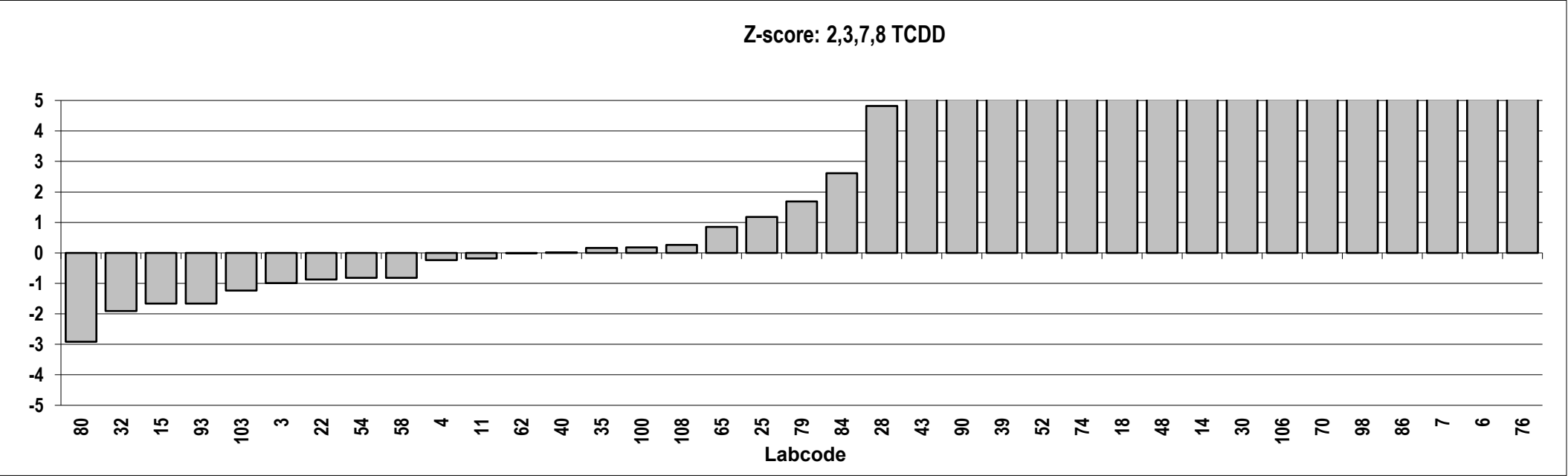
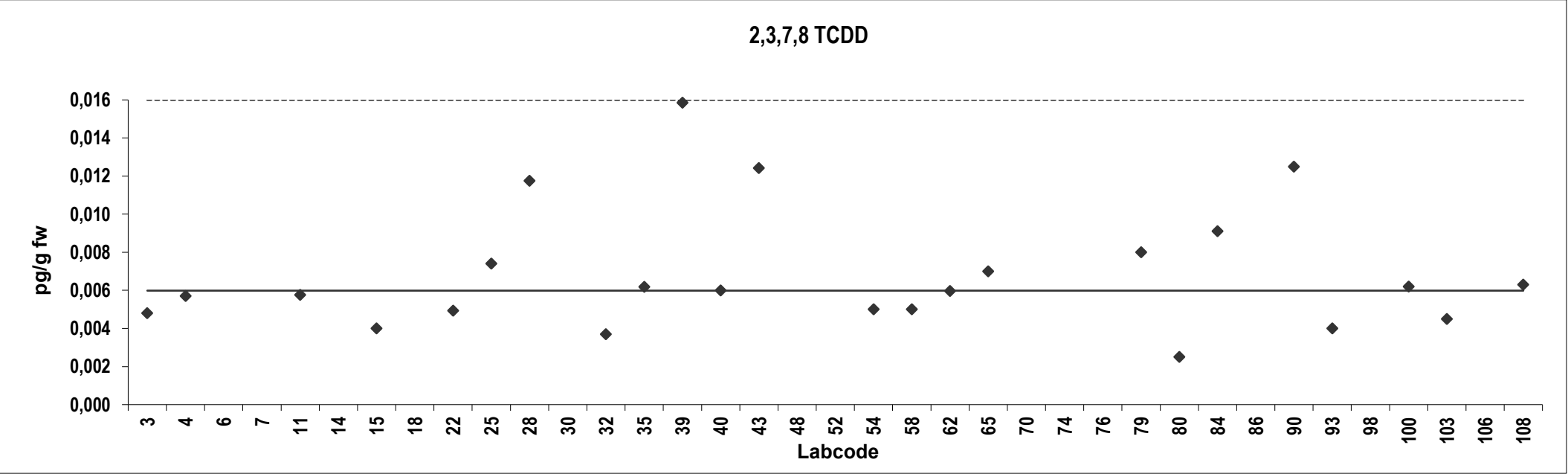


Salmon
Congener: 2,3,7,8 TCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0048	-0,99				
4	0,0057	-0,24				
6	0,15	121	Outlier			
7	0,11	83	Outlier,ND			
11	0,0058	-0,18				
14	0,038	26	Outlier			
15	0,0040	-1,7	ND			
18	0,020	12	Outlier,ND			
22	0,0049	-0,87				
25	0,0074	1,2				
28	0,012	4,8				
30	0,040	28	Outlier,ND			
32	0,0037	-1,9				
35	0,0062	0,16	ND			
39	0,016	8,2				
40	0,0060	0,012	ND			
43	0,012	5,4				
48	0,030	20	Outlier,ND			
52	0,016	8,4	Outlier,ND			
54	0,0050	-0,82				
58	0,0050	-0,82				
62	0,0060	-0,012				
65	0,0070	0,85	ND			
70	0,063	48	Outlier,ND			
74	0,017	8,9	Outlier,ND			
76	0,39	317	Outlier			
79	0,0080	1,7				
80	0,0025	-2,9				
84	0,0091	2,6	ND			
86	0,074	57	Outlier,ND			
90	0,013	5,4				
93	0,0040	-1,7	ND			
98	0,064	48	Outlier,ND			
100	0,0062	0,18				
103	0,0045	-1,2				
106	0,060	45	Outlier			
108	0,0063	0,26				

Consensus statistics

Consensus median, pg/g	0,0060
Median all values pg/g	0,0080
Consensus mean, pg/g	0,0069
Standard deviation, pg/g	0,0033
Relative standard deviation, %	48
No. of values reported	37
No. of values removed	13
No. of reported non-detects	15

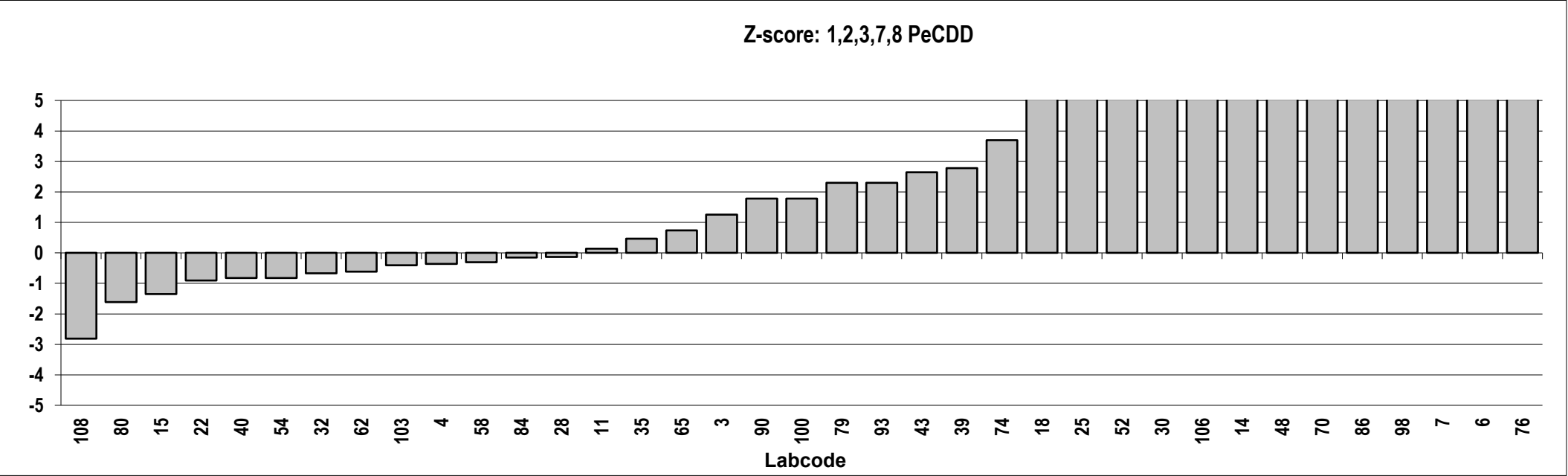
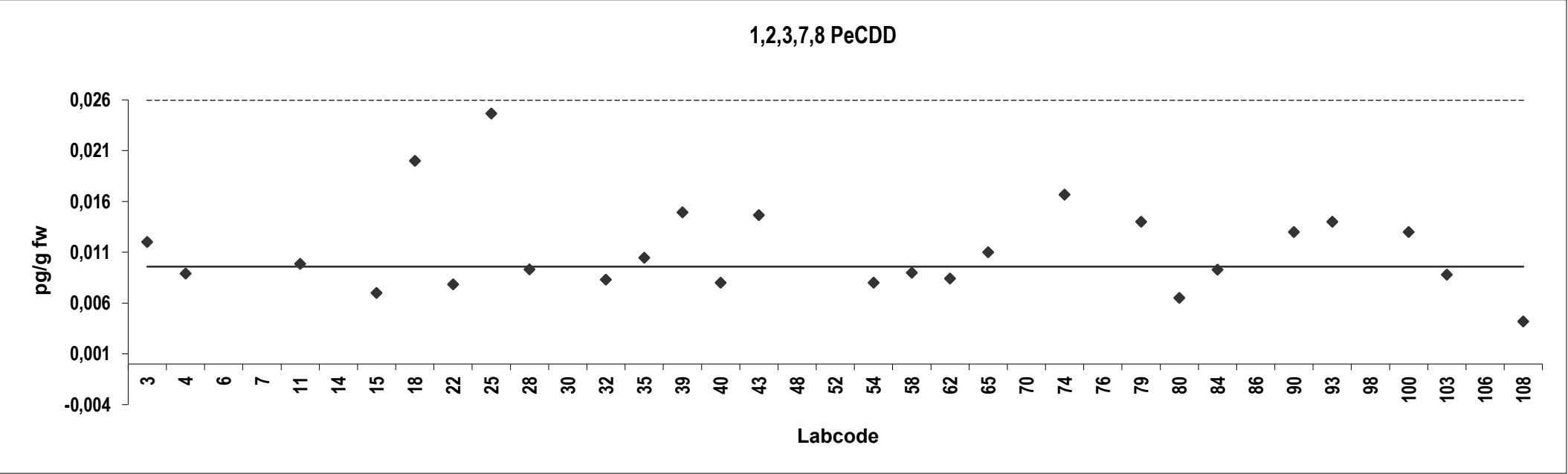


Salmon
Congener: 1,2,3,7,8 PeCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,012	1,3				
4	0,0089	-0,36				
6	0,27	138	Outlier			
7	0,15	75	Outlier,ND			
11	0,0098	0,14				
14	0,044	18	Outlier			
15	0,0070	-1,3	ND			
18	0,020	5,4	ND			
22	0,0078	-0,91				
25	0,025	7,9				
28	0,0093	-0,14	ND			
30	0,040	16	Outlier,ND			
32	0,0083	-0,67				
35	0,010	0,46	ND			
39	0,015	2,8				
40	0,0080	-0,83				
43	0,015	2,6				
48	0,056	24	Outlier,ND			
52	0,032	12	Outlier,ND			
54	0,0080	-0,83				
58	0,0090	-0,31				
62	0,0084	-0,61				
65	0,011	0,74				
70	0,063	28	Outlier,ND			
74	0,017	3,7	ND			
76	0,60	310	Outlier			
79	0,014	2,3				
80	0,0065	-1,6				
84	0,0093	-0,15	ND			
86	0,074	34	Outlier,ND			
90	0,013	1,8				
93	0,014	2,3	ND			
98	0,093	43	Outlier,ND			
100	0,013	1,8				
103	0,0088	-0,41				
106	0,040	16	Outlier			
108	0,0042	-2,8	ND			

Consensus statistics

Consensus median, pg/g	0,010
Median all values pg/g	0,013
Consensus mean, pg/g	0,011
Standard deviation, pg/g	0,0045
Relative standard deviation, %	40
No. of values reported	37
No. of values removed	11
No. of reported non-detects	15

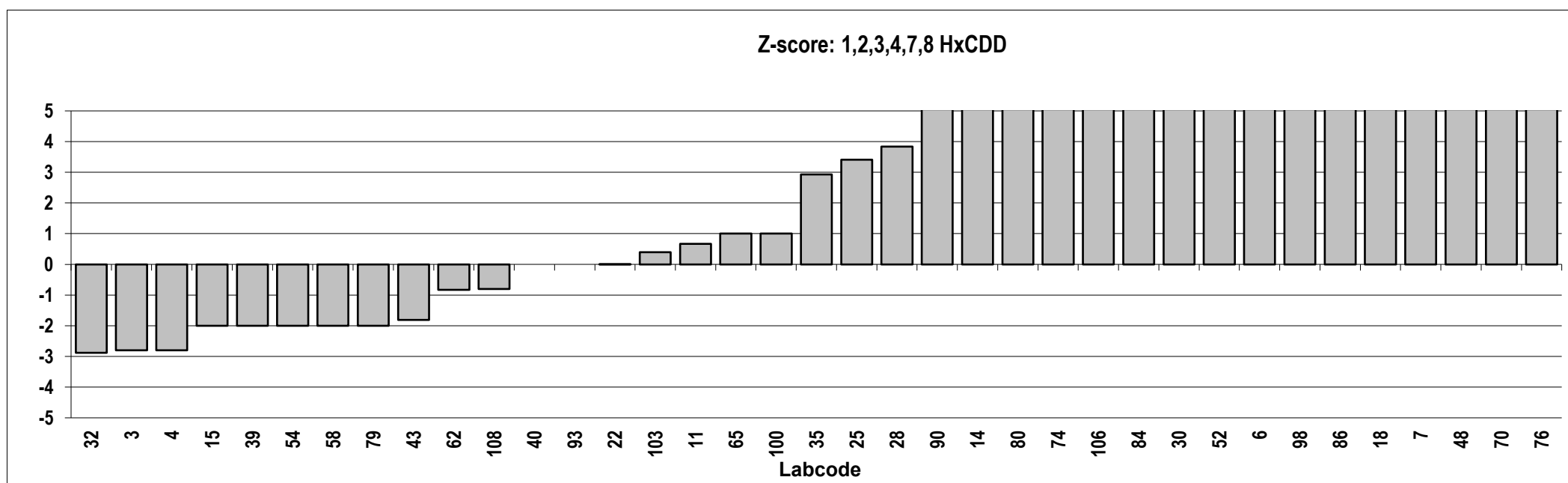
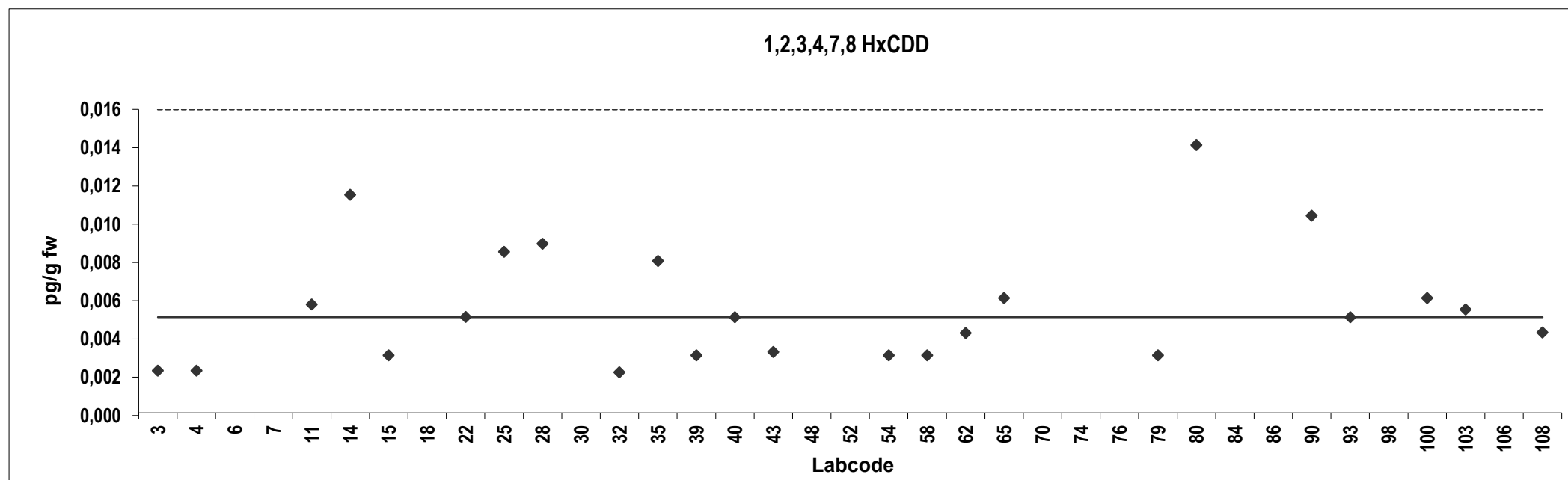


Salmon
Congener: 1,2,3,4,7,8 HxCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0022	-2,8				
4	0,0022	-2,8	ND			
6	0,065	60	Outlier			
7	0,12	115	Outlier,ND			
11	0,0057	0,67	ND			
14	0,011	6,4				
15	0,0030	-2,0	ND			
18	0,10	95	Outlier,ND			
22	0,0050	0,0059				
25	0,0084	3,4				
28	0,0088	3,8	ND			
30	0,050	45	Outlier,ND			
32	0,0021	-2,9				
35	0,0079	2,9	ND			
39	0,0030	-2,0	ND			
40	0,0050	0,0000	ND			
43	0,0032	-1,8				
48	0,12	115	Outlier,ND			
52	0,056	51	Outlier,ND			
54	0,0030	-2,0				
58	0,0030	-2,0				
62	0,0042	-0,83				
65	0,0060	1,0	ND			
70	0,13	125	Outlier,ND			
74	0,017	12	Outlier,ND			
76	1,7	1706	Outlier			
79	0,0030	-2,0				
80	0,014	9,0				
84	0,023	18	Outlier,ND			
86	0,099	94	Outlier,ND			
90	0,010	5,3	ND			
93	0,0050	0,0000	ND			
98	0,067	62	Outlier,ND			
100	0,0060	1,0				
103	0,0054	0,40				
106	0,020	15	Outlier,ND			
108	0,0042	-0,80	ND			

Consensus statistics

Consensus median, pg/g	0,0050
Median all values pg/g	0,0079
Consensus mean, pg/g	0,0055
Standard deviation, pg/g	0,0032
Relative standard deviation, %	58
No. of values reported	37
No. of values removed	13
No. of reported non-detects	22

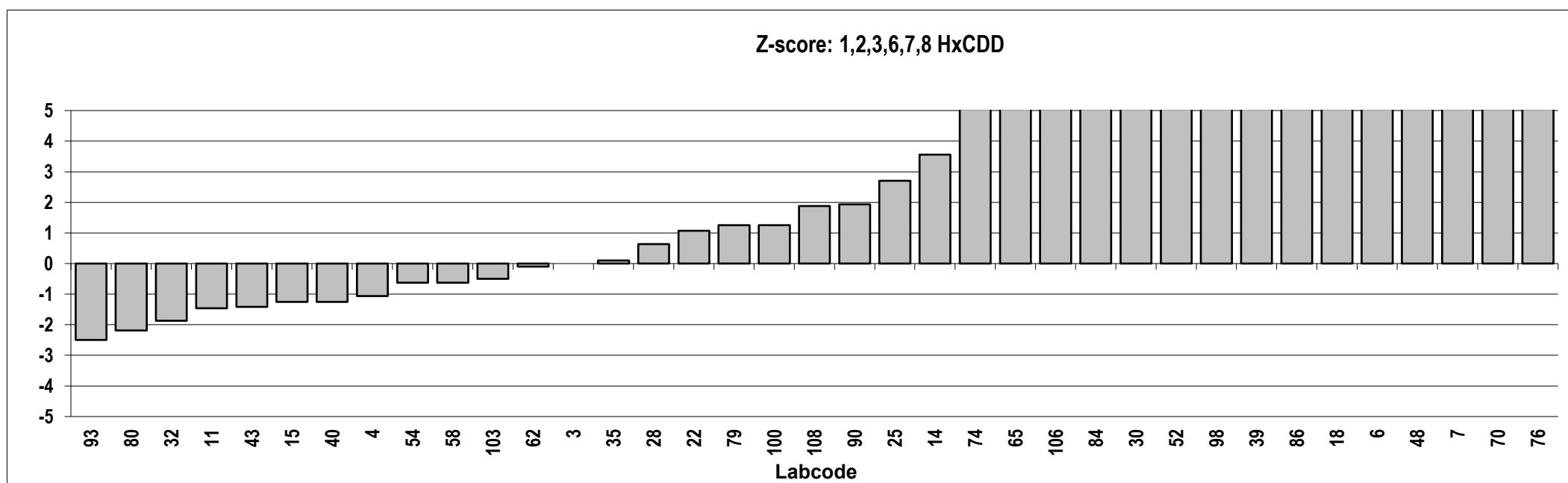
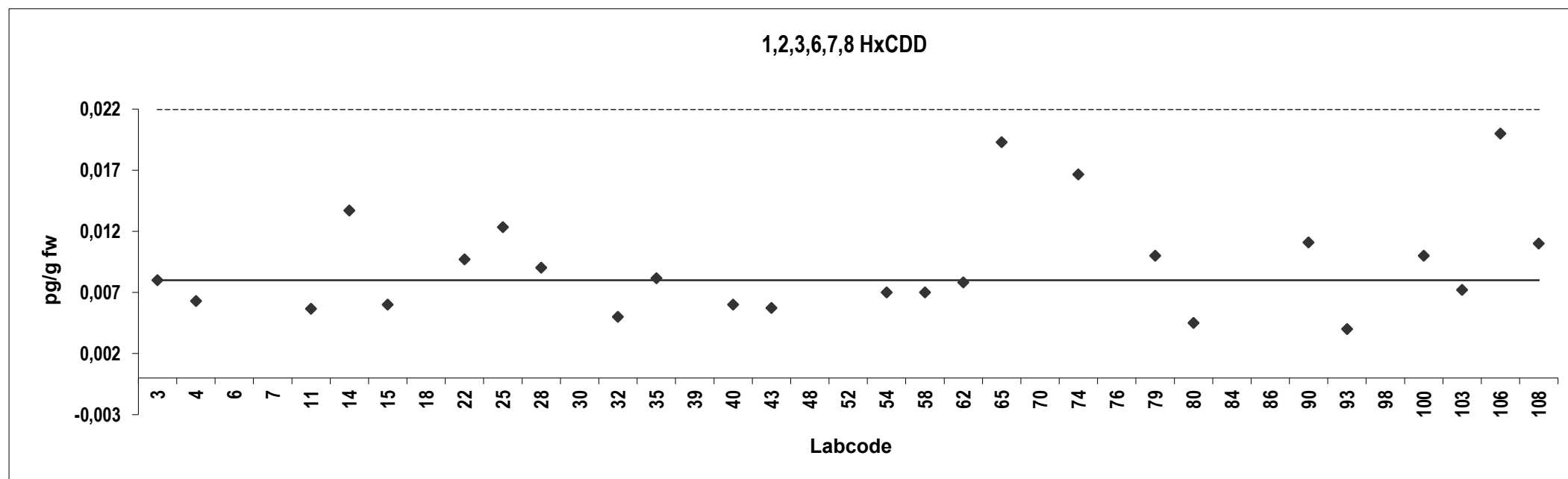


Salmon
Congener: 1,2,3,6,7,8 HxCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0080	0,0000				
4	0,0063	-1,1				
6	0,11	61	Outlier			
7	0,12	73	Outlier,ND			
11	0,0057	-1,5	ND			
14	0,014	3,6				
15	0,0060	-1,3				
18	0,10	58	Outlier,ND			
22	0,0097	1,1				
25	0,012	2,7				
28	0,0090	0,64	ND			
30	0,050	26	Outlier,ND			
32	0,0050	-1,9				
35	0,0082	0,10	ND			
39	0,081	46	Outlier			
40	0,0060	-1,3				
43	0,0057	-1,4				
48	0,11	64	Outlier,ND			
52	0,055	30	Outlier,ND			
54	0,0070	-0,63				
58	0,0070	-0,63				
62	0,0078	-0,10				
65	0,019	7,1				
70	0,13	76	Outlier,ND			
74	0,017	5,4	ND			
76	1,5	933	Outlier			
79	0,010	1,3				
80	0,0045	-2,2	ND			
84	0,023	9,2	Outlier,ND			
86	0,099	57	Outlier,ND			
90	0,011	1,9	ND			
93	0,0040	-2,5	ND			
98	0,056	30	Outlier,ND			
100	0,010	1,3				
103	0,0072	-0,50				
106	0,020	7,5	ND			
108	0,011	1,9				

Consensus statistics

Consensus median, pg/g	0,0080
Median all values pg/g	0,011
Consensus mean, pg/g	0,0092
Standard deviation, pg/g	0,0043
Relative standard deviation, %	47
No. of values reported	37
No. of values removed	12
No. of reported non-detects	17

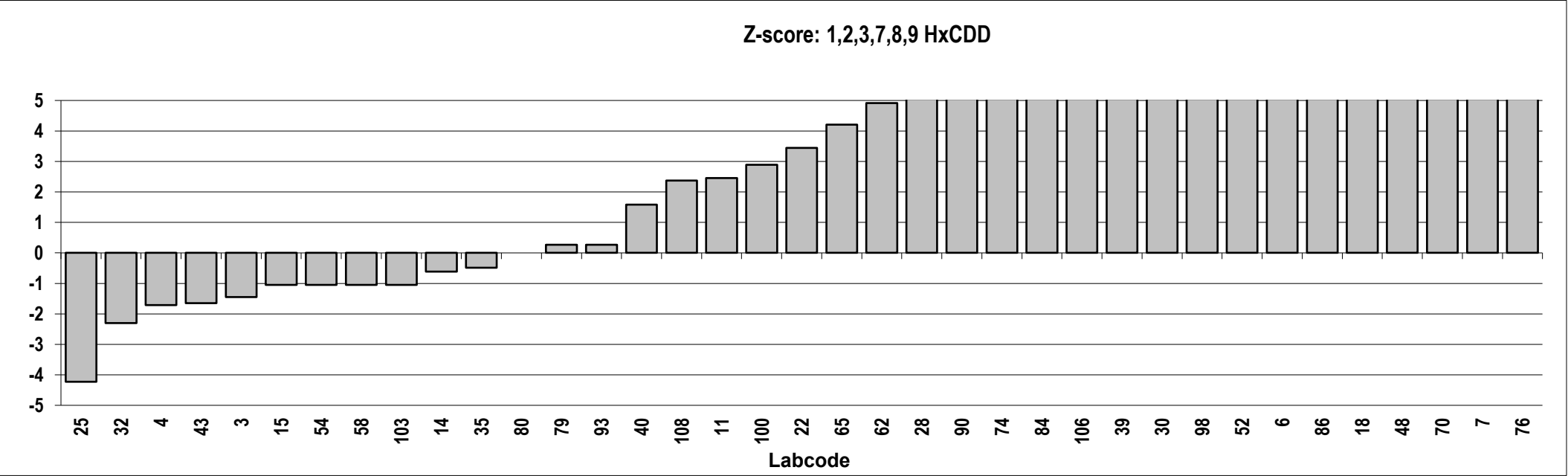
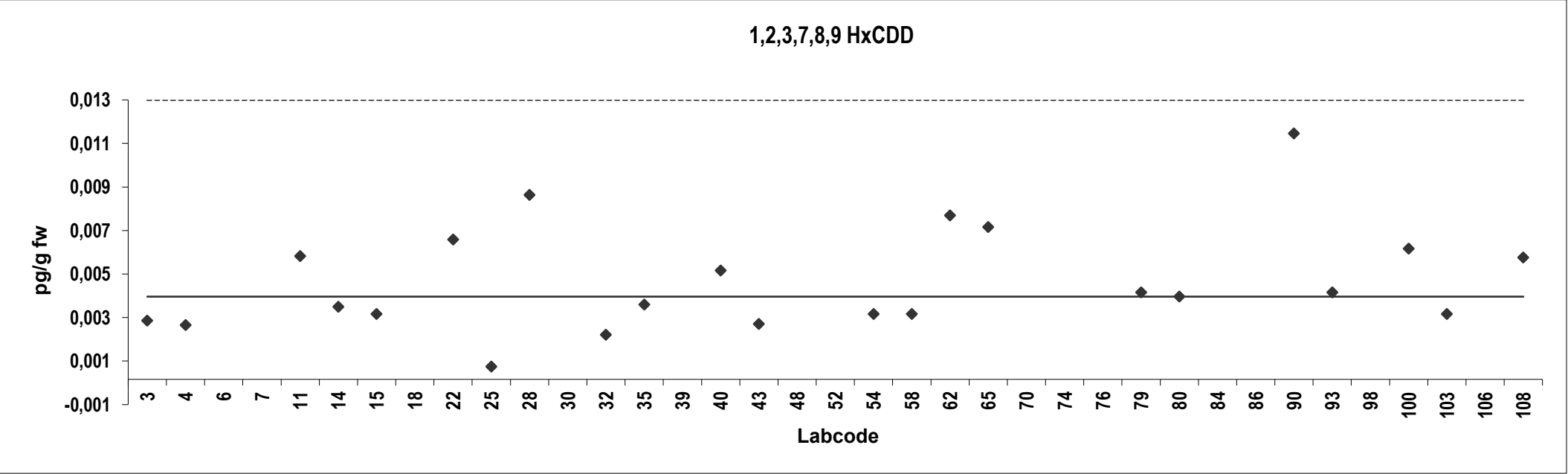


Salmon
Congener: 1,2,3,7,8,9 HxCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0027	-1,4				
4	0,0025	-1,7	ND			
6	0,075	94	Outlier			
7	0,13	167	Outlier,ND			
11	0,0057	2,5	ND			
14	0,0033	-0,62	ND			
15	0,0030	-1,1	ND			
18	0,10	127	Outlier,ND			
22	0,0064	3,4				
25	0,00059	-4,2				
28	0,0085	6,1	ND			
30	0,050	61	Outlier,ND			
32	0,0021	-2,3				
35	0,0034	-0,49	ND			
39	0,040	47	Outlier			
40	0,0050	1,6	ND			
43	0,0025	-1,6				
48	0,13	166	Outlier,ND			
52	0,064	79	Outlier,ND			
54	0,0030	-1,1	ND			
58	0,0030	-1,1	ND			
62	0,0075	4,9				
65	0,0070	4,2	ND			
70	0,13	166	Outlier,ND			
74	0,017	17	Outlier,ND			
76	1,5	2031	Outlier			
79	0,0040	0,26				
80	0,0038	0,0000	ND			
84	0,019	19	Outlier,ND			
86	0,099	125	Outlier,ND			
90	0,011	9,9	ND			
93	0,0040	0,26	ND			
98	0,062	76	Outlier,ND			
100	0,0060	2,9	ND			
103	0,0030	-1,1	ND			
106	0,020	21	Outlier,ND			
108	0,0056	2,4				

Consensus statistics

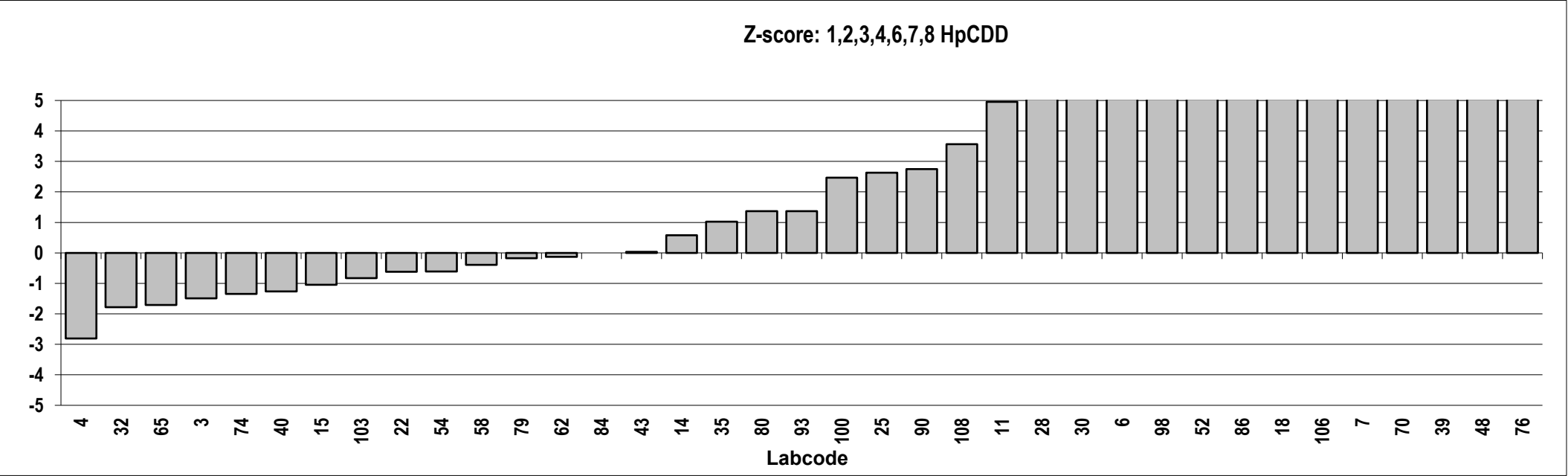
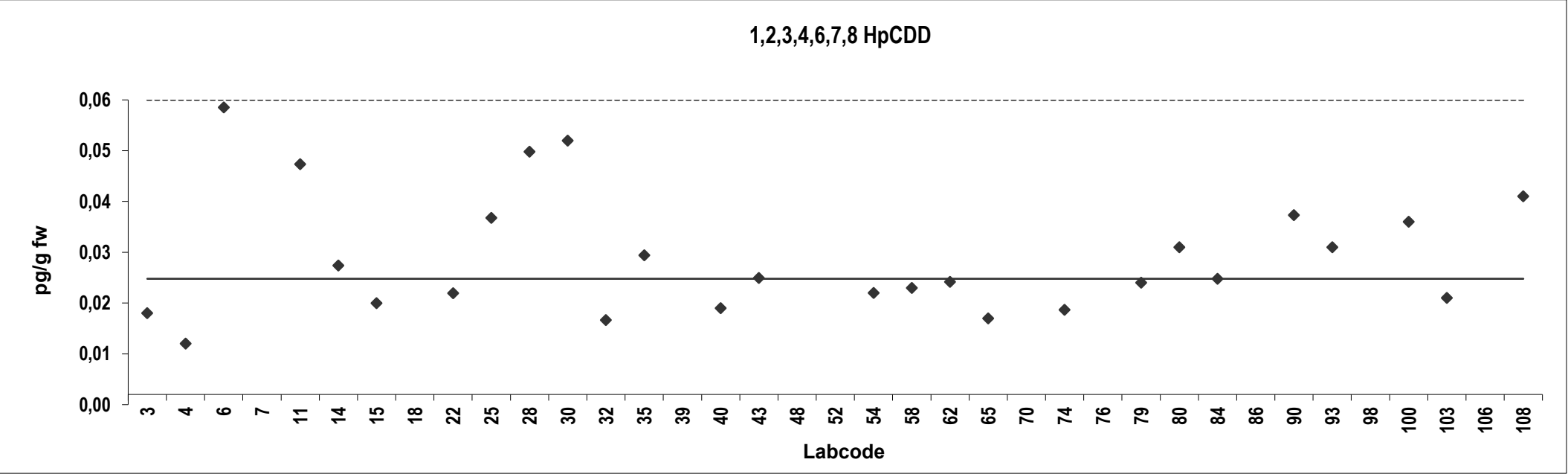
Consensus median, pg/g	0,0038
Median all values pg/g	0,0064
Consensus mean, pg/g	0,0045
Standard deviation, pg/g	0,0024
Relative standard deviation, %	54
No. of values reported	37
No. of values removed	14
No. of reported non-detects	26



