

1 **Direct determination of lead in bones using slurry sampling high-**
2 **resolution continuum source electrothermal atomic absorption**
3 **spectrometry**

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5 Lenka Husáková*, Tereza Šídová, Lucie Ibrahimová, Monika Svíželová, Tomáš Mikysek

6 *Department of Analytical Chemistry, Faculty of Chemical Technology, University of*

7 *Pardubice, Pardubice, Studentska 573 HB/D, CZ-532 10, Czech Republic*

8

9 * Corresponding author: Tel. +420 466 037 029 fax: +420 466 037 068

10 E-mail address: Lenka.Husakova@upce.cz

11

12 **Abstract**

13 Straightforward, quick, sensitive and reliable method is introduced for determination
14 of lead in bones using slurry sampling and high-resolution continuum source electrothermal
15 atomic absorption spectrometry (HR-CS-ETAAS). The spectral interference caused by the
16 molecular absorption of PO molecules with rotational fine structure coinciding with the
17 analyte absorption at the most sensitive resonance line 217.001 nm in time was identified and
18 successfully corrected by applying a mathematical correction algorithm using the spectrum
19 obtained by vaporization of hydroxyapatite. The slurry preparation and measuring conditions
20 were determined by means of a response surface methodology. Experiments were designed
21 according to a 2^{7-4} replicate ($n = 3$) fractional factorial design for seven factors (particle size,
22 glycerol and HNO_3 concentration, sonication time, concentration of chemical modifier,
23 pyrolysis and atomization temperature) each at three different levels, including central points.
24 The optimized conditions were 100 mg of a ground sample with particle size $< 315 \mu\text{m}$,
25 dilution in a liquid-phase composed by 10 % w/w glycerol, 5.0 % w/w nitric acid solutions,
26 sonication time of 2 min and final slurry volume of 10.0 mL. Detection limit of $9.1 \mu\text{g kg}^{-1}$
27 and characteristic mass 7.6 pg were achieved using the suggested method under the optimized
28 experimental conditions. Sufficient analyte stabilization was achieved by using $1 \mu\text{g Pd}$ and
29 $50 \mu\text{g citric acid}$. Accurate data were obtained with the use of matrix-free calibration. The
30 accuracy of the method was established by analysing NIST SRM 1486 Bone Meal. Further,
31 the results acquired for ten river otter samples by slurry sampling were compared with those
32 determined after microwave-assisted digestion by inductively coupled plasma time of flight
33 mass spectrometry (TOF-ICP-MS) to assess the accuracy of the method. The results obtained
34 by the two methods were compared using a paired t -test (at 95% confidence level) and
35 showed no significant difference. The precision of the introduced method was better than
36 5.5 %.

- 37 **Keywords:** Lead determination; Bone analysis; High-resolution continuum source AAS;
- 38 Electrothermal atomic absorption spectrometry; Matrix effects; Design of experiments;
- 39 Microwave-assisted digestion; Inductively coupled plasma mass spectrometry

40

41 **1. Introduction**

42 Lead determination in bone and teeth samples has a wide-ranging importance in the clinical ¹,
43 ², environmental ³⁻⁵, forensic ^{6,7} and archaeological ^{8,9} sciences as these calcified tissues are
44 especially useful for pollution assessments ^{3,4,10}, nutritional and clinical status evaluations ^{1,2},
45 toxicology analysis ⁸ or for identification purposes ^{11,12}. The bone Pb levels of individuals
46 who were not exposed to Pb may reach ^{1,4,13-16} up to 20 mg kg⁻¹, however, the concentration
47 is generally lower ^{1,15,16}, only occasionally exceeding 10 mg kg⁻¹. Several analytical methods
48 such as X-ray fluorescence (XRF) ^{13,14,17}, proton-induced X-ray emission (PIXE) ¹,
49 inductively coupled plasma optical emission spectrometry (ICP-OES) ^{18,19}, two-jet plasma
50 atomic emission spectrometry ²⁰, inductively coupled plasma mass spectrometry (ICP-MS)
51 alone ^{2,11,12,21} or combined with laser ablation (LA-ICP-MS) ^{9,22}, hydride generation (HG-
52 ICP-MS) ²³, atomic fluorescence spectrometry (AFS) ²⁴ or electrothermal atomic absorption
53 spectrometry (ETAAS) ^{1,3,6,12,15,16,21,25-32} previously assessed the validity and limitations for
54 the purpose of quantification of Pb in both human and animal samples. Among these methods,
55 ETAAS ^{1,5,10,25,27,31-36} still represents the most commonly used one in routine analytical
56 practice because of its high sensitivity, favourable detection limits and relatively low cost of
57 the instrumentation. However, due to the complex matrix, laborious sample preparation,
58 relatively low analytical concentration and high volatility of some lead species, the analysis of
59 bone or teeth samples is difficult ^{1,33,34,37}.

60 Serious analytical problems related to spectral interference from phosphate matrix
61 during ETAAS determination of Pb in bone and teeth samples at the main analytical lines:
62 217.001 and/or 283.306 nm, were reported by a number of authors ^{5,27,35}. To alleviate these
63 problems attributed to PO molecular absorption, instrumentation with several different
64 background correction systems have been employed ^{3,5,6,10,15,16,21,25-27,30-33,36,38,39}. Among

65 conventional line source atomic absorption spectrometers (LS AAS), Zeeman systems using
66 both transverse^{21, 31, 33, 36} and longitudinal^{3, 25, 32, 39} arrangement were found^{3, 5, 16, 21, 25, 30, 33,}
67^{34, 36, 39} to be the most reliable for this purpose. However, it has been shown that under the
68 influence of the magnetic field the rotational lines of molecular spectrum of the gaseous
69 phosphorus monoxide (PO) have split thus making the analysis of samples with high
70 phosphorus content when using Zeeman effect background corrections prone to errors^{39, 40}.

