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# Validation of a Metallomics Analysis of Placenta Tissue by Inductively-Coupled Plasma Mass Spectrometry

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## Abstract

Trace elements can play an important role in maternal health and fetal development, and deficiencies in some essential minerals including zinc and copper have been correlated in some individuals to the development of birth defects and adverse health outcomes later in life. The exact etiology of conditions like preeclampsia and the effects of fetal exposure to toxic metals has not been determined, making the assessment of trace element levels crucial to the elucidation of the causes of conditions like preeclampsia. Previous studies analyzing serum and placenta tissue have produced conflicting findings, suggesting the need for a robust, validated sample preparation and analysis method for the determination of trace elements in placenta. In this report, an acid digestion method and analysis by ICP-MS for a broad metallomics/mineralomics panel of trace elements is developed and validated over three experimental days for inter- and intraday precision and accuracy, linear range, matrix impact, and dilution verification. Spike recovery experiments were performed for the essential elements chromium (Cr), iron (Fe), cobalt (Co), nickel (Ni), copper (Cu), and zinc (Zn), and the toxic elements arsenic (As), cadmium (Cd), and lead (Pb) at levels equal to and in excess of native concentrations in control placenta tissue. The validated method will be essential for the development of scientific studies of maternal health and toxic metal exposure effects in childhood.

## Introduction

Trace elements play a role in a variety of biological processes in humans ranging from the transport of small molecules to the catalysis of redox reactions in cells.[1] While dietary intake of essential minerals is a requirement to sustain these biological processes, a variety of factors can result in improper absorption or utilization of the metals. This deficiency can

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### Ethical statement

This manuscript does not contain samples that were obtained from clinical studies and no personally identifiable patient data is included. Sample collection procedures followed the Helsinki Declaration guidelines regarding informed consent of human volunteers.

The authors declare that they have no conflict of interest.

result in adverse health effects to the individual, and these health effects can become doubly threatening when the individual is pregnant. In the same way, an exposure to an excess of an essential mineral or to a toxic metal can result in illnesses resulting from toxicity. For example, an excess of iron such as in hemochromatosis or excessive iron supplementation can result in increased cellular oxidative damage.[2]

Previous studies of health outcomes such as preeclampsia, low birth weight, intra-uterine growth restriction (IUGR), and childhood neurological effects in the fetus have focused on the analysis of serum and plasma from the mother, but the placenta has a more direct role in the development and sustenance of the fetus and can be viewed as a biological indicator of exposure of the fetus to toxic compounds before birth. Placenta tissue has previously been used as a matrix for studying fetal exposure to contaminants including toxic metals and for relation to the development of diseases.[3–5] The results of these investigations have provided conflicting findings regarding correlations between exposure and toxic metal concentrations in placenta tissue.

One potential explanation for the inconsistent findings of studies of trace metals in placenta may arise from the analytical method used to monitor the elemental concentrations. Inconsistent analytical methods between studies can result in variability in elemental concentrations, especially for trace elements that approach the limits of detection and quantification for the methods. This is especially true for the analysis of organ tissue, which can exhibit uneven spatial distribution, making the exact sampling and preparation methods crucial to the outcome of the study. In order to draw sound scientific conclusions from bioanalysis research projects, it is important to have a robust validated analytical method. [6,7] While some methods may focus on the analysis of a single analyte to investigate causal relationships between the analyte and the health outcome, biological systems are often more complicated and require multifactorial analyses to elucidate the complex interrelationships of a broad panel of analytes, or “-omics” type approaches. “Metallomics” is an encompassing term for the study of metals and their function in cells that has seen increased visibility and attention in recent years.[8–11] An additional subset of metallomics has recently been described that focuses on the essential nutritional metals or minerals that is referred to as “mineralomics.” [12,13] It is especially important to carefully validate broad panel “omics” methods to ensure that accurate results are achieved for all elements included in the analytical method.

The purpose of this study was to develop and validate an analytical method for a suite of toxic and essential metals in placenta tissue to support studies of the relationships between trace elements and health outcomes in pregnancy. The primary analyte of interest was zinc (Zn), but the method was also validated for the essential elements chromium (Cr), iron (Fe), cobalt (Co), nickel (Ni), and copper (Cu), and the toxic elements arsenic (As), cadmium (Cd), and lead (Pb) at varying concentrations. The validation was performed by bioanalytical method guidelines to demonstrate interday (same day) and intraday (different day) accuracy and precision for these elements in the human placental matrix. The method was shown to be valid for the elements of interest at levels comparable to or above the native concentrations in the tissues.