Salmon
Congener: 1,2,3,4,6,7,8 HpCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,016	-1,5				
4	0,010	-2,8				
6	0,057	7,4				
7	0,13	23	Outlier,ND			
11	0,045	5,0	ND			
14	0,025	0,58				
15	0,018	-1,0				
18	0,10	17	Outlier,ND			
22	0,020	-0,62				
25	0,035	2,6				
28	0,048	5,5				
30	0,050	6,0	ND			
32	0,015	-1,8				
35	0,027	1,0	ND			
39	0,14	26	Outlier			
40	0,017	-1,3				
43	0,023	0,036				
48	0,25	50	Outlier,ND			
52	0,090	15	Outlier,ND			
54	0,020	-0,61				
58	0,021	-0,39				
62	0,022	-0,13				
65	0,015	-1,7				
70	0,13	24	Outlier,ND			
74	0,017	-1,3	ND			
76	1,6	343	Outlier			
79	0,022	-0,17				
80	0,029	1,4				
84	0,023	0,0000	ND			
86	0,099	17	Outlier,ND			
90	0,035	2,7				
93	0,029	1,4	ND			
98	0,069	10	Outlier,ND			
100	0,034	2,5				
103	0,019	-0,83				
106	0,12	21	Outlier			
108	0,039	3,6				

Consensus statistics	
Consensus median, pg/g	0,023
Median all values pg/g	0,029
Consensus mean, pg/g	0,027
Standard deviation, pg/g	0,012
Relative standard deviation, %	45
No. of values reported	37
No. of values removed	10
No. of reported non-detects	13

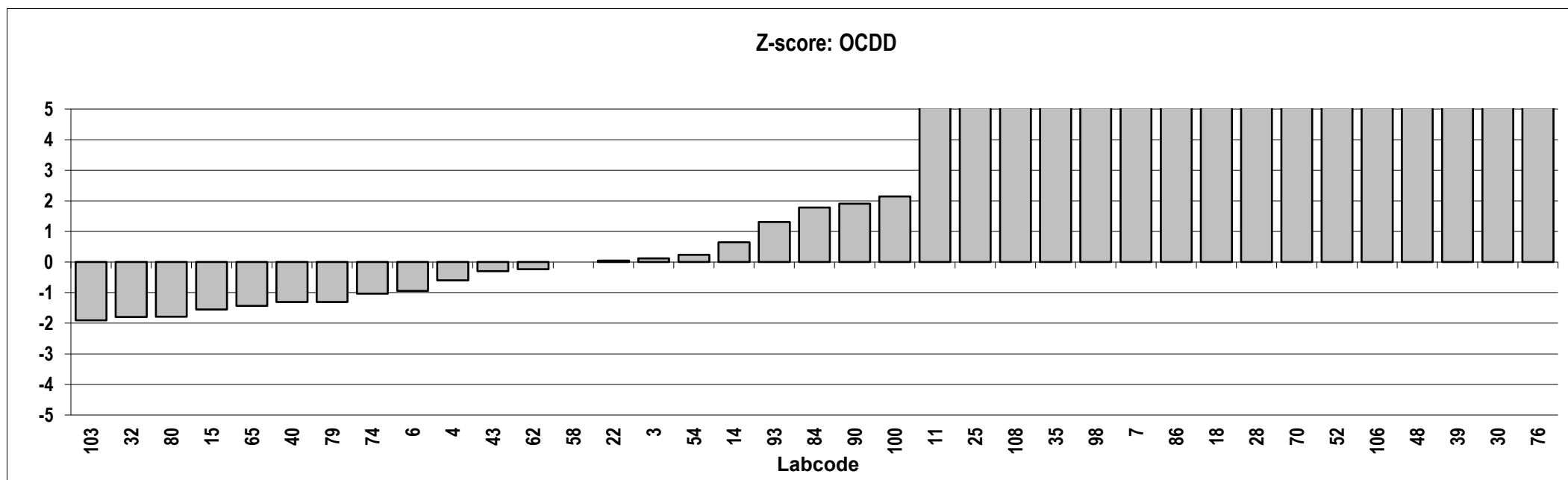
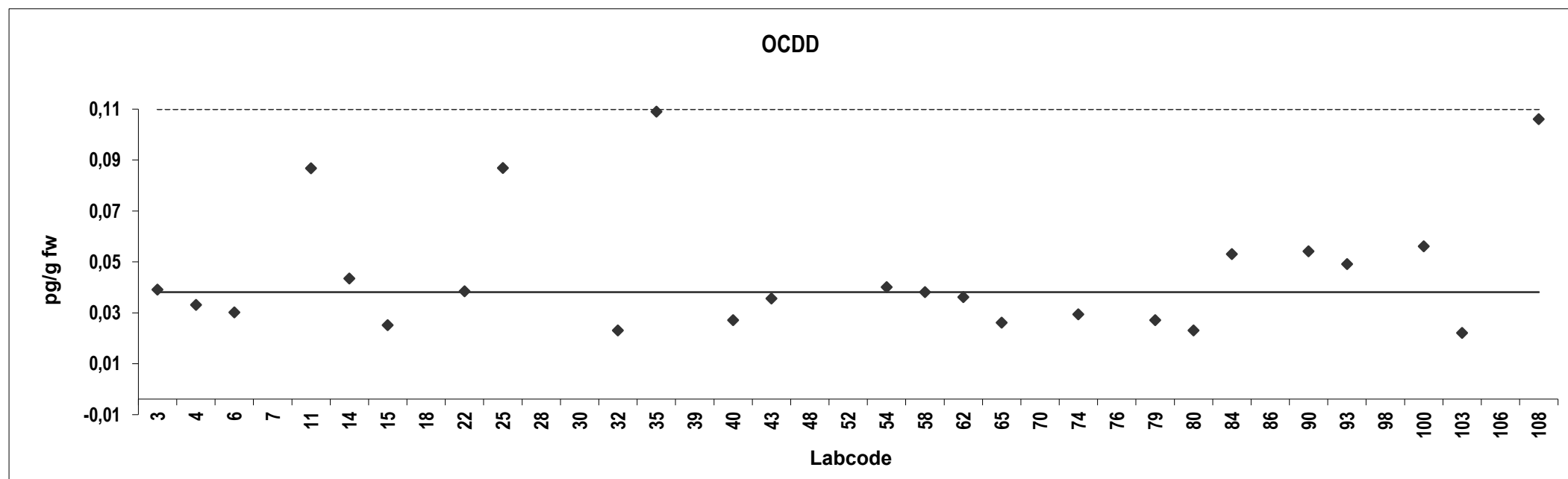


Salmon
Congener: OCDD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,043	0,12				
4	0,037	-0,60				
6	0,034	-0,94				
7	0,16	14	Outlier,ND			
11	0,091	5,8	ND			
14	0,047	0,64				
15	0,029	-1,5				
18	0,20	19	Outlier,ND			
22	0,042	0,046				
25	0,091	5,8				
28	0,26	26	Outlier			
30	2,0	233	Outlier,ND			
32	0,027	-1,8				
35	0,11	8,4	ND			
39	0,65	73	Outlier			
40	0,031	-1,3				
43	0,040	-0,30				
48	0,55	60	Outlier,ND			
52	0,35	36	Outlier,ND			
54	0,044	0,24				
58	0,042	0,0000				
62	0,040	-0,23				
65	0,030	-1,4				
70	0,31	32	Outlier,ND			
74	0,033	-1,0	ND			
76	3,6	421	Outlier			
79	0,031	-1,3				
80	0,027	-1,8				
84	0,057	1,8	ND			
86	0,20	19	Outlier,ND			
90	0,058	1,9				
93	0,053	1,3	ND			
98	0,13	10	Outlier,ND			
100	0,060	2,1				
103	0,026	-1,9				
106	0,54	59	Outlier			
108	0,11	8,1				

Consensus statistics

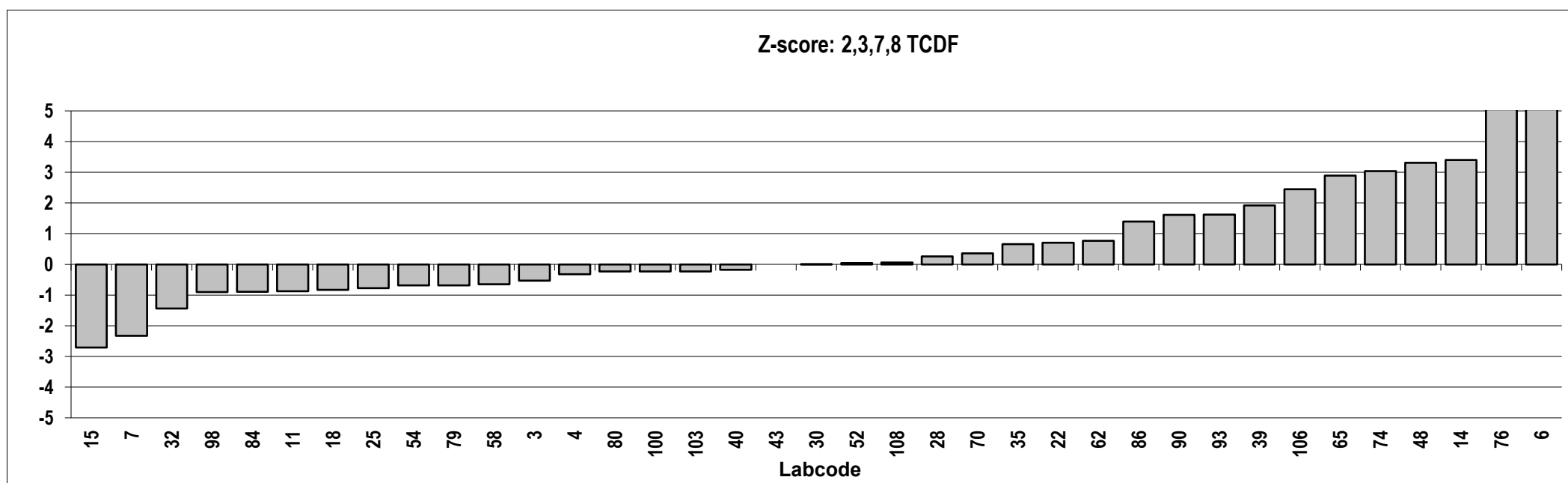
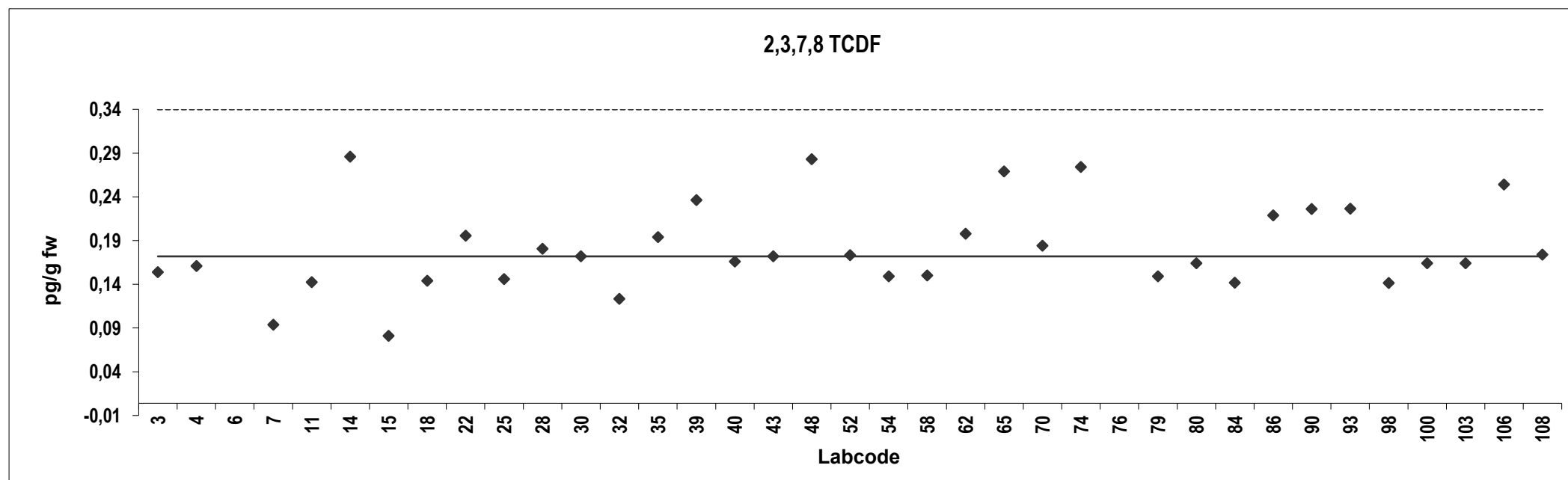
Consensus median, pg/g	0,042
Median all values pg/g	0,057
Consensus mean, pg/g	0,049
Standard deviation, pg/g	0,025
Relative standard deviation, %	51
No. of values reported	37
No. of values removed	12
No. of reported non-detects	13



Salmon
Congener: 2,3,7,8 TCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,15	-0,53				
4	0,16	-0,33				
6	0,37	6,0	Outlier			
7	0,090	-2,3	ND			
11	0,14	-0,87				
14	0,28	3,4				
15	0,077	-2,7				
18	0,14	-0,83				
22	0,19	0,71				
25	0,14	-0,78				
28	0,18	0,26				
30	0,17	0,0025				
32	0,12	-1,4				
35	0,19	0,66				
39	0,23	1,9				
40	0,16	-0,18				
43	0,17	0,0000				
48	0,28	3,3				
52	0,17	0,038				
54	0,15	-0,68				
58	0,15	-0,65				
62	0,19	0,77				
65	0,27	2,9				
70	0,18	0,36				
74	0,27	3,0				
76	0,36	5,7	Outlier			
79	0,15	-0,68				
80	0,16	-0,24				
84	0,14	-0,89				
86	0,21	1,4				
90	0,22	1,6				
93	0,22	1,6				
98	0,14	-0,90	ND			
100	0,16	-0,24				
103	0,16	-0,24				
106	0,25	2,4				
108	0,17	0,062				

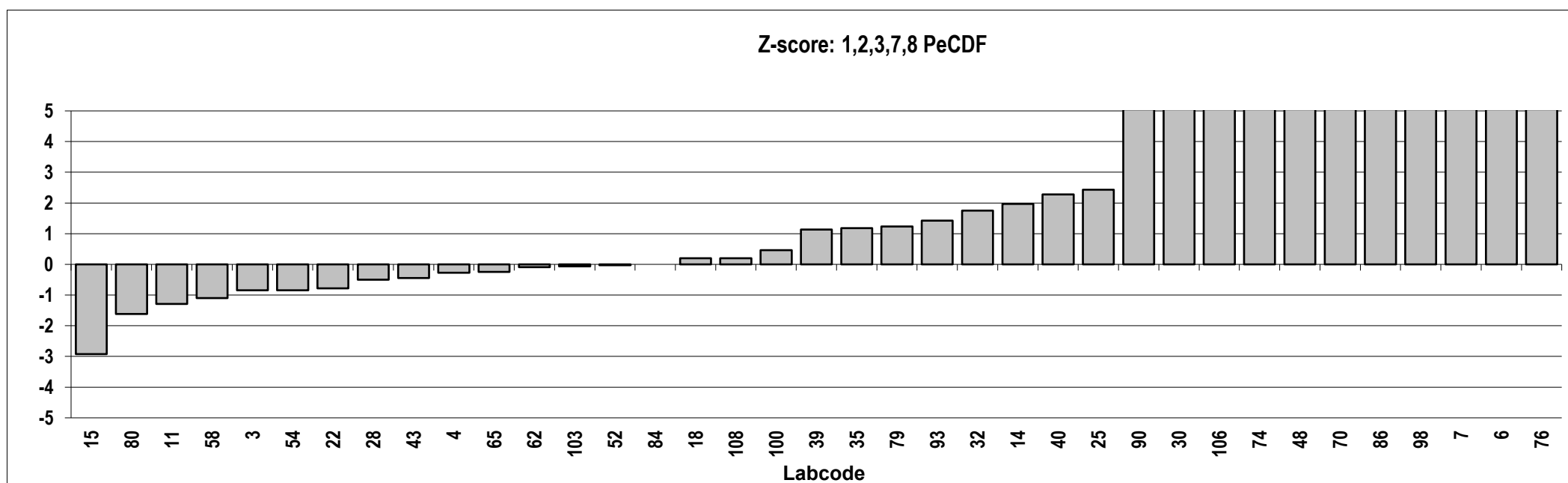
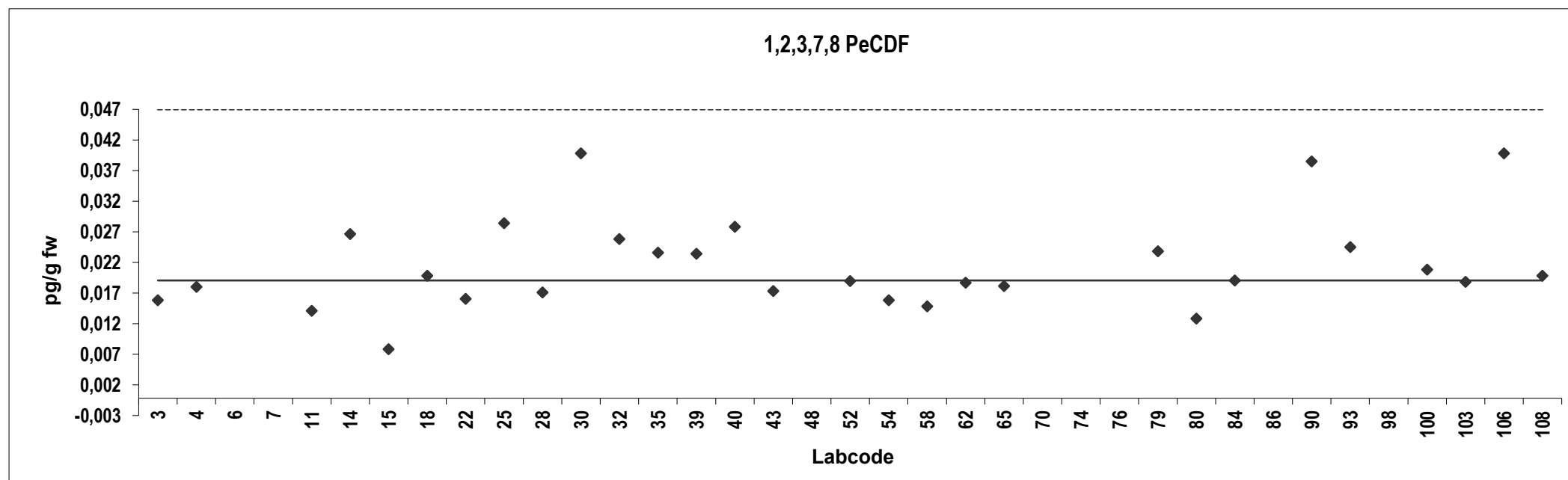
Consensus statistics	
Consensus median, pg/g	0,17
Median all values pg/g	0,17
Consensus mean, pg/g	0,18
Standard deviation, pg/g	0,050
Relative standard deviation, %	28
No. of values reported	37
No. of values removed	2
No. of reported non-detects	2



Salmon
Congener: 1,2,3,7,8 PeCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,016	-0,84				
4	0,018	-0,27				
6	0,16	36	Outlier			
7	0,15	34	Outlier,ND			
11	0,014	-1,3				
14	0,027	2,0				
15	0,0080	-2,9				
18	0,020	0,20	ND			
22	0,016	-0,78				
25	0,029	2,4				
28	0,017	-0,50				
30	0,040	5,4	ND			
32	0,026	1,8				
35	0,024	1,2				
39	0,024	1,1				
40	0,028	2,3				
43	0,018	-0,44				
48	0,054	9,0	Outlier,ND			
52	0,019	-0,028				
54	0,016	-0,84				
58	0,015	-1,1				
62	0,019	-0,094				
65	0,018	-0,24				
70	0,063	11	Outlier,ND			
74	0,050	8	Outlier			
76	1,3	328	Outlier			
79	0,024	1,2				
80	0,013	-1,6				
84	0,019	0,0000				
86	0,074	14	Outlier,ND			
90	0,039	5,1				
93	0,025	1,4				
98	0,088	18	Outlier,ND			
100	0,021	0,46				
103	0,019	-0,061				
106	0,040	5,4				
108	0,020	0,20				

Consensus statistics	
Consensus median, pg/g	0,019
Median all values pg/g	0,024
Consensus mean, pg/g	0,022
Standard deviation, pg/g	0,0077
Relative standard deviation, %	35
No. of values reported	37
No. of values removed	8
No. of reported non-detects	7

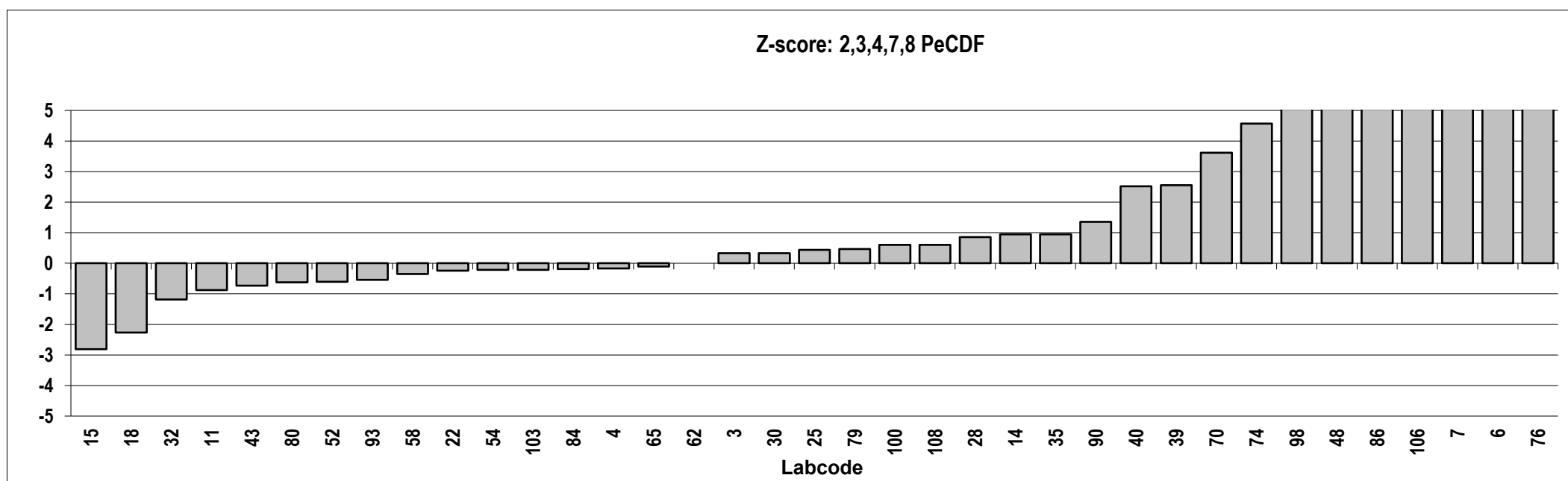
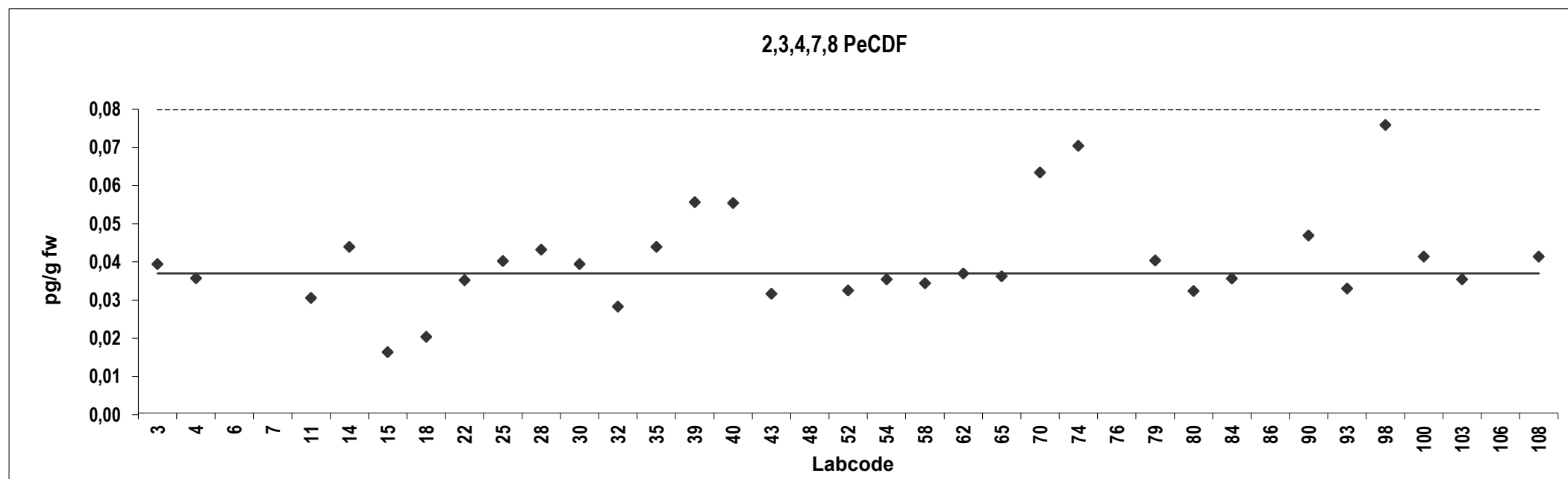


Salmon
Congener: 2,3,4,7,8 PeCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,039	0,33				
4	0,035	-0,17				
6	0,19	22	Outlier			
7	0,13	12	Outlier,ND			
11	0,030	-0,88				
14	0,044	0,95				
15	0,016	-2,8				
18	0,020	-2,3	ND			
22	0,035	-0,24				
25	0,040	0,44				
28	0,043	0,86				
30	0,039	0,33				
32	0,028	-1,2				
35	0,044	0,95				
39	0,055	2,6				
40	0,055	2,5				
43	0,031	-0,73				
48	0,088	7,0	Outlier,ND			
52	0,032	-0,61				
54	0,035	-0,21				
58	0,034	-0,35				
62	0,037	0,0000				
65	0,036	-0,11				
70	0,063	3,6	ND			
74	0,070	4,6				
76	1,3	178	Outlier			
79	0,040	0,47				
80	0,032	-0,62				
84	0,035	-0,19				
86	0,099	8,5	Outlier,ND			
90	0,047	1,4				
93	0,033	-0,54				
98	0,075	5,3	ND			
100	0,041	0,61				
103	0,035	-0,21				
106	0,11	10	Outlier			
108	0,041	0,61				

Consensus statistics

Consensus median, pg/g	0,037
Median all values pg/g	0,040
Consensus mean, pg/g	0,040
Standard deviation, pg/g	0,013
Relative standard deviation, %	32
No. of values reported	37
No. of values removed	6
No. of reported non-detects	6

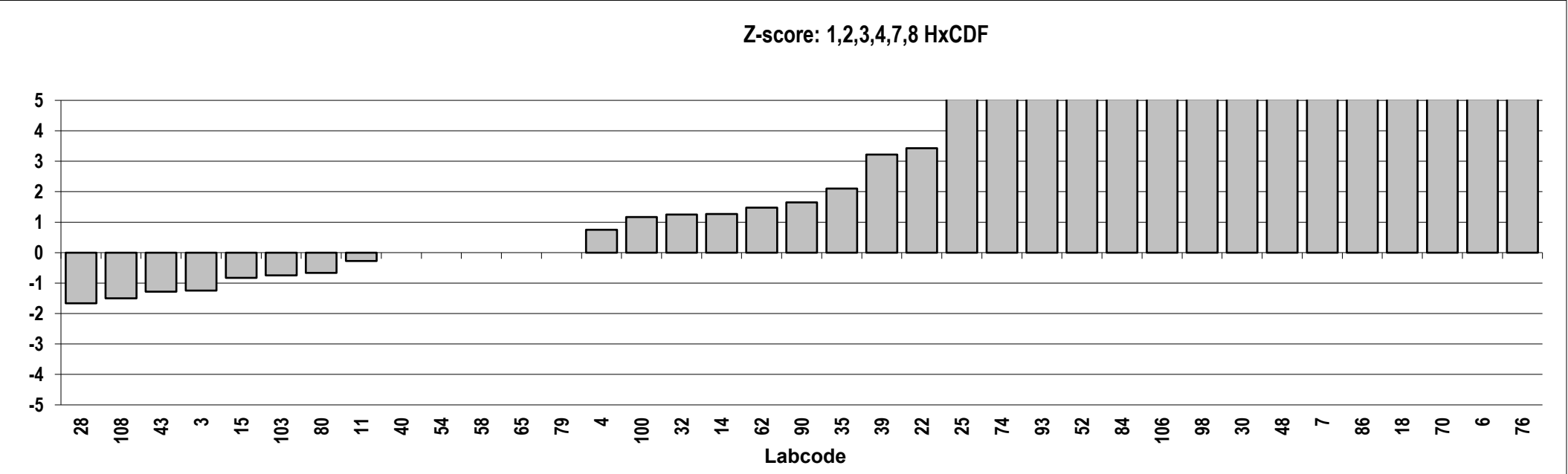
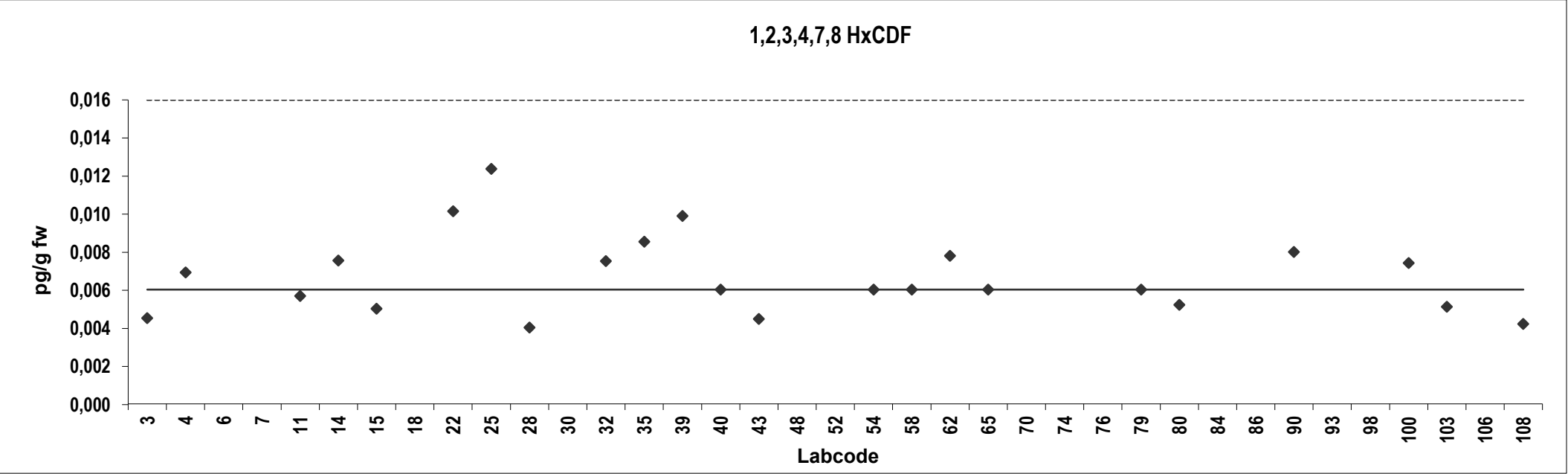


Salmon
Congener: 1,2,3,4,7,8 HxCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0045	-1,3				
4	0,0069	0,75				
6	0,24	195	Outlier			
7	0,089	69	Outlier,ND			
11	0,0057	-0,28	ND			
14	0,0075	1,3				
15	0,0050	-0,83				
18	0,10	78	Outlier,ND			
22	0,010	3,4				
25	0,012	5,3				
28	0,0040	-1,7	ND			
30	0,050	37	Outlier,ND			
32	0,0075	1,3				
35	0,0085	2,1	ND			
39	0,0099	3,2				
40	0,0060	0,0000	ND			
43	0,0045	-1,3				
48	0,070	53	Outlier,ND			
52	0,022	13	Outlier,ND			
54	0,0060	0,0000				
58	0,0060	0,0000				
62	0,0078	1,5				
65	0,0060	0,0000	ND			
70	0,13	103	Outlier,ND			
74	0,017	8,9	Outlier,ND			
76	1,6	1299	Outlier			
79	0,0060	0,0000				
80	0,0052	-0,67				
84	0,023	14	Outlier,ND			
86	0,099	78	Outlier,ND			
90	0,0080	1,7	ND			
93	0,021	13	Outlier,ND			
98	0,040	28	Outlier,ND			
100	0,0074	1,2				
103	0,0051	-0,75				
106	0,030	20	Outlier			
108	0,0042	-1,5				

Consensus statistics

Consensus median, pg/g	0,0060
Median all values pg/g	0,0080
Consensus mean, pg/g	0,0067
Standard deviation, pg/g	0,0021
Relative standard deviation, %	31
No. of values reported	37
No. of values removed	14
No. of reported non-detects	17