71 Many chemical modifiers act as efficient thermal stabilizers of phosphorus containing
72 species thus delaying and depressing the background signal^{33, 38, 41}. Lead can be then
73 determined in bones or in phosphates matrix using LS ETAAS^{25, 27, 33, 34, 38, 41} via matrix
74 modification. In many cases, modifiers such as Pd(nitrate)²⁷, Pd/Mg(nitrates)^{25, 34}, W-Rh
75 permanent modifier²⁵ had to be applied to further attenuate the interferences produced by
76 phosphate matrix as well as to increase the thermal stability of Pb to yield reliable results for
77 digested^{25, 27} or solid²⁵ calcified tissues. Application of phosphate modifiers has also been
78 widely recommended for thermal stabilization of Pb during the analysis of bone digestates^{1, 3,}
79³¹, as calcium hydroxyapatite, i.e. the main component of bone matrix and an endogenous
80 source of phosphate, was shown^{1, 3, 33} to be less efficient for this purpose. Although an
81 excessive background absorption may be associated with the use of phosphate modifiers^{39, 41},
82 these drawbacks can be controlled by using phosphate based mixed modifiers e.g. with Pd⁴¹,
83 Ca^{21, 33} or Mg^{32, 33} or by employing an adequate amount of the phosphate modifier³⁹.

84 Most of the studies presented in the literature^{1-3, 5, 6, 11, 12, 16, 21, 26, 27, 29, 30, 32, 35, 36} report
85 that the determination of metals in bone and teeth samples is usually carried out by methods
86 involving some type of sample digestion using nitric acid and its mixtures. In several cases,
87 except for wet digestion, preliminary time-consuming dry-ashing step is additionally required
88^{10, 26, 35} which may however significantly affect the recovery of Pb^{27, 42}.

89 On the other hand, direct introduction of solid samples or slurries into the graphite
90 furnace may be more efficient as it reduces the speed and price of the analysis, risk of analyte
91 loss and contamination and does not usually involve the use of toxic and/or concentrated
92 chemicals making it more in compliance with green chemistry trends⁴³⁻⁴⁷. In addition, it may
93 significantly increase the detection power that is very important in this particular case as
94 reliable analysis of Pb in tissues of non-exposed population still constitutes an analytical
95 challenge^{14, 18}. In this term, direct solid sample analysis may be an alternative to methods
96 employing concentration/separation steps^{15, 29, 37}. While these methods are usually time
97 consuming, and prone to serious systematic errors²⁸, they may help to reduce the matrix
98 effects. On the other hand, spectral interferences caused by molecular absorption due to
99 diatomic molecules with pronounced fine structure⁴⁸ may be the major obstacle in the
100 application of direct solid samples analysis due to the limited background correction
101 capability especially for the LS AAS²⁸.

102 In contrast to the conventional AAS systems, HR-CS-AAS enables both detailed
103 observations of the structured background signals as well as the efficient correction due to its
104 unsurpassed background correction capabilities⁴⁹⁻⁵². Previously, a number of new methods
105 for trace elements analysis in a wide variety of complex matrices by HR-CS-ETAAS using
106 direct solid sampling were elaborated without any interference^{43, 49, 50, 53}. However, according
107 to the best our knowledge to this date no method has been published for the purpose of direct
108 determination of Pb in bone solids by this technique.

109 In this work, the development of such reliable and environmentally friendly method
110 suited for routine direct analysis in tissues of non-exposed animals, which constitutes an
111 analytical challenge is described. The ‘visibility’ of the spectral environment around the
112 analytical line with the employment of the least squares background correction (LSBC) for
113 elimination of the fine structure of PO band directly hampering the determination of Pb at the

114 resonance line 217.001 nm together with the response surface methodology (RSM) employed
115 for optimization of experimental conditions has been shown to significantly facilitate the
116 method development.

117

118 **2. Experimental**

119 **2.1 Instrumentation**

120 The analyses were performed using the model contraAA 600 high-resolution continuum
121 source atomic absorption spectrometer (Analytik Jena AG, Jena, Germany) equipped with
122 transversely heated graphite furnace and autosampler MPE 60. Xenon short-arc lamp with a
123 nominal power of 300 W, operating in a hot-spot mode was the light source. The double
124 monochromator (prism and echelle grating) combined with a CCD array detector with 588
125 pixels ensures the unique resolution with a spectral bandwidth of below 2 pm/pixel. Peak area
126 absorbance values were measured. The primary analytical line 217.001 nm for Pb was used
127 for the measurement. Pyrolytically coated graphite tubes (Analytik Jena, Part no. 407–
128 152.023 Schunk, Germany, Batch-No: 103074348-46/13) with preinstalled PIN platform and
129 integrated contacts for transverse heating were used. Argon (99.999 %, Linde Gas, Inc.,
130 Czech Republic) was used as the sheating gas. The internal gas flow in the graphite tube was
131 interrupted during the atomization step and also during one part of the pyrolysis stage.

132 The microwave-assisted digestion of samples was performed in the SpeedwaveTM
133 Xpert (Berghof, Germany) microwave system with the maximum total output of the
134 microwave generator 2000 W. The sample throughput of the Xpert system was increased by
135 using the Multitube System (MT) employed with the DAK-100X digestion vessels⁵⁴. This
136 arrangement allows simultaneous digestion of three samples in one DAK-100X PTFE vessel
137 by placing three MT PFA tubes into each of the vessels.

138 Inductively coupled plasma time of flight mass spectrometer Optimass8000 (GBC

139 Scientific Equipment Pty. Ltd., Australia)⁵⁵ was used in several cases for comparative
140 measurements.

141 Samples of slurries were sonicated using a Sonorex Super RK52 ultrasonic bath (35
142 kHz, RF-power 240 W; BANDELIN electronic GmbH & Co. KG, Germany).

143 The diffraction patterns (Cu K α , $\lambda = 1.5418 \text{ \AA}$) were recorded on powdered samples
144 using a D8 Advance diffractometer (Bruker AXS, Germany) with Bragg-Brentano Θ - Θ
145 goniometer (radius 217.5 mm) equipped with Ni-beta filter and LynxEye detector. The scan
146 was performed at room temperature from 5 to 70° (2 Θ) in 0.01° step with a counting time of
147 2 s per step.

148 The scanning electron microscopy (SEM) images were recorded with a VEGA3
149 TESCAN model under high vacuum at 20 kV accelerating voltage. Energy-dispersive X-ray
150 spectroscopy (EDX) was used to characterize the presence of major elements in bone samples
151 using Bruker Nano GmbH, X Flash Detector 410 model, Germany.