## Methods and Materials

### Samples and Materials

Two whole human placenta samples were purchased from Bioreclamation IVT (Baltimore, MD) and were stored at a nominal temperature of  $-80\text{ }^{\circ}\text{C}$  until time of use. Samples were collected from healthy volunteers and were used as received for preparation. Sample collection procedures followed the Helsinki Declaration guidelines regarding informed consent of human volunteers. Samples were fortified and calibration standards were prepared with the elements of interest by spiking with stock standards that were prepared with National Institute of Standards and Technology (NIST)-traceable standards of As, Cd, Co, Cr, Cu, Ni, Pb (all  $10\text{ }\mu\text{g/mL}$ ), Fe, and Zn (both  $1000\text{ }\mu\text{g/mL}$ ) purchased from High Purity Standards (Charleston, SC). Prepared samples and calibration standards were spiked with NIST-traceable Bi, In, Pr, and Y standards ( $10\text{ }\mu\text{g/mL}$ ) as internal standards that were also purchased from High Purity Standards. Samples were prepared with concentrated, 70% (v/v) nitric acid ( $\text{HNO}_3$ , Optima purity, Thermo-Fisher) and high purity 30% (v/v) hydrogen peroxide ( $\text{H}_2\text{O}_2$ , Ultrex purity, non-stabilized, Thermo-Fisher Scientific, Waltham, MA). Approximately  $18\text{ M}\Omega\text{ cm}^{-1}$  water was used in all sample preparation and analysis steps.

### Validation guidelines

Digested samples were prepared from a homogenized mixture of two whole human placenta and were used to prepare multiple replicates of samples that were spiked at seven concentration levels representing a range of concentrations relative to native levels (Table S1, Supplemental Information). Samples were prepared in triplicate at all concentration levels and on three separate, non-consecutive analytical days, and multiple replicates ( $n = 6$ ) were prepared at the two lowest nominal concentration levels to challenge detection and quantitation limits. The preparation of samples in at least triplicate was performed to assess interday (same day) accuracy and precision, and the repetition of analyses on at least three non-consecutive analytical days was intended to assess intraday (different day) accuracy and precision. The limit of detection (LOD), and lower limit of quantification (LLOQ), and selectivity were assessed by analysis of six replicates of the lowest concentration spike levels on the first analytical day. In addition, on the first analytical day, a series of matrix standards were prepared for comparison to solvent standards in order to assess matrix effects on analyte signal. On the second analytical day, three replicate samples were spiked and prepared at nominal concentrations exceeding analyte linear ranges to perform a dilution verification check.

### Instrumentation

All samples were prepared in a Class 100 clean hood to prevent contamination by atmospheric particulates. After digestion, the samples were analyzed for trace minerals using the Thermo (Bremen, Germany) X-Series I inductively-coupled plasma mass spectrometer (ICP-MS) equipped with a concentric glass nebulizer, collision cell technology (CCT) and Peltier-cooled glass spray chamber. Digestion of samples was performed using a DigiPrep graphite block digestion unit (SCP Science, Quebec) equipped for 15 mL digestion vessels.

## Digestion method

A scheme of the sample preparation method is shown in Figure 1 and details are provided in Table S2 (Supplemental Information). In order to promote an even distribution of tissue and promote optimal acid digestion of samples throughout, tissue samples were massed and homogenized by mixture with 2 times the sample weight of concentrated Optima-purity nitric acid in a heated circulating water bath and stirring until smooth. Nominal 0.500 mL homogenate aliquots were transferred to acid-washed 15-mL digestion vessels and directly spiked with elemental intermediate stock solutions at concentrations described in Table S1. After spiking, 0.500 mL of HNO<sub>3</sub> and 0.100 mL of H<sub>2</sub>O<sub>2</sub> were added to each vessel and the cap was placed loosely on the vessel. All vessels were placed in the graphite block digestion unit at 100 °C for 60 minutes and the digestion vessels were removed to allow to cool to room temperature. Samples were spiked with internal standard stock solution to a final concentration of 2 ng/mL to provide a measure of matrix effects on analyte ionization in the plasma and provide an indicator of instrument performance throughout the analysis, and the solutions were diluted to a final volume of 10 mL with DI water.