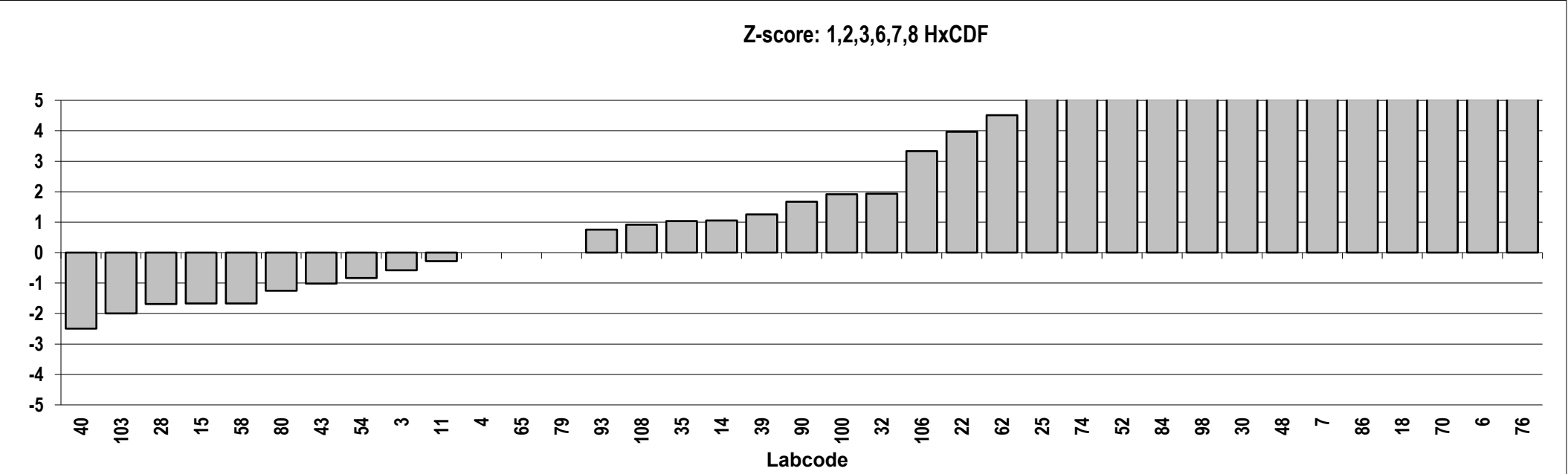
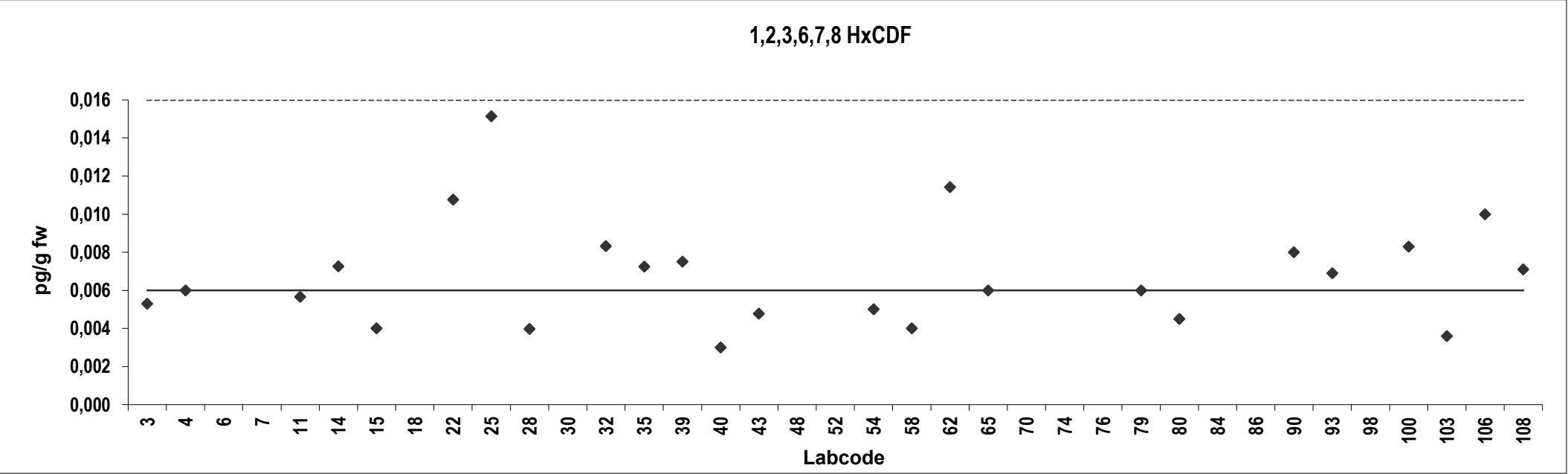


Salmon
Congener: 1,2,3,6,7,8 HxCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0053	-0,58				
4	0,0060	0,0000				
6	0,17	140	Outlier			
7	0,097	76	Outlier,ND			
11	0,0057	-0,28	ND			
14	0,0073	1,1				
15	0,0040	-1,7				
18	0,10	78	Outlier,ND			
22	0,011	4,0				
25	0,015	7,6				
28	0,0040	-1,7	ND			
30	0,050	37	Outlier,ND			
32	0,0083	1,9				
35	0,0072	1,0				
39	0,0075	1,3				
40	0,0030	-2,5	ND			
43	0,0048	-1,0				
48	0,068	52	Outlier,ND			
52	0,022	13	Outlier,ND			
54	0,0050	-0,83				
58	0,0040	-1,7				
62	0,011	4,5				
65	0,0060	0,0000	ND			
70	0,13	103	Outlier,ND			
74	0,017	8,9	Outlier,ND			
76	1,5	1232	Outlier			
79	0,0060	0,0000				
80	0,0045	-1,3				
84	0,023	14	Outlier,ND			
86	0,099	78	Outlier,ND			
90	0,0080	1,7	ND			
93	0,0069	0,75				
98	0,039	28	Outlier,ND			
100	0,0083	1,9				
103	0,0036	-2,0				
106	0,010	3,3	ND			
108	0,0071	0,92				

Consensus statistics

Consensus median, pg/g	0,0060
Median all values pg/g	0,0080
Consensus mean, pg/g	0,0068
Standard deviation, pg/g	0,0028
Relative standard deviation, %	41
No. of values reported	37
No. of values removed	12
No. of reported non-detects	16

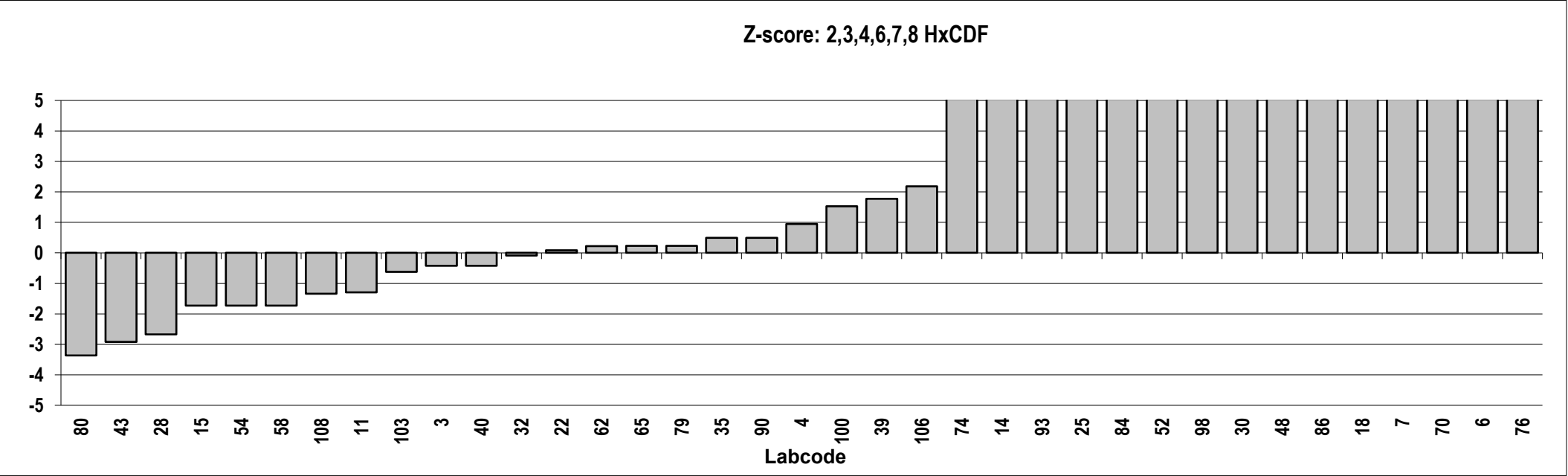
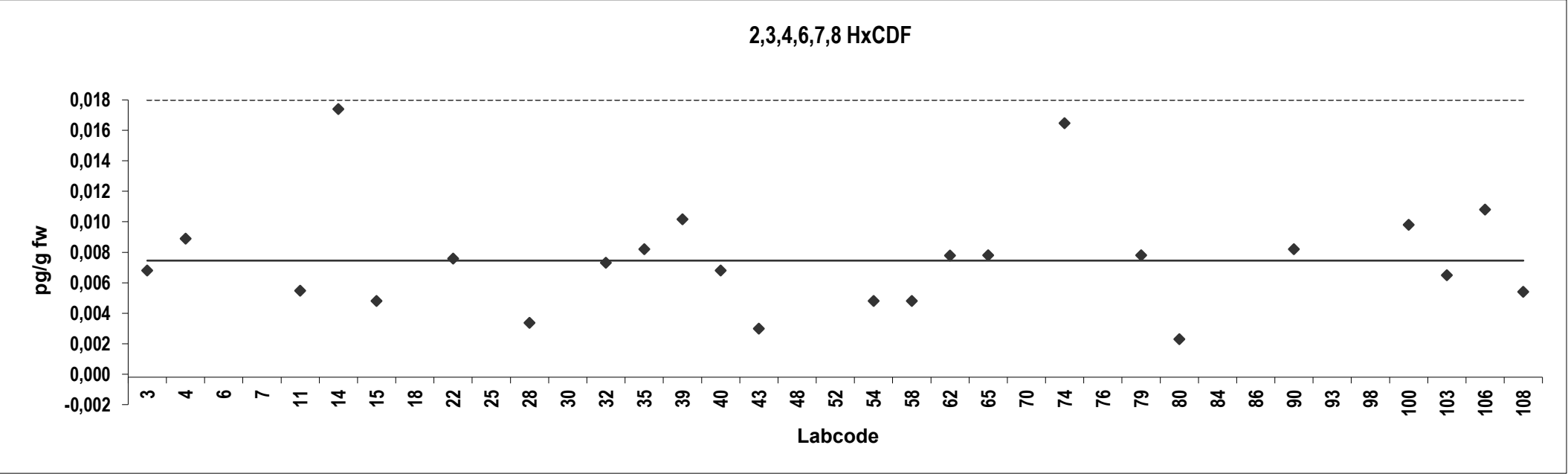


Salmon
Congener: 2,3,4,6,7,8 HxCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0070	-0,43				
4	0,0091	0,95				
6	0,18	110	Outlier			
7	0,11	66	Outlier,ND			
11	0,0057	-1,3	ND			
14	0,018	6,5				
15	0,0050	-1,7				
18	0,10	60	Outlier,ND			
22	0,0078	0,087				
25	0,022	9,6	Outlier			
28	0,0036	-2,7	ND			
30	0,050	28	Outlier,ND			
32	0,0075	-0,087				
35	0,0084	0,49				
39	0,010	1,8				
40	0,0070	-0,43				
43	0,0032	-2,9				
48	0,073	43	Outlier,ND			
52	0,024	11	Outlier,ND			
54	0,0050	-1,7				
58	0,0050	-1,7				
62	0,0080	0,22				
65	0,0080	0,23	ND			
70	0,13	80	Outlier,ND			
74	0,017	5,9	ND			
76	1,6	1073	Outlier			
79	0,0080	0,23				
80	0,0025	-3,4				
84	0,023	9,9	Outlier,ND			
86	0,099	60	Outlier,ND			
90	0,0084	0,49	ND			
93	0,021	8,7	Outlier,ND			
98	0,038	20	Outlier,ND			
100	0,010	1,5				
103	0,0067	-0,62				
106	0,011	2,2	ND			
108	0,0056	-1,3				

Consensus statistics

Consensus median, pg/g	0,0077
Median all values pg/g	0,0091
Consensus mean, pg/g	0,0078
Standard deviation, pg/g	0,0036
Relative standard deviation, %	46
No. of values reported	37
No. of values removed	13
No. of reported non-detects	16

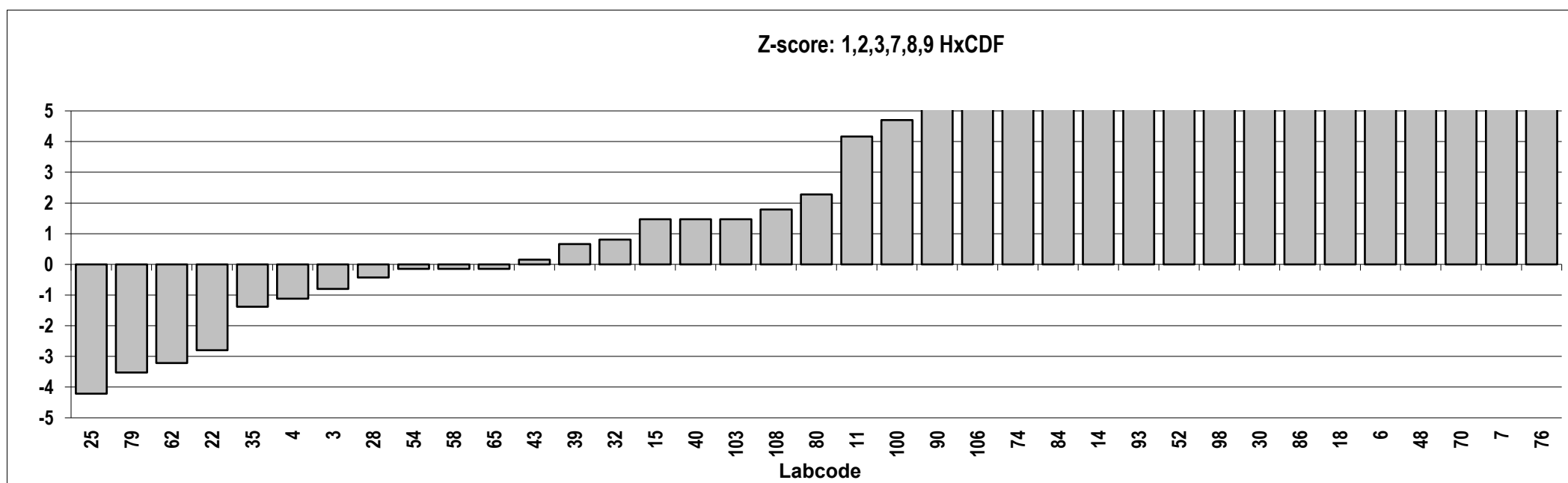
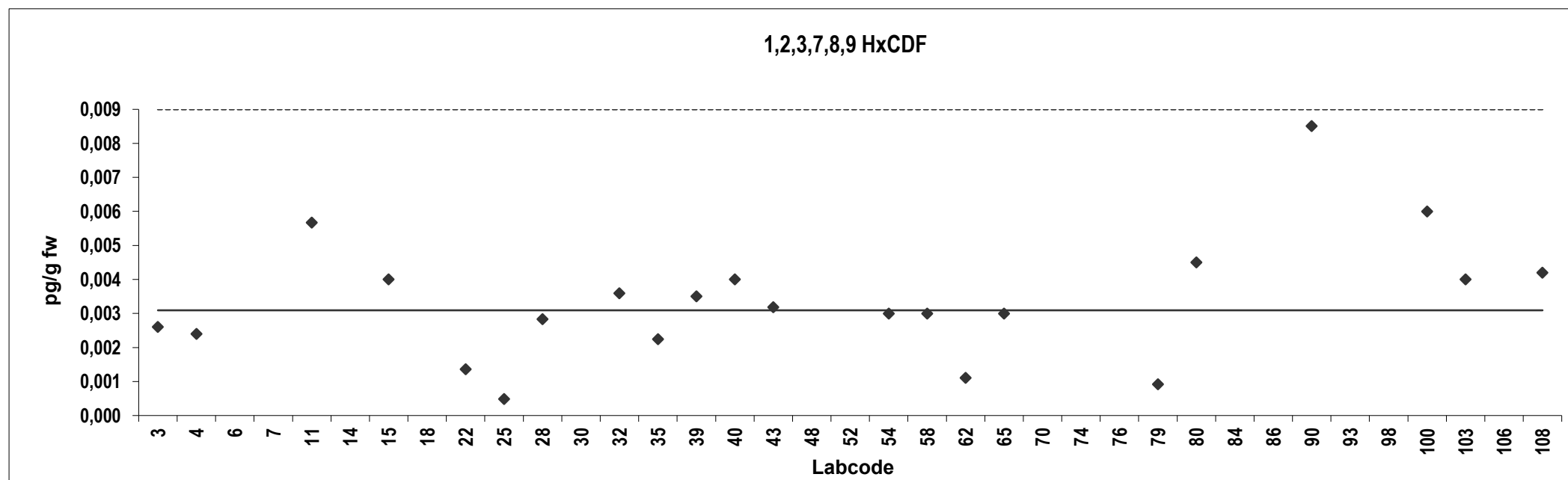


Salmon
Congener: 1,2,3,7,8,9 HxCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0026	-0,80				
4	0,0024	-1,1	ND			
6	0,11	175	Outlier			
7	0,18	284	Outlier,ND			
11	0,0057	4,2	ND			
14	0,024	34	Outlier			
15	0,0040	1,5	ND			
18	0,10	157	Outlier,ND			
22	0,0014	-2,8	ND			
25	0,00049	-4,2				
28	0,0028	-0,42	ND			
30	0,050	76	Outlier,ND			
32	0,0036	0,80				
35	0,0022	-1,4	ND			
39	0,0035	0,66	ND			
40	0,0040	1,5	ND			
43	0,0032	0,15				
48	0,13	205	Outlier,ND			
52	0,032	47	Outlier,ND			
54	0,0030	-0,15	ND			
58	0,0030	-0,15	ND			
62	0,0011	-3,2	ND			
65	0,0030	-0,15	ND			
70	0,13	205	Outlier,ND			
74	0,017	22	Outlier,ND			
76	1,6	2658	Outlier			
79	0,00092	-3,5				
80	0,0045	2,3				
84	0,023	32	Outlier,ND			
86	0,099	155	Outlier,ND			
90	0,0085	8,8	ND			
93	0,028	40	Outlier,ND			
98	0,045	68	Outlier,ND			
100	0,0060	4,7				
103	0,0040	1,5	ND			
106	0,013	16	Outlier,ND			
108	0,0042	1,8	ND			

Consensus statistics

Consensus median, pg/g	0,0031
Median all values pg/g	0,0045
Consensus mean, pg/g	0,0034
Standard deviation, pg/g	0,0018
Relative standard deviation, %	54
No. of values reported	37
No. of values removed	15
No. of reported non-detects	27

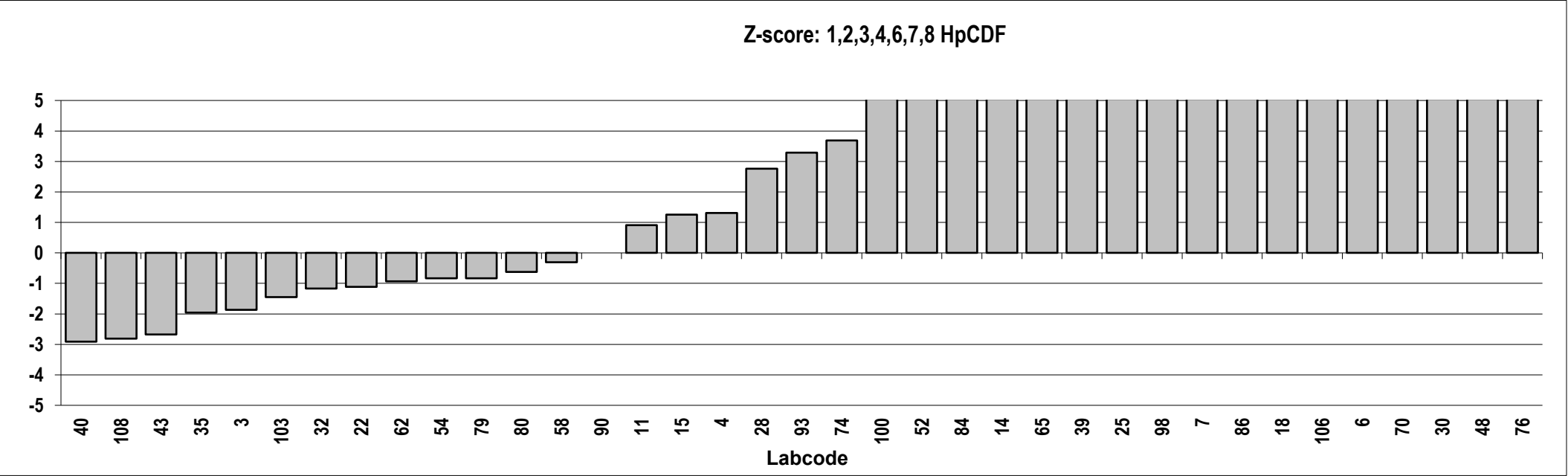
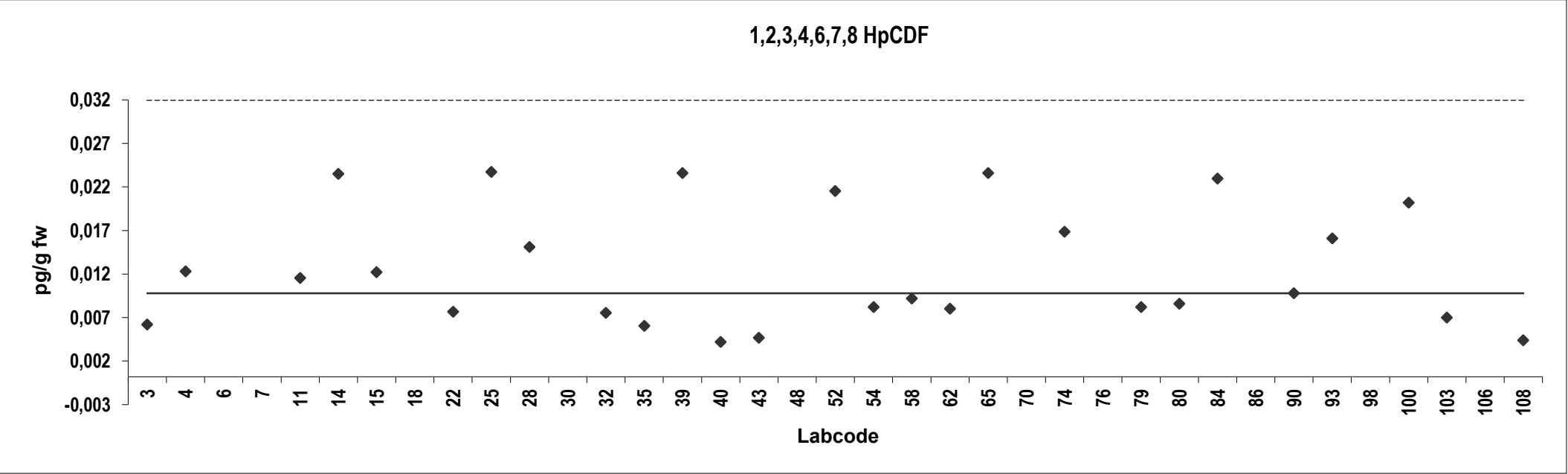


Salmon
Congener: 1,2,3,4,6,7,8 HpCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0060	-1,9				
4	0,012	1,3				
6	0,11	54	Outlier			
7	0,083	38	Outlier,ND			
11	0,011	0,91	ND			
14	0,023	7,1				
15	0,012	1,3				
18	0,10	47	Outlier,ND			
22	0,0075	-1,1				
25	0,024	7,3				
28	0,015	2,8				
30	0,15	73	Outlier,ND			
32	0,0074	-1,2				
35	0,0058	-2,0	ND			
39	0,023	7,2				
40	0,0040	-2,9				
43	0,0045	-2,7				
48	0,15	73	Outlier,ND			
52	0,021	6,1	ND			
54	0,0080	-0,83				
58	0,0090	-0,31				
62	0,0078	-0,93				
65	0,023	7,2				
70	0,13	63	Outlier,ND			
74	0,017	3,7	ND			
76	1,7	901	Outlier			
79	0,0080	-0,83				
80	0,0084	-0,62				
84	0,023	6,9	ND			
86	0,099	47	Outlier,ND			
90	0,0096	0,0000	ND			
93	0,016	3,3				
98	0,037	14	Outlier,ND			
100	0,020	5,4				
103	0,0068	-1,5				
106	0,11	52	Outlier			
108	0,0042	-2,8	ND			

Consensus statistics

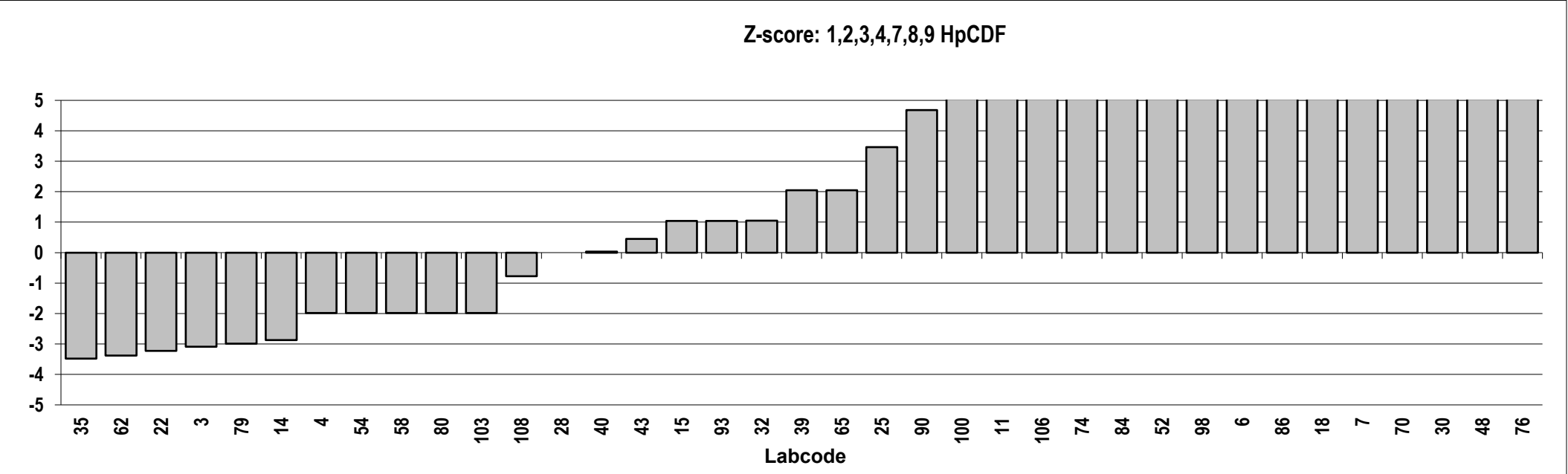
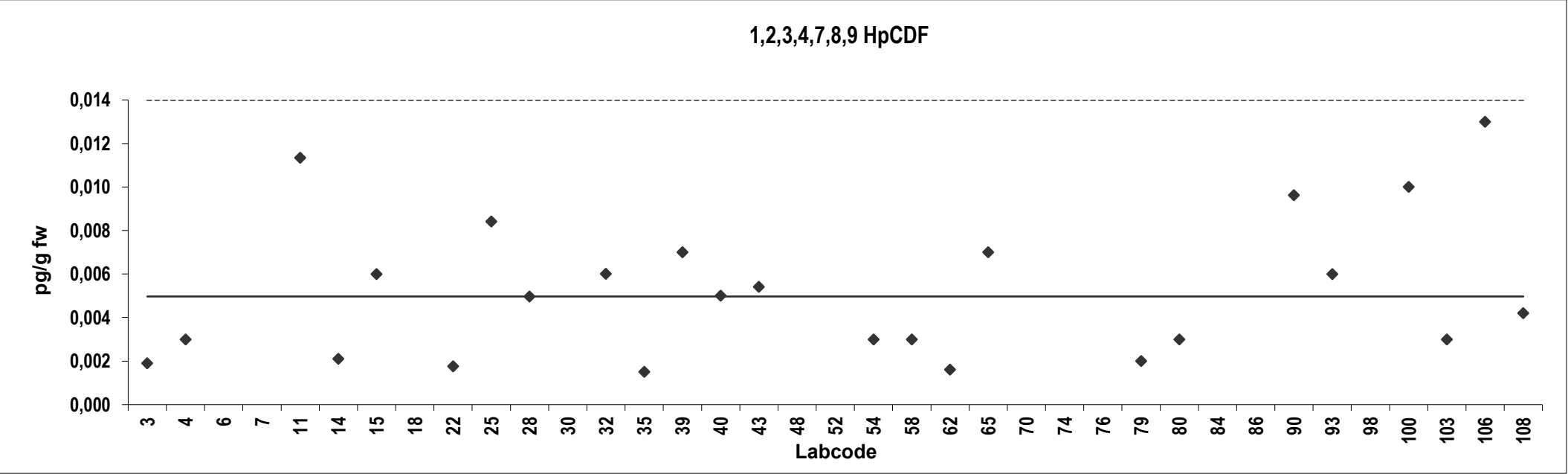
Consensus median, pg/g	0,010
Median all values pg/g	0,016
Consensus mean, pg/g	0,013
Standard deviation, pg/g	0,0069
Relative standard deviation, %	55
No. of values reported	37
No. of values removed	10
No. of reported non-detects	14



Salmon
Congener: 1,2,3,4,7,8,9 HpCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0019	-3,1				
4	0,0030	-2,0				
6	0,061	56	Outlier			
7	0,11	108	Outlier,ND			
11	0,011	6,4	ND			
14	0,0021	-2,9	ND			
15	0,0060	1,0				
18	0,10	96	Outlier,ND			
22	0,0018	-3,2	ND			
25	0,0084	3,5				
28	0,0050	0,0000	ND			
30	0,15	146	Outlier,ND			
32	0,0060	1,1				
35	0,0015	-3,5	ND			
39	0,0070	2,0	ND			
40	0,0050	0,032	ND			
43	0,0054	0,45				
48	0,18	176	Outlier,ND			
52	0,039	34	Outlier,ND			
54	0,0030	-2,0	ND			
58	0,0030	-2,0	ND			
62	0,0016	-3,4	ND			
65	0,0070	2,0	ND			
70	0,13	126	Outlier,ND			
74	0,017	12	Outlier,ND			
76	1,6	1628	Outlier			
79	0,0020	-3,0				
80	0,0030	-2,0				
84	0,023	18	Outlier,ND			
86	0,099	95	Outlier,ND			
90	0,0096	4,7	ND			
93	0,0060	1,0	ND			
98	0,045	40	Outlier,ND			
100	0,010	5,1	ND			
103	0,0030	-2,0	ND			
106	0,013	8,1	ND			
108	0,0042	-0,77	ND			

Consensus statistics	
Consensus median, pg/g	0,0050
Median all values pg/g	0,0070
Consensus mean, pg/g	0,0052
Standard deviation, pg/g	0,0033
Relative standard deviation, %	63
No. of values reported	37
No. of values removed	12
No. of reported non-detects	27

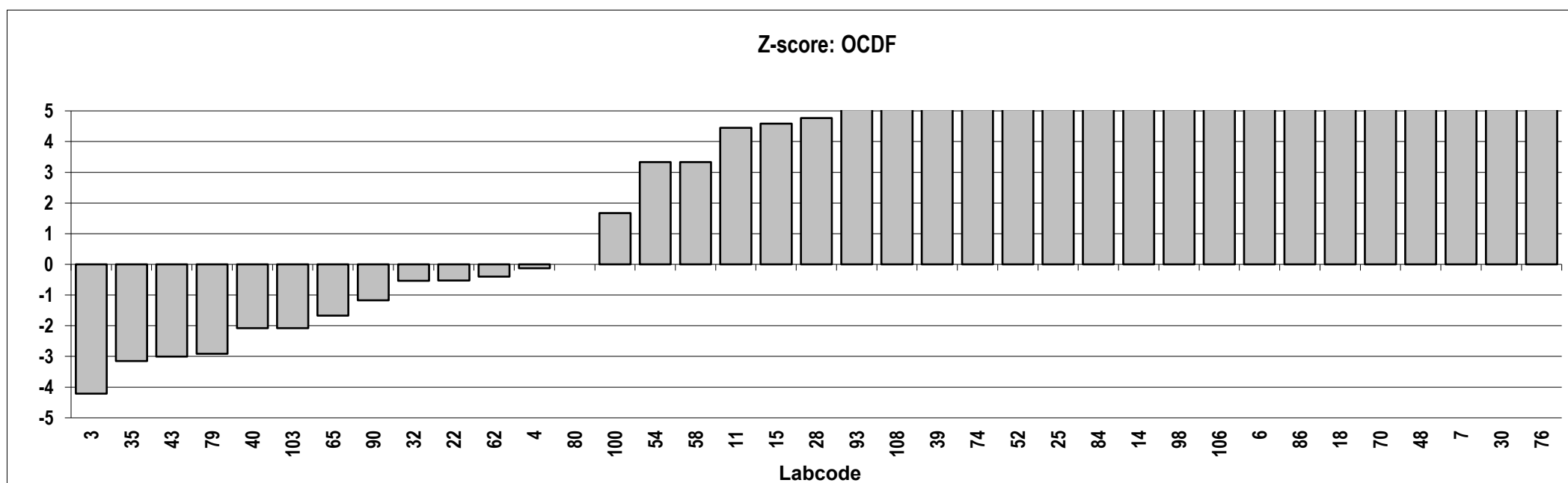
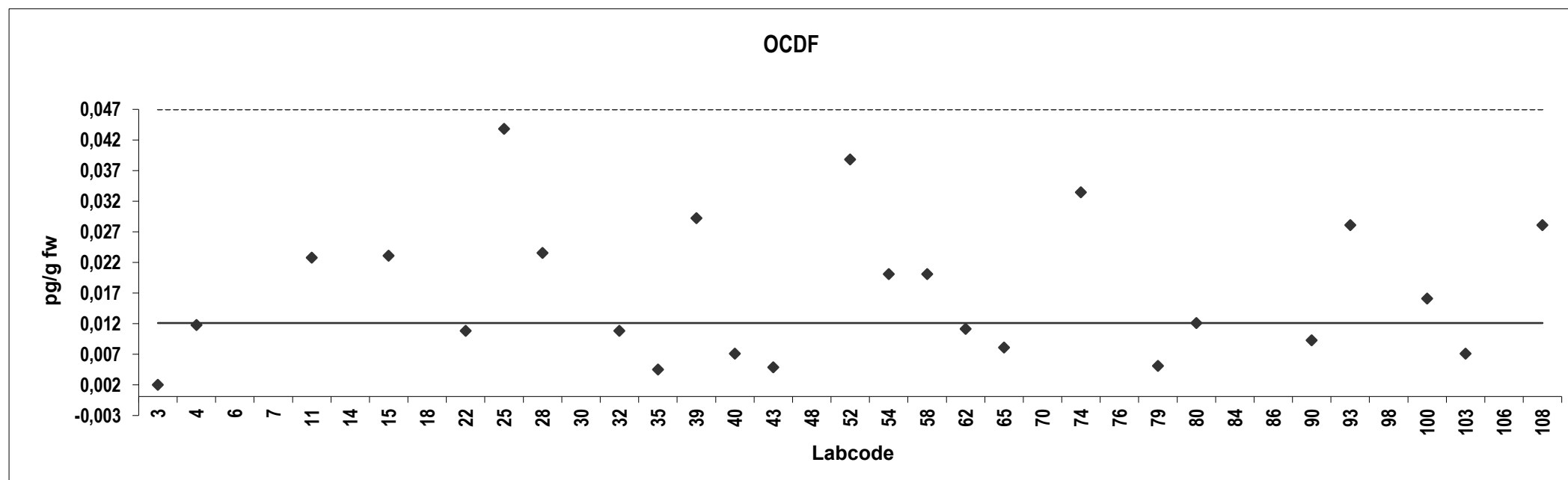


