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**REPORTS ON THE RESULTS OF THE SEVENTH INTERCALIBRATION
EXERCISE ON TRACE METALS IN BIOTA (PART 1)**

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**ICES SEVENTH ROUND INTERCALIBRATION FOR TRACE METALS
IN BIOLOGICAL TISSUES
ICES 7/TM/BT (PART 1)**

by

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ABSTRACT

Three marine biological tissue samples (lobster hepatopancreas, scallops and plaice) were distributed to 57 ICES laboratories, 51 of which submitted results.

Values were collated for the six metals, copper, zinc, arsenic, cadmium, mercury and lead. Most laboratories perform well for copper, zinc and cadmium, although there is an indication of a possible matrix problem for copper in plaice. Most of those that analyze for mercury are able to do so even at low (0.06 mg/kg) concentrations. Many laboratories have problems with lead, even at the 5 mg/kg level, and with low level cadmium (0.06 mg/kg).

Only about 40 percent of the respondents analyzed the materials for As, an element which appears to present problems. The distribution of results is poor (21% RSD) at the 25 mg/kg level and deteriorates further (35% RSD) at 5 mg/kg.

Results for chromium, manganese, iron, cobalt, nickel and selenium are also tabulated.

RÉSUMÉ

On a distribué trois échantillons de tissus biologiques marins (hépatopancréas de homard, pétoncles et plie) à 57 laboratoires de CIEM, dont 51 ont soumis des résultats.

Des valeurs ont été compilées pour les 6 métaux, cuivre, zinc, arsenic, cadmium, mercure et plomb. La plupart des laboratoires réussissent bien pour le zinc, le cuivre et le cadmium, bien que l'on décèle une problème de matrice pour le cuivre dans la plie. La plupart de ceux qui analysent pour le mercure peuvent le faire même à de basses concentrations (0.06 mg/kg). Plusieurs laboratoires ont des difficultés avec le plomb, même à 5 mg/kg, et avec le cadmium à 0.06 mg/kg.

Seulment 40% des répondants ont déterminé l'arsenic dans les échantillons. Cet élément semble présenter des problèmes. La distribution des résultats est pauvre (21% CV) à 25 mg/kg et se détériore encore plus (35% CV) à 5 mg/kg.

On donne également des résultats pour le chrome, le manganèse, le fer, le cobalt, le nickel et le sélénium.

NOTE

This report differs but little from the preliminary report (Berman, 1984). An arithmetical error by Lab 9 for zinc has been corrected. Lab 13's results for chromium, cobalt and selenium were wrongly interpreted. This has resulted in the recalculation of the means for these metals (pages 20 to 21). Errors were corrected in the calculations of the means for nickel (page 21). Tables II to XIV have been revised to reflect the number of data sets incorporated in the means. A set of charts indicating laboratory precision and bias for each sample for each of the six core metals has been appended. Some typographical and other minor changes have been made.

INTRODUCTION

An ongoing series of ICES intercomparison exercises has been conducted since 1971 in order to examine results of analytical procedures used for the measurement of trace metals in marine biological tissues. The participants in these exercises were laboratories associated with the ICES fish and shellfish baseline and monitoring programs, those nominated by the Joint Monitoring Group of the Oslo/Paris Commissions, as well as others that had expressed an interest in participating in ICES projects. A gradual improvement in the level of agreement between laboratories for the trace metals copper, zinc and mercury was seen over the years, whereas little or no improvement was noted for cadmium and lead (Topping and Holden, 1978; Holden and Topping, 1981) by the end of the fifth exercise in 1979.

The sixth ICES intercomparison study in 1980 was concerned only with cadmium and lead. It was concluded from this study that most laboratories could produce reliable cadmium and lead values at the 1 and 2 mg/kg levels respectively, but only a small minority could analyze for lead at about 0.5 mg/kg (Topping, 1982). As a result, the ICES Marine Chemistry Working Group (MCWG) in 1981 appointed a sub-group whose task was to identify and try to resolve problems concerning the determination of lead in biological tissues. This sub-group confirmed the presence of matrix problems in the analysis using graphite furnace atomic absorption spectrometry and recommended procedures which should enable confident analyses at the greater than 1 mg/kg level (Harms *et al.*, 1982). Harms has recently expanded this work to sub mg/kg levels (Harms, 1985).

In 1982 the MCWG recommended a seventh intercomparison exercise for trace metals in biological tissues in preparation for the 1985 baseline studies of contaminant levels in marine organisms (MCWG, 1982) and in 1983 set up a scheme for a two part study whereby the first phase would be carried out in 1983 prior to the baseline study and the second simultaneously with the baseline study in 1985 (MCWG, 1983). The elements of interest were copper, zinc, arsenic, cadmium, mercury and lead.

PREPARATION OF SAMPLES

Three samples were prepared for the first part of this intercomparison study. The first (Sample A), from the National Research Council of Canada, was prepared from lobster hepatopancreas which had been homogenized, spray dried and acetone extracted to remove the majority of the lipids. The second (Sample B) was freeze dried, ground scallops adductor muscle. The third (Sample C), prepared by U. Harms, Bundesforschungsanstalt für Fischerei, Federal Republic of Germany, was freeze dried plaice. It was hoped that these three materials would provide an ample range of metal concentrations as well as a variety of fish and shellfish matrices.

The materials were thoroughly blended and bottled in 60-ml bottles in quantities of 15, 15 and 20 g respectively. Randomly selected bottles were analyzed at the National Research Council of Canada (Laboratory 49) to assess the interbottle homogeneity. The results of these tests are listed with all the other laboratory data on pages 29 to 52. The difference between the data of Lab 49 and all others is that the individual values in each set of results from this laboratory are from samples taken from different bottles. The materials appear to be homogeneous with respect to copper, zinc, arsenic, cadmium and mercury, but there may be problems with lead in Samples B and C.

SAMPLE DISTRIBUTION AND RECEIPT OF RESULTS

The samples were sent by air post in the latter part of August 1983 to the 57 laboratories that had indicated a desire to participate in this intercomparison exercise (Table XVII, pages 25 to 28). Six replicate analyses of each material for each of the six metals of interest was requested along with data regarding the analytical procedures used in the analyses. Results were requested on a "dry weight" basis. The deadline for the receipt of results was set at December 15, 1983.

Three recipients reported that their samples may have been opened by customs officials prior to delivery. One set had to be replaced. A reminder of the deadline was sent on December 7.

Over 40 sets of results were received by the end of December. It then became apparent that the Christmas mail and other post office problems were delaying the receipt of results. (One set mailed in late December did not arrive until early in March.) A number was assigned to each of the respondents (Table XVII). The final number of respondents was fifty-one, but the results from Labs 50 and 51 arrived too late to be included in the calculated data. These results are listed, however, with all the rest of the data.

RESULTS

It is pleasing to note that a good majority of the respondents supplied most of the data requested of them. These included sample drying techniques, decomposition and dissolution procedures, analytical methodologies, calibration techniques and limits of detection for the analyte elements.

All results, except a few very obvious outliers in some sets, were tabulated (pages 29 to 52). The number of significant figures in submitted data was reduced in many cases to a maximum of three for tabulation and computation.

1. "DRY WEIGHT"

There appear to be almost as many procedures for "drying" a biological tissue sample as laboratories analyzing the materials. Happily, most methods seem to yield similar results. The percent weight loss reported by each laboratory for the three sample and the method used by each are tabulated in Table I on page 4. A *t* test at the 95 percent confidence level was applied to each set of results and the means for each sample were calculated after the successive rejection of the outliers which are indicated by ** in Table I. The final means and standard deviations are listed in Table II below. The final columns in each of the tables indicate the number of laboratories contributing to the mean and the number rejected before calculation of the final mean.

It is obvious that for these materials that simply drying the samples for a few hours at temperatures between 100 and 115 degrees Celsius is apparently sufficient in order to reach comparable sample weights. Lower temperatures, even for prolonged periods, may not be sufficient, and higher temperatures may cause excessive weight losses. The use of dessicants without vacuum appears to lead to lower than average weight losses.

TABLE II
MEAN WEIGHT LOSSES - percent

Sample A	3.96	± 0.86	(22%)	31, 9
Sample B	3.83	± 0.83	(22%)	31, 8
Sample C	8.81	± 0.78	(?)	36, 5

TABLE I

WEIGHT LOSS - percent

<u>Lab</u>	<u>A</u>	<u>B</u>	<u>C</u>	<u>Procedure</u>
1	4.80	5.80*	2.90*	110 C for 18 hours
2	4.40	4.96	9.01	105 C
3	3.95	4.18	8.17	100-110 C for 4 hours
4	3.48	3.38	7.73	100-105 for 24 hours
5	3.30	3.20	8.60	No data
6	----	----	9.10	105 C for 2 hours
7	3.04	2.97	8.12	105 C for 2 hours
8	4.70	4.29	8.93	105 C for 24 hours
9	4.94	4.40	9.44	100 C for 23 hours
10	3.90	2.77	9.23	Vacuum dessicant for 48 hours
11	4.35	3.78	9.15	110 C for 3 hours
12	3.48	3.70	8.31	95 C for 12 hours
13	3.00	2.50	7.50	80 C for 24 hours
14	2.70	2.70	6.70*	105 C for 5 hours
15	----	----	----	Dessicant for 3 months
16	7.92*	8.92*	9.39	120 C for 3 days
17	4.17	4.00	8.17	105 C overnight
18	4.00	3.90	8.92	105 C overnight
19	5.59	4.56	9.09	105 C for 16 hours
20	2.01	1.10	6.24	65 C for 12 hours
21	----	----	----	Dessicant
22	4.67	2.93	8.96	? C for 2 hours
23	4.45	5.01	7.68	105-110 C for 18 hours
24	----	----	----	No data
25	3.00	1.80*	7.00*	70 C for 6 hours
26	3.04	3.71	7.34	75 C for 15 hours
27	3.92	4.12	9.01	105 C for 4 hours
28	4.56	3.53	9.35	105 C for 16 hours
29	6.39*	6.03*	9.82	105 C for 5 hours
30	----	----	----	Freeze dried
31	4.90	4.77	8.70	103 C for 1 hour
32	7.50*	9.00*	10.0	105 C for 2 hours
33	4.22	4.05	8.69	No method
34	5.60	4.90	8.50	100-105 C for 24 hours
35	----	----	----	60 C for 16 hours
36	10.3*	3.72	9.91	105 C for 2 hours
37	----	----	----	Not dried
38	6.29*	5.32	10.2	150 C for 5 days
39	2.54	----	7.49	105 C
40	4.84	4.51	9.39	100 C for 24 or 48 hours
41	3.18	3.18	8.68	Mg(ClO ₄) ₂ for 72 hours
42	2.73	2.73	6.57*	No method
43	8.80*	4.87	9.53	105 C to constant weight
44	9.07*	7.62*	9.90	130 C for 48 hours
45	7.86*	1.57*	7.50	Lyophilized for 24 hours
46	----	----	----	No data
47	----	----	----	Silica gel for 3 months
48	1.76*	1.00*	6.59*	Silica gel for 48 hours
49	----	----	----	No drying data
50	3.27	2.34	8.87	Silica gel for 48 hours
51	----	----	----	No drying data

---- No data available

* Rejected by t test

The samples were all bottled in an area with reduced humidity and each of the sample sets should have the same original moisture content. However, these materials will pick up moisture from the atmosphere, and if weighings are not made soon after the bottles are opened the weight loss on drying will increase. This may explain some of the high weight loss results noted in Table I on page 4.

For the few laboratories that did not report their results on a dry weight basis, their values were factored by the mean weight losses shown in Table II to make their analytical results comparable to the others.

2. METALS

An evaluation has been made for the results for the six core metals of this study (copper, zinc, arsenic, cadmium, mercury and lead). The data has also been tabulated for those elements where five or more sets of results were received (chromium, manganese, iron, cobalt, nickel and selenium).

An attempt has been made to represent all the data received for the six metals on the individual graphs for each sample:

The range of results from each laboratory and their mean is plotted. A number of laboratories submitted more than one set of results for various elements in the samples. These have been plotted sequentially and the term "labs" in this section refers to the number of sets of results, not the number of individual participating laboratories.

A range continued by a forward arrow → beyond the right margin of the graph indicates that the range exceeds this boundary.

A forward arrow → beginning at the right margin indicates that all values submitted exceed the boundary.

A backward arrow ← at the right margin indicates results less than a limit of detection beyond the boundary.

A backward arrow ← within the graph indicates results less than the marked limit of detection.

A backward arrow ← from the left margin indicates results less than the boundary value or a limit of detection which is less than this boundary.

Means marked by a "■" are those that have been retained after a rejection test and are incorporated into the overall result. Means marked by a "+" were rejected.

The results of the various calculations are beside each graph. The total range for all values reported is presented. A mean has been calculated for all quantitative results.

A *t* test at the 95 percent confidence level was applied to the means of all the values submitted. Means were successively rejected until a homogeneous set of results was obtained. This, of course, implies a normal distribution of results, which may not be a valid assumption. An overall mean, standard deviation and relative standard deviation (RSD) were calculated for the remaining values. This mean and standard deviation are plotted on each graph. The small discrepancy sometimes apparent in the RSD is the result of rounding off significant figures in reporting the mean and standard deviation.

The final means are the consensus values for the concentrations of the six metals in the three samples for this exercise. And, indeed, they may be good estimates of the real concentrations. There is enough variety of methodologies for all the metals, except possibly for arsenic, to come to this conclusion. Also, Laboratory 49 attempted to analyze each sample by three independent methods including isotope dilution mass spectrometry (except for arsenic which is a monoisotopic element). The means of the results from this laboratory, which are the top three values plotted on each graph, are never far from the consensus values.

The means of the results for the six metals in Samples B and C have been plotted against each other for those labs that submitted results for both samples. This type of plot, if the concentrations of the metals are not too different, can indicate those laboratories which consistently produce lower or higher values than the overall mean. Also, those that produce inconsistent results are readily obvious.

Copper

Summaries of the results for copper are listed beside the graphs on page 7 and the overall final means and standard deviations are shown in Table III below.

The results for Sample A are quite satisfactory with only four sets of results rejected by the *t* test. However, even at this relatively high concentration of copper there are sets of values with unacceptably high scatter (above 20 percent RSD).

The response for Sample B is good. There are seven rejected sets, most of which are higher than the mean, indicating possible contamination problems.

TABLE III

COPPER CONCENTRATIONS - mg/kg

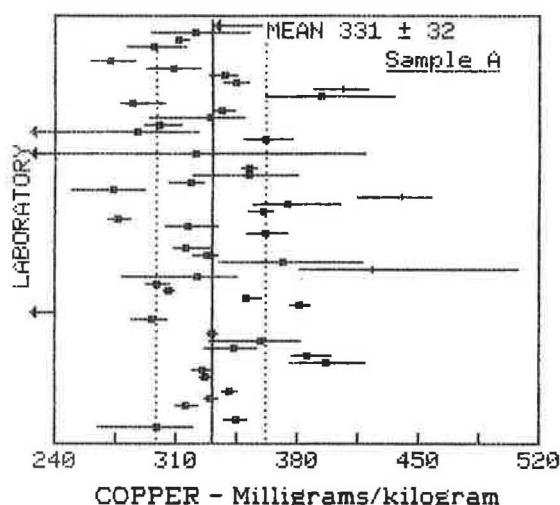
Sample A	331	± 32	(10%)	45, 4
Sample B	3.7	± 0.6	(16%)	41, 7
Sample C	3.1	± 0.8	(27%)	42, 7

The copper results for Sample C are rather puzzling. The concentrations of copper in Samples B and C are almost identical, yet the scatter of results between laboratories is significantly higher for Sample C. There is no evidence of interbottle inhomogeneity suggesting that the problem may be one of the plaice matrix itself. Seven sets of results were rejected, all of which were higher than the mean. The relative standard deviation of 27 percent for this sample is high, indicating a factor of three between the lowest and highest accepted means.

The most common method of sample decomposition is an open beaker digestion with nitric acid or a combination of nitric acid with hydrogen peroxide or perchloric acid. A third of the labs use some form of closed vessel acid decomposition (BOMB) and about 15 percent dry ash the samples.

About half the labs used flame atomic absorption spectrometry (FAAS) and another third, graphite furnace atomic absorption spectrometry (GFAAS), in their measurement procedures. Inductively coupled plasma atomic emission spectrometry (ICP) is not yet employed by many marine laboratories and only a few seem to use electroanalytical, neutron activation or x-ray fluorescence methods.

COPPER DATA

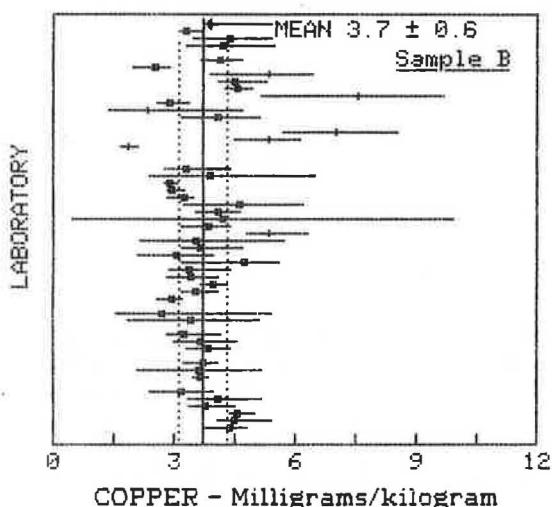


49 labs submitted results ranging from 110 to 508 mg/kg.

All labs submitted quantitative values with a mean of 333 ± 45 mg/kg.

A *t* test rejected the results of 4 of these labs.

The mean of the results of the remaining 45 labs is 331 ± 32 mg/kg. The relative standard deviation is 10 percent.

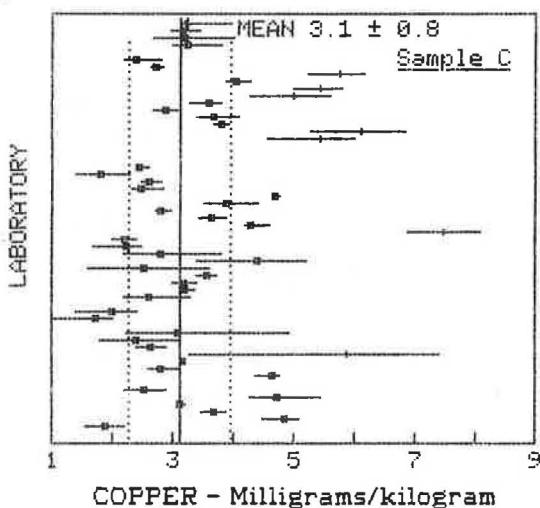


48 labs submitted results ranging from 0.48 to 9.9 mg/kg.

All labs submitted quantitative values with a mean of 3.9 ± 1.0 mg/kg.

A *t* test rejected the results of 7 of these labs.

The mean of the results of the remaining 41 labs is 3.7 ± 0.6 mg/kg. The relative standard deviation is 16 percent.



49 labs submitted results ranging from 1.0 to 8.1 mg/kg.

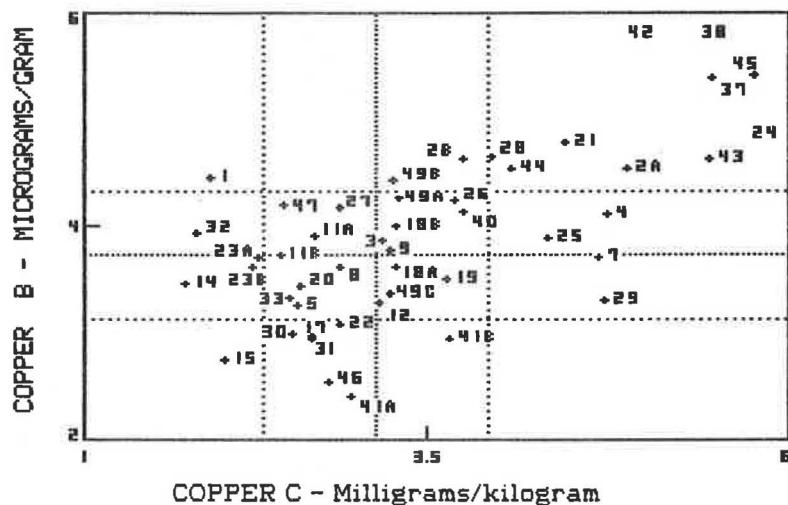
All labs submitted quantitative values with a mean of 3.5 ± 1.3 mg/kg.

A *t* test rejected the results of 7 of these labs.

The mean of the results of the remaining 42 labs is 3.1 ± 0.8 mg/kg. The relative standard deviation is 27 percent.

There is no apparent correlation between performance and methodology. Three out of four of the rejected sets for Sample A used GFAAS, an overly sensitive technique for such a high copper concentration, potentially more prone to matrix and contamination problems than FAAS. Five of the seven sets rejected for both Samples B and C were from the same laboratories.

COPPER B vs COPPER C



The plot of Sample B against Sample C shows no erratic results, but a bias towards high values on the part of some labs. Lab numbers which do not have a "+" next to them have values beyond the boundaries of the graph in the same quadrant in which the numbers are found.

The submitted copper data are listed on pages 29 to 33 along with abbreviated methodologies.

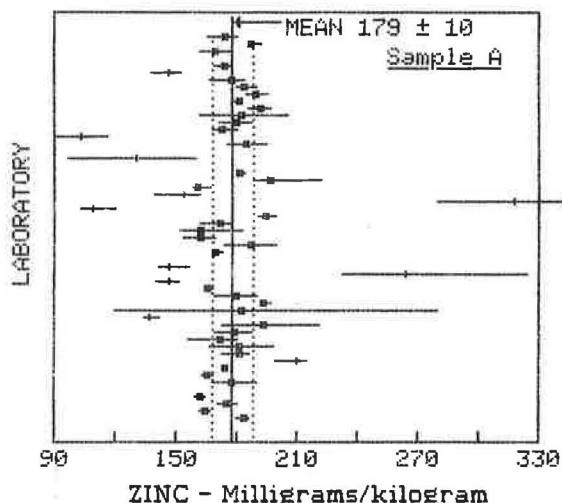
Zinc

The data for zinc are the best of all the metals. Summaries of the results are listed beside the graphs on page 9 and the overall final means and standard deviations are shown in Table IV on page 10.

Sample preparation for zinc analysis is generally identical to that for copper. Seventy percent of the labs used FAAS methods and only ten percent GFAAS. This is because of the inherently high sensitivity of zinc by atomic absorption where flame techniques are quite adequate for the zinc concentrations in these samples.

As in the case of copper, there is no apparent correlation between laboratory performance and methodology. There are eleven, eight and six sets of rejected results for the three samples respectively, but these relatively high numbers are probably due to the quite tight concurrence of the majority of the labs about the consensus values.

ZINC DATA

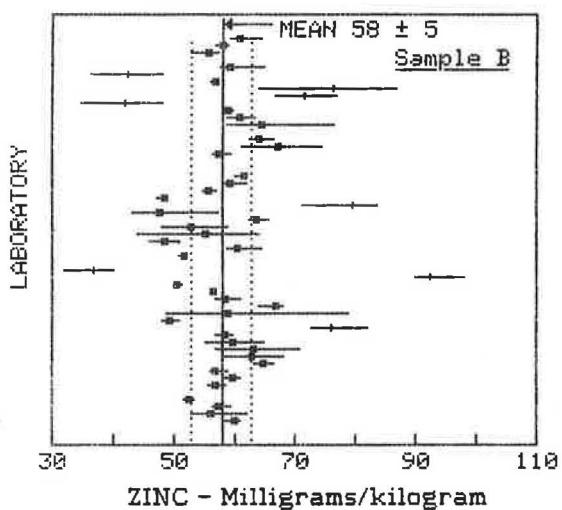


49 labs submitted results ranging from 90.9 to 335 mg/kg.

All labs submitted quantitative values with a mean of 177 ± 32 mg/kg.

A *t* test rejected the results of 11 of these labs.

The mean of the results of the remaining 38 labs is 179 ± 10 mg/kg. The relative standard deviation is 6 percent.

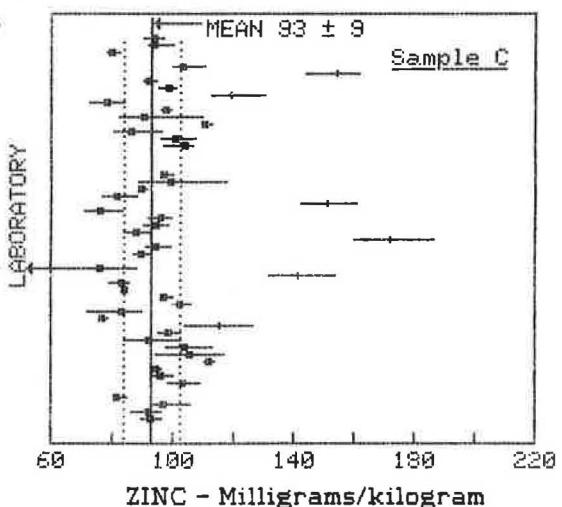


48 labs submitted results ranging from 32 to 98 mg/kg.

All labs submitted quantitative values with a mean of 58 ± 12 mg/kg.

A *t* test rejected the results of 8 of these labs.

The mean of the results of the remaining 40 labs is 58 ± 5 mg/kg. The relative standard deviation is 9 percent.



48 labs submitted results ranging from 59 to 187 mg/kg.

All labs submitted quantitative values with a mean of 98 ± 23 mg/kg.

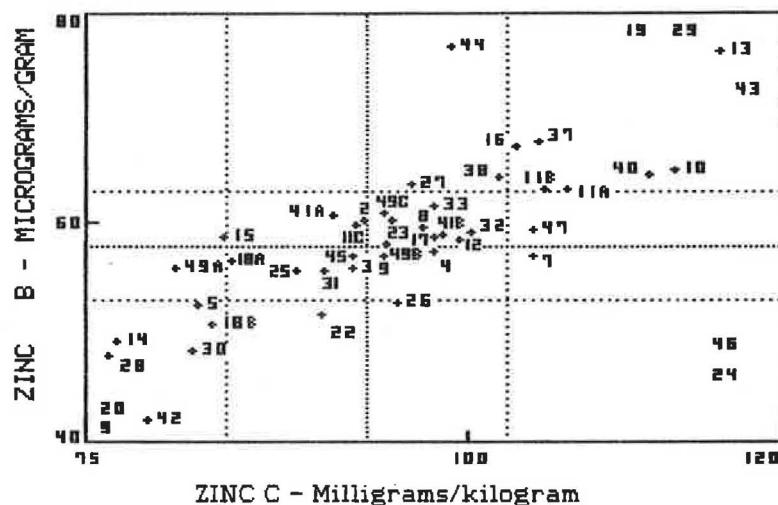
A *t* test rejected the results of 6 of these labs.