## Sample analysis

Sample analysis was performed by quadrupole ICP-MS (Table 1). Calibration ranges were prepared to contain both the native concentrations of the trace elements and the spiked levels of all elements. All elements were analyzed in kinetic energy discrimination (KED) mode with 10% H<sub>2</sub>/He collision cell gas to minimize polyatomic interferences. System suitability was monitored throughout the analysis by measurement of multiple matrix QC samples over a range of concentrations. Blank samples were measurement immediately after calibration to confirm a lack of analyte carryover. For the calibration, standards were accepted as passing if their back-calculated concentration was found to be within ±15% of the nominal concentration (±20% of the nominal concentration for the lowest concentration standard). At least 8 calibration standards were required to pass in order to accept the calibration curve as valid for the analytical day.

## Calculations

Analytical recovery of spiked samples was calculated using Eq. 1 for bias (relative error):

$$\%RE = \left( \frac{C_{meas} - C_{MxB}}{C_{nom}} \times 100\% \right) - 100\% \quad (1)$$

Where  $C_{meas}$  represents the measured concentration of each element,  $C_{MxB}$  represents the average measured concentration of the element in five replicates of matrix blank samples, and  $C_{nom}$  represents the nominal concentration for the element at that concentration level. Precision was calculated as the relative standard deviation (RSD) of repeated analyses of the same concentration level, shown in Eq. 2:

$$RSD = \frac{\sigma_n}{C_{meas}} \times 100\% \quad (2)$$

Where  $\sigma_n$  represents the standard deviation of all measurements.

Matrix impact was calculated for each element as the ratio of the slope of the calibration curve obtained from matrix standards ( $m_{Mx}$ ) to the slope of the calibration curve obtained from solvent standards ( $m_{SolV}$ ), Eq. 3:

$$MRF_{slope} = \frac{m_{Mx}}{m_{SolV}} \times 100\% \quad (3)$$

## Results

### Precision and accuracy

Results for intraday precision and accuracy for all three validation days are presented in Tables 2 and 3. For analytes (Table 2 and Table 3), precision and accuracy were required to be 20% and within  $\pm 20\%$ , respectively. These pre-defined acceptance criteria were achieved for all analytes and concentration levels on all three validation days, indicating that the validated analytical method was robust and fit for purpose.

### Analytical limits

Analytical measurement limits for the analyte suite are presented in Table 4. The lower limit of quantitation (LLOQ) was conservatively defined as the lowest nominal matrix QC level meeting the above described *a priori* defined precision and accuracy acceptance criteria on each of three validation days. To further challenge the analytical method, six replicates at the lowest two nominal concentration levels were prepared on the first validation day (Table S1). The limit of detection (LOD) for each analyte was then calculated as three times the standard deviation of the determined concentration at the LLOQ level on the first validation day. A total of six replicates were used to establish these figures of merit for all elements except for As and Cr, which had three replicates because it was necessary to shorten the linear range to meet precision and accuracy acceptance criteria. The upper limit of quantitation (ULOQ) for each analyte was defined as the highest level matrix standard included in the linear regression on all validation days. The reported preparation and analytical method has provided validated analytical ranges between one and two orders of magnitude depending on the identity and endogenous levels of each element.

### Dilution verification and matrix impact

Precision and accuracy data from the dilution verification matrix samples (Table S1) are presented in Table 5. Precision (%RSD) and accuracy (%RE) were required to be 20% and within  $\pm 20\%$ , respectively, for all other analytes. These pre-defined acceptance criteria were easily achieved, indicating the feasibility of diluting samples to have determined concentrations that fall within validated linear ranges. The pre-defined acceptance criterion was a slope ratio within  $\pm 20\%$  for all analytes in the suite. This criterion was achieved on the first validation day indicating that the overall matrix impact was not significant, so all precision and accuracy calculations were conducted using solvent curves.

