	a	7.8	7.7	7.6	7.6	7.6	8.1	6.3	6.5	7.2	7.0	18.2	18.6	19.4	18.8	19.1	34.4	34.4	34.5	34.6	34.8
NI-OF26-SDB7-FF	12.5%	a	7.7	7.7	7.6	7.7	7.7	7.5	6.3	6.1	7.2	6.7	18.8	18.3	19.4	18.6	19.2	33.1	33.3	33.4	33.6	33.6
	50%	a	7.7	7.6	7.5	7.6	7.6	7.5	5.5	5.6	6.1	6.1	18.6	18.4	19.4	18.6	19.3	32.7	32.8	32.8	33.1	33.1
	100%	a	7.8	7.5	7.5	7.5	7.6	7.8	4.8	5.0	5.7	5.6	18.1	18.6	19.4	18.7	19.3	31.6	31.8	31.9	32.0	32.1
NI-OF26-SDB7-COMP	12.5%	a	7.8	7.7	7.6	7.7	7.7	7.6	6.4	6.3	7.0	6.6	19.3	18.2	19.3	18.3	19.1	33.4	33.3	33.4	33.6	33.6
	50%	a	7.9	7.7	7.6	7.7	7.7	7.8	6.3	6.0	7.0	6.8	19.1	18.1	19.3	18.2	18.9	32.8	32.8	32.9	33.1	33.3
	100%	a	8.0	7.8	7.6	7.7	7.6	8.1	5.9	6.0	6.6	6.5	18.6	18.0	19.1	18.1	18.8	32.0	32.2	32.5	32.5	32.7
NI-OF23A-SDB7-PRE	100%	a	7.7	7.7	7.7	7.7	7.8	7.5	6.7	6.3	7.4	6.5	19.0	18.6	19.4	18.7	19.3	33.7	33.9	33.9	34.1	33.9
NI-BAY26-SDB7-PRE	100%	a	7.7	7.7	7.7	7.7	7.8	7.3	6.6	6.5	7.2	6.5	19.3	18.3	19.3	18.4	19.3	33.2	33.8	33.9	34.1	34.0
NI-OF23A-SDB7-DUR	100%	a	7.7	7.7	7.7	7.7	7.6	7.3	6.6	6.4	7.3	5.8	18.4	18.3	19.3	18.6	19.0	33.4	33.4	33.4	33.2	33.1
NI-BAY26-SDB7-DUR	100%	a	7.7	7.5	7.7	7.8	7.7	7.5	6.9	6.6	7.3	6.1	18.4	18.3	19.3	18.6	19.1	33.5	33.5	33.5	33.4	33.4
Scripps Control	0	a	7.8	7.8	7.6	7.8	7.7	7.7	6.9	6.6	7.5	7.2	18.6	18.3	19.3	18.3	19.1	31.6	31.5	31.7	31.7	31.9
Scripps Cu Ref. Tox.	50 µg/l	a	7.8	7.8	7.6	7.8	7.8	7.8	7.0	6.7	7.7	7.4	18.8	18.3	19.3	18.3	19.0	31.5	31.5	31.6	31.6	31.6
	100 µg/l	a	7.8	7.8	7.6	7.8	7.8	7.7	7.0	6.6	7.7	7.6	18.8	18.3	19.3	18.4	19.1	31.5	31.5	31.6	31.5	31.6
	200 µg/l	a	7.8	7.8	7.6	7.8	7.7	7.7	7.1	6.6	7.9	7.4	18.8	18.2	19.1	18.4	19.0	31.5	31.6	31.8	31.9	31.6
	400 µg/l	a	7.9	7.8	7.7	7.8	7.7	7.7	7.2	7.1	7.9	7.3	18.8	18.3	19.1	18.6	19.0	31.5	31.6	31.8	31.8	31.6
Salt Control	n/a	a	7.9	7.7	7.5	7.6	7.5	7.4	6.1	6.4	7.0	6.6	19.4	18.3	19.2	18.6	19.0	31.4	31.5	31.6	31.4	31.5

MUSSELS (*M. galloprovincialis*)

SAMPLE ID	Effluent Concentration (% or µg/l Cu)	Rep.	pH (SU)			D.O. (mg/l)			Temperature (°C)			Salinity (‰)		
			0	24	48	0	24	48	0	24	48	0	24	48
NI-OF23A-SDB7-PRE	100%	f	7.7	7.8	7.7	7.8	8.6	8.4	16.0	15.3	15.1	33.9	33.9	33.8
NI-BAY26-SDB7-PRE	100%	f	7.7	7.8	7.7	8.0	8.4	8.3	16.0	15.2	15.6	33.6	33.3	33.0
NI-OF23A-SDB7-DUR	100%	f	7.7	7.8	7.8	7.9	8.7	8.5	16.0	15.2	14.9	33.5	33.5	33.5
NI-BAY26-SDB7-DUR	100%	f	7.7	7.8	7.8	8.0	8.3	8.4	16.0	15.3	15.0	33.7	33.4	33.3
Scripps Control	0	f	8.1	8.0	7.7	7.5	8.2	8.2	16.0	15.5	15.2	29.4	29.3	29.0
Scripps Cu Ref. Tox.	2.9 µg/l	f	7.9	7.9	7.9	7.2	8.3	8.5	15.8	15.2	15.0	29.5	29.6	29.7
	4.1 µg/l	f	7.8	7.8	7.7	7.9	8.4	8.4	15.8	15.2	15.3	29.5	29.6	29.7
	5.9 µg/l	f	7.8	7.8	7.7	7.8	8.3	8.4	15.9	15.2	15.4	29.5	29.6	29.6
	8.4 µg/l	f	7.8	7.8	7.7	7.9	8.6	8.2	15.9	15.2	15.2	29.6	29.5	29.6
	12 µg/l	f	7.8	7.8	7.7	7.9	8.5	8.3	15.8	15.0	15.1	29.6	29.5	29.6
	17.2 µg/l	f	7.8	7.8	7.8	7.7	8.2	8.5	15.8	15.0	15.1	29.6	29.6	29.7

Appendix D

Chemistry

Table of Data Qualifiers

Flag	Application
B	Analyte concentration found in the sample at a concentration <5x the level found in the procedural blank
D	Dilution Run. Initial run outside linear range of instrument
E	Estimate, result is greater than the highest concentration level in the calibration
H	Surrogate diluted out. Used when surrogate recovery is affected by excessive dilution of the sample extract
J	Analyte detected below the sample-specific Reporting Limit (RL)
ME	Significant matrix Interference – Estimated value
MI	Significant Matrix Interference – value could not be determined or estimated
n	Quality Control (QC) value is outside the accuracy or precision Data Quality Objective (DQO), but meets the contingency criteria
N	Quality Control (QC) value is outside the accuracy or precision Data Quality Objective (DQO)
NA	Not applicable
T	Holding Time (HT) exceeded
U	Analyte not detected at 3:1 signal:noise ratio. The sample-specific method detection limit (MDL) reported

Table of Acronyms

Acronym	Definition
CVAF	Cold Vapor Atomic Fluorescence
FIAS	Field Injection Atomic Spectroscopy
GFAA	Graphite Furnace-Atomic Absorption
ICP-MS	Inductively Coupled Plasma – Mass Spectrometry
ICP-OES	Inductively Coupled Plasma – Optical Emission Spectrometry
LCS	Laboratory Control Sample
MB	Matrix Blank
MSL	Battelle/Marine Sciences Laboratory
NS	Not Spiked
NS&T	National Standards and Trends
OPR	On-going Precision and Recovery
PB	Procedural Blank
PD	Percent Difference
RIS	Recovery Internal Standard
RPD	Relative Percent Difference
SRM	Standard Reference Material

Appendix D1

NAV

SDB1- 11/7/2002

SDB2- 2/24/2003

SDB4- 10/17/2004

SDB45- 10/26/2004

SDB1-11/7/2002

METALS

MSL	CLIENT CODE	PARAMETER	Ag		Cu		Pb		Hg		Zn
CHEMISTRY CODE			(µg/L)		(µg/L)		(µg/L)		(µg/L)		(µg/L)
1919-1	NAV-OF9-SDB1-COMP (T)	Total metals	0.215		123		23.5		0.0441		969
1919-13	NAV-OF9-SDB1-COMP (D)	Dissolved metals	0.0113	J	59.6		0.437		0.00676		776
1919-2	NAV-OF11-SDB1-DUR (T)	Total metals	0.100	J	136		6.50		0.266		407
1919-14	NAV-OF11-SDB1-DUR (D)	Dissolved metals	0.0250	J	58.8		0.536		0.0116		499
1919-3 R1	NAV-OF14-SDB1-DUR (T)	Total metals	0.0688	J	75.7		8.93		0.0732		573
1919-3 R2	NAV-OF14-SDB1-DUR (T)	Total metals	--		--		--		0.0732		--
1919-3 MEAN	NAV-OF14-SDB1-DUR (T)	Total metals	--		--		--		0.0732		--
1919-15	NAV-OF14-SDB1-DUR (D)	Dissolved metals	0.0125	J	51.9		0.602		0.0123		366
1919-4 R1	NAV-BAY9-SDB1-PRE (T)	Total metals	0.0225	J	4.53		0.235		0.00358		9.06
1919-4 R2	NAV-BAY9-SDB1-PRE (T)	Total metals	0.0275	J	4.50		0.223		--		9.42
1919-4 MEAN	NAV-BAY9-SDB1-PRE (T)	Total metals	0.0250	J	4.52		0.229		--		9.24
1919-16	NAV-BAY9-SDB1-DUR (D)	Dissolved metals	0.0225	J	3.42		0.0749	J	0.00457		8.55
1919-5	NAV-BAY9-SDB1-DUR (T)	Total metals	0.0163	J	4.91		0.200		0.00292		10.4
1919-17	NAV-BAY9-SDB1-DUR (D)	Dissolved metals	0.0175	J	3.52		0.0897	J	0.00457		10.7
1919-9	NAV-BAY9-SDB1-AFT (T)	Total metals	0.0188	J	4.79		0.151		0.00244		10.4
1919-21	NAV-BAY9-SDB1-AFT (D)	Dissolved metals	0.0175	J	4.10		0.0771	J	0.00322		10.8
1919-6	NAV-BAY11-SDB1-DUR (T)	Total metals	0.0163	J	4.72		0.219		0.00329		10.1
1919-18	NAV-BAY11-SDB1-DUR (D)	Dissolved metals	0.0301	J	3.55		0.0716	J	0.00459		10.9
1919-10	NAV-BAY11-SDB1-AFT (T)	Total metals	0.0238	J	4.59		0.140		0.00197		9.91
1919-22	NAV-BAY11-SDB1-AFT (D)	Dissolved metals	0.0163	J	3.52		0.0794	J	0.00284		10.3
1919-7	NAV-BAY14-SDB1-DUR (T)	Total metals	0.0200	J	5.21		0.217		0.00279		11.3
1919-19	NAV-BAY14-SDB1-DUR (D)	Dissolved metals	0.0213	J	3.66		0.0733	J	0.00403		10.1
1919-11 R1	NAV-BAY14-SDB1-AFT (T)	Total metals	0.0225	J	4.86		0.169		0.00214		10.3
1919-11 R2	NAV-BAY14-SDB1-AFT (T)	Total metals	--		--		--		0.00205		--
1919-11 MEAN	NAV-BAY14-SDB1-AFT (T)	Total metals	--		--		--		0.00210		--
1919-23	NAV-BAY14-SDB1-AFT (D)	Dissolved metals	0.0188	J	3.97		0.0847	J	0.00387		10.2
1919-8 R1	NAV-BAY14A-SDB1-DUR (T)	Total metals	0.0225	J	4.69		0.208		0.00228		10.2
1919-8 R2	NAV-BAY14A-SDB1-DUR (T)	Total metals	0.0213	J	4.69		0.201		--		8.90
1919-8 MEAN	NAV-BAY14A-SDB1-DUR (T)	Total metals	0.0219	J	4.69		0.205		--		9.55
1919-20	NAV-BAY14A-SDB1-DUR (D)	Dissolved metals	0.0188	J	3.59		0.0889	J	0.00331		9.28
1919-12	NAV-BAY14A-SDB1-AFT (T)	Total metals	0.0238	J	5.08		0.162		0.00174		10.4
1919-24	NAV-BAY14A-SDB1-AFT (D)	Dissolved metals	0.0125	J	3.93		0.0722	J	0.00328		9.76

METALS QA/QC

PROJECT: Contaminant Analyses of Storm water and San Diego Bay Seawater
PARAMETER: Metals
LABORATORY: Battelle Marine Sciences Laboratory, Sequim, Washington
MATRIX: Seawater

SAMPLE CUSTODY AND PROCESSING: Twenty-four seawater samples (12 total and 12 dissolved) for metals analysis were received on 11/13/02. All samples were received in good condition (i.e., all sample containers were intact). Dissolved metals were received in polyethylene containers, which are not suitable for Hg analysis. The client was informed of the potentially compromised dissolved Hg data.

Samples were assigned a Battelle Central File (CF) identification number (1919) and were entered into Battelle's log-in system.

The following lists information on sample receipt and processing activities:

Lab Sample IDs:	1907-1 through -24
Description:	Seawater/seawater samples
Collection dates	11/07/02, 11/09/02, 11/10/02
Laboratory arrival date	11/13/02
Cooler temperature on arrival	12.5°C ⁽¹⁾
CVAA analysis (Hg)	12/03/02, 12/04/02
Fe-Pd preconcentration	12/18/02
ICP-MS analysis (Cu, Pb, Zn)	12/30/02, 01/17/03 reruns
GFAA analysis (Ag)	12/31/02, 01/02/03

⁽¹⁾ Cooler temperature outside the acceptable range of 4±2°C, however samples were not compromised as they were already acidified, see note on Log-in Checklist.

DATA QUALITY OBJECTIVES:

Analyte	Analytical Method	Range of Recovery	Relative Precision	SRM Accuracy	Project Reporting Limits (µg/L) ^a	Achieved Detection Limits (µg/L) ^b
Ag	GFAA	75-125%	±20%	±20%	0.1	0.01
Cu	ICP-MS	75-125%	±20%	±20%	0.1	0.028
Hg	CVAF	75-125%	±20%	±20%	0.001	0.00016
Pb	ICP-MS	75-125%	±20%	±20%	0.1	0.0059
Zn	ICP-MS	75-125%	±20%	±20%	0.1	0.11

(a) Extracted from the statement of work

(b) Achieved MDLs from Fe/Pd seawater MDL, Ag from GFAA report, Hg from 2002 MDL Study

METHODS: Five metals were analyzed: silver (Ag), copper (Cu), lead (Pb), mercury (Hg), and zinc (Zn).

Samples were preconcentrated using iron (Fe) and palladium (Pd) according to Battelle SOP MSL-I-025, *Methods of Sample Preconcentration*, which is derived from EPA Method 1640. Samples were then submitted for analysis for Cu, Pb and Zn by inductively coupled plasma-mass spectrometry (ICP-MS) following Battelle SOP MSL-I-022, *Determination of Elements in Aqueous and Digestate Samples by ICP-MS*, derived from EPA Method 1638. Six samples identified as OF9-SDB1, OF11-SDB1, OF14-SDB1, both dissolved and total fractions, were analyzed at a ten fold dilution directly by ICP-MS for Cu and Zn, as the preconcentrated samples were outside the calibration range of the instrument.

These results are reported.

METHODS:

Ag was analyzed in the Fe-Pd preconcentrate by graphite furnace atomic absorption (GFAA) following Battelle SOP MSL-I-029, *Determination of Metals in Aqueous and Digestate Samples by GFAA*, derived from EPA Method 200.9.

Hg was analyzed directly (with no preconcentration step) using cold-vapor atomic fluorescence (CVAF) spectroscopy according to Battelle SOP MSL-I-013, *Total Mercury in Aqueous Samples by CVAF*, which directly follows EPA Method 1631.

All results were reported in units of $\mu\text{g/L}$. Data are not blank corrected.

HOLDING TIMES:

The recommended holding times for metals analyses are 28 days from sample collection for Hg analysis and 6 months for analysis of all other metals. All samples were analyzed within their respective holding times.

DETECTION LIMITS:

Laboratory-achieved detection limits reported are from the Fe/Pd seawater MDL for Cu, Pb and Zn; GFAA daily analysis for Ag; and the 2002 MDL Study for Hg. MDLs were less than target reporting limits for all metals. The data are flagged by the following criteria:

- U Analyte not detected above the laboratory achieved MDL, which is reported
- J Analyte detected above the MDL, but below the reporting limit
- # Data quality precision or accuracy outside the criteria of $\pm 20\%$ or recoveries outside criteria of $\pm 25\%$

METHOD BLANKS:

Three method blanks were analyzed for all metals. Blank concentrations for all metals were below or less than three times the project reporting limits. Data were not blank corrected.

**LABORATORY
CONTROL SAMPLE
ACCURACY:**

Four blank spike (LCS) samples were analyzed with the set of water samples. Recoveries were reported for samples spiked at approximately $0.005 \mu\text{g/L}$ for Hg and $10 \mu\text{g/L}$ for Ag, Cu, Pb, and Zn in the preconcentrated samples. LCS samples analyzed with the reanalysis of Cu and Zn by direct ICP-MS were spiked at $3.2 \mu\text{g/L}$ for Cu and $29.4 \mu\text{g/L}$ for Zn. LCS recoveries ranged from 82% to 118% and were within the QC acceptance criteria of 75% to 125% for all metals.

**MATRIX SPIKE
ACCURACY:**

Two samples were selected as a matrix spike sample for each analyte. Recoveries were reported for samples spiked at approximately $0.02 \mu\text{g/L}$ Hg and $10 \mu\text{g/L}$ for Ag, Cu, Pb, and Zn. Matrix spike recoveries ranged from 77% to 104% and were within QC acceptance criteria of 75%-125% recovery for all metals, except one replicate for Zn (64%). Acceptable accuracy for Zn analysis is demonstrated in the three alternate MS samples.