152 CoolSafe 4-15 L bench-top freeze dryer (LaboGene, Denmark) for drying of samples
153 and enhancing both stability of a dry powder and analyte in a dry state as well was used
154 throughout this study.

155 Wig-L-Bug 30 (Crescent Dental Mfg. Co.) vibration mill was used for production of
156 powders from individual samples. Particle size distribution was measured by a
157 Mastersizer 2000/MU (Malvern Instruments, Ltd., GB).

158

159 **2.2 Reagents and standards**

160 Lead stock solution with 1 g L⁻¹ Pb in 3% HNO₃ was obtained from Analytika, Ltd.
161 (Czech Republic). Nitric acid (65%, w/w) of Selectipur quality and glycerol (99.6%, p.a.)
162 were purchased from (Lach-Ner, Neratovice, Czech Republic). Laboratory grade Triton X-
163 100 (4-(1,1,3,3-Tetramethylbutyl)phenyl-polyethylene glycol) and synthetic hydroxyapatite

164 (99.8%, trace metals basis) were obtained from Sigma-Aldrich (USA). Hydrogen peroxide
165 (Trace Select, $\geq 30\%$, w/w) was purchased from Fluka (Switzerland). The solution of 1 g L^{-1}
166 of Pd in $10\% \text{ v/v HCl}$ was bought from SCP Science (Canada). Solution of 25 g L^{-1} of citric
167 acid (Lachema, Brno, Czech Republic) was prepared by dissolving this pro-analysis grade salt
168 in water. Solutions were prepared using ultrapure water of $0.055 \mu\text{S cm}^{-1}$ conductivity
169 obtained using the Milli-Q® (Millipore, USA) water purification system.

170

171 **2.3 Quality assurance and quality control**

172 Commercially supplied quality control material SRM® 1486 Bone Meal intended
173 primarily to evaluate analytical methods used for the determination of selected major, minor,
174 and trace elements in bone and in material of a similar matrix was purchased from National
175 Institute of Standards & Technology, USA. Particle size smaller than $355 \mu\text{m}$ was ensured by
176 the material certificate.

177

178 **2.4 Sample collection, storing, handling and preparation**

179 Ten rib samples from Eurasian otters, mostly killed in traffic accidents, were obtained
180 from the ALKA Wildlife organization, which is a group of experts on ecology and
181 conservation of wildlife in the Czech Republic. In addition for method development purposes,
182 rabbit bones were obtained from a local farm.

183 At first, all adhering tissues and tendons were removed from the samples (Fig. S1a,b)
184 with a ceramic knife made from zirconium dioxide. After that, bones were cut into smaller
185 pieces (Fig. S1c,d) and marrow deposits were removed by a stainless steel scraper.

186 Around 1 g of the sample was placed into the containers (Fig. S1c) wherein the
187 material was dried. The containers were tightly closed, and the material was subsequently
188 freeze dried in the closed container under the conditions developed and tested successfully

189 with animal bone tissues in order to allow for sample variability. Before the freeze drying
190 process, the samples were placed in a deep freezer at $-80\text{ }^{\circ}\text{C}$ for 24 hours to provide a
191 necessary conditioning for low temperature drying. Total drying times of about 29 hours were
192 employed, with the room temperature held in the first drying stage for about 1 hour and a total
193 chamber pressure of 0 Pa, followed by a 24-hour drying stage at $-111\text{ }^{\circ}\text{C}$ and 0 Pa and a final
194 drying stage at room temperature and 0 Pa pressure for 4 hours.

195 The dried samples were placed in a stainless steel vial (1" height x 1/2" diameter) with
196 stainless steel ball pestle (1/4" diameter) (Fig. S1d) and ground in a vibration mill. The
197 powder (Fig. S1d) was then sieved manually by using sieves with hole sizes 315, 160 and 54
198 μm . Ground bone samples were kept in sealed plastic vials and stored at $-20\text{ }^{\circ}\text{C}$ until analysis.
199

200 **2.5 Procedure for slurry analysis**

201 The slurries were prepared using the following procedure. An accurately weighed
202 amount of about 100.0 mg of sample (fractions $< 315, 160$ or $54\text{ }\mu\text{m}$) was transferred into a 10
203 mL calibrated flask and three drops of 2% (w/w) Triton X-100 solution in ethanol were added
204 to ensure wetting of the sample. Appropriate amounts of glycerol and/or nitric acid were
205 added to yield a final solution containing 0, 10, 20 % (w/w) glycerol and 0, 2.5 and 5 % (w/w)
206 HNO_3 . The slurries were then sonicated for 2, 6 or 10 min in an ultrasonic bath. Thereafter,
207 approximately 1 mL of the slurry was transferred into the autosampler cup and then only
208 manual shaking was performed a right before the injection. Finally, 25 μL of the slurry plus
209 0–10 μL of the chemical modifier mixture containing 0.5 g L^{-1} of Pd and 25 g L^{-1} of citric
210 acid were injected into the graphite furnace by the auto-sampler and subjected to the heating
211 program presented in Table 1. The preparation of the reagent blank was subjected to the
212 procedure as outlined above, to correct for any possible trace amount of the analyte in the
213 reagents used for slurry preparation. The slurries, similarly as all the other samples

214 investigated in this study, were prepared in triplicate.

215 The concentrations of five standard solutions used to obtain the calibration curves
216 ranged from 0 to 40 $\mu\text{g L}^{-1}$ of Pb. The calibrations and standard additions were controlled by
217 the instrument software. Next to the aqueous calibration, two standard additions were made.
218

219 **2.6 Procedure for microwave digestion**

220 For comparative purposes, an adapted microwave-assisted procedure⁵⁶ was used.
221 Sample mass of 100.0 mg was weighed into the MT-tubes and 1 mL of 65 % (w/w) HNO_3
222 and 1 mL of 30% H_2O_2 (w/w) were added. These MT-tubes were placed into the outer
223 digestion vessel containing 15 mL of HNO_3 (65%, w/w) and H_2O_2 (30%, w/w) mixture (1:1,
224 v/v). This ensured that the level of the HNO_3 - H_2O_2 mixture was higher in the outer vessel
225 than in the PFA tubes. The vapour pressures were thus compensated and the evaporation of
226 the solution from the PFA tubes was forestalled⁵⁴. Samples were digested according to
227 following 5-steps program: (i) 10 min at 130 °C and 20% power (ramp 5 min), (ii) 10 min at
228 160 °C and 40% power (ramp 5 min), (iii) 15 min at 200 °C and 60% power (ramp 5 min)
229 (iv–v) 5 min at 50 °C and 0% power (ramp 1 min). The resulting solutions were diluted to
230 10 mL with deionised water.