## Discussion

### Digestion method

In ICP-MS methods and in bioanalytical methods, the sample preparation technique is of the utmost importance to ensure that consistent, reliable analytical results are obtained. It is necessary to ensure complete solubilization of the elements of interest uniformly throughout the sample solution and that minimal particulate matter remains in the solution, which could potentially clog the nebulizer of the instrument.

The first step of the sample preparation technique, homogenization of the tissue prior to digestion, has been successfully applied to previous method validation efforts for other analytes and tissues.[14] Homogenization assists in the degradation of the tissue in solution by providing more exposed surface area for acid digestion to occur. The process also helps to provide a uniform distribution of the element throughout the solution, as opposed to previous studies, which cut small masses of tissue from the whole organ.[4,5,15] This sample preparation method may not account for the possibility of the localization of elements within the mass of tissue, potentially resulting in artificially high or low results for the elements of interest.

Another difference in the method applied here is that digestion of the tissue is occurring through the use of a graphite digestion unit rather than microwave digestion. While microwave assisted digestion is appropriate for some types of samples, graphite heating unit digestion is a less-expensive, gentler digestion technique that is fit for the intended purpose described in this validation[16] The application of cost-effective, graphite heating digestion units can result in a less violent digestion than microwave digestion and a lower probability of sample loss due to frothing or bursting of digestion vessels. Further, the homogenization step serves to perform a predigestion of the sample, breaking down much of the organic matter before the addition of hydrogen peroxide, which produces more frothing than nitric acid alone. Previous studies have used cryogenic grinding of biological samples to homogenize the sample prior to microwave digestion with some success, but this method of homogenization requires the purchase of specialized equipment, which was not required for this reported method.[17]

### Method figures of merit

The LLOQ, LLOD, and ULOQ for each analyte, presented as  $\mu\text{g}$  element/g tissue, were robustly established and confirmed over three validation days. One previous study reported limits of detection for Cd, As, and Pb of 0.007 ng/g, 0.041 ng/g, and 0.01 ng/g, respectively, although the method used to determine the limits was not described.[15] Another investigation reported the LOD for the elements of interest as Fe = 0.0016 ng/g, Co = 0.002 ng/g, Cu = 0.0002 ng/g, Zn = 0.0058 ng/g, As = 0.02 ng/g, Cd = 0.001 ng/g, and Pb = 0.003 ng/g.[5] However, these values were calculated as 3 times the standard deviation of blank samples, but an LLOQ as defined above may be a more realistic estimate of values that can be consistently achieved due to the requirement of meeting predefined acceptance criteria for both accuracy and precision. Another study reported limits of detection of 0.4 ng/mL for Pb, 0.08 ng/mL for Cd, 4 ng/mL for Zn, and 1 ng/mL for Cu. These values appear to be

more consistent with those obtained in the current study. However, it is unclear whether any of these studies take into account dilution of the samples as a result of the sample preparation method, a phenomenon that is taken into account with the results reported here. Further, the values determined in this validation are supported by a more robust method of determination of the limits owing to the use of multiple measurements of quantifiable samples. This method of determination is also the explanation of the relatively high limits determined for Fe and Zn, elements that suffer from appreciable polyatomic interferences (Table S3) and lower sensitivity in KED mode. However, it is unlikely that samples of this type will ever exhibit concentrations that challenge the limit of detection owing to the high native concentrations of the elements in tissues.

### Broad panel analyses

Broad panel analyses such as the one described here can have a significant impact in elucidating the causal relationships that can produce harmful health outcomes. In the study of complex conditions such as preeclampsia where the exact etiological factors that result in their development are uncharacterized, a holistic approach to the complex interrelationships between different compounds, in this case minerals, can be informative. The approach described here can also provide a “shotgun” approach to toxicological studies, producing a maximum amount of information to determine correlations between health outcomes later in life and early life exposures to toxic metals or trace mineral deficiencies.

Another advantage to the “omics” approach described here is that the method described uses a minimal amount of sample for each measurement. “Omics” approaches often make use of minimal amounts of tissue or biological fluid in order to maximize the number of analyses that can be performed and minimize the invasiveness of sample collection, allowing the collection of a vast amount of data from a small amount of tissue. This approach can be particularly effective for performing research on samples that are stored in biological sample repositories, where the sample mass may be limited.