The mean of the results of the remaining 42 labs is 93 ± 9 mg/kg. The relative standard deviation is 10 percent.

TABLE IV
ZINC CONCENTRATIONS - mg/kg

Sample A	179	\pm 10	(6%)	38,11
Sample B	58	\pm 5	(9%)	40, 8
Sample C	93	\pm 9	(10%)	42, 6

ZINC B vs ZINC C



The plot of Sample B against Sample C shown above indicates only two sets of erratic results, two sets biased low and six or seven biased high. Five of the eight sets of results rejected for Sample B and of the six sets rejected for Sample C were from the same labs. However, only one of these five labs was common to the five whose copper results were rejected.

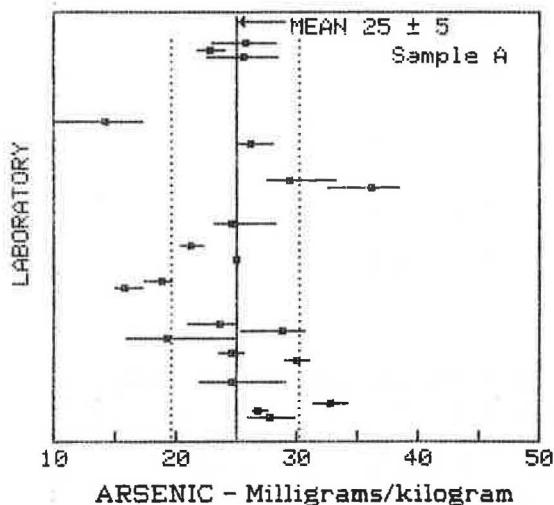
The submitted data for zinc are listed on pages 33 to 37.

Arsenic

Only forty percent of the responding laboratories submitted results for arsenic. It appears that the analysis of biological materials for arsenic is in a questionable state, second only to lead for poor interlaboratory comparisons.

Summaries of the results are listed beside the graphs on page 11 and the overall final means and standard deviations are shown in Table V on page 12.

ARSENIC DATA

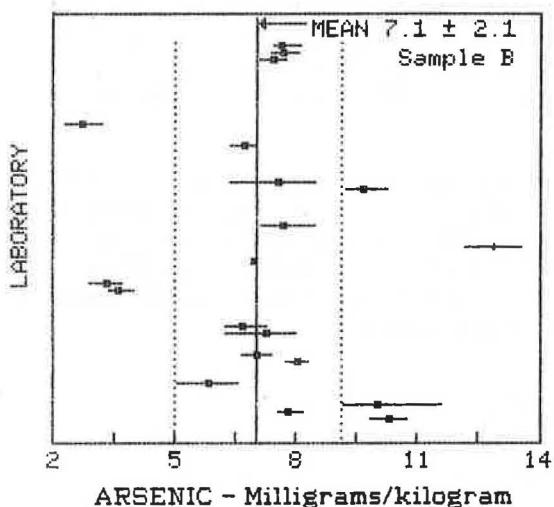


21 labs submitted results ranging from 10.2 to 38.5 mg/kg.

All labs submitted quantitative values with a mean of 25 ± 5 mg/kg.

A *t* test rejected the results of none of these labs.

The relative standard deviation is 21 percent.

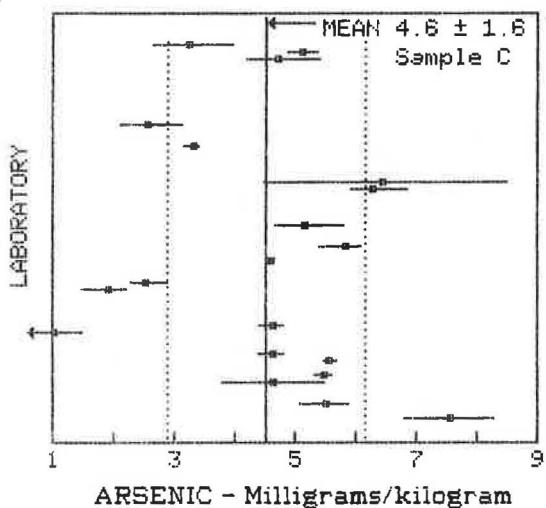


20 labs submitted results ranging from 2.32 to 13.6 mg/kg.

All labs submitted quantitative values with a mean of 7.4 ± 2.4 mg/kg.

A *t* test rejected the results of 1 of these labs.

The mean of the results of the remaining 19 labs is 7.1 ± 2.1 mg/kg. The relative standard deviation is 29 percent.



20 labs submitted results ranging from 0.85 to 8.5 mg/kg.

All labs submitted quantitative values with a mean of 4.6 ± 1.6 mg/kg.

A *t* test rejected the results of none of these labs.

The relative standard deviation is 35 percent.

TABLE V
ARSENIC CONCENTRATIONS - mg/kg

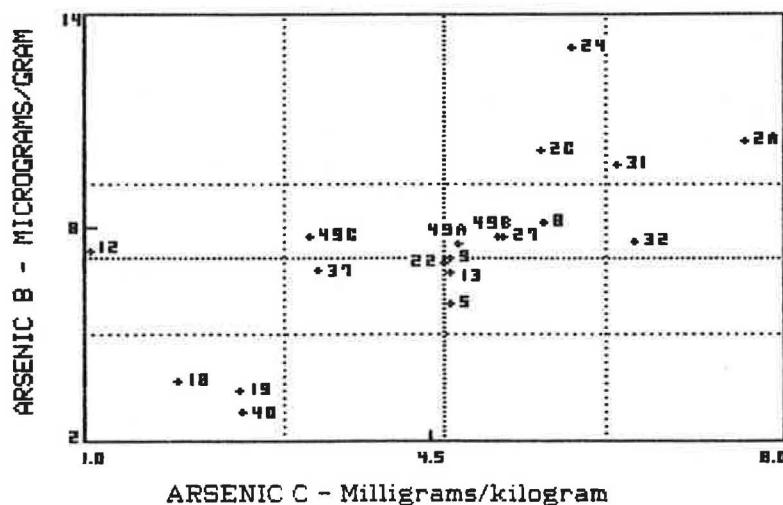
Sample A	25 ± 5	(21%)	21, 0
Sample B	7.1 ± 2.1	(29%)	19, 1
Sample C	4.6 ± 1.6	(35%)	20, 0

Methods of sample preparation are almost equally divided between (i) closed vessel decompositions with nitric acid alone or in combination with perchloric or sulphuric acids, (ii) open beaker decompositions with nitric acid or nitric acid in combination with sulphuric acid, or (iii) dry ashing. One lab carried out an oxygen combustion of the sample and two labs which used instrumental neutron activation analysis (INAA) did not treat the sample at all.

Slightly over half the labs used a hydride generation (HG) technique followed by atomic absorption spectrometry (AAS) in order to measure the arsenic, except in one case where spectrophotometry was used. The majority of the others used GFAAS, generally adding nickel to the sample prior to electrothermal atomization. Three labs employed INAA and one, ICP atomic emission spectrometry.

There are no apparent relationships between methodologies and performance. The close grouping of a number of laboratories about the means are representative of almost all methods of sample decomposition and arsenic measurement. The poor precision between laboratories, especially for Samples B and C, is disappointing. The 35 percent RSD for Sample C represents an almost six-fold difference between the lowest and highest samples. Because of the high general scatter between labs only one set of results in all three samples was identified as an outlier.

ARSENIC B vs ARSENIC C



The plot of Sample B against Sample C on page 12 shows little indication of erraticism, just a general spread from low to high results with some bunching near the means.

The submitted data for arsenic are listed on pages 38 to 40.

Cadmium

Cadmium results for Samples A and B are equivalent in quality to those achieved for zinc, especially considering the rather low concentrations of cadmium relative to those of zinc. However, the interlaboratory precision deteriorates markedly for Sample C, which is not too surprising considering the low concentration of 0.06 mg/kg cadmium in this sample.

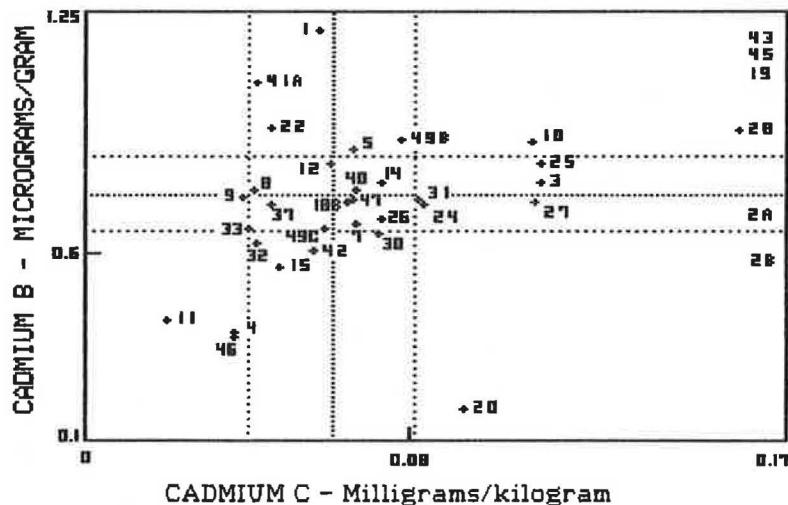
Summaries of the results are listed beside the graphs on page 14 and the overall final means and standard deviations are shown below in Table VI.

TABLE VI

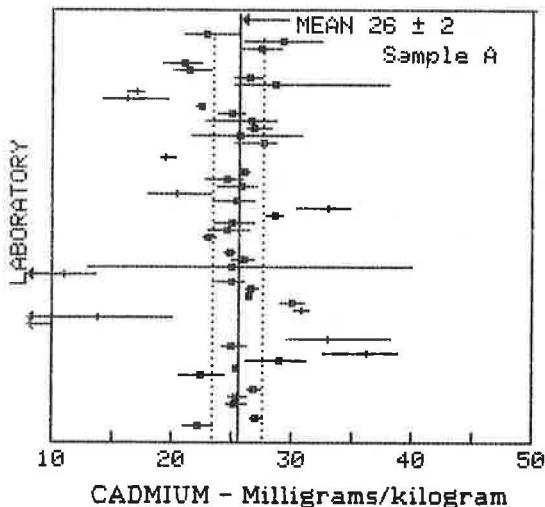
CADMUM CONCENTRATIONS - mg/kg

Sample A	26	\pm 2	(8%)	37,11
Sample B	0.75	\pm 0.10	(13%)	37,10
Sample C	0.06	\pm 0.02	(38%)	31, 6

CADMIUM B vs CADMIUM C



CADMIUM DATA

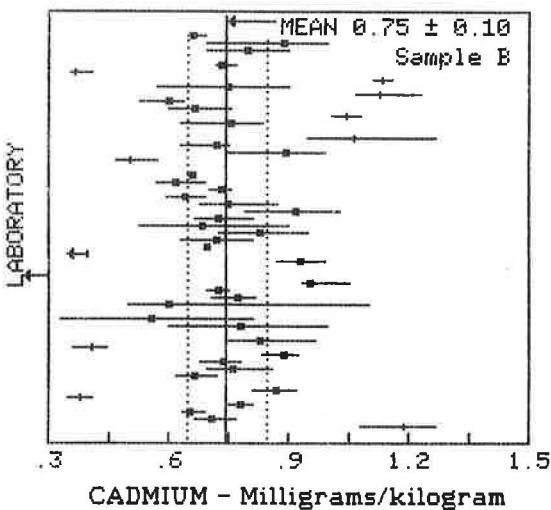


48 labs submitted results ranging from 5.2 to 40 mg/kg.

All labs submitted quantitative values with a mean of 25 ± 5 mg/kg.

A *t* test rejected the results of 11 of these labs.

The mean of the results of the remaining 37 labs is 26 ± 2 mg/kg. The relative standard deviation is 8 percent.

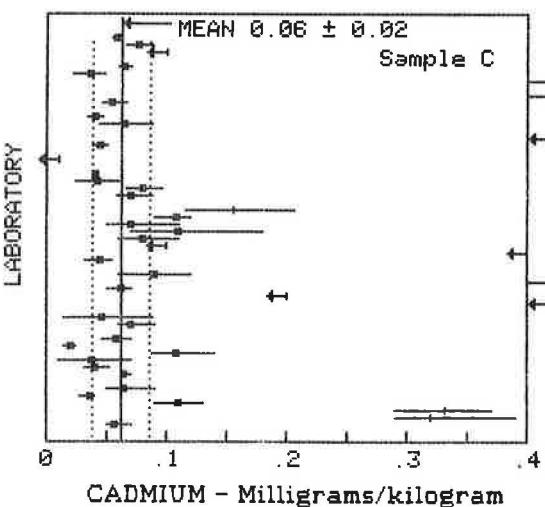


48 labs submitted results ranging from 0.12 to 1.27 mg/kg.

47 labs submitted quantitative values over the same range with a mean of 0.75 ± 0.20 mg/kg.

A *t* test rejected the results of 10 of these labs.

The mean of the results of the remaining 37 labs is 0.75 ± 0.10 mg/kg. The relative standard deviation is 13 percent.



44 labs submitted results ranging from 0.01 to 0.85 mg/kg.

37 labs submitted quantitative values over the same range with a mean of 0.13 ± 0.17 mg/kg.

A *t* test rejected the results of 6 of these labs.

The mean of the results of the remaining 31 labs is 0.06 ± 0.02 mg/kg. The relative standard deviation is 38 percent.

Methods of sample preparation are generally the same as for copper and zinc. About 40 and 30 percent of the participants used FAAS and GFAAS respectively to measure cadmium in Sample A. However, these percentages were approximately reversed for Sample B, and fifty percent of the analysts used GFAAS for cadmium in Sample C.

More than half of all results rejected for Samples A and B were obtained using GFAAS. Also, almost all the results biased both high and low with respect to Sample B, shown below in the plot of Sample B against Sample C, resulted from GFAAS measurements. However, the majority of the same labs produced acceptable values for the much lower concentration of cadmium in Sample C.

The results of three of the nine labs using FAAS without prior concentration of the cadmium for Sample C were rejected, and another three were biased rather high for this sample as seen in the plot on page 18. FAAS may not be sensitive enough to provide reliable results for cadmium at less than 0.1 mg/kg without some form of preconcentration unless great care is taken.

It is also apparent that inductively coupled plasma atomic emission spectrometry is not sensitive enough to analyze for cadmium at this level without preconcentration.

The submitted data for cadmium are listed on pages 40 to 45.

Mercury

About three quarters of the responding laboratories submitted results for mercury. In spite of the inherently low concentrations of mercury in these samples the great majority of labs that analyze for mercury are apparently capable of producing good results.

Summaries of the results are listed beside the graphs on page 16 and the overall final means and standard deviations are shown in Table VII below.

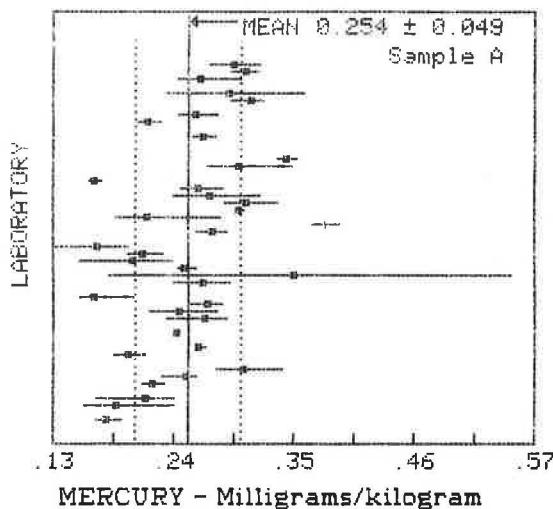
TABLE VII
MERCURY CONCENTRATIONS - mg/kg

Sample A	0.254 ± 0.049	(19%)	36, 1
Sample B	0.081 ± 0.012	(10%)	31, 4
Sample C	0.056 ± 0.009	(16%)	30, 5

The majority of labs (57 percent) prefer to digest the sample with nitric acid or a mixture of nitric and sulphuric and/or perchloric acids, sometimes followed by an oxidant such as hydrogen peroxide, potassium permanganate or vanadium pentoxide. About 30 percent of the participants used a closed vessel decomposition (BOMB) with nitric acid or the same mixed acids as the former group. A few labs destroyed the organic material using oxygen combustion followed by acid dissolution of the residue.

All but three of the labs used some form of cold vapour atomic absorption spectrometry (CVAAS) to determine the mercury. Only a few of these collected the mercury on gold before evolution of the metal into the absorption cell. One lab used INAA, another an electroanalytical procedure and a third isotope dilution mass spectrometry.

MERCURY DATA

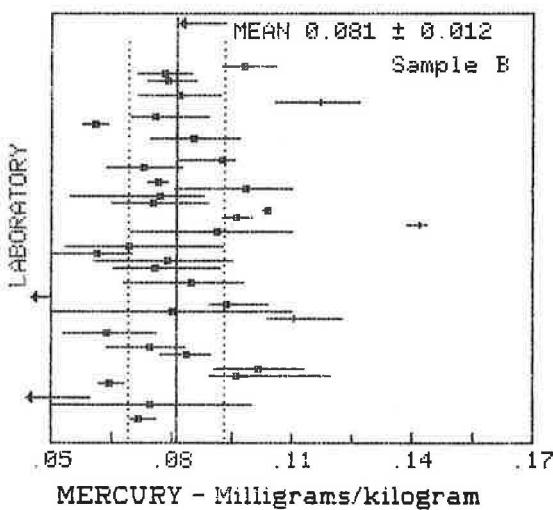


37 labs submitted results ranging from 0.13 to 0.549 mg/kg.

All labs submitted quantitative values with a mean of 0.258 ± 0.052 mg/kg.

A *t* test rejected the results of 1 of these labs.

The mean of the results of the remaining 36 labs is 0.254 ± 0.049 mg/kg. The relative standard deviation is 19 percent.

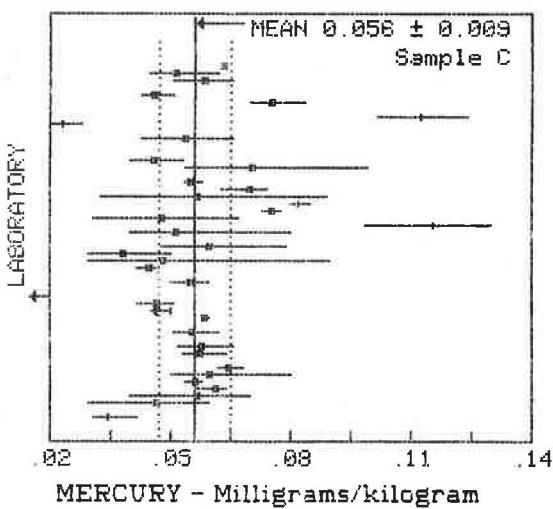


36 labs submitted results ranging from less than 0.01 to 0.144 mg/kg.

35 labs submitted quantitative values with a range of 0.03 to 0.144 and a mean of 0.084 ± 0.019 mg/kg.

A *t* test rejected the results of 4 of these labs.

The mean of the results of the remaining 31 labs is 0.081 ± 0.012 mg/kg. The relative standard deviation is 15 percent.



37 labs submitted results ranging from less than 0.01 to 0.13 mg/kg.

35 labs submitted quantitative values with a range of 0.021 to 0.13 and a mean of 0.059 ± 0.018 mg/kg.

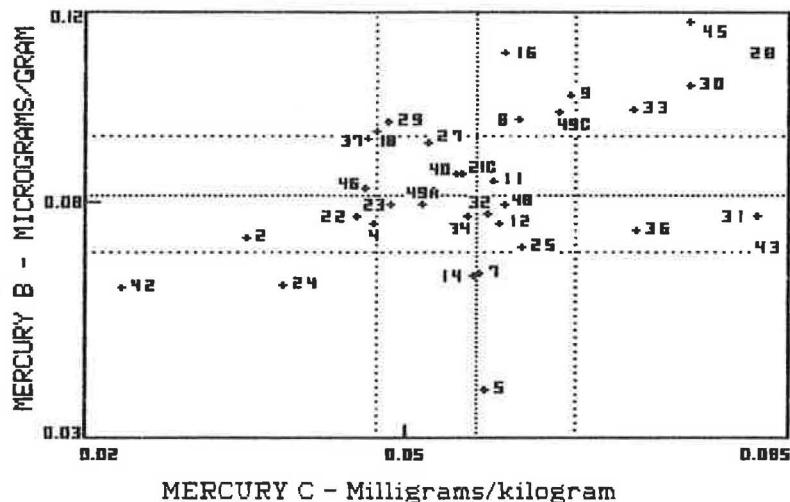
A *t* test rejected the results of 5 of these labs.

The mean of the results of the remaining 30 labs is 0.056 ± 0.009 mg/kg. The relative standard deviation is 16 percent.

There are no apparent relationships between methodologies and performance.

The plot of Sample B against Sample C below again shows little indication of erraticism. There is a generally a greater bias to high values rather than to low, but overall the results are excellent considering the low mercury concentrations.

MERCURY B vs MERCURY C



The submitted data for mercury are listed on pages 45 to 48.

Lead

About 85 percent of the labs submitted results for lead in Sample A. This number decreased to 65 percent and 75 percent for Samples B and C respectively with the changes in lead concentrations.

The results for lead are disappointing, showing no improvement over the 6th Intercomparison Exercise. Even at the 6 mg/kg level the relative standard deviation of 36 labs (6 having been rejected) is 30 percent, a figure indicating a four-fold difference between the lowest and highest means of the group.

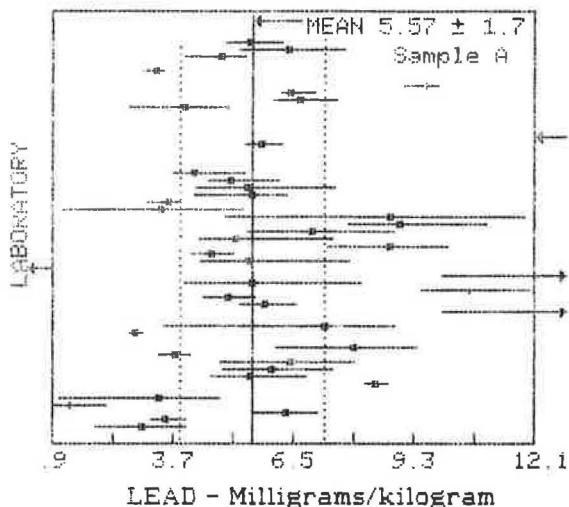
Although there is a possibility of some interbottle inhomogeneity for lead in Samples B and C this probably accounts for only a minor fraction of the interlab variance for these samples. The RSD of 41 percent for both of these samples represents a greater than nine-fold factor between low and high values.

Summaries of the results are listed beside the graphs on page 18 and the overall final means and standard deviations are shown on page 19 in Table VIII.

Samples for lead analyses were generally prepared as for copper, zinc and cadmium.

About half the labs employed GFAAS for lead in Sample A and 30 percent used FAAS. A few labs used electroanalytical methods and some others inductively coupled plasma atomic emission spectrometry (ICP-AES). Three labs concentrated the lead before measurement by FAAS

LEAD DATA

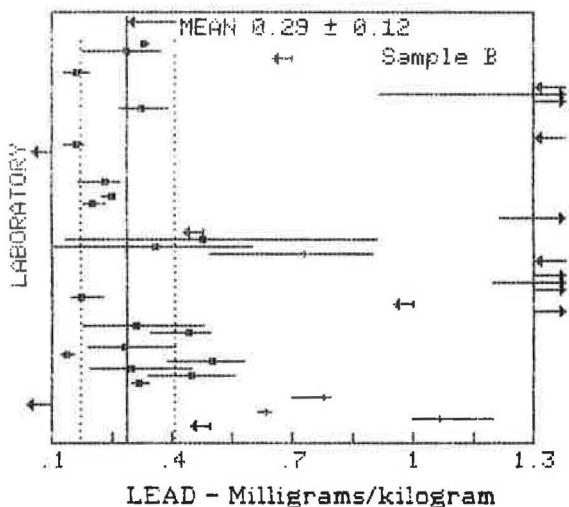


43 labs submitted results ranging from 0.6 to 63 mg/kg.

42 labs submitted quantitative values with the same range and a mean of 6.65 ± 5.94 mg/kg.

A *t* test rejected the results of 6 of these labs.

The mean of the results of the remaining 36 labs is 5.57 ± 1.70 mg/kg. The relative standard deviation is 30 percent.

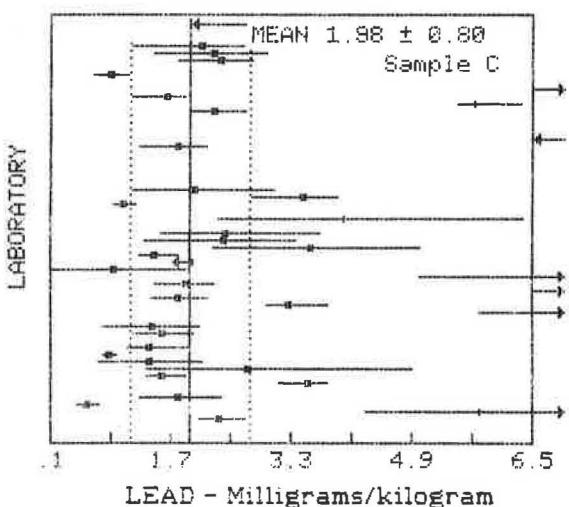


39 labs submitted results ranging from 0.065 to less than 25 mg/kg.

31 labs submitted quantitative values with a range of 0.065 to 18 and a mean of 1.41 ± 2.90 mg/kg.

A *t* test rejected the results of 11 of these labs.

The mean of the results of the remaining 20 labs is 0.29 ± 0.12 mg/kg. The relative standard deviation is 41 percent.



38 labs submitted results ranging from 0.1 to less than 25 mg/kg.

36 labs submitted quantitative values with a range of 0.1 to 18.8 and a mean of 3.29 ± 3.38 mg/kg.

A *t* test rejected the results of 7 of these labs.

The mean of the results of the remaining 29 labs is 1.98 ± 0.80 mg/kg. The relative standard deviation is 41 percent.