**REPLICATE
PRECISION:**

Replicate precision was assessed by duplicate sample analysis and expressed as the relative percent deviation (RPD) of replicate results. RPDs ranged from 0% to 20% and were within the QC acceptance criteria of 20% for all metals.

**STANDARD
REFERENCE
MATERIAL
ACCURACY:**

Three SRMs were analyzed with this set of samples. CASS-4, SLRS-3 and 1640 were analyzed for ICP-MS metals and SRM 1641d was analyzed for Hg. SRM accuracy was expressed as the percent difference (PD) between the measured and certified or laboratory consensus value within the certified range.

The SRM CASS-4 is a nearshore seawater reference material, which is not certified for Ag or Hg. The certified values for CASS-4 are generally less than five times the laboratory achieved detection limit, therefore not a useful indicator of data set accuracy. However, CASS-4 was analyzed with this set because a alternate seawater SRM is not available. Laboratory consensus values were determined as CASS-4 is certified at concentrations near the detection limit. Laboratory consensus values were determined from multiple analyses of CASS-4 conducted over the past year. The target QC acceptance criterion is 20% PD, which was achieved from either end of the certified or consensus value range for all metals, except Pb (153%, 33%) and one replicate of Zn (182%). Acceptable analysis accuracy for Pb and Zn is demonstrated in the four alternate SRM recoveries.

The SRM SLRS-3 is a riparian reference material, which is not certified for Ag or Hg. The percent differences ranged from 1% to 18% and were within the QC acceptance criteria of 20% for all certified metals.

SRM 1640 and 1641 are freshwater reference materials used as instrument check samples. Data accuracy for mercury is evaluated in SRM 1641. SRM percent differences ranged from 0% to 14% and were within QC acceptance criteria of 20% (PD) for all metals.

METALS QA/QC (CONT.)

MSL	CHEMISTRY CODE	CLIENT CODE	PARAMETER	Ag (µg/L)		Cu (µg/L)		Pb (µg/L)		Hg (µg/L)		Zn (µg/L)	
DETECTION LIMITS													
Project Reporting Limit				0.1		0.1		0.1		0.001		0.1	
Laboratory Achieved MDL ⁽¹⁾				0.010		0.028		0.0059		0.00016		0.11	
METHOD BLANKS													
blkr1				0.010	U	0.127		0.0429	J	0.000594	J	0.174	
blkr2				0.010	U	0.131		0.0248	J	0.000299	J	0.137	
blank trm r1 (reruns)						0.028	U					0.11	U
Mean				0.010	U	0.0953	J	0.0339	J	0.000446	J	0.140	
LABORATORY CONTROL SAMPLE (LCS) ACCURACY													
OPR120202run1 (Hg)	LCS R1			9.02		9.31		8.46		0.00523		10.1	
OPR120202run2 (Hg)	LCS R2			9.24		9.18		8.23		0.00532		9.87	
OPR120203run1 (Hg)	LCS R3 (reruns)			--		3.56		--		0.00593		32.3	
OPR120203run2 (Hg)	LCS R4 (reruns)			--		3.51		--		0.00590		32.4	
OPR120202run1 (Hg)	LCS R1		% Rec (10 or Hg 0.005 ppb)	90%		92%		84%		98%		99%	
OPR120202run2 (Hg)	LCS R2		% Rec (10 or Hg 0.005 ppb)	92%		90%		82%		100%		97%	
OPR120203run1 (Hg)	LCS R3 (reruns)		% Rec (3.2 Cu, 29.4 Zn or Hg 0.005 ppb)			110%				118%		110%	
OPR120203run2 (Hg)	LCS R4 (reruns)		% Rec (3.2 Cu, 29.4 Zn or Hg 0.005 ppb)			106%				112%		110%	
MATRIX SPIKE ACCURACY													
1919-4	NAV-BAY9-SDB1-PRE		Total metals	0.0225	J	4.53		0.235		0.00358		9.06	
1919-4 MS				NS		NS		NS		0.0210		NS	
1919-4 MSD				NS		NS		NS		0.0203		NS	
1919-4 MS			% Rec (0.017ppb)							102%			
1919-4 MSD			% Rec (0.017ppb)							98%			
1919-5	NAV-BAY9-SDB1-DUR		Total metals	0.0163	J	4.91		0.200		NS		10.4	
1919-5 MS				9.81		13.8		8.22		NS		18.8	
1919-5 MSD				10.35		14.4		8.46		NS		19	
1919-5 MS			% Rec (10ppb)	98%		89%		80%				84%	
1919-5 MSD			% Rec (10ppb)	103%		95%		83%				86%	
1919-7	NAV-BAY14-SDB1-DUR		Total metals	0.0200	J	5.21		0.217		0.00279		11.3	
1919-7 MS				NS		NS		NS		0.0198		NS	
1919-7 MSD				NS		NS		NS		0.0209		NS	
1919-7 MS			% Rec (0.018ppb)							97%			
1919-7 MSD			% Rec (0.018ppb)							100%			
1919-8	NAV-BAY14A-SDB1-DUR		Total metals	0.0225	J	4.69		0.208		0.00228		10.2	
1919-8 MS				NS		NS		NS		0.0217		NS	
1919-8 MSD				NS		NS		NS		0.0203		NS	
1919-8 MS			% Rec (0.018ppb)							105%			
1919-8 MSD			% Rec (0.018ppb)							102%			
1919-11	NAV-BAY14-SDB1-AFT		Total metals	0.0225	J	4.86		0.169		NS		10.3	
1919-11 MS				9.46		13		7.85		NS		18.9	
1919-11 MSD				10.4		13.9		8.28		NS		16.7	
1919-11 MS			% Rec (10ppb)	94%		81%		77%				86%	
1919-11 MSD			% Rec (10ppb)	104%		90%		81%				64%	#
REPLICATE PRECISION													
1919-3 R1	NAV-OF14-SDB1-DUR (T)		Total metals	0.0688	J	73.6		8.93		0.0732		588	
1919-3 R2	NAV-OF14-SDB1-DUR (T)		Total metals	--		--		--		0.0732		--	
1919-3 MEAN	NAV-OF14-SDB1-DUR (T)		Total metals	--		--		--		0.0732		--	
			RPD							0%			
1919-4r1	NAV-BAY9-SDB1-PRE		Total metals	0.0225	J	4.53		0.235				9.06	
1919-4r2	NAV-BAY9-SDB1-PRE		Total metals	0.0275	J	4.5		0.223				9.42	
1919-4 MEAN	NAV-BAY-SDB1-PRE		Total metals	0.0250	J	4.515		0.229				9.24	
			RPD	20%		1%		5%				4%	
1919-8r1	NAV-BAY14A-SDB1-DUR		Total metals	0.0225	J	4.69		0.208				10.2	
1919-8r2	NAV-BAY14A-SDB1-DUR		Total metals	0.0213	J	4.69		0.201				8.90	
1919-8 MEAN	NAV-BAY14A-SDB1-DUR		Total metals	0.0219	J	4.69		0.205				9.55	
			RPD	6%		0%		3%				14%	
1919-11 R1	NAV-BAY14-SDB1-AFT		Total metals	0.0225	J	4.86		0.169		0.00214		10.3	
1919-11 R2	NAV-BAY14-SDB1-AFT		Total metals	--		--		--		0.00205		--	
1919-11 MEAN	NAV-BAY14-SDB1-AFT		Total metals	--		--		--		0.00210		--	

METALS QA/QC (CONT.)

MSL			Ag	Cu	Pb	Hg	Zn
CHEMISTRY CODE	CLIENT CODE	PARAMETER	(µg/L)	(µg/L)	(µg/L)	(µg/L)	(µg/L)
STANDARD REFERENCE MATERIAL ACCURACY							
cass4r1			--	0.692	0.0541	J	1.16
cass4r2			--	0.676	0.0285	J	0.526
		SRM Certified or Laboratory Consensus Value	NC	0.592	0.0214	J	NC
		Range		±0.055	±0.01		±0.055
PD CASS-4 r1				17%	153%	#	182% #
PD CASS-4 r2				14%	33%	#	13%
slrs3r1			--	1.36	0.0747	J	1.20
slrs3r2			--	1.38	0.0691	J	1.23
		SRM Certified Value	NC	1.35	0.068	NC	1.04
		Range		±0.07	±0.007		±0.09
PD SLRS3 r1				1%	10%		15%
PD SLRS3 r2				2%	2%		18%
1640Direct R1 or 1641 R1 for Hg			--	86.5	26.8	1624	60.7
1640Direct R2 or 1641 R2 for Hg			--	76.6	24.0	1603	56.5
1640 TRM			--	86.0	--	--	53.0
certified value			--	85.2	27.9	1590	53.2
Range			--			±40	
PD 1640 Direct R1				2%	4%	2%	14%
PD 1640 Direct R2				10%	14%	1%	6%
PD 1640 TRM				1%			0%

(1)= Fe/Pd MDL Study, Ag from Graphite Furnace report, and Hg from 2002 MDL Study; NC = Analyte not certified; NS= Analyte not spike; # = Data quality outside the accuracy criteria of ±20% or precision/MS recovery criteria of ±25%; U= Analyte not detected above the laboratory achieved MDL, which is reported; J = Analyte detected above the MDL, but below the reporting limit.

PAHS

CLIENT SAMPLE ID										NAV- OF 9- SDB1- COMP	NAV- OF 11- SDB1- COMP	NAV- OF 14- SDB1- COMP	NAV- BAY9- SDB1- PRE	NAV- BAY9- SDB1- DUR	NAV- BAY9- SDB1-AFT	NAV- BAY11- SDB1- DUR	NAV- BAY11- SDB1-AFT	NAV- BAY14- SDB1- DUR	NAV- BAY14- SDB1-AFT	NAV- BAY14A- SDB1- DUR	NAV- BAY14A- SDB1-AFT
Battelle Sample ID	V9897	V9898	V9899	V9900	V9901	V9902	V9903	V9904	V9905	V9906	V9907	V9908									
Battelle Batch ID	02-634	02-634	02-634	02-634	02-634	02-634	02-634	02-634	02-634	02-634	02-634	02-634									
Associated Blank	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB	AB383PB									
QC Type	N	N	N	N	N	N	N	N	N	N	N	N									
Data File	A0409.D	A0412.D	A0413.D	A0414.D	A0416.D	A0417.D	A0418.D	A0419.D	A0422.D	A0421.D	A0423.D	A0423.D									
Extraction Date	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02	11/14/02									
Acquired Date	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02	11/26/02									
Matrix	Water	Water	Water	Water	Water	Water	Water	Water	Water	Water	Water	Water									
Sample Size	1 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L	2.64 L									
Dilution Factor	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667									
P/V	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL	1 mL									
Min Reporting Limit	16.7	6.31	6.31	6.29	7.58	6.34	6.34	6.34	6.34	6.34	6.29	6.31									
Amount Units	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L									
Naphthalene	11.70	5.47	5.47	2.07	17.10	1.58	1.58	1.58	1.58	1.58	1.58	1.58									
C1-Naphthalenes	7.61	2.76	2.76	3.34	27.00	0.51	0.51	0.51	0.51	0.51	0.51	0.51									
C2-Naphthalenes	1.33	0.50	0.50	0.50	43.50	0.51	0.51	0.51	0.51	0.51	0.51	0.51									
C3-Naphthalenes	1.33	0.50	0.50	0.50	19.00	0.51	0.51	0.51	0.51	0.51	0.51	0.51									
C4-Naphthalenes	1.33	0.50	0.50	0.50	5.66	0.51	0.51	0.51	0.51	0.51	0.51	0.51									
Biphenyl	0.83	0.32	0.32	0.31	4.13	0.31	0.31	0.31	0.31	0.31	0.31	0.31									
Acenaphthylene	3.70	1.27	1.27	0.76	2.10	0.76	0.76	0.76	0.76	0.76	0.76	0.76									
Acenaphthene	1.07	0.41	0.41	0.44	5.55	0.42	0.42	0.42	0.42	0.42	0.42	0.42									
Fluorene	5.51	2.41	2.41	2.18	4.07	1.78	1.78	1.78	1.78	1.78	1.78	1.78									
C1-Fluorenes	0.98	0.37	0.37	0.37	3.60	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
C2-Fluorenes	0.98	0.37	0.37	0.37	0.45	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
C3-Fluorenes	0.98	0.37	0.37	0.37	0.45	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
Phenanthrene	56.20	10.30	10.30	2.39	7.50	1.36	1.36	1.36	1.36	1.36	1.36	1.36									
Anthracene	3.08	1.98	1.98	0.26	3.07	0.76	0.76	0.76	0.76	0.76	0.76	0.76									
C1-Phenanthrenes/Anthracenes	36.20	11.80	11.80	0.26	5.76	0.27	0.27	0.27	0.27	0.27	0.27	0.27									
C2-Phenanthrenes/Anthracenes	105.00	35.80	35.80	0.26	0.32	0.27	0.27	0.27	0.27	0.27	0.27	0.27									
C3-Phenanthrenes/Anthracenes	89.90	16.30	16.30	0.26	0.32	0.27	0.27	0.27	0.27	0.27	0.27	0.27									
Dibenzothiophene	24.80	14.00	14.00	0.35	1.18	0.35	0.35	0.35	0.35	0.35	0.35	0.35									
C1-Dibenzothiophenes	64.20	33.40	33.40	0.35	5.55	0.35	0.35	0.35	0.35	0.35	0.35	0.35									
C2-Dibenzothiophenes	104.00	40.70	40.70	0.35	0.42	0.35	0.35	0.35	0.35	0.35	0.35	0.35									
C3-Dibenzothiophenes	113.00	29.30	29.30	0.35	0.42	0.35	0.35	0.35	0.35	0.35	0.35	0.35									
Fluoranthene	99.30	13.50	13.50	9.17	18.00	11.00	11.00	11.00	11.00	11.00	11.00	11.00									
Pyrene	129.00	17.00	17.00	4.66	9.91	5.85	5.85	5.85	5.85	5.85	5.85	5.85									
C1-Fluoranthenes/Pyrenes	52.30	10.70	10.70	0.36	8.47	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
C2-Fluoranthenes/Pyrenes	73.30	14.60	14.60	0.36	0.43	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
C3-Fluoranthenes/Pyrenes	84.30	12.10	12.10	0.36	0.43	0.37	0.37	0.37	0.37	0.37	0.37	0.37									
Benzo(a)anthracene	22.90	2.71	2.71	0.52	2.93	0.53	0.53	0.53	0.53	0.53	0.53	0.53									
Chrysene	81.60	10.40	10.40	0.28	4.57	0.28	0.28	0.28	0.28	0.28	0.28	0.28									
C1-Chrysenes	68.10	11.00	11.00	0.28	0.33	0.28	0.28	0.28	0.28	0.28	0.28	0.28									
C2-Chrysenes	118.00	0.28	0.28	0.28	0.33	0.28	0.28	0.28	0.28	0.28	0.28	0.28									
C3-Chrysenes	0.73	0.28	0.28	0.28	0.33	0.28	0.28	0.28	0.28	0.28	0.28	0.28									
C4-Chrysenes	0.73	0.28	0.28	0.28	0.33	0.28	0.28	0.28	0.28	0.28	0.28	0.28									
Benzo(b)fluoranthene	56.70	5.00	5.00	0.32	0.39	0.33	0.33	0.33	0.33	0.33	0.33	0.33									
Benzo(k)fluoranthene	42.20	3.41	3.41	0.33	0.39	0.33	0.33	0.33	0.33	0.33	0.33	0.33									
Benzo(e)pyrene	63.90	6.61	6.61	0.34	0.41	0.35	0.35	0.35	0.35	0.35	0.35	0.35									
Benzo(a)pyrene	52.90	4.16	4.16	0.49	0.59	0.50	0.50	0.50	0.50	0.50	0.50	0.50									
Perylene	26.70	0.53	0.53	0.53	0.63	0.53	0.53	0.53	0.53	0.53	0.53	0.53									
Indeno(1,2,3-cd)pyrene	64.60	4.00	4.00	0.69	0.83	0.70	0.70	0.70	0.70	0.70	0.70	0.70									
Dibenz(a,h)anthracene	10.80	0.79	0.79	0.79	0.95	0.80	0.80	0.80	0.80	0.80	0.80	0.80									
Benzo(g,h,i)perylene	91.80	6.20	6.20	0.43	0.51	0.43	0.43	0.43	0.43	0.43	0.43	0.43									
Total Priority Pollutant PAHs	731.78	92.22	92.22	26.67	74.80	27.25	27.25	27.25	27.25	27.25	27.25	27.25									
Total PAH	1763.10	331.29	331.29	30.01	198.65	58.99	58.99	58.99	58.99	58.99	58.99	58.99									

PAHs (CONT.)

CLIENT SAMPLE ID	NAV- OF9- SDB1- COMP	NAV- OF11- SDB1- COMP	NAV- OF14- SDB1- COMP	NAV- BAY9- SDB1-PRE	NAV- BAY9- SDB1- DUR	NAV- BAY9- SDB1-AFT	NAV- BAY11- SDB1- DUR	NAV- BAY11- SDB1-AFT	NAV- BAY14- SDB1- DUR	NAV- BAY14- SDB1-AFT	NAV- BAY14A- SDB1- DUR	NAV- BAY14A- SDB1-AFT
Surrogate Recoveries (%)												
Naphthalene-d8	56	60	56	56	52	57	60	56	47	52	61	54
Phenanthrene-d10	71	71	59	62	60	65	66	67	57	65	64	63
Chrysene-d12	81	77	66	72	73	77	79	83	72	73	78	74

PAHs QA/QC

PROJECT: Contamination Analysis of Stormwater and San Diego Seawater
PARAMETER: PAH
LABORATORY: Battelle, Duxbury, MA
MATRIX: Waters
SAMPLE CUSTODY: Water samples were collected between November 7 – 10, 2002, shipped on November 12, 2002, and received at Battelle Duxbury on November 13, 2002. All samples were received in good condition. The cooler temperature on arrival was 0.8°C. Samples were stored refrigerated until processing.