Salmon
Congener: OCDF

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,0019	-4,2				
4	0,012	-0,13				
6	0,16	61	Outlier			
7	0,89	368	Outlier,ND			
11	0,023	4,4	ND			
14	0,083	29	Outlier			
15	0,023	4,6				
18	0,20	78	Outlier,ND			
22	0,011	-0,53	ND			
25	0,044	13				
28	0,023	4,8				
30	2,0	828	Outlier,ND			
32	0,011	-0,54				
35	0,0044	-3,2	ND			
39	0,029	7,1				
40	0,0070	-2,1	ND			
43	0,0048	-3,0				
48	0,34	137	Outlier,ND			
52	0,039	11	ND			
54	0,020	3,3	ND			
58	0,020	3,3	ND			
62	0,011	-0,40	ND			
65	0,0080	-1,7	ND			
70	0,31	124	Outlier,ND			
74	0,033	8,9	ND			
76	3,8	1599	Outlier			
79	0,0050	-2,9				
80	0,012	0,0000	ND			
84	0,057	19	Outlier,ND			
86	0,20	78	Outlier,ND			
90	0,0092	-1,2				
93	0,028	6,7	ND			
98	0,12	44	Outlier,ND			
100	0,016	1,7				
103	0,0070	-2,1	ND			
106	0,13	49	Outlier			
108	0,028	6,7	ND			

Consensus statistics

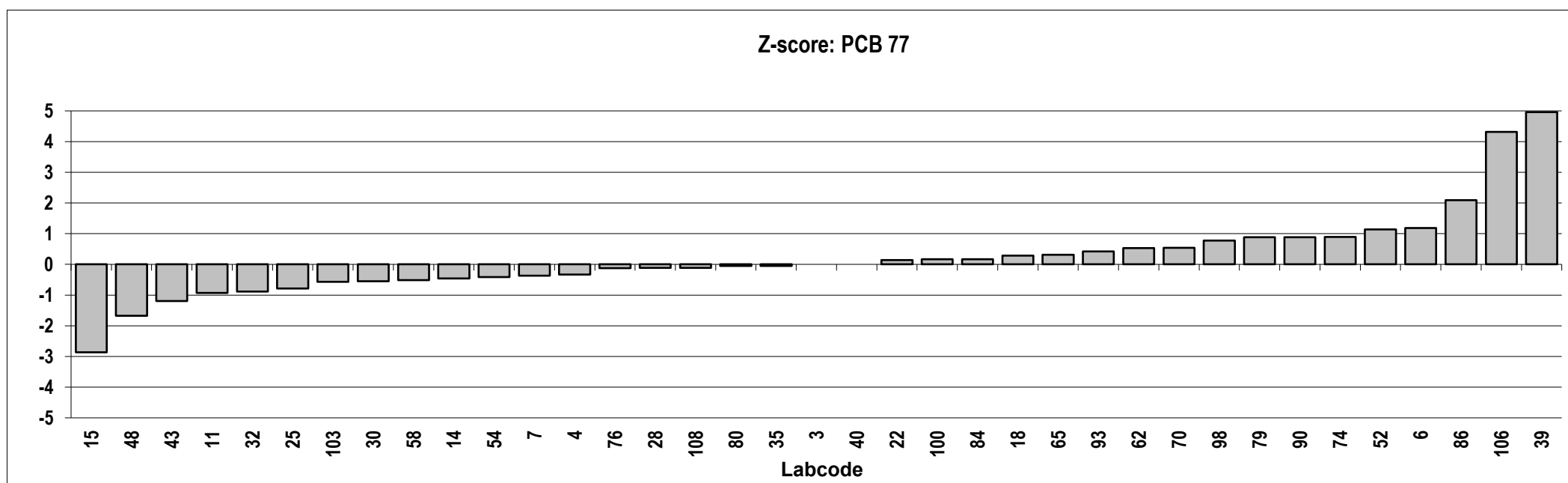
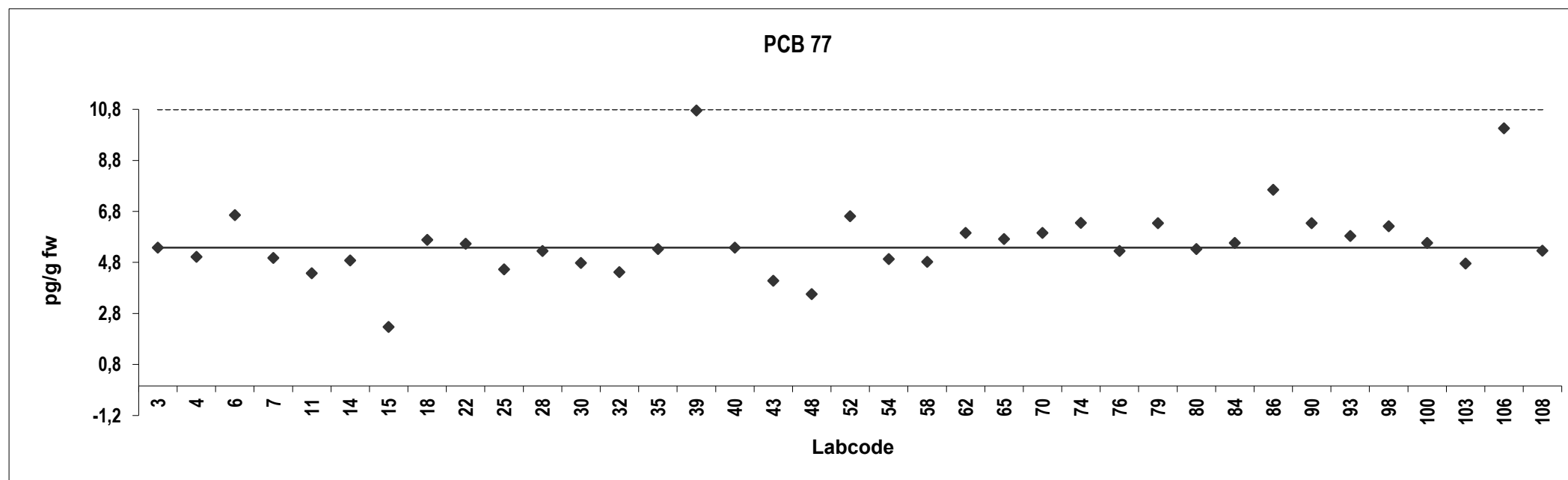
Consensus median, pg/g	0,012
Median all values pg/g	0,023
Consensus mean, pg/g	0,017
Standard deviation, pg/g	0,011
Relative standard deviation, %	67
No. of values reported	37
No. of values removed	12
No. of reported non-detects	22



Salmon
Congener: PCB 77

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	5,4	0,0000				
4	5,1	-0,33				
6	6,7	1,2				
7	5,0	-0,37				
11	4,4	-0,93				
14	4,9	-0,46				
15	2,3	-2,9				
18	5,7	0,29				
22	5,6	0,14				
25	4,6	-0,79				
28	5,3	-0,12				
30	4,8	-0,55				
32	4,5	-0,89				
35	5,4	-0,049				
39	11	5,0				
40	5,4	0,0000				
43	4,1	-1,2				
48	3,6	-1,7				
52	6,7	1,1				
54	5,0	-0,42				
58	4,9	-0,52				
62	6,0	0,53				
65	5,8	0,32				
70	6,0	0,54				
74	6,4	0,89				
76	5,3	-0,12				
79	6,4	0,88				
80	5,4	-0,055				
84	5,6	0,17				
86	7,7	2,1				
90	6,4	0,89				
93	5,9	0,42				
98	6,3	0,78				
100	5,6	0,17				
103	4,8	-0,57				
106	10	4,3				
108	5,3	-0,11				

Consensus statistics	
Consensus median, pg/g	5,4
Median all values pg/g	5,4
Consensus mean, pg/g	5,6
Standard deviation, pg/g	1,5
Relative standard deviation, %	27
No. of values reported	37
No. of values removed	0
No. of reported non-detects	0

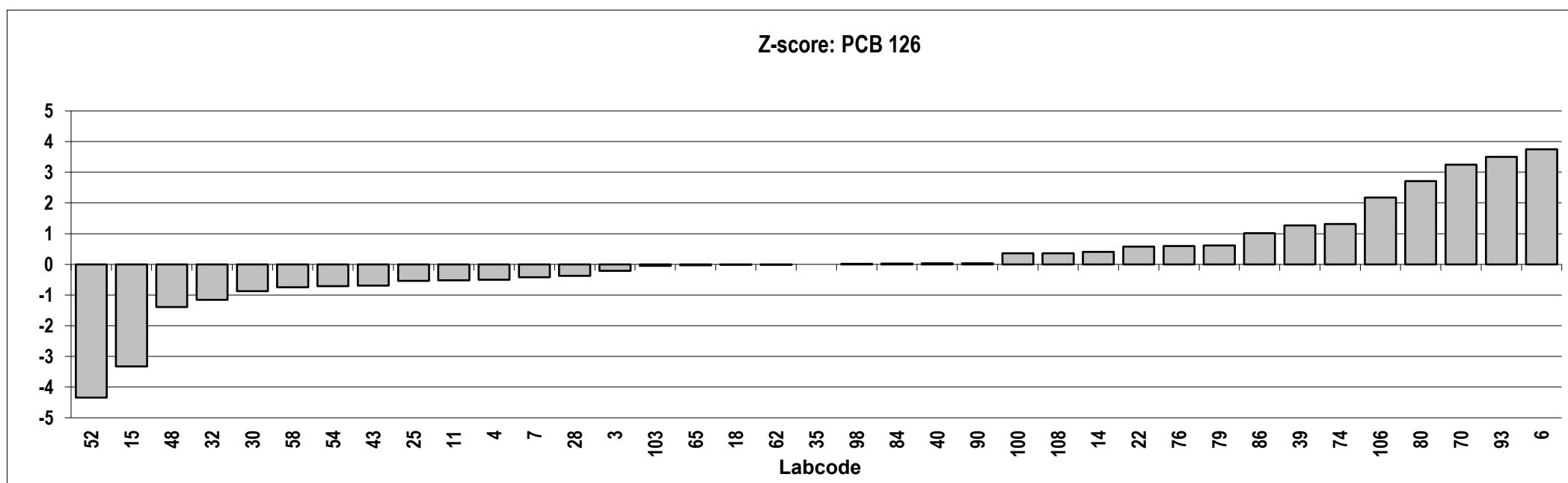
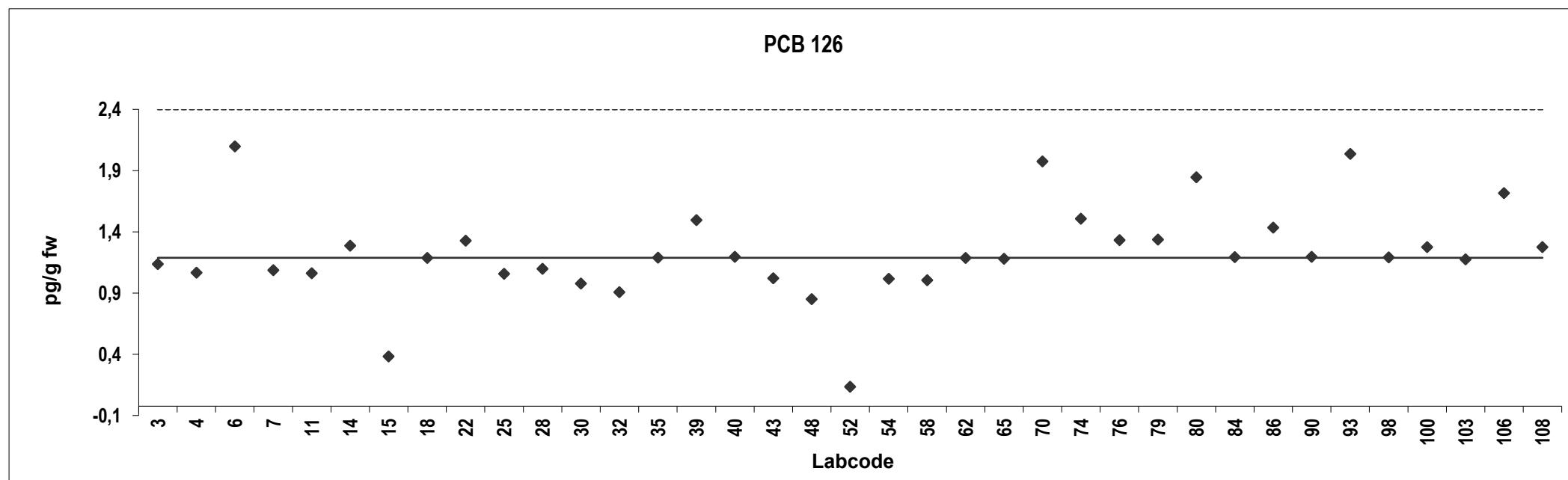


Salmon
Congener: PCB 126

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	1,2	-0,21				
4	1,1	-0,50				
6	2,1	3,7				
7	1,1	-0,42				
11	1,1	-0,52				
14	1,3	0,40				
15	0,41	-3,3				
18	1,2	-0,0083				
22	1,4	0,58				
25	1,1	-0,54				
28	1,1	-0,37				
30	1,0	-0,87				
32	0,93	-1,2				
35	1,2	0,0000				
39	1,5	1,3				
40	1,2	0,033				
43	1,0	-0,69				
48	0,88	-1,4				
52	0,16	-4,3	ND			
54	1,0	-0,71				
58	1,0	-0,75				
62	1,2	-0,0083				
65	1,2	-0,033				
70	2,0	3,3	ND			
74	1,5	1,3				
76	1,4	0,60				
79	1,4	0,61				
80	1,9	2,7				
84	1,2	0,027				
86	1,5	1,0				
90	1,2	0,033				
93	2,1	3,5				
98	1,2	0,014				
100	1,3	0,36				
103	1,2	-0,050				
106	1,7	2,2				
108	1,3	0,36				

Consensus statistics

Consensus median, pg/g	1,2
Median all values pg/g	1,2
Consensus mean, pg/g	1,3
Standard deviation, pg/g	0,38
Relative standard deviation, %	31
No. of values reported	37
No. of values removed	0
No. of reported non-detects	2

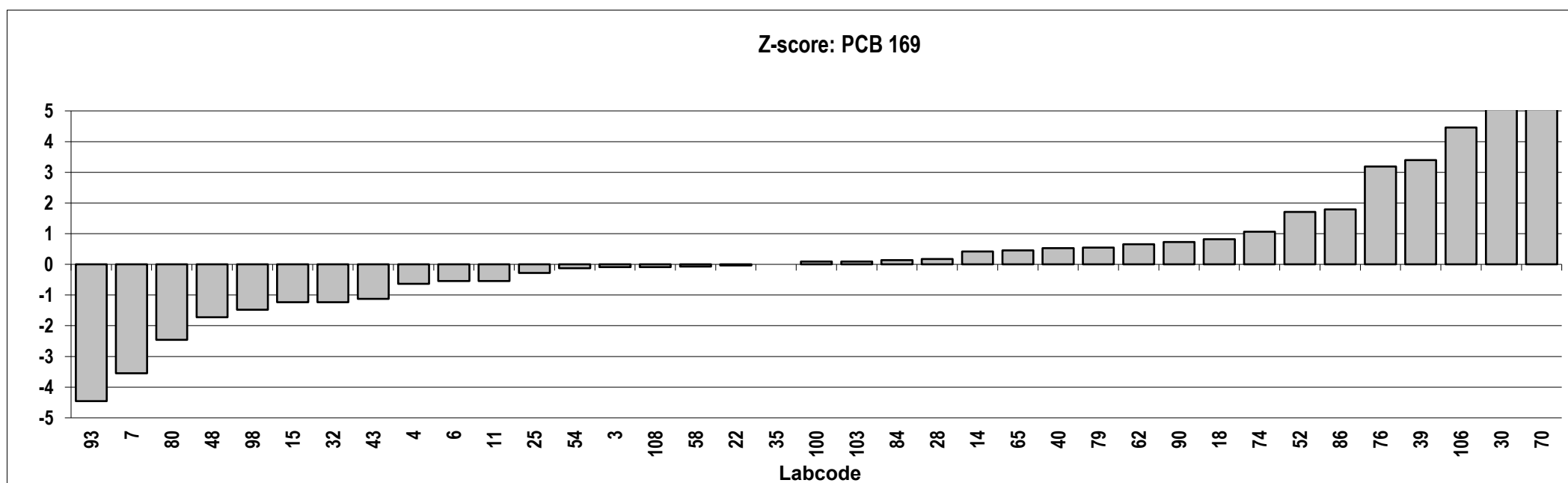
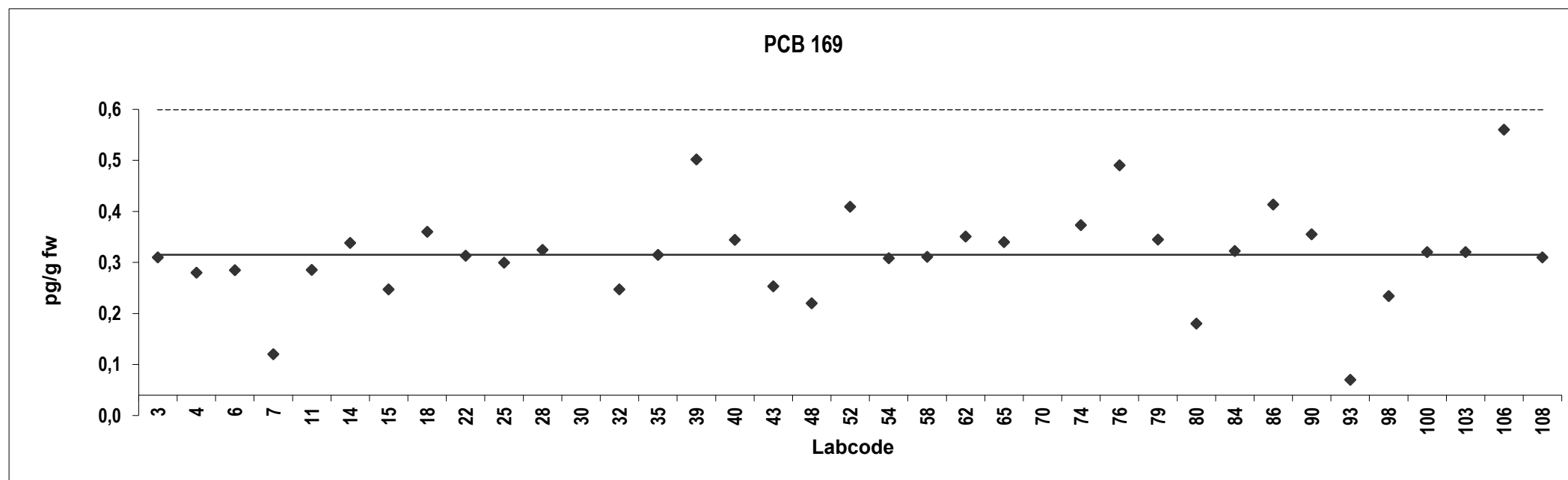


Salmon
Congener: PCB 169

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,27	-0,091				
4	0,24	-0,64				
6	0,25	-0,54				
7	0,080	-3,5				
11	0,25	-0,54				
14	0,30	0,42				
15	0,21	-1,2	ND			
18	0,32	0,82				
22	0,27	-0,033				
25	0,26	-0,28				
28	0,28	0,17				
30	1,0	13	Outlier,ND			
32	0,21	-1,2				
35	0,28	0,0000				
39	0,46	3,4				
40	0,30	0,53				
43	0,21	-1,1				
48	0,18	-1,7	ND			
52	0,37	1,7	ND			
54	0,27	-0,13				
58	0,27	-0,073				
62	0,31	0,65				
65	0,30	0,45				
70	2,0	31	Outlier,ND			
74	0,33	1,1	ND			
76	0,45	3,2				
79	0,31	0,55				
80	0,14	-2,5				
84	0,28	0,13				
86	0,37	1,8				
90	0,32	0,73				
93	0,030	-4,5	ND			
98	0,19	-1,5				
100	0,28	0,091				
103	0,28	0,091				
106	0,52	4,5				
108	0,27	-0,091				

Consensus statistics

Consensus median, pg/g	0,28
Median all values pg/g	0,28
Consensus mean, pg/g	0,28
Standard deviation, pg/g	0,095
Relative standard deviation, %	34
No. of values reported	37
No. of values removed	2
No. of reported non-detects	7

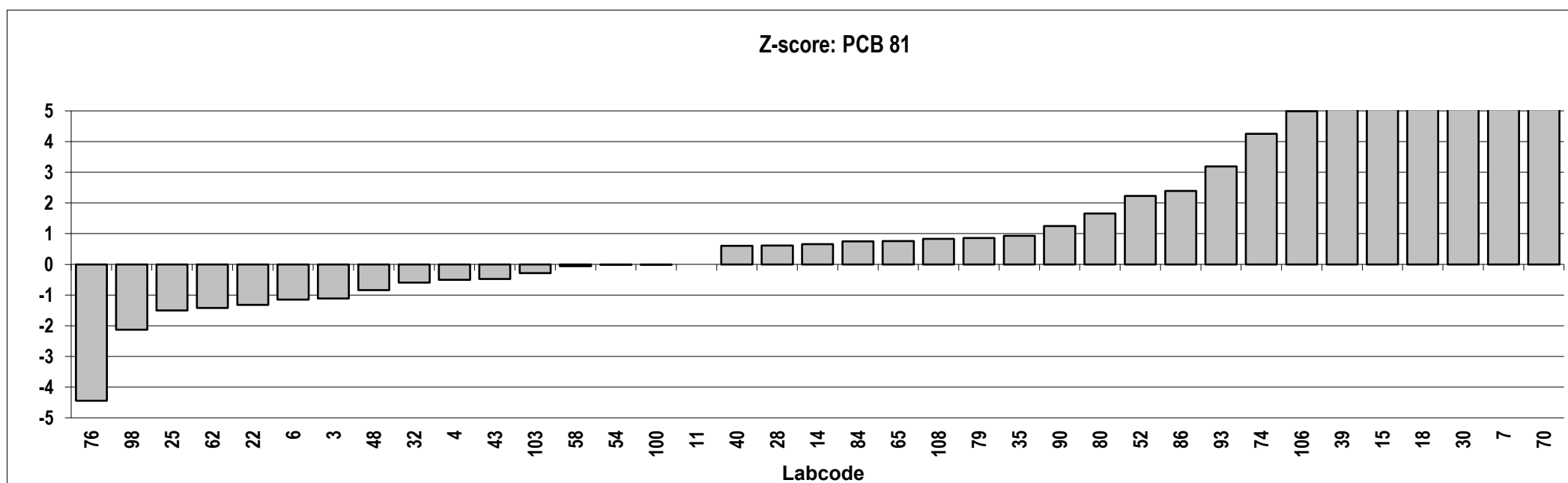
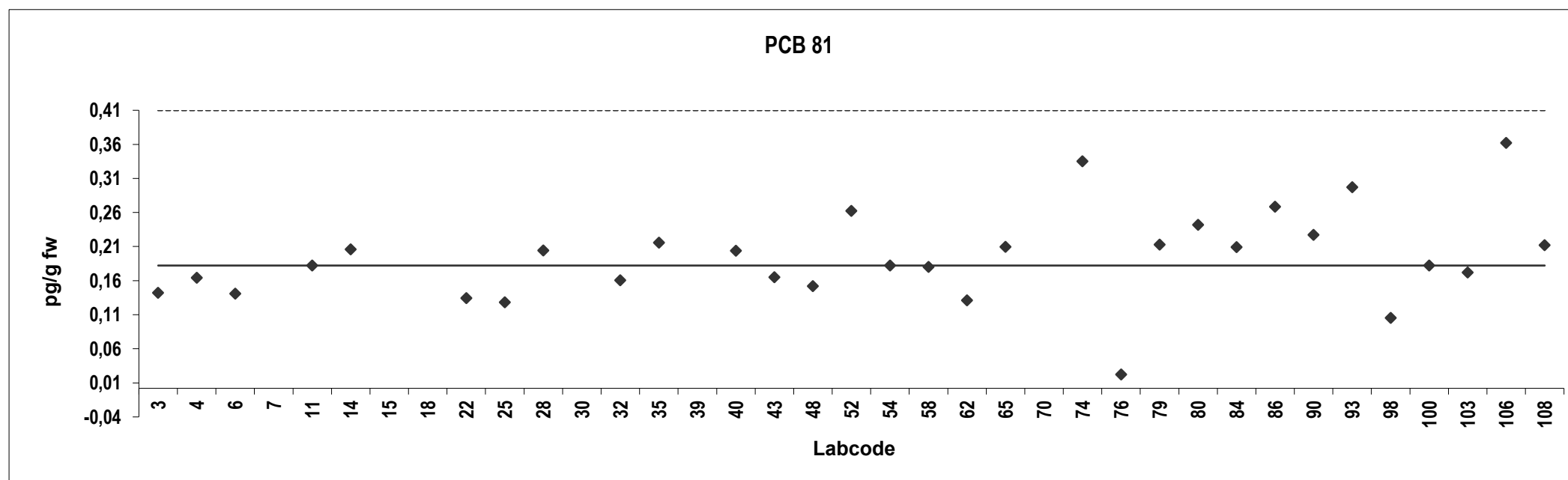


Salmon
Congener: PCB 81

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,14	-1,1				
4	0,16	-0,50				
6	0,14	-1,1				
7	1,4	34	Outlier			
11	0,18	0,0000				
14	0,20	0,66				
15	0,74	15	Outlier			
18	1,0	23	Outlier,ND			
22	0,13	-1,3				
25	0,13	-1,5				
28	0,20	0,61				
30	1,0	23	Outlier,ND			
32	0,16	-0,60				
35	0,21	0,93				
39	0,51	9,0	Outlier			
40	0,20	0,61				
43	0,16	-0,47				
48	0,15	-0,84				
52	0,26	2,2				
54	0,18	-0,0044				
58	0,18	-0,060				
62	0,13	-1,4				
65	0,21	0,76				
70	2,0	51	Outlier,ND			
74	0,33	4,3	ND			
76	0,020	-4,4				
79	0,21	0,86				
80	0,24	1,7				
84	0,21	0,75				
86	0,27	2,4				
90	0,23	1,2				
93	0,30	3,2				
98	0,10	-2,1	ND			
100	0,18	-0,0044				
103	0,17	-0,28				
106	0,36	5,0				
108	0,21	0,83				

Consensus statistics

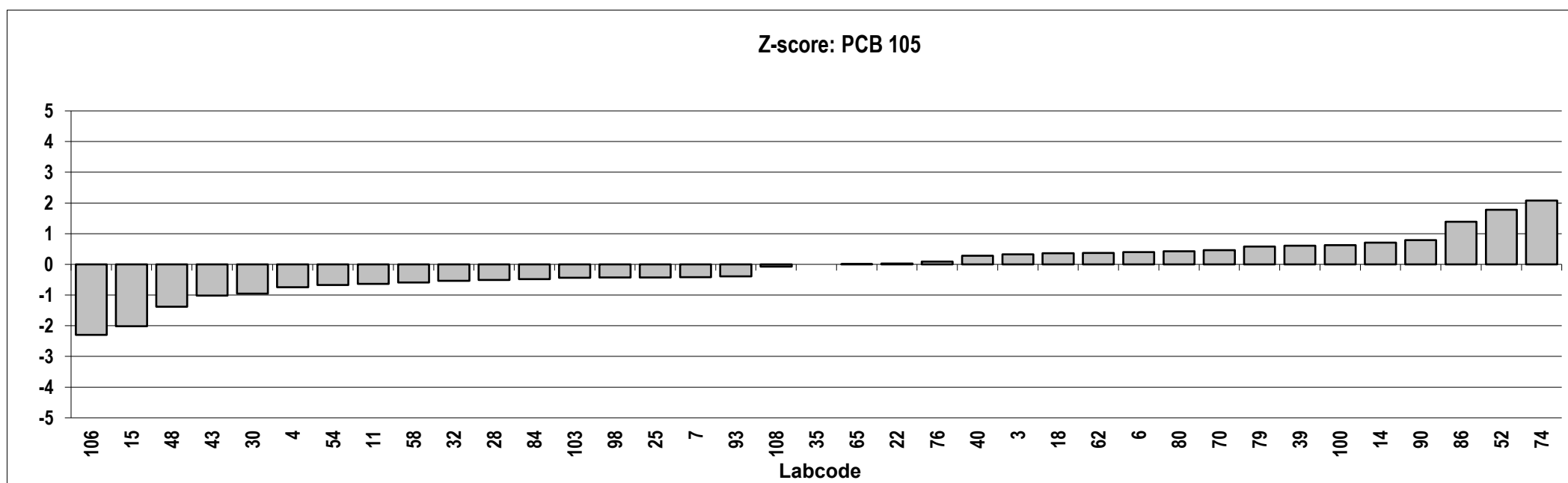
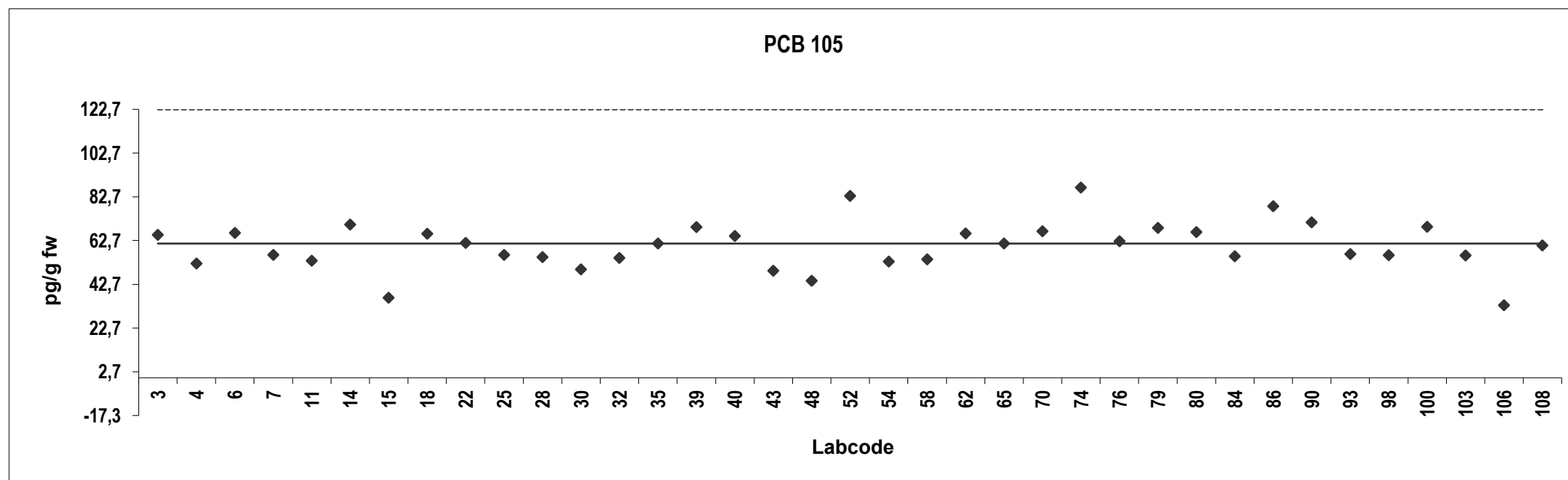
Consensus median, pg/g	0,18
Median all values pg/g	0,20
Consensus mean, pg/g	0,19
Standard deviation, pg/g	0,067
Relative standard deviation, %	35
No. of values reported	37
No. of values removed	6
No. of reported non-detects	5



Salmon
Congener: PCB 105

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	65	0,33				
4	52	-0,75				
6	66	0,40				
7	56	-0,42				
11	54	-0,64				
14	70	0,70				
15	37	-2,0				
18	66	0,36				
22	62	0,028				
25	56	-0,43				
28	55	-0,51				
30	50	-0,96				
32	55	-0,54				
35	61	0,0000				
39	69	0,61				
40	65	0,28				
43	49	-1,0				
48	44	-1,4				
52	83	1,8				
54	53	-0,67				
58	54	-0,59				
62	66	0,37				
65	61	0,0052				
70	67	0,46				
74	87	2,1				
76	62	0,088				
79	69	0,58				
80	67	0,43				
84	56	-0,48				
86	78	1,4				
90	71	0,79				
93	57	-0,39				
98	56	-0,43				
100	69	0,62				
103	56	-0,44				
106	33	-2,3				
108	61	-0,071				