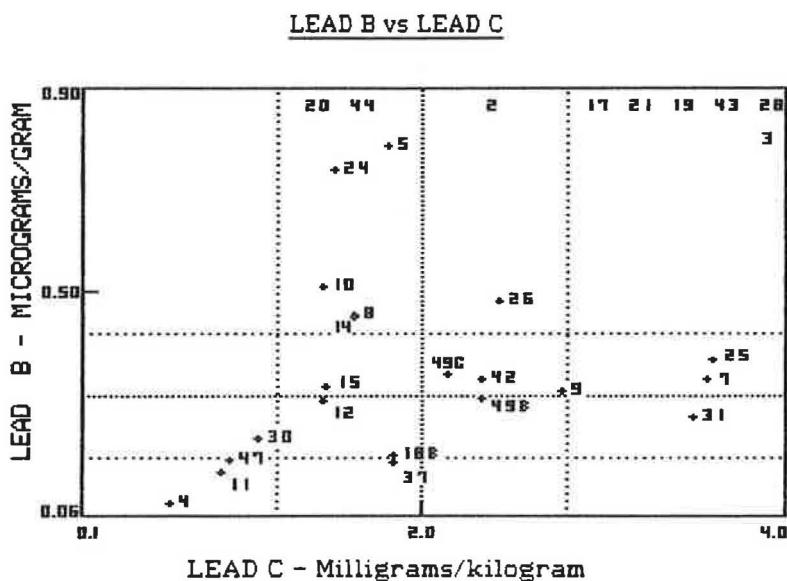
or GFAAS. The proportion of those reporting quantitative results using FAAS declines with lead concentration to about 20 percent in Samples B and C.

TABLE VIII
LEAD CONCENTRATIONS - mg/kg

Sample A	5.57 ± 1.70	(30%)	36, 6
Sample B	0.29 ± 0.12	(41%)	20, 11
Sample C	1.98 ± 0.80	(41%)	29, 7

There is no clear relationship between methodology and performance for Sample A. Three of the six rejected sets were obtained using FAAS.

Five of the eleven and eight sets rejected respectively for Samples B and C were from FAAS data. These represent about two-thirds of all the FAAS results for these samples.



The tendency toward high biased results is obvious from the plot of Sample B against Sample C on page 21. Four of the six values in the upper right quadrant come from FAAS results. On the other hand, four of the five high results for Sample B (but acceptable for C) are GFAAS values.

The submitted data for lead are listed on pages 49 to 52.

Other Metals

Results for 14 other metals were received from a number of laboratories: sodium (1), magnesium (1), phosphorus (1), potassium (1), calcium (1), chromium (5), manganese (12), iron (13), cobalt (7), nickel (11), selenium (5), bromine (1), strontium (2) and silver (2). There were not sufficient data for any of these metals to provide for an adequate analysis. Means and standard deviations were calculated for all metals where there were at least 5 sets of data (chromium, manganese, iron, cobalt, nickel and selenium). These are listed below in Tables IX to XIV and the submitted data for those metals are listed on pages 53 to 57.

There was no attempt to eliminate even obvious outliers from individual sets of results. The results of lab 21 were arbitrarily deleted from the nickel calculations.

TABLE IX

CHROMIUM CONCENTRATIONS - mg/kg

n = 5

Sample A	2.40	±	1.22	(51%)
Sample B	0.61	±	0.37	(60%)
Sample C	1.45	±	0.54	(37%)

TABLE X

MANGANESE CONCENTRATIONS - mg/kg

n = 12

Sample A	20.4	±	3.8	(19%)
Sample B	1.96	±	2.12	(108%)
Sample C	14.6	±	4.90	(34%)

TABLE XI

IRON CONCENTRATIONS - mg/kg

n = 13

Sample A	167	±	29.9	(18%)
Sample B	23.3	±	11.4	(49%)
Sample C	136	±	26.7	(20%)

TABLE XII

COBALT CONCENTRATIONS - mg/kg

n = 7

Sample A	0.67	±	0.36	(54%)
Sample B	0.20	±	0.12	(59%)
Sample C	0.07	±	0.02	(30%)

TABLE XIII

NICKEL CONCENTRATIONS - mg/kg

n = 10

Sample A	2.69	±	0.97	(36%)
Sample B	0.38	±	0.19	(49%)
Sample C	0.74	±	0.37	(50%)

TABLE XIV

SELENIUM CONCENTRATIONS - mg/kg

n = 5

Sample A	6.9	±	1.0	(14%)
Sample B	0.63	±	0.33	(53%)
Sample C	2.46	±	0.34	(14%)

3. SUMMARIES

The results for the six core metals in the three samples are summarized in Table XV on page 23.

Table XVI on page 24 is an attempt to show the pattern of results achieved by each laboratory for the six metals in the three samples. The symbols in the table represent

- ** Results reported and accepted by a *t* test.
- LD Results reported, but less than a limit of detection.
- RT Results reported, but rejected by a *t* test.
- No results reported.

The list of participating laboratories (Table XVII) is on pages 25 to 28.

A set of charts indicating individual laboratory precision and bias for each of the six core metals in each sample is appended (pages 58 to 75).

4. CONCLUSIONS

This exercise has demonstrated that the good majority of the participants can produce comparable and accurate data for the trace metals copper, zinc, cadmium and mercury at levels found in fish and shellfish tissues.

A number of laboratories have problems with cadmium at the less than 0.1 mg/kg level, but at least half the participants do fairly well even at this low level.

The analysis for arsenic seems to present problems even to the minority of laboratories that submitted arsenic results. The reason for this is not evident from this study but is probably related to sample preparation and the efficiency of arsine generation. About half of the arsenic participants produce comparable and probably accurate results.

Lead continues to be a problem element. Most laboratories using flame atomic absorption spectrometry have troubles at the two mg/kg level. The majority of the others produce an unsatisfactory spread of values (RSD = 41%) at this concentration. Also, only a small number of laboratories can successfully cope with a sample containing lead at a sub mg/kg concentration. It would appear that the plaice sample (C) may have been contaminated with respect to lead during preparation.

Comparable "dry weights" can be achieved with samples such as used in this study by drying for a few hours in an oven at 105 to 115 degrees Celcius.

5. ACKNOWLEDGEMENTS

The authors gratefully acknowledge the help of Dr. Graham Topping (DAF Marine Laboratory, Aberdeen) in the planning of this exercise. We would also like to thank Dr. Uwe Harms (Bundesforschungsanstalt für Fischerei, Hamburg), and Dr. Jack Uthe and Dr. Chiu Long Chou (Fisheries and Oceans Canada, Halifax) for their contributions, as well as Vincent Clancy of this laboratory for his help in packaging and circulating the materials.

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TABLE XV

SUMMARY OF RESULTS
milligrams per Kilogram

SAMPLE A

	<u>Copper</u>	<u>Zinc</u>	<u>Arsenic</u>	<u>Cadmium</u>	<u>Mercury</u>	<u>Lead</u>
Total range	110-508	90.9-335	10.2-38.5	5.2-40	0.13-0.549	0.6-63
No. of labs	49	49	21	48	37	43
Quant. values	110-508	90.9-335	10.2-38.5	5.2-40	0.13-0.549	0.6-63
No. of labs	49	49	21	48	37	42
Mean	333±45	177±32	25±5	25±5	0.258±0.052	6.65±5.94
After χ^2 test	236-437	153-222	10.2-38.5	19.3-40	0.13-0.549	1.1-11.9
No. of labs	45	38	21	37	36	36
Mean	331±32	179±10	25±5	26±2	0.254±0.049	5.57±1.70
RSD	10%	6%	21%	8%	19%	30%

SAMPLE B

	<u>Copper</u>	<u>Zinc</u>	<u>Arsenic</u>	<u>Cadmium</u>	<u>Mercury</u>	<u>Lead</u>
Total range	0.48-9.9	32-98	2.32-13.6	0.12-1.27	<0.01-0.144	0.065-<25
No. of labs	48	48	20	48	36	39
Quant. values	0.48-9.9	32-98	2.32-13.6	0.12-1.27	0.03-0.144	0.065-18
No. of labs	48	48	20	47	35	31
Mean	3.9±1.0	58±12	7.4±2.4	0.75±0.20	0.084±0.019	1.41±2.90
After χ^2 test	0.48-9.9	47.1-79	2.32-11.6	0.33-1.1	0.05-0.12	0.065-0.91
No. of labs	41	40	19	37	31	20
Mean	3.7±0.6	58±5	7.1±2.1	0.75±0.10	0.081±0.012	0.29±0.12
RSD	16%	9%	29%	13%	15%	41%

SAMPLE C

	<u>Copper</u>	<u>Zinc</u>	<u>Arsenic</u>	<u>Cadmium</u>	<u>Mercury</u>	<u>Lead</u>
Total range	1.0-8.1	59-187	0.85-8.5	0.01-0.85	<0.01-0.13	0.1-<25
No. of labs	49	48	20	44	37	38
Quant. values	1.0-8.1	59-187	0.85-8.5	0.01-0.85	0.021-0.13	0.1-18.8
No. of labs	49	48	20	37	35	36
Mean	3.5±1.3	98±23	4.6±1.6	0.13±0.17	0.059±0.018	3.29±3.38
After χ^2 test	1.0-5.46	59-117	0.85-8.5	0.02-0.18	0.031-0.0995	0.1-5.0
No. of labs	42	42	20	31	30	29
Mean	3.1±0.6	93±9	4.6±1.6	0.06±0.02	0.056±0.009	1.98±0.80
RSD	27%	10%	35%	38%	16%	41%

TABLE XVI

PATTERN OF RESULTS

Lab	Copper	Zinc	Arsenic	Cadmium	Mercury	Lead
1	** ** **	-----	-----	** RT **	-----	** LD --
2A	** ** **	** ** **	** ** **	** ** RT	** ** RT	-----
2B	-- ** **	--- ---	** ** --	-- ** RT	-----	** RT **
2C	-----	-----	** ** **	-----	-----	-----
3	** ** **	** ** **	-----	** ** **	-----	** RT RT
4	** ** **	** ** **	-----	** RT **	** ** **	RT ** **
5	** ** **	** ** **	** ** **	** ** **	** RT **	** RT **
6	-----	-----	-----	-----	-----	-----
7	** ** **	** ** **	-----	** ** **	** ** **	** ** **
8	** ** **	** ** **	** ** **	** ** **	** ** **	** ** **
9	** ** **	** ** **	** ** **	** ** **	** ** **	** ** **
10	** -- RT	RT ** **	-----	RT ** **	-----	** ** **
11A	** ** **	** ** **	** -- --	** RT **	** ** **	** ** **
11B	** ** **	-----	-----	-----	-----	-----
12	** ** **	** ** **	** ** **	RT ** **	** ** **	-----
13	-----	** RT RT	** ** **	-----	-----	-----
14	** ** **	RT ** **	-----	RT ** **	** ** **	** ** **
15	RT ** **	** ** **	-----	RT ** **	-----	** ** **
16	-- --	** ** **	-----	RT -- --	** RT **	-----
17	** ** **	** ** **	-----	** ** LD	** ** LD	RT RT RT
18A	** ** **	** ** **	** ** **	** ** LD	** ** **	** LD **
18B	** ** **	RT ** **	-----	** ** **	-----	** ** **
19	** ** **	RT RT RT	** ** **	** ** RT	** LD LD	RT RT RT
20	RT ** **	RT RT **	-----	RT RT **	-----	** RT **
21A	** ** **	-----	-----	** -- --	** -- --	RT RT RT
21B	-----	-----	-----	-----	** -- --	-----
22	** ** **	** ** **	** ** **	** ** **	** ** **	RT -- **
23A	** ** **	** ** **	-----	** LD LD	** ** **	** LD LD
23B	-----	-----	-----	-- ** LD	-----	-- LD LD
24	** RT RT	** ** RT	** RT **	** ** **	** ** **	** RT **
25	** ** **	** -- --	-----	** ** **	** ** **	** ** **
26	** ** **	-----	-----	** ** **	-----	** ** **
27A	** ** **	** ** **	** ** **	** ** **	** ** **	** LD **
27B	-----	-----	-----	-----	-----	** -- --
28	** ** **	RT ** **	-----	RT ** RT	RT RT RT	** RT RT
29	RT ** **	RT RT RT	-----	** ** --	** ** **	** -- --
30	** ** **	RT ** **	-----	RT ** **	** ** **	** ** **
31	** ** **	** ** **	** ** **	** ** **	** ** RT	** ** **
32	** ** **	** ** **	** ** **	** ** **	** ** **	** -- --
33A	** ** **	** ** **	-----	** ** **	** ** **	** -- --
33B	-----	-----	-----	-----	-----	** -- --
34	-----	-----	-----	-----	** ** **	-----
35	-- --	RT -- --	-----	RT RT LD	-----	-----
36	-- RT --	-- ** --	-----	-- ** --	** ** **	-- LD --
37	-- RT RT	** ** **	** ** **	** ** **	** ** **	** ** **
38	-- RT RT	RT ** **	-----	** RT LD	-----	LD LD LD
39	-- --	-- --	-----	-- --	-----	-----
40	-- --	-- --	-----	-- --	-----	-----
41A	-- RT **	-----	-----	-- RT **	-----	-----
41B	-- RT **	-----	-----	-- RT **	-----	-----
42	-- RT RT	-- RT **	-----	RT ** **	** ** RT	** ** **
43	RT ** RT	-- RT RT	-----	RT RT RT	** ** RT	** RT RT
44	-- RT **	-- RT **	-----	-- RT **	-----	** RT **
45	-- RT RT	-- RT **	-----	-- RT RT	-- RT **	RT LD RT
46	-- RT RT	RT RT RT	-----	-- RT **	** ** **	-----
47	-- RT RT	-- RT RT	-----	-- RT **	** ** **	-----
48	-----	-----	-----	-----	** ** **	-----
49A	-- RT RT	-- RT RT	** ** **	-- RT RT	** RT **	** LD **
49B	-- RT RT	-- RT RT	** ** **	-- RT RT	** RT **	** LD **
49C	-- RT RT	-- RT RT	** ** **	-- RT RT	** RT **	** LD **

TABLE XVII

PARTICIPATING LABORATORIES

BELGIUM	7 Instituut voor Scheikundig Onderzoek Museumlaan 5 B 1980 Tervuren	R. De Borger J.R. Istanas
	13 Instituut voor Nucleaire Wetenschappen Rijksuniversiteit-Gent Proeftuinstraat 86 9000 Gent	R. Dams
	45 Analytische Scheikunde Faculteit der Toegepaste Wetenschappen Vrije Universiteit Brussel Pleinlaan 2 1050 Brussel	I. Elskens
CANADA	2 Institute for Environmental Studies University of Toronto Toronto, Ontario M5S 1A1	Jon Van Loon
	10 Atlantic Research Laboratory National Research Council Halifax, Nova Scotia B3H 3Z1	Roger Guevremont
	16 Centre Champlain des Sciences de la Mer Pêches et Océans B.P. 15500 901, Cap Diamant Québec, P.Q. G1K 7Y7	Daniel Cossa
	37 Fisheries Research Laboratory P.O. Box 550 Halifax, Nova Scotia B3J 2S7	Chiu Long Chou Jack Utche
	49 Division of Chemistry National Research Council Montreal Road Ottawa, Ontario K1A 0R9	Shier Berman Victor Boyko Armand Desauvigniers Alex Mykytiuk Michael Siu
	Institute of Scientific Research University of Québec Rimouski P.Q.	Daniel Cossa
DENMARK	6 Danish Isotope Centre Skelbaeksgade 2 DK 1717 Copenhagen V	Iver Drabaek
	33 Marine Pollution Laboratory Kavalergarden 6 DK-2920 Charlottenlund	Arne Jensen
	36 Water Quality Institute 11, Agern Alle DK-2970 Horsholm	Merete Reuss
	40 National Food Institute 19, Mørkholøj Bygade DK-2860 Soborg	Allan Andersen Gitte Rasmussen

FINLAND	50	National Board of Waters Research Laboratory Kylasaarenkatu 10 00550 Helsinki	Kirsti Haapala
FRANCE	24	Institut Scientifique et Technique des Pêches Maritimes Rue de l'Ille d'Yeu B.P. 1049 44037 Nantes Cedex	Yves Thibaud
GERMAN DEMOCRATIC REPUBLIC	14	Institut für Meereskunde Schliessfach 38 Seestrasse 15 DDR-2530 Rostock-Warnemünde	L. Brügmann
	42	Hygiene Institut Rostock Abteilung Lebensmittel und Ernährungshygiene Gertrudenstrasse 9 DDR-2500 Rostock	G. Manthey
GERMAN FEDERAL REPUBLIC	9	Bodenseewerk Perkin-Elmer & Co GmbH Postfach 1120 D-7770 Überlingen	Bernhard Welz
	11	Bundsforschungsanstalt für Fischerei Labor für Radioökologie der Gewässer Wustand 2 D-2000 Hamburg 55	Uwe Harms
	22	Bundesanstalt für Gewässerkunde Kaiserin Augusta Anlagen 15-17 D-5400 Koblenz	W. Speer
	35	Institut für Meereskunde an der Universität Kiel Dusternbrooker Weg 20 D 2300 Kiel 1	H. Fischer
IRELAND	46	Fisheries Research Centre Abbotstown Castleknock Co. Dublin	Mairin O Sullivan
NETHERLANDS	5	Government Institute of Sewage and Waste Water Treatment Hereweg 99a 9721 AA Groningen	W. Wilts
	31	Rijks-Kwaliteitsinstituut Postbus 230 6700 AE Wageningen	M. Heuver
	32	Division for Nutrition and Food Research TNO P.O. Box 183 1970 AD IJmuiden	J.B. Luten
	48	Netherlands Institute for Fishery Investigations P.O. Box 68 1970 AA IJmuiden	H. Pieters
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	18	Central Institute for Industrial Research P.O. Box 350 Blindern Oslo 3	Beate Enger Lilla Madsen
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COPPER DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
1			HNO₃-GFAAS						
	299	6.2	320	305	265	304	305	297	
	4.43	8.7	3.73	4.35	4.85	4.43	4.7	4.49	
	1.91	12.8	1.69	2.23	1.87	1.99	2.09	1.58	
2A			BOMB/HNO₃/HClO₄-ICP						
	346	1.1	345	339	345	351	347	347	
	4.53	10.0	5.4	4.1	4.6	4.4	4.4	4.3	
	4.87	4.4	4.8	5.0	5.1	4.8	5.0	4.5	
2B			BOMB/HNO₃/HClO₄-GFAAS						
	----	----	No data						
	4.62	5.2	4.6	5.0	4.4	4.5	4.8	4.4	
	3.70	3.4	3.9	3.7	3.7	3.5	3.7	3.7	
3			HNO₃/HClO₄/HCl-FAAS						
	316	1.4	318	311	323	317	316	311	
	3.83	11.9	3.7	4.5	3.4	3.5	4.3	3.6	
	3.13	1.6	3.2	3.1	3.1	3.1	3.1	3.2	
4			HNO₃-FAAS						
	330	0.8	330	332	334	327	328	330	
	4.08	20.2	3.38	3.98	3.50	3.42	5.02	5.20	
	4.74	10.1	4.35	4.29	4.71	5.46	5.17	4.44	
5			BOMB/HNO₃-FAAS(A)-GFAAS(B/C)						
	341	0.9	341	341	346	343	341	337	
	3.23	18.9	4.0	3.1	3.7	2.4	2.7	3.5	
	2.53	11.9	2.9	2.5	2.9	2.3	2.4	2.2	
6			NO DATA						
7			DRY ASH/HNO₃/H₂O₂-FAAS						
	328	0.9	331	332	327	327	325	326	
	3.68	3.8	3.51	3.85	3.83	3.59	3.59	3.69	
	4.67	3.4	4.38	4.75	4.61	4.78	4.75	4.78	
8			HNO₃/H₂O₂-FAAS						
	326	1.3	327	332	324	329	324	320	
	3.60	32.3	4.1	2.1	2.7	4.4	5.2	3.1	
	2.83	7.3	3.1	3.0	2.9	2.6	2.8	2.6	
9			BOMB/HNO₃-GFAAS						
	398	5.5	376	420	397				
	3.74	11.5	3.26	3.86	4.09				
	3.18	0.9	3.2	3.2	3.15				
10			HNO₃/HClO₄-ICP						
	387	2.2	401	379	390	388	387	378	
	----	----	No data						
	5.89 ^m	24.4	5.45	3.30	5.97	6.80	6.43	7.41	
11A			BOMB/HNO₃-EXTN-FAAS						
	344	3.2	351	349	327	357	335	347	
	3.88	10.6	3.3	3.9	4.4	3.5	4.1	4.1	
	2.65	7.8	2.8	2.5	2.8	2.9	2.4	2.5	
11B			BOMB/HNO₃-FAAS						
	359	4.7	358	382	330	362	364	358	
	3.70	15.5	3.0	3.9	3.2	4.6	3.9	3.6	
	2.42	18.9	2.3	2.5	1.8	2.7	3.1	2.1	

COPPER DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
12			BOMB/HNO ₃ /HClO ₄ -FAAS						
	332	0.7	330	333	331	329	335	333	
	3.25	16.9	2.85	3.00	2.89	2.91	3.67	4.18	
	3.12	29.9	2.90	2.98	2.97	2.64	2.27	4.94	
13			NO DATA						
14			BOMB/HNO ₃ -GFAAS						
	296	2.6	303	297	286	288	298	305	
	3.43	41.3	4.1	5.1	2.7	1.9	4.8	2.0	
	1.73	21.5	2.0	2.0	1.0	1.8	1.8	1.8	
15			HNO ₃ /H ₂ O ₂ -GFAAS						
	163 ^{RT}	31.1	140	110	220	110	220	180	
	2.72	50.0	2.5	5.4	2.5	2.0	2.3	1.6	
	2.02	19.2	1.7	2.3	2.2	2.4	2.1	1.4	
16			BOMB/HNO ₃ -FAAS						
	382	1.1	377	376	384	387	383	383	
	----	----	No data						
	----	----	No data						
17			HNO ₃ /H ₂ O ₂ -FAAS						
	352	1.2	350	350	350	350	350	360	
	2.95	9.3	2.6	3.1	3.1	2.6	3.1	3.2	
	2.62	17.4	2.2	2.7	3.3	2.7	2.2		
18A			DRY ASH/HNO ₃ -ICP						
	307	0.7	305	307	309	309	307	304	
	3.59	8.6	3.67	3.68	3.32	3.56	3.23	4.10	
	3.23	3.6	3.37	3.35	3.13	3.12	3.13	3.27	
18B			DRY ASH/HNO ₃ -EXTN(B&C)-FAAS						
	299	1.6	303	306	294	298	301	294	
	3.97	5.8	3.94	4.07	3.86	3.93	3.66	4.35	
	3.24	4.8	3.42	3.36	3.28	3.02	3.25	3.08	
19			DRY ASH/HCl-FAAS						
	324	7.8	345	309	280	345	328	335	
	3.48	13.7	3.25	2.83	3.35	4.09	3.98	3.35	
	3.59	3.7	3.74	3.63	3.52	3.52	3.74	3.41	
20			HNO ₃ -GFAAS						
	424 ^{RT}	12.6	472	508	382	383	393	404	
	3.42	19.2	3.0	4.1	4.4	2.9	3.0	3.1	
	2.55	31.4	1.6	2.3	3.4	2.5	1.9	3.6	
21			BOMB/HNO ₃ /H ₂ SO ₄ (A)-HNO ₃ /HClO ₄ (B/C)-ASV						
	373	8.7	388	418	373	349	336		
	4.77	28.5	3.2	5.5	5.6				
	4.43	21.0	3.4	5.2	4.7				
22			HNO ₃ /HCl-FAAS(A)/GFAAS(B/C)						
	328	1.5	334	320	331	331	330	325	
	3.07	22.2	3.0	3.6	2.6	4.0	3.1	2.1	
	2.83	20.3	2.8	2.3	2.9	3.8	3.0	2.2	
23A			HNO ₃ -FAAS-0.25 g samples						
	316	2.7	311	311	309	316	317	332	
	3.68	15.5	4.7	3.3	4.0	3.4	3.2	3.5	
	2.25	14.3	2.2	2.1	1.7	2.5	2.5	2.5	

COPPER DATA

COPPER DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
36	-----	-----	HNO ₃ /H ₂ SO ₄ /H ₂ O ₂ -EXTN-GFAAS					
	1.90 ^{RT}	8.7	1.79	2.11	1.83	2.09	1.84	1.71
	-----	-----	No data					
37	363	3.1	350	372	378	368	357	354
	5.37 ^{RT}	12.5	5.44	5.82	6.16	5.68	4.53	4.59
	5.46 ^{RT}	9.0	5.57	6.00	5.33	4.57	5.54	5.76
38	288	13.0	320	323	299	237	263	
	7.04 ^{RT}	13.4	5.71	7.10	7.12	8.53	7.30	6.45
	6.16 ^{RT}	9.4	6.85	6.44	6.11	6.10	5.28	
39	301	2.5	300	296	300	293	314	306
	-----	-----	No data					
	3.80	2.6	3.71	3.85	3.75	3.87	3.94	3.70
40	330	7.2	349	295	307	348	350	333
	4.09	15.7	3.74	4.13	4.33	5.13	3.20	4.02
	3.71	7.1	3.40	3.76	3.74	4.11	3.56	
41A	338	1.2	341	337	338	333	334	344
	2.38 ^{RT}	49.9	4.7	1.6	2.2	2.1	2.3	1.4
	2.90	4.9	2.9	2.9	3.1	3.0	2.8	2.7
41B	286	3.4	284	304	290	280	281	278
	2.92	13.4	3.4	2.9	3.4	2.6	2.6	2.6
	3.60	5.3	3.8	3.6	3.8	3.3	3.6	3.5
42	394	9.8	382	437	363			
	7.57 ^{RT}	29.9	5.2	7.8	9.7			
	5.03 ^{RT}	13.2	5.2	4.3	5.6			
43	407 ^{RT}	2.7	408	400	421	416	405	390
	4.62	5.1	4.61	4.60	4.45	4.92	4.83	4.29
	5.46 ^{RT}	5.0	5.43	5.02	5.35	5.80	5.47	5.67
44	346	1.6	345	339	349	352		
	4.53	12.1	4.14	4.59	5.28	4.10		
	4.05	4.7	4.28	3.91	3.88	4.13		
45	338	1.8	330	336	345	343	337	
	5.40 ^{RT}	17.2	3.95	6.22	4.76	5.55	6.46	5.44
	5.78 ^{RT}	5.8	6.09	6.19	5.65	5.77	5.25	5.71
46	310	3.8	322	325	309	305	304	294
	2.53	13.7	2.28	2.02	2.84	2.43	2.72	2.90
	2.76	2.3	2.73	2.68	2.78	2.85	2.74	

COPPER DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
47			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	272	3.7	266	261	282	287	267	271	
	4.18	9.6	4.71	4.21	4.38	3.69	4.35	3.72	
	2.44	8.7	2.82	2.40	2.28	2.50	2.22	2.39	
48			NO DATA						
49A			$\text{HNO}_3/\text{HClO}_4$ -ICP						
	298	4.7	308	293	316	282	290		
	4.23	23.5	4.5	5.5	3.3	3.6			
	3.25	11.6	3.2	3.0	3.0	3.8			
49B			$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS(A)-GFAAS(B/C)						
	313	0.9	310	315	312	310	312	317	
	4.40	15.7	4.5	4.5	5.4	3.5	4.1		
	3.20	16.3	3.0	2.7	2.9	2.9	4.0	3.7	
49C			HNO_3 -EP-IDSSMS						
	322	7.6	296	329	353	312			
	3.35	6.2	3.69	3.12	3.42	3.44	3.22	3.22	
	3.19	6.7	2.99	3.22	3.47	3.06			
50			HNO_3 -GFAAS						
	342	3.3	341	347	346	321	348		
	3.58	28.2	2.5	4.1	2.8	5.0	3.0		
	2.02	4.1	2.1	1.9	2.0	2.0	2.1		
51			No procedure						
	357	1.6	360	360	350				
	---	---	No data						
	2.53	13.6	2.4	2.6	2.8	1.9	2.7	2.8	