However, the use of the broad panel analysis method by ICP-MS can potentially suffer from challenges to some of the analytes of interest. Polyatomic interferences (Table S3) can result in difficulties in the consistent determination of the concentration of analytes without the use of corrective measures. In this analysis, all elements were analyzed in kinetic energy discrimination mode (KED mode, also known as collision cell mode or CCT mode) with H<sub>2</sub>/He gas to minimize the presence of polyatomic interferences, which could pose significant analytical challenges for a number of the elements in biological matrices, shown in Table S3.[18] Analysis in KED mode also results in decreased sensitivity for all elements compared to standard mode analysis. This issue is a particular concern for the analysis of Cr (ArC and CaC polyatomic interferences) and Co, where the native levels of the metals are very low, to the point of approaching the limits of detection. The exogenous toxic metal arsenic also suffers from a polyatomic interference in high salt solutions due to the formation of the ArCl species in the plasma, necessitating its analysis in KED mode.[19] It is possible that the use of higher resolution analysis modes, such as by sector-field ICP-MS (SF-ICP-MS) could provide higher sensitivity and lower limits of detection for challenging elements.[20]



In some cases, it may also be possible to analyze an alternate isotope that does not suffer from the same interferences or has interferences that contain lesser abundant isotopes of their own elements. This can allow the analyst to obtain higher sensitivity for the element of interest (if the isotope is sufficiently abundant that the signal is greater than that obtained with the primary isotope in KED mode). Conversely, lead and cadmium do not suffer from any commonly-observed polyatomic interferences in ICP-MS, making it possible to quantify the elements without the use of KED mode, such as in a multimode analytical run, allowing lower limits of detection and quantification for those analytes. However, this could in some cases present challenges resulting from the need to consume increased volumes of the sample as the instrument would perform two sets of scans, one in standard mode and another in KED mode. It is also possible to use interference correction equations to account for the formation of polyatomic interferences.[21] However, this practice is less accurate than other forms of interference exclusion, as it assumes constant rates of formation of polyatomic interferences, and requires thorough and repeated characterization of the correction factors necessary to accurately calculate the analyte counts throughout the analysis. This approach may be better suited to laboratories that do not have instruments equipped with collision cells, but can also over- or under-compensate depending on the levels of analyte and interference.

Finally, it is important to note that the methods described here have only definitively been shown to produce accurate analytical results for the elements reported here. The analysis of other elements, such as the alkali metals, has not been definitively demonstrated and will require additional work to verify the appropriateness of these digestion and analysis methods prior to their use.

## Applications

The analytical approach validated in this study was primarily developed to address correlations between the occurrence of preeclampsia and zinc levels in the placenta. However, the broad panel analysis validated here can be applied to a variety of health equity issues, some of which have been described in previous reports citing correlations between trace element concentrations and health outcomes. The studies reported thus far have touched on the potential roles for specific analytes in the prenatal or early childhood development of illnesses, but a detailed link to prenatal exposure and mineral nutrition status has not been explored and the method described here may be able to provide a more detailed link between these factors.

Preeclampsia is a serious multisystemic medical condition in pregnant women that is generally observed after 20 weeks of gestation and is characterized by new-onset elevated blood pressure and proteinuria, among a variety of other symptoms.[22,23] Advanced cases of preeclampsia can result in serious medical complications in the mother including stroke and heart disorders, as well as fetal outcomes including IUGR.[24] The development of the placenta has been found to be linked to some of the maternal symptoms of preeclampsia.[22] The condition is currently one of the leading causes of mother and infant morbidity and mortality, and is especially prevalent in the developing world, where malnutrition leads to decreased dietary intake of many essential nutrients.[25,26] The occurrence of preeclampsia



has also been linked to the development of noncommunicable diseases later in life.[27] The only effective cure for preeclampsia is delivery or abortion, making early manifestation of the disease a challenge to maintain maternal health until the fetus can mature sufficiently to survive outside of the womb. Early detection and the development of effective treatment regimens requires an understanding of the etiology of the disease.