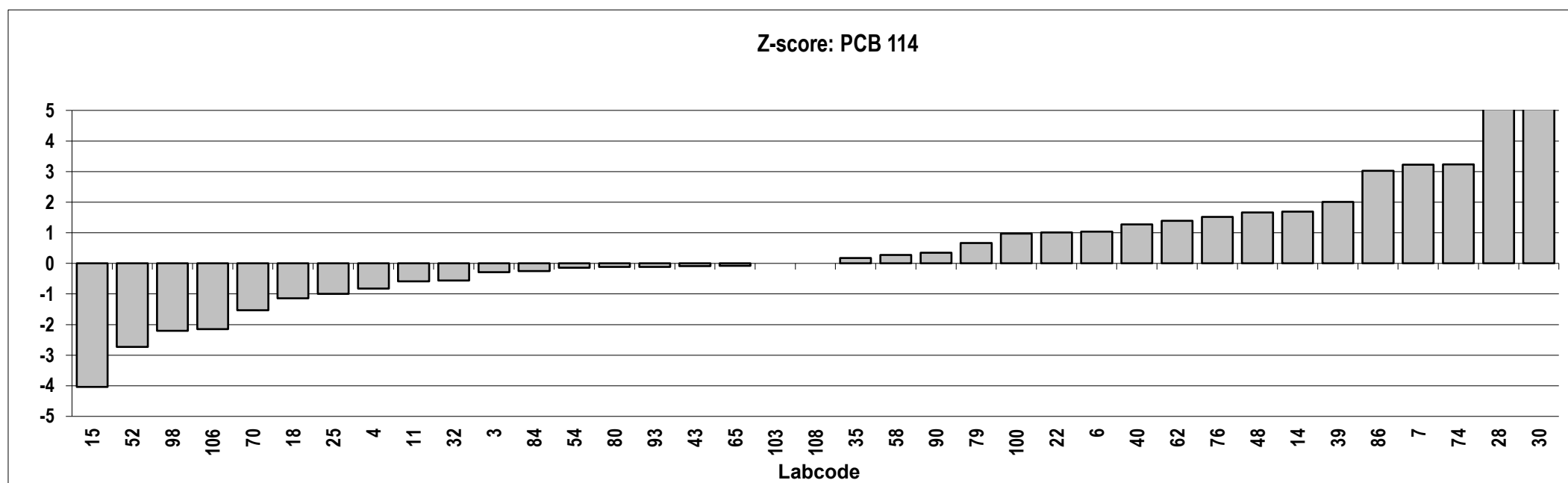
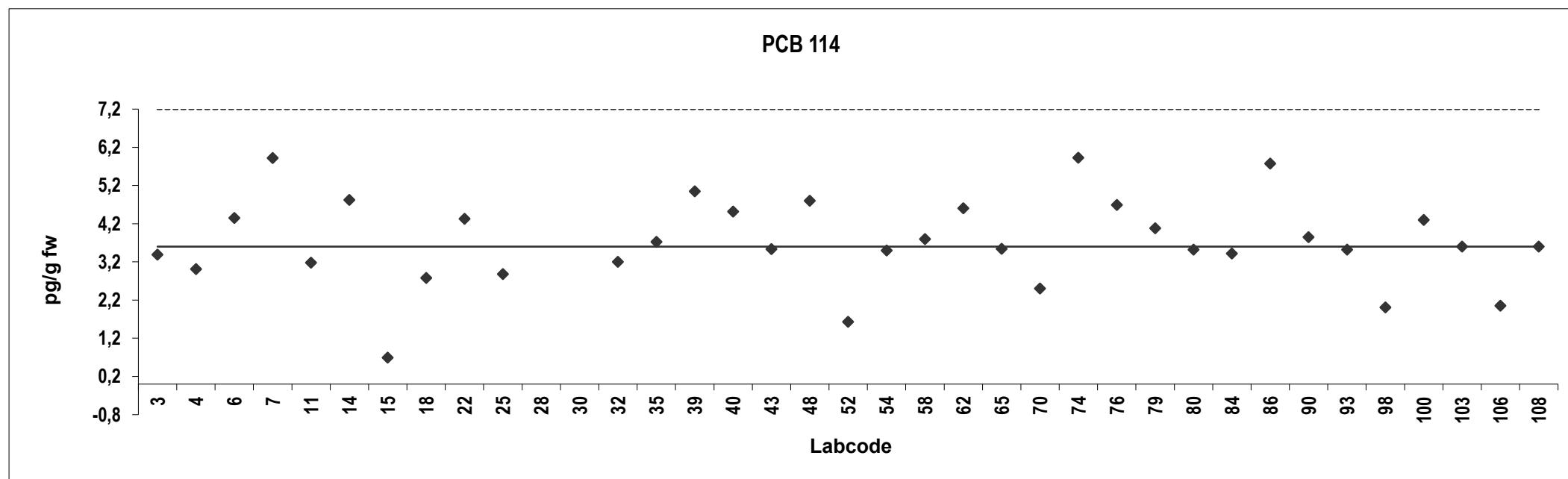
Consensus statistics	
Consensus median, pg/g	61
Median all values pg/g	61
Consensus mean, pg/g	60
Standard deviation, pg/g	11
Relative standard deviation, %	18
No. of values reported	37
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PCB 114

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	3,4	-0,29				
4	3,0	-0,82				
6	4,3	1,0				
7	5,9	3,2				
11	3,2	-0,59				
14	4,8	1,7				
15	0,69	-4,0				
18	2,8	-1,1				
22	4,3	1,0				
25	2,9	-1,0				
28	9,5	8,3	Outlier,ND			
30	10	8,9	Outlier,ND			
32	3,2	-0,56				
35	3,7	0,17				
39	5,0	2,0				
40	4,5	1,3				
43	3,5	-0,089				
48	4,8	1,7	ND			
52	1,6	-2,7				
54	3,5	-0,14				
58	3,8	0,28				
62	4,6	1,4				
65	3,5	-0,075				
70	2,5	-1,5				
74	5,9	3,2				
76	4,7	1,5				
79	4,1	0,67				
80	3,5	-0,11				
84	3,4	-0,25				
86	5,8	3,0				
90	3,9	0,35				
93	3,5	-0,11				
98	2,0	-2,2				
100	4,3	1,0				
103	3,6	0,0000				
106	2,1	-2,2				
108	3,6	0,0000				

Consensus statistics	
Consensus median, pg/g	3,6
Median all values pg/g	3,6
Consensus mean, pg/g	3,7
Standard deviation, pg/g	1,2
Relative standard deviation, %	31
No. of values reported	37
No. of values removed	2
No. of reported non-detects	3

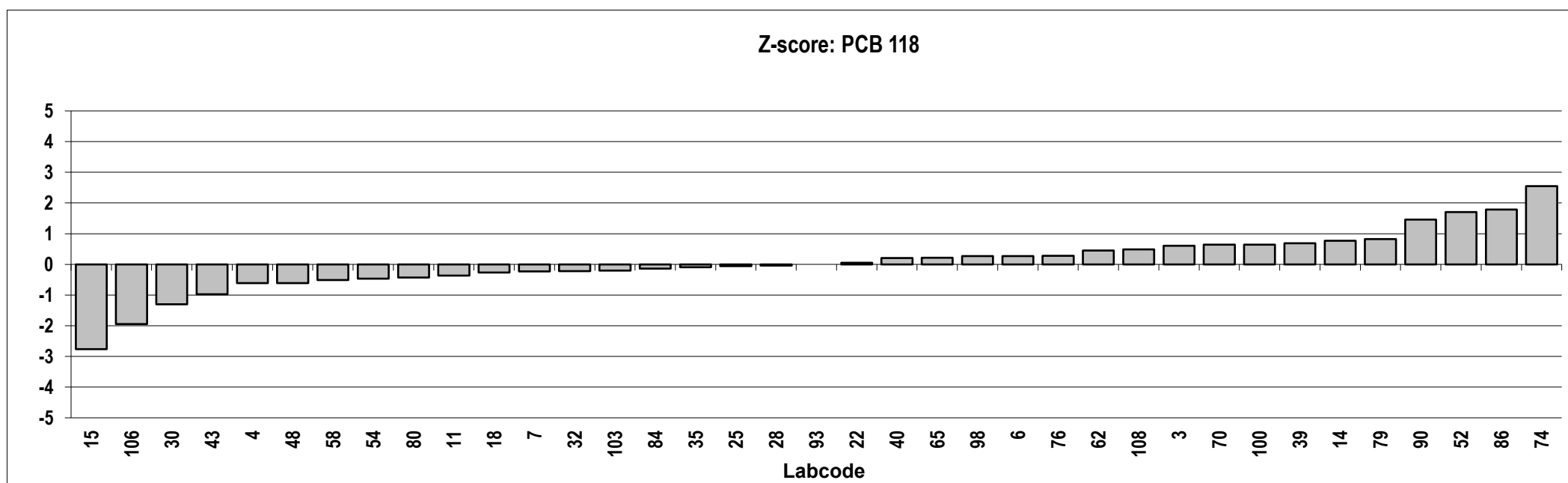
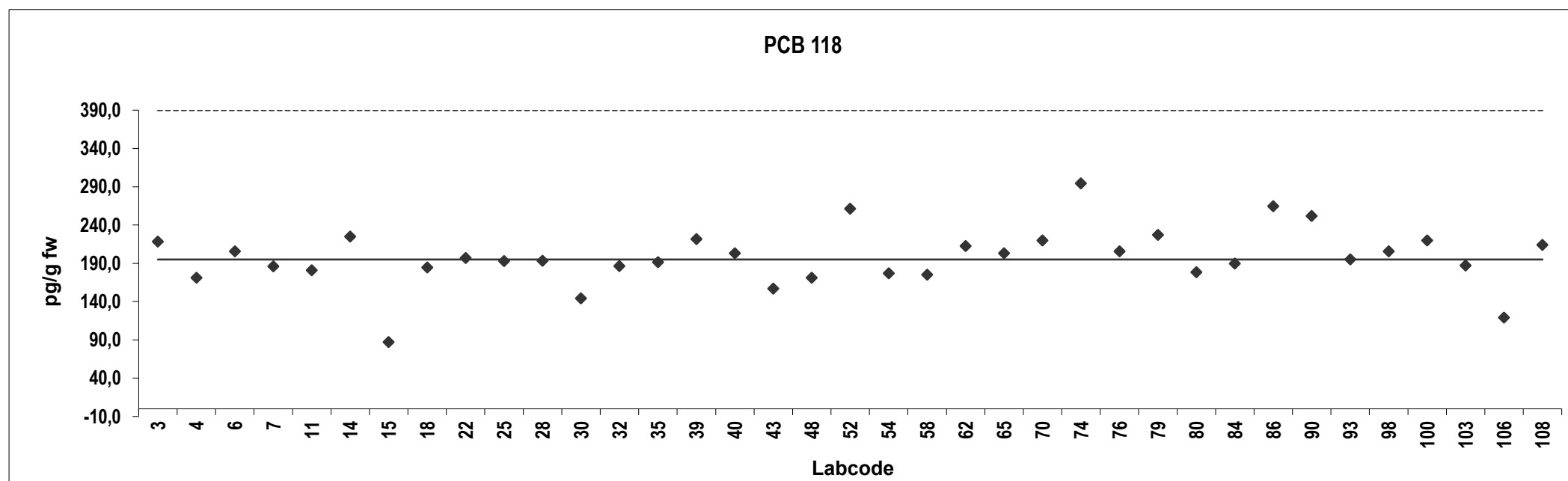


Congener: PCB 118

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	219	0,60				
4	171	-0,62				
6	206	0,27				
7	186	-0,23				
11	181	-0,36				
14	225	0,77				
15	87	-2,8				
18	184	-0,27				
22	197	0,052				
25	193	-0,056				
28	193	-0,044				
30	144	-1,3				
32	186	-0,22				
35	191	-0,092				
39	222	0,68				
40	203	0,21				
43	157	-0,98				
48	171	-0,62				
52	261	1,7				
54	177	-0,46				
58	175	-0,51				
62	213	0,45				
65	203	0,21				
70	220	0,64				
74	294	2,6				
76	206	0,28				
79	227	0,82				
80	178	-0,43				
84	190	-0,14				
86	265	1,8				
90	252	1,5				
93	195	0,0000				
98	206	0,27				
100	220	0,64				
103	187	-0,21				
106	119	-1,9				
108	214	0,49				

Consensus statistics

Consensus median, pg/g	195
Median all values pg/g	195
Consensus mean, pg/g	198
Standard deviation, pg/g	38
Relative standard deviation, %	19
No. of values reported	37
No. of values removed	0
No. of reported non-detects	0

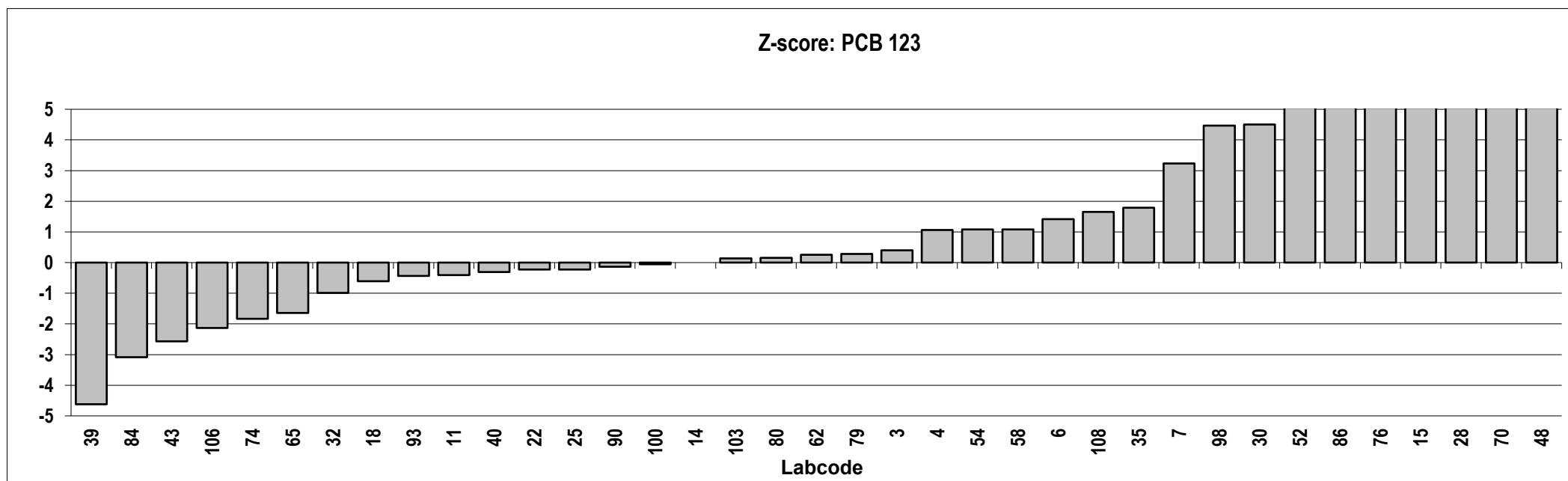
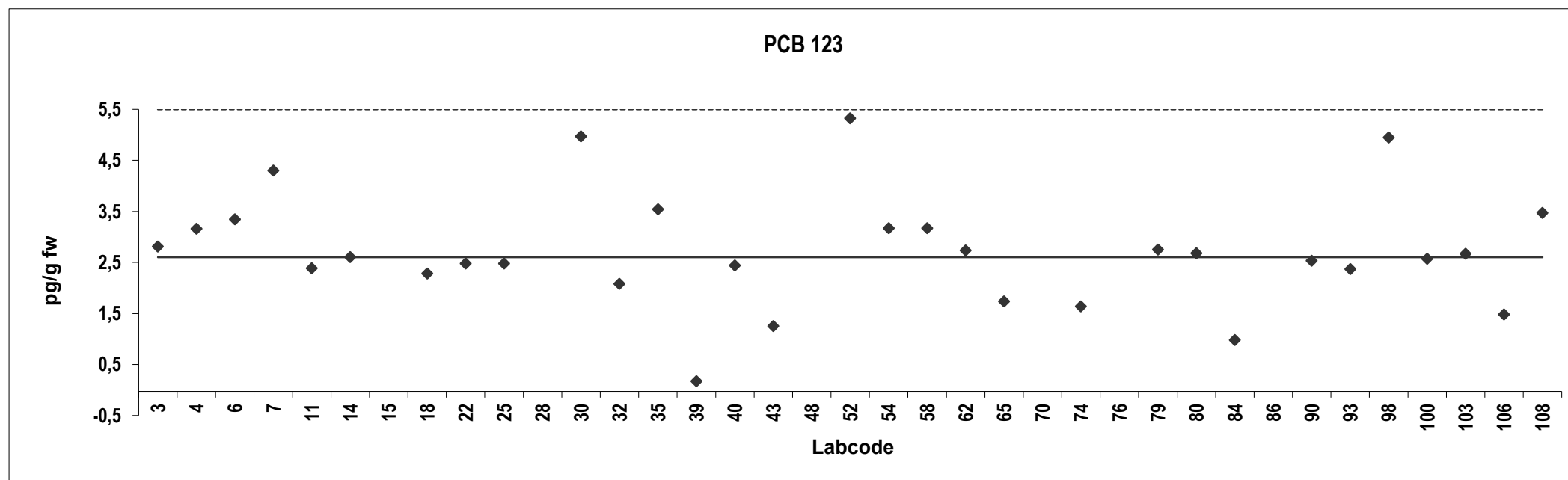


Salmon
Congener: PCB 123

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	2,8	0,40				
4	3,2	1,1				
6	3,4	1,4				
7	4,3	3,2				
11	2,4	-0,41				
14	2,6	0,0000				
15	8,9	12	Outlier			
18	2,3	-0,61				
22	2,5	-0,23				
25	2,5	-0,23				
28	10	14	Outlier,ND			
30	5,0	4,5	ND			
32	2,1	-0,99				
35	3,6	1,8				
39	0,20	-4,6	ND			
40	2,5	-0,30				
43	1,3	-2,6				
48	47	85	Outlier			
52	5,4	5,2				
54	3,2	1,1				
58	3,2	1,1				
62	2,8	0,25				
65	1,8	-1,6				
70	10	14	Outlier,ND			
74	1,7	-1,8	ND			
76	8,5	11	Outlier			
79	2,8	0,29				
80	2,7	0,15				
84	1,0	-3,1				
86	7,7	9,6	Outlier			
90	2,6	-0,13				
93	2,4	-0,44				
98	5,0	4,5				
100	2,6	-0,057				
103	2,7	0,13				
106	1,5	-2,1				
108	3,5	1,7				

Consensus statistics

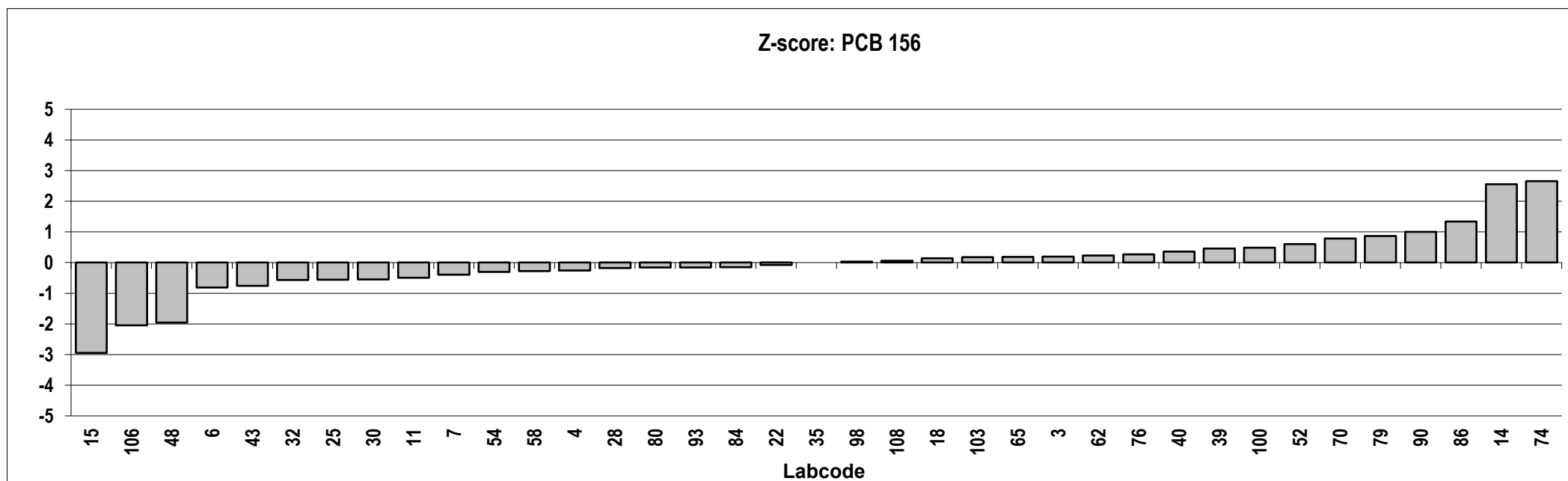
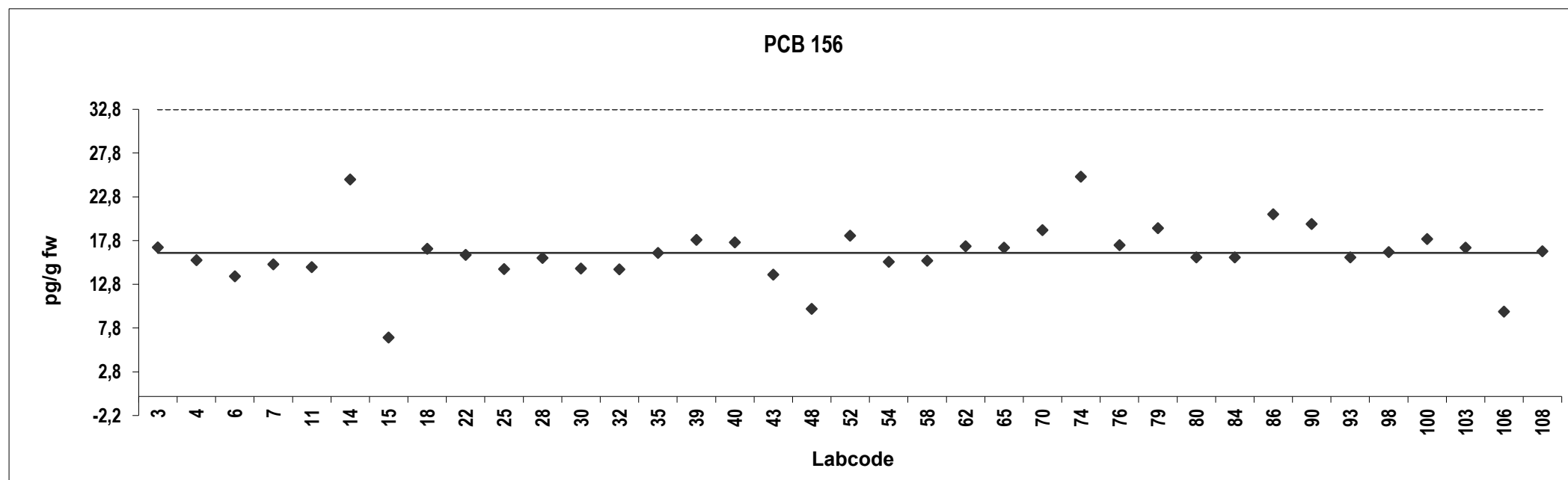
Consensus median, pg/g	2,6
Median all values pg/g	2,8
Consensus mean, pg/g	2,8
Standard deviation, pg/g	1,1
Relative standard deviation, %	41
No. of values reported	37
No. of values removed	6
No. of reported non-detects	5



Salmon
Congener: PCB 156

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	17	0,19	ND			
4	16	-0,26				
6	14	-0,81				
7	15	-0,40				
11	15	-0,50				
14	25	2,6				
15	6,7	-2,9				
18	17	0,14				
22	16	-0,073				
25	15	-0,56				
28	16	-0,17				
30	15	-0,55				
32	15	-0,57				
35	16	0,0000				
39	18	0,46				
40	18	0,36				
43	14	-0,76				
48	10	-2,0				
52	18	0,60				
54	15	-0,31				
58	16	-0,28				
62	17	0,23				
65	17	0,18				
70	19	0,79				
74	25	2,7				
76	17	0,27				
79	19	0,86				
80	16	-0,16				
84	16	-0,15				
86	21	1,3				
90	20	1,0				
93	16	-0,16				
98	17	0,029				
100	18	0,48				
103	17	0,18				
106	9,7	-2,0				
108	17	0,058				

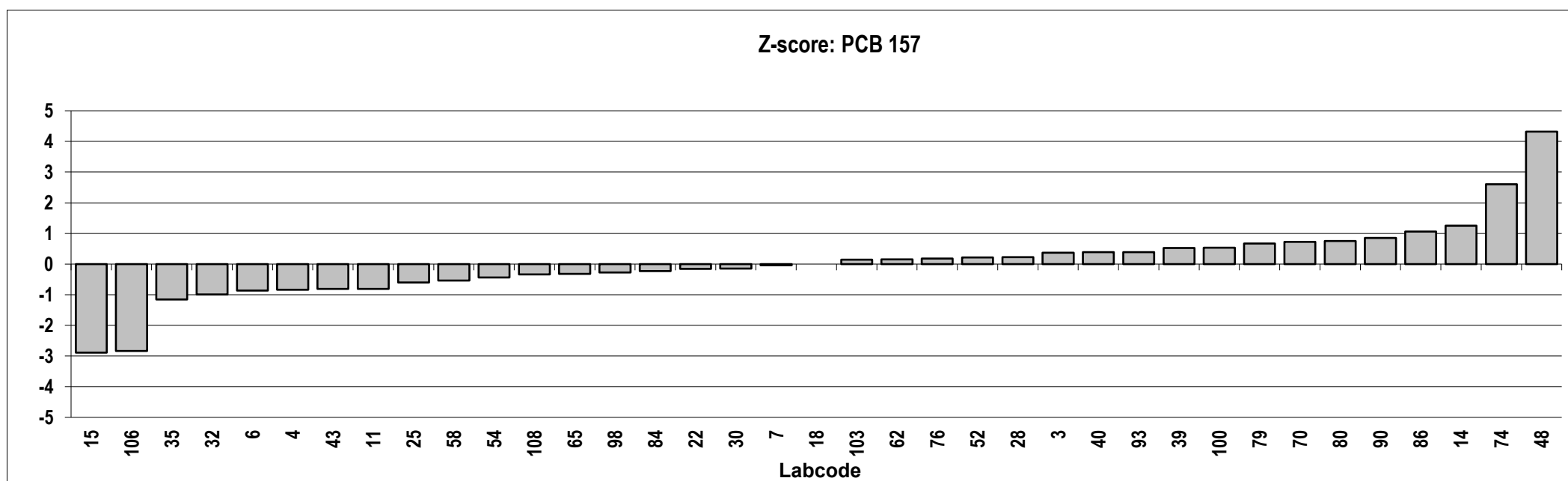
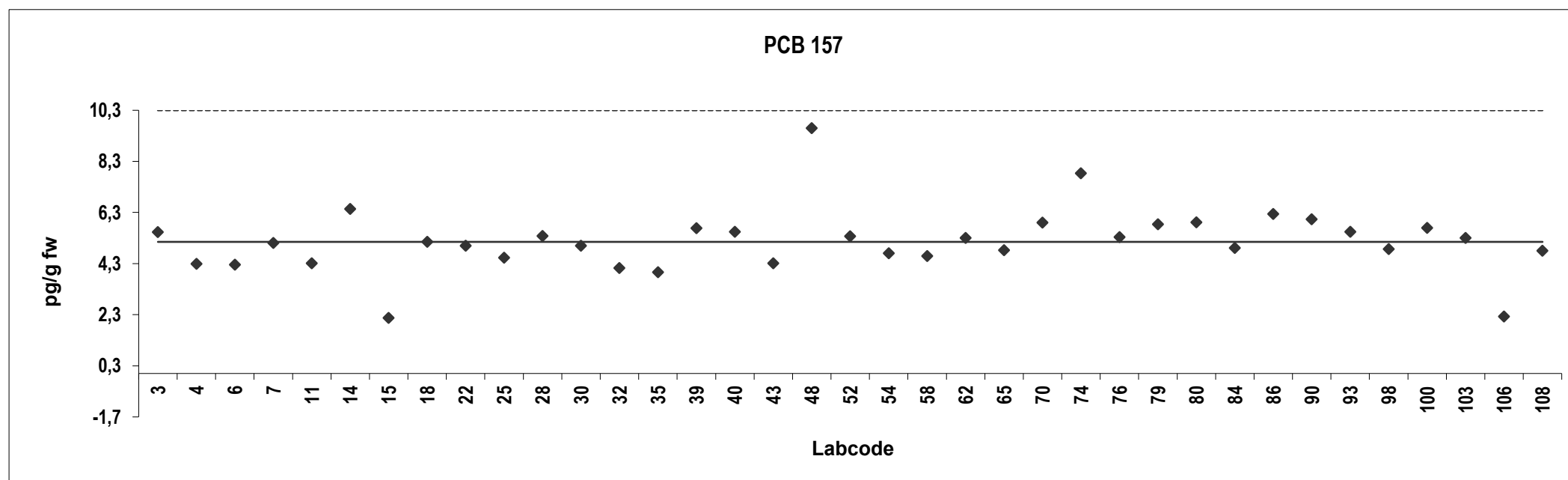
Consensus statistics	
Consensus median, pg/g	16
Median all values pg/g	16
Consensus mean, pg/g	16
Standard deviation, pg/g	3,4
Relative standard deviation, %	21
No. of values reported	37
No. of values removed	0
No. of reported non-detects	1



Salmon
Congener: PCB 157

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	5,5	0,37				
4	4,3	-0,83				
6	4,3	-0,87				
7	5,1	-0,039				
11	4,3	-0,81				
14	6,4	1,3				
15	2,2	-2,9				
18	5,2	0,0000				
22	5,0	-0,15				
25	4,5	-0,60				
28	5,4	0,23				
30	5,0	-0,15	ND			
32	4,1	-0,99				
35	4,0	-1,2				
39	5,7	0,52				
40	5,6	0,39				
43	4,3	-0,81				
48	9,6	4,3	ND			
52	5,4	0,21				
54	4,7	-0,44				
58	4,6	-0,53				
62	5,3	0,15				
65	4,8	-0,32				
70	5,9	0,73				
74	7,8	2,6				
76	5,3	0,18				
79	5,8	0,67				
80	5,9	0,75				
84	4,9	-0,23				
86	6,2	1,1				
90	6,0	0,85				
93	5,6	0,39				
98	4,9	-0,27				
100	5,7	0,53				
103	5,3	0,15				
106	2,2	-2,8				
108	4,8	-0,34				

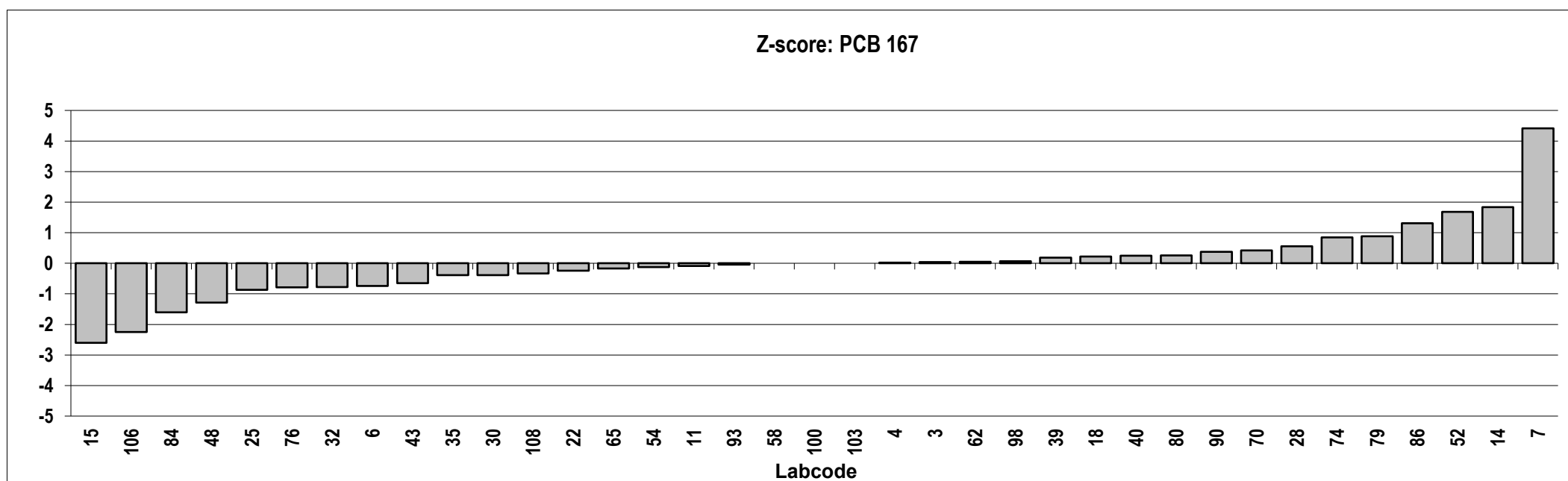
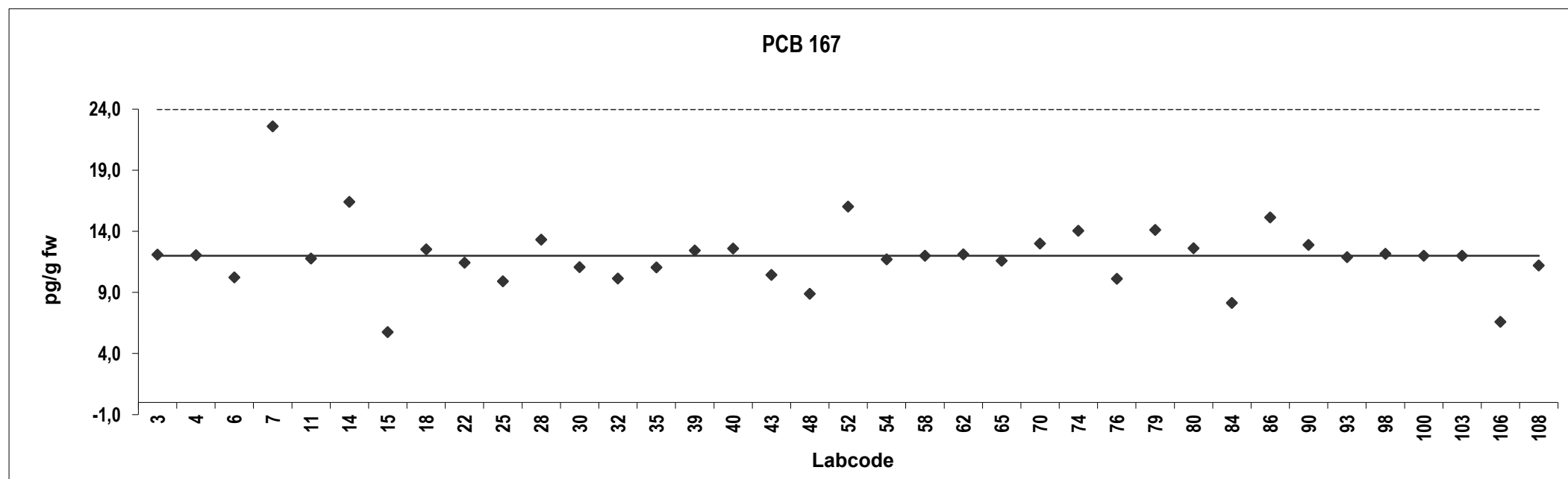
Consensus statistics	
Consensus median, pg/g	5,2
Median all values pg/g	5,2
Consensus mean, pg/g	5,2
Standard deviation, pg/g	1,3
Relative standard deviation, %	24
No. of values reported	37
No. of values removed	0
No. of reported non-detects	2