ZINC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
1			NO DATA						
2			BOMB/ $\text{HNO}_3/\text{HClO}_4$ -ICP						
	184	1.1	186	185	185	184	184	180	
	60.3	2.0	61	58	61	61	61	60	
	92.8	2.4	90	92	94	91	96	94	
3			$\text{HNO}_3/\text{HClO}_4/\text{HCl}$ -FAAS						
	165	1.1	163	164	164	167	167	167	
	56.0	5.8	62	55	57	54	53	55	
	92.2	3.6	96	93	87	90	92	95	
4			HNO_3 -FAAS						
	176	1.9	179	180	171	175	177	174	
	57.5	1.9	57.8	57	56.8	57.4	56.6	59.6	
	97.4	4.5	93.5	93.8	97.3	105.7	97.5	96.6	
5			BOMB/ HNO_3 -FAAS						
	162	1.1	162	165	161	163	160	163	
	52.6	1.1	52.3	52.5	53.2	51.7	53.1	52.8	
	82.3	1.7	82	81	82	82	82	85	

ZINC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
6			NO DATA						
7			DRY ASH/HNO ₃ /H ₂ O ₂ -FAAS						
	178	4.0	190	179	179	168	176	178	
	57.1	1.8	57.7	58.6	57.1	56.9	56.4	55.7	
	103.7	4.2	98.7	108	105	102	99.4	109	
8			HNO ₃ /H ₂ O ₂ -FAAS						
	166	1.1	169	167	165	164	167	166	
	59.7	1.7	61	58	60	60	59	60	
	96.7	2.2	95	94	98	96	97	100	
9			BOMB/HNO ₃ -FAAS						
	174	0.6	175	174	173				
	57.0	2.8	59.0	56.0	56.5				
	94.5	1.3	96.0	94.5	93.5				
10			HNO ₃ /HClO ₄ -ICP						
	210 ^{IT}	2.5	213	200	212	215	211	211	
	65.0	1.8	65.4	64.4	65.6	66.7	63.3	64.3	
	112.7	1.1	112	111	113	112	114	114	
11A			BOMB/HNO ₃ -EXTN-FAAS						
	181	3.1	180	177	186	173	185	187	
	63.2	5.3	58	62	68	64	62	65	
	105.8	7.5	111	102	101	109	95	117	
11B			BOMB/HNO ₃ -FAAS						
	182	6.7	188	168	198	185	167	188	
	63.3	7.9	62	67	63	71	57	60	
	104.5	6.1	113	102	112	98	101	101	
11C			BOMB/HNO ₃ -ICP						
	172	4.8	171	176	180	175	173	156	
	59.9	6.8	61.4	65	63.7	58.1	55.4	55.7	
	92.3	8.7	101	91	103	84	85	90	
12			BOMB/HNO ₃ /HClO ₄ -FAAS						
	179	3.7	170	188	185	175	181	178	
	58.7	1.8	59	57	59.9	58.2	59.6	58.2	
	99.1	3.3	98.3	103	98.5	96.4	95.2	103	
13			PELLET-INAA						
	194	10.5	173	221	190	217	181	180	
	76.2 ^{IT}	4.9	82	76	77	73	73		
	115.7 ^{IT}	9.4	127	122	127	106	108	104	
14			BOMB/HNO ₃ -FAAS						
	138 ^{IT}	1.9	139	135	136	138	136	142	
	49.3	2.1	48	51	50	49	49	49	
	77.0	1.4	77	76	79	77	76	77	
15			HNO ₃ /H ₂ O ₂ -FAAS						
	183	35.6	160	120	150	140	250	280	
	58.8	19.3	53	58	49	50	79	64	
	83.8	7.6	90	83	72	86	88	84	
16			BOMB/HNO ₃ -FAAS						
	194	1.2	193	192	192	197	196	192	
	67.2	2.4	68	64	68	68	68	67	
	102.7	2.0	103	100	103	103	101	106	

ZINC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
17			$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS						
	180	3.5	170	180	180	180	190	180	
	58.7	2.8	58	57	60	61	57	59	
	97.3	2.1	100	96	96	96	96	100	
18A			DRY ASH/ HNO_3 -ICP						
	166	0.6	166	166	168	166	166	165	
	56.5	0.6	56.7	56.5	56.2	57	56.4	56	
	84.5	0.6	84.3	85.5	84.6	84.6	84.1	84.1	
18B			DRY ASH/ HNO_3 -FAAS						
	146 ^{WT}	2.6	152	148	148	144	146	141	
	50.8	1.1	51.5	51.1	50.5	51.1	50	50.5	
	83.3	2.6	84	86	83.3	84	82.8	79.5	
19			DRY ASH/HCl-FAAS						
	265 ^{WT}	15.8	237	324	313	248	233	235	
	92.4 ^{WT}	3.6	98	92	92	90	90		
	141.7 ^{WT}	5.0	143	141	154	141	132	139	
20			HNO_3 -FAAS						
	147 ^{WT}	3.3	147	156	145	142	145	150	
	36.8 ^{WT}	8.5	40	37	39	36	32		
	76.0	13.9	88	76	69	80	59	84	
21			NO DATA						
22			HNO_3/HCl -FAAS						
	170	1.0	169	169	170	170	173	168	
	51.7	0.9	52.1	50.8	51.7	51.9	51.5	52	
	90.2	2.1	91.9	89.9	92	91.5	88.3	87.7	
23A			HNO_3 -FAAS-0.25 g samples						
	188	5.5	200	185	175	196	177	193	
	60.4	3.6	64.4	59.4	58.9	58.9	59.3	61.5	
	94.8	3.2	94.4	93.7	92.3	96.6	91.8	99.9	
23B			NO DATA						
24			BOMB/ $\text{HNO}_3/\text{H}_2\text{SO}_4$ -FAAS						
	162	4.1	170	163	164	167	154	154	
	48.7	3.4	49	49	49	51	46	48	
	172.5 ^{WT}	7.2	186	187	164	177	161	160	
25			$\text{HNO}_3/\text{HClO}_4$ -FAAS						
	163	6.6	162	160	155	153	165	183	
	55.6	13.1	58	56	44	56	64		
	88.7	4.1	89	89	85	84	92	93	
26			DRY ASH/ HNO_3 -DCP						
	172	3.5	178	176	175	168	174	166	
	52.8	7.5	50	59	55.8	48.2	52.3	51.8	
	95.0	2.9	90.9	93.8	94	97.1	95.8	98.6	
27A			HNO_3 -FAAS						
	195	1.9	199	193	193	200	196	191	
	63.7	1.9	65.7	62.6	63.6	63.7	62.5	64.3	
	96.0	3.0	98.4	97.4	99.7	94	92.7	93.8	
27B			NO DATA						
28			HNO_3 -GFAAS						
	110 ^{WT}	5.6	104	105	112	110	108	121	
	48.0	10.7	57.4	48.1	47.1	43.3	43.6	48.3	
	76.5	5.4	71.8	73.5	75.7	75.9	78.9	83.4	

ZINC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
29			$\text{HNO}_3\text{-FAAS}$						
	319 ^{RT}	6.5	314	335	334	280	329	321	
	79.9 ^{RT}	6.0	83.6	76.9	82.5	82	71.4	82.9	
	151.3 ^{RT}	4.7	161	149	155	156	144	143	
30			DRY ASH/HCl/ $\text{HNO}_3\text{-FAAS}$						
	154 ^{RT}	6.0	145	140	157	160	161	162	
	48.4	1.4	48.7	48.9	48.6	49.1	47.4	47.8	
	81.9	5.5	84.6	83.7	88.7	78.9	77.8	77.4	
31			DRY ASH/ $\text{HNO}_3\text{/HCl-FAAS}$						
	162	2.9	167	159	159				
	55.7	1.7	55.2	56.8	55.1				
	90.4	0.9	89.7	91.3	90.1				
32			BOMB/ $\text{HNO}_3\text{-HCl-GFAAS}$						
	198	6.2	190	192	222	196	198	189	
	59.2	2.8	62	59	59	58	58		
	99.7	10.5	101	118	103	89	91	96	
33			$\text{HNO}_3\text{/HClO}_4\text{-FAAS}$						
	182	0.9	180	182	184	182	183	180	
	61.7	1.3	62	62	60	62	62	62	
	97.3	1.8	100	96	96	97	99	96	
34			NO DATA						
35			$\text{HNO}_3\text{/HClO}_4\text{-XRF}$						
	131 ^{RT}	16.6	132	137	116	160	144	98	
	----	----	No data						
	----	----	No data						
36			$\text{HNO}_3\text{/H}_2\text{SO}_4\text{/H}_2\text{O}_2\text{-EXTN-GFAAS}$						
	----	----	No data						
	57.6	1.8	56.8	59.5	57.6	56.6	57.6	57.4	
	----	----	No data						
37			NO DATA-AAS						
	185	3.4	188	185	195	176	183	186	
	67.6	6.9	67.6	67.8	70.1	74.7	63.9	61.4	
	104.1	3.5	103.8	103.4	97.4	105.6	106.9	107.2	
38			$\text{HNO}_3\text{/H}_2\text{O}_2\text{-FAAS}$						
	104 ^{RT}	11.7	117	110	90.9	96.2			
	64.3	2.2	62.7	63.1	64.3	64.2	64.9	66.7	
	101.5	4.4	99.9	107.6	104.5	99	96.4		
39			$\text{HNO}_3\text{/H}_2\text{O}_2\text{-FAAS}$						
	174	2.3	177	180	172	173	172	169	
	----	----	No data						
	86.9	5.9	84.4	96.5	85.8	85.4	88	81.5	
40			$\text{HNO}_3\text{/H}_2\text{SO}_4\text{/H}_2\text{O}_2\text{-EXTN-FAAS}$						
	180	3.4	188	178	180	184	172		
	64.5	10.1	66.5	58.9	60.3	61.2	76.6	63.2	
	111	1.3	110	110	113	111			
41A			BOMB/ $\text{HNO}_3\text{-GFAAS}$						
	183	9.7	183	172	206	163	173	204	
	60.8	2.6	60.2	60	60.6	63.6	61.4	59.1	
	91.0	10.6	86	83	89	110	88	90	

ZINC DATA

ARSENIC DATA

ARSENIC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
17			NO DATA						
18			HNO ₃ -HG-AAS						
	15.9	5.3	15.9	15.3	15.3	15.1	16.3	17.3	
	3.6	7.1	4.0	3.4	3.6	3.5	3.9	3.4	
	1.9	13.4	2.2	1.9	1.9	2.2	1.9	1.5	
19			DRY ASH/HCl/KI-HG-AAS						
	18.8	3.9	17.5	19.1	19.6	18.5	19.1	19.1	
	3.4	9.6	2.9	3.5	3.7	3.0	3.5	3.5	
	2.6	9.2	2.6	2.3	2.9	2.5	2.7	2.3	
20			NO DATA						
21			NO DATA						
22			NO PREP-INAA						
	25.0	---	25.0						
	7.0	---	7.0						
	4.6	---	4.6						
23A			NO DATA						
23B			NO DATA						
24			BOMB/HNO ₃ /H ₂ SO ₄ -GFAAS						
	21.2	4.3	22.2	21.0	20.4				
	13.0 ^{RT}	5.5	13.1	13.6	12.2				
	5.9	6.9	6.1	5.4	6.1				
25			NO DATA						
26			NO DATA						
27A			HNO ₃ /H ₂ SO ₄ -HG-AAS						
	24.7	7.7	24.1	28.2	25.6	23.5	23.2	23.8	
	7.7	6.4	7.2	7.2	8.5	7.7	8.0	7.8	
	5.2	8.9	5.1	5.8	5.7	5.0	4.8	4.7	
27B			NO DATA						
28			NO DATA						
29			NO DATA						
30			NO DATA						
31			DRY ASH/HNO ₃ /HCl-HG-AAS						
	36.3	8.6	37.6	38.5	32.7				
	9.7	5.6	9.48	10.3	9.28				
	6.3	7.6	6.84	6.10	5.95				
32			DRY ASH/HNO ₃ -HG-SPEC						
	29.5	6.6	29.3	28.6	29.3	28.6	27.8	33.3	
	7.6	10.6	7.2	6.4	7.9	7.9	8.5		
	6.5	23.9	4.5	6.2	8.5	7.4	5.7		
33			NO DATA						
34			NO DATA						
35			NO DATA						
36			NO DATA						
37			HNO ₃ -Ni/GFAAS						
	26.4	3.9	28.0	25.5	27.0	25.9	26.6	25.3	
	6.8	4.2	6.45	6.77	6.38	6.91	6.94	7.09	
	3.3	2.6	3.17	3.38	3.35	3.37	3.31	3.42	

ARSENIC DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
38			NO DATA						
39			NO DATA						
40			$\text{HNO}_3/\text{H}_2\text{SO}_4/\text{H}_2\text{O}_2$ -HG-AAS						
	14.2	23.3	10.7	17.4	13.1	16.8	17.1	10.2	
	2.7	13.7	2.32	2.58	2.42	3.21	2.72	3.16	
	2.6	17.4	2.58	2.15	2.15	2.39	3.12	3.14	
41			NO DATA						
42			NO DATA						
43			NO DATA						
44			NO DATA						
45			NO DATA						
46			NO DATA						
47			NO DATA						
48			NO DATA						
49A			$\text{HNO}_3/\text{H}_2\text{O}_2$ -Ni/GFAAS						
	25.7	8.0	27.1	24.1	22.8	25.2	28.5	26.2	
	7.5	3.6	7.07	7.79	7.51	7.40	7.78	7.41	
	4.7	9.2	4.2	5.4	4.6	4.6	5.1	4.5	
49B			$\text{HNO}_3/\text{H}_2\text{SO}_4$ -HG-AAS						
	23.0	3.8	21.9	22.7	23.2	24			
	7.7	3.8	7.40	7.58	7.81	8.08			
	5.1	3.9	4.90	5.20	5.36	5.04			
49C			DRY ASH/MG(NO_3) ₂ -GC						
	25.9	8.4	28.2	23.1	25.2	25.0	28.0		
	7.7	3.7	7.62	7.47	7.75	8.17	7.48		
	3.2	15.0	3.06	3.42	3.97	2.66	3.11		
50			NO DATA						
51			NO DATA						

CADMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
1			HNO_3 -GFAAS						
	22.4	3.8	22.1	22.6	23.4	22.1	21.0	23.1	
	1.19 ⁿ	5.8	1.17	1.18	1.18	1.26	1.27	1.08	
	0.06	14.4	0.06	0.06	0.05	0.07	0.05	0.05	
2A			BOMB/ $\text{HNO}_3/\text{HClO}_4$ -ICP						
	27.2	2.0	27.8	26.7	26.8	27.3	27.8	26.6	
	0.71	5.1	0.73	0.77	0.71	0.71	0.67	0.68	
	0.32 ⁿ	11.7	0.39	0.33	0.31	0.29	0.29	0.31	
2B	-----	-----	BOMB/ $\text{HNO}_3/\text{HClO}_4$ -GFAAS						
			No data						
	0.66	3.0	0.69	0.65	0.65	0.65	0.64		
	0.33 ⁿ	8.0	0.33	0.37	0.33	0.33	0.35	0.29	

CADMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
3			$\text{HNO}_3/\text{HClO}_4/\text{HCl}$ -FAAS					
	25.3	2.3	26.2	24.6	25.8	25.0	25.1	25.3
	0.78	2.9	0.81	0.79	0.77	0.75	0.77	0.80
	0.11	11.5	0.11	0.11	0.11	0.09	0.11	0.13
4			HNO_3 -GFAAS					
	25.5	1.7	25.8	25.5	26.2	25.4	24.9	25.4
	0.38 ^{RT}	6.1	0.38	0.35	0.39	0.36	0.41	0.40
	0.04	12.1	0.036	0.028	0.039	0.039	0.040	0.036
5			BOMB/ HNO_3 -FAAS(A)-GFAAS(B/C)					
	26.9	1.3	27.3	26.9	26.8	27.4	26.7	26.5
	0.87	4.8	0.85	0.88	0.85	0.92	0.81	0.91
	0.07	25.3	0.05	0.05	0.06	0.06	0.09	0.08
6			NO DATA					
7			DRY ASH/ $\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS(A)-GFAAS(B/C)					
	22.5	7.2	21.6	24.5	20.4	23.7	21.1	23.6
	0.67	5.2	0.65	0.68	0.66	0.72	0.69	0.62
	0.07	4.8	0.069	0.070	0.063	0.065	0.062	0.066
8			$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS(A)-GFAAS(B/C)					
	25.5	0.7	25.7	25.4	25.4	25.6	25.7	25.3
	0.76	7.3	0.78	0.72	0.86	0.76	0.76	0.70
	0.04	18.0	0.052	0.036	0.033	0.044	0.041	
9			O_2 COMBUSTION- HNO_3 -GFAAS					
	29.1	6.8	27.8	26.3	31.3	28.1	30.3	30.8
	0.74	5.3	0.73	0.78	0.77	0.74	0.68	
	0.04	60.4	0.01	0.04	0.07	0.06	0.02	0.03
10			$\text{HNO}_3/\text{HClO}_4$ -ICP					
	36.3 ^{RT}	6.3	36.3	32.7	38.1	38.9	37.1	34.6
	0.89	4.1	0.859	0.924	0.929	0.906	0.838	0.882
	0.11	18.6	0.108	0.095	0.140	0.094	0.088	0.123
11			BOMB/ HNO_3 -EXTN-GFAAS					
	25.1	3.0	25.2	25.4	25.2	24.3	24.3	26.3
	0.41 ^{RT}	6.7	0.414	0.448	0.426	0.426	0.402	0.366
	0.02	17.5	0.018	0.015	0.023	0.021	0.025	0.020
12			BOMB/ $\text{HNO}_3/\text{HClO}_4$ -GFAAS					
	33.1 ^{RT}	9.5	38.3	33.3	29.7	31.8	35.0	30.7
	0.83	11.2	0.88	0.97	0.89	0.76	0.75	0.75
	0.06	13.8	0.064	0.062	0.046	0.055	0.070	0.061
13			NO DATA					
14			BOMB/ HNO_3 -GFAAS					
	8.6 ^{RT}	7.9	9.1	7.9	9.3	9.1	7.7	8.6
	0.78	18.8	0.9	0.8	1.0	0.6	0.7	0.7
	0.07	16.3	0.09	0.07	0.07	0.08	0.06	0.06
15			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS					
	13.9 ^{RT}	48.0	20.0	12.0	19.0	5.2	7.3	20.0
	0.56	37.4	0.81	0.33	0.70	0.37	0.42	0.73
	0.05	66.3	0.075	0.031	0.061	0.088	0.016	0.014
16			BOMB/ HNO_3 -FAAS					
	30.9 ^{RT}	1.7	30.2	30.3	30.9	31.4	31.4	31.2
	----	----	No data					
	----	----	No data					

CADMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
17			$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS						
	30.0	2.1	30	29	30	30	30	31	
	0.6	40.8	0.5	0.5	0.5	0.5	0.5	1.1	
	----	----	Less than 0.5						
18A			DRY ASH/ HNO_3 -ICP						
	26.4	0.5	26.6	26.5	26.5	26.4	26.4	26.2	
	0.78	6.1	0.72	0.81	0.79	0.71	0.80	0.82	
	----	----	Less than 0.2						
18B			DRY ASH/ HNO_3 -EXTN(B/C)-FAAS						
	26.7	1.4	27.1	26.3	26.6	26.8	27.2	26.3	
	0.73	2.5	0.70	0.75	0.74	0.72	0.74	0.72	
	0.06	12.9	0.07	0.06	0.07	0.07	0.06	0.05	
19			DRY ASH/HCl-FAAS						
	25.2	3.8	26.1	24.6	23.5	25.5	25.6	25.6	
	0.96	4.7	0.94	0.94	0.94	1.05	0.94	0.94	
	0.55 ^{RT}	0	0.55	0.55	0.55	0.55	0.55	0.55	
20			HNO_3 -GFAAS						
	11.0 ^{RT}	16.7	11.18	13.72	12.61	9.30	9.16	10.19	
	0.18 ^{RT}	28.9	0.17	0.15	0.12	0.25	0.22		
	0.09	28.0	0.10	0.08	0.12	0.12	0.06	0.07	
21			BOMB/ $\text{HNO}_3/\text{H}_2\text{SO}_4$ (A)- $\text{HNO}_3/\text{HClO}_4$ (B/C)-ASV						
	25.0	45.8	30	28	40	14	13		
	----	----	No data						
	----	----	No data						
22			HNO_3/HCl -FAAS(A)/GFAAS(B/C)						
	26.0	2.8	26.9	25.1	26.4	25.3	26.5	25.8	
	0.93	5.3	0.99	0.88	0.93	0.87	0.98	0.93	
	0.05	18.2	0.043	0.033	0.053	0.054	0.050	0.040	
23A			HNO_3 -FAAS-0.25 g samples						
	25.0	1.2	24.4	25.0	25.3	25.0	25.1	25.1	
	----	----	Less than 0.4						
	----	----	Less than 0.4						
23B			HNO_3 -FAAS-1 g samples						
	----	----	No data						
	0.70	0	0.7	0.7	0.7	0.7	0.7		
	----	----	Less than 0.1						
24			BOMB/ $\text{HNO}_3/\text{H}_2\text{SO}_4$ -GFAAS						
	23.2	1.8	22.8	23.3	22.6	23.6	23.6	23.3	
	0.73	10.0	0.63	0.73	0.65	0.79	0.74	0.81	
	0.08	21.1	0.08	0.09	0.07	0.06	0.11	0.08	
25			$\text{HNO}_3/\text{HClO}_4$ -FAAS						
	24.8	5.6	26.0	25.5	23.7	23.1	23.8	26.4	
	0.83	9.8	0.85	0.95	0.79	0.84	0.73		
	0.11	41.1	0.18	0.15	0.09	0.10	0.07	0.07	
26			DRY ASH/ HNO_3 -DCP(A)-GFAAS(B/C)						
	25.1	4.7	25.7	26.9	24.0	23.6	25.2	25.0	
	0.68	19.4	0.57	0.66	0.90	0.53	0.75	0.70	
	0.07	29.8	0.06	0.07	0.11	0.06	0.05	0.08	

CADMIUM DATA

CADMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
40			$\text{HNO}_3/\text{H}_2\text{SO}_4/\text{H}_2\text{O}_2$ -EXTN-FAAS						
	26.6	9.0	27.5	22.9	24.3	28.1	28.7	28.1	
	0.76	9.5	0.634	0.768	0.725	0.834	0.784	0.811	
	0.07	30.0	0.056	0.089	0.044	0.073			
41A			BOMB/ HNO_3 -GFAAS						
	25.2	3.1	25.8	25.0	26.0	24.7	23.9	25.6	
	1.05 ^{RT}	2.6	1.05	1.07	1.08	1.01	1.02	1.05	
	0.04	12.3	0.045	0.047	0.045	0.043	0.037	0.034	
41B			HNO_3 -FAAS						
	22.5	1.1	22.5	22.7	22.6	22.2	22.7	22.1	
	0.67	9.2	0.68	0.76	0.60	0.69	0.60	0.69	
	----	----	No data						
42			HNO_3 -DRY ASH/ HNO_3/HCl -ASV						
	16.2 ^{RT}	14.7	19.6	16.0	15.0	14.2			
	0.60	10.1	0.63	0.64	0.53				
	0.06	14.8	0.057	0.066	0.046	0.054			
43			DRY ASH/ HNO_3 -FAAS						
	17.1 ^{RT}	3.1	16.3	17.0	17.7	17.7	16.9	17.0	
	1.13 ^{RT}	5.8	1.08	1.23	1.08	1.13	1.18	1.07	
	0.53 ^{RT}	7.6	0.59	0.56	0.48	0.50	0.54	0.52	
44			DRY ASH/ HNO_3 -FAAS						
	28.6	21.8	38.0	25.5	25.3	25.7			
	0.75	18.1	0.77	0.76	0.90	0.57			
	----	----	No data						
45			BOMB/ $\text{HNO}_3/\text{HCl}/\text{H}_2\text{O}_2$ -FAAS						
	26.5	3.0	26.8	27.5	25.3	26.4	26.7		
	1.14 ^{RT}	1.6	1.12	1.11	1.15	1.14	1.16	1.14	
	0.80 ^{RT}	5.4	0.77	0.75	0.81	0.84	0.85	0.76	
46			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	21.4	6.6	23.3	22.5	20.9	20.4	20.0		
	0.37 ^{RT}	6.8	0.357	0.357	0.408	0.359			
	0.04	34.8	0.048	0.023	0.028	0.046			
47			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	21.2	4.9	21.6	19.3	21.3	21.2	22.5	21.2	
	0.74	3.2	0.72	0.72	0.72	0.76	0.77	0.72	
	0.07	8.4	0.06	0.07	0.07	0.06	0.07	0.06	
48			NO DATA						
49A			$\text{HNO}_3/\text{HClO}_4$ -ICP						
	27.5	4.3	26.8	28.6	25.9	26.9	27.5	29.0	
	0.80	12.5	0.8	0.9	0.7				
	----	----	Less than 0.1						
49B			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	29.3	8.1	26.1	27.9	30.1	29.8	32.4		
	0.89	13.7	0.70	0.79	1.00	0.97	0.91	0.99	
	0.08	11.9	0.074	0.066	0.089	0.083	0.072		
49C			HNO_3 -EP-IDSSMS						
	22.8	6.6	25.5	22.7	22.8	21.0	23.0	21.9	
	0.66	2.4	0.659	0.691	0.651	0.668	0.648	0.655	
	0.06	4.4	0.055	0.062	0.058	0.057	0.059		

CADMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
50			HNO ₃ -GFAAS					
	25.8	4.2	26	26	26	24	27	
	0.43 ^{RT}	11.7	0.44	0.46	0.40	0.50	0.37	
	0.03 ^{RT}	9.7	0.030	0.033	0.037	0.029	0.032	
51			No procedure					
	26.0	---	27	25				
	----	----	No data					
	0.05	30.6	0.04	0.05	0.05	0.02	0.05	0.06