231 Determination of Pb in the mineralized samples by a reference TOF-ICP-MS method
232 was done using aqueous standard calibration curve with Rh as an internal standard⁵⁶.

233

234 **2.7 Experimental design and statistical data treatment**

235 The response surface methodology⁵⁷⁻⁵⁹ was used to estimate the main effects of the
236 selected variables onto analytical response and to find the optimal conditions for the sample
237 preparation and measuring conditions as well. The variables selected in this study were: (1)
238 glycerol concentration (% , w/w), (2) HNO_3 concentration (% , w/w), (3) time of sonification,

239 (4) size of particles, (5) pyrolysis temperature, (6) atomization temperature, (7) concentration
240 of chemical modifier.

241 The Box, Hunter, and Hunter^{57, 58} two-level ($2^{7-4} + 3C$, $n = 3$) fractional factorial
242 design (FFD)^{57, 58} was used for designing the experimental data. Thus, only 33 experiments
243 were run instead of 384 required for a full factorial design (2^7 , $n = 3$) as the number of
244 experiments in FFD is given by ($2^{k-p} \times n + C \times n$), where k is the number of variables, C is the
245 number of central points, n number of replicates and p a whole number that indicates how
246 fractionated the experimental design is. All factors were evaluated at two levels, low (denoted
247 as -1) and high (denoted as $+1$). The central point of the design space (middle value denoted
248 as 0), i.e. the experiment, in which all the parameters have a value which is the average
249 between the low and high level, was also added to the experiment. The investigated maximum
250 and minimum levels of the variables are shown in Table 2. The responses for each experiment
251 calculated as the mean of integrated absorbance obtained for Pb in SRM® 1486 Bone Meal
252 are summarized in Table S1.

253 The experimental data were processed using the Statistica 12 computer program
254 (StatSoft, Inc., USA), Minitab 18.1 (Minitab Inc., USA) and QC Expert™ 2.5, TriloByte
255 Statistical Software, (Pardubice, Czech Republic).

256

257 **3. Results and discussion**

258 **3.1 Evaluation of spectral interference on Pb determination**

259 Spectral interference may be recognized and controlled by using the visibility of the
260 spectral environment at high resolution in HR-CS-AAS as a diagnostic tool⁴⁸. The
261 wavelength-resolved absorbance spectrum as collected on contrAA® 600 during the
262 atomization of a slurry of NIST SRM 1486 Bone Meal shown in Fig. 1a demonstrates the
263 well-structured background in the surroundings of the most sensitive analytical line of Pb at

264 217.001 nm. This complex background may be attributed to the diatomic molecule PO caused
265 by vaporization of bone matrix as this in agreement with the reference spectrum recorded with
266 hydroxyapatite (Fig. 1b). The structured background caused by PO molecules is thus
267 recognized to be the main spectral interference in the determination of Pb in bone samples by
268 ETAAS. No other potential interferences in the neighborhood of the lead line at 217.001 nm
269 by e.g. a secondary aluminium line at 216.883 nm and a secondary iron line at 217.130 nm
270 were observed, although these have been reported ⁶⁰ previously for determination of Pb in
271 various biological samples.

272 The presence of structured background absorption caused by PO makes the
273 determination extremely difficult or virtually impossible using LS ETAAS, as none of the
274 currently available background correction systems of such instruments is able to correct
275 reliably for this kind of absorption ⁴⁰. However, as it is evident from the literature ^{40, 48, 60}, this
276 interference can be fully controlled by HR-CS-ETAAS during the determination of several
277 analytes in various matrices under optimized conditions. It is important to highlight, that
278 chemical composition of each sample is responsible for the fine structured background ^{33, 38-40}
279 and even for the same analyte in a similar matrix the situation may be quite different.
280 The HR-CS-ETAAS instrument software enables to correct the fine-structured background
281 caused by the interfering molecules by means of the least-squares background correction
282 (LSBC) using reference spectra ^{48, 50}. However, when the PO bands do not overlap with the
283 analytical line it is not necessary to employ LSBC because of the high spectral and time
284 resolution of the equipment the atomic absorbance signal can be resolved in time from the
285 molecular structures by the setting integration limits adjusted to integrate only the atomic
286 signal ^{48, 60}. While, using this approach the interference-free determination can be achieved for
287 Pb at 217.001 nm line using direct solid sample analysis in a variety of biological materials ⁶⁰,
288 the situation for determination of Pb in bones is more complex. In this case a direct

289 coincidence in time between the atomic and molecular absorption is observed (Fig. S2) using
290 the atomization temperature of 1500 °C and higher. Although, according to the literature ⁴⁸,
291 the phosphates in biological materials, which are the source of interfering PO molecules, are
292 volatilized only above 1700 °C it can be seen from Fig.S2 that background absorption with
293 pronounced fine structure is observed even at lower temperatures. These observations are also
294 in accordance to those published previously by Borges at al. ⁶⁰. Lowering the atomization
295 temperature to less than 1500°C may lead to problems associated with broad transient signal
296 resulting in worse precision and longer atomization step.

297 Reference spectrum belonging to the molecule causing the spectral interference is
298 necessary for successful elimination of the fine structured background by its subtraction using
299 LSBC ⁴⁰. In the literature, PO spectra obtained by vaporization of $\text{NH}_4\text{H}_2\text{PO}_4$ ⁴⁰ or
300 $(\text{NH}_4)_2\text{HPO}_4$ are most widely employed as a reference. However, thermal behaviour of these
301 compounds ⁶¹ is significantly different compared to bone phosphate and thus for generation of
302 reference spectra hydroxyapatite was used.