A number of previous studies have shown a statistical correlation between micronutrient levels in the mother and the occurrence of preeclampsia.[28] Zinc is an essential trace micronutrient that plays a role in signaling proteins and catalytic enzymes, and its deficiency has been linked to a wide variety of adverse health outcomes.[29] Maternal plasma zinc concentrations have been found to naturally decrease during pregnancy, and some studies have found that individuals suffering from preeclampsia exhibit zinc levels that are decreased relative to normal pregnancies, with lower levels observed in more severe cases. [30–32] One study demonstrated a correlation between decreased zinc absorption and iron and folic acid dietary supplements that are commonly prescribed to pregnant women.[33] Maternal serum levels of other elements, including copper, zinc, and iron were shown to be lower in preeclamptic women than in normal pregnant women.[34] The ability to mitigate the occurrence of preeclampsia through the administration of dietary mineral supplements is a subject of some debate.[18,35] Other studies have shown that administration of multivitamins containing essential micronutrients preconception can result in a significant decrease in the rate of occurrence of preeclampsia and other adverse birth outcomes.[36] An appropriate analytical method that provides consistent concentrations of zinc and other micronutrients could shed some light on the etiology of the disease, providing novel strategies for preventing the development of the disease.

However, beyond the study of preeclampsia, there are a variety of other conditions and health outcomes that can be studied with the broad panel approach validated here.[37] Low maternal iron levels have been suggested to influence birth weight, and iron and folic acid supplementation has previously been correlated to low maternal zinc uptake.[35] Prenatal iron deficiency has also been linked to neurocognitive developmental effects in early childhood.[38] Vitamin B<sub>12</sub> (cobalamin, a cobalt compound) deficiencies in maternal diets have been linked to low fetal growth and other adverse symptoms including cleft palate.[39] Nutritional deficiencies are very frequently observed in developing countries, making dietary uptake and maternal health an urgent health equity issue.[40]

Additionally, exposures to toxic metals may be of greater concern in developing countries as a result of less stringent controls on environmental release, less developed testing and treatment systems, and the availability of consumer products containing toxic metals.[41] Maternal occupational exposure to nickel has been found to be linked to the development of musculoskeletal birth defects.[38] Elevated lead levels are observed in children in developing countries, which can have a strongly adverse effect on neurocognitive development.[42] Arsenic exposure from the water supply has been observed in some eastern Asian countries including Bangladesh, which can result in neurocognitive effects and other health outcomes.[43]

## Conclusions

The analysis of biological tissues for trace metals can provide invaluable information regarding the etiology of diseases and other outcomes, as well as providing an assessment of environmental exposure to contaminants that may be correlated to adverse health outcomes later in life. The accurate determination of the levels of trace elements requires a robust sample preparation and analysis method that provides reliable, precise results. The method of sample homogenization and acid digestion detailed in this report has been validated for inter- and intraday accuracy and precision and matrix effects on the analytical signal and has demonstrated a high degree of reproducibility of results. The results suggest that the method can provide strong analytical results in support of studies to assess maternal and natal health and their relationship to the environmental exposome.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Abbreviations

<b>ICP-MS</b>	Inductively Coupled Plasma-Mass spectrometry
<b>NIST</b>	National Institute of Standards and Technology
<b>ELOQ</b>	Estimated limit of quantitation
<b>LOD</b>	Limit of Detection
<b>KED</b>	Kinetic Energy Discrimination
<b>SF-ICP-MS</b>	Sector field inductively coupled plasma mass spectrometry
<b>IUGR</b>	Intra-uterine growth restriction