MERCURY DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
1			NO DATA					
2			BOMB/HNO ₃ /HClO ₄ -CVAAS					
	0.180	5.5	0.192	0.171	0.192	0.178	0.171	0.174
	0.072	3.2	0.072	0.070	0.076	0.071	0.070	0.073
	0.035 ^{RT}	10.8	0.033	0.031	0.034	0.035	0.034	0.042
3			NO DATA					
4			HNO ₃ -CVAAS					
	0.188	14.4	0.18	0.18	0.18	0.19	0.16	0.24
	0.075	27.6	0.07	0.05	0.10	0.1	0.07	0.06
	0.047	26.0	0.05	0.04	0.06	0.04	0.03	0.06
5			BOMB/HNO ₃ -CVAAS					
	0.215	10.9	0.22	0.22	0.22	0.22	0.24	0.17
	0.040 ^{RT}	27.4	0.04	0.04	0.06	0.04	0.03	0.03
	0.057	18.2	0.04	0.06	0.06	0.05	0.06	0.07
6			NO PREP-INAA					
	----	----	No data					
	----	----	No data					
	0.062	4.2	0.0629	0.0628	0.0569	0.0635	0.0609	0.0639
7			H ₂ SO ₄ /H ₂ O ₂ -CVAAS					
	0.220	3.5	0.221	0.222	0.215	0.232	0.212	
	0.064	4.0	0.062	0.062	0.066	0.064	0.068	
	0.056	2.7	0.057	0.056	0.054	0.058	0.057	
8			HNO ₃ -CVAAS					
	0.252	5.3	0.26	0.26	0.26	0.24	0.26	0.23
	0.097	12.5	0.10	0.12	0.09	0.09	0.09	0.09
	0.060	21.1	0.08	0.06	0.05	0.05	0.05	0.07
9			O ₂ COMBUSTION-HNO ₃ -CVAAS					
	0.305	6.5	0.30	0.28	0.34	0.30	0.30	0.31
	0.102	8.5	0.091	0.107	0.092	0.105	0.103	0.113
	0.065	3.7	0.066	0.064	0.063	0.062	0.068	
10			NO DATA					
11			BOMB/HNO ₃ -CVAAS					
	0.199	5.5	0.207	0.214	0.201	0.186	0.200	0.187
	0.084	6.0	0.087	0.081	0.090	0.081	0.077	0.088
	0.058	7.1	0.053	0.064	0.056	0.061	0.055	0.057

MERCURY DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
12			$\text{HNO}_3/\text{H}_2\text{SO}_4/\text{H}_2\text{O}_2$ -CVAAS					
	0.263	2.0	0.26	0.27	0.26	0.26	0.26	0.27
	0.075	10.5	0.064	0.066	0.081	0.077	0.077	0.083
	0.058	9.2	0.053	0.062	0.066	0.057	0.052	0.058
13			NO DATA					
14			BOMB/ $\text{HNO}_3/\text{KMNO}_4$ -CVAAS					
	0.243	0.5	0.242	0.242	0.245	0.243	0.242	0.244
	0.064	14.1	0.060	0.056	0.076	0.053	0.068	0.071
	0.056	6.9	0.051	0.058	0.062	0.055	0.056	0.053
15			NO DATA					
16			$\text{H}_2\text{SO}_4/\text{HNO}_3$ -CVAAS					
	0.269	7.8	0.272	0.279	0.253	0.285	0.235	0.290
	0.111 ^N	6.5	0.115	0.104	0.109	0.111	0.123	0.104
	0.059	1.4	0.058	0.058	0.059	0.059	0.058	0.060
17			$\text{HNO}_3/\text{H}_2\text{O}_2$ -CVAAS					
	0.245	9.2	0.25	0.22	0.26	0.23	0.23	0.28
	0.08	25.0	0.11	0.09	0.05	0.08	0.08	0.07
	----	----	Less than 0.05					
18			HNO_3 -CVAAS					
	0.272	4.6	0.260	0.272	0.281	0.28	0.285	0.254
	0.094	5.9	0.104	0.090	0.090	0.098	0.092	0.092
	0.047	6.8	0.045	0.046	0.049	0.051	0.042	0.048
19			$\text{HNO}_3/\text{HClO}_4$ -CVAAS					
	0.169	10.7	0.156	0.168	0.204	0.157	0.159	0.170
	----	----	Less than 0.01					
	----	----	Less than 0.01					
20			NO DATA					
21A			BOMB/ $\text{H}_2\text{SO}_4/\text{HNO}_3$ -ASV					
	0.352	40.7	0.229	0.373	0.481	0.181	0.549	0.300
	----	----	No data					
	----	----	No data					
21B			BOMB/ HNO_3 -CVAAS					
	0.268	6.7	0.274	0.291	0.272	0.275	0.251	0.242
	0.085	14.8	0.076	0.091	0.068	0.098	0.093	
	0.055	6.5	0.057	0.052	0.055	0.055	0.050	0.060
22			$\text{HNO}_3/\text{H}_2\text{SO}_4/\text{KMNO}_4/\text{K}_2\text{S}_2\text{O}_8$ -CVAAS					
	0.251	2.9	0.246	0.253	0.247	0.245	0.252	0.260
	0.076	12.3	0.079	0.074	0.068	0.092	0.078	0.066
	0.045	4.7	0.047	0.042	0.047	0.044	0.044	0.047
23A			$\text{H}_2\text{SO}_4/\text{KMNO}_4$ -CVAAS					
	0.203	16.7	0.224	0.238	0.228	0.169	0.206	0.155
	0.079	16.3	0.090	0.076	0.095	0.083	0.061	0.069
	0.048	45.5	0.031	0.043	0.051	0.030	0.090	0.045
23B			NO DATA					
24			$\text{H}_2\text{SO}_4/\text{HNO}_3/\text{KMNO}_4$ -CVAAS					
	0.212	6.3	0.20	0.22	0.20	0.20	0.23	0.22
	0.062	15.9	0.05	0.07	0.06	0.05	0.07	0.07
	0.038	19.6	0.03	0.05	0.03	0.04	0.04	0.04

MERCURY DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
25			O2-H ₂ SO ₄ /KMNO ₄ -CVAAS					
	0.172	14.5	0.13	0.16	0.20	0.19	0.18	0.17
	0.070	20.8	0.093	0.076	0.068	0.054	0.072	0.055
	0.060	18.8	0.068	0.055	0.079	0.057	0.054	0.048
26			NO DATA					
27			O2-H ₂ SO ₄ /KMNO ₄ -CVAAS					
	0.276	4.1	0.27	0.28	0.29	0.28	0.26	
	0.092	18.8	0.11	0.11	0.07	0.08	0.10	0.08
	0.052	28.5	0.08	0.05	0.05	0.04	0.05	0.04
28			H ₂ SO ₄ /HNO ₃ -CVAAS					
	0.380 ^{IT}	2.5	0.377	0.371	0.383	0.393	0.387	0.369
	0.142 ^{IT}	1.6	0.139	0.140	0.142	0.144	0.144	
	0.116 ^{IT}	9.8	0.122	0.116	0.099	0.120	0.130	0.106
29			H ₂ SO ₄ /HNO ₃ -CVAAS					
	0.218	20.2	0.190	0.190	0.189	0.188	0.283	0.267
	0.097	2.5	0.098	0.096	0.100	0.097	0.095	0.093
	0.048	28.3	0.051	0.044	0.067	0.058	0.031	0.036
30			No procedure					
	0.302	---	0.298	0.305				
	0.104	---	0.103	0.105				
	0.076	---	0.073	0.078				
31			BOMB/HNO ₃ -CVAAS					
	0.307	8.3	0.287	0.336	0.299			
	0.076	16.2	0.065	0.089	0.073			
	0.082 ^{IT}	3.1	0.080	0.085	0.082			
32			BOMB/HNO ₃ -CVAAS					
	0.273	10.5	0.29	0.32	0.24	0.25	0.27	0.27
	0.077	17.5	0.055	0.088	0.088	0.077	0.077	
	0.057	31.5	0.033	0.089	0.055	0.055	0.055	0.055
33			HNO ₃ /H ₂ SO ₄ /V ₂ O ₅ -Au-CVAAS					
	0.264	5.0	0.286	0.264	0.254	0.267	0.247	0.264
	0.099	11.0	0.094	0.110	0.081	0.098	0.101	0.11
	0.070	5.9	0.063	0.071	0.068	0.073	0.073	0.074
34			HNO ₃ /H ₂ O ₂ -CVAAS					
	0.169	2.1	0.169	0.167	0.165	0.170	0.175	0.166
	0.077	2.4	0.076	0.074	0.075	0.079	0.077	0.078
	0.055	2.7	0.054	0.055	0.055	0.055	0.058	0.054
35			NO DATA					
36			HNO ₃ /H ₂ SO ₄ -CVAAS					
	0.301	9.3	0.348	0.272	0.281	0.29	0.318	0.296
	0.073	9.1	0.0703	0.0783	0.0716	0.0639	0.0718	0.0829
	0.070	23.1	0.0995	0.0678	0.0683	0.0575	0.0539	0.0757
37			HNO ₃ /H ₂ SO ₄ -CVAAS					
	0.344	1.8	0.340	0.346	0.354	0.336	0.341	0.346
	0.093	6.0	0.082	0.096	0.092	0.096	0.096	0.096
	0.046	8.9	0.046	0.046	0.040	0.046	0.053	0.046
38			NO DATA					

MERCURY DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
39			NO DATA					
40			HNO ₃ /H ₂ SO ₄ -CVAAS					
	0.269	3.6	0.261	0.279	0.279	0.258	0.273	0.261
	0.086	8.8	0.085	0.085	0.097	0.075	0.081	0.090
	0.054	16.1	0.066	0.063	0.053	0.049	0.043	0.051
41			NO DATA					
42			COLD STANNITE RED/Au-CVAAS					
	0.218	2.9	0.214	0.218	0.220	0.209	0.228	0.216
	0.061	3.3	0.062	0.061	0.062	0.064	0.058	0.060
	0.024 ^{RT}	11.0	0.024	0.023	0.028	0.024	0.021	0.021
43			HNO ₃ /HClO ₄ -Au-CVAAS					
	0.261	5.2	0.245	0.273	0.250	0.280	0.262	0.255
	0.076	9.4	0.078	0.070	0.070	0.089	0.073	0.076
	0.113 ^{RT}	7.4	0.113	0.102	0.124	0.110	0.107	0.121
44			NO DATA					
45			BOMB/HNO ₃ /HF-H ₂ SO ₄ /KMNO ₄ -CVAAS					
	0.311	3.4	0.323	0.313	0.319	0.305	0.312	0.293
	0.117 ^{RT}	6.6	0.116	0.127	0.113	0.125	0.118	0.106
	0.076	6.8	0.071	0.084	0.074	0.076	0.070	0.078
46			No procedure					
	0.293	14.7	0.234	0.308	0.262	0.291	0.36	0.3
	0.082	8.9	0.084	0.078	0.079	0.088	0.092	0.072
	0.046	7.0	0.044	0.043	0.043	0.046	0.048	0.051
47			NO DATA					
48			BOMB/HNO ₃ -CVAAS					
	0.265	7.3	0.302	0.257	0.268	0.261	0.259	0.245
	0.079	5.2	0.076	0.080	0.074	0.086	0.079	0.078
	0.059	8.6	0.058	0.066	0.061	0.059	0.051	0.056
49A			HNO ₃ /HClO ₄ -CVAAS					
	0.307	3.5	0.308	0.318	0.313	0.315	0.293	0.295
	0.079	6.9	0.083	0.072	0.085	0.078	0.075	
	0.051	12.9	0.052	0.062	0.05	0.047	0.045	
49B			NO DATA					
49C			HNO ₃ /HClO ₄ /H ₂ SO ₄ -EP-IDSSMS					
	0.295	6.7	0.284	0.321	0.311	0.273	0.288	
	0.098	6.9	0.096	0.106	0.093			
	0.064	---	0.063	0.064				
50			HNO ₃ /H ₂ SO ₄ -CVAAS					
	0.433 ^{RT}	4.3	0.43	0.42	0.41	0.46	0.43	0.45
	0.197 ^{RT}	5.1	0.21	0.20	0.20	0.18	0.19	0.20
	0.178 ^{RT}	4.5	0.17	0.17	0.19	0.18	0.19	0.20
51			NO DATA					

LEAD DATA

Lab	Mean	RSD							
1			HNO ₃ -GFAAS						
	3.00	25.4	2.31	3.38	4.02	1.94	3.34	3.01	
	-----	-----	Less than 0.5						
	-----	-----	No data						
2			BOMB/HNO ₃ /HClO ₄ -GFAAS						
	3.53	8.5	3.2	4.0	3.4	3.8	3.4	3.4	
	1.07 ^{RT}	7.7	1.1	1.1	1.2	1.0	1.0	1.0	
	2.37	10.6	2.1	2.3	2.4	2.7	2.6	2.1	
3			HNO ₃ /HClO ₄ /HCl-GFAAS						
	6.35	11.4	7.0	5.6	6.9	5.9	7.1	5.6	
	0.64 ^{RT}	1.6	0.62	0.64	0.64	0.65	0.64	0.63	
	5.80 ^{RT}	21.5	4.7	6.9	4.3	5.1	6.6	7.2	
4			HNO ₃ -GFAAS						
	1.33 ^{RT}	34.9	1.54	2.14	0.95	1.37	1.10	0.90	
	0.08	17.1	0.076	0.065	0.093	0.075	0.099		
	0.60	19.4	0.61	0.64	0.66	0.46	0.75	0.46	
5			BOMB/HNO ₃ -GFAAS						
	3.42	36.0	1.1	3.5	4.8	3.7	3.6	3.8	
	0.78 ^{RT}	5.7	0.8	0.8	0.8	0.8	0.7		
	1.80	22.8	2.4	1.3	1.5	2.1	1.9	1.6	
6			NO DATA						
7			DRY ASH/HNO ₃ /H ₂ O ₂ -GFAAS						
	8.44	2.1	8.48	8.23	8.49	8.25	8.45	8.72	
	0.32	5.7	0.30	0.30	0.34	0.33	0.32	0.34	
	3.54	7.1	3.14	3.56	3.37	3.59	3.78	3.80	
8			HNO ₃ /H ₂ O ₂ -GFAAS						
	5.54	16.1	5.0	6.1	6.8	4.6	5.2		
	0.45	17.7	0.56	0.34	0.46	0.42	0.46		
	1.60	13.3	1.6	1.9	1.4	1.7	1.4		
9			O ₂ COMBUSTION-HNO ₃ -GFAAS						
	6.05	14.5	7.4	6.3	5.5	6.5	5.7	4.9	
	0.30	30.9	0.45	0.22	0.27	0.36	0.20	0.31	
	2.74	51.3	2.7	3.1	1.6	1.4	4.9		
10			HNO ₃ /HClO ₄ -ICP						
	6.45	20.2	6.92	6.51	7.95	7.52	4.95	4.83	
	0.51	13.8	0.566	0.392	0.517	0.463	0.581	0.526	
	1.43	38.8	1.96	1.05	1.63	1.05	0.76	2.12	
11			BOMB/HNO ₃ -EXTN-GFAAS						
	3.80	6.9	4.1	3.6	4.0	3.8	3.4	3.9	
	0.14	7.7	0.133	0.127	0.151	0.144	0.140	0.156	
	0.88	8.0	0.937	0.853	0.844	0.807	0.843	0.992	
12			BOMB/HNO ₃ /HClO ₄ -GFAAS						
	7.94	13.8	9.41	6.14	8.06	7.4	8.41	8.2	
	0.28	26.2	0.30	0.19	0.29	0.25	0.41	0.25	
	1.43	21.9	1.51	1.98	1.46	1.14	1.36	1.13	
13			NO DATA						
14			BOMB/HNO ₃ -GFAAS						
	2.83	4.8	2.8	3.0	2.8	3.0	2.7	2.7	
	0.45	12.1	0.35	0.49	0.43	0.44	0.50	0.46	
	1.60	17.2	2.0	1.2	1.7	1.6	1.4	1.7	

LEAD DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
15			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS					
	7.26	30.0	3.5	8.6	7.7	7.6	8.9	
	0.31	31.3	0.18	0.28	0.32	0.31	0.48	0.29
	1.45	28.4	0.82	1.4	2.1	1.4	1.4	1.6
16			NO DATA					
17			$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS					
	16.8 ^{RT}	12.7	15	16	16	17	16	21
	8.70 ^{RT}	9.3	9.4	9.4	8.0	8.0	7.9	9.5
	7.25 ^{RT}	17.9	7.3	5.8	8.7	8.7	7.2	5.8
18A			DRY ASH/ HNO_3 -ICP					
	5.86	10.1	6.6	5.5	5.3	5.5	6.4	
	----	----	Less than 1					
	3.28	9.5	3.1	3.0	3.8	3.3	3.2	
18B			DRY ASH/ HNO_3 -EXTN-FAAS					
	5.03	9.9	5.36	5.08	4.60	4.47	5.64	
	0.18	18.5	0.15	0.15	0.19	0.19	0.15	0.23
	1.82	17.6	1.47	1.52	1.87	2.2	2.05	
19			DRY ASH/HCl-FAAS					
	10.6 ^{RT}	7.9	10.6	10.2	12.0	9.5	10.5	11.0
	3.48 ^{RT}	5.9	3.67	3.67	3.14	3.56	3.35	3.46
	17.4 ^{RT}	6.4	18.3	17.5	18.8	17.4	16.0	16.2
20			HNO_3 -GFAAS					
	5.58	32.4	4.3	4.0	4.1	7.5	8.1	5.5
	2.26 ^{RT}	51.1	3.5	1.2	1.3	3.5	1.8	
	1.90	16.0	2.1	1.5	2.3	1.9	1.6	2
21			BOMB/ $\text{HNO}_3/\text{H}_2\text{SO}_4$ (A)- $\text{HNO}_3/\text{HClO}_4$ (B/C)-ASV					
	40.0 ^{RT}	58.3	14	56	63	27		
	14.3 ^{RT}	23.2	14	18	15	10		
	8.67 ^{RT}	37.1	5	10	11			
22			HNO_3/HCl -GFAAS					
	0.64 ^{RT}	8.6	0.7	0.6	0.6	0.7	0.6	
	----	----	No data					
	0.95	74.7	0.1	0.9	1.5	0.2	1.9	1.1
23A			HNO_3 -FAAS-0.25 g SAMPLES					
	5.48	22.5	5.3	7.8	4.8	5.8	4.8	4.4
	----	----	Less than 2					
	----	----	Less than 2					
23B			HNO_3 -FAAS-1 g SAMPLES					
	----	----	No data					
	----	----	Less than 0.5					
	----	----	Less than 0.5					
24			BOMB/ $\text{HNO}_3/\text{H}_2\text{SO}_4$ -GFAAS					
	4.65	8.7	4.8	4.2	5.1	4.3	4.4	5.1
	0.73 ^{RT}	20.5	0.7	0.5	0.7	0.9	0.9	0.7
	1.50	11.9	1.5	1.3	1.8	1.4	1.6	1.4
25			$\text{HNO}_3/\text{HClO}_4$ -FAAS					
	8.77	12.4	8.1	9.1	7.3	9.8	8.2	10.1
	0.36	54.1	0.3	0.1	0.5	0.3	0.6	
	3.57	26.3	2.8	2.3	3.6	3.9	5.0	3.8

LEAD DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
26			DRY ASH/HNO ₃ -DCP(A)-GFAAS(B/C)						
	5.17	22.4	4.37	4.38	4.51	5.13	5.19	7.41	
	0.48	55.2	0.14	0.43	0.41	0.91	0.35	0.64	
	2.41	38.0	3.16	1.70	1.70	3.37	1.35	3.17	
27A			HNO ₃ -FAAS						
	6.95	22.0	5.8	5.4	7.7	8.9	8.3	5.6	
	----	----	Less than 0.48						
	2.45	28.4	1.6	2.3	3.7	2.2	2.6	2.3	
27B			HNO ₃ -GFAAS						
	8.98	17.0	7.8	7.8	9.3	11			
	----	----	No data						
	----	----	No data						
28			HNO ₃ -GFAAS						
	8.78	37.7	4.94	4.97	11.8	11.2	7.88	11.9	
	2.15 ^{RT}	31.4	1.58	1.22	2.21	2.11	2.95	2.81	
	4.03 ^{RT}	33.9	4.21	2.35	3.78	3.13	6.38	4.32	
29			HNO ₃ -EXTN-FAAS						
	3.43	42.9	3.57	4.54	5.37	3.12	1.13	2.82	
	----	----	No data						
	----	----	No data						
30			DRY ASH/HCl/HNO ₃ -FAAS						
	3.59	9.4	3.86	3.34	3.09	3.84	3.92	3.48	
	0.21	10.8	0.227	0.237	0.188	0.205	0.178	0.205	
	1.08	11.4	0.985	0.963	1.02	1.25	1.05	1.22	
31			DRY ASH/HNO ₃ /HCl-ASV						
	5.60	21.5	6.38	6.20	4.21				
	0.25	6.9	0.26	0.23	0.26				
	3.47	17.0	2.80	3.68	3.92				
32			DRY ASH/HNO ₃ -ASV						
	5.48	19.6	5.3	4.3	7.5	5.5	5.0	5.3	
	----	----	No data						
	2.03	33.7	2.5	1.7	3.1	1.6	1.2	2.1	
33A			HNO ₃ /HClO ₄ -FAAS						
	5.07	12.0	4.5	5.1	5.0	6.2	4.6	5.0	
	0.24	15.4	0.25	0.26	0.27	0.24	0.22	0.17	
	----	----	No data						
33B			HNO ₃ /HClO ₄ -GFAAS						
	4.25	14.3	3.7	4.3	4.1	5.4	3.8	4.2	
	----	----	No data						
	----	----	No data						
34			NO DATA						
35			NO DATA						
36			HNO ₃ /H ₂ SO ₄ /H ₂ O ₂ -EXTN-GFAAS						
	----	----	No data						
	----	----	Less than 0.1						
	----	----	No data						
37			No procedure						
	5.82	5.5	5.95	5.47	5.40	6.23	5.97	5.88	
	0.16	9.6	0.181	0.168	0.168	0.157	0.158	0.135	
	1.82	21.7	1.77	1.41	2.15	2.07	1.29	2.21	

LEAD DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
38	-----	-----	$\text{HNO}_3/\text{H}_2\text{O}_2$ -FAAS					
	-----	-----	Less than 25					
	-----	-----	Less than 25					
	-----	-----	Less than 25					
39			NO DATA					
40			NO DATA					
41			NO DATA					
42			$\text{DRY ASH}/\text{HNO}_3/\text{HCl}$ -ASV					
	4.03	29.6	5.0	4.4	2.7			
	0.32	18.9	0.39	0.31	0.27			
	2.30	12.8	2.2	2.0	2.3	2.7		
43			$\text{DRY ASH}/\text{HNO}_3$ -FAAS					
	6.67	7.2	6.07	6.66	7.52	6.79	6.57	6.42
	2.53 ^{NT}	12.0	2.54	3.11	2.33	2.32	2.52	2.33
	5.74 ^{NT}	5.5	6.34	5.52	5.54	5.75	5.53	5.74
44			$\text{DRY ASH}/\text{HNO}_3$ -FAAS(A)/EXTN-FAAS(B/C)					
	6.45	5.9	6.26	6.28	7.02	6.23		
	1.32 ^{NT}	40.3	1.92	0.92	1.11			
	1.64	23.3	1.2	1.91	1.8			
45			$\text{BOMB}/\text{HNO}_3/\text{HCl}$ -FAAS					
	9.60 ^{NT}	3.6	9.9	9.7	9.7	9.1		
	-----	-----	Less than 4.5					
	12.1 ^{NT}	17.1	12.0	10.8	10.6	12.0	16.1	10.9
46			NO DATA					
47			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS					
	3.35	4.8	3.36	3.05	3.40	3.32	3.41	3.53
	0.16	16.7	0.17	0.19	0.13	0.13	0.19	0.17
	0.92	19.8	0.78	1.07	0.70	1.13	0.79	1.04
48			NO DATA					
49A			$\text{HNO}_3/\text{HClO}_4$ -ICP					
	4.85	18.3	4.7	5.3	5.4	4.0		
	-----	-----	Less than 0.7					
	2.38	21.3	2.1	2.8	2.8	1.8		
49B			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS					
	6.40	16.7	5.8	6.8	5.3	7.7		
	0.28	30.3	0.21	0.18	0.31	0.36	0.97	
	2.30	26.3	1.5	2.7	1.9	2.4	3.0	
49C			HNO_3 -EP-IDSSMS					
	5.53	8.7	5.25	5.99	6.23	5.30	4.97	5.43
	0.33	---	0.322	0.340				
	2.12	32.3	2.53	2.54	2.66	2.51	1.21	1.27
50			HNO_3 -GFAAS					
	2.50	8.7	2.5	2.5	3.1	2.4	2.0	
	-----	-----	Less than 0.1					
	6.60 ^{NT}	38.6	6.8	3.5	6.3	9.7		
51			NO DATA					

CHROMIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
13			INAA						
	3.5	---	3.5						
	1.0	---	1.0						
	1.8	---	1.8						
15			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	1.9	59.7	0.59	1.6	2.4	0.83	2.1	3.6	
	0.27	45.7	0.09	0.19	0.22	0.39	0.32	0.40	
	1.1	57.1	0.53	2.00	0.50	1.80	0.88	1.00	
18			DRY ASH/ HNO_3 -ICP						
	4.1	6.0	3.89	4.14	4.05	3.97	4.53	3.87	
	1.08	11.9	0.99	1.11	0.90	1.27	1.07	1.15	
	1.6	4.2	1.70	1.61	1.69	1.53	1.60	1.56	
27			HNO_3 -FAAS						
	2.0	4.0	2.0	2.1	1.9	2.1	2.0	2.1	
	0.31	9.2	0.30	0.32	0.28	0.29	0.32	0.36	
	1.0	10.7	1.04	1.20	1.06	1.05	1.04	0.85	
49			$\text{HNO}_3/\text{HClO}_4$ -ICP						
	1.3	33.0	1.1	1.1	2.0	1.1	1.0		
	0.73	30.7	0.7	1.1	0.6	0.6	0.5	0.9	
	2.2	8.3	2.4	2.0	2.1	2.3			