Salmon
Congener: PCB 167

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	12	0,042	ND			
4	12	0,021				
6	10	-0,74				
7	23	4,4				
11	12	-0,092				
14	16	1,8				
15	5,7	-2,6				
18	13	0,22				
22	11	-0,24				
25	9,9	-0,87				
28	13	0,55				
30	11	-0,39				
32	10	-0,78				
35	11	-0,39				
39	12	0,18				
40	13	0,25				
43	10	-0,65				
48	8,9	-1,3				
52	16	1,7				
54	12	-0,13				
58	12	0,0000				
62	12	0,050				
65	12	-0,17				
70	13	0,42				
74	14	0,85				
76	10	-0,79				
79	14	0,88				
80	13	0,25				
84	8,1	-1,6				
86	15	1,3				
90	13	0,38				
93	12	-0,042				
98	12	0,069				
100	12	0,0000				
103	12	0,0000				
106	6,6	-2,3				
108	11	-0,33				

Consensus statistics	
Consensus median, pg/g	12
Median all values pg/g	12
Consensus mean, pg/g	12
Standard deviation, pg/g	2,8
Relative standard deviation, %	23
No. of values reported	37
No. of values removed	0
No. of reported non-detects	1

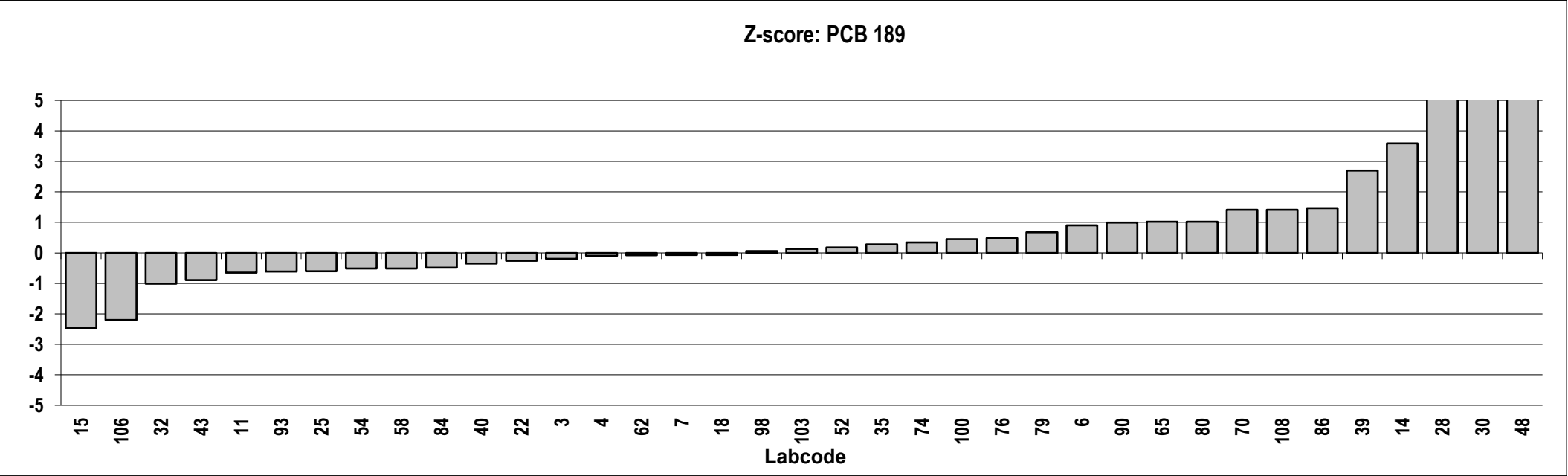
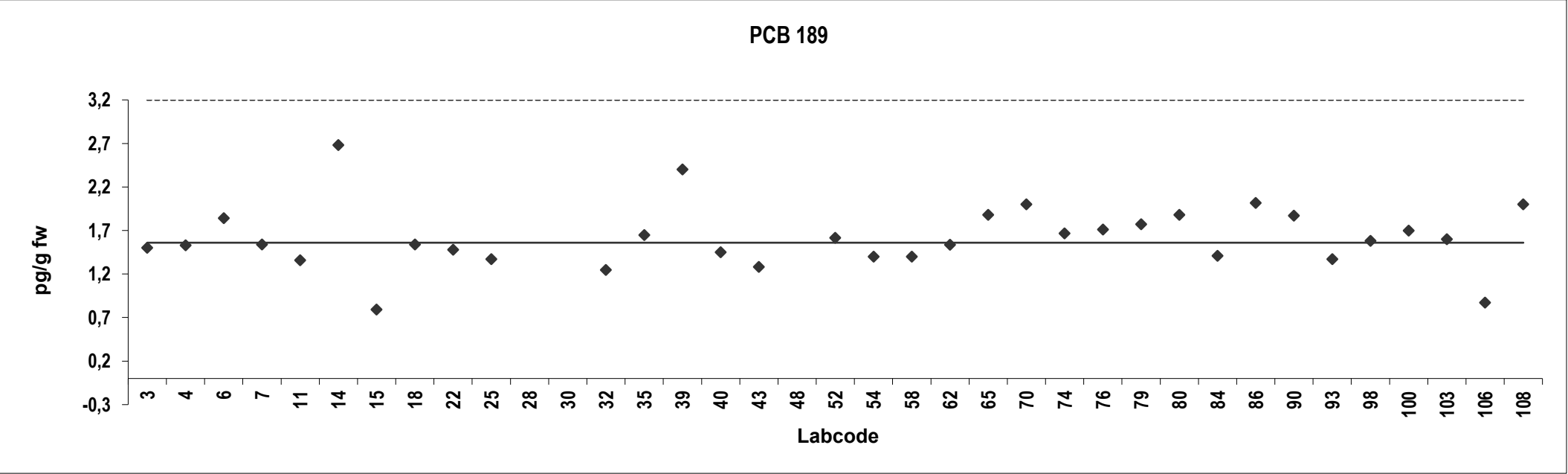


Salmon
Congener: PCB 189

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	1,5	-0,19				
4	1,5	-0,10				
6	1,8	0,90				
7	1,5	-0,063				
11	1,4	-0,65				
14	2,7	3,6				
15	0,79	-2,5				
18	1,5	-0,063				
22	1,5	-0,26				
25	1,4	-0,60				
28	3,4	6,0	Outlier,ND			
30	5,0	11	Outlier,ND			
32	1,2	-1,0				
35	1,6	0,28				
39	2,4	2,7				
40	1,5	-0,35				
43	1,3	-0,89				
48	12	33	Outlier,ND			
52	1,6	0,18				
54	1,4	-0,51				
58	1,4	-0,51				
62	1,5	-0,076				
65	1,9	1,0				
70	2,0	1,4	ND			
74	1,7	0,34	ND			
76	1,7	0,49				
79	1,8	0,68				
80	1,9	1,0				
84	1,4	-0,49				
86	2,0	1,5				
90	1,9	0,99				
93	1,4	-0,61				
98	1,6	0,063				
100	1,7	0,45				
103	1,6	0,13				
106	0,87	-2,2				
108	2,0	1,4				

Consensus statistics

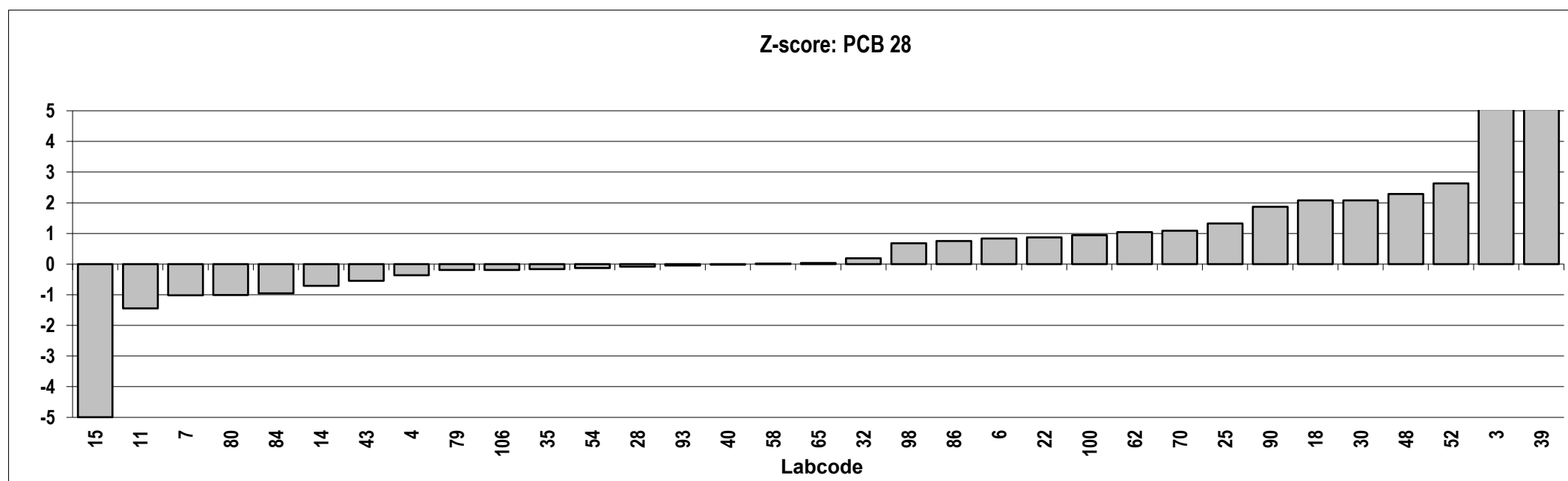
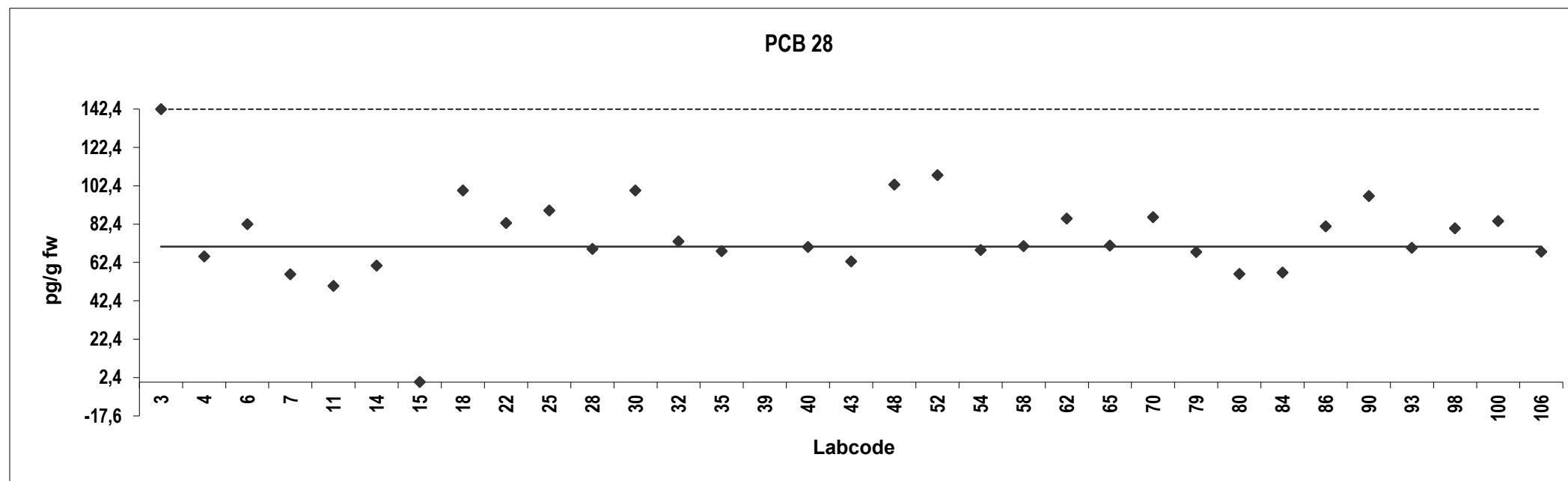
Consensus median, pg/g	1,6
Median all values pg/g	1,6
Consensus mean, pg/g	1,6
Standard deviation, pg/g	0,37
Relative standard deviation, %	23
No. of values reported	37
No. of values removed	3
No. of reported non-detects	5



Salmon
Congener: PCB 28

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	142	5,1				
4	66	-0,37				
6	82	0,83				
7	56	-1,0				
11	50	-1,4				
14	61	-0,71				
15	0,031	-5,0				
18	100	2,1	ND			
22	83	0,87				
25	89	1,3				
28	69	-0,083				
30	100	2,1	ND			
32	73	0,19				
35	68	-0,16				
39	379	22	Outlier			
40	70	-0,016				
43	63	-0,54				
48	103	2,3				
52	108	2,6				
54	69	-0,13				
58	71	0,016				
62	85	1,0				
65	71	0,037				
70	86	1,1				
79	68	-0,20				
80	56	-1,0				
84	57	-1,0				
86	81	0,75				
90	97	1,9				
93	70	-0,048				
98	80	0,68				
100	84	0,94				
106	68	-0,19				

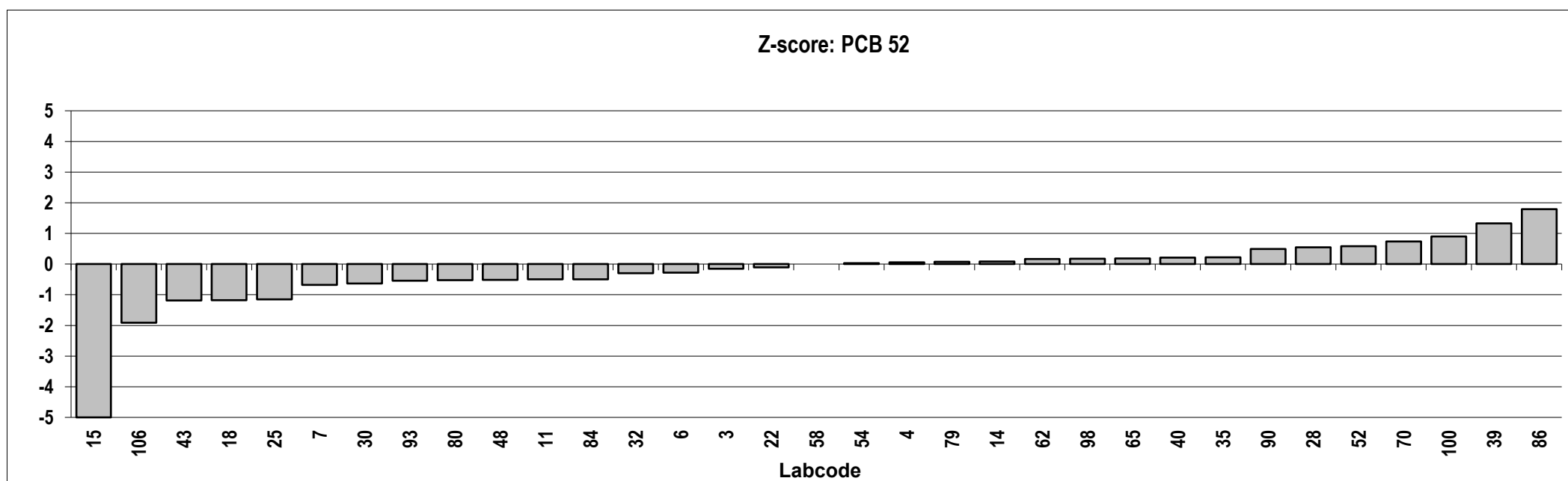
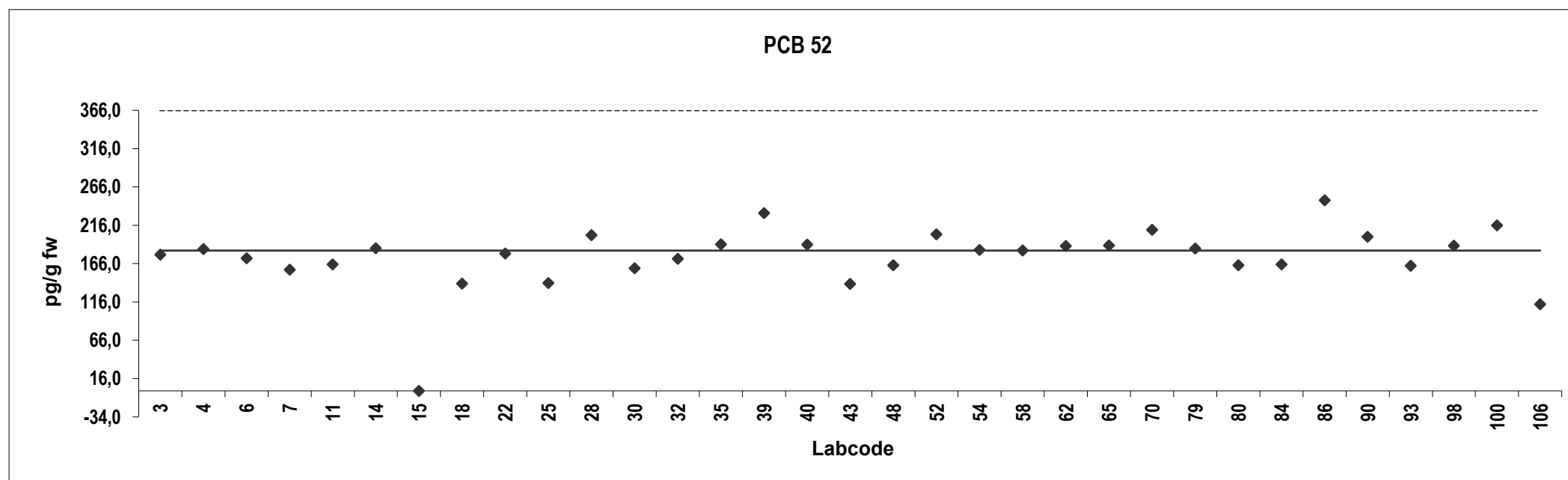
Consensus statistics	
Consensus median, pg/g	71
Median all values pg/g	71
Consensus mean, pg/g	76
Standard deviation, pg/g	23
Relative standard deviation, %	31
No. of values reported	33
No. of values removed	1
No. of reported non-detects	2



Salmon
Congener: PCB 52

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	178	-0,15				
4	185	0,055				
6	173	-0,28				
7	158	-0,68				
11	165	-0,50				
14	186	0,082				
15	0,083	-5,0				
18	140	-1,2				
22	179	-0,11				
25	141	-1,2				
28	203	0,55				
30	160	-0,63				
32	172	-0,30				
35	191	0,22				
39	232	1,3				
40	191	0,22				
43	139	-1,2				
48	164	-0,52				
52	204	0,58				
54	184	0,027				
58	183	0,0000				
62	189	0,17				
65	190	0,18				
70	210	0,74				
79	186	0,078				
80	164	-0,52				
84	165	-0,50				
86	249	1,8				
90	201	0,49				
93	163	-0,55				
98	189	0,17				
100	216	0,90				
106	113	-1,9				

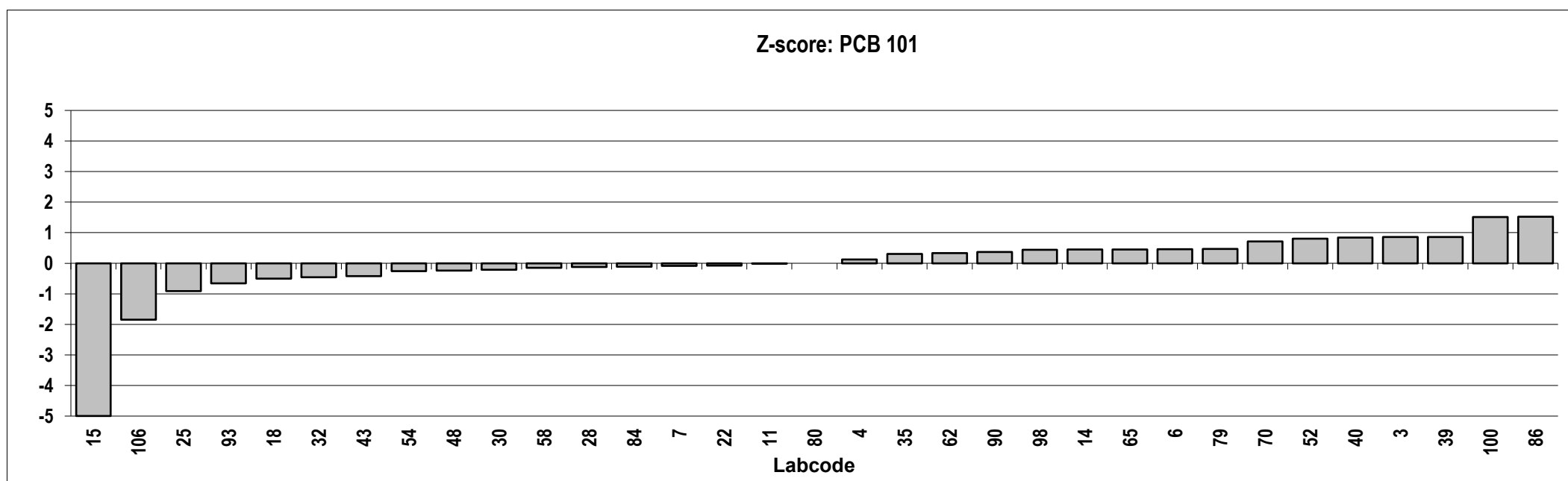
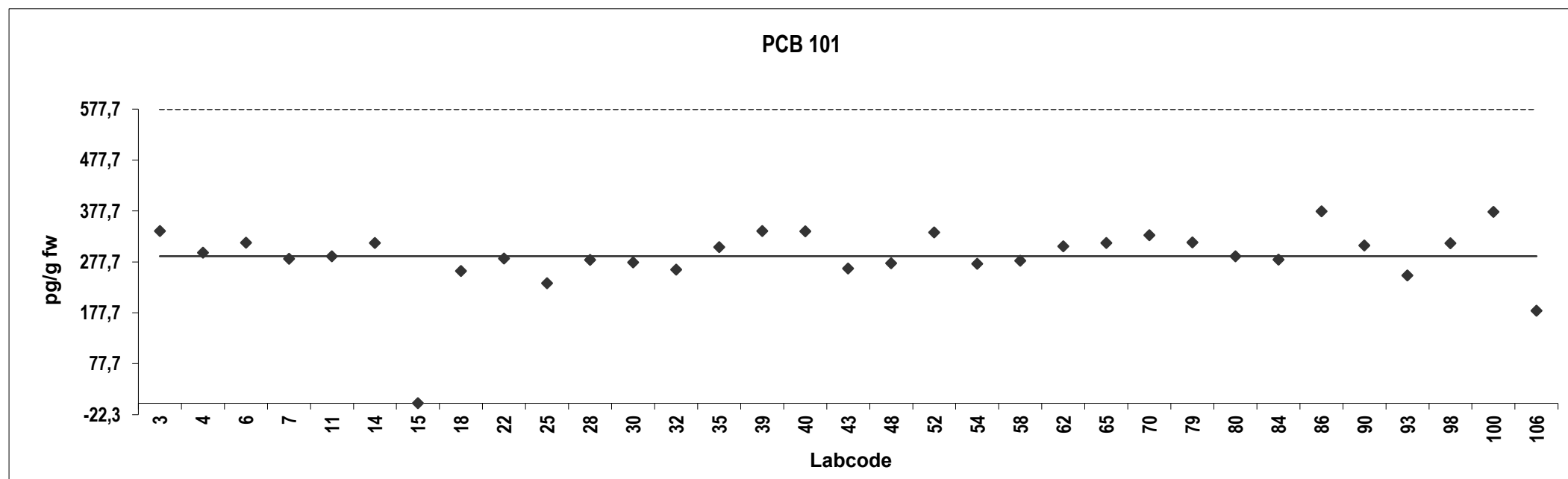
Consensus statistics	
Consensus median, pg/g	183
Median all values pg/g	183
Consensus mean, pg/g	175
Standard deviation, pg/g	41
Relative standard deviation, %	24
No. of values reported	33
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PCB 101

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	338	0,85				
4	296	0,12				
6	315	0,46				
7	284	-0,084				
11	289	-0,0042				
14	315	0,45				
15	0,14	-5,0				
18	260	-0,50				
22	284	-0,075				
25	236	-0,92				
28	282	-0,12				
30	277	-0,21				
32	263	-0,45				
35	307	0,31				
39	338	0,86				
40	338	0,84				
43	265	-0,42				
48	275	-0,24				
52	336	0,81				
54	274	-0,26				
58	280	-0,15				
62	308	0,33				
65	315	0,45				
70	330	0,71				
79	316	0,47				
80	289	0,0000				
84	282	-0,11				
86	377	1,5				
90	310	0,37				
93	251	-0,65				
98	314	0,44				
100	376	1,5				
106	182	-1,8				

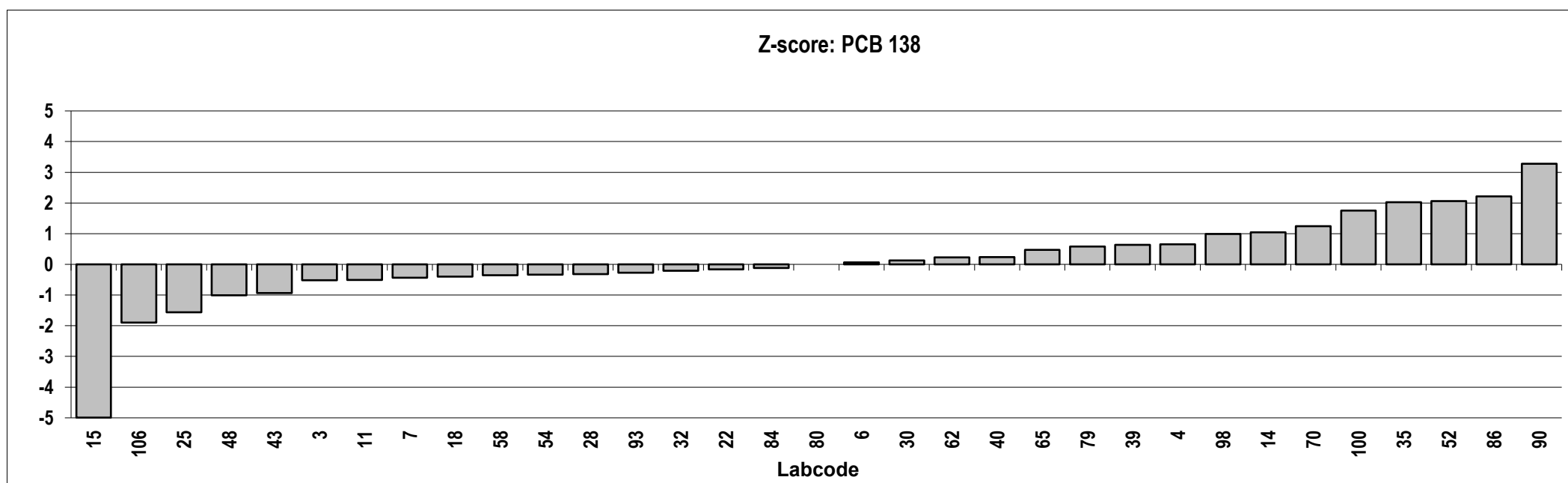
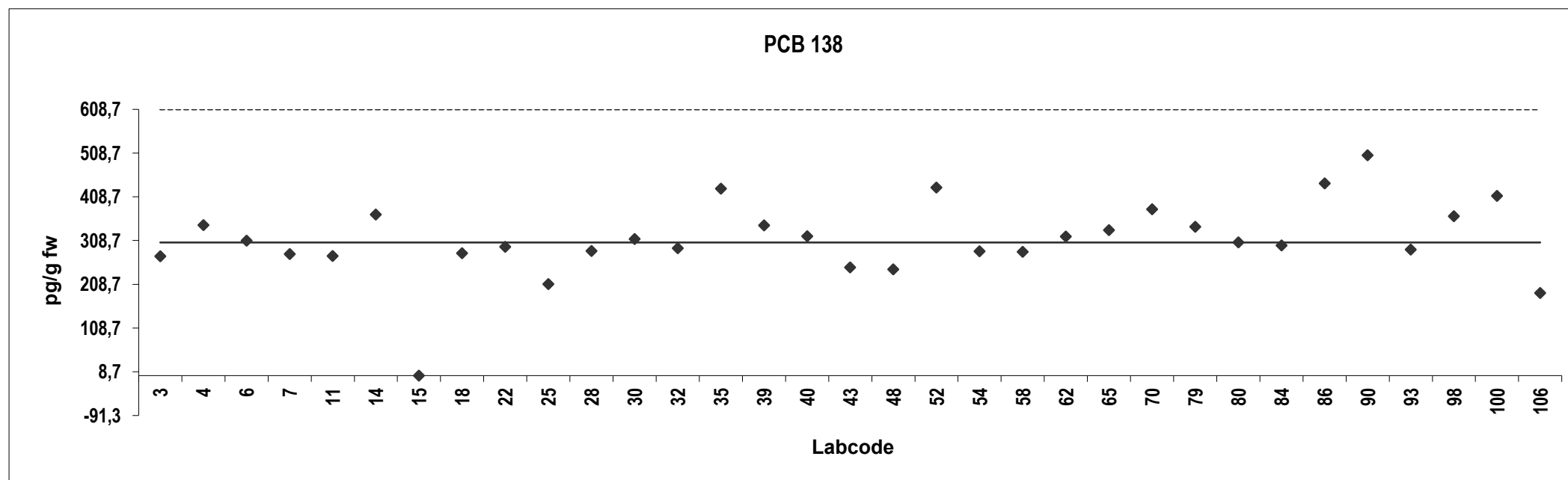
Consensus statistics	
Consensus median, pg/g	289
Median all values pg/g	289
Consensus mean, pg/g	288
Standard deviation, pg/g	64
Relative standard deviation, %	22
No. of values reported	33
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PCB 138

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	273	-0,52				
4	344	0,65				
6	308	0,063				
7	278	-0,43				
11	273	-0,51				
14	368	1,0				
15	0,20	-5,0				
18	280	-0,40				
22	295	-0,16				
25	209	-1,6				
28	285	-0,32				
30	312	0,13				
32	291	-0,21				
35	427	2,0				
39	343	0,64				
40	318	0,23				
43	247	-0,94				
48	243	-1,0				
52	430	2,1				
54	284	-0,33				
58	283	-0,35				
62	318	0,23				
65	333	0,47				
70	380	1,2				
79	340	0,58				
80	304	0,0000				
84	297	-0,11				
86	439	2,2				
90	504	3,3				
93	288	-0,27				
98	365	0,99				
100	411	1,8				
106	189	-1,9				

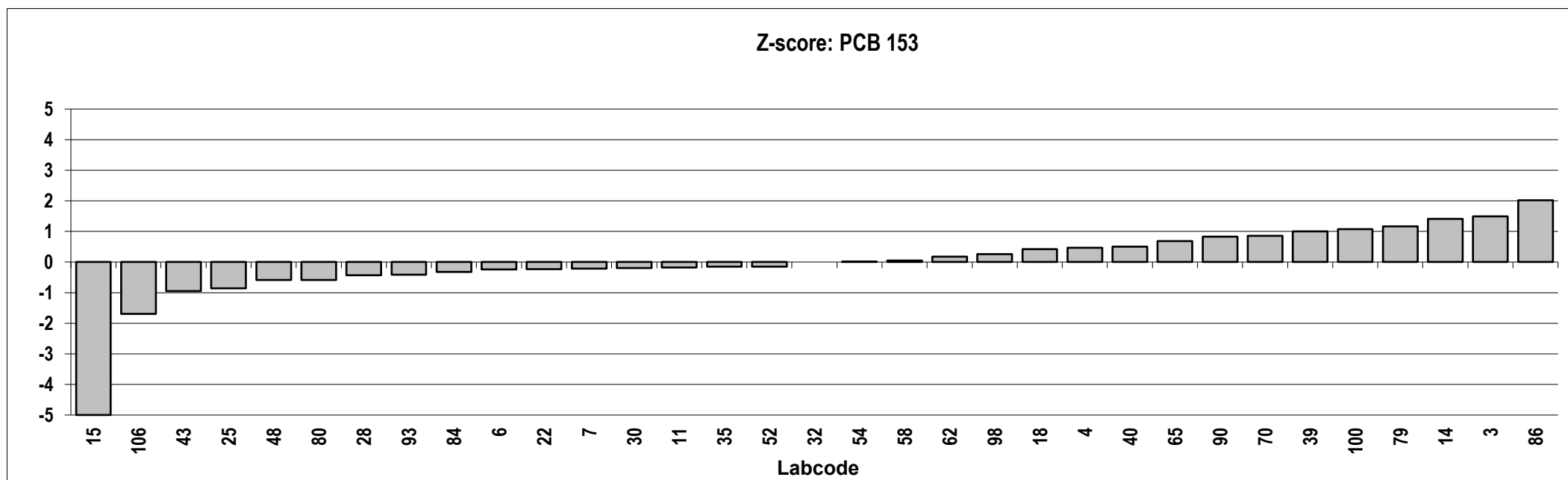
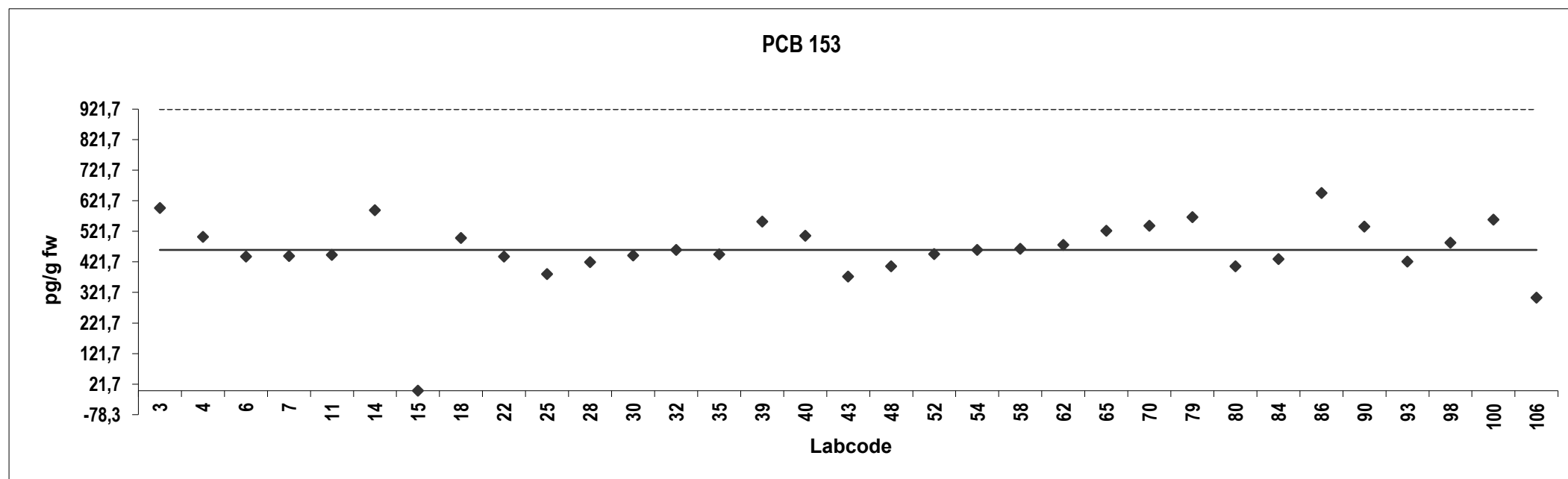
Consensus statistics	
Consensus median, pg/g	304
Median all values pg/g	304
Consensus mean, pg/g	311
Standard deviation, pg/g	87
Relative standard deviation, %	28
No. of values reported	33
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PCB 153