303 The PO reference spectra were recorded using 20–250 µg of hydroxyapatite which
304 was introduced into the graphite furnace via the injection of 2–25 µL of 1.0 % (w/w)
305 $\text{Ca}_5(\text{OH})(\text{PO}_4)_3$ slurry together with the chemical modifier. The reference spectra were stored
306 in the method for subsequent subtraction from the sample spectra. During the analysis of real
307 samples the reference spectra recorded using 200 µg of hydroxyapatite removed essentially all
308 the molecular absorption structures, as is shown in Fig. 1b. The robustness of the method to
309 correct for structured background is documented (Fig. S3) for up to 250 µg of $\text{Ca}_5(\text{OH})(\text{PO}_4)_3$,
310 which corresponded to the hypothetical situation that the slurry samples of 1 % (w/w)
311 concentration prepared according to the description in section 2.5 would contain nothing but
312 hydroxyapatite. The robustness of the method for the correction of spectral effect is important
313 as the composition of bone matrix may vary among the analysed samples as can be seen from

314 the results of XRD (Fig. S4) or EDX (Table S2) analysis. As can be seen from the data
315 presented in Table S2, the bone matrix is rich not only in phosphorus but also in Ca, Mg or
316 Na. As the presence of Ca and Mg significantly affects the thermal behavior of interfering
317 molecules and also delays the appearance of Pb^{33, 39, 40}, a strong and fine-structured
318 background coinciding in time and wavelength with the analytical line of Pb 217.001 nm may
319 be attributed just to a coexistence of a high concentration of P and Ca, eventually also with
320 Mg.

321

322 **3.2 Optimization of the experimental conditions for slurry analysis**

323 Variables including particle size, pyrolysis and atomization temperature, volume of
324 chemical modifier, concentration of stabilizing and extracting agent, and ultrasonic agitation
325 and their influence onto the analytical results was investigated using the response surface
326 methodology for data designed by Box Hunter & Hunter^{57, 58}. Each independent variable,
327 being previously reported^{44, 62} as the most important in terms of influencing the accuracy of
328 the analytical procedure when using the slurry technique, was tested at a high (+) and a low
329 (–) level, as shown in Table 2. The ranges of values are in agreement with practice in slurry
330 sampling analysis and our preliminary experiments⁵².

331 The Pareto chart depicted in Fig. 2, where the vertical line that corresponds to the 95%
332 limit indicating statistical significance, visualizes the effects and significance of the variables.
333 This figure reveals that the glycerol and nitric acid concentration, pyrolysis temperature and
334 chemical modifier appeared to have a significant effect, while the other variables (extraction
335 time, particle size and atomization temperature) were not significant factors in the studied
336 range.

337 Response surface regression analysis employed to describe the data presented in
338 Table S1 revealed that a second- order polynomial model as shown in the following equation

339 was the best to fit the data: $y = 0.1853 (1.17 \times 10^{-2}) + 0.01417 (1.04 \times 10^{-3}) x_1 + 0.011953$
340 $(9.3 \times 10^{-4}) x_2 - 0.000123(9 \times 10^{-6}) x_3 - 0.004575 (5.80 \times 10^{-4}) x_4 - 0.000733 (3.1 \times 10^{-5}) x_1^2$,
341 where y stands for the predicted integrated absorbance, x_1 through x_4 stand for the settings of
342 the glycerol and nitric acid concentration (x_1 , resp. x_2), pyrolysis temperature (x_3) and volume
343 of the chemical modifier (x_4). Standard deviations of the estimates are given in parentheses.
344 The equation illustrates the reduced quadratic model obtained from the analysis by
345 eliminating the terms found statistically insignificant. The elimination of statistically
346 insignificant terms helped to increase the capability for precise predictions from the model.
347 The results of analysis of variance which is essential to test the significance and adequacy of
348 the model are presented in Table S3. Significance was evaluated by determining the
349 probability level that the F -statistic calculated from the data is less than 5%. Data given in this
350 table demonstrates that the model is significant at the 5% confidence level since p values are
351 smaller than 0.05. The large p value for lack of fit (>0.05) presented in Table S3 shows that
352 the model is valid. The model adequacies were checked by R^2 , adjusted- R^2 and predicted- R^2 .
353 The coefficient of determination (R^2) of the model was 0.983, which indicated a good fit
354 between predicted values and the experimental data points. In addition, this implies that
355 98.3 % of the variations for analytical response are explained by the independent variables,
356 and this also means that the model does not explain only about 1.7 % of variation. Both R^2
357 and predicted R^2 values obtained (see Table S3) proved the goodness of fit of the regression
358 model.

359 As is depicted on Fig. 2, the analytical response is significantly improved by
360 increasing the HNO_3 concentration and, at the same time by decreasing the concentration of
361 glycerol and volume of chemical modifier or pyrolysis temperature. While an increase in the
362 glycerol concentration up to 10% has a promoting effect onto the analytical response, the
363 effect of a higher investigated concentration was the opposite. When using 10% glycerol

364 concentration, the suspension can be stabilized for more than 10 s, which is long enough to
365 ensure a representative aliquot of the sample to be introduced into the cuvette after its manual
366 shaking right before the injection. The presence of nitric acid in the slurry solution within all
367 the investigated range affected significantly the rate of the analyte extraction and improved
368 the precision of the measurement as well. On the other hand, the further effect of sonication
369 treatment at different time intervals has not been shown to have a significant promoting effect
370 onto sensitivity. Sufficient analyte extraction and slurry homogenization was thus achieved
371 even at the lowest sonication times.

372 In the analysis of bones, the particle size in the range from approx. < 54 to 315 μm did not
373 affect the analytical response (see Fig. 2), although in general the benefit from smaller
374 particles sizes, especially for refractory and more dense samples is frequently mentioned in
375 the literature⁴⁴ as it can positively influence the accuracy of the measurement. For otter and
376 rabbit samples only 3 min of grinding resulted in a fine and totally ground powder ensuring
377 that 90% of total volume was composed by particles lower than 175 μm (see Fig. S5,6). The
378 particle diameter medians of both samples were about 50 μm (Fig. S5).