MANGANESE DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
10			$\text{HNO}_3/\text{HClO}_4$ -ICP						
	27.6	2.2	27.5	26.5	27.9	28.1	28.1	27.5	
	1.48	7.2	1.66	1.54	1.39	1.4	1.4	1.5	
	19.6	5.4	21.7	19.3	19.1	19.0	19.0	19.2	
11A			BOMB/ HNO_3 -GFAAS						
	24.6	5.8	23	24.3	23.8	24.4	25	27.2	
	1.37	10.0	1.3	1.4	1.4	1.6	1.2	1.3	
	19.8	12.6	21.6	22.2	22.4	16.8	17.7	18.3	
11B			BOMB/ HNO_3 -ICP						
	21.8	7.5	21.5	19.8	24.6	22.2	22.2	20.7	
	----	----	No data						
	14.2	7.3	15.0	14.1	15.8	13.2	13.2	13.9	
14			BOMB/ HNO_3 -FAAS						
	16.8	4.5	17	16	18	16	17	17	
	2.60	16.5	2.6	2.1	3.1	2.1	3.0	2.7	
	13.2	5.7	14	12	14	13	13	13	
15			$\text{HNO}_3/\text{H}_2\text{O}_2$ -GFAAS						
	16.2	49.3	11	19	10	12	29		
	0.74	22.6	0.95	0.53	0.62	0.83	0.77		
	6.60	52.8	5.4	5.9	4.2	2.6	9.3	12	
18			DRY ASH/ HNO_3 -ICP						
	19.9	0.7	19.8	19.9	20.1	19.9	19.8	19.7	
	1.12	2.8	1.14	1.11	1.09	1.08	1.16	1.14	
	12.8	0.9	12.9	12.9	12.7	12.8	12.7	13.0	

MANGANESE DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
23A			HNO ₃ -FAAS-0.25 g samples						
	22.0	3.1	21.4	22.1	21.5	21.5	23.2	22.1	
	1.25	16.6	1.5	1.0	1.0	1.3	1.4	1.3	
	15.4	2.8	14.8	14.9	15.8	15.7	15.5	15.7	
23B			HNO ₃ -FAAS-1 g samples						
	----	----	No data						
	0.75	14.0	0.7	0.7	0.6	0.8	0.8	0.9	
	12.2	2.2	12.4	12.6	12.0	12.1	11.9	12.0	
26			DRY ASH/HNO ₃ -DCP						
	21.2	2.9	22.2	21.3	21.3	20.3	21.3	21.0	
	1.50	1.9	1.49	1.52	1.51	1.48	1.54	1.46	
	18.8	3.4	19.1	18.1	18.0	19.5	18.7	19.4	
38			HNO ₃ /H ₂ O ₂ -FAAS						
	14.7	11.0	15.7	14.5	12.2	14.8	16.5		
	7.80	0.3	7.81	7.77	7.79	7.78	7.79	7.84	
	8.60	4.6	8.33	8.77	8.36	8.34	9.23		
49			HNO ₃ /HClO ₄ -ICP						
	18.0	3.2	18.4	17.6	17.8	18.7	17.3		
	0.98	20.9	0.9	1.2	0.8	0.8	1.2		
	10.7	2.5	10.3	10.8	10.6	10.9			
51			No procedure						
	22.3	22.1	19	20	28				
	----	----	No data						
	22.8	10.1	20	25	23	21	22	26	

IRON DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
10			HNO ₃ /HClO ₄ -ICP						
	222	3.3	218	210	223	232	225	222	
	29.5	13.5	25.6	25.8	30.4	30.6	35.2		
	173	2.9	170	171	176	176	166	180	
11A			BOMB/HNO ₃ -FAAS						
	200	6.4	212	180	210	209	191	197	
	13.9	12.0	14.7	13.8	14.7	16.1	12.7	11.4	
	142	5.3	133	145	145	139	136	154	
11B			BOMB/HNO ₃ -ICP						
	157	5.0	158	148	156	158	170	150	
	15.0	20.5	11.3	17.6	19.7	13.9	13.8	13.6	
	125	8.2	132	127	136	118	111		
14			BOMB/HNO ₃ -FAAS						
	140	3.0	140	148	137	139	136	139	
	26.7	16.0	24	32	30	29	24	21	
	154	8.3	165	145	172	141	158	144	
15			HNO ₃ /H ₂ O ₂ -GFAAS						
	132	24.2	92	170	120	120	160		
	23.8	35.3	17	24	25	37	16		
	123	10.3	120	110	120	140			

IRON DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
16			BOMB/HNO ₃ -FAAS					
	177	1.9	175	173	178	176	180	182
	----	----	No data					
	160	9.3	154	149	160	152	186	
18			DRY ASH/HNO ₃ -ICP					
	162	2.7	168	156	165	161	159	160
	19.7	7.1	18.8	19.4	18.5	18.7	21.3	21.7
	124	1.6	125	123	121	124	123	127
23A			HNO ₃ -FAAS-0.25 g samples					
	166	6.4	152	156	170	173	166	180
	19.5	14.8	18.2	20.9	19.6	23.8	19.1	15.1
	141	9.6	134	162	124	144	134	150
23B			NO DATA					
26			DRY ASH/HNO ₃ -DCP					
	159	6.1	177	158	160	150	159	151
	16.5	25.8	22.4	13.5	21.4	14.5	15.0	12.4
	142	3.7	149	143	132	148	143	141
35			HNO ₃ /HClO ₄ -XRF					
	136	18.8	153	134	118	173	139	100
	----	----	No data					
	----	----	No data					
44			DRY ASH/HNO ₃ -FAAS					
	212	6.8	200	209	206	233		
	52.2	12.7	46.0	47.5	60.1	55.1		
	168	2.0	163	169	168	171		
49			HNO ₃ /HClO ₄ -ICP					
	132	4.5	134	135	139	131	123	
	16.4	16.9	16.2	19.9	16.2	12.1	15.0	18.7
	104	4.9	99	102	105	111		
51			No procedure					
	180	5.6	190	180	170			
	----	----	No data					
	81.0	10.9	74	78	96	78	73	87

COBALT DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
11			BOMB/HNO ₃ -EXTN-GFAAS					
	0.60	5.2	0.58	0.60	0.56	0.54	0.53	0.53
	0.17	10.2	0.17	0.14	0.17	0.19	0.16	0.18
	0.08	9.7	0.069	0.069	0.080	0.088	0.074	0.080
13			INAA					
	0.95	---	0.95					
	0.26	---	0.26					
	0.12	---	0.12					
15			HNO ₃ /H ₂ O ₂ -GFAAS					
	0.50	33.7	0.45	0.65	0.73	0.55	0.22	0.59
	0.20	51.1	0.32	0.16	0.09	0.20	0.11	0.33
	0.07	19.1	0.075	0.083	0.088	0.064	0.060	0.054

COBALT DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
21			BOMB/HNO ₃ /H ₂ SO ₄ -ASV						
	1.30	42.9	1.5	0.7	1.6	1.9	0.7		
	----	-----	No data						
	----	-----	No data						
47			HNO ₃ /H ₂ O ₂ -GFAAS						
	0.70	4.6	0.66	0.71	0.75	0.70	0.73	0.68	
	0.14	17.7	0.11	0.15	0.17	0.15	0.13	0.11	
	0.07	18.8	0.05	0.08	0.06	0.08	0.06	0.06	
49			HNO ₃ /HClO ₄ -ICP						
	0.30	16.1	0.3	0.3	0.3	0.4	0.4		
	0.34	57.3	0.5	0.2	0.2	0.2	0.6		
	0.10	0	0.1	0.1	0.1				
51			No procedure						
	0.54	6.5	0.54	0.58	0.51				
	----	-----	No data						
	0.04	18.8	0.05	0.04	0.05	0.03	0.05	0.04	

NICKEL DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>							
3			HNO ₃ /HClO ₄ /HCl-FAAS						
	3.48	7.1	3.6	3.2	3.9	3.4	3.3	3.5	
	0.47	9.2	0.49	0.51	0.51	0.43	0.43	0.42	
	1.23	7.2	1.27	1.25	1.21	1.08	1.24	1.35	
14			BOMB/HNO ₃ -FAAS						
	1.17	10.4	1.2	1.3	1.3	1.1	1.1	1.0	
	0.10	0	0.1	0.1	0.1	0.1	0.1	0.1	
	1.42	10.4	1.6	1.2	1.5	1.3	1.5	1.4	
15			HNO ₃ /H ₂ O ₂ -GFAAS						
	3.10	36.3	1.3	2.2	3.5	4.0	3.4	4.2	
	0.35	66.7	0.12	0.35	0.18	0.76	0.24	0.44	
	0.68	48.9	0.43	0.65	0.21	1.10	0.70	0.98	
18			DRY ASH/HNO ₃ -ICP						
	3.26	8.5	3.03	3.76	3.21	3.04	3.37	3.13	
	0.44	6.8	0.42	0.42	0.44	0.45	0.43	0.50	
	0.68	5.0	0.72	0.64	0.64	0.68	0.71	0.69	
21			BOMB/HNO ₃ /H ₂ SO ₄ (A)-HNO ₃ /HClO ₄ (B/C)-ASV						
	15.2	43.5	19	25	12	10	10		
	7.8	----	7.8						
	8.1	----	8.1						
23A			HNO ₃ -FAAS-0.25 g samples						
	3.05	48.0	2.5	2.8	2.4	2.5	6.0	2.1	
	----	-----	Less than 0.8						
	----	-----	Less than 0.8						
23B			HNO ₃ -FAAS-1 g samples						
	----	-----	No data						
	0.38	57.1	0.2	0.7	0.3	0.2	0.5		
	0.30	21.1	0.3	0.4	0.3	0.2	0.3	0.3	

NICKEL DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
25			$\text{HNO}_3/\text{HClO}_4\text{-FAAS}$					
	1.90	16.3	1.8	1.9	1.8	1.6	1.8	2.5
	0.36	31.7	0.3	0.2	0.4	0.4	0.5	
	0.50	16.7	0.60	0.45	0.40	0.60	0.50	0.45
27			$\text{HNO}_3\text{-FAAS}$					
	2.64	2.2	2.74	2.59	2.58	2.61	2.63	2.66
	---	----	Less than 0.29					
	0.61	5.7	0.60	0.55	0.65	0.59	0.63	0.61
49			$\text{HNO}_3/\text{HClO}_4\text{-ICP}$					
	2.54	10.6	2.5	2.3	3.0	2.5	2.4	
	0.57	26.6	0.3	0.7	0.5	0.6	0.6	0.7
	0.68	7.4	0.7	0.7	0.6	0.7		
51			No procedure					
	3.47	4.4	3.3	3.5	3.6			
	---	----	No data					
	0.56	15.2	0.58	0.59	0.41	0.60	0.62	

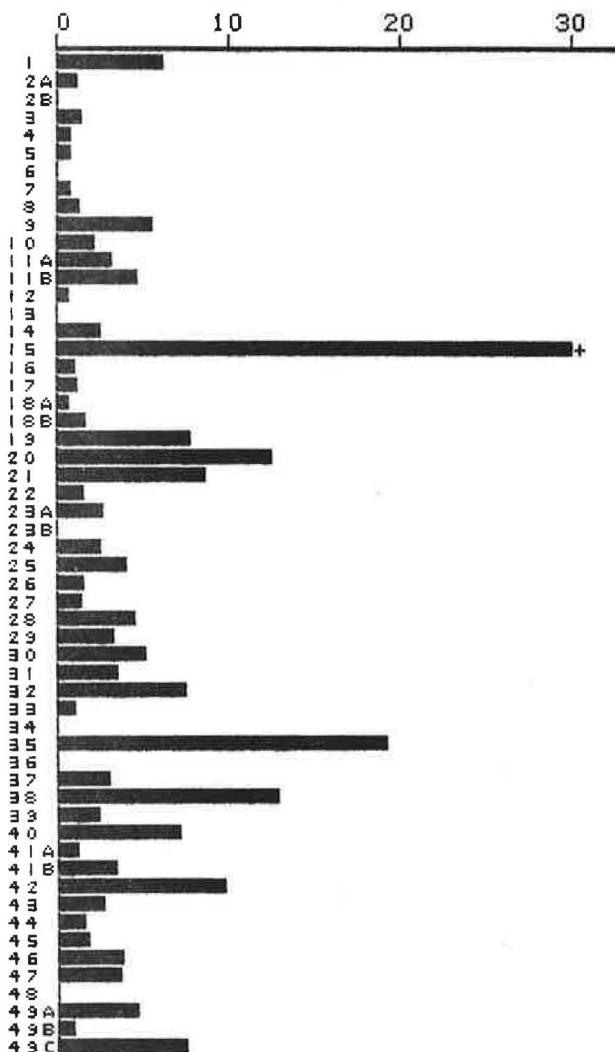
SELENIUM DATA

<u>Lab</u>	<u>Mean</u>	<u>RSD</u>						
9			$\text{O}_2 \text{ COMBUSTION-}\text{HNO}_3\text{-HG-AAS}$					
	6.2	5.7	6.4	6.0	6.3	5.5	6.4	6.3
	0.70	11.7	0.75	0.84	0.67	0.69	0.65	0.61
	2.6	6.6	2.8	2.6	2.6	2.7	2.7	2.9
12			$\text{BOMB/HNO}_3/\text{HClO}_4\text{-NI-GFAAS}$					
	6.9	4.5	6.45	7.28	6.66	6.92	7.09	7.12
	0.77	7.9	0.86	0.83	0.70	0.74	0.76	0.74
	2.1	7.5	1.90	1.97	1.94	2.23	2.23	2.19
13			INAA					
	7.7	----	7.7					
	1.4	----	1.4					
	3.0	----	3.0					
40			$\text{HNO}_3/\text{H}_2\text{SO}_4\text{-HG-AAS}$					
	8.3	7.8	8.27	8.00	7.24	9.22	8.48	8.30
	0.14	0	0.14	0.14	0.14	0.14	0.14	0.14
	2.8	9.7	2.53	2.87	2.45	2.82	2.78	3.21
48			$\text{BOMB/HNO}_3\text{-FLUOR}$					
	6.1	3.5	6.32	6.02	6.15	6.15	6.38	5.78
	0.86	5.6	0.86	0.93	0.82	0.84		
	2.3	5.8	2.35	2.35	2.16	2.37	2.40	2.08

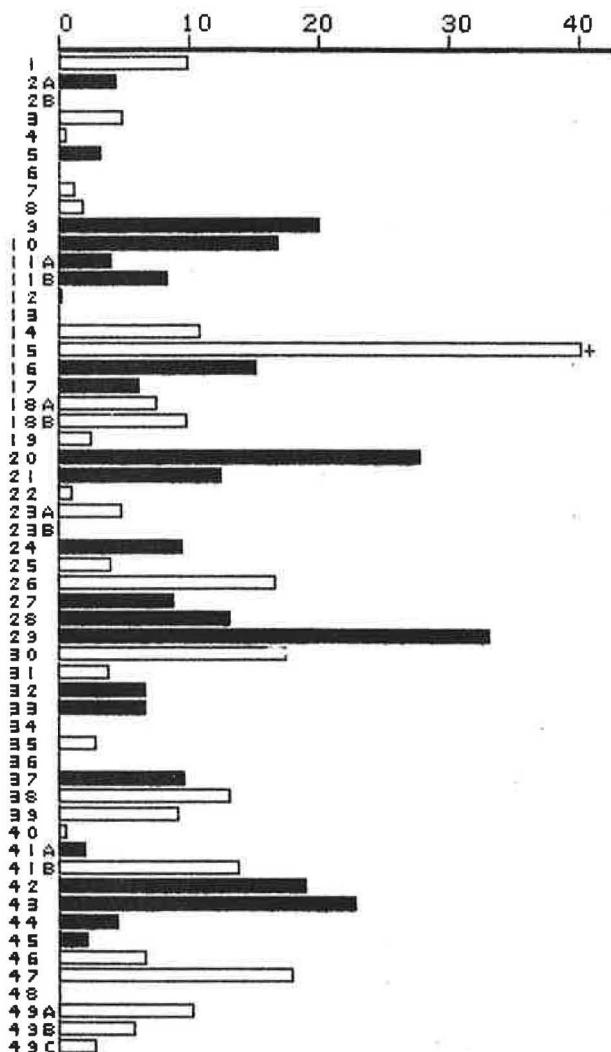
COPPER - SAMPLE A

LOBSTER HEPATOPANCREAS - 331 mg/Kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