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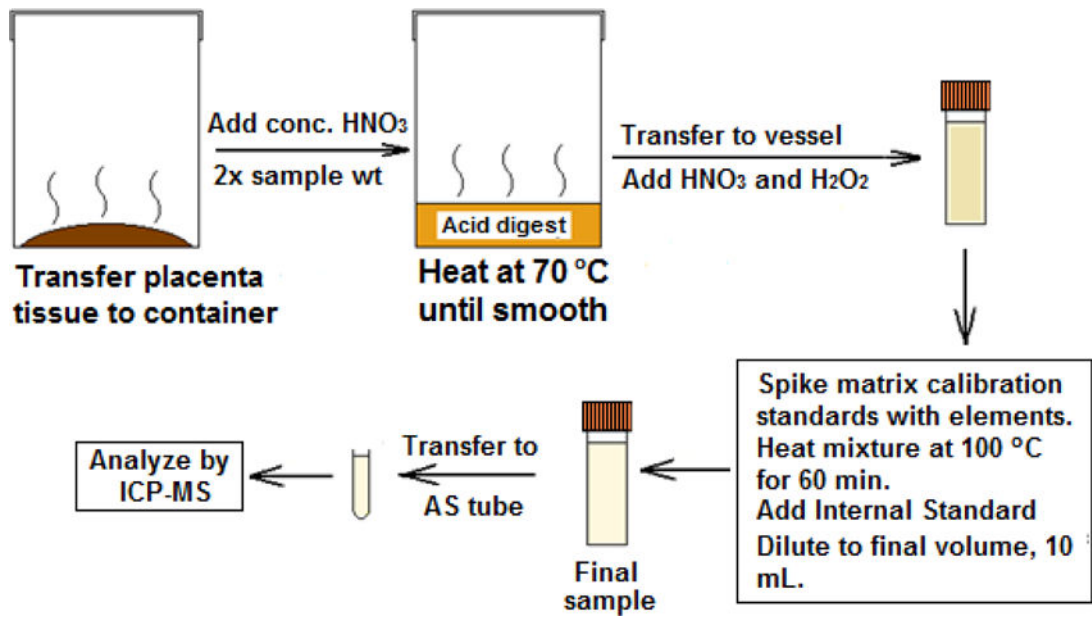
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**Figure 1.**  
A flow diagram showing the sample preparation method developed and validated here.

**Table 1**

Instrumental parameters for the ICP-MS analysis of digested placenta samples

<b>Instrument</b>	Thermo X Series I ICP-MS
<b>Software</b>	PlasmaLab ICP-MS Software, Version 2.3.0.161
<b>Nebulizer Type</b>	Glass Concentric
<b>Spray Chamber</b>	Peltier cooled
<b>Collision Cell Gas</b>	10% H <sub>2</sub> /He
<b>Injector</b>	1.8 mm
<b>Analyte Masses</b>	<sup>52</sup> Cr, <sup>56</sup> Fe, <sup>59</sup> Co, <sup>60</sup> Ni, <sup>63</sup> Cu, <sup>66</sup> Zn, <sup>75</sup> As, <sup>111</sup> Cd, <sup>208</sup> Pb
<b>Internal Standard Masses</b>	<sup>89</sup> Y, <sup>115</sup> In, <sup>141</sup> Pr, <sup>209</sup> Bi
<b>Cones</b>	Ni HPI

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Table 2

Intraday Precision and accuracy at all concentration levels for As, Cd, Co, Cr, and Cu

	Nominal [X] <sup>μ</sup> (μg/g)	Day 1 RE (%)	Day 1 RSD (%)	Day 2 RE (%)	Day 2 RSD (%)	Day 3 RE (%)	Day 3 RSD (%)
<b>As</b>							
	0.00855	-14	4.9	3	16	6	5.2
	0.0214	4	2.3	10	6.9	14	1.9
	0.0427	12	1.7	15	5.0	14	2.1
	0.150	15	1.9	16	1.2	19	0.86
	0.214	20	1.5	15	0.68	20	1.0
<b>Cd</b>							
	0.00214	4	14	5	10	2	13
	0.00427	3	4.3	4	3.9	2	12
	0.00855	-2	3.4	-0.1	3.5	7	0.71
	0.0214	1	1.8	-0.4	1.6	1	2.5
	0.0427	-1	0.84	-0.4	0.44	-0.1	2.2
	0.150	-4	0.78	-2	1.2	-0.2	0.56
	0.214	-1	0.58	-2	1.5	0	1.8
<b>Co</b>							
	0.00427	3	5.4	-2	2.7	-15	1.8
	0.00855	-2	1.1	-1	1.8	-4	3.5
	0.0214	-2	2.1	-6	1.8	-8	1.1
	0.0427	-6	1.3	-4	2.0	-6	0.4
	0.150	-8	1.5	-8	0.065	-8	0.66
	0.214	-0.2	0.20	-8	0.42	-8	0.11
<b>Cr</b>							
	0.00855	-12	20	-17	3.8	3	12
	0.0214	14	20	-9	2.5	-5	1.2
	0.0427	-7	2.3	-10	0.61	-3	3.7
	0.150	-8	0.91	-9	0.39	-6	0.39
	0.214	12	11	-9	0.32	-7	0.37
<b>Cu</b>							

Nominal [X] <sup>a</sup> (µg/g)	Day 1 RE (%)	Day 1 RSD (%)	Day 2 RE (%)	Day 2 RSD (%)	Day 3 RE (%)	Day 3 RSD (%)
0.427	-10	3.7	-17	4.4	-17	2.8
1.07	-12	0.99	-11	1.1	-12	0.66
3.21	-11	1.5	-10	1.7	-7	5.1
4.27	-4	2.6	-10	2.3	-10	0.61

<sup>a</sup>[X] = analyte concentration.