379 Palladium with citric acid chemical modifier, currently being well established in our
380 laboratory⁶³⁻⁶⁵, was applied to promote the thermal stabilization of the analyte. As it can be
381 seen from Figure S7, already 1 μg of Pd with 50 μg of citric stabilizes Pb up to 1300° C,
382 similarly as some of other chemical modifiers, which were previously suggested for this
383 purpose^{41, 66}. Although the presence of a chemical modifier during the real sample analysis
384 impacts positively the sensitivity and the precision (see Table S1), higher amounts of
385 chemical modifier leads to gradual decrease of the analytical response meaning that the lowest
386 investigated amounts are adequate for the analyte stabilization. This behavior is probably the
387 consequence of the over-stabilization of Pb by the modifier⁶⁷.

388 3D surface plots for overall response desirability were built to show the effects of two

389 variables within their studied ranges and to visualize the tendency of important factors to
390 influence the analytical response (see Fig. 3). The desirability values for the analytical
391 response (integrated absorbance) were set 0 for minimum (0.0017), 0.5 for middle (0.07465)
392 and 1.0 for maximum (0.1476). These values were set based on the experimental data
393 presented in Table S1. The individual desirability score of 1.0 selected as the target value for
394 the optimization of the dependent factors shows its optimum value. The optimum conditions
395 for the most important factors were as follows: 10 and 5 % (m/m) of glycerol and HNO₃,
396 respectively, pyrolysis temperature 1100 °C and 2 μL of chemical modifier mixture
397 containing 0.5 g L⁻¹ of Pd and 25 g L⁻¹ of citric acid. The optimal values for atomization
398 temperature, sonication time, particle size and/or grinding time were set at 1600 °C, 2 min, <
399 315 μm and 3 min, respectively.

400

401 **3.3 Figures of merit**

402 The slope of the standard additions curve for Pb concentration was found to be quite
403 comparable with the slope obtained by the external calibration technique over a range where
404 the response is linear since under the optimized conditions they differ less than 10 %. These
405 results demonstrate that the quantification of Pb in investigated samples by slurry sampling
406 can be performed using the external calibration technique with aqueous standard solutions.
407 This yields a model described by the following equation: $Q_A = 1.20 \times 10^{-2} (2.64 \times 10^{-4}) [Pb] -$
408 $6.43 \times 10^{-5} (8.39 \times 10^{-6}) [Pb]^2$ (where Q_A = integrated absorbance and $[Pb]$ is the concentration of
409 the analyte in μg L⁻¹; standard deviation of the estimate is given in parentheses). R-squared
410 which indicates the percentage of variation that can be explained by the regression equation
411 equals 99.99 %.

412 The limit of detection (LOD) and limit of quantification (LOQ) expressed as the
413 concentration given by an integrated absorbance corresponding to three times and ten times of

414 the standard deviation of ten measurements of a sample blank were $0.091 \mu\text{g L}^{-1}$ and
415 $0.30 \mu\text{g L}^{-1}$, respectively. These values are equivalent to $9.1 \mu\text{g kg}^{-1}$ and $30 \mu\text{g kg}^{-1}$ of Pb in
416 the original sample and are sufficiently low for quantification of lead concentrations in human
417 and animal bones even for a non-exposed population ^{4, 68, 69}.

418 The characteristic mass, which gives an integrated absorbance of 0.0044 s, was found to be
419 7.6 pg. This value is coherent with those of 5.6–8 pg presented elsewhere ^{40, 48, 60} for Pb at the
420 most sensitive analytical line of 217.001 nm using HR-CS-ETAAS with transversely heated
421 graphite tube atomizer.

422 The figures of merit obtained in this work at 217.001 nm ($m_0 = 7.6 \text{ pg}$, LOD $9.1 \mu\text{g kg}^{-1}$) are
423 better than those reported in the literature for Pb determination in bone and teeth samples by
424 LS ETAAS at 283.3 nm. The values range between 12.9–57.2 pg and 22–600 $\mu\text{g kg}^{-1}$ for
425 characteristic mass value and limit of detection, respectively, using direct analysis of the
426 samples ²⁵ or acid digestion as the sample treatment ^{1, 26, 27, 31-33, 39}.

427 The line 283.3 nm, although less sensitive, is usually preferred ^{1, 27} in routine analytical
428 practice because it is less interfered by PO molecular structures ^{48, 66}. The proposed method
429 achieved nearly the same LOD value as those of $10 \mu\text{g kg}^{-1}$ reported by Borges et al. ⁶⁰ for
430 determination of Pb at 217.001 nm in various biological samples like human hair, bovine
431 muscle, dogfish liver, pig kidney, lobster hepatopancreas, oyster tissue and bovine blood by
432 high-HR-CS-ETAAS with direct solid sampling.

433

434 **3.4 Accuracy and precision**

435 The reference material NIST® SRM® 1486 Bone Meal, with a certified content of
436 lead being $1.335 \pm 0.014 \text{ mg kg}^{-1}$, was used to study the accuracy of the introduced method.
437 Three replicates of this material were analyzed and the mean value with 95% confidence level
438 obtained was $1.34 \pm 0.08 \text{ mg kg}^{-1}$. As confirmed by a *t*-test (95% confidence level), no

439 significant difference was found between the certified and the determined values (*t*-test, 95%
440 confidence level).

441 The method accuracy was also checked by analyzing real samples and comparison of
442 the data obtained by the proposed method with those acquired after microwave assisted
443 digestion followed by TOF-ICP-MS analysis (see Table 3). The concentration of Pb in the real
444 samples varied from 0.28 to 2.05 mg kg⁻¹. The *p*-value for the paired *t*-test being 0.702
445 suggests that at the significance level of 0.05 the compared results obtained by both methods
446 showed no statistical difference.

447 A linear regression analysis of the results obtained with TOF-ICP-MS and HR-CS-
448 ETAAS using the method of weighted least squares⁷⁰ yielded a slope of 0.914 (95%
449 confidence interval CI 0.76–1.07), intercept of 0.061 (95% CI -0.018–0.140) and R² = 0.990.
450 These results demonstrate that the calculated slope and intercept do not differ significantly
451 from the values of 1 and 0, respectively, and that the results achieved by both procedures are
452 comparable.

453 The precision of the method was assessed in terms of intra-day and inter-day
454 comparison. The analysis of the slurry samples three times during the same day was done to
455 assess the intra-day precision. Inter-day precision was calculated after the analysis of the same
456 samples on three different days during one week. Within each series, every sample was
457 analysed in three replicates to assess the relative standard deviation (RSD). The precision of
458 the method was found satisfactory as the RSD values of intra-day and inter-day studies were
459 typically found to be below 5.5 %, which can be seen in Table 3.