— POSITIVE BIAS

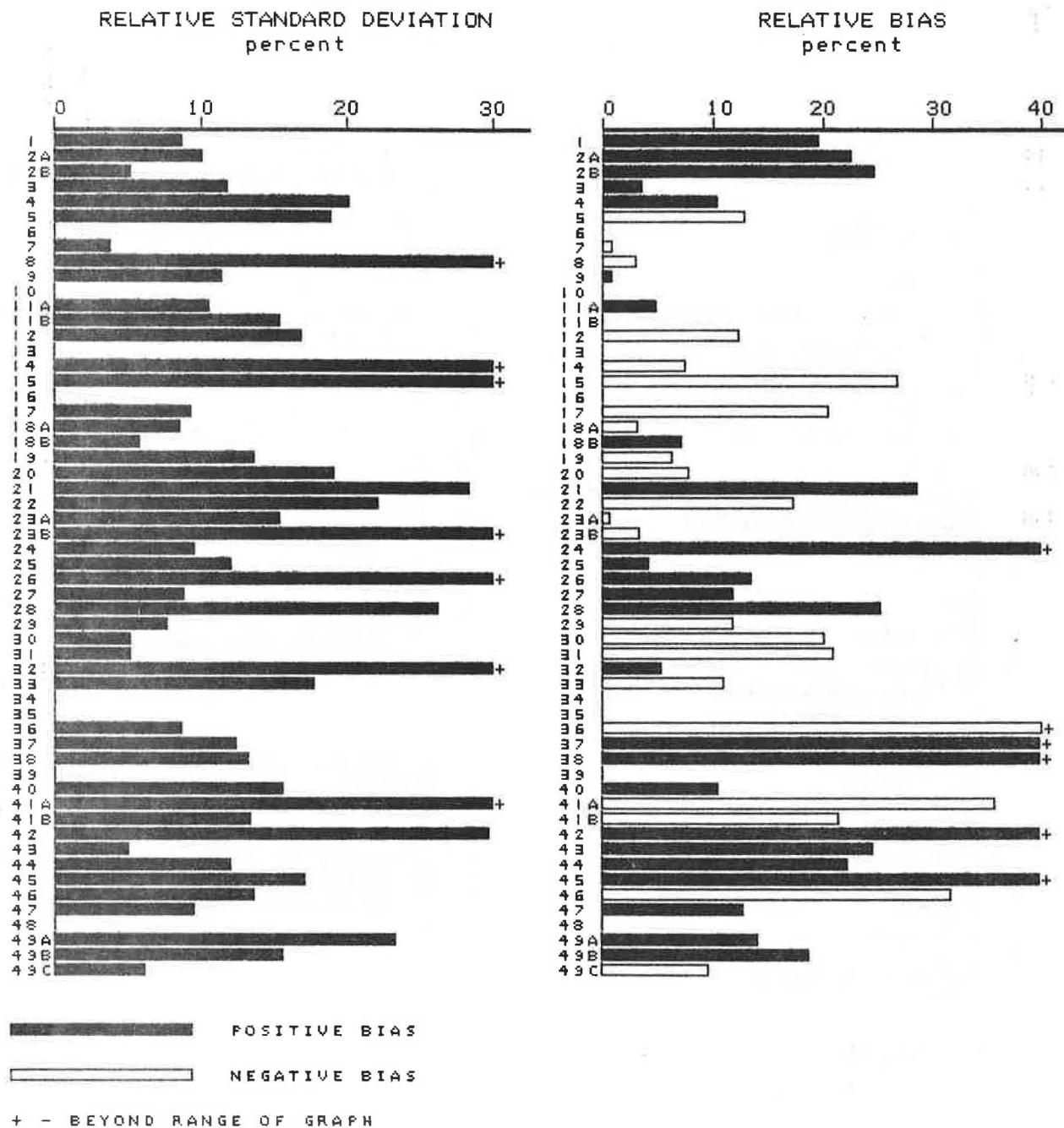
— NEGATIVE BIAS

* - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

COPPER - SAMPLE B

SCALLOPS ADDUCTOR MUSCLE - 3.7 mg/kg



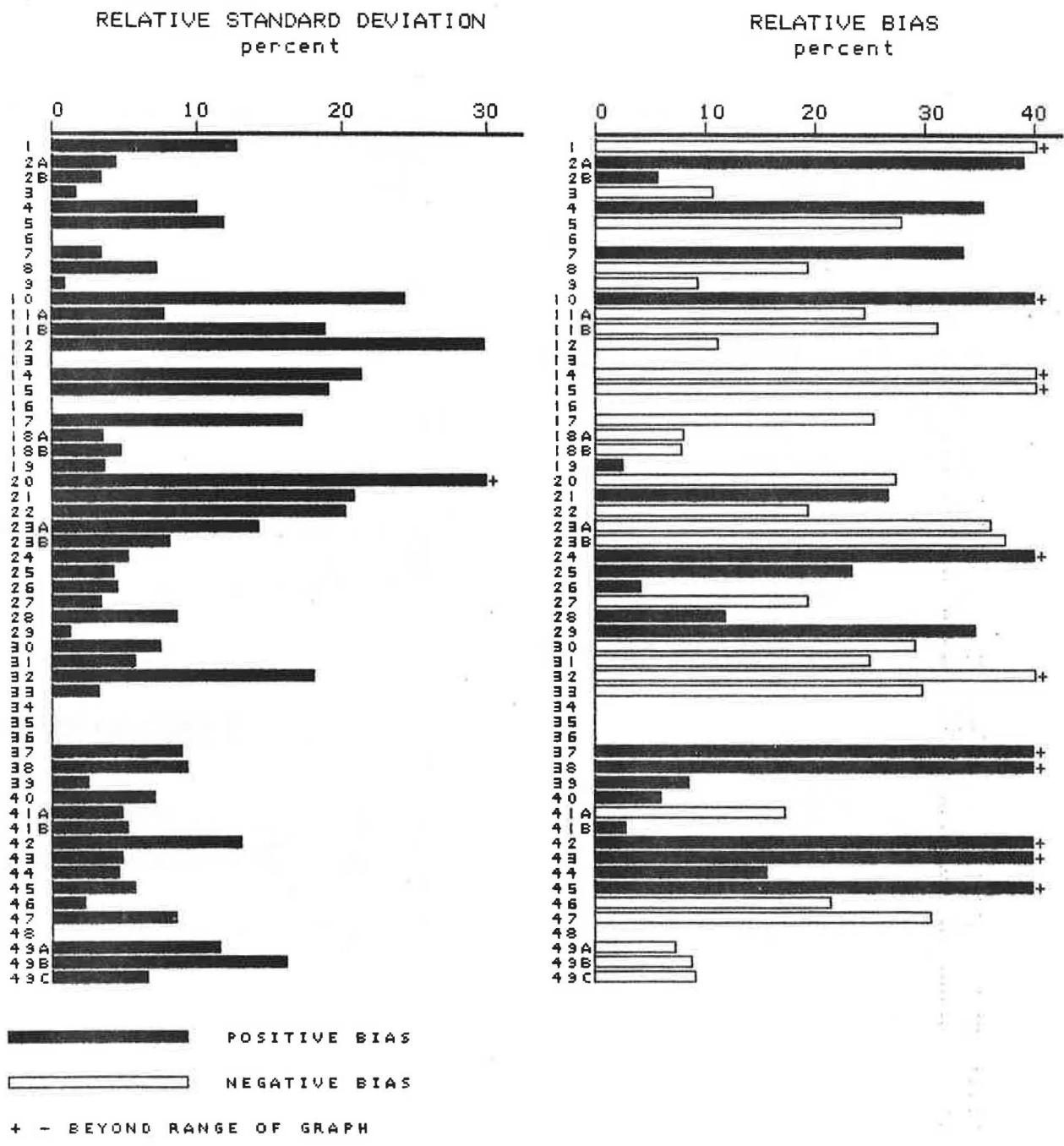
THERE ARE NO RESULTS FROM

THERE ARE NO RESULTS FROM A

+ - BEYOND RANGE OF GRAPH

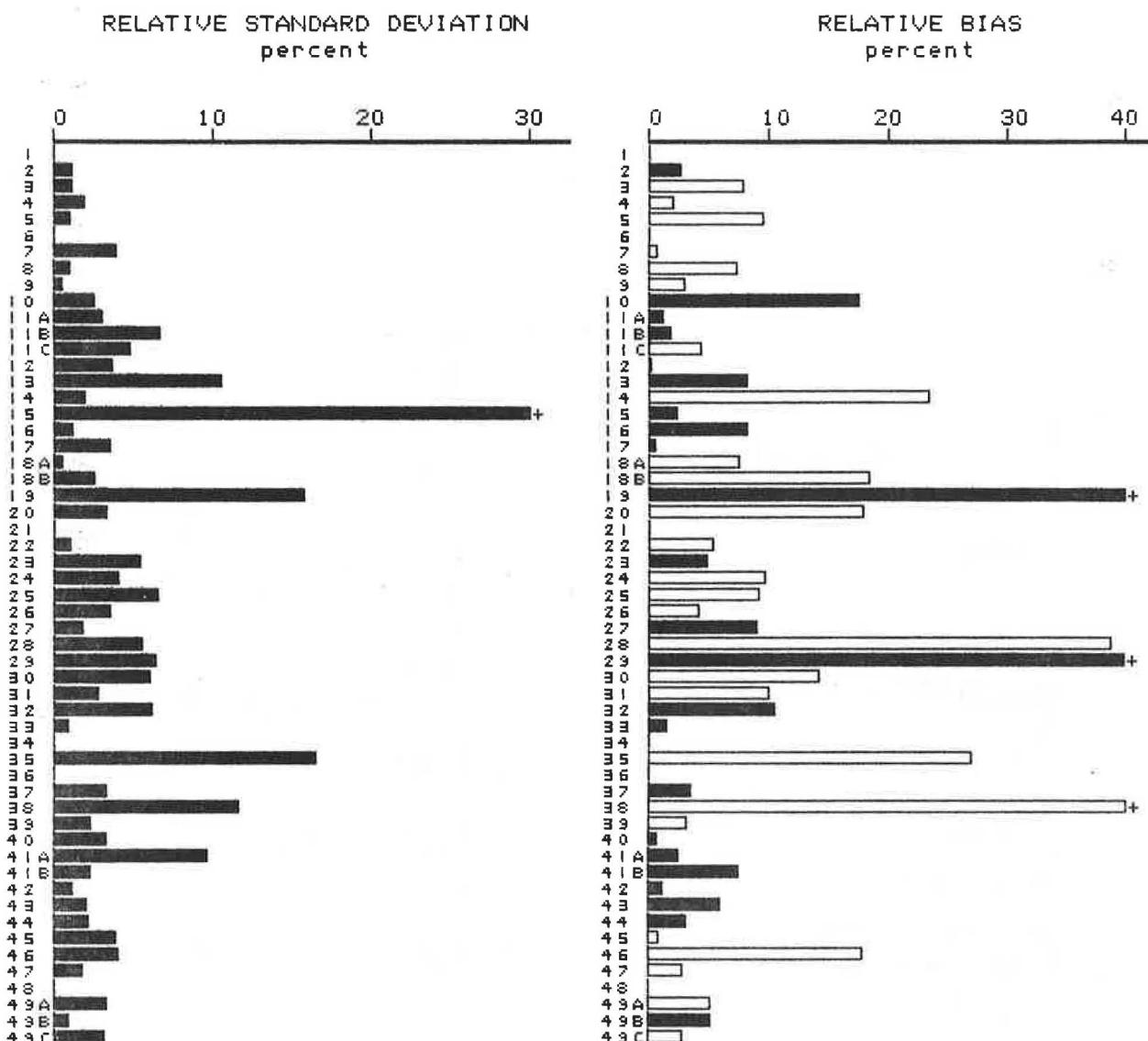
COPPER - SAMPLE C

PLAICE MUSCLE - 3.1 mg/kg



ZINC - SAMPLE A

LOBSTER HEPATOPANCREAS - 179 mg/Kg



— POSITIVE BIAS

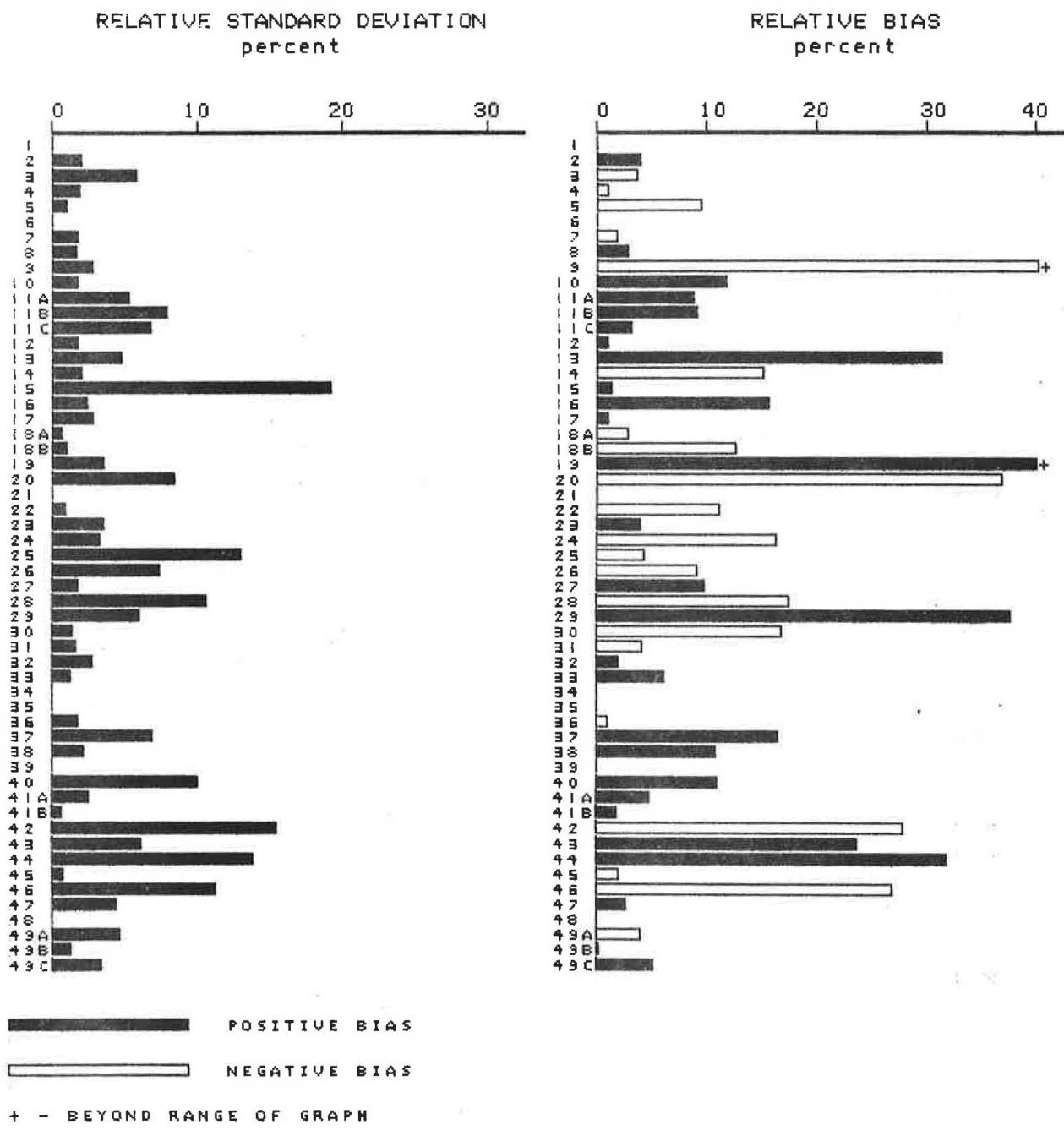
— NEGATIVE BIAS

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

ZINC - SAMPLE B

SCALLOPS ADDUCTOR MUSCLE - 58 mg/Kg

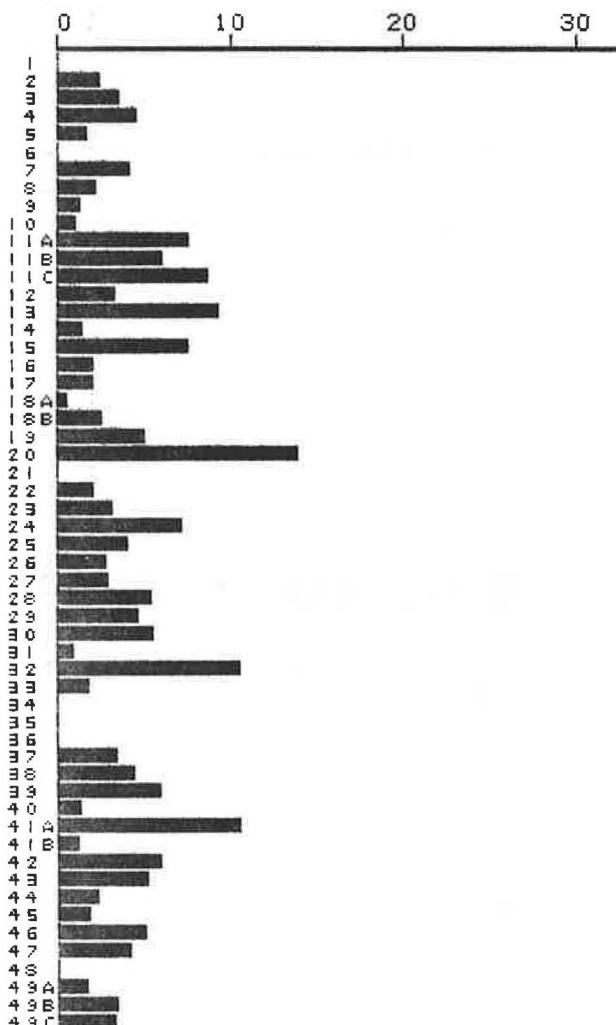


THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

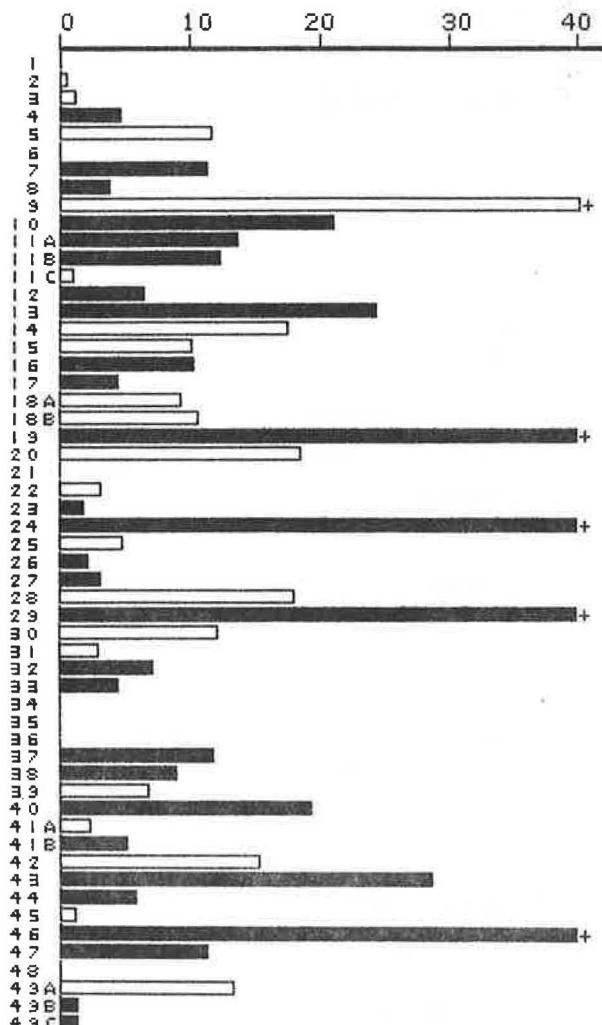
ZINC - SAMPLE C

PLAICE MUSCLE - 93 mg/kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



— POSITIVE BIAS

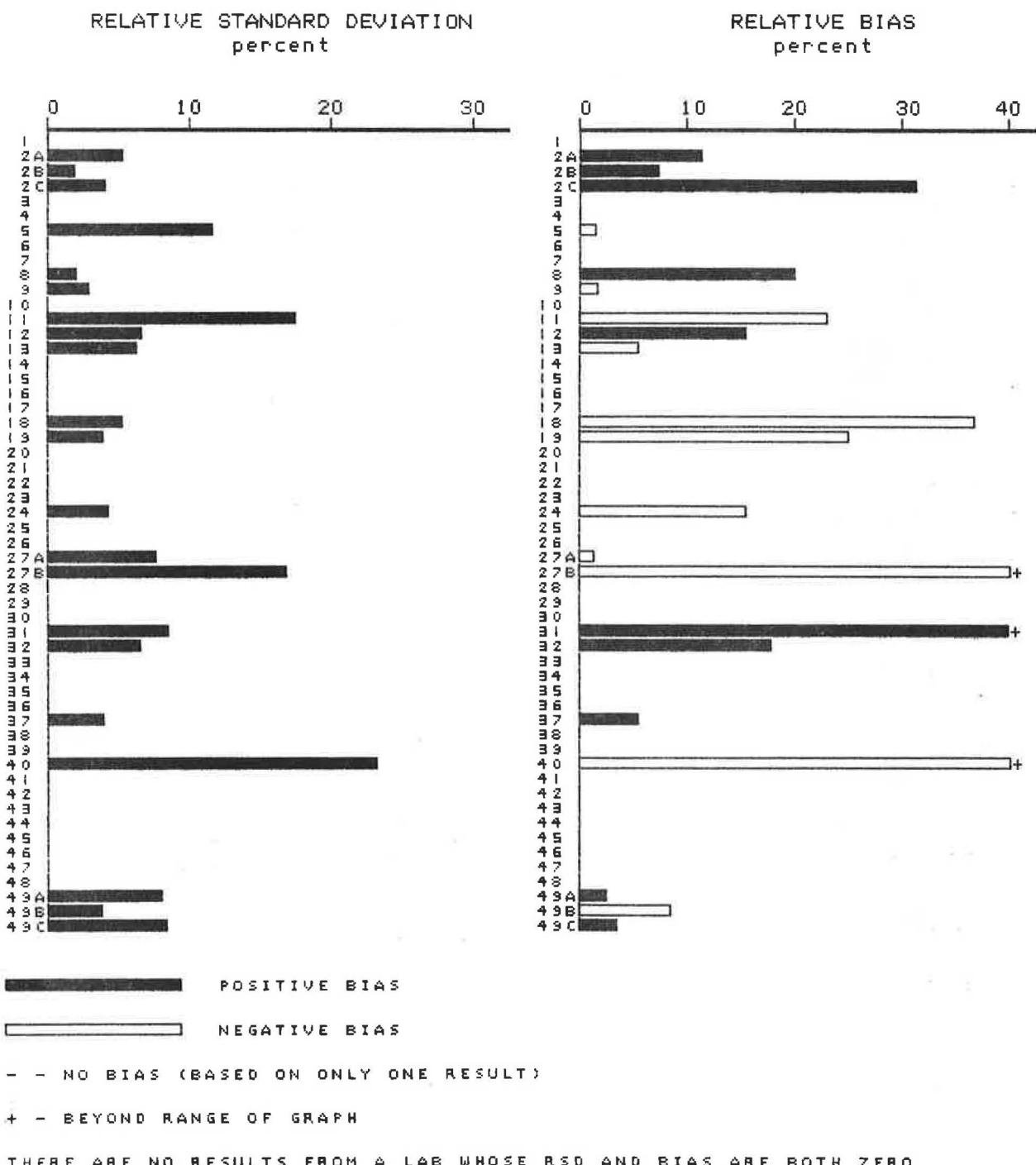
— NEGATIVE BIAS

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

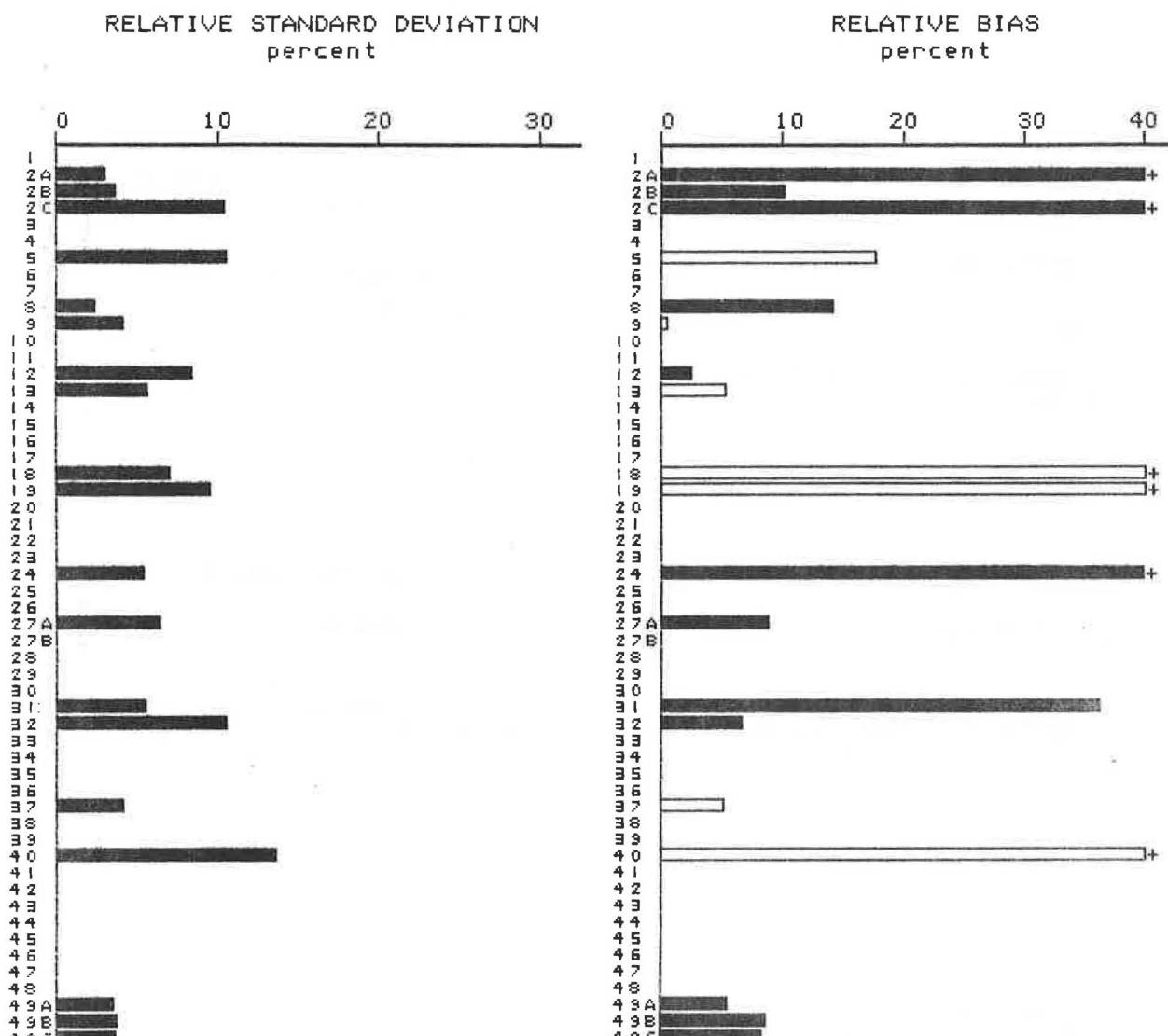
ARSENIC - SAMPLE A

LOBSTER HEPATOPANCREAS - 25 mg/Kg



ARSENIC - SAMPLE B

SCALLOPS ADDUCTOR MUSCLE - 7.1 mg/Kg



— POSITIVE BIAS

— NEGATIVE BIAS

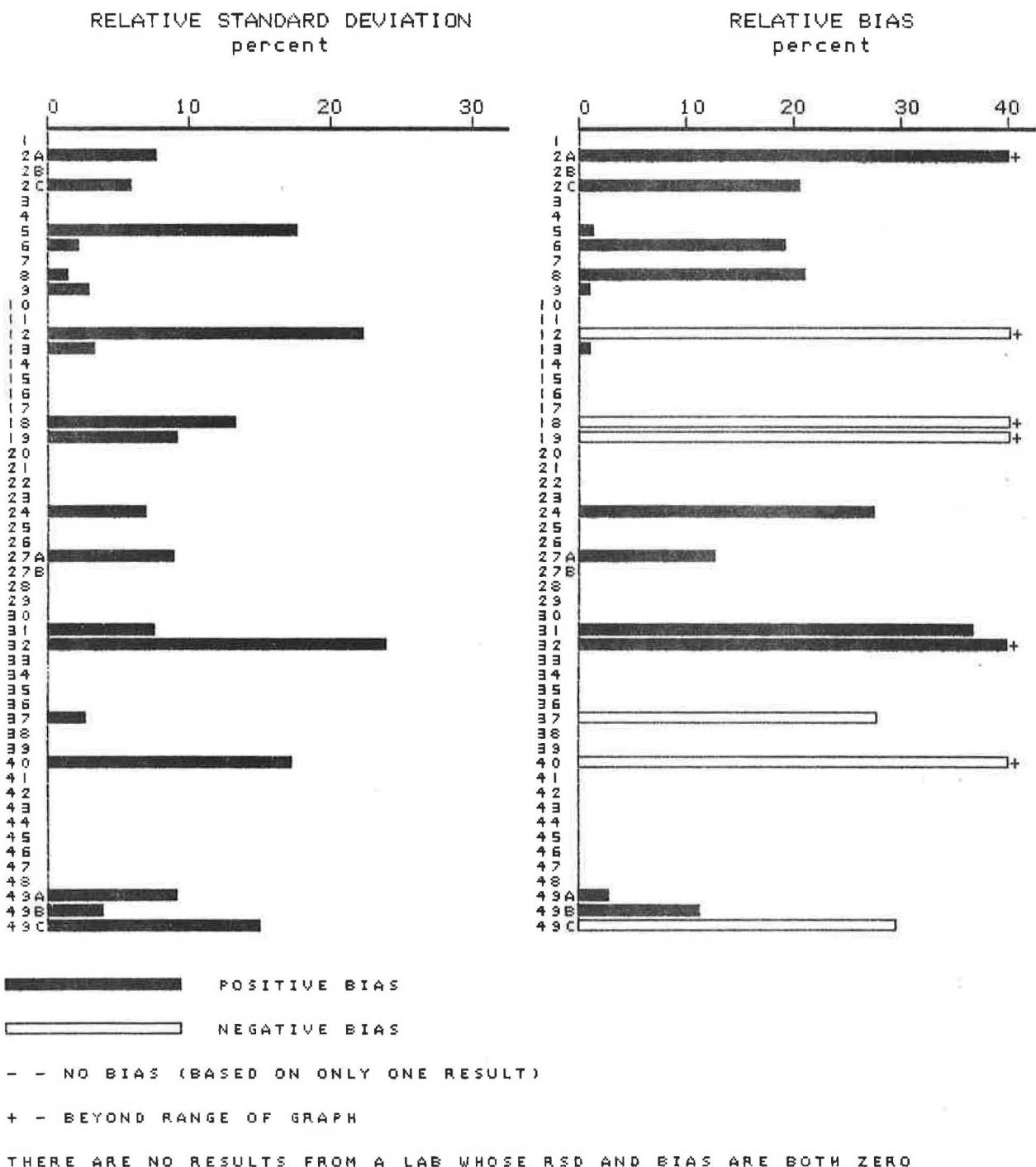
-- NO BIAS (BASED ON ONLY ONE RESULT)

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

ARSENIC - SAMPLE C

PLAICE MUSCLE - 4.6 mg/kg



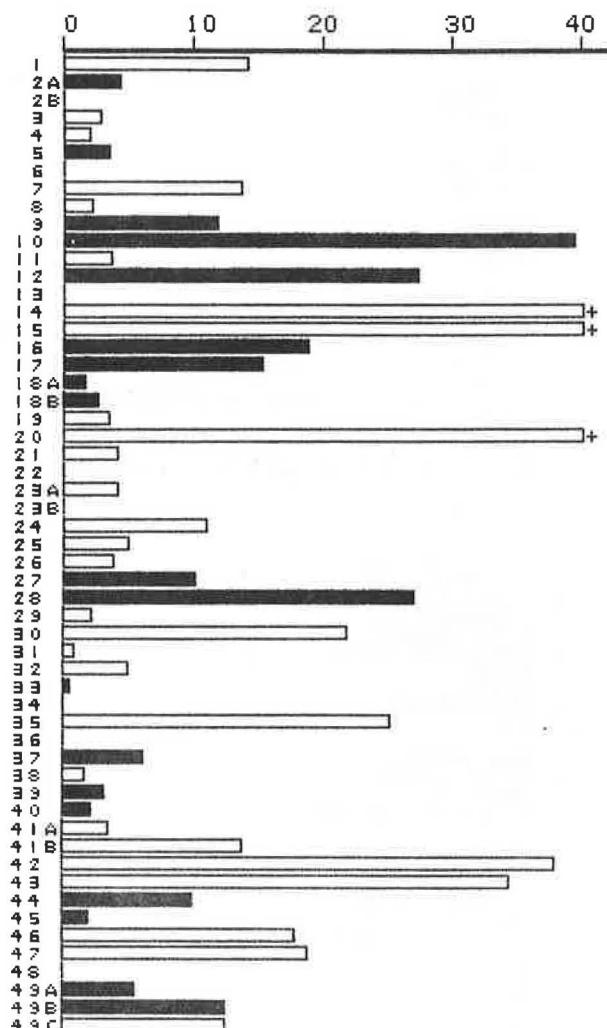
CADMIUM - SAMPLE A

LOBSTER HEPATOPANCREAS - 26 mg/Kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