**Table 3**

Intraday Precision and accuracy at all concentration levels for Fe, Ni, Pb, and Zn

	Nominal [X] <sup>μ</sup> (μg/g)	Day 1 RE (%)	Day 1 RSD (%)	Day 2 RE (%)	Day 2 RSD (%)	Day 3 RE (%)	Day 3 RSD (%)
Fe	21.4	-11	13	-15	6.6	-14	4.6
	32.1	-10	4.2	-12	2.1	-11	3.2
	42.7	-12	3.5	-12	2.1	-8	0.89
	64.1	-9	1.5	-10	1.6	-7	1.4
	85.5	-8	1.1	-10	1.6	-8	0.61
	128	-9	2.2	-10	0.62	-10	1.0
	214	2	1.3	-11	0.45	-10	0.81
Ni	0.0855	-6	1.8	-6	2.7	-8	0.26
	0.214	-5	8.0	-9	1.2	-8	0.60
	0.321	-8	1.8	-8	0.61	-9	1.6
	0.427	-6	0.88	-9	0.13	-7	0.65
	1.07	-7	0.43	-8	0.26	-7	0.26
	3.210	-8	1.1	-10	0.42	-8	0.67
	4.27	-2	0.83	-10	0.74	-9	0.30
Pb	0.00427	-12	8.4	5	13	-17	1.8
	0.00855	-11	1.9	6	2.7	-11	1.3
	0.0214	-7	0.69	-1	0.35	-6	1.2
	0.0427	-5	0.59	-1	1.8	-4	0.75
	0.150	-5	0.21	-4	0.47	-5	0.14
	0.214	-4	1.7	-4	0.074	-4	0.44
Zn	4.27	1	10	-4	7.7	0.4	4.6
	8.55	0.2	3.7	-2	1.8	-0.2	0.75
	12.8	-2	3.1	-3	3.1	0.9	0.55
	17.1	-3	0.68	-4	0.59	0.4	1.2

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Nominal [X] <sup>a</sup> (µg/g)	Day 1 RE (%)	Day 1 RSD (%)	Day 2 RE (%)	Day 2 RSD (%)	Day 3 RE (%)	Day 3 RSD (%)
21.4	-4	1.3	-4	0.74	0.7	0.84
32.1	-4	1.6	-2	0.51	2	4.6
42.7	3	1.3	-2	2.2	14	3.5

<sup>a</sup>[X] = analyte concentration.

**Table 4**

Analytical figures of merit

Analyte	LLOQ <sup>a</sup> (µg/g)	ULOQ <sup>b</sup> (µg/g)	LOD <sup>c</sup> (µg/g)
As	0.00855	0.427	0.001
Cd	0.00214	0.427	0.001
Co	0.00427	0.427	0.0007
Cr	0.00855	0.427	0.005
Cu	0.427	8.55	0.04
Fe	21.4	214	8
Ni	0.0855	8.55	0.04
Pb	0.00427	0.427	0.0009
Zn	4.27	42.7	1

<sup>a</sup> Lower limit of quantitation (LLOQ) defined as lowest matrix QC level with %RE within ±15% and %RSD < 20% on all three validation days.

<sup>b</sup> Upper limit of quantitation (ULOQ) defined as highest matrix standard included in regression.

<sup>c</sup> Limit of detection (LOD) calculated as three times the standard deviation of determined concentrations for matrix QC replicates (n = 3 or 6) on the first validation day.

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**Table 5**

## Dilution verification

Analyte	RE (%)	RSD (%)
As	1	0.40
Cd	1	1.4
Co	-3	0.96
Cr	-7	0.75
Cu	10	1.2
Fe	-1	4.0
Ni	-1	0.17
Pb	2	0.36
Zn	11	0.6

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