460

461 **4. Conclusions**

462 It was demonstrated in this work that under optimized experimental conditions, an
463 interference-free, accurate and precise determination of Pb in bone samples can be

464 successfully performed using HR-CS-ETAAS, slurry sampling analysis, Pd with citric acid
465 chemical modifier and calibration using aqueous standard solutions. The fine-structured
466 background observed at 217.001 nm caused by the presence of PO molecules, was completely
467 corrected by employing LSBC. The spectrum obtained by vaporization of hydroxyapatite was
468 required to reproduce the matrix and generate the correct reference spectrum. The attained
469 detection limit of $9.1 \mu\text{g kg}^{-1}$ was low enough to perform reliable Pb determinations in tissues
470 of non-exposed animals. The results gained using the described method are comparable to
471 those achieved using a method employing microwave assisted acid digestion. Another
472 important advantage of this method is the use of diluted nitric acid to extract the analyte from
473 the bone samples, which reduces safety problems and analysis costs. It may be expected that
474 the procedure will furthermore allow simple reliable analysis of some other trace elements in
475 calcified tissues strongly sensitive to the presence of phosphate matrix without any extensive
476 sample preparation. In comparison with other reported methods in the references which were
477 used for direct analysis of solid samples to determine Pb in calcified tissue samples, such as
478 XRF, PIXE or LA-ICP-MS, the presented method offers a better limit of detection and good
479 analytical precision for low concentration levels of the analyte, does not need to prepare
480 pellets, has a small dependence on the size and structure of the particles to be analyzed, is
481 interference-free, inexpensive and does not need an internal standard. On the other hand, it
482 lacks when compared in terms of linear dynamic range or capabilities for performing depth
483 profiling analysis, elemental mapping or in vivo measurements.

484

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490

491 **Figure captions**

492 **Fig. 1.** Wavelength-resolved absorbance spectra in the vicinity of the 217.001 nm analytical
493 line using HR-CS-ETAAS recorded for NIST SRM® 1486 Bone Meal (a) without correction
494 and (b) after correction using LSBC and PO as a reference spectrum in the presence of
495 1 µg Pd + 50 µg citric acid. Pyrolysis and atomization temperatures were 1000 and 2400 °C,
496 respectively.

497

498 **Fig. 2** Pareto chart of the standardized effects in the fractional factorial design 2^{7-4} for the
499 study of variables (1) glycerol concentration, (2) HNO₃ concentration, (3) sonification time,
500 (4) particle size, (5) pyrolysis temperature (Tp), (6) atomization temperature (Ta) and (7)
501 volume of chemical modifier. The L and Q letters indicate linear and quadratic effect of the
502 factor, respectively.

503

504 **Fig. 3.** Response surfaces from 2^{7-4} design for the desirability produced the best absorbance
505 (target is maximized) of Pb in the bone slurry as a function of glycerol and nitric acid
506 concentration, pyrolysis temperature (Tp) and volume of the chemical modifier.

507

508

509 **References**

- 510 1. M. A. Deibel, J. M. Savage, J. D. Robertson, W. D. Ehmann and W. R. Markesbery, *J*
511 *Radioan Nucl Ch Ar*, 1995, 195, 83-89.
- 512 2. T. R. Helliwell, S. A. Kelly, H. P. J. Walsh, L. Klenerman, J. Haines, R. Clark and N.
513 B. Roberts, *Bone*, 1996, 18, 151-157.
- 514 3. V. Spevackova and J. Smid, *Spectrochim Acta B*, 1999, 54, 865-871.
- 515 4. F. Barbosa, J. E. Tanus-Santos, R. F. Gerlach and P. J. Parsons, *Environ Health Persp*,
516 2005, 113, 1669-1674.
- 517 5. I. Baranowska, K. Czernicki and R. Aleksandrowicz, *Sci Total Environ*, 1995, 159,
518 155-162.
- 519 6. S. I. Al-Qattan and M. A. Elfawal, *J Forensic Leg Med*, 2010, 17, 325-328.
- 520 7. S. C. Zapico and D. H. Ubelaker, *Ageing Res Rev*, 2013, 12, 605-617.
- 521 8. E. Sguazza, D. Gibelli, M. Caligara, D. Di Candia, P. M. Galimberti and C. Cattaneo,
522 *Archaeometry*, 2016, 58, 152-158.
- 523 9. M. J. Kohn, J. Morris and P. Olin, *J Archaeol Sci*, 2013, 40, 1689-1699.
- 524 10. L. Gerhardsson, A. Akantis, N. G. Lundstrom, G. F. Nordberg, A. Schutz and S.
525 Skerfving, *J Trace Elem Med Bio*, 2005, 19, 209-215.
- 526 11. T. A. Hinnens, R. Hughes, P. M. Outridge, W. J. Davis, K. Simon and D. R. Woolard,
527 *J Anal Atom Spectrom*, 1998, 13, 963-970.
- 528 12. V. G. Thomas, A. M. Scheuhammer and D. E. Bond, *Sci Total Environ*, 2009, 407,
529 3494-3502.
- 530 13. A. C. Todd, P. J. Parsons, S. Carroll, C. Geraghty, F. A. Khan, S. Tang and E. L.
531 Moshier, *Phys Med Biol*, 2002, 47, 673-687.
- 532 14. T. M. Ambrose, M. Al-Lozi and M. G. Scott, *Clin Chem*, 2000, 46, 1171-1178.
- 533 15. E. Hac, W. Czarnowski, T. Gos and J. Krechniak, *Sci Total Environ*, 1997, 206, 249-
534 254.
- 535 16. J. Scancar, R. Milacic, M. Benedik and P. Bukovec, *Clin Chim Acta*, 2000, 293, 187-
536 197.
- 537 17. A. C. Todd, E. L. Moshier, S. Carroll and S. W. Casteel, *Environ Health Persp*, 2001,
538 109, 1115-1119.
- 539 18. M. Grotti, M. L. Abemoschi, S. Dalla Riva, F. Soggia and R. Frache, *Anal Bioanal*
540 *Chem*, 2005, 381, 1395-1400.
- 541 19. H. W. Kuo, S. M. Kuo, C. H. Chou and T. C. Lee, *Sci Total Environ*, 2000, 255, 45-
542 54.
- 543 20. N. P. Zaksas, T. T. Sultangazieva and V. A. Gerasimov, *Anal Bioanal Chem*, 2008,
544 391, 687-693.
- 545 21. K. M. Hetter, D. J. Bellis, C. Geraghty, A. C. Todd and P. J. Parsons, *Anal Bioanal*
546 *Chem*, 2008, 391, 2011-2021.
- 547 22. W. Castro, J. Hoogewerff, C. Latkoczy and J. R. Almirall, *Forensic Sci Int*, 2010, 195,
548 17-27.
- 549 23. V. Yilmaz, Z. Arslan and L. Rose, *Anal Chim Acta*, 2013, 761, 18-26.
- 550 24. B. Beltran, L. O. Leal, L. Ferrer and V. Cerda, *J Anal Atom Spectrom*, 2015, 30, 1072-
551 1079.
- 552 25. D. Santos, F. Barbosa, S. S. de Souza and F. J. Krug, *J Anal Atom Spectrom*, 2003, 18,
553 939-945.
- 554 26. L. E. Wittmers, A. Alich and A. C. Aufderheide, *Am J Clin Pathol*, 1981, 75, 80-85.
- 555 27. K. S. Subramanian, J. W. Connor and J. C. Meranger, *Arch Environ Con Tox*, 1993,
556 24, 494-497.