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	598	1,5				
4	504	0,47				
6	439	-0,24				
7	441	-0,22				
11	444	-0,18				
14	591	1,4				
15	0,24	-5,0				
18	500	0,42				
22	439	-0,23				
25	381	-0,86				
28	421	-0,43				
30	443	-0,20				
32	461	0,0000				
35	447	-0,15				
39	553	1,0				
40	507	0,50				
43	373	-0,95				
48	407	-0,58				
52	447	-0,15				
54	461	0,0015				
58	465	0,045				
62	477	0,18				
65	524	0,68				
70	540	0,86				
79	569	1,2				
80	407	-0,58				
84	431	-0,33				
86	647	2,0				
90	537	0,83				
93	423	-0,41				
98	485	0,26				
100	560	1,1				
106	305	-1,7				

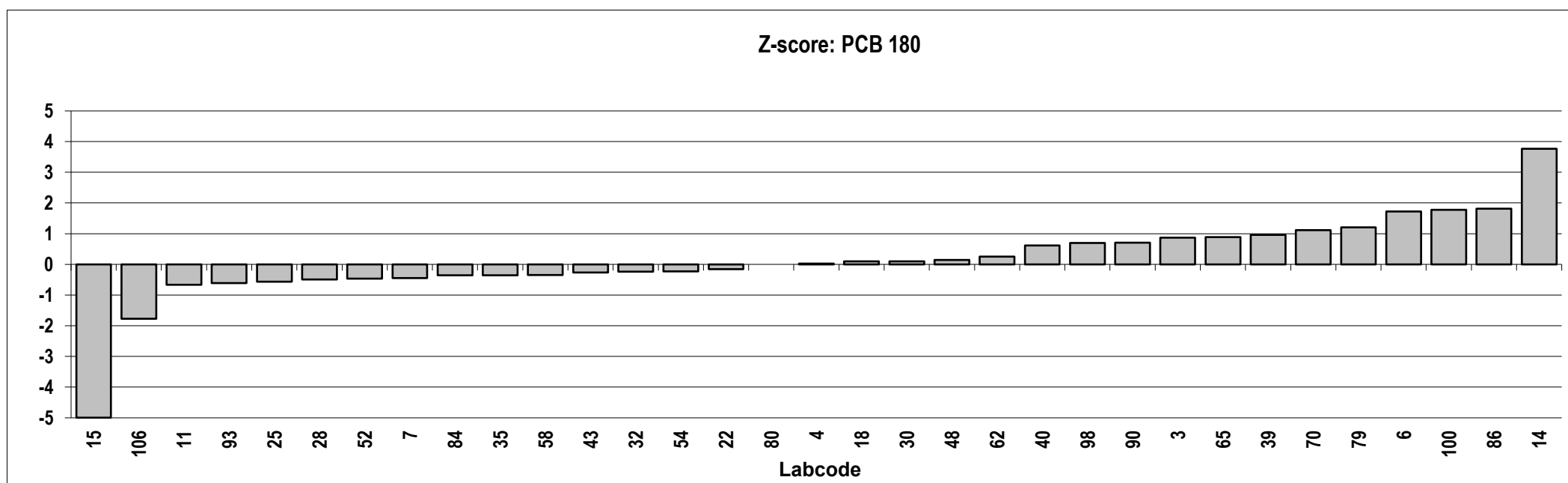
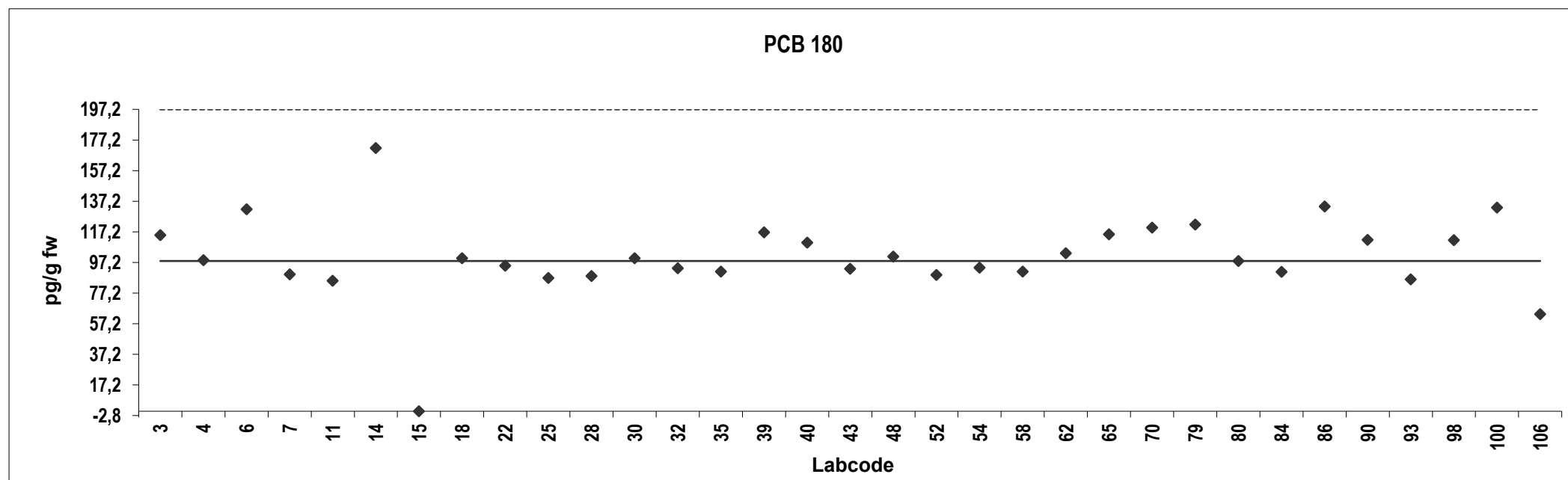
Consensus statistics	
Consensus median, pg/g	461
Median all values pg/g	461
Consensus mean, pg/g	461
Standard deviation, pg/g	110
Relative standard deviation, %	24
No. of values reported	33
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PCB 180

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	115	0,87				
4	99	0,023				
6	132	1,7				
7	89	-0,45				
11	85	-0,66				
14	172	3,8				
15	0,045	-5,0				
18	100	0,095	ND			
22	95	-0,16				
25	87	-0,56				
28	88	-0,50				
30	100	0,095	ND			
32	93	-0,24				
35	91	-0,35				
39	117	0,96				
40	110	0,61				
43	93	-0,26				
48	101	0,15				
52	89	-0,46				
54	94	-0,23				
58	91	-0,35				
62	103	0,26				
65	116	0,89				
70	120	1,1				
79	122	1,2				
80	98	0,0000				
84	91	-0,36				
86	134	1,8				
90	112	0,71				
93	86	-0,61				
98	112	0,69				
100	133	1,8				
106	63	-1,8				

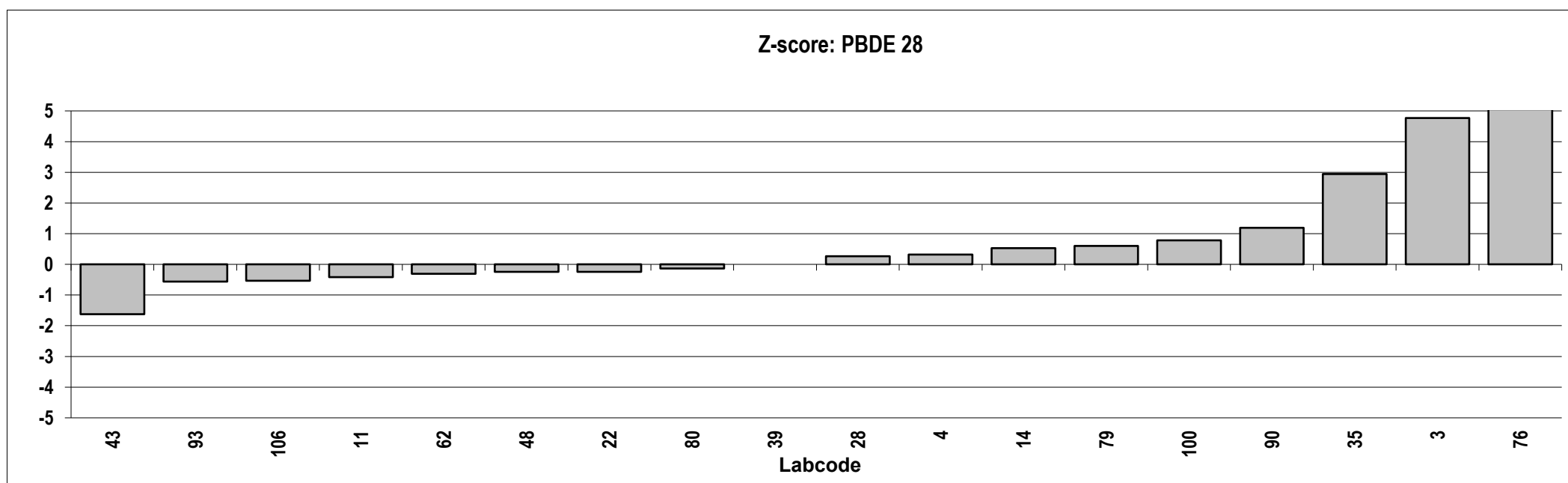
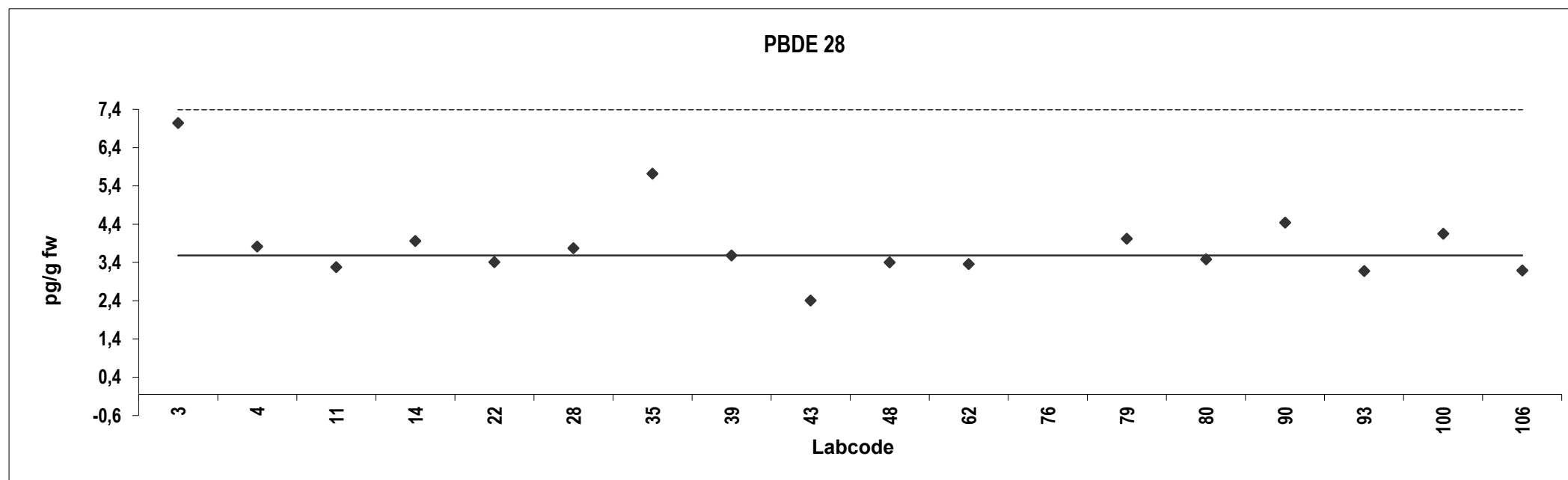
Consensus statistics	
Consensus median, pg/g	98
Median all values pg/g	99
Consensus mean, pg/g	101
Standard deviation, pg/g	27
Relative standard deviation, %	27
No. of values reported	33
No. of values removed	0
No. of reported non-detects	2



Salmon
Congener: PBDE 28

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	7,1	4,8	Outlier			
4	3,9	0,32				
11	3,3	-0,42				
14	4,0	0,52				
22	3,5	-0,24				
28	3,8	0,26				
35	5,8	2,9				
39	3,6	0,0000				
43	2,5	-1,6				
48	3,5	-0,25				
62	3,4	-0,31				
76	35	43				
79	4,1	0,60				
80	3,5	-0,14				
90	4,5	1,2				
93	3,2	-0,56				
100	4,2	0,79				
106	3,2	-0,54				

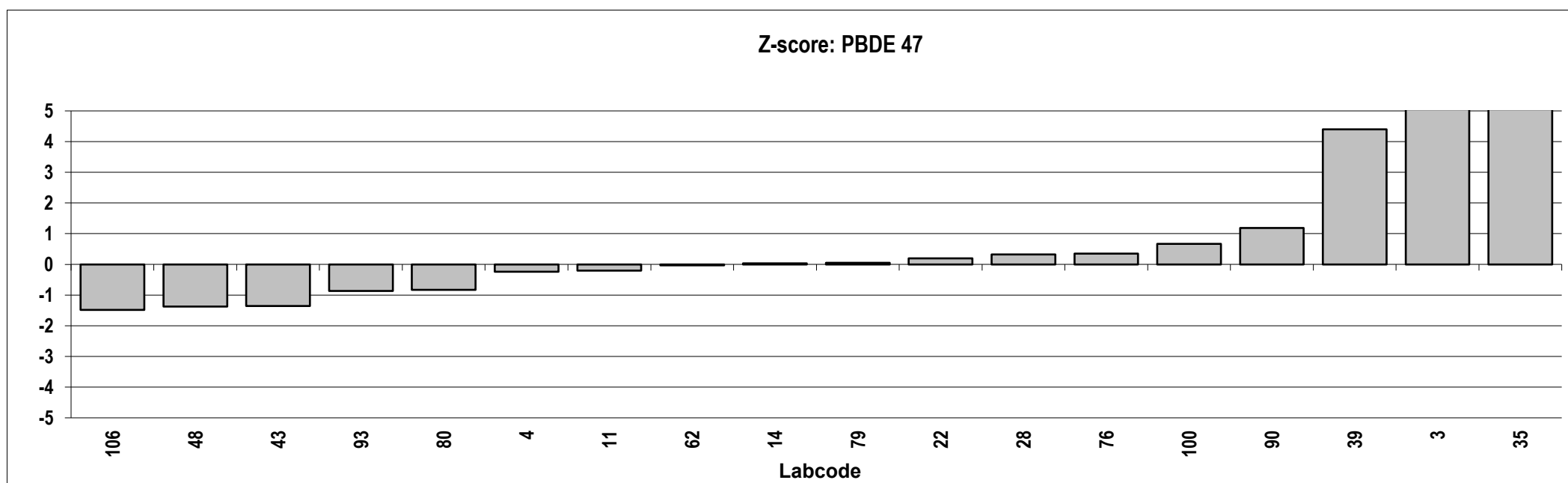
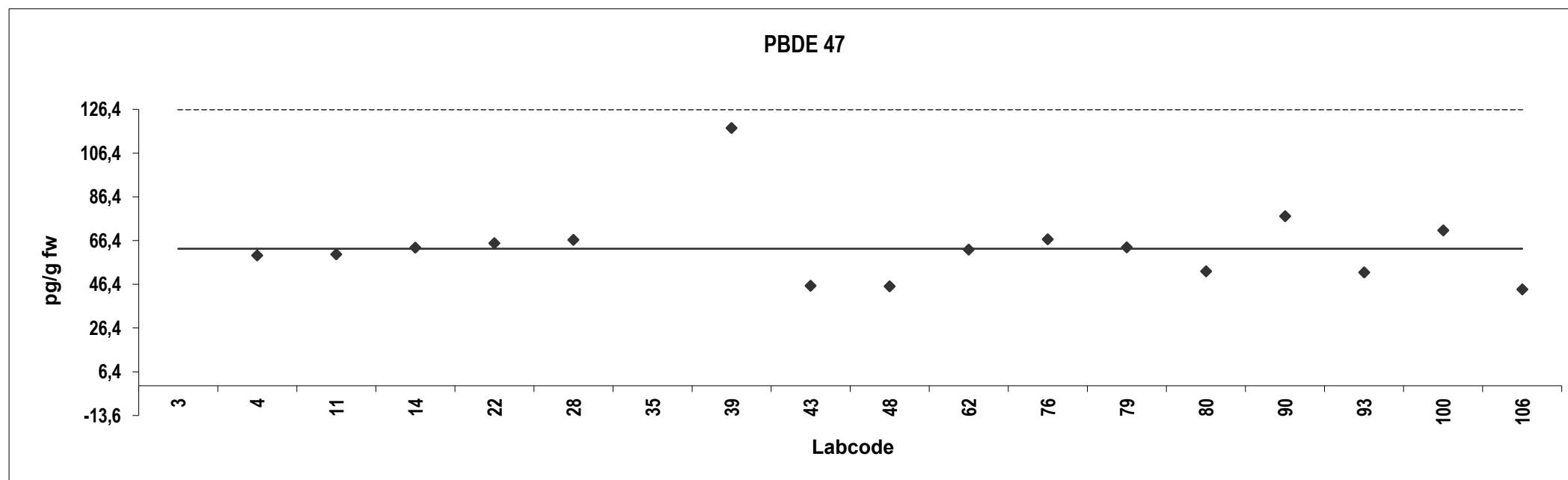
Consensus statistics	
Consensus median, pg/g	3,6
Median all values pg/g	3,7
Consensus mean, pg/g	3,9
Standard deviation, pg/g	1,1
Relative standard deviation, %	27
No. of values reported	18
No. of values removed	1
No. of reported non-detects	0



Salmon
Congener: PBDE 47

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	127	5,1	Outlier			
4	60	-0,24				
11	60	-0,20				
14	63	0,036				
22	65	0,20				
28	67	0,32				
35	143	6,4	Outlier			
39	118	4,4				
43	46	-1,4				
48	45	-1,4				
62	62	-0,036				
76	67	0,35				
79	63	0,050				
80	52	-0,83				
90	78	1,2				
93	52	-0,86				
100	71	0,67				
106	44	-1,5				

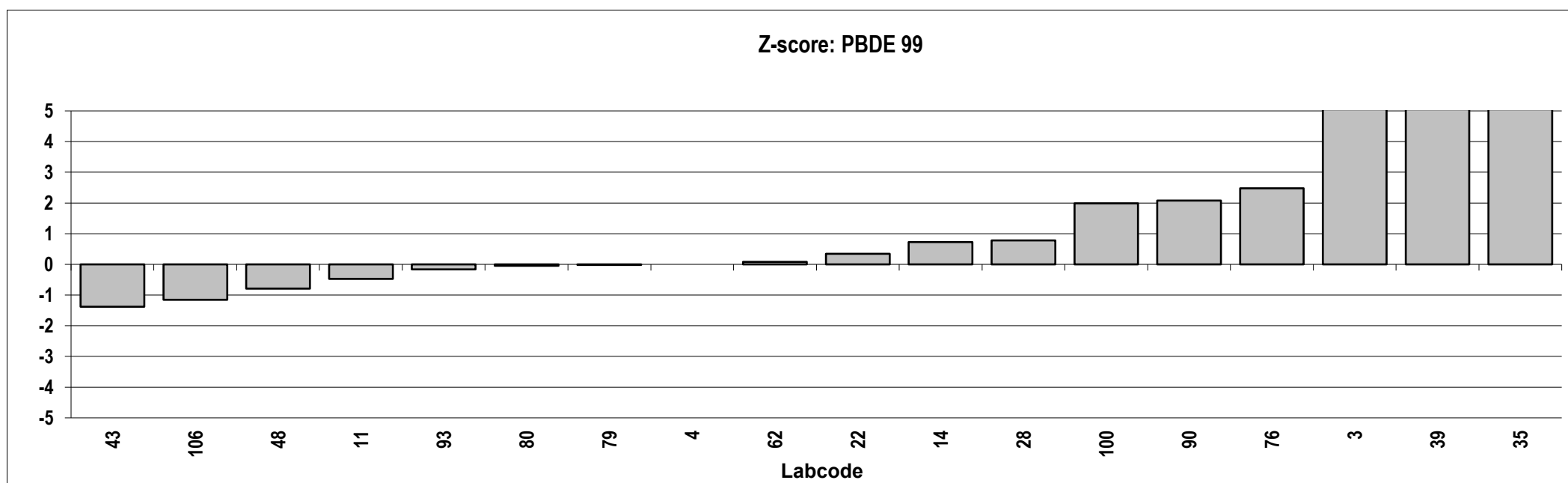
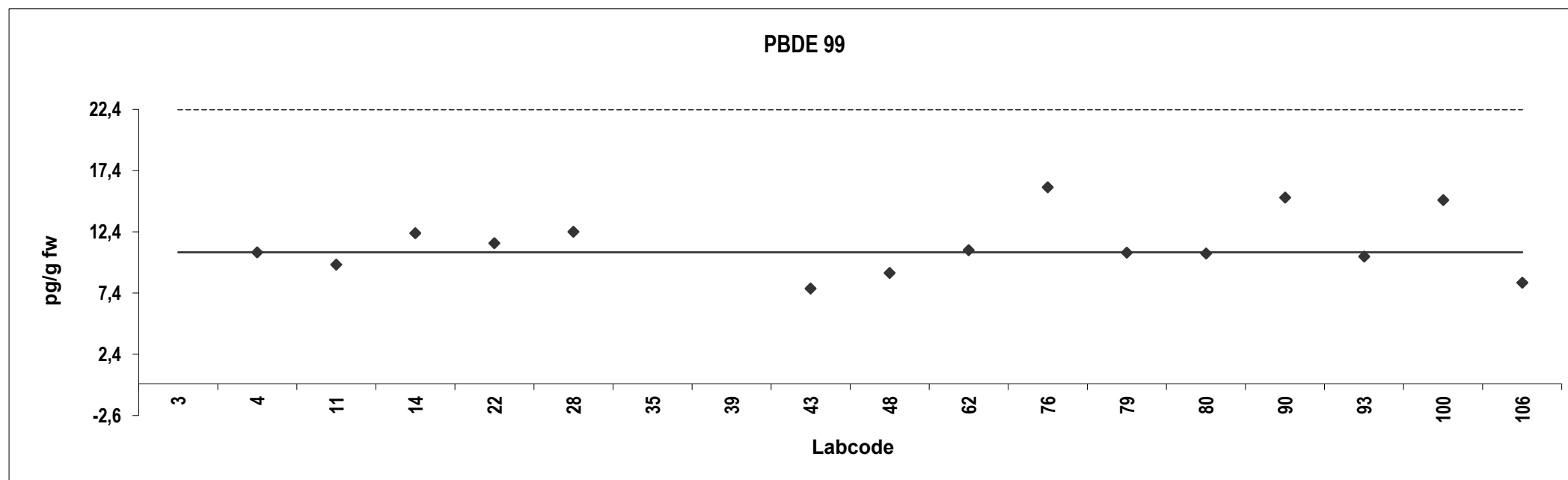
Consensus statistics	
Consensus median, pg/g	63
Median all values pg/g	63
Consensus mean, pg/g	63
Standard deviation, pg/g	17
Relative standard deviation, %	28
No. of values reported	18
No. of values removed	2
No. of reported non-detects	0



Salmon
Congener: PBDE 99

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	23	5,5	Outlier			
4	11	0,0000				
11	9,7	-0,47				
14	12	0,73				
22	11	0,35				
28	12	0,78				
35	55	21	Outlier			
39	42	14	Outlier			
43	7,8	-1,4				
48	9,0	-0,79				
62	11	0,084				
76	16	2,5				
79	11	-0,018				
80	11	-0,042				
90	15	2,1				
93	10	-0,16				
100	15	2,0				
106	8,3	-1,2				

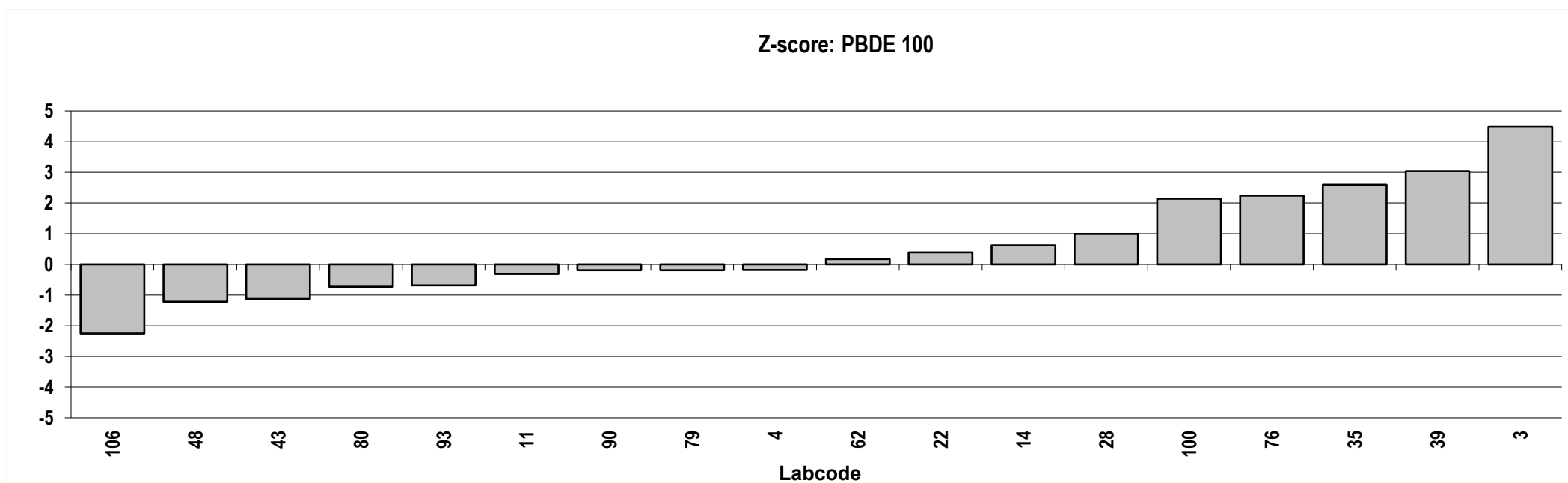
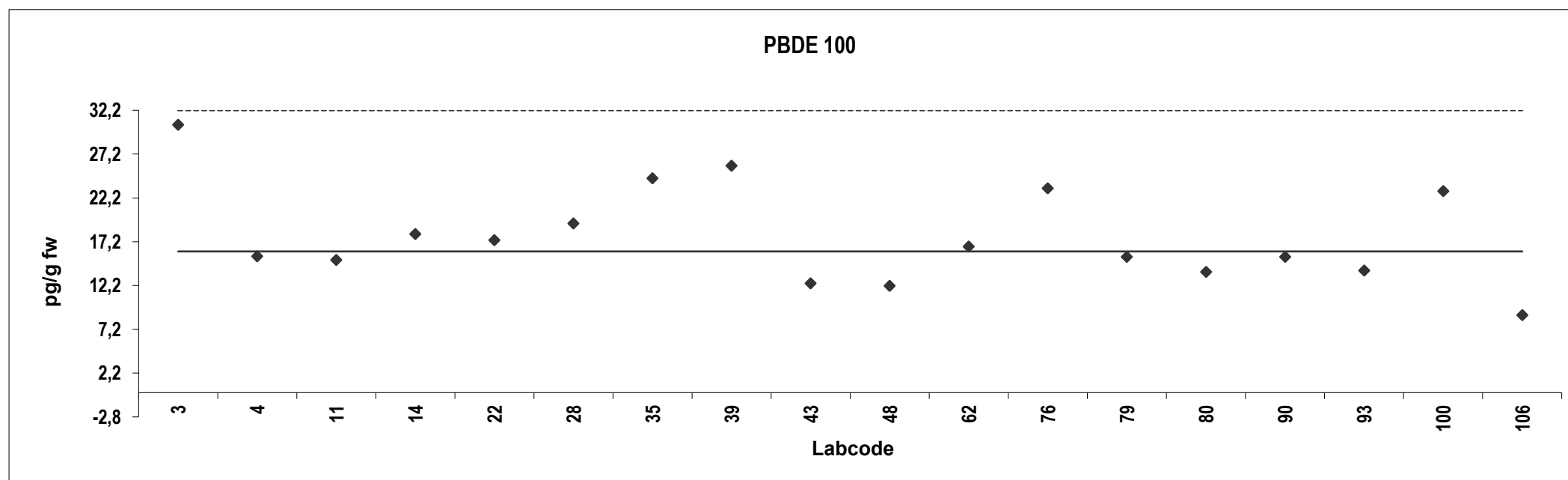
Consensus statistics	
Consensus median, pg/g	11
Median all values pg/g	11
Consensus mean, pg/g	11
Standard deviation, pg/g	2,5
Relative standard deviation, %	22
No. of values reported	18
No. of values removed	3
No. of reported non-detects	0



Salmon
Congener: PBDE 100

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	31	4,5				
4	16	-0,17				
11	15	-0,30				
14	18	0,62				
22	17	0,40				
28	19	0,99				
35	24	2,6				
39	26	3,0				
43	12	-1,1				
48	12	-1,2				
62	17	0,17				
76	23	2,2				
79	16	-0,19				
80	14	-0,72				
90	16	-0,19				
93	14	-0,68				
100	23	2,1				
106	8,8	-2,3				

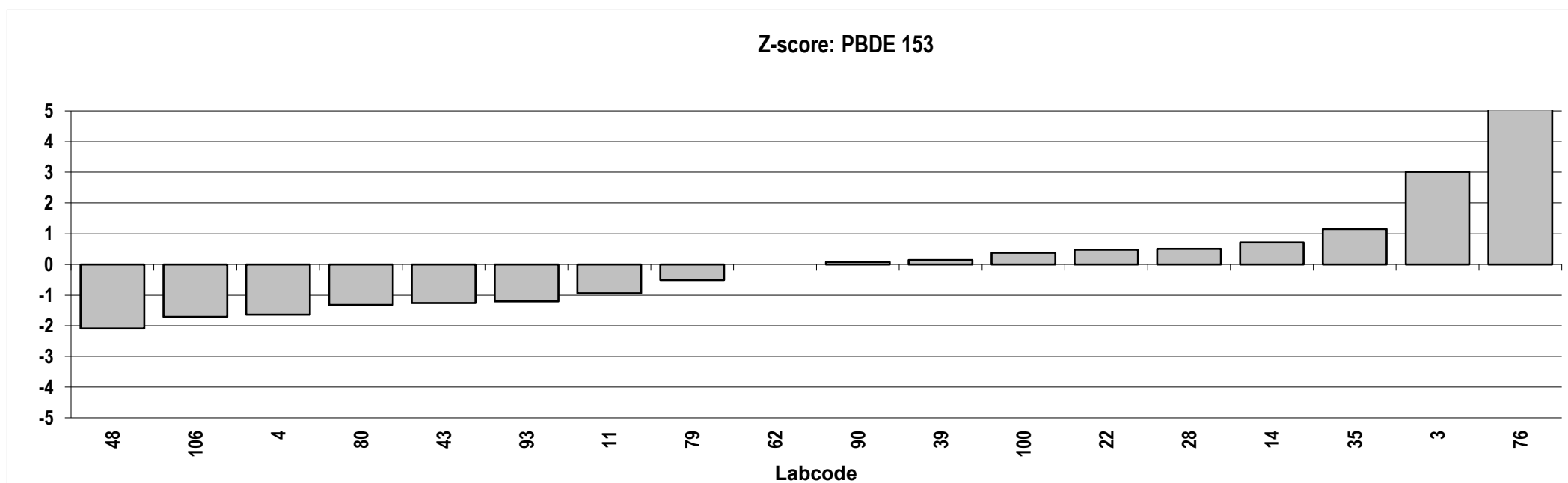
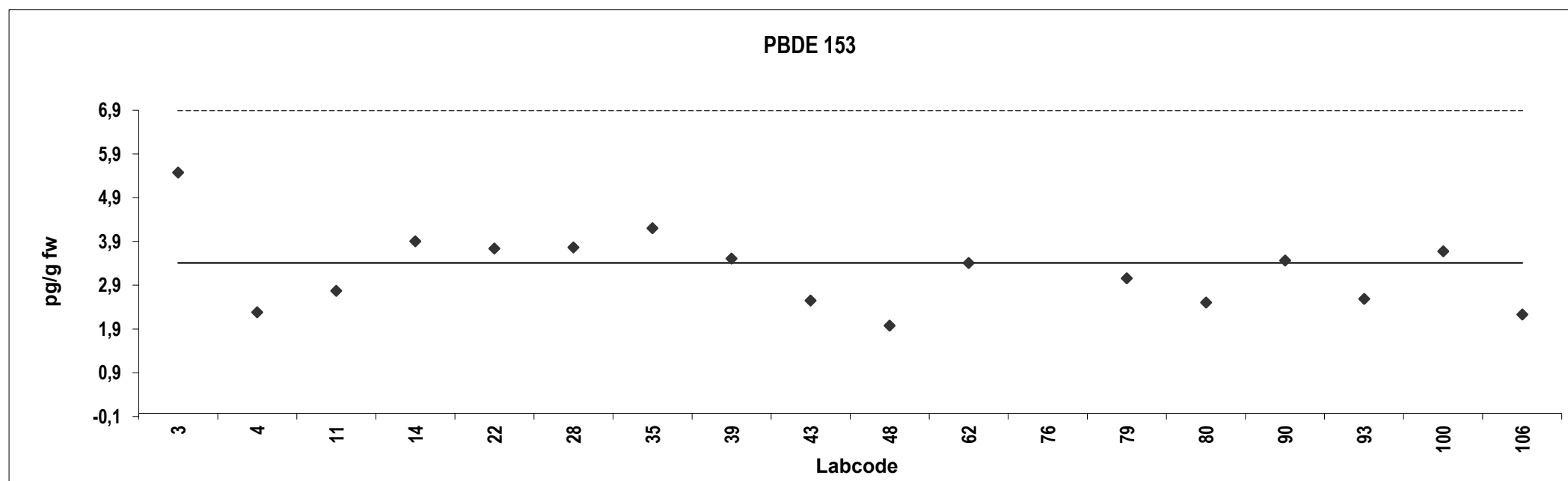
Consensus statistics	
Consensus median, pg/g	16
Median all values pg/g	16
Consensus mean, pg/g	18
Standard deviation, pg/g	5,6
Relative standard deviation, %	31
No. of values reported	18
No. of values removed	0
No. of reported non-detects	0



Salmon
Congener: PBDE 153

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	5,5	3,0	Outlier			
4	2,3	-1,6				
11	2,8	-0,93				
14	3,9	0,72				
22	3,8	0,48				
28	3,8	0,51				
35	4,2	1,2				
39	3,5	0,14				
43	2,6	-1,3				
48	2,0	-2,1				
62	3,4	0,0000				
76	7,8	6,4				
79	3,1	-0,51				
80	2,5	-1,3				
90	3,5	0,077				
93	2,6	-1,2				
100	3,7	0,38				
106	2,3	-1,7				

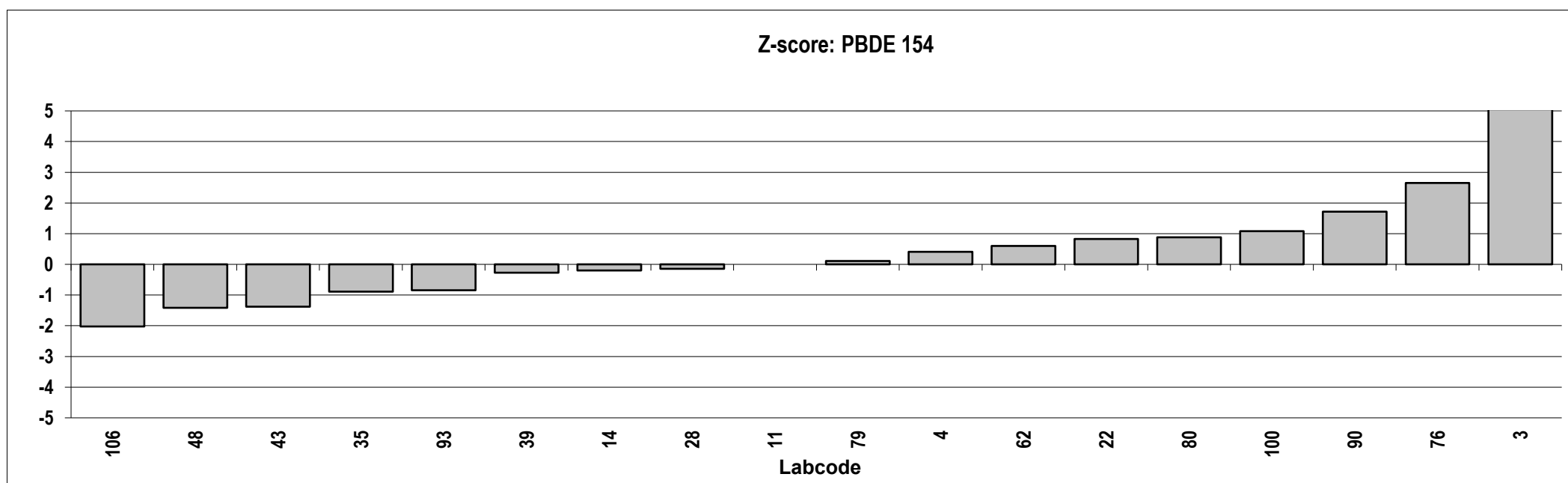
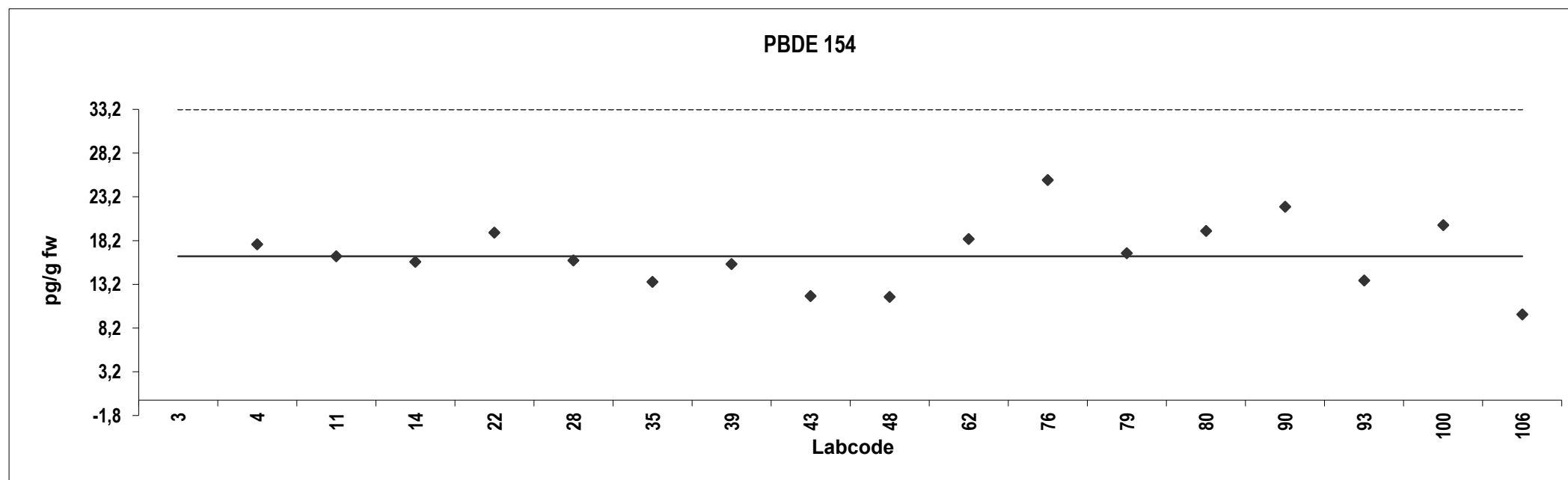
Consensus statistics	
Consensus median, pg/g	3,4
Median all values pg/g	3,5
Consensus mean, pg/g	3,3
Standard deviation, pg/g	0,88
Relative standard deviation, %	27
No. of values reported	18
No. of values removed	1
No. of reported non-detects	0



Salmon
Congener: PBDE 154

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	34	5,3	Outlier			
4	18	0,41				
11	16	0,0000				
14	16	-0,20				
22	19	0,82				
28	16	-0,15				
35	14	-0,89				
39	16	-0,27				
43	12	-1,4				
48	12	-1,4				
62	18	0,60				
76	25	2,7				
79	17	0,11				
80	19	0,88				
90	22	1,7				
93	14	-0,85				
100	20	1,1				
106	9,8	-2,0				

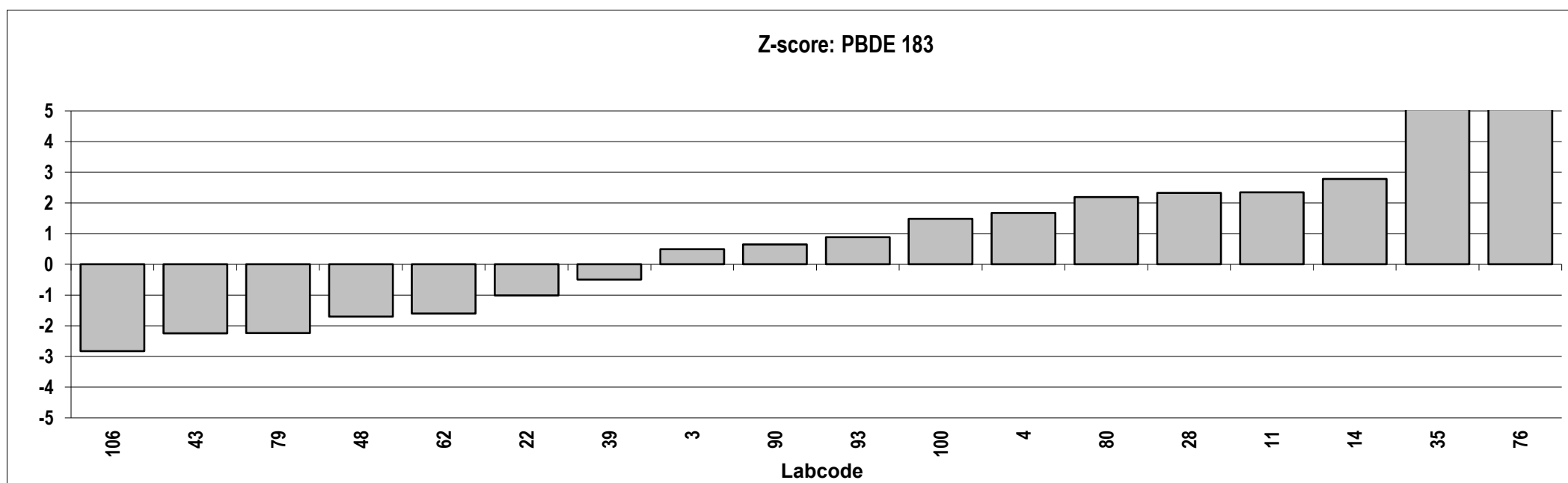
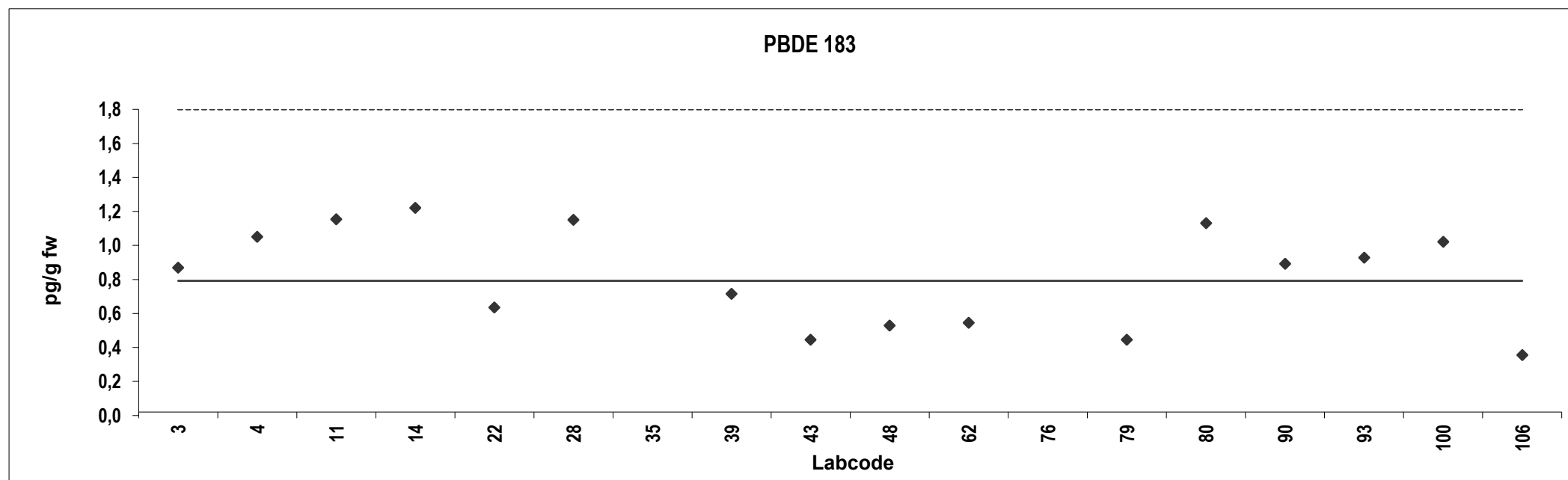
Consensus statistics	
Consensus median, pg/g	16
Median all values pg/g	17
Consensus mean, pg/g	17
Standard deviation, pg/g	3,9
Relative standard deviation, %	24
No. of values reported	18
No. of values removed	1
No. of reported non-detects	0



Salmon
Congener: PBDE 183

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	0,85	0,50				
4	1,0	1,7				
11	1,1	2,3	ND			
14	1,2	2,8	ND			
22	0,62	-1,0				
28	1,1	2,3				
35	1,8	6,6	Outlier			
39	0,69	-0,50				
43	0,42	-2,2				
48	0,51	-1,7				
62	0,52	-1,6				
76	3,0	14	Outlier			
79	0,43	-2,2				
80	1,1	2,2				
90	0,87	0,65				
93	0,91	0,88				
100	1,0	1,5	ND			
106	0,34	-2,8	ND			

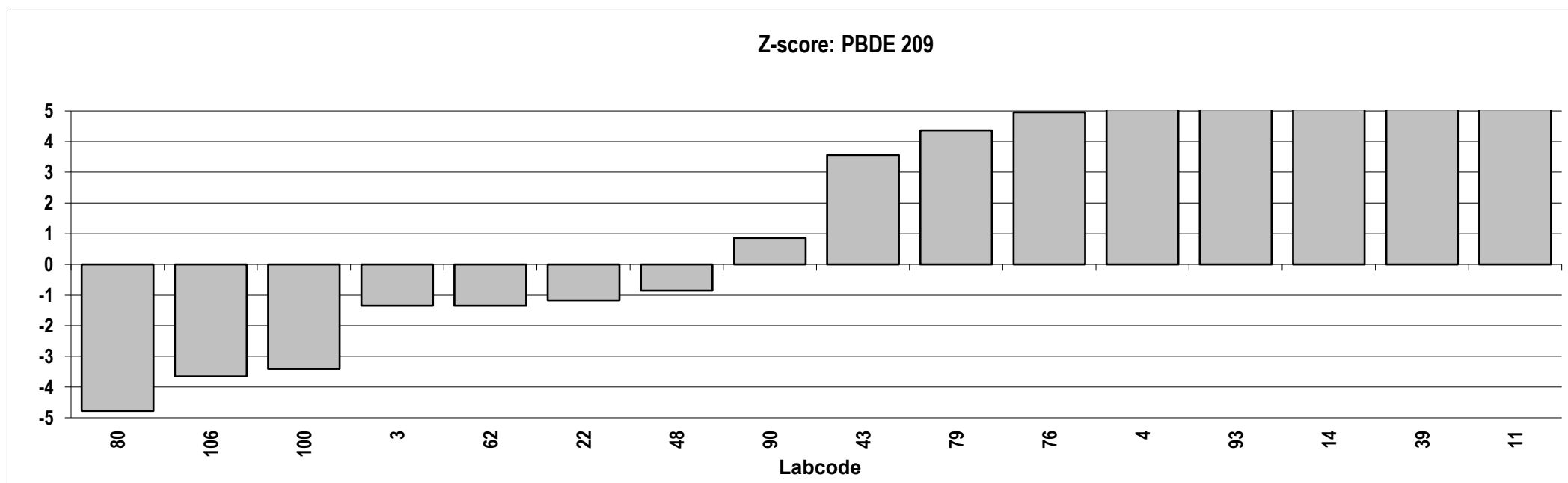
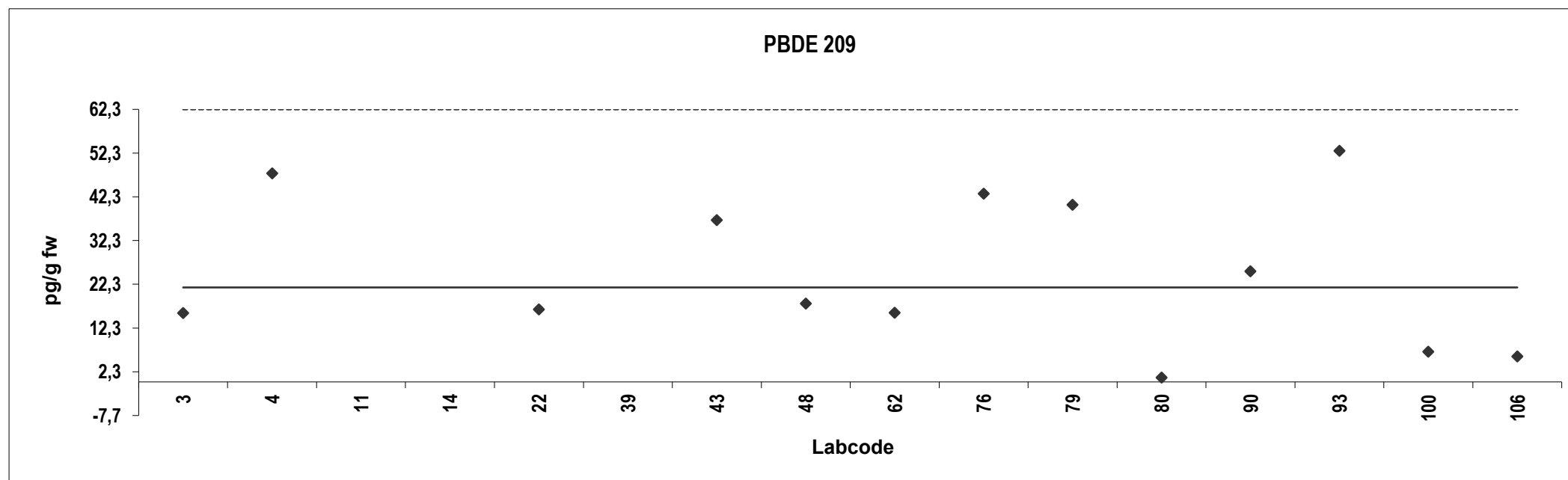
Consensus statistics	
Consensus median, pg/g	0,77
Median all values pg/g	0,89
Consensus mean, pg/g	0,80
Standard deviation, pg/g	0,29
Relative standard deviation, %	37
No. of values reported	18
No. of values removed	2
No. of reported non-detects	4



Salmon
Congener: PBDE 209

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
3	16	-1,3				
4	48	6,0				
11	379	83	Outlier			
14	68	11	Outlier			
22	17	-1,2				
39	234	49	Outlier			
43	37	3,6				
48	18	-0,86				
62	16	-1,3				
76	43	5,0				
79	40	4,4				
80	0,96	-4,8				
90	25	0,86				
93	53	7,2				
100	6,9	-3,4				
106	5,8	-3,7	ND			

Consensus statistics	
Consensus median, pg/g	22
Median all values pg/g	31
Consensus mean, pg/g	25
Standard deviation, pg/g	17
Relative standard deviation, %	69
No. of values reported	16
No. of values removed	3
No. of reported non-detects	1

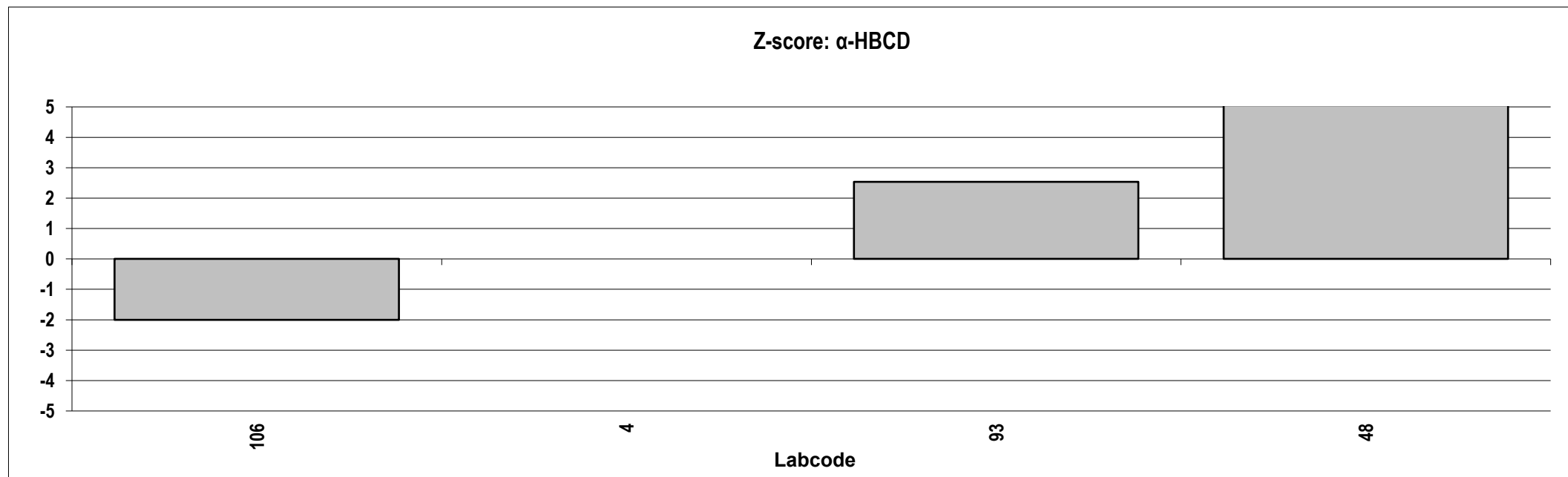
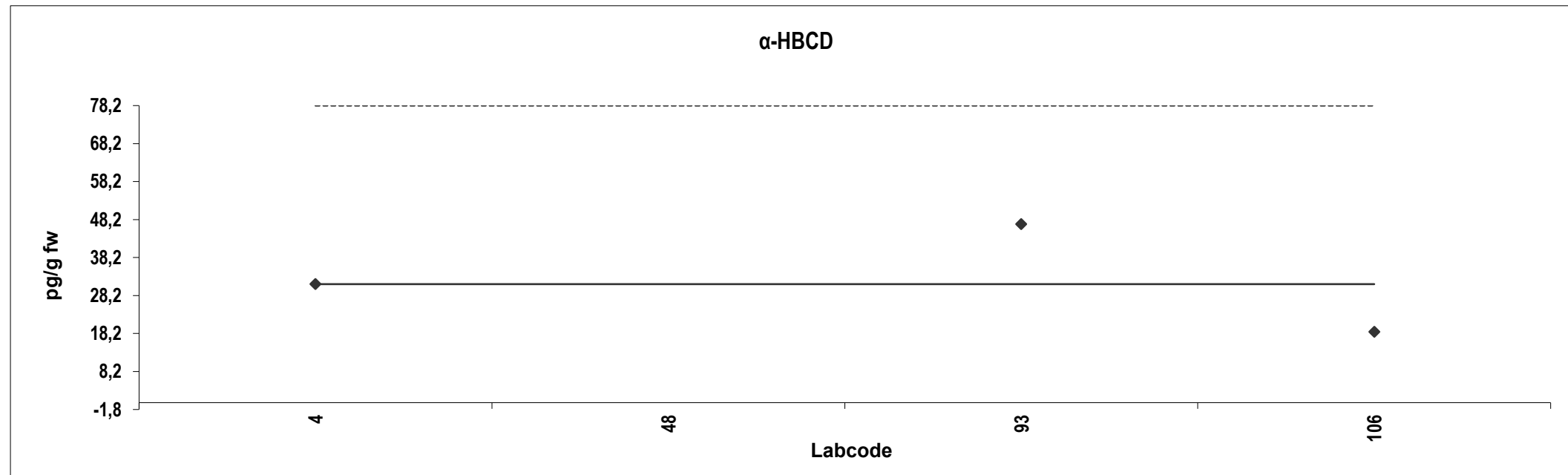


Congener: α -HBCD

[illegible]

Consensus statistics

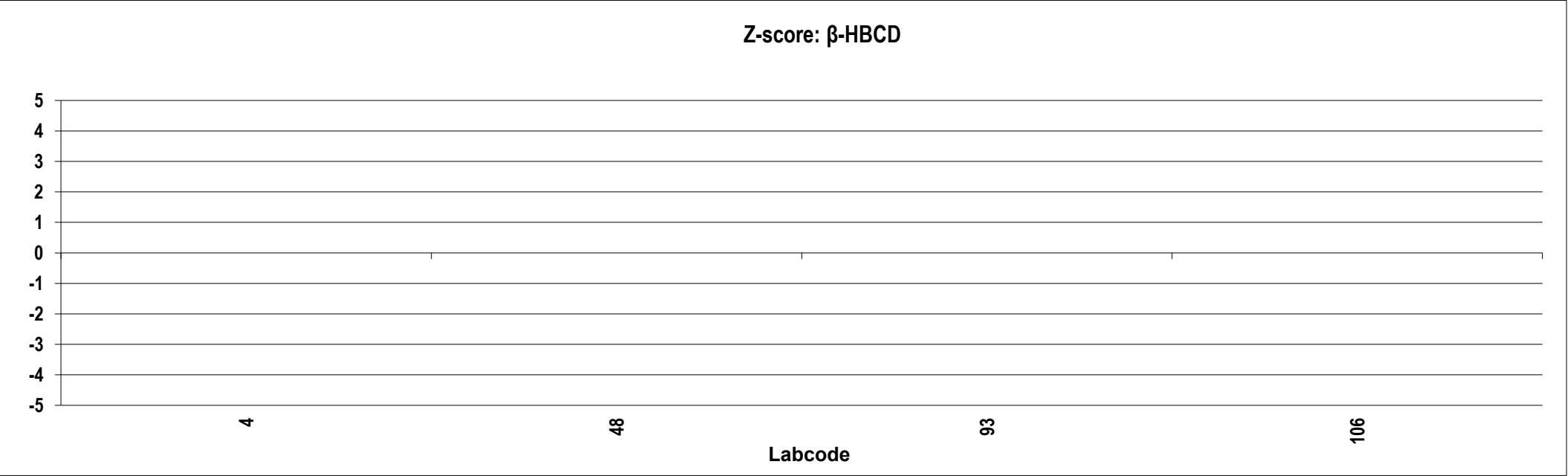
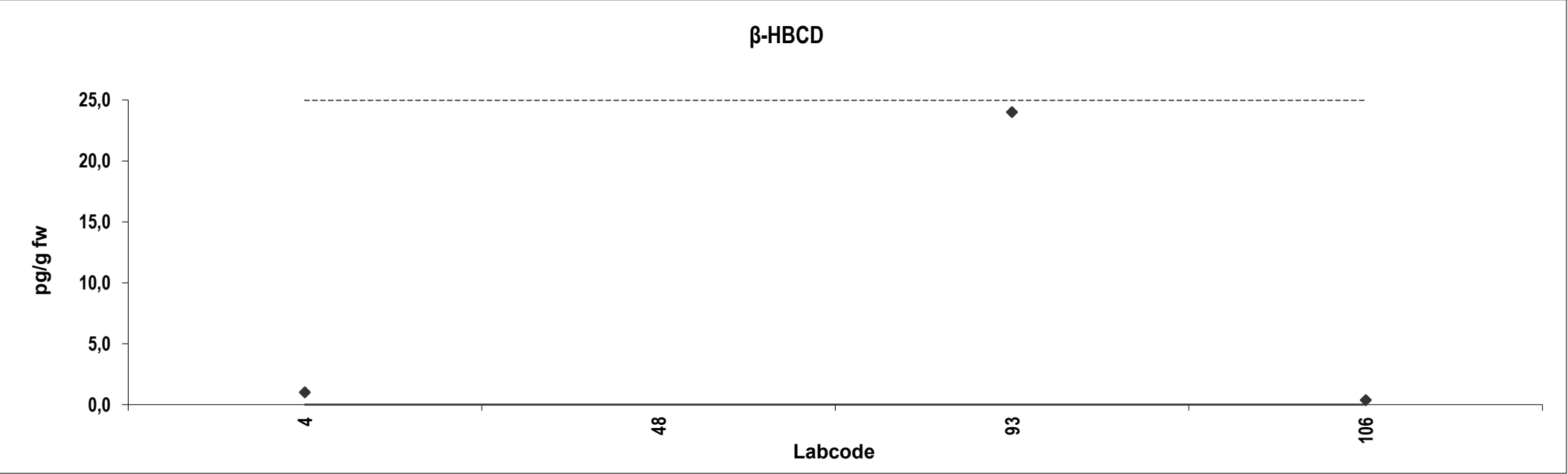
Consensus median, pg/g	31
Median all values pg/g	39
Consensus mean, pg/g	32
Standard deviation, pg/g	14
Relative standard deviation, %	44
No. of values reported	4
No. of values removed	1
No. of reported non-detects	0



Salmon
Congener: β -HBCD

Lab code	Conc. pg/g fw.	Z-score	Notes	Lab code	Conc. pg/g fw.	Notes
4	1,0	**	ND			
48	200	**	Outlier,ND			
93	24	**	ND			
106	0,37	**	ND			

Consensus statistics	
Consensus median, pg/g	**
Median all values pg/g	13
Consensus mean, pg/g	8,5
Standard deviation, pg/g	13
Relative standard deviation, %	159
No. of values reported	4
No. of values removed	1
No. of reported non-detects	4

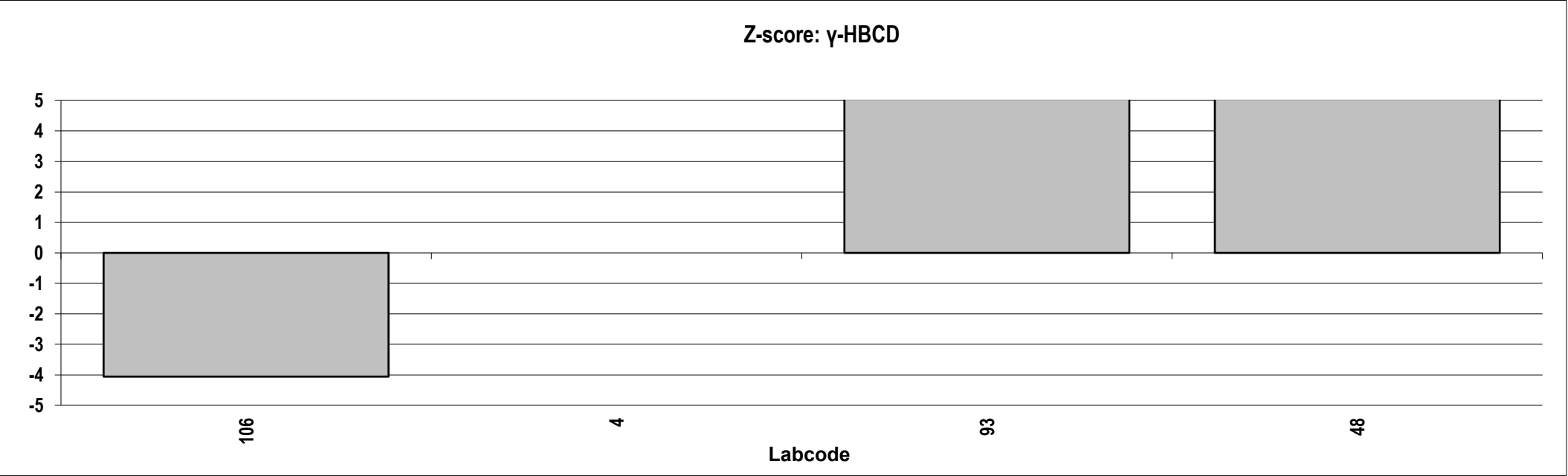
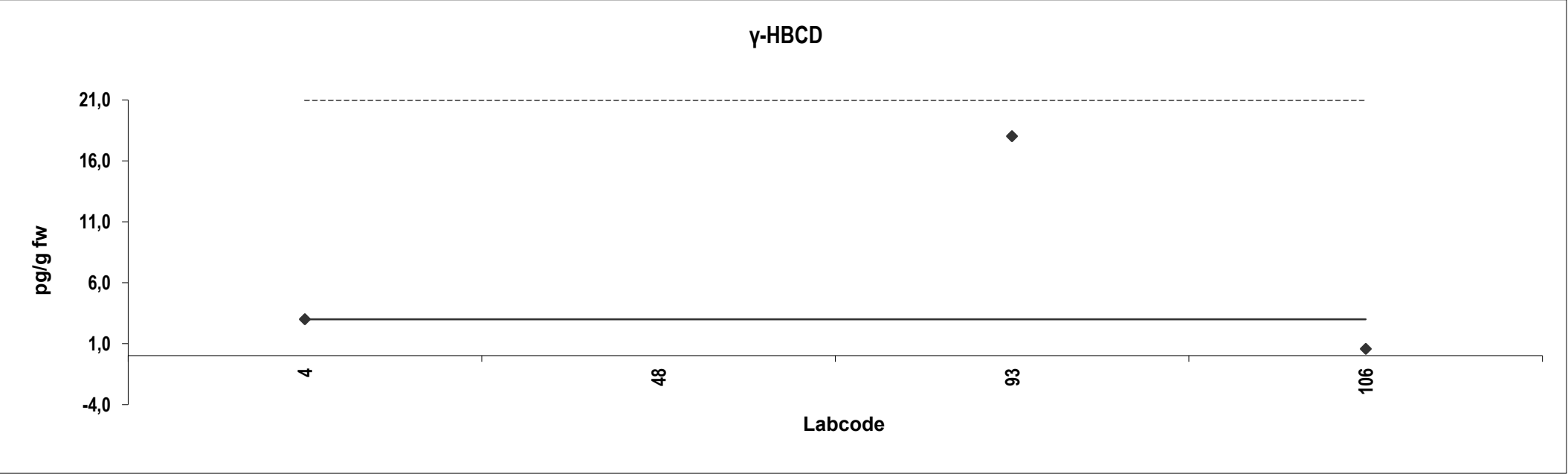


Congener: γ -HBCD

Lab code	Conc. pg/g fw.	Z-score	Notes
4 48 93 106	3,0 200 18 0,56	0,0000 331 25 -4,1	Outlier,ND ND ND

Consensus statistics

Consensus median, pg/g	3,0
Median all values pg/g	10
Consensus mean, pg/g	7,2
Standard deviation, pg/g	9,4
Relative standard deviation, %	132
No. of values reported	4
No. of values removed	1
No. of reported non-detects	3



Salmon

Congener: total HBCD

[illegible]