— POSITIVE BIAS

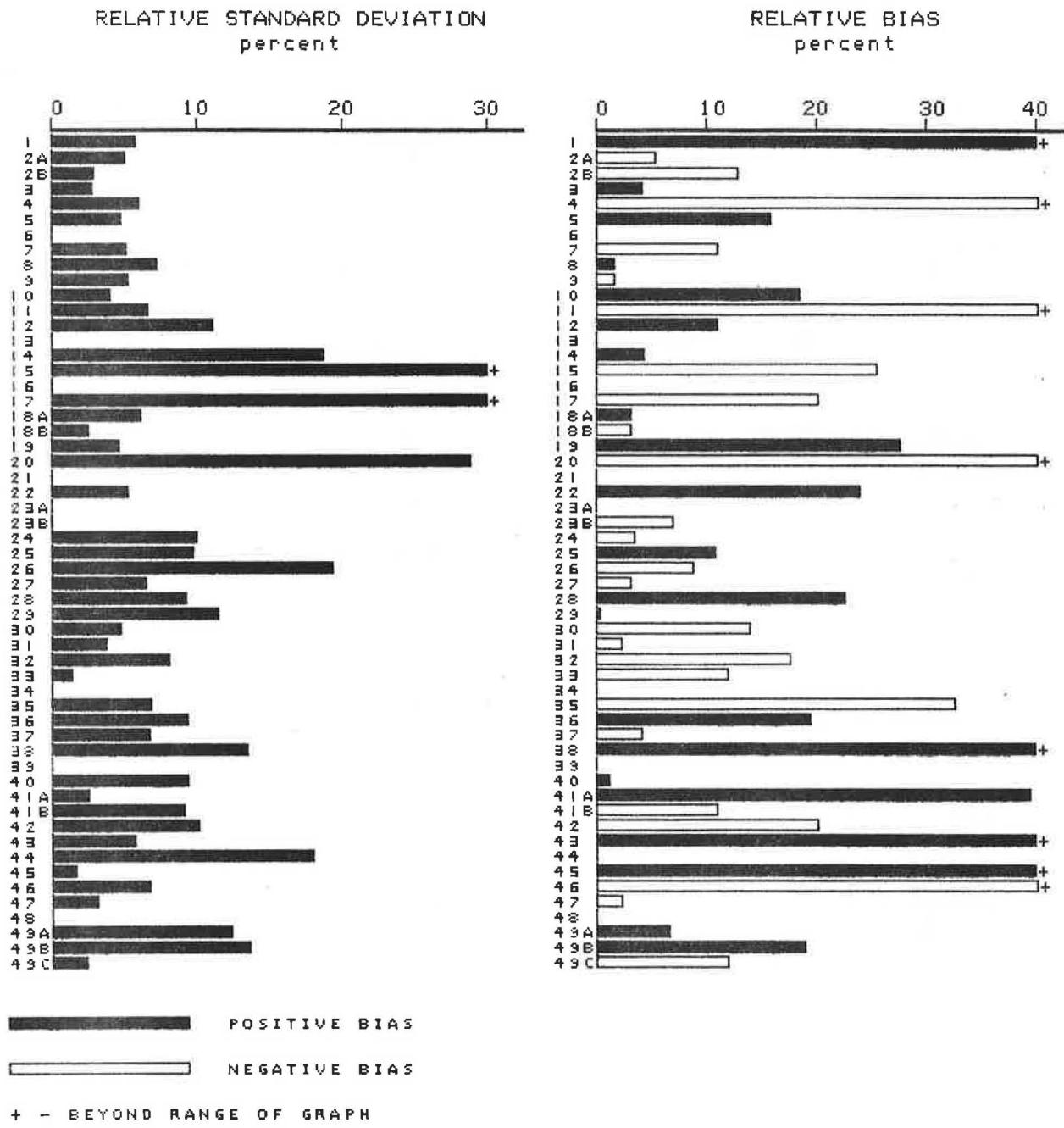
— NEGATIVE BIAS

+ — BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

CADMIUM - SAMPLE B

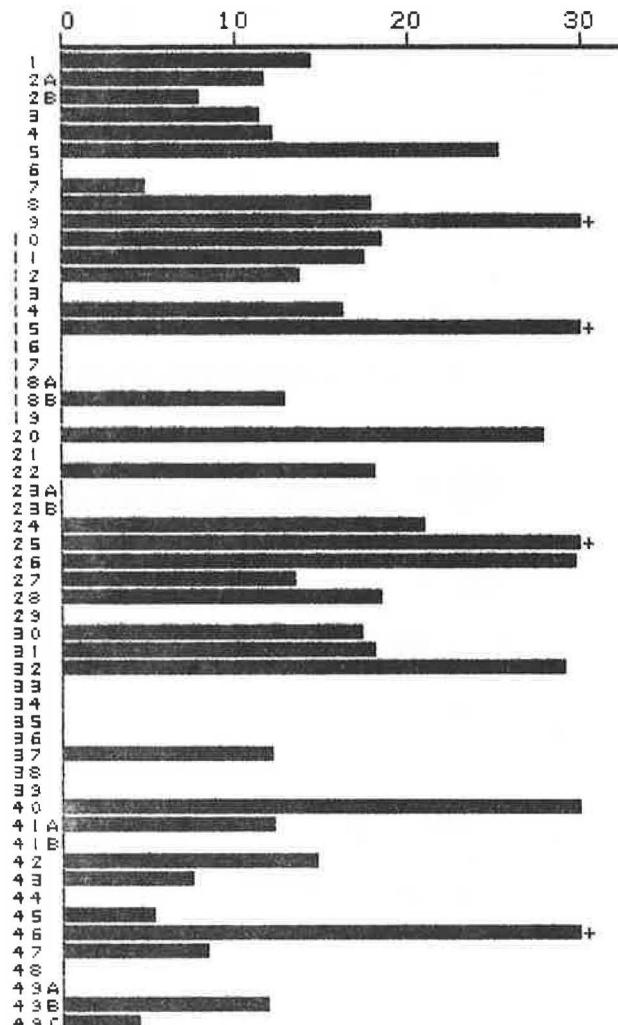
SCALLOPS ADDUCTOR MUSCLE - 0.75 mg/Kg



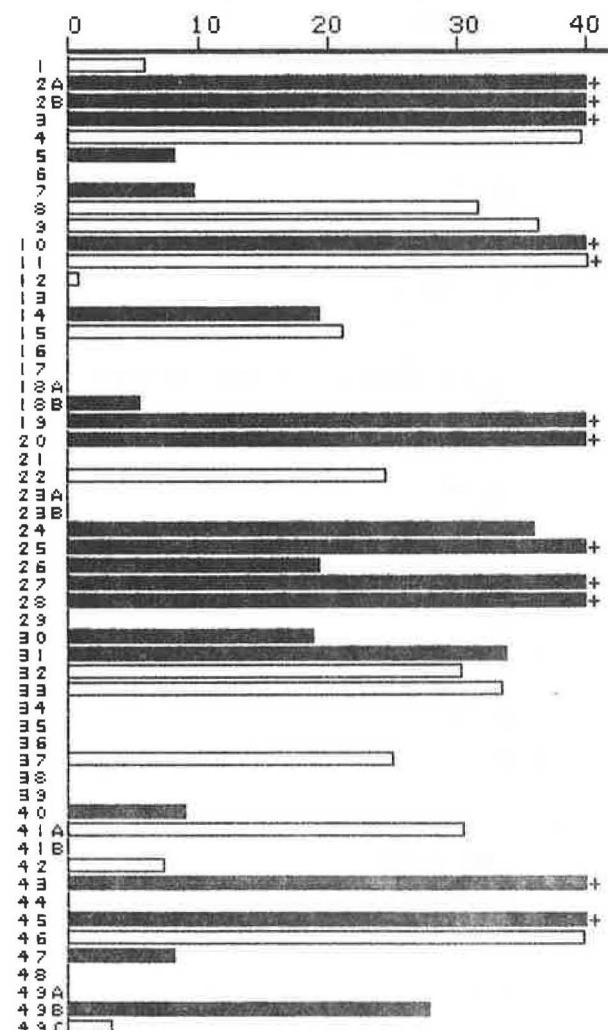
CADMIUM - SAMPLE C

PLAICE MUSCLE - 0.06 mg/kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



— POSITIVE BIAS

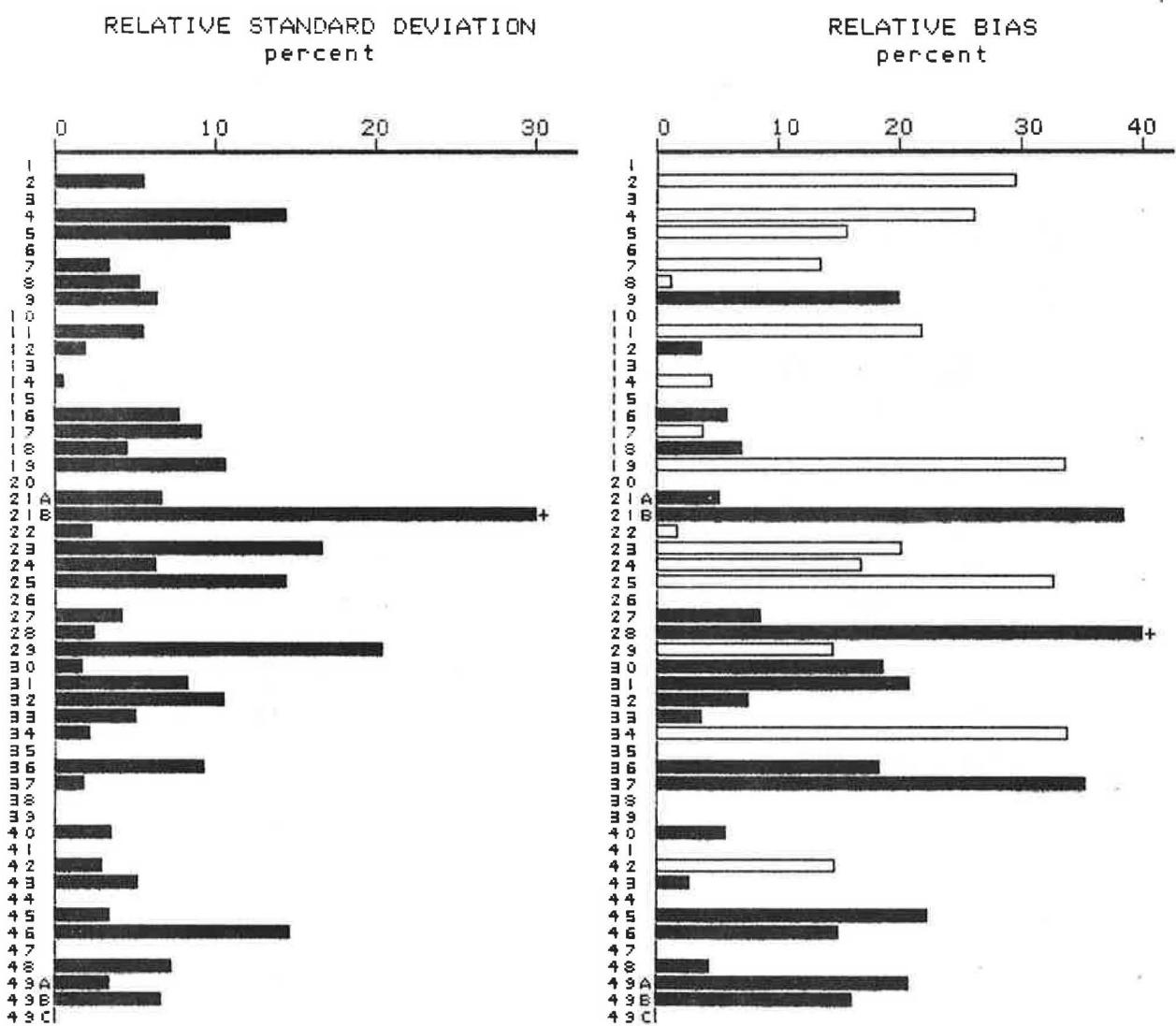
— NEGATIVE BIAS

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

MERCURY - SAMPLE A

LOBSTER HEPATOPANCREAS - 0.254 mg/Kg



— POSITIVE BIAS

— NEGATIVE BIAS

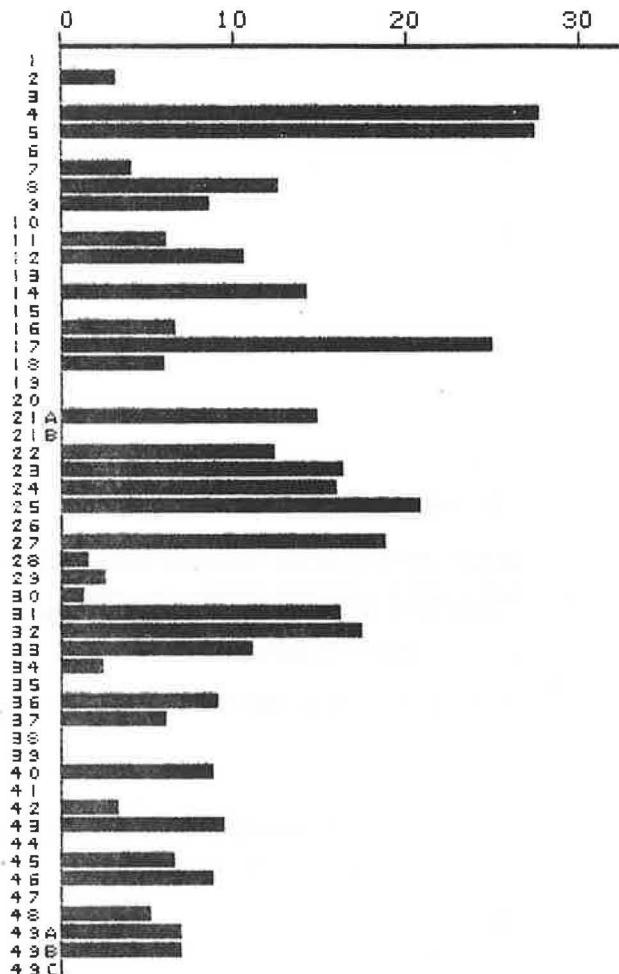
+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

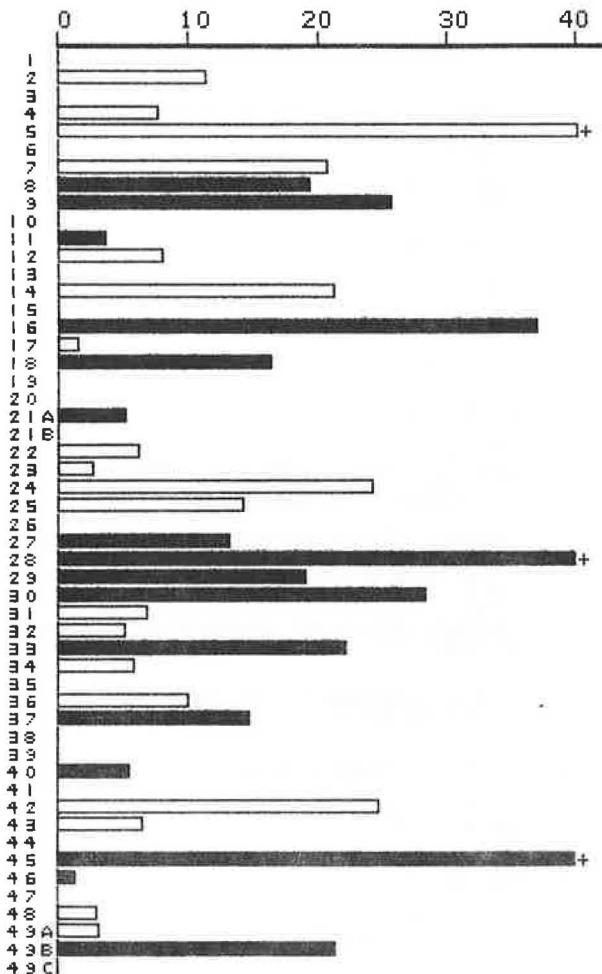
MERCURY - SAMPLE B

SCALLOPS ADDUCTOR MUSCLE - 0.081 mg/Kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



===== POSITIVE BIAS

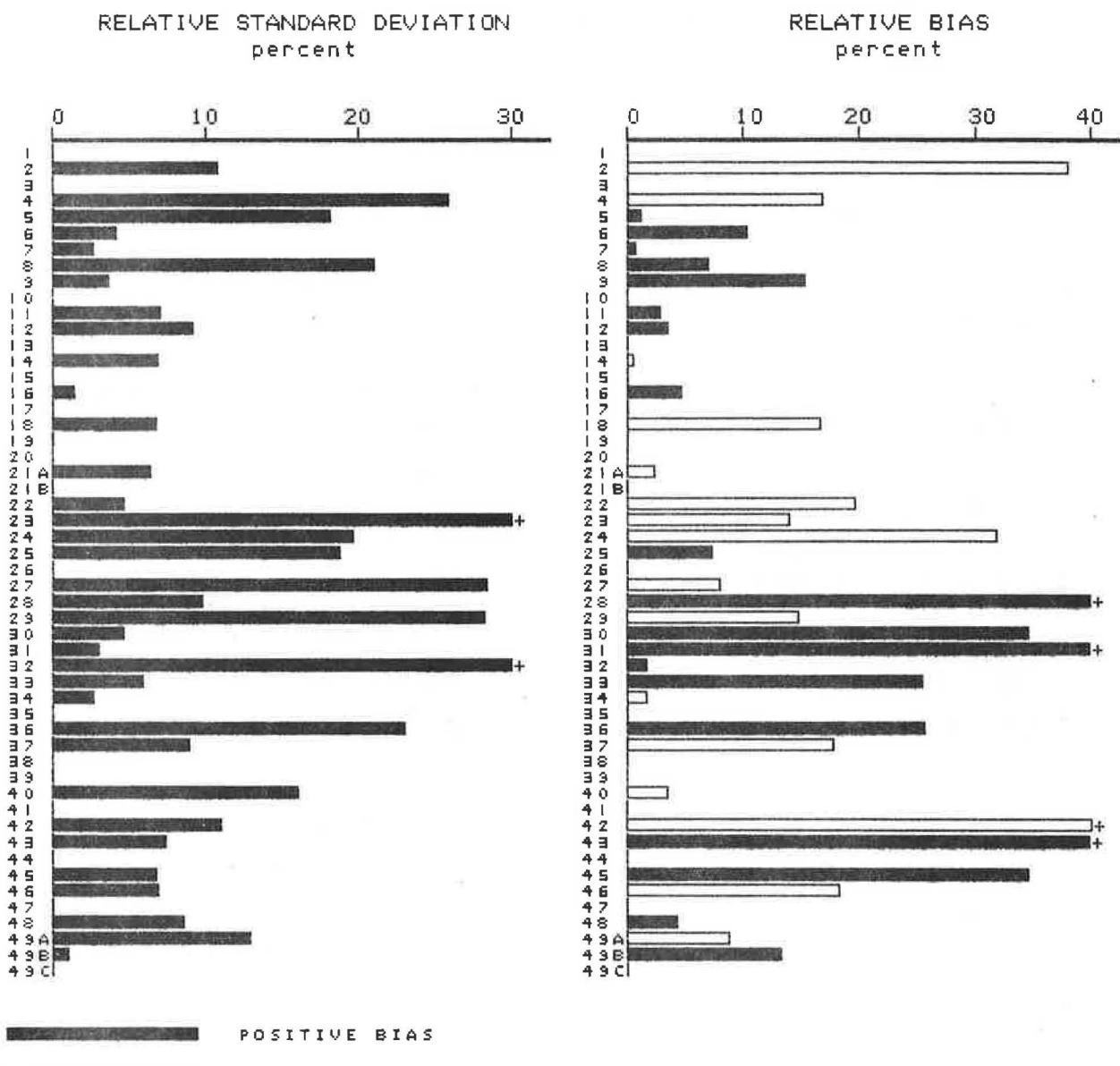
===== NEGATIVE BIAS

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

MERCURY - SAMPLE C

PLAICE MUSCLE - 0.056 mg/Kg



THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

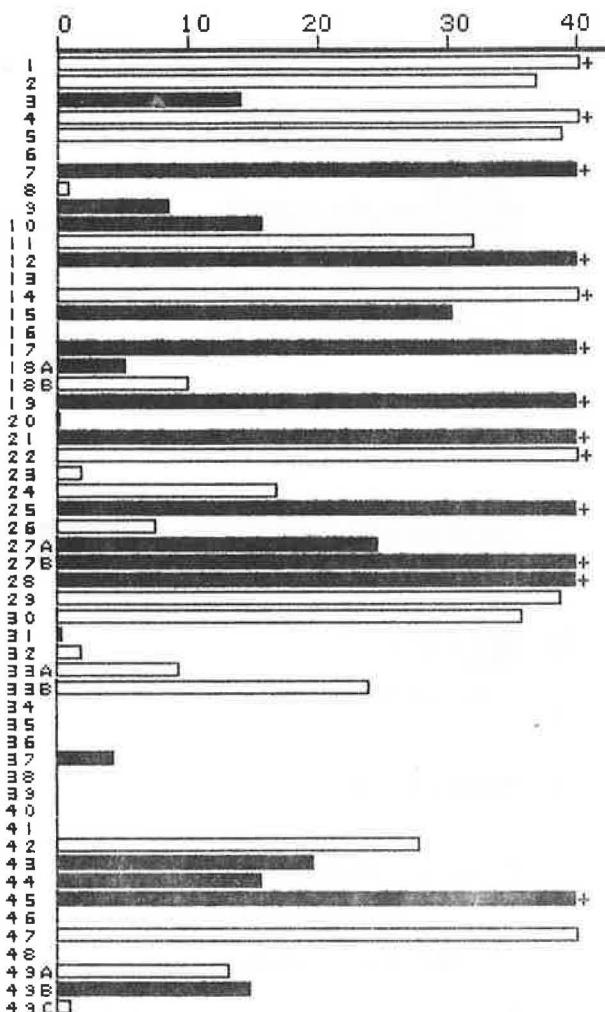
LEAD - SAMPLE A

LOBSTER HEPATOPANCREAS - 5.57 mg/Kg

RELATIVE STANDARD DEVIATION
percent



RELATIVE BIAS
percent



— POSITIVE BIAS

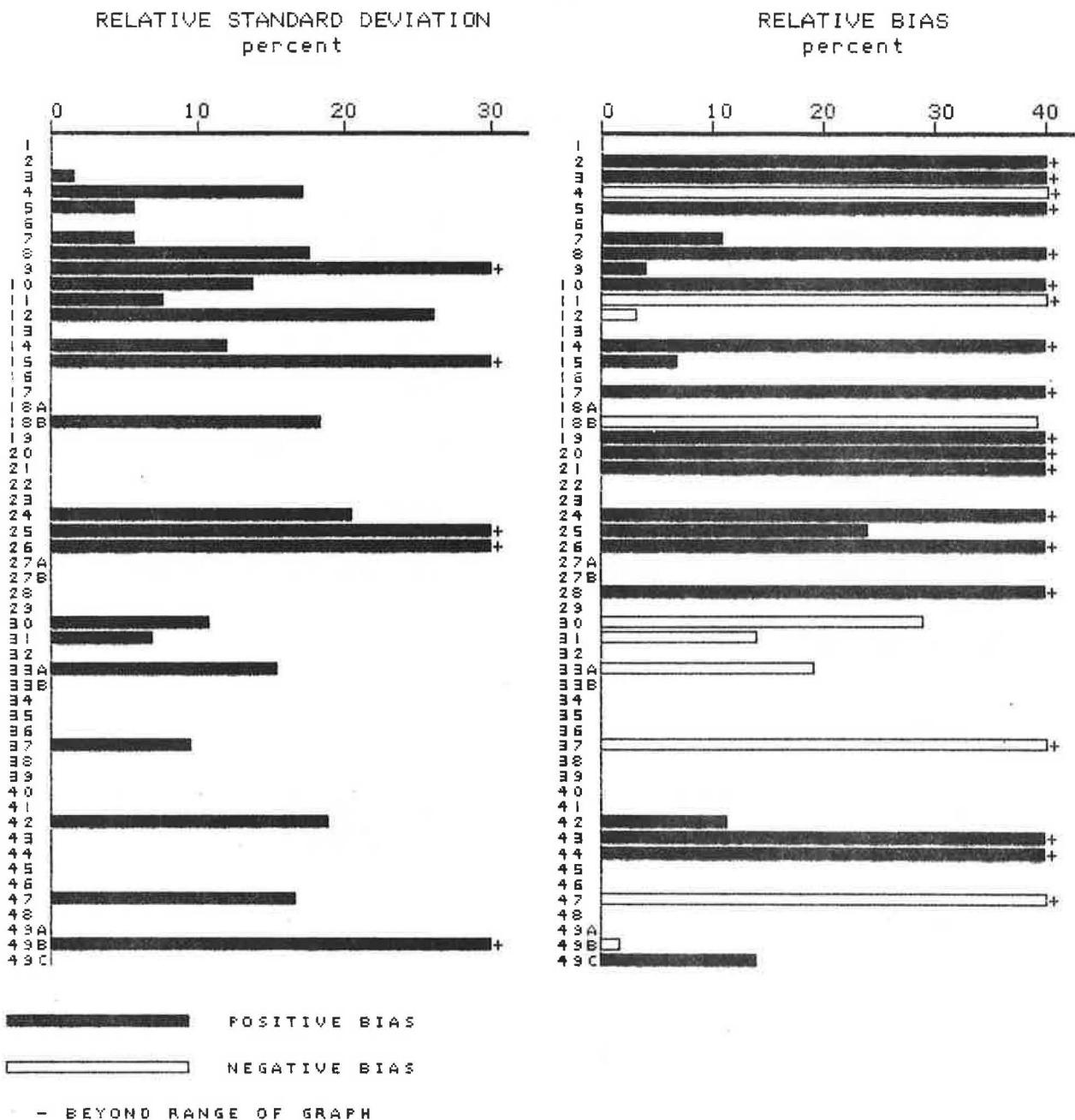
— NEGATIVE BIAS

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

LEAD - SAMPLE B

SCALLOPS ADDUCTOR MUSCLE - 0.29 mg/Kg

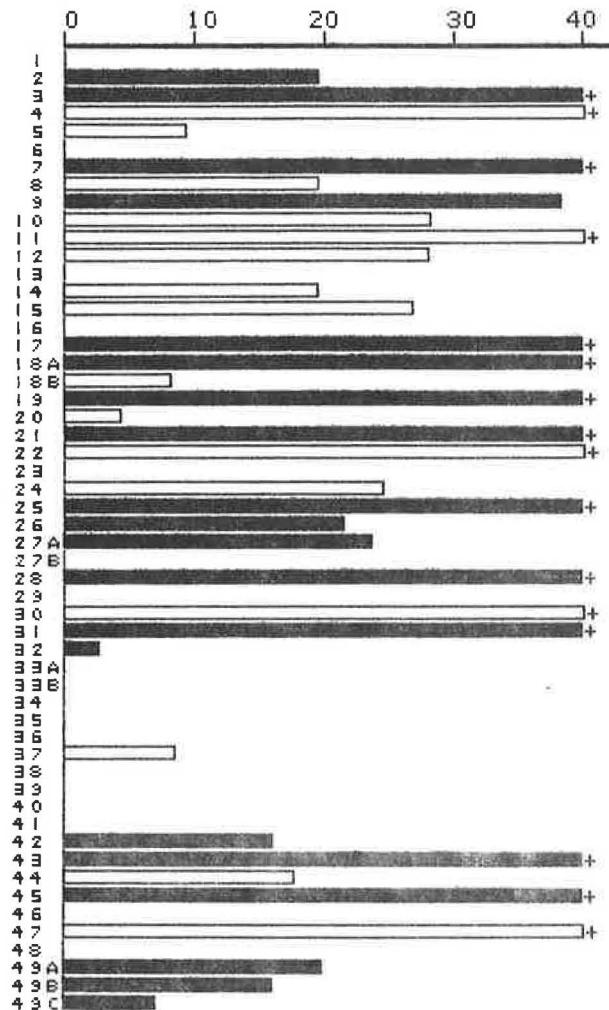
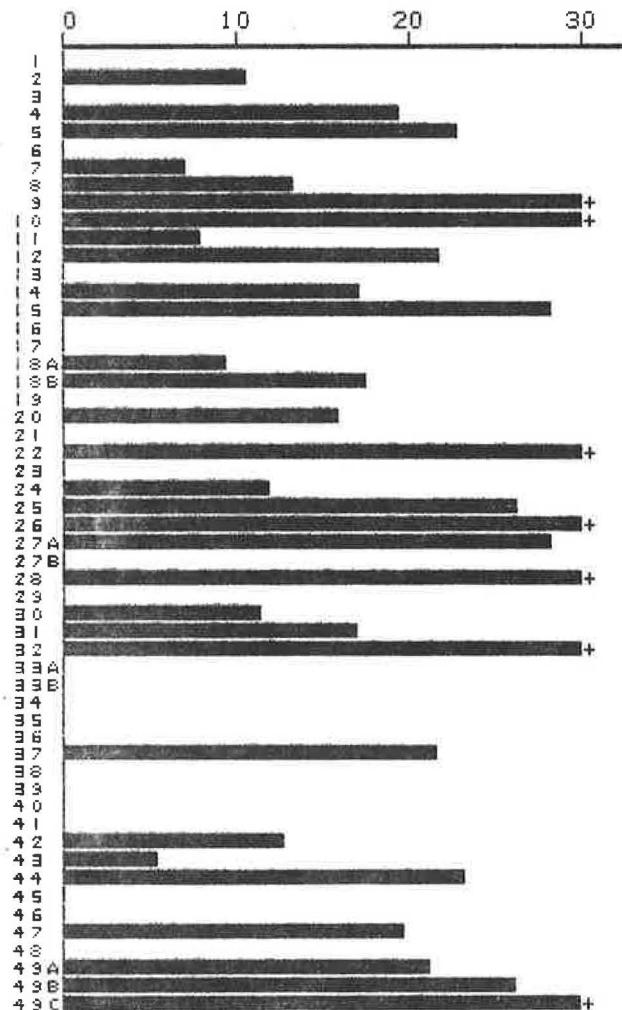


LEAD - SAMPLE C

PLAICE MUSCLE - 1.98 mg/Kg

RELATIVE STANDARD DEVIATION
percent

RELATIVE BIAS
percent



— POSITIVE BIAS —

— NEGATIVE BIAS —

+ - BEYOND RANGE OF GRAPH

THERE ARE NO RESULTS FROM A LAB WHOSE RSD AND BIAS ARE BOTH ZERO

Indication of spine colours

Reports of the Advisory Committee on Fishery Management	Red
Reports of the Advisory Committee on Marine Pollution	Yellow
Fish Assessment Reports	Grey
Pollution Studies	Green
Others	Black

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