- 557 28. F. J. Langmyhr and I. Kjuus, *Anal Chim Acta*, 1978, 100, 139-144.
- 558 29. H. Mosbaek, P. E. Holm and J. C. Tjell, *J Anal Atom Spectrom*, 2003, 18, 1489-1492.
- 559 30. M. Kim, C. Kim and I. Song, *Food Addit Contam*, 2003, 20, 149-153.
- 560 31. H. B. Pereira, A. S. Luna, F. W. Herms and R. C. de Campos, *J Brazil Chem Soc*,
561 2004, 15, 487-490.
- 562 32. R. A. de Sousa, C. M. Sabarense, G. L. P. Prado, K. Metze and S. Cadore, *Talanta*,
563 2013, 104, 90-96.
- 564 33. Y. Y. Zong, P. J. Parsons and W. Slavin, *J Anal Atom Spectrom*, 1996, 11, 25-30.
- 565 34. I. Iavicoli, G. Carelli, N. Castellino and G. Schlemmer, *Fresen J Anal Chem*, 2001,
566 370, 1100-1104.
- 567 35. J. Simon and T. Liese, *Fresen Z Anal Chem*, 1983, 314, 483-486.
- 568 36. A. C. Todd, P. J. Parsons, S. D. Tang and E. L. Moshier, *Environ Health Persp*, 2001,
569 109, 1139-1143.
- 570 37. C. Y. Zhang, Y. N. Wang, X. R. Cheng, H. B. Xia and P. Liang, *J Chin Chem Soc-
571 Taip*, 2011, 58, 919-924.
- 572 38. Y. Y. Zong, P. J. Parsons and W. Slavin, *Spectrochim Acta B*, 1994, 49, 1667-1680.
- 573 39. Y. Y. Zong, P. J. Parsons and W. Slavin, *Spectrochim Acta B*, 1998, 53, 1031-1039.
- 574 40. A. R. Borges, E. M. Becker, L. L. Francois, A. de Jesus, M. G. R. Vale, B. Welz, M.
575 B. Dessuy and J. B. de Andrade, *Spectrochim Acta B*, 2014, 101, 213-219.
- 576 41. D. L. Tsalev, V. I. Slaveykova and P. B. Mandjukov, *Spectrochim Acta Rev*, 1990, 13,
577 225-274.
- 578 42. H. A. McKenzie and L. E. Smythe, *Quantitative Trace Analysis of Biological
579 Materials: Principles and Methods for Determination of Trace Elements and Trace
580 Amounts of Some Macro Elements*, Elsevier, Amsterdam, 1988.
- 581 43. M. Resano, M. Aramendia and M. A. Belarra, *J Anal Atom Spectrom*, 2014, 29, 2229-
582 2250.
- 583 44. N. J. Miller-Ihli, *Anal Chem*, 1992, 64, A964-A968.
- 584 45. P. T. Anastas and M. M. Kirchhoff, *Accounts Chem Res*, 2002, 35, 686-694.
- 585 46. S. L. C. Ferreira, M. Miro, E. G. P. da Silva, G. D. Matos, P. S. dos Reis, G. C.
586 Brandao, W. N. L. dos Santos, A. T. Duarte, M. G. R. Vale and R. G. O. Araujo, *Appl
587 Spectrosc Rev*, 2010, 45, 44-62.
- 588 47. J. Sardans, F. Montes and J. Penuelas, *Spectrochim Acta B*, 2010, 65, 97-112.
- 589 48. B. Welz, H. Becker-Ross, S. Florek and U. Heitmann, *High-Resolution Continuum
590 Source AAS: The Better Way to Do Atomic Absorption Spectrometry*, 2006.
- 591 49. R. G. O. Araujo, B. Welz, F. Vignola and H. Becker-Ross, *Talanta*, 2009, 80, 846-
592 852.
- 593 50. B. Welz, M. G. R. Vale, E. R. Pereira, I. N. B. Castilho and M. B. Dessuy, *J Brazil
594 Chem Soc*, 2014, 25, 799-821.
- 595 51. H. Becker-Ross, S. Florek and U. Heitmann, *J Anal Atom Spectrom*, 2000, 15, 137-
596 141.
- 597 52. L. Husakova, I. Urbanova, M. Safrankova and T. Sidova, *Talanta*, 2017, 175, 93-100.
- 598 53. W. Boschetti, M. Orlando, M. Dullius, M. B. Dessuy, M. G. R. Vale, B. Welz and J.
599 B. de Andrade, *J Anal Atom Spectrom*, 2016, 31, 1269-1277.
- 600 54. L. Husakova, I. Urbanova, T. Sidova, T. Cahova, T. Faltys and J. Sramkova, *Int J
601 Environ an Ch*, 2015, 95, 922-935.
- 602 55. L. Husakova, I. Urbanova, L. Audