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Nitrate_plus_Nitrite:Bottle

Precision statement for replicate samples drawn from a single Niskin bottle:

The pooled standard deviation for Nitrate_plus_Nitrite:Bottle for the range 0.0 to 43.1 $\mu\text{mol/l}$ was 0.17, $k = 24$ (2 outliers removed) where k is the number of pairs of duplicates.

The pooled standard deviation of pairs of samples (S_p) was calculated by:

$$S_p = \text{SQRT}\{\text{sum}(d^2)/2k\}$$

where k is the number of pairs and d is the difference between pairs.

Nitrate data from 1000 to 4000 m agrees with results averaged over 16 profiles at P26 from 1995 to 2001 to within 0.13 %.

Duplicate samples from a single Niskin bottle

Event Number	Sample Number	Station	Pressure dbar	Nitrate 1 $\mu\text{mol/l}$	Nitrate 2 $\mu\text{mol/l}$	Rejected yes / no	Comment
1	8	SI03	52.0	28.5	28.4		
4	42	P2	24.9	22.2	22.4		
8	60	P4	15.7	0.1	0.0		
14	108	P8	150.9	26.8	26.9		
19	129	P12	1250.9	46.3	44.9	yes	Bad duplicate for nitrate
22	164	P12	30.5	7.5	7.4		
27	194	P16	174.7	27.9	27.9		
28	217	P16	2.5	4.7	4.7		
35	238	P20	198.7	31.4	33.3	yes	Bad duplicate for nitrate
47	319	P26	19.9	10.7	10.6		
48	344	P26	126.8	30.4	30.6		
51	351	P26	300.4	43.1	43.1		
55	397	P26	75.1	17.4	17.5		
59	448	P33	123.9	32.9	32.1		
64	470	R17	125.1	29.6	29.8		
69	491	R13	151.4	25.6	25.5		
76	508	R9N	400.0	36.5	36.2		
85	573	SS1	245.3	34.1	34.1		
88	590	SS3	124.4	20.5	20.6		
90	601	SS5	127.0	29.8	29.8		
93	612	SS7	75.7	28.2	28.3		
94	620	Ri1	200.4	29.2	29.1		
101	635	Ri4	175.3	28.4	29.0		
106	649	M2	100.1	29.9	30.1		
107	658	M3	200.1	31.4	31.5		
108	679	M4	25.2	21.2	21.0		

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Phosphate:Bottle

Precision statement for replicate samples drawn from a single Niskin bottle:

The pooled standard deviation for Phosphate:Bottle for the range 0.30 to 3.31 $\mu\text{mol/l}$ was 0.005,
k = 26 (0 outlier removed) where k is the number of pairs of duplicates.

The pooled standard deviation of pairs of samples (S_p) was calculated by:

$$S_p = \text{SQRT}\{\text{sum } (d^*d)/2k\}$$

where k is the number of pairs and d is the difference between pairs.

Phosphate data from 1000 to 4000 m agrees with results averaged over 16 profiles at P26 from 1995 to 2001
to within 1.78 %.

Duplicate samples from a single Niskin bottle

Event Number	Sample Number	Station	Pressure dbar	Phosphate 1 $\mu\text{mol/l}$	Phosphate 2 $\mu\text{mol/l}$	Rejected yes / no	Comment
1	8	SI03	52.0	2.35	2.35		
4	42	P2	24.9	1.77	1.78		
8	60	P4	15.7	0.31	0.30		
14	108	P8	150.9	2.04	2.04		
19	129	P12	1250.9	3.29	3.31		
22	164	P12	30.5	0.98	0.98		
27	194	P16	174.7	1.97	1.96		
28	217	P16	2.5	0.72	0.72		
35	238	P20	198.7	2.41	2.42		
47	319	P26	19.9	1.07	1.07		
48	344	P26	126.8	2.19	2.20		
51	351	P26	300.4	2.93	2.93		
55	397	P26	75.1	1.48	1.48		
59	448	P33	123.9	2.27	2.27		
64	470	R17	125.1	2.20	2.21		
69	491	R13	151.4	1.93	1.93		
76	508	R9N	400.0	2.64	2.64		
85	573	SS1	245.3	2.59	2.58		
88	590	SS3	124.4	1.71	1.71		
90	601	SS5	127.0	2.37	2.37		
93	612	SS7	75.7	2.38	2.38		
94	620	Ri1	200.4	2.56	2.56		
101	635	Ri4	175.3	2.54	2.55		
106	649	M2	100.1	2.44	2.43		
107	658	M3	200.1	2.48	2.48		
108	679	M4	25.2	1.71	1.71		

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Silicate:Bottle

Precision statement for replicate samples drawn from a single Niskin bottle:

The pooled standard deviation for Silicate:Bottle for the range 9.4 to 158.1 $\mu\text{mol/l}$ was 0.45,
k = 26 (0 outlier removed) where k is the number of pairs of duplicates.

The pooled standard deviation of pairs of samples (S_p) was calculated by:

$$S_p = \text{SQRT}\{\text{sum } (d^*d)/2k\}$$

where k is the number of pairs and d is the difference between pairs.

Silicate data from 1000 to 4000 m agrees with results averaged over 16 profiles at P26 from 1995 to 2001 to within 0.59 %

Duplicate samples from a single Niskin bottle

Event Number	Sample Number	Station	Pressure dbar	Silicate 1 $\mu\text{mol/l}$	Silicate 2 $\mu\text{mol/l}$	Rejected yes / no	Comment
1	8	SI03	52.0	51.7	51.7		
4	42	P2	24.9	33.4	33.3		
8	60	P4	15.7	10.0	10.0		
14	108	P8	150.9	34.4	34.4		
19	129	P12	1250.9	158.1	158.1		
22	164	P12	30.5	9.5	9.4		
27	194	P16	174.7	46.8	46.7		
28	217	P16	2.5	10.9	10.8		
35	238	P20	198.7	61.7	61.8		
47	319	P26	19.9	16.8	16.9		
48	344	P26	126.8	50.1	50.2		
51	351	P26	300.4	85.7	86.9		
55	397	P26	75.1	25.6	25.7		
59	448	P33	123.9	49.7	50.8		
64	470	R17	125.1	50.8	49.3		
69	491	R13	151.4	39.8	40.5		
76	508	R9N	400.0	60.7	60.9		
85	573	SS1	245.3	57.7	57.0		
88	590	SS3	124.4	31.4	29.4		
90	601	SS5	127.0	51.6	51.9		
93	612	SS7	75.7	52.8	52.8		
94	620	Ri1	200.4	58.7	58.7		
101	635	Ri4	175.3	58.4	58.8		
106	649	M2	100.1	55.7	55.1		
107	658	M3	200.1	57.9	58.0		
108	679	M4	25.2	39.7	39.7		