QA/QC DATA QUALITY OBJECTIVES:

	Reference Method	Surrogate Recovery	Procedural Blank	LCS/MS Recovery	MS/MSD Relative Precision	Achieved Detection Limit (ng/L)
PAH	General NS&T	40-120% Recovery	Less than 5X MDL	40-120% Recovery (for at least 80% of analytes; analyte conc. in MS must be >10x background)	►30% RPD (analyte conc. in MS must be >10x background)	PAH Naphthalene ~ 17.0 Other PAHs ~0.5

METHOD: Water samples were extracted for PAHs following general NS&T methodologies. A volume of ~2 L of sample was extracted three times with dichloromethane using separatory funnel techniques. The combined extract was dried over anhydrous sodium sulfate, concentrated and processed through alumina column and HPLC/GPC. The extract was concentrated, fortified with RIS and submitted for GC/MS analysis. Water extracts were analyzed directly using gas chromatography/mass spectrometry (GC/MS) following general NS&T methods. Sample data were quantified by the method of internal standards, using the Recovery Internal Standard (RIS) compounds.

HOLDING TIMES: Water samples for PAH were stored refrigerated until extraction. There is a 7-day holding time associated with these samples.

Samples were prepared for analysis in one analytical batch and were extracted within 7 days of sample collection and analyzed within 40 days of extraction.

<u>Batch</u>	<u>Extraction Date</u>	<u>Analysis Date</u>
02-634	11/14/02	11/26/02 - 11/26/02

BLANKS: A procedural blank (PB) was prepared with each analytical batch. Blanks were analyzed to ensure the sample extraction and analysis methods were free of contamination.

02-634 – The blank was void of contamination.

Comments – None.

LABORATORY CONTROL SAMPLE: A laboratory control sample (LCS) was prepared with each analytical batch. The percent recoveries of PAH analytes were calculated to measure data quality in terms of accuracy.

02-634 – All PAHs were recovered within the laboratory control limits specified by the method (40 – 120%) ranging from 58 – 82% recovery.

Comments – None.

MATRIX SPIKES:

A matrix spike (MS) sample was prepared with each analytical batch. The percent recoveries of PAH analytes were calculated to measure data quality in terms of accuracy.

02-634 – All PAHs were recovered within the laboratory control limits specified by the method (40 – 120%) and ranged from 61 – 89% recovery for the matrix spike. All PAHs were recovered within the laboratory control limits specified by the method (40 – 120%) and ranged from 59 – 86% recovery for the matrix spike duplicate.

Comments – None.

SURROGATES:

Three surrogate compounds were added prior to extraction, including Naphthalene-d8, Phenanthrene-d10, and Chrysene-d12. The recovery of each surrogate compound was calculated to measure data quality in terms of accuracy (extraction efficiency).

02-634 – Surrogate recovery for all PAH surrogate compounds were within the laboratory control limits specified by the method (40 – 120% recovery).

Comments – None.

REPLICATES:

Replicate samples for waters were prepared with each analytical batch as an MS and MSD. The RPD between replicate analyses for PAH analytes is calculated to measure data quality in terms of precision.

02-634 – All PAH analytes were recovered within the laboratory control limits specified by the method (<30%) and ranged from 0 – 6.6 % RPD.

Comments – None.

PAHs QA/QC (CONT.)

CLIENT SAMPLE ID	LAB CONTROL SAMPLE			MATRIX SPIKE- NAV-OF9-SDB1- COMP	MATRIX SPIKE DUPLICATE- NAV- OF9-SDB1-COMP	PROCEDURAL BLANK		
Battelle Sample ID	AB384LCS			V9897MS	V9897MSD	AB383PB		
Battelle Batch ID	02-634			02-634	02-634	02-634		
Associated Blank	AB383PB			AB383PB	AB383PB	NA		
QC Type	LCS			MS	MSD	PB		
Data File	A0408.D			A0410.D	A0411.D	A0407.D		
Extraction Date	14-Nov-02			11/14/02	11/14/02	14-Nov-02		
Acquired Date	26-Nov-02			11/26/02	11/26/02	26-Nov-02		
Matrix	Water			Water	Water	Water		
Sample Size	2 L			0.81 L	0.81 L	2 L		
Dilution Factor	1.667			1.667	1.667	1.667		
PIV	1 mL			1 mL	1 mL	1 mL		
Min Reporting Limit	8.34			20.6	20.6	8.34		
Amount Units	ng/L		Rec% Q	ng/L	ng/L	ng/L		
Naphthalene	291.00		58	763.00	744.00	0.66		U
C1-Naphthalenes	0.66 U	NA		950.00	942.00	0.66		U
C2-Naphthalenes	0.66 U	NA		1.64 U	1.64 U	0.66		U
C3-Naphthalenes	0.66 U	NA		1.64 U	1.64 U	0.66		U
C4-Naphthalenes	0.66 U	NA		1.64 U	1.64 U	0.66		U
Biphenyl	313.00		62	833.00	833.00	0.42		U
Acenaphthylene	309.00		62	854.00	828.00	0.41		U
Acenaphthene	321.00		64	885.00	863.00	0.54		U
Fluorene	332.00		66	962.00	925.00	0.49		U
C1-Fluorenes	0.49 U	NA		1.21 U	1.21 U	0.49		U
C2-Fluorenes	0.49 U	NA		1.21 U	1.21 U	0.49		U
C3-Fluorenes	0.49 U	NA		1.21 U	1.21 U	0.49		U
Phenanthrene	370.00		74	1090.00	1060.00	0.38		U
Anthracene	330.00		66	948.00	909.00	0.35		U
C1-Phenanthrenes/Anthracenes	0.35 U	NA		70.00	56.00	0.35		U
C2-Phenanthrenes/Anthracenes	0.35 U	NA		140.00	141.00	0.35		U
C3-Phenanthrenes/Anthracenes	0.35 U	NA		132.00	122.00	0.35		U
Dibenzothiophene	3.84 J	NA		43.00	38.80	0.46		U
C1-Dibenzothiophenes	0.46 U	NA		86.90	78.00	0.46		U
C2-Dibenzothiophenes	0.46 U	NA		131.00	118.00	0.46		U
C3-Dibenzothiophenes	0.46 U	NA		131.00	120.00	0.46		U
Fluoranthene	399.00		80	1160.00	1120.00	0.44		U
Pyrene	397.00		79	1180.00	1150.00	0.48		U
C1-Fluoranthenes/Pyrenes	0.48 U	NA		64.60	72.10	0.48		U
C2-Fluoranthenes/Pyrenes	0.48 U	NA		98.80	105.00	0.48		U
C3-Fluoranthenes/Pyrenes	0.48 U	NA		97.90	100.00	0.48		U
Benzo(a)anthracene	399.00		80	1070.00	1050.00	0.69		U
Chrysene	392.00		78	1120.00	1090.00	0.37		U
C1-Chrysenes	0.37 U	NA		88.30	90.60	0.37		U
C2-Chrysenes	0.37 U	NA		151.00	153.00	0.37		U
C3-Chrysenes	0.37 U	NA		0.90 U	0.90 U	0.37		U
C4-Chrysenes	0.37 U	NA		0.90 U	0.90 U	0.37		U
Benzo(b)fluoranthene	404.00		81	1120.00	1060.00	0.43		U
Benzo(k)fluoranthene	423.00		85	1100.00	1100.00	0.43		U
Benzo(e)pyrene	369.00		75	1020.00	969.00	0.45		U
Benzo(a)pyrene	380.00		76	1010.00	965.00	0.65		U
Perylene	343.00		69	972.00	920.00	0.70		U
Indeno(1,2,3-c,d)pyrene	394.00		79	1060.00	1020.00	0.91		U
Dibenz(a,h)anthracene	412.00		82	1110.00	1050.00	1.04		U
Benzo(g,h,i)perylene	313.00		63	905.00	853.00	0.57		U
Total Priority Pollutant PAHs				16337.00	15787.00			
Surrogate Recoveries (%)								
Naphthalene-d8	61			60	61	61		
Phenanthrene-d10	66			77	73	66		
Chrysene-d12	79			85	82	81		

PCBs

CLIENT SAMPLE ID:	NAV-OF9-SDB1-COMP		NAV-OF11-SDB1-COMP		NAV-OF14-SDB1-COMP	
Battelle Sample ID:	V9897		V9898		V9899	
Client Description:	Seawater/ Stormwater		Seawater/ Stormwater		Seawater/ Stormwater	
Battelle Batch ID:	02-634		02-634		02-634	
Sample Volume (L):	1.000		2.640		2.630	
Units:	ng/L		ng/L		ng/L	
CI2 08	2.197		0.100	U	0.100	U
CI3 18	0.100	U	0.100	U	0.100	U
CI3 28	0.969		0.100	U	0.100	U
CI4 44	1.484		0.100	U	0.243	
CI4 49	2.929		0.100	U	0.100	U
CI4 52	2.274		0.100	U	0.100	U
CI4 66	0.812		0.100	U	0.100	U
CI5 77	0.100	U	0.100	U	0.100	U
CI5 87	1.110		0.195		0.376	
CI5 101	1.880		0.517		0.624	
CI5 105	0.600		0.272		0.296	
CI5 118	1.617		0.473		0.383	
CI6 126	0.100	U	0.100	U	0.100	U
CI6 128	1.010		0.155		0.348	
CI6 138	3.824		0.568		1.472	
CI6 153	3.627		0.514		1.625	
CI6 156	0.100	U	0.100	U	0.100	U
CI7 169	0.100	U	0.100	U	0.100	U
CI7 170	1.285		0.100	U	0.264	
CI7 180	3.166		0.212		1.026	
CI7 183	1.317		0.147		0.344	
CI7 184	0.100	U	0.100	U	0.100	U
CI7 187	1.382		0.096	J	0.636	
CI8 195	0.432		0.058	J	0.139	
CI9 206	0.182		0.047	J	0.120	
CI10 209	1.472		0.100	U	0.100	U
Total PCB	33.568		3.253		7.896	
Surrogate Recoveries:						
CI3(34)	84		84		69	
CI5(112)	87		82		70	

PCBs QA/QC

PROJECT: Contamination Analysis of Stormwater and San Diego Seawater
PARAMETER: PCB
LABORATORY: Battelle, Duxbury, MA
MATRIX: Waters
SAMPLE CUSTODY: Water samples were collected between November 7 – 10, 2002, shipped on November 12, 2002, and received at Battelle Duxbury on November 13, 2002. All samples were received in good condition. The cooler temperature on arrival was 0.8 C. Samples were stored refrigerated until processing.

QA/QC DATA QUALITY OBJECTIVES:

	Reference Method	Surrogate Recovery	Procedural Blank	LCS/MS Recovery	MS/MSD Relative Precision	Achieved Detection Limit (ng/L)
PCB	General NS&T	40-120% Recovery	Less than 5X MDL	40-120% Recovery (for at least 80% of analytes; analyte conc. in MS must be >10x background)	▶30% RPD (analyte conc. in MS must be >10x background)	PCB ~ 0.1

METHOD: Water samples were extracted for PCBs following general NS&T methodologies. A volume of ~2 L of sample was extracted three times with dichloromethane using separatory funnel techniques. The combined extract was dried over anhydrous sodium sulfate, concentrated and processed through alumina column and HPLC/GPC. The extract was concentrated, fortified with RIS, solvent exchanged, and submitted for GC/ECD analysis. Water extracts were analyzed directly using gas chromatography/electron capture detector (GC/ECD) following general NS&T methods. Sample data were quantified by the method of internal standards, using the Recovery Internal Standard (RIS) compounds.

HOLDING TIMES: Water samples for PCB were stored refrigerated until extraction. There is a 7-day holding time associated with these samples.

Samples were prepared for analysis in one analytical batch and were extracted within 7 days of sample collection.

Batch	Extraction Date	Analysis Date
02-634	11/14/02	12/11/02 -1/7/03

BLANKS: The original instrumental runs of the procedural blank in December yielded a cross-contamination of the procedural blank due to a calibration standard run just prior to the procedural blank. Archive extracts of the blank and samples were run in January (outside the 40 day extract holding time), and the blank was void of contamination. This data is reported. Therefore the extract analysis was outside the 40-day holding time for extracts. The QC data is acceptable so there is no impact on the reported data.

A procedural blank (PB) was prepared with each analytical batch. Blanks were analyzed to ensure the sample extraction and analysis methods were free of contamination.

02-634 – No analytes were detected in the blank.

LABORATORY CONTROL **Comments – None.**
 A laboratory control sample (LCS) was prepared with each analytical batch. The percent recoveries of PCB analytes were calculated to measure data quality in terms of accuracy.

SAMPLE:

02-634 – All PCBs were recovered within the laboratory control limits specified by the method (40 – 120 and ranged from 51 – 108% recovery).

Comments – None.

MATRIX SPIKES:

A matrix spike (MS) sample was prepared with each analytical batch. The percent recoveries of PCB analytes were calculated to measure data quality in terms of accuracy.

02-634 – All PCBs were recovered within the laboratory control limits specified by the method (40 – 120%) and ranged from 58 – 115% recovery for the matrix spike. All PCBs were recovered within the laboratory control limits specified by the method (40 – 120%) and ranged from 56 – 105% recovery for the matrix spike duplicate.

Comments – None.

SURROGATES:

Two surrogate compounds were added prior to extraction, including PCB34 and PCB112. The recovery of each surrogate compound was calculated to measure data quality in terms of accuracy (extraction efficiency).

02-634 – Surrogate recovery for all PCB surrogate compounds was within the laboratory control limits specified by the method (40 – 120% recovery).

Comments – None.

REPLICATES:

Replicate samples for waters were prepared with each analytical batch as an MS and MSD. The RPD between replicate analyses for PCB analytes is calculated to measure data quality in terms of precision.

02-634 – All PCB analytes were recovered within the laboratory control limits specified by the method (<30%) and ranged from 0 – 5 % RPD.

Comments – None.

PCBs QA/QC (CONT.)

CLIENT SAMPLE ID:	LAB CONTROL SAMPLE	MATRIX SPIKE-NAV-OF9-SDB1-COMP	MATRIX SPIKE-DUPLICATE-NAV-OF9-SDB1-COMP	PROCEDURAL BLANK
Battelle Sample ID:	AB384LCS	V9897MS	V9897MSD	AB383PB
Client Description:	NA	Seawater/Stormwater	Seawater/Stormwater	NA
Battelle Batch ID:	02-634	02-634	02-634	02-634
Sample Volume (L):	NA	0.810	0.810	2.000
Units:	ng	% Recovery	% Recovery	ng/L
C12 08	32.495	108	38.761	0
C13 18	15.448	51	20.784	4
C13 28	21.306	71	30.401	6
C14 44	23.283	78	31.348	6
C14 49	22.610	75	33.163	6
C14 52	23.756	79	32.879	6
C14 66	22.233	74	30.833	6
C15 77	29.721	99	39.109	9
C15 87	26.402	88	36.024	1
C15 101	23.748	79	32.124	3
C15 105	24.910	83	34.363	2
C15 118	25.245	84	32.764	3
C16 126	23.713	79	34.350	2
C16 128	25.318	84	33.682	4
C16 138	25.607	85	36.781	1
C16 153	25.238	84	37.542	3
C16 156	NS	NA	NS	NA
C17 169	26.460	88	35.542	2
C17 170	25.218	84	34.149	3
C17 180	25.032	83	41.735	3
C17 183	25.920	86	34.618	4
C17 184	28.021	93	36.625	3
C17 187	25.135	84	34.066	3
C18 195	23.998	80	32.072	2
C19 206	17.842	59	22.982	5
C10 209	22.546	75	29.570	5
Total PCB	611.206	NA	836.266	NA
Surrogate Recoveries:				
C13(34)	70	80	57	75
C15(112)	82	88	85	83

TSS

SAMPLE LABEL	TSS (mg/L)
NAV-OF9-SDB1-COMP	170.267
NAV-OF11-SDB1-COMP	122.600
NAV-OF14-SDB1-COMP	126.800
NAV-BAY-PRE	0.72
NAV-BAY9-SDB1-PRE	1.44
NAV-BAY9-SDB1-DUR	0.72
NAV-BAY9-SDB1-AFT	0.65
NAV-BAY11-SDB1-PRE	1.38
NAV-BAY11-SDB1-DUR	1.00
NAV-BAY11-SDB1-AFT	0.52
NAV-BAY14-SDB1-PRE	0.85
NAV-BAY14-SDB1-DUR	1.21
NAV-BAY14-SDB1-AFT	0.65
NAV-BAY14A-SDB1-PRE	1.18
NAV-BAY14A-SDB1-DUR	0.93
NAV-BAY14A-SDB1-AFT	0.67

SDB2- 2/24/2003

METALS

MSL Code	Instrument: Sponsor ID:	GFAA Ag (µg/L)	ICP-MS Al (µg/L)	FIAS As (µg/L)	ICP-MS Cd (µg/L)	ICP-MS Cr (µg/L)	ICP-MS Cu (µg/L)	ICP-MS Fe (µg/L)	CVAF Hg (µg/L)	ICP-MS Mn (µg/L)	ICP-MS Ni (µg/L)	ICP-MS Pb (µg/L)	FIAS Se (µg/L)	ICP-MS Sr (µg/L)	ICP-MS Zn (µg/L)
1979-7	NAV-PR6-SDB2-FF (T)	0.192 J	320 J	1.49	5.49	3.33	84.7	515	0.00555 J	22.4	7.72	20.5	0.671	0.715	521
1979-22	NAV-PR6-SDB2-FF (T)	0.0150 J	39.6 J	1.23	4.97	1.30	69.4	22.9	0.00273 J	14.4	5.22	11.8	0.367	0.0859 J	458
1979-9	NAV-PR6-SDB2-COMP (T)	0.247 J	1025 J	1.78	2.27	7.19	104	1417	0.0213 J	31.5	11.2	23.4	0.102 J	1.13	391
1979-24	NAV-PR6-SDB2-COMP (T)	0.00809 J	15.1 J	1.18	0.303	1.12	14.2	17.6	0.00219 J	5.94	1.88	0.533	0.247	0.0603 J	80.8
1979-8	NAV-PR6-SDB2-FF (T)	0.0522 J	179 J	1.66	1.37	4.24	183	428	0.0188 J	84.2	17.2	4.06	1.08	0.205 J	314
1979-23	NAV-PR6-SDB2-FF (T)	0.0286 J	30.4 J	1.41	1.23	3.58	177	161	0.0133 J	81.5	17.2	0.879	1.33	0.289 J	288
1979-10	NAV-PR6-SDB2-COMP (T)	0.132 J	722 J	1.36	1.12	6.62	66.2	1149	0.0189 J	32.9	7.33	14.6	0.161 J	0.816	249
1979-25	NAV-PR6-SDB2-COMP (T)	0.0119 J	32.2 J	1.04	0.265	1.69	33.0	27.5	0.00412 J	14.1	4.11	0.281	0.257	0.101 J	78.2

MSL Code	Instrument:	GFAA	ICP-MS	FIAS	ICP-MS	Cd (µg/L)	ICP-MS	Cr (µg/L)	ICP-MS	Cu (µg/L)	ICP-MS	Fe (µg/L)	CVAF	ICP-MS	Mn (µg/L)	ICP-MS	Ni (µg/L)	ICP-MS	Pb (µg/L)	ICP-MS	Se (µg/L)	FIAS	ICP-MS	Sr (µg/L)	ICP-MS	Zn (µg/L)
1979-1	Sponsor ID:	Ag (µg/L)	Al (µg/L)	As (µg/L)	Cd (µg/L)	Cr (µg/L)	Cu (µg/L)	Fe (µg/L)	Hg (µg/L)	Mn (µg/L)	Ni (µg/L)	Pb (µg/L)	Se (µg/L)	Sn (µg/L)	Th (µg/L)	U (µg/L)	V (µg/L)	W (µg/L)	X (µg/L)	Y (µg/L)	Zn (µg/L)					
1979-16	NAV-OF9-SDB2-COMP (D)	0.0203 J	16.6 J	0.695	0.388	1.22	25.8	18.5	0.00367 J	28.7	6.95	0.369	0.132 J	0.165 J	218											
1979-4	NAV-OF9-SDB2-COMP (D)	0.185 J	1050 J	1.42	0.659	8.56	36.1	1610	0.0151 J	50.0	7.83	15.9	0.0860 J	1.12	233											
1979-19	NAV-OF9-SDB2-COMP (D)	0.010 J	8.25 J	0.820	0.366	2.32	9.88	62.1	0.00186 J	11.8	2.83	0.156	0.0352 J	0.286 J	112											
1979-2	NAV-OF11-SDB2-COMP (D)	0.0175 J	1690 J	1.18	1.23	5.55	68.4	2290	0.0508 J	34.9	5.27	22.4	0.169 J	0.924	555											
1979-17	NAV-OF11-SDB2-COMP (D)	0.0293 J	18.0 J	1.33	0.776	4.70	46.9	1390	0.0541 J	55.5	14.1	0.108 J	0.872	298												
1979-5	NAV-OF11-SDB2-COMP (D)	0.0130 J	18.7 J	0.814	0.669	1.20	15.1	68.7	0.00314 J	25.1	2.21	0.247	0.0703 J	0.227 J	179											
1979-20	NAV-BAY11-SDB2-COMP (D)	0.0293 J	74.9 J	1.15	0.105	1.96	4.73	129	0.00216 J	10.7	2.06	0.428	0.0435 J	0.201 J	23.5											
1979-11	NAV-BAY11-SDB2-COMP (D)	0.010 J	13.7 J	1.13	0.100	0.219 J	3.16	88.5	0.00973 J	9.01	1.17	0.0789	0.0352 J	0.228 J	21.6											
1979-26	NAV-OF14-SDB2-COMP (D)	0.229 J	2640 J	2.92	2.59	13.7	72.6	3940	0.0536 J	131	15.7	43.8	0.149 J	1.44	797											
1979-3	NAV-OF14-SDB2-COMP (D)	0.0267 J	10.5 J	0.781	0.983	0.804 J	22.1	18.6	0.00374 J	31.5	5.78	0.916	0.0873 J	0.0945 J	310											
1979-18	NAV-OF14-SDB2-COMP (D)	0.0860 J	1270 J	2.02	0.673	7.24	26.9	1870	0.0314 J	56.5	5.34	15.0	0.0352 J	0.345 J	200											
1979-21	NAV-OF14-SDB2-COMP (D)	0.010 J	39.9 J	1.24	0.533	1.73	7.23	70.8	0.00177 J	15.9	1.80	0.330	0.0352 J	0.124 J	110											
1979-12	NAV-BAY14-SDB2-COMP (D)	0.0324 J	107 J	1.17	0.109	1.75	5.01	152	0.00229 J	12.5	1.93	0.623	0.0539 J	0.253 J	24.7											
1979-27	NAV-BAY14-SDB2-COMP (D)	0.0111 J	2.32 J	1.11	0.106	0.242 J	3.53	125	0.00102 J	10.0	1.21	0.137	0.0640 J	0.235 J	24.9											

METALS (CONT.)

MSL CHEMISTRY CODE	CLIENT CODE	PARAMETER	GFAA Ag (µg/L)	TCP-MS Cu (µg/L)	TCP-MS Pb (µg/L)	CVAF Hg (µg/L)	TCP-MS Zn (µg/L)
1919-25	NAV-BAY9-SDB2-PRE (T)	Total metals	0.0254 J	5.70	0.0828 J	0.00102	11.8
1919-39	NAV-BAY9-SDB2-PRE (D)	Dissolved metals	0.0227 J	3.89	0.0541 J	0.0130	5.23
1919-29	NAV-BAY9-SDB2-DUR (T)	Total metals	0.0581 J	6.13	0.0602 J	0.00904	12.7
1919-43	NAV-BAY9-SDB2-DUR (D)	Dissolved metals	0.0138 J	3.92	0.0592 J	0.00134	3.09
1919-31 r1	NAV-BAY9-SDB2-AFT (T)	Total metals	0.0241 J	5.00	0.0988 J	0.000979 J	16.0
1919-31 r2	NAV-BAY9-SDB2-AFT (T)	Total metals	---	---	0.0892 J	0.00123	15.0
1919-45	NAV-BAY9-SDB2-AFT (D)	Dissolved metals	0.0226 J	4.25	0.100	0.000876 J	14.8
1919-26	NAV-BAY11-SDB2-PRE (T)	Total metals	0.0310 J	5.82	0.516	0.00210	16.5
1919-40	NAV-BAY11-SDB2-PRE (D)	Dissolved metals	0.0329 J	3.74	0.318	0.00129	14.9
1919-32	NAV-BAY11-SDB2-AFT (T)	Total metals	0.0151 J	5.10	0.151	---	12.5
1919-46	NAV-BAY11-SDB2-AFT(D)	Dissolved metals	0.0303 J	4.13	0.629	0.00227	20.5
1919-27	NAV-BAY14-SDB2-PRE (T)	Total metals	0.0368 J	4.86	0.0541 J	0.00139	18.2
1919-41	NAV-BAY14-SDB2-PRE (D)	Dissolved metals	0.0246 J	3.87	0.0772 J	0.000830 J	15.2
1919-33	NAV-BAY14-SDB2-AFT (T)	Total metals	0.0196 J	5.24	0.110	0.00101	4.82
1919-47	NAV-BAY14-SDB2-AFT (D)	Dissolved metals	0.0240 J	4.24	0.106	---	4.79
1919-28 r1	NAV-BAY14A-SDB2-PRE (T)	Total metals	0.0235 J	4.97	0.159	0.00158	10.6
1919-28 r2	NAV-BAY14A-SDB2-PRE (T)	Total metals	0.0283 J	5.00	0.116	0.00115	2.83
1919-42	NAV-BAY14A-SDB2-PRE (D)	Dissolved metals	0.0312 J	4.31	0.558	0.0101	14.2
1919-30	NAV-BAY14A-SDB2-DUR (T)	Total metals	0.0280 J	5.89	0.293	0.00166	14.9
1919-44	NAV-BAY14A-SDB2-DUR (D)	Dissolved metals	0.0178 J	3.80	0.290	0.00134	13.8
1919-34	NAV-BAY14A-SDB2-AFT (T)	Total metals	0.0201 J	4.95	---	0.00129	---
1919-48	NAV-BAY14A-SDB2-AFT (D)	Dissolved metals	0.0186 J	3.95	0.293	0.00176	15.8

METALS QA/QC

PROJECT: SPAWARS Task 11, San Diego Bay Stormwater
PARAMETER: Metals
LABORATORY: Battelle Marine Sciences Laboratory, Sequim, Washington
MATRIX: Seawater and Freshwater

SAMPLE CUSTODY AND PROCESSING: Eighteen seawater and twelve freshwater samples were received in on 03/03/03. All samples were received in good condition (i.e., all sample containers were intact). Samples were assigned a Battelle Central File (CF) identification number (1979) and were entered into Battelle's sample tracking system.

The following lists information on sample receipt and processing activities:

Chemistry Lab ID	1979-1 through -30
Collection dates	02/25/03
Laboratory arrival dates	03/03/03
Cooler temperatures, on arrival	NA – Samples arrived preserved
Fe/Pd Preconcentration (seawater)	03/14/03
FIAS (As – seawater)	03/14/03
FIAS (Se – seawater)	03/17/03
GFAA (Ag – seawater)	03/20/03
CVAA analyses (Hg)	03/13/03, 03/14/03, 03/18/03
ICP-MS analyses:	
Fe/Pd Seawater (Cd, Cr, Cu, Ni, Pb)	03/18/03
Direct Seawater (Al, Fe, Mn, Sn, Zn)	03/27/03
Freshwater (Ag, Al, As, Cd, Cr, Cu, Fe, Mn, Ni, Pb, Se, Sn, Zn)	03/24/03
Rerun Freshwater (Al, Fe)	04/11/03

QA/QC DATA QUALITY OBJECTIVES:

Analyte	Analytical Method Seawater	Analytical Method Freshwater	Range of Recovery	SRM Accuracy	Relative Precision	Detection Limits (µg/L)		
						Target MDL ⁽¹⁾	Achieved MDL Seawater ⁽²⁾	Achieved MDL Freshwater ⁽²⁾
Silver	GFAA	ICP-MS	50-150%	≤20%	≤50%	0.50	0.010	0.0038
Aluminum	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	50.0	0.823	0.823
Arsenic	FIAS	ICP-MS	50-150%	≤20%	≤30%	0.50	0.0275	0.0087
Cadmium	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.05	0.0094	0.0008
Chromium	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	1.00	1.00	0.024
Copper	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.05	0.05	0.0029
Iron	ICP-MS	ICP-MS	50-150%	≤20%	≤50%	10.0	0.983	0.983
Mercury	CVAA	CVAA	50-150%	≤25%	≤30%	0.01	0.00014	0.00014
Manganese	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.50	0.50	0.003
Nickel	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.05	0.05	0.0114
Lead	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.05	0.0035	0.0044
Selenium	FIAS	ICP-MS	50-150%	≤20%	≤30%	0.20	0.0352	0.0991
Tin	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.50	0.0024	0.0024
Zinc	ICP-MS	ICP-MS	50-150%	≤20%	≤30%	0.50	0.50	0.0493

(1) As stated in the Statement of Work for Chemical Analysis of Marine and Estuarine Samples 15 May 2001.

(2) Reported from the 2003 MDL study.

METHODS:

Battelle MSL analyzed both seawater and freshwater samples for fourteen metals: silver (Ag), aluminum (Al), arsenic (As), cadmium (Cd), chromium (Cr), copper (Cu), iron (Fe), mercury (Hg), manganese (Mn), nickel (Ni), lead (Pb), selenium (Se), tin (Sn) and zinc (Zn). The samples were submitted for analyses by four analytical methods: GFAA, ICP-MS, FIAS and CVAA.

Seawater samples were preconcentrated using iron (Fe) and palladium (Pd) in accordance with the Battelle SOP MSL-I-025, *Methods of Sample Preconcentration*, which is derived from EPA Method 1640. The sample preconcentration was submitted for analysis by ICP-MS and GFAA.

Seawater samples were analyzed by inductively coupled plasma-mass spectrometry (ICP-MS) in accordance with Battelle SOP MSL-I-022, *Determination of Elements in Aqueous and Digestate Samples by ICP-MS*. This method is based on two EPA Methods: 200.8 and 1638. Analytes reported from the preconcentrated seawater samples include: Cd, Cr, Cu, Ni, and Pb. Analytes reported from the direct analysis of the seawater samples include: Al, Fe, Mn, Sn, and Zn. Freshwater samples were analyzed directly by ICP-MS for all analytes, except Hg.

Ag was analyzed in the Fe-Pd preconcentrate by graphite furnace atomic absorption (GFAA) following Battelle SOP MSL-I-029, *Determination of Metals in Aqueous and Digestate Samples by GFAA*, which is derived from EPA Method 200.9.

Seawater samples were analyzed by hydride generation flow injection atomic spectroscopy (FIAS) for As and Se according to Battelle SOP MSL-I-030 *Determination of Metals in Aqueous and Digestate Samples by HGAA-FIAS*.

Seawater and freshwater samples were analyzed by cold-vapor atomic fluorescence spectroscopy (CVAf) for Hg according to Battelle SOP MSL-I-013, *Total Mercury in Aqueous Samples by CVAf*, which is derived from EPA Method 1631.

All results are reported in units of µg/L.

HOLDING TIMES:

The holding times for metals analyses are 90 days from sample collection for Hg analysis, and 6 months from sample collection for analysis of all other metals. The holding times for all metals were achieved.

DETECTION LIMITS:

Target detection limits (TDL) were achieved for all analytes. Achieved method detection limits are reported from the 2003 MDL study. Sample concentrations were evaluated and flagged to the following criteria:

- U Analyte not detected at or above the detection limit, MDL reported
- J Analyte detected above MDL, but below TDL
- * Duplicate out of QC criteria
- e SRM recovery out of QC criteria
- w Spike recovery out of QC criteria due to inappropriate spiking level
- # Continuing calibration recovered outside of acceptable method criteria

NOTE ON Hg QA/QC SAMPLES:

Seawater and freshwater samples were analyzed concurrently for Hg. The QC samples are reported in both the seawater and freshwater tables.

METHOD BLANKS:	Seawater: A minimum of one method blank was analyzed with each analysis batch. Metals concentrations in the method blanks were below the TDL, with the exception of one method blank for Ni and Cu. All sample concentrations for Ni and Cu are greater than five times the detected blank. No corrective action was required. The data were not blank-corrected. Freshwater: A minimum of one method blank was analyzed with each analysis batch. All metals concentrations in the method blanks were below the TDL. The data were not blank-corrected.
BLANK SPIKE or OPR ACCURACY:	Seawater: A minimum of one blank spike or on-going precision and recovery (OPR) sample was analyzed with each analysis batch. Recoveries were reported for spikes at approximate concentrations of 0.005 µg/L for Hg; 5 µg/L for As and Se; and 10 µg/L for Ag, Cd, Cr, Cu, Ni, and Pb. BS recoveries among all metals analyzed ranged from 82% to 107% and were within the QC acceptance criteria of 50% to 150% for all metals. Freshwater: A minimum of one blank spike or on-going precision and recovery (OPR) sample was analyzed with each analysis batch. Recoveries were reported for spikes at approximate concentrations of 0.005 µg/L for Hg; 10 µg/L for Cr, Mn, Ni, Cu, Zn, As, Se, Ag, Cd, Sn, and Pb; and 100 µg/L for Al and Fe. BS recoveries among all metals analyzed ranged from 91% to 119% and were within the QC acceptance criteria of 50% to 150% for all metals.
MATRIX SPIKE ACCURACY:	Seawater: A minimum of one matrix spike was analyzed with each analysis batch. Recoveries were reported for spikes at approximate concentrations of 0.01 µg/L for Hg; 5 µg/L for As and Se; 10 µg/L for Cr, Ni, Cu, Ag, Cd, Sn, and Pb; and 100 µg/L for Al, Fe, Mn and Zn. Matrix spike recoveries among all metals analyzed ranged from 83% to 117% and were within the QC acceptance criteria of 50% to 150% for all metals, with the exception of one MS for Al (240%) and two replicates for Fe (0%, 220%). Low recoveries for the matrix spikes are due to an inappropriate spiking level relative to the native sample concentration. Spiking levels were less than 10% of the native sample concentration, therefore not appropriate for evaluating matrix spike accuracy. Acceptable MS accuracy for Al and Fe was demonstrated in the alternate matrix spike samples. Freshwater: A minimum of one matrix spike was analyzed with each analysis batch. Recoveries were reported for spikes at approximate concentrations of 0.01 µg/L for Hg; 10 µg/L for Cr, Mn, Ni, Cu, As, Se, Ag, Cd, Sn, and Pb; and 100 µg/L for Al, Fe and Zn. Matrix spike recoveries among all metals analyzed ranged from 94% to 118% and were within the QC acceptance criteria of 50% to 150% for all metals.
REPLICATE PRECISION:	Analytical precision for each analysis batch was evaluated by the analysis of laboratory duplicates and expressed as the relative percent deviation (RPD) of duplicate results. Seawater: A minimum of one set of laboratory duplicates was analyzed with each analysis batch. Precision for all metals, except Fe, ranged from 0% to 18% RPD and were within the QC limits of ≤ 30%. RPD values for Fe were 9% and 32% and were within the QC limits of ≤ 50%. Freshwater: A minimum of one set of laboratory duplicates was analyzed with each analysis batch. Precision for all metals ranged from 1% to 19% RPD and

**STANDARD
REFERENCE
MATERIAL
ACCURACY:**

were within the QC limits of $\leq 30\%$.

Accuracy of recovery of SRM analytes was expressed as the percent difference (PD) between the measured and certified SRM concentrations. The target QC criterion is $\leq 20\%$ PD.

Seawater: Standard reference material analyzed for seawater samples include: SRM 1640, SRM CASS-4, and SRM 1641 for Hg. The SRM 1640 is not certified for Sn and the certified value for Fe is not at a level appropriate for data quality evaluation. Percent differences for SRM 1640 and SRM 1641 ranged from 0% to 17% and were within the QC criterion.

The SRM CASS-4 is a low-level seawater reference material. Analytes of interest certified in CASS-4 are less than 10 times the laboratory achieved MDL for all metals except Cu. Currently, there is not seawater SRM certified at a practical quantification level for all analytes of interest. The SRM CASS-4 was analyzed with the preconcentrated seawater samples, and applies only to the metals obtained from this method (Ag, Cr, Ni, Cu, Cd, Pb). Percent differences for analytes within the QC criteria for CASS-4 include As (9%) and Cd (15%). The required preconcentration procedure for low level seawater samples includes the addition of chelating agents to induce precipitation of metals under specific conditions. Subsequently, reagents added to the samples should be of the purest quality to result in zero addition of metals to the samples. The current reagents available contain traces of Cr, Cu and Ni. Correcting CASS-4 results for reagent contributions provide PD values within the QC criterion for Cr (9%), Ni (2%), and Cu (1%). Since CASS-4 is not certified for Ag or Hg and is not certified at practical levels for a majority of the analytes of interest, the alternate SRM (1640 or 1641, respectively) should be used to evaluate the accuracy of this data set. The data were not blank corrected, as the sample concentrations are greater than five times the detected blank for these analytes.

Freshwater: Standard reference material analyzed for freshwater samples include: SRM 1640, SLRS-3 for Fe, and SRM 1641 for Hg. The SRM 1640 is not certified for Sn and the certified value for Fe is not at a level appropriate for data quality evaluation. Percent differences for all SRMs ranged from 0% to 19% and were within the QC acceptance criterion for all metals, with the following exceptions. One replicate of 1640 for Se (28%) and one replicate of 1640 for Zn (21%). In both cases, an alternate replicate of SRM 1640 was analyzed within the batch, which demonstrated acceptable accuracy for Se (0% PD) and Zn (3% PD).

METALS QA/QC (CONT.)

MSL Code	Instrument: Sponsor ID	Rep	GFAA Ag (µg/L)	ICP-MS Al (µg/L)	FIAS As (µg/L)	ICP-MS Cd (µg/L)	ICP-MS Cr (µg/L)	ICP-MS Cu (µg/L)	ICP-MS Fe (µg/L)	ICP-MS Hg (µg/L)	CVAF Hg (µg/L)	ICP-MS Mn (µg/L)	ICP-MS Ni (µg/L)	ICP-MS Pb (µg/L)	FIAS Se (µg/L)	ICP-MS Sn (µg/L)	ICP-MS Zn (µg/L)
METHOD BLANK																	
Method Blank	Hg-03/13/03		NA	NA	NA	NA	NA	NA	NA	NA	0.0014 U	NA	NA	NA	NA	NA	NA
Method Blank	Hg-03/14/03		NA	NA	NA	NA	NA	NA	NA	NA	0.0014 U	NA	NA	NA	NA	NA	NA
Method Blank	Hg-03/18/03		NA	NA	NA	NA	NA	NA	NA	NA	0.0014 U	NA	NA	NA	NA	NA	NA
1979-BLK TRM r1	ICP-MS		0.0038 U	NA	0.0087 U	0.0008 U	0.245 J	0.0029 U	NA	NA	NA	0.003 U	0.0114 U	0.0044 U	0.0951 U	0.0185 J	0.0493 U
1979-BLK TRM r2	ICP-MS		0.0038 U	NA	0.00929 J	0.0008 U	0.321 J	0.0029 U	NA	NA	NA	0.003 U	0.0114 U	0.0044 U	0.0951 U	0.00810 J	0.0493 U
1979-dissolved Blank	ICP-MS		0.00463 J	0.823 U	0.0087 U	0.0039 J	0.024 U	0.0029 U	0.983 U	NA	NA	0.003 U	0.0259 J	0.0044 U	0.0951 U	0.0103 J	0.0493 U
Blank trm r1	ICP-MS (Al, Fe)		NA	0.823 U	NA	NA	NA	NA	0.983 U	NA	NA	NA	NA	NA	NA	NA	NA
METHOD DETECTION LIMIT			0.0038	0.823	0.0087	0.0008	0.024	0.0029	0.983	0.0014	0.0014	0.003	0.0114	0.0044	0.0991	0.0024	0.0493
Project Target Detection Limit			0.50	50.0	0.50	0.05	1.00	0.05	10.0	0.01	0.01	0.5	0.5	0.05	0.20	0.50	0.50
STANDARD REFERENCE MATERIAL																	
1979-1640 TRM	ICP-MS		7.49	58.2	28.3	23.1	41.5	87.8	NA	NA	NA	132	29.2	27.7	21.9	1.54	54.9
1640 Direct	ICP-MS		7.63	53.7	30.8	25.3	40.4	89.9	NA	NA	NA	127	29.2	27.6	28.2	1.56	64.4
1640 TRM	ICP-MS (Al, Fe)		NA	50.6	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
1640	certified value		7.6	52.0	26.7	22.8	38.6	85.2	34.3	NC	NC	122	27.4	27.9	22.0	NC	53.2
	range		±0.25	±1.5	±0.73	±0.96	±1.6	±1.2	±1.6	NA	NA	±1.1	±0.8	±0.14	±0.51	NC	±1.1
	% difference		2%	12%	6%	1%	8%	3%	NA	NA	NA	9%	7%	1%	0%	NA	5%
	% difference		0%	3%	15%	11%	5%	6%	NA	NA	NA	9%	7%	1%	28%	NA	21%
STRS-3 (Fe)	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
STRS-3 (Fe)	ICP-MS		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
STRS-3 (Fe)	ICP-MS (Al, Fe)		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
	certified value		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
	range		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
1641-0031203	Hg-03/13/03		NA	NA	NA	NA	NA	NA	NA	1565	NA	NA	NA	NA	NA	NA	NA
1641-0031303	Hg-03/14/03		NA	NA	NA	NA	NA	NA	NA	1466	NA	NA	NA	NA	NA	NA	NA
1641-0031703	Hg-03/18/03		NA	NA	NA	NA	NA	NA	NA	1573	NA	NA	NA	NA	NA	NA	NA
1641d	certified value		NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC
	range		NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC	NC
	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
ICV,CCV RESULTS																	
ICV	ICP-MS or Hg 1		102%	102%	103%	100%	103%	102%	104%		93%	103%	103%	101%	104%	101%	102%
CCV	ICP-MS or Hg 1		103%	113%	107%	102%	109%	105%	110%		100%	110%	106%	100%	104%	102%	107%
CCV	ICP-MS or Hg 1		104%	113%	105%	102%	108%	106%	115%		98%	109%	108%	98%	104%	102%	105%
CCV	ICP-MS or Hg 1		103%	113%	105%	100%	108%	106%	113%		101%	109%	106%	98%	104%	101%	105%
CCV	ICP-MS or Hg 1		101%	114%	104%	100%	108%	103%	111%		94%	108%	104%	98%	101%	100%	105%
ICV	ICP-MS (Al, Fe) or Hg 2		NA	97%	NA	NA	NA	NA	92%		94%	NA	NA	NA	NA	NA	NA
CCV	ICP-MS (Al, Fe) or Hg 2		NA	101%	NA	NA	NA	NA	96%		92%	NA	NA	NA	NA	NA	NA
CCV	ICP-MS (Al, Fe) or Hg 2		NA	NA	NA	NA	NA	NA	NA		94%	NA	NA	NA	NA	NA	NA
CCV	ICP-MS (Al, Fe) or Hg 2		NA	NA	NA	NA	NA	NA	NA		97%	NA	NA	NA	NA	NA	NA
ICV	Hg 3		NA	NA	NA	NA	NA	NA	100%		100%	NA	NA	NA	NA	NA	NA
CCV	Hg 3		NA	NA	NA	NA	NA	NA	101%		101%	NA	NA	NA	NA	NA	NA
CCV	Hg 3		NA	NA	NA	NA	NA	NA	103%		103%	NA	NA	NA	NA	NA	NA
CCV	Hg 3		NA	NA	NA	NA	NA	NA	NA		98%	NA	NA	NA	NA	NA	NA

METALS QA/QC (CONT.)

MEL	Instrument:	GFAA	ICP-MS	FIAS	ICP-MS	ICP-MS	ICP-MS	ICP-MS	ICP-MS	ICP-MS	ICP-MS	ICP-MS	FIAS	ICP-MS	ICP-MS	ICP-MS
	Sponsor I.D.	Ag (µg/L)	Al (µg/L)	As (µg/L)	Cd (µg/L)	Cr (µg/L)	Cu (µg/L)	Fe (µg/L)	Hg (µg/L)	Mn (µg/L)	Ni (µg/L)	Pb (µg/L)	Se (µg/L)	Si (µg/L)	Zn (µg/L)	
BLANK SPIKE RESULTS																
	Amount Spiked	10	100	10	10	10	10	100	0.00497	10	10	10	10	10	10	10
1979-blk TRM11 or BLANK031203		0.0038 U	0.823 U	0.0087 U	0.0008 U	0.245 J	0.0029 U	36.7	0.00491 J	0.003 U	0.0114 U	0.0044 U	0.0891 U	0.0185 U	0.0493 U	
1979-blk spike r1 or OPR031203Jun1		10.6	114	9.40	9.86	12.1	10.9	149	0.05869	11.7	11	10.8	9.56	11.8	10.2	
Amount Recovered		10.6	114	9.40	9.86	11.9	10.9	112	0.05827	7-Nov	11	10.8	9.56	11.8	10.2	
Percent Recovery		106%	114%	94%	100%	119%	109%	112%	105%	117%	110%	108%	95%	116%	102%	
	Amount Spiked	100	100	10	10	10	10	100	0.00497	10	10	10	10	10	10	
1979-blk TRM12 or BLANK031203		0.0038 U	0.823 U	0.00929 J	0.0008 U	0.321 J	0.0029 U	36.5	0.00491 J	0.003 U	0.0114 U	0.0044 U	0.0891 U	0.00810 U	0.0493 U	
1979-blk spike r2 or OPR031203Jun2		10.7	113	9.30	9.89	12.1	10.9	150	0.05845	11.8	11	10.6	9.05	11.7	9.76	
Amount Recovered		10.7	113	9.29	9.89	11.8	10.9	114	0.05803	11.8	11	10.6	9.05	11.7	9.76	
Percent Recovery		107%	113%	93%	99%	118%	109%	114%	101%	118%	110%	105%	91%	117%	98%	
Amount Spiked		NS	NS	NS	NS	NS	NS	NS	0.00487	NS	NS	NS	NS	NS	NS	
BLANK031303									0.000172 J							
OPR031303Jun1		Amount Recovered	NA	NA	NA	NA	NA	NA	0.00473			NA	NA	NA	NA	
	Percent Recovery	NA	NA	NS	NS	NS	NA	NA	97%			NA	NA	NA	NA	
	Amount Spiked	NS	NS	NS	NS	NS	NS	NS	0.00487	NS	NS	NS	NS	NS	NS	
BLANK031303									0.000172 J							
OPR031303Jun2		Amount Recovered	NA	NA	NA	NA	NA	NA	0.0502							
	Percent Recovery	NA	NA	NS	NS	NS	NA	NA	0.00485	NA	NA	NA	NA	NA	NA	
	Amount Spiked	NS	NS	NS	NS	NS	NS	NS	100%	NA	NA	NA	NA	NA	NS	
BLANK031403									0.00491							
OPR031403Jun1		Amount Recovered	NA	NA	NA	NA	NA	NA	0.00528							
	Percent Recovery	NA	NA	NS	NS	NS	NA	NA	0.00508			NA	NA	NA	NA	
	Amount Spiked	NS	NS	NS	NS	NS	NS	NS	103%	NA	NA	NA	NA	NA	NS	
BLANK031403									0.00491							
OPR031403Jun2		Amount Recovered	NA	NA	NA	NA	NA	NA	0.00202 J							
	Percent Recovery	NA	NA	NS	NS	NS	NA	NA	0.00547							
	Amount Spiked	NS	NS	NS	NS	NS	NS	NS	0.00527	NA	NA	NA	NA	NA	NA	
MATRIX SPIKE RESULTS									107%	NA	NA	NA	NA	NA	NA	
1979-15		Amount Spiked	NS	NS	NS	NS	NS	NS	0.0161	NS	NS	NS	NS	NS	NS	
	NAV-OP24-SDB2-FF								0.00679 J							
	MS								0.0223							
	Amount Recovered	NA	NA	NA	NA	NA	NA	NA	0.0155	NA	NA	NA	NA	NA	NA	
	Percent Recovery	NA	NA	NA	NA	NA	NA	NA	96%	NA	NA	NA	NA	NA	NA	
	Amount Spiked	NS	NS	NS	NS	NS	NS	NS	0.0157	NS	NS	NS	NS	NS	NS	
1979-15		NAV-OP24-SDB2-FF							0.00679 J							
	MSD								0.0215							
	Amount Recovered	NA	NA	NA	NA	NA	NA	NA	0.0147	NA	NA	NA	NA	NA	NA	
	Percent Recovery	NA	NA	NA	NA	NA	NA	NA	94%	NS	NS	NA	NA	NA	NA	
	Amount Spiked	10	100	10	10	10	10	100	NS	10	10	10	10	10	100	
1979-24		NAV-PRS-SDB2-Comp	0.00809 J	15.1	1.18	0.303	1.12	17.6	0.533	1.88	0.533	0.247	0.0603 J	0.0603 J	80.8	
	MS	10.6	130	11.4	10.7	12.6	10.3	130	12.3	12.8	10.5	11.0	11.6	11.6	185	
	Amount Recovered	10.6	116	10.2	10.4	11.5	10.1	112	NA	11.4	10.9	10.8	11.5	11.5	104	
	Percent Recovery	106%	116%	102%	104%	115%	101%	112%	NA	114%	109%	109%	108%	115%	104%	
	Amount Spiked	10	100	10	10	10	10	100	NS	10	10	10	10	10	100	
1979-24		NAV-PRS-SDB2-Comp	0.00809 J	15.1	1.18	0.303	1.12	17.6	0.533	1.88	0.533	0.247	0.0603 J	0.0603 J	80.8	
	MSD	10.7	129	11.7	10.8	12.7	10.3	132	11.9	12.9	10.9	11.1	11.9	11.9	184	
	Amount Recovered	10.7	114	10.5	10.5	11.6	10.3	114	NA	11.4	11	10.4	10.9	11.8	103	
	Percent Recovery	107%	114%	105%	105%	116%	103%	114%	NA	114%	110%	104%	108%	118%	103%	
REPLICATE RESULTS																
1979-23		NAV-PR6-SDB2-FF	0.0266	1.41	1.23	3.58	177	161	0.0133	81.5	17.2	0.879	1.33	0.289 J	288	
1979-23		2 NAV-PR6-SDB2-FF	NA	NA	NA	NA	NA	NA	0.0132	NA	NA	NA	NA	NA	NA	
	% difference	NA	NA	NA	NA	NA	NA	NA	1%	NA	NA	NA	NA	NA	NA	
1979-24		NAV-PR3-SDB2-Comp	0.00809	15.1	1.18	0.303	1.12	17.6	0.00219 J	5.94	2	0.533	0.247	0.0603 J	80.8	
2 NAV-PR3-SDB2-Comp		0.00670	14.8	1.10	0.295	1.14	13.6	16.2	NA	1.91	0.502	0.0591 U	0.0688 J	79.7		
	% difference	19%	2%	7%	3%	2%	4%	8%	NA	1%	2%	6%	NA	13%		
= Seawater MDLs reported from the 2003 MDL Study; U= Analyte not detected at or above detection limit; MDL reported; NC = SRM not certified; NA= Not analyzed or applicable; B = Sample results are less than 5x the blank; J= Analyte detected above the MDL, but below the TDL; % difference																

t = Seawater MDLs reported from the 2003 MDL Study; U = Analyte not detected at or above detection limit, MDL reported; NC = SRM not certified; NA = Not analyzed or applicable; B = Sample results are less than 5x the blank; J = Analyte detected above the MDL, but below the TDL; e = SRM recovery out of QC criteria; w = Spike recovery out of QC criteria due to inappropriate spiking level; # = continuing calibration recovered outside of acceptable method criteria.

METALS QA/QC (CONT.)

MST Code	Instrument: Sponsor ID	Rep	GFAA Ag (µg/L)	ICP-MS Al (µg/L)	FIAS As (µg/L)	ICP-MS Cd (µg/L)	ICP-MS Cr (µg/L)	ICP-MS Cu (µg/L)	ICP-MS Fe (µg/L)	CVAE Hg (µg/L)	ICP-MS Mn (µg/L)	ICP-MS Ni (µg/L)	ICP-MS Pb (µg/L)	FIAS Se (µg/L)	ICP-MS Sn (µg/L)	ICP-MS Zn (µg/L)
METHOD BLANK																
bik TRM r1	ICP-MS Direct		NA	0.823 U	NA	NA	NA	NA	0.983 U	NA	0.003 U	NA	NA	NA	0.00578 U	0.5 U
bik TRM r2	ICP-MS Direct		NA	0.823 U	NA	NA	NA	NA	0.983 U	NA	0.003 U	NA	NA	NA	0.00578 U	0.5 U
1979-bik	Fe/Pd ICP-MS or GFAA-Ag		0.0174 J	NA	NA	0.0094 U	0.0913 U	0.151	NA	0.00014 U	NA	0.105	0.0203 J	NA	NA	NA
Method Blank	Hg- 03113003		NA	NA	NA	NA	NA	NA	NA	0.00014 U	NA	NA	NA	NA	NA	NA
Method Blank	Hg- 03114003		NA	NA	NA	NA	NA	NA	NA	0.00014 U	NA	NA	NA	NA	NA	NA
Method Blank	Hg- 03118003		NA	NA	NA	NA	NA	NA	NA	0.00014 U	NA	NA	NA	NA	NA	NA
BLANK	FIAS - As		NA	NA	0.0275 U	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
BLANK	FIAS - Se		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	0.0352 U	NA	NA
METHOD DETECTION LIMIT ¹			0.010	0.823	0.0275	0.0094	0.0218	0.0540	0.983	0.00014	0.003	0.0286	0.0035	0.00352	0.0024	0.5
Project Target Detection Limit																
STANDARD REFERENCE MATERIAL																
1640 Direct	ICP-MS Direct		NA	51.8	NA	NA	NA	NA	NA	NA	123	NA	NA	NA	1.58	62.3
1640 TRM	ICP-MS Direct		NA	48.8	NA	NA	NA	NA	NA	NA	117	NA	NA	NA	1.52	50.6
1640 Direct	Fe/Pd ICP-MS or GFAA-Ag		6.96	NA	NA	24.7	39.4	87.2	NA	NA	NA	28.2	28.0	NA	NA	NA
1640 Direct	Fe/Pd ICP-MS or GFAA-Ag		NA	NA	NA	24.5	37.1	83.0	NA	NA	NA	26.7	27.3	NA	NA	NA
1640	FIAS - Se		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
1640	certified value		7.6	52.0	26.7	22.8	38.6	85.2	34.3	NC	122	27.4	27.9	22.0	NC	53.2
1640	range		±0.25	±1.5	±0.73	±0.96	±1.6	±1.2	±1.6	NC	±1.1	±0.8	±0.14	±0.51	NC	±1.1
1640	% difference		NA	0%	NA	NA	NA	NA	NA	NA	1%	NA	NA	NA	NA	17%
1640	% difference		NA	6%	NA	8%	NA	2%	NA	NA	4%	NA	NA	NA	NA	5%
1640	% difference		NA	3%	NA	8%	4%	3%	NA	NA	NA	3%	NA	NA	NA	NA
1640	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
1640	% difference		NA	NA	NA	NA	NA	0.748	NA	NA	NA	0.425	0.0265	NA	NA	NA
1640	FIAS - As		NA	1.01	NA	NA	NA	NA	0.71	NA	2.78	0.314	0.0098	NC	NA	NA
CASS-4	certified value		NC	NC	1.11	0.026	0.144	0.592	0.71	NA	2.78	0.314	0.0098	NC	NA	NA
CASS-4	range		NC	NC	±0.16	±0.003	±0.029	±0.055	±0.058	NA	±0.19	±0.030	±0.0038	NC	NC	±0.057
CASS-4	% difference		NA	NA	9%	15%	54%	27%	NA	NA	NA	35%	170%	NC	NA	0.381
CASS-4	% difference		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
1641-d031203	Hg		NA	NA	9%	NA	NA	NA	NA	1565	NA	NA	NA	NA	NA	NA
1641-d031303	Hg		NA	NA	NA	NA	NA	NA	NA	1465	NA	NA	NA	NA	NA	NA
1641-d031703	Hg		NA	NA	NA	NA	NA	NA	NA	1573	NA	NA	NA	NA	NA	NA
1641-d	certified value		NC	NC	NC	NC	NC	NC	NC	1590	NC	NC	NC	NC	NC	NC
1641-d	range		NC	NC	NC	NC	NC	NC	NC	±4.00	NC	NC	NC	NC	NC	NC
1641-d	% difference		NA	NA	NA	NA	NA	NA	NA	2%	NA	NA	NA	NA	NA	NA
1641-d	% difference		NA	NA	NA	NA	NA	NA	NA	8%	NA	NA	NA	NA	NA	NA
1641-d	% difference		NA	NA	NA	NA	NA	NA	NA	1%	NA	NA	NA	NA	NA	NA
ICV/CCV RESULTS																
ICV	ICP-MS Direct or Hg			101%					109%	93%	104%				108%	105%
CCV	ICP-MS Direct or Hg			98%					99%	100%	105%				104%	105%
CCV	ICP-MS Direct or Hg			106%					113%	98%	104%				105%	107%
CCV	ICP-MS Direct or Hg			101%					101%	101%	104%				101%	100%
ICV	ICP-MS Direct or Hg		102%			102%	103%	102%	112%	94%	100%		100%		101%	98%
CCV	Fe/Pd ICP-MS or Hg		104%			100%	98%	98%		92%		102%	102%			
CCV	Fe/Pd ICP-MS or Hg		101%			101%	98%	96%		97%		97%	105%			
CCV	Fe/Pd ICP-MS or Hg		101%			100%	98%	96%		97%		96%	102%			
ICV	Fe/Pd ICP-MS		NA			99%	95%	95%		NA		94%	100%			
CCV	FIAS-As or Hg				103%					100%						
CCV	FIAS-As or Hg				98%					101%						
CCV	FIAS-As or Hg				96%					98%						
ICV	FIAS-Se				99%					98%						
CCV	FIAS-Se															
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[illegible]

METALS QA/QC (CONT.)

Reap	NSL Code	Intrinsic Sponsor ID.	GFAA Au (µg/L)	ICP-MS Al (µg/L)	FIAS As (µg/L)	ICP-MS Cd (µg/L)	ICP-MS Cr (µg/L)	ICP-MS Cu (µg/L)	ICP-MS Fe (µg/L)	CYAF Hg (µg/L)	ICP-MS Mn (µg/L)	ICP-MS Ni (µg/L)	ICP-MS Pb (µg/L)	ICP-MS Se (µg/L)	FIAS Sn (µg/L)	ICP-MS Zn (µg/L)
REPLICATE RESULTS																
			0.168 J	1840	1.58	0.987	6.56	54.2	2390	0.0773	92.6	12.5	22.7	0.187 J	1.00	433
2	NAV-CF8-SDB2-FF	1979-1		1790	NA	NA	NA	NA	2180	0.074	87.8	NA	NA	1.02	422	
	% difference		NA	3%	NA	NA	NA	1%	9%	NA	5%	NA	NA	2%	3%	
			0.0324 J	107	1.17	0.109	1.76	5.01	152	0.00290 J	12.5	1.93	0.822	0.0359 J	NA	24.7
2	NAV-BAY14-SDB2-D	1979-12		NA	1.20	0.113	1.74	4.98	NA	NA	NA	1.94	0.602	0.032 U	NA	NA
	% difference		18%		3%	0%	1%	0%	NA	NA	NA	1%	3%	NA	NA	NA
			0.0203 J	16.8 J	0.695	0.388	1.22	25.8	18.5	0.00387 J	28.7	6.95	0.369	0.132 J	0.165 J	218
2	NAV-CF8-SDB2-FF	1979-16		NA	16.4 J	NA	NA	NA	25.5	NA	29.9	NA	NA	NA	0.149 J	229
	% difference		NA	1%	NA	NA	NA	NA	32%	NA	4%	NA	NA	NA	10%	5%

1 = Senwater MDLs reported from the 2003 MDL Study. U = Analyte not detected at or above detection limit, MDL reported; NC = SRM not certified; NA = Not analyzed or applicable; B = Sample results are less than 5x the blank; J = Analyst detected above the MDL, but below the TDL, e = SRM recovery out of QC criteria, w = Spike recovery out of QC criteria due to inappropriate spiking level, # = continuing calibration recovered outside of acceptable method criteria.

WIL		CHEMISTRY CODE		CLIENT CODE		PARAMETER		CRM-MS		CRM-MS		CRM-MS		CRM-MS		CRM-MS	
DETECTION LIMITS								AsI (µg/L)		Cu (µg/L)		Pb (µg/L)		Hg (µg/L)		Zn (µg/L)	
Project Reservoir Limit								0.1		0.1		0.1		0.1		0.1	
Laboratory Achieved MDL								0.025		0.025		0.0055		0.0016		0.11	
Blank (1)								0.0145		0.312		0.0233		0.00326		0.382	
Blank (2)								0.0255		0.0156		0.004018		0.027		0.27	
								0.0198		0.068		0.0259		0.00093		0.82	
LABORATORY CONTROL SAMPLE (LCS) ACCURACY								Mean									
LCS 1								0.0136		0.63		0.0245		0.003036		1.72	
LCS 2								9.30		10.4		8.67		0.00526		11.6	
LABORATORY CONTROL SAMPLE (LCS) RECOVERY								NS		NS		NS		NS		NS	
Amount Recovered								9.25		9.77		8.55		0.00495		9.87	
Amount Spiked								NS		NS		NS		NS		NS	
Blank (1)								Amount Recovered		Amount Recovered		Amount Recovered		Amount Recovered		Amount Recovered	
OPR030603-ws2 (Hg)								NA		NA		NA		NA		NA	
								NA		NA		NA		NA		NA	
								NA		NA		NA		NA		NA	
								Amount Spiked		Amount Spiked		Amount Spiked		Amount Spiked		Amount Spiked	
Blank (2)								0.0152		0.45		0.0236		0.00566		1.07	
SB Blank (2)								9.30		10.1		8.26		0.00566		11.5	
OPR031003-w1 (Hg)								9.24		9.45		8.25		0.00592		10.2	
LCS 2								52%		95%		83%		100%		102%	
								NS		NS		NS		NS		NS	
								Amount Spiked		Amount Spiked		Amount Spiked		Amount Spiked		Amount Spiked	
Blank (2)								NA		NA		NA		NA		NA	
OPR031003-ws2 (Hg)								NA		NA		NA		NA		NA	
								NA		NA		NA		NA		NA	
								Amount Recovered		Amount Recovered		Amount Recovered		Amount Recovered		Amount Recovered	
NA-TEX SPIKE ACCURACY								NA		NA		NA		NA		NA	
NA-19-27 NSD								Total metals		Total metals		Total metals		Total metals		Total metals	
								0.0306		4.86		0.152		--		13.1	
								9.24		14.5		8.72		--		21.8	
								% Rec (10ppb)		91%		93%		85%		82%	
								% Rec (10ppb)		85%		86%		85%		84%	
								Total metals		0.031		0.02		--		22.2	
NA-19-37								9.55		12.3		8.95		--		22.2	
NA-19-37 NSD								9.24		12.3		8.95		--		22.2	
								% Rec (10ppb)		55%		95%		85%		85%	
								% Rec (10ppb)		52%		95%		85%		85%	
								Total metals		95%		53%		0.0010		85%	
NA-19-38 1								NA		NA		NA		NA		NA	
NA-19-38 NSD								NA		NA		NA		NA		NA	
								% Rec (0.012ppb)		--		--		0.012		--	
								% Rec (0.012ppb)		--		--		152%		--	
								Total metals		--		--		160%		--	

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PAHs

CLIENT SAMPLE ID	NAV- PR5-SDB2-FF	NAV- PR5-SDB2- COMP	NAV- PR6-SDB2-FF	NAV- PR6-SDB2- COMP
Battelle Sample ID	U7089	U7091	U7090	U7092
Battelle Batch ID	03-0203	03-0203	03-0203	03-0203
Data File	A1884.D	A1886.D	A1885.D	A1887.D
Extraction Date	03/04/03	03/04/03	03/04/03	03/04/03
Acquired Date	03/20/03	03/20/03	03/20/03	03/20/03
Matrix	Water	Water	Water	Water
Sample Size (L)	2.62	2.64	2.66	2.66
Dilution Factor	1.667	1.667	1.667	1.667
PIV (mL)	0.3	0.3	0.3	0.3
Min Reporting Limit	0.95	0.95	0.94	0.94
Amount Units	ng/L	ng/L	ng/L	ng/L
Naphthalene	6.33 B	545.51	7.70	10.74
C1-Naphthalenes	4.17	2246.10	6.50	8.58
C2-Naphthalenes	5.15	1886.51	9.47	13.41
C3-Naphthalenes	4.38	784.87	8.81	6.71
C4-Naphthalenes	0.51 U	301.05	0.50 U	0.50 U
2-Methylnaphthalene	3.62	2088.87	4.74	8.10
1-Methylnaphthalene	2.60	1449.34	4.68	5.34
2,6-Dimethylnaphthalene	1.27	860.28	1.50	2.66
2,3,5-Trimethylnaphthalene	0.82 J	150.84	0.72 J	1.20
Biphenyl	2.18	472.71	7.14	4.17
Acenaphthylene	1.25	15.67	1.05	2.69
Acenaphthene	2.42	29.29	1.94	1.67
Fluorene	3.27	94.48	2.90	3.50
C1-Fluorenes	1.95 B	108.44	4.84	4.77
C2-Fluorenes	10.13 B	63.38	59.86	24.62
C3-Fluorenes	14.05 B	81.06	24.82 B	62.85
Phenanthrene	29.86	54.65	15.88 B	34.46
Anthracene	1.79 B	4.14	1.00 B	2.72
C1-Phenanthrenes/Anthracenes	21.63 B	49.05	10.27 B	28.98 B
C2-Phenanthrenes/Anthracenes	56.76 B	59.66 B	22.81 B	49.46 B
C3-Phenanthrenes/Anthracenes	54.90 B	44.74 B	11.21 B	38.07 B
C4-Phenanthrenes/Anthracenes	24.81 B	15.45 B	0.28 U	14.00 B
1-Methylphenanthrene	5.84 B	12.86	3.65 B	8.78
Dibenzothiophene	7.87	11.22	19.20	9.80
C1-Dibenzothiophenes	14.21	19.15	30.88	20.81
C2-Dibenzothiophenes	49.50 B	41.42 B	59.11 B	64.18 B
C3-Dibenzothiophenes	48.76 B	39.87 B	37.56 B	52.76 B
Fluoranthene	29.72	48.56	9.44 B	34.66
Pyrene	31.48 B	56.25	8.35 B	38.24 B
C1-Fluoranthenes/Pyrenes	34.42 B	24.86 B	9.16 B	21.00 B
C2-Fluoranthenes/Pyrenes	27.19 B	26.14 B	7.14 B	26.09 B
C3-Fluoranthenes/Pyrenes	17.20 B	24.23 B	5.11 B	25.04 B
Benzo(a)anthracene	3.58	12.42	0.75 J	4.00
Chrysene	21.51	30.86	4.84	30.12
C1-Chrysenes	14.05 B	23.82	4.46 B	20.74
C2-Chrysenes	17.70 B	33.07 B	0.28 U	27.39 B
C3-Chrysenes	13.41 B	31.09 B	0.28 U	27.34 B
C4-Chrysenes	0.28 U	0.28 U	0.28 U	0.28 U
Benzo(b)fluoranthene	10.84	18.72	2.12	10.57
Benzo(k)fluoranthene	7.08	13.96	1.21	6.19
Benzo(e)pyrene	10.01	16.44	1.93 B	10.36
Benzo(a)pyrene	4.99	11.68	0.72 J	4.32
Perylene	1.48	3.60	0.53 U	1.86
Indeno(1,2,3-cd)pyrene	6.91	12.53	1.13 B	6.08
Dibenz(a,h)anthracene	1.23	2.44	0.78 U	1.17
Benzo(g,h,i)perylene	15.16	25.43	2.48	13.59
Total Priority Pollutant PAHs	177.42	976.59	61.51	204.72
Naphthalene-d8	60	64	63	60
Phenanthrene-d10	75	76	76	79
Chrysene-d12	81	85	80	91

PAHs (CONT.)

CLIENT SAMPLE ID	NAV-BAY9-SDB2-PRE	NAV-BAY9-SDB2-DUR	NAV-BAY9-SDB2-AFT	NAV-BAY11-SDB2-PRE	NAV-BAY11-SDB2-DUR	NAV-BAY11-SDB2-AFT	NAV-BAY14-SDB2-PRE
Battelle Sample ID	U7067	U7071	U7075	U7068	U7072	U7076	U7069
Battelle Batch ID	03-0200	03-0200	03-0200	03-0200	03-0200	03-0200	03-0200
Data File	A1850.D	A1857.D	A1861.B.D	A1854.D	A1858.D	A1863.D	A1855.D
Extraction Date	03/03/03	03/03/03	03/03/03	03/03/03	03/03/03	03/03/03	03/03/03
Acquired Date	03/18/03	03/18/03	03/18/03	03/18/03	03/18/03	03/18/03	03/18/03
Matrix	Water	Water	Water	Water	Water	Water	Water
Sample Size (L)	1.3	2.64	2.65	2.64	2.66	2.64	2.62
Dilution Factor	1.667	1.667	1.667	1.667	1.667	1.667	1.667
PIV (mL)	0.3	0.3	0.3	0.3	0.3	0.3	0.3
Min Reporting Limit	1.92	0.95	0.94	0.95	0.94	0.95	0.95
Amount Units	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L
Naphthalene	3.89 B	1.74 B	1.48 B	1.43 B	1.36 B	1.26 B	1.26 B
C1-Naphthalenes	2.45 B	0.91 JB	0.66 JB	1.27 B	0.70 JB	0.75 JB	0.73 JB
C2-Naphthalenes	1.02 U	2.90	0.50 U	0.50 U	0.50 U	2.46	2.18
C3-Naphthalenes	1.02 U	2.10	1.62	0.50 U	0.50 U	2.15	1.41
C4-Naphthalenes	1.02 U	0.50 U	0.50 U	0.50 U	0.50 U	0.50 U	0.51 U
2-Methylnaphthalene	2.00 B	0.83 JB	0.61 JB	1.12 B	0.64 JB	0.77 JB	0.67 JB
1-Methylnaphthalene	1.69 JB	0.57 JB	0.35 JB	0.94 JB	0.31 JB	0.55 JB	0.32 JB
2,6-Dimethylnaphthalene	0.73 U	0.70 J	0.36 U	0.93 J	0.36 U	1.03	0.78 J
2,3,5-Trimethylnaphthalene	0.77 U	0.78 J	0.38 U	1.37	0.37 U	0.64 J	0.48 J
Biphenyl	0.60 J	0.89 J	0.31 U	0.79 J	0.31 U	0.57 J	0.41 J
Acenaphthylene	1.37 J	0.89 J	0.65 J	1.74	0.63 J	1.04	0.66 J
Acenaphthene	6.51	6.44	3.08	4.74	4.91	10.63	3.13
Fluorene	2.63	2.70	1.03	2.03	1.84	3.98	0.90 J
C1-Fluorenes	0.75 U	1.38	1.22	0.37 U	0.37 U	1.79	0.99
C2-Fluorenes	0.75 U	0.37 U	0.37 U	0.37 U	0.37 U	2.71	0.37 U
C3-Fluorenes	0.75 U	0.37 U	0.37 U	0.37 U	0.37 U	0.37 U	0.37 U
Phenanthrene	2.74	7.01	1.11 B	1.87	3.32	5.05	0.89 JB
Anthracene	1.27 J	1.93	1.08	2.00	1.60	2.23	0.83 J
C1-Phenanthrenes/Anthracenes	0.58 U	4.31	2.31	0.28 U	2.98	4.02	0.29 U
C2-Phenanthrenes/Anthracenes	0.58 U	3.18	1.79	0.28 U	1.81	2.51	0.29 U
C3-Phenanthrenes/Anthracenes	0.58 U	3.52	1.48	0.28 U	1.97	1.76	0.29 U
C4-Phenanthrenes/Anthracenes	0.58 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	0.29 U
1-Methylphenanthrene	0.67 U	1.02	0.47 J	1.39	0.57 J	0.81 J	0.33 U
Dibenzothiophene	0.90 J	1.52	0.69 J	1.62	0.78 J	1.23	0.36 J
C1-Dibenzothiophenes	0.71 U	1.96	0.89 J	0.35 U	0.83 J	1.03	0.35 U
C2-Dibenzothiophenes	0.71 U	3.64	1.99	0.35 U	1.91	2.57	0.35 U
C3-Dibenzothiophenes	0.71 U	3.23	1.15	0.35 U	1.33	1.12	0.35 U
Fluoranthene	10.18	20.21	10.58	10.30	13.99	25.73	6.95
Pyrene	4.90	13.05	5.34	5.41	8.43	14.77	3.31
C1-Fluoranthenes/Pyrenes	0.68 U	4.93	2.23	1.70	3.50	5.70	1.70
C2-Fluoranthenes/Pyrenes	0.68 U	3.42	0.33 U	0.33 U	1.91	1.76	0.34 U
C3-Fluoranthenes/Pyrenes	0.68 U	2.92	0.33 U	0.33 U	0.33 U	0.33 U	0.34 U
Benzo(a)anthracene	1.07 U	2.63	0.75 J	1.62	1.28	3.30	0.34 J
Chrysene	0.56 U	5.87	1.60	2.14	2.34	4.52	0.78 J
C1-Chrysenes	0.56 U	3.33	0.28 U	0.28 U	1.60	1.79	0.28 U
C2-Chrysenes	0.56 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U
C3-Chrysenes	0.56 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U
C4-Chrysenes	0.56 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U
Benzo(b)fluoranthene	0.66 U	3.49	1.15	2.56	1.93	2.73	0.33 U
Benzo(k)fluoranthene	0.67 U	2.44	1.08	1.91	1.45	2.48	0.33 U
Benzo(e)pyrene	0.70 U	3.04	1.06	2.08	1.27	1.62	0.88 J
Benzo(a)pyrene	1.00 U	2.19	0.74 J	1.85	1.15	1.92	0.50 U
Perylene	1.07 U	0.67 J	0.53 U	1.51	0.53 U	0.53 U	0.53 U
Indeno(1,2,3-cd)pyrene	0.82 JB	2.10 B	0.66 JB	2.13 B	0.85 JB	0.81 JB	0.35 JB
Dibenz(a,h)anthracene	1.60 U	0.58 J	0.79 U	1.21	0.78 U	0.79 U	0.80 U
Benzo(g,h,i)perylene	2.31 B	2.63 B	0.92 JB	2.49 B	1.22 B	1.10 B	0.66 JB
Total Priority Pollutant PAHs	36.62	75.89	31.26	45.41	46.29	81.56	20.07
Naphthalene-d8	65	53	54	62	54	63	70
Phenanthrene-d10	82	72	70	73	71	72	74
Chrysene-d12	92	87	86	83	76	88	79

PAHs (CONT.)

CLIENT SAMPLE ID	NAV-BAY14-SDB2-DUR	NAV-BAY14-SDB2-AFT	NAV-BAY14A-SDB2-PRE	NAV-BAY14A-SDB2-DUR	NAV-BAY14A-SDB2-AFT	
Battelle Sample ID	U7073	U7077	U7070	U7074	U7078	
Battelle Batch ID	03-0200	03-0200	03-0200	03-0200	03-0200	
Data File	A1861.D	A1864.D	A1856.D	A1861A.D	A1866.D	
Extraction Date	03/03/03	03/03/03	03/03/03	03/03/03	03/03/03	
Acquired Date	03/18/03	03/19/03	03/18/03	03/18/03	03/19/03	
Matrix	Water	Water	Water	Water	Water	
Sample Size (L)	2.65	2.66	2.64	2.66	2.64	
Dilution Factor	1.667	1.667	1.667	1.667	1.667	
PIV (mL)	0.3	0.3	0.3	0.3	0.3	
Min Reporting Limit	0.94	0.94	0.95	0.94	0.95	
Amount Units	ng/L	ng/L	ng/L	ng/L	ng/L	
Naphthalene	1.30 B	1.18 B	1.38 B	1.27 B	1.31 B	
C1-Naphthalenes	0.76 JB	0.57 JB	0.76 JB	0.70 JB	0.64 JB	
C2-Naphthalenes	0.50 U	0.50 U	1.45	0.50 U	0.50 U	
C3-Naphthalenes	1.55	1.08	1.11	1.13	1.50	
C4-Naphthalenes	0.50 U	0.50 U	0.50 U	0.50 U	0.50 U	
2-Methylnaphthalene	0.60 JB	0.48 JB	0.73 JB	0.60 JB	0.65 JB	
1-Methylnaphthalene	0.46 JB	0.27 JB	0.42 JB	0.34 JB	0.30 JB	
2,6-Dimethylnaphthalene	0.36 U	0.36 U	0.73 J	0.36 U	0.36 U	
2,3,5-Trimethylnaphthalene	0.38 U	0.37 U	0.38 U	0.37 U	0.38 U	
Biphenyl	0.25 J	0.24 J	0.28 J	0.31 U	0.34 J	
Acenaphthylene	0.54 J	0.49 J	0.54 J	0.41 J	0.64 J	
Acenaphthene	2.46	2.36	2.53	2.45	3.99	
Fluorene	0.82 J	0.85 J	0.78 J	0.70 J	1.28	
C1-Fluorenes	0.37 U	0.37 U	0.37 U	1.03	1.12	
C2-Fluorenes	0.37 U	0.37 U	0.37 U	0.37 U	0.37 U	
C3-Fluorenes	0.37 U	0.37 U	0.37 U	0.37 U	0.37 U	
Phenanthrene	2.16	1.18 B	0.79 JB	1.54 B	1.37 B	
Anthracene	1.09	0.78 J	0.68 J	1.22	1.09	
C1-Phenanthrenes/Anthracenes	2.93	2.14	1.41	2.08	2.49	
C2-Phenanthrenes/Anthracenes	2.31	1.85	0.28 U	1.45	2.30	
C3-Phenanthrenes/Anthracenes	1.85	0.28 U	0.28 U	2.01	1.59	
C4-Phenanthrenes/Anthracenes	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	
1-Methylphenanthrene	0.60 J	0.48 J	0.39 J	0.43 J	0.54 J	
Dibenzothiophene	0.88 J	0.60 J	0.44 J	0.72 J	0.68 J	
C1-Dibenzothiophenes	1.48	0.72 J	0.35 U	1.13	0.78 J	
C2-Dibenzothiophenes	2.81	1.81	0.35 U	1.62	1.74	
C3-Dibenzothiophenes	1.86	0.35 U	0.35 U	1.13	1.45	
Fluoranthene	12.93	10.43	7.22	10.95	11.65	
Pyrene	8.05	5.43	3.11	6.32	5.89	
C1-Fluoranthenes/Pyrenes	4.06	2.58	1.26	3.47	2.48	
C2-Fluoranthenes/Pyrenes	1.98	0.33 U	0.33 U	0.33 U	0.33 U	
C3-Fluoranthenes/Pyrenes	0.33 U	0.33 U	0.33 U	0.33 U	0.33 U	
Benzo(a)anthracene	1.37	0.62 J	0.34 J	1.08	0.69 J	
Chrysene	3.40	1.54	0.69 J	2.24	1.41	
C1-Chrysenes	2.64	0.28 U	0.28 U	0.28 U	0.28 U	
C2-Chrysenes	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	
C3-Chrysenes	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	
C4-Chrysenes	0.28 U	0.28 U	0.28 U	0.28 U	0.28 U	
Benzo(b)fluoranthene	2.98	1.34	0.33 U	2.17	1.23	
Benzo(k)fluoranthene	2.47	1.02	0.33 U	1.75	0.89 J	
Benzo(e)pyrene	2.03	1.46	0.34 U	1.59	1.00	
Benzo(a)pyrene	1.84	0.49 U	0.49 U	1.50	0.74 J	
Perylene	0.53 U	0.53 U	0.53 U	0.53 U	0.53 U	
Indeno(1,2,3-cd)pyrene	1.43 B	0.69 JB	0.45 JB	0.97 B	0.56 JB	
Dibenz(a,h)anthracene	0.79 U	0.78 U	0.79 U	0.78 U	0.79 U	
Benzo(g,h,i)perylene	1.54 B	0.68 JB	0.54 JB	1.27 B	0.48 JB	
Total Priority Pollutant PAHs	44.36	28.59	19.04	35.84	33.23	
Naphthalene-d8	54	57	69	55	65	
Phenanthrene-d10	70	69	77	70	72	
Chrysene-d12	85	88	81	85	86	

PAHs QA/QC

PROJECT: SPAWAR TO0011, Contaminant Analysis of Stormwater and San Diego Bay
PARAMETER: Seawater
LABORATORY: PAH
MATRIX: Battelle, Duxbury, MA
SAMPLE CUSTODY: Water
 The water samples were collected February 25, 2003. They were received in Duxbury on February 28, 2003 in good condition in six coolers. The cooler temperature on arrival ranged from 0.2 °C to 1.3 °C. Samples were stored at 4 °C until processing.

QA/QC DATA QUALITY OBJECTIVES:

	Reference Method	Surrogate Recovery	LCS/MS Recovery	Sample Replicate Relative Precision	Procedural Blank
PAH	General NS&T	30-130% Recovery	LCS: 40-120% Recovery for at least 80% of analytes MS: 50-150% Recovery for at least 70% of analytes; analyte conc. in MS must be >5x background	≤30% RSD analyte conc. in MS must be <5x background	<3X MDL

METHOD: Water samples were extracted for PAH following general NS&T methods. Full water samples were spiked with surrogates and extracted three times with dichloromethane using separatory funnel techniques. The combined extract was dried over anhydrous sodium sulfate, concentrated, processed through alumina cleanup column, concentrated, and further purified by GPC/HPLC. The post-HPLC extract was concentrated, fortified with Recovery Internal Standard (RIS) and split for analysis. Extracts were analyzed using gas chromatography/mass spectrometry (GC/MS) with the MS operating in the selected ion monitoring (SIM) mode, following general NS&T methods. Sample data were quantified by the method of internal standards, using the RIS compounds.

HOLDING TIMES: Samples were prepared for analysis in one analytical batch.

Samples were extracted with in the 7-day holding time for waters. Extracts were analyzed within the 40-day holding time for extracts

Batch	Extraction Date	Analysis Date
03-0203	3/4/2003	3/19/2003 – 3/20/2003

BLANKS: A procedural blank (PB) was prepared with each analytical batch. Blanks were analyzed to ensure the sample extraction and analysis methods were free of contamination.

03-0203 – Several target analytes were detected at concentrations greater than 3X the MDL.

Comments – All samples are appropriately flagged. The contamination in the blank does not appear to have the same PAH homologue pattern as the samples indicating that the contamination is likely isolated to the blank and that the samples are not impacted by the blank contamination. This is supported by the fact that no alkyl homologues were detected in the LCS (blank spike) sample – the LCS is prepared in the same manner as the blank, with the addition of a spike of the target analytes of interest (in this case, the parent PAH).

LABORATORY CONTROL Note: The 2003 MDL for substituted naphthalenes were updated.
 A laboratory control sample (LCS) was prepared with each analytical batch. The percent recoveries of target PAH were calculated to measure data quality in terms of accuracy.

SAMPLE:

03-0203 – All target analytes were recovered within the laboratory control limits specified by the client.

Comments – None.

**MATRIX
SPIKE/MATRIX
SPIKE
DUPLICATE:**

A matrix spike (MS)/matrix spike duplicate (MSD) pair was prepared with each analytical batch. The percent recoveries of target PAH were calculated to measure data quality in terms of accuracy; the relative percent difference between the pair was calculated to measure data quality in terms of precision.

03-0203 – All target analytes were recovered within the laboratory control limits specified by the client. The relative percent differences between the MS and MSD recoveries were within the laboratory control limits for all target PAH.

Comments – None.

SURROGATES:

Three surrogate compounds were added prior to extraction, including naphthalene-d8, phenanthrene-d10, and chrysene-d12. The recovery of each surrogate compound was calculated to measure data quality in terms of accuracy (extraction efficiency).

03-0203 – All surrogate percent recoveries were within the laboratory control limits specified by the client.

Comments – None.

PAHs QA/QC (CONT.)

CLIENT SAMPLE ID	LABORATORY CONTROL SAMPLE			MATRIX SPIKE- NAV-BAY9-SDB2- PRE		MATRIX SPIKE DUPLICATE-NAV-BAY9- SDB2-PRE			PROCEDURAL BLANK	
Battelle Sample ID	BB584LCS			U7067MS		U7067MSD			BB583PB	
Battelle Batch ID	03-0200			03-0200		03-0200			03-0200	
Data File	A1849.D			A1851.D		A1852.D			A1848.D	
Extraction Date	03/03/03			3/3/2003		03/03/03			03/03/03	
Acquired Date	03/18/03			3/18/2003		3/7698			03/18/03	
Matrix	Water			Water		Water			Water	
Sample Size (L)	2			0.65		0.65			2	
Dilution Factor	1.667			1.667		1.667			1.667	
PIV (mL)	0.3			0.30		0.30			0.3	
Min Reporting Limit	1.25			3.85		3.85			1.25	
Amount Units	ng	Rec%	Q	ng/L	Rec%	ng/L		Rec%	ng/L	
Naphthalene	368.77		74	1019.92		66	1010.15		65	1.89
C1-Naphthalenes	0.66	U	NA	2.04	U	NA	2.04	U	NA	1.14
C2-Naphthalenes	0.66	U	NA	2.04	U	NA	2.04	U	NA	0.66
C3-Naphthalenes	0.66	U	NA	2.04	U	NA	2.04	U	NA	0.66
C4-Naphthalenes	0.66	U	NA	2.04	U	NA	2.04	U	NA	0.66
2-Methylnaphthalene	371.20		74	983.20		64	982.48		64	0.95
1-Methylnaphthalene	347.39		69	946.06		61	940.09		61	0.63
2,6-Dimethylnaphthalene	354.90		71	916.90		60	894.33		58	0.47
2,3,5-Trimethylnaphthalene	378.08		76	1001.89		65	975.25		63	0.50
Biphenyl	373.48		74	958.19		62	931.34		60	0.42
Acenaphthylene	384.65		77	1086.17		70	1084.45		70	0.41
Acenaphthene	369.53		74	1021.95		66	1027.95		66	0.54
Fluorene	399.58		80	1119.79		73	1121.97		73	0.49
C1-Fluorenes	0.49	U	NA	1.51	U	NA	1.51	U	NA	0.49
C2-Fluorenes	0.49	U	NA	1.51	U	NA	1.51	U	NA	0.49
C3-Fluorenes	0.49	U	NA	1.51	U	NA	1.51	U	NA	0.49
Phenanthrene	416.86		83	1205.14		78	1189.70		77	0.53
Anthracene	418.43		84	1193.24		77	1170.26		76	0.35
C1-Phenanthrenes/Anthracenes	0.38	U	NA	1.15	U	NA	1.15	U	NA	0.38
C2-Phenanthrenes/Anthracenes	0.38	U	NA	1.15	U	NA	1.15	U	NA	0.38
C3-Phenanthrenes/Anthracenes	0.38	U	NA	1.15	U	NA	1.15	U	NA	0.38
C4-Phenanthrenes/Anthracenes	0.38	U	NA	1.15	U	NA	1.15	U	NA	0.38
1-Methylphenanthrene	409.14		82	1204.04		78	1194.14		78	0.43
Dibenzothiophene	0.46	U	NA	1.43	U	NA	1.43	U	NA	0.46
C1-Dibenzothiophenes	0.46	U	NA	1.43	U	NA	1.43	U	NA	0.46
C2-Dibenzothiophenes	0.46	U	NA	1.43	U	NA	1.43	U	NA	0.46
C3-Dibenzothiophenes	0.46	U	NA	1.43	U	NA	1.43	U	NA	0.46
Fluoranthene	447.78		90	1342.42		87	1330.47		86	0.39
Pyrene	460.98		92	1406.44		91	1361.94		88	0.48
C1-Fluoranthenes/Pyrenes	0.44	U	NA	1.35	U	NA	1.35	U	NA	0.44
C2-Fluoranthenes/Pyrenes	0.44	U	NA	1.35	U	NA	1.35	U	NA	0.44
C3-Fluoranthenes/Pyrenes	0.44	U	NA	1.35	U	NA	1.35	U	NA	0.44
Benzo(a)anthracene	466.38		93	1434.42		93	1368.53		89	0.69
Chrysene	467.52		93	1447.34		94	1385.71		90	0.37
C1-Chrysenes	0.37	U	NA	1.13	U	NA	1.13	U	NA	0.37
C2-Chrysenes	0.37	U	NA	1.13	U	NA	1.13	U	NA	0.37
C3-Chrysenes	0.37	U	NA	1.13	U	NA	1.13	U	NA	0.37
C4-Chrysenes	0.37	U	NA	1.13	U	NA	1.13	U	NA	0.37
Benzo(b)fluoranthene	456.45		91	1398.10		91	1354.66		88	0.43
Benzo(k)fluoranthene	458.55		92	1408.51		92	1333.50		87	0.43
Benzo(e)pyrene	404.89		82	1251.12		82	1188.80		78	0.45
Benzo(a)pyrene	440.30		88	1348.53		88	1269.82		82	0.65
Perylene	399.28		80	1197.11		78	1158.92		75	0.70
Indeno(1,2,3-cd)pyrene	422.43		84	1254.97		81	1235.79		80	0.80
Dibenz(a,h)anthracene	456.49		91	1395.33		91	1356.43		88	1.04
Benzo(g,h,i)perylene	342.98		69	1035.28		67	1012.59		66	1.31
Total Priority Pollutant PAHs										
Naphthalene-d8	74			69			72			76
Phenanthrene-d10	77			72			73			75
Chrysene-d12	90			88			86			91

PCBs

CLIENT SAMPLE ID:	NAV- PR5-SDB2- FF	NAV- PR5-SDB2- COMP	NAV- PR6-SDB2- FF	NAV- PR6-SDB2- COMP	NAV- OF9-SDB2- FF	NAV- OF9-SDB2- COMP	NAV- OF11-SDB2- FF	NAV- OF11-SDB2- COMP	NAV- OF14-SDB2- FF	NAV- OF14-SDB2- COMP
Battelle Sample ID:	U7089	U7091	U7090	U7092	U7083	U7086	U7084	U7087	U7085	U7088
Battelle Batch ID:	03-0203	03-0203	03-0203	03-0203	03-0203	03-0203	03-0203	03-0203	03-0203	03-0203
Data File:	sc0382.49.1	sc0382.51.1	sc0382.50.1	sc0382.52.1	sc0382.39.1	sc0382.44.1	sc0382.42.1	sc0382.47.1	sc0382.45.1	sc0382.48.1
Extraction Date:	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03	3/04/03
Acquired Date:	3/19/03	3/19/03	3/19/03	3/19/03	3/18/03	3/18/03	3/18/03	3/18/03	3/18/03	3/18/03
Matrix:	Water	Water	Water	Water	Water	Water	Water	Water	Water	Water
Sample Volume (L):	2.620	2.640	2.660	2.660	1.350	2.660	2.660	2.660	2.660	2.620
Dilution Factor:	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667	1.667
Pre Injection Volume (µL):	300	300	300	300	300	300	300	300	300	300
Minimum Reporting Limit (ng/L):	0.191	0.189	0.188	0.188	0.370	0.188	0.188	0.188	0.188	0.191
Units:	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L	ng/L
C12 08	1.833 U	11.781 NC	1.805 U	1.805 U	3.557 U	1.805 U	1.805 U	1.805 U	0.277 NC	15.762 U
C13 18	0.159 U	0.157 U	7.223 U	0.156 U	2.993 U	2.485 U	6.150 U	0.156 U	0.156 U	0.159 U
C13 28	0.198 U	0.197 U	0.195 U	0.195 U	0.385 U	0.195 U	0.195 U	0.195 U	0.195 U	0.198 U
C14 44	0.634 U	0.521 U	0.663 U	0.546 U	0.323 U	2.330 U	4.776 NC	0.649 U	1.120 U	0.166 U
C14 49	0.170 U	0.169 U	0.168 U	0.168 U	0.330 U	0.168 U	11.503 NC	0.168 U	0.168 U	0.170 U
C14 52	0.164 U	0.163 U	0.162 U	0.162 U	0.319 U	1.088 NC	0.162 U	0.162 U	0.162 U	0.164 U
C14 66	0.171 U	0.169 U	0.168 U	0.168 U	0.331 U	0.168 U	0.168 U	0.168 U	0.168 U	0.171 U
C14 77	0.243 U	0.489 NC	0.239 U	1.887 NC	1.361 NC	1.565 NC	3.689 NC	1.640 NC	1.375 U	1.171 NC
C15 87	0.177 NC	0.350 NC	0.127 U	0.152 NC	0.704 U	0.894 U	4.045 NC	0.232 NC	0.848 U	0.940 NC
C15 101	0.131 U	0.922 NC	0.129 U	1.659 NC	2.055 NC	2.818 U	11.255 U	1.184 U	3.290 U	0.131 U
C15 105	0.066 U	0.065 U	0.065 U	0.065 U	0.353 NC	1.513 U	2.188 NC	0.167 NC	1.153 U	0.215 NC
C15 114	0.113 U	0.112 U	0.111 U	0.111 U	0.219 U	0.111 U	1.623 U	0.111 U	0.111 U	0.113 U
C15 118	2.355 U	1.733 NC	3.206 NC	3.620 NC	0.808 NC	1.753 NC	5.469 NC	0.333 NC	3.117 NC	0.374 NC
C15 123	0.113 U	0.112 U	0.111 U	0.111 U	0.219 U	0.111 U	0.111 U	0.111 U	0.111 U	0.113 U
C15 126	0.141 U	0.140 U	0.139 U	0.139 U	0.273 U	0.139 U	0.139 U	0.139 U	0.139 U	0.141 U
C16 128	0.160 U	0.485 NC	0.508 NC	0.893 NC	1.336 NC	1.265 NC	4.836 NC	0.597 NC	2.128 NC	1.203 NC
C16 138	0.321 NC	0.814 NC	0.133 NC	0.504 U	2.362 U	3.719 NC	17.547 NC	1.316 NC	3.099 NC	1.803 NC
C16 153	0.122 U	0.231 U	0.120 U	1.069 U	1.883 U	5.412 U	21.543 NC	1.226 U	3.269 U	1.728 U
C16 156	0.137 U	0.136 U	0.135 U	0.130 U	0.265 U	0.226 U	0.786 U	0.135 U	0.612 U	0.137 U
C16 157	0.137 U	0.136 U	0.135 U	0.135 U	0.265 U	0.135 U	0.135 U	0.135 U	0.135 U	0.137 U
C16 167	0.137 U	0.426 U	0.135 U	0.389 U	1.059 U	2.020 U	5.665 U	0.889 U	3.229 U	0.137 U
C16 169	0.109 U	0.109 U	0.238 NC	0.108 U	0.212 U	0.108 U	0.108 U	0.108 U	0.108 U	0.109 U
C17 170	0.116 U	0.115 U	0.290 NC	0.114 U	0.861 U	1.424 U	8.071 NC	0.420 U	0.473 U	0.216 U
C17 180	0.110 U	0.109 U	0.427 U	0.108 U	2.074 U	3.483 U	18.727 NC	1.264 U	1.565 U	0.110 U
C17 183	0.107 U	0.106 U	0.105 U	0.105 U	0.207 U	0.466 U	5.751 NC	0.105 U	0.105 U	0.107 U
C17 184	0.105 U	0.104 U	0.104 U	0.104 U	0.204 U	0.104 U	0.104 U	0.104 U	0.119 NC	0.105 U
C17 187	0.099 U	0.182 U	0.097 U	0.384 U	0.694 U	0.887 U	7.818 U	0.345 U	0.466 U	0.247 U
C17 189	0.108 U	0.107 U	0.106 U	0.106 U	0.209 U	0.106 U	0.106 U	0.106 U	0.106 U	0.108 U
C18 195	0.124 U	0.123 U	0.122 U	0.122 U	0.241 U	0.236 NC	1.453 NC	0.061 NC	0.122 U	0.095 NC
C19 206	0.139 U	0.138 U	0.371 U	0.137 U	0.270 U	0.137 U	2.957 U	0.137 U	0.457 U	0.139 U
C10 209	0.702 U	0.772 U	0.593 U	0.143 U	1.427 NC	0.143 U	6.227 U	0.143 U	1.794 U	0.145 U
Total PCB	4.190	18.281	13.652	10.846	18.911	31.563	144.788	9.432	25.162	23.753
Surrogate Recoveries:										
C13(34)	80	79	75	83	86	81	91	91	85	82
C15(112)	79	80	71	80	78	76	93	81	78	82

PCBs QA/QC

PROJECT: SPAWAR TO0011, Contaminant Analysis of Stormwater and San Diego Bay Seawater
PARAMETER: PCB Congener
LABORATORY: Battelle, Duxbury, MA
MATRIX: Water
SAMPLE CUSTODY: The water samples were collected on February 25, 2003. They were received in Duxbury on February 28, 2003 in good condition in six coolers. The cooler temperature on arrival ranged from 0.2 °C to 1.3 °C. Samples were stored at 4 °C until processing.

QA/QC DATA QUALITY OBJECTIVES:

	Reference Method	Surrogate Recovery	LCS/MS Recovery	Sample Replicate Relative Precision	Procedural Blank
PAH	General NS&T	30-130% Recovery	LCS: 40-120% Recovery for at least 80% of analytes MS: 50-150% Recovery for at least 70% of analytes; analyte conc. in MS must be >5x background	≤30% RSD analyte conc. in MS must be <5x background	<3X MDL

METHOD: Water samples were extracted for PCB Congener following general NS&T methods. Full water samples were spiked with surrogates and extracted three times with dichloromethane using separatory funnel techniques. The combined extract was dried over anhydrous sodium sulfate, concentrated, processed through alumina cleanup column, concentrated, and further purified by GPC/HPLC. The post-HPLC extract was concentrated, fortified with Recovery Internal Standard (RIS) and split for analysis. Extracts were analyzed using gas chromatography/electron capture detection (GC/ECD), following general NS&T methods. Sample data were quantified by the method of internal standards, using the RIS compounds.

HOLDING TIMES: Samples were prepared for analysis in one analytical batch.

Samples were extracted with in the 7-day holding time for waters. Extracts were analyzed within the 40-day holding time for extracts

<u>Batch</u>	<u>Extraction Date</u>	<u>Analysis Date</u>
03-0203	3/4/2003	3/17/2003 – 3/22/2003

BLANKS: A procedural blank (PB) was prepared with each analytical batch. Blanks were analyzed to ensure the sample extraction and analysis methods were free of contamination.

03-0203 – No analytes identified at greater than 3X the MDL.

LABORATORY CONTROL SAMPLE: **Comments** – None.
A laboratory control sample (LCS) was prepared with each analytical batch. The percent recoveries of target PCB Congeners were calculated to measure data quality in terms of accuracy.

03-0203 – All target analytes were recovered within the laboratory control limits specified by the client.

MATRIX SPIKE/MATRIX SPIKE DUPLICATE: **Comments – None.**
A matrix spike (MS)/matrix spike duplicate (MSD) pair was prepared with each analytical batch. The percent recoveries of target PCB Congeners were calculated to measure data quality in terms of accuracy; the relative percent difference between the pair was calculated to measure data quality in terms of precision.

03-0203 – All target analytes were recovered within the laboratory control limits specified by the client. The relative percent differences between the MS and MSD recoveries were within the laboratory control limits for all target PCB Congeners.

SURROGATES: **Comments – None.**
Two surrogate compounds were added prior to extraction, including PCB34 and PCB112. The recovery of each surrogate compound was calculated to measure data quality in terms of accuracy (extraction efficiency).

03-0203 – All surrogate percent recoveries were within the laboratory control limits specified by the client.

Samples: **Comments – None.**
The condition of the confirmation column was in question after the analysis. It was decided to report all “hits” from the primary column regardless if confirmed or not confirmed. The analytes are appropriately flagged if reported, but not confirmed.

PCBs QA/QC (CONT.)

CLIENT SAMPLE ID:	LABORATORY CONTROL SAMPLE		MATRIX SPIKE-NAV-OF9-SDB2-FF		MATRIX SPIKE DUPLICATE-NAV-OF9-SDB2-FF				PROCEDURAL BLANK
Battelle Sample ID:	BB593LCS		U7083MS		U7083MSD				BB592PB
Battelle Batch ID:	03-0203		03-0203		03-0203				03-0203
Data File:	sc0382,38,1		sc0382,40,1		sc0382,41,1				sc0382,37,1
Extraction Date:	3/04/03		3/04/03		3/04/03				3/04/03
Acquired Date:	3/17/03		3/18/03		3/18/03				03/17/03
Matrix:	Water		Water		Water				Water
Sample Volume (L):	2.000		1.175		1.175				2.000
Dilution Factor:	1.667		1.667		1.667				1.667
Pre Injection Volume (µL):	300		300		300				300
Minimum Reporting Limit (ng/L):	0.250		0.426		0.426				0.250
Units:	ng	% Recovery	ng/L	% Recovery	ng/L	% Recovery	% RPD		ng/L
C12 08	20.509	68	17.345	68	20.449	80	16		2.401 U
C13 18	21.498	72	19.854	66	20.959	70	6		0.208 U
C13 28	35.064	117	21.372	84	25.258	99	17		0.260 U
C14 44	25.535	85	21.262	83	23.531	92	10		0.218 U
C14 49	25.348	84	24.907	97	25.302	99	2		0.223 U
C14 52	24.280	81	20.705	81	21.345	84	3		0.215 U
C14 66	27.632	92	22.603	89	25.267	99	11		0.224 U
C14 77	24.023	80	20.028	73	22.256	82	11		0.318 U
C15 87	24.470	82	20.388	77	21.738	82	7		0.169 U
C15 101	25.400	85	21.352	76	23.998	86	13		0.172 U
C15 105	26.157	87	20.806	80	21.916	84	5		0.086 U
C15 114	NA	NA	NS	NA	NS	NA	NA		0.148 U
C15 118	23.286	78	19.969	75	21.834	82	9		0.130 U
C15 123	NA	NA	NS	NA	NS	NA	NA		0.148 U
C15 126	28.227	94	19.566	77	20.869	82	6		0.184 U
C16 128	26.487	88	23.105	85	21.485	79	8		0.210 U
C16 138	25.310	84	22.841	80	24.760	88	9		0.199 U
C16 153	22.656	76	22.155	79	23.904	86	8		0.160 U
C16 156	NA	NA	NS	NA	NS	NA	NA		0.179 U
C16 157	NA	NA	NS	NA	NS	NA	NA		0.179 U
C16 167	NA	NA	NS	NA	NS	NA	NA		0.179 U
C16 169	28.949	96	23.429	91	25.035	98	7		0.143 U
C17 170	25.778	86	22.468	85	24.052	91	7		0.152 U
C17 180	25.907	86	23.798	85	26.011	94	10		0.144 U
C17 183	25.158	84	21.384	84	22.322	87	4		0.140 U
C17 184	23.828	79	19.885	78	20.606	80	4		0.138 U
C17 187	23.085	77	19.872	75	20.867	79	5		0.129 U
C17 189	NA	NA	NS	NA	NS	NA	NA		0.141 U
C18 195	25.317	84	20.354	80	21.844	86	7		0.163 U
C19 206	23.978	80	20.081	79	21.885	86	9		0.183 U
C10 209	23.396	78	18.226	66	19.594	71	8		0.190 U
Total PCB	631.280	NA	527.755	NA	567.085	NA	NA		0.000
Surrogate Recoveries:									
C13(34)	84		90		99				78
C15(112)	78		74		77				78

SDB4- 10/17/2004

METALS

SAMPLE ID	DISSOLVED Cu (µg/L)	TOTAL Cu (µg/L)	DISSOLVED Zn (µg/L)	TOTAL Zn (µg/L)
NAV-BAY SDB4 PRE	3.0	3.5	9.5	12
NAV-OF11-SDB4-FF	89	244	2453	3631
NAV-BAY14-SDB4-DUR	14	21	182	238

TSS

SAMPLE LABEL	TSS (mg/L)
NAV-OF14-SDB4-FF	838.75
NAV-BAY14-SDB4-DUR	20.66

SDB45- 10/26/2004

METALS

MSL Code	Rep	Sponsor I.D.	Al (µg/L) ICP-OES	Fe (µg/L) ICP-OES	Cr (µg/L) ICP-MS	Mn (µg/L) ICP-OES	Ni (µg/L) ICP-MS	Cu (µg/L) ICP-MS
SAMPLE RESULTS								
2173*27		NAV-OF14-SD45-FF	1322	2138	6.93	66.2	7.19	45.3
2173*28		NAV-OF14-SD45-FF (F)	14.7	26.4	2.22	29.2	3.67	18.9
2173*29		NAV-OF14-SD45-COMP	2618	4481	12.9	71.5	4.81	38.0
2173*30		NAV-OF14-SD45-COMP (F)	17.7	25.0	9.99	13.2	1.66	9.89
2173*31		NAV-OF14-SD45-Blank (F)	3.36 U	2.51 U	0.724 U	0.025 U	0.009 U	0.008 U

MSL Code	Rep	Sponsor I.D.	Zn (µg/L) ICP-MS	As (µg/L) ICP-MS	Se (µg/L) ICP-MS	Ag (µg/L) ICP-MS	Cd (µg/L) ICP-MS	Sn (µg/L) ICP-MS	Pb (µg/L) ICP-MS	Hg (µg/L) CVAF
SAMPLE RESULTS										
2173*27		NAV-OF14-SD45-FF	362	3.20	1.30	0.0741	1.18	0.663	21.7	0.0629
2173*28		NAV-OF14-SD45-FF (F)	175	2.04	0.848	0.00601	0.492	0.50 U	0.493	0.00597
2173*29		NAV-OF14-SD45-COMP	220	2.39	0.530	0.0632	0.871	0.536	21.6	0.0694
2173*30		NAV-OF14-SD45-COMP (F)	68.4	1.72	0.356	0.00378	0.244	0.50 U	0.441	0.00330
2173*31		NAV-OF14-SD45-Blank (F)	0.0713	0.015 U	0.101 U	0.002 U	0.002 U	0.50 U	0.00266	0.000510

SAMPLE ID	DISSOLVED COPPER (µg/L)	TOTAL COPPER (µg/L)	DISSOLVED ZINC (µg/L)	TOTAL ZINC (µg/L)
NAV-BAY-SD45-PRE	3.94	6.97	7.79	8.42
NAV-BAY14-SD45-DUR 1	4.32	7.05	10.21	12.79
NAV-BAY14-SD45-DUR 2	4.32	6.19	9.15	9.26
NAV-BAY14-SD45-DUR 3	4.55	6.82	9.06	9.51
NAV-BAY14-SD45-DUR 4	3.32	5.89	7.70	8.60
NAV-BAY14-SD45-AFT 1	3.41	6.05	8.07	8.98
NAV-BAY14-SD45-AFT 2	3.66	5.97	7.89	8.98
NAV-BAY14-SD45-AFT 3	3.98	6.26	10.08	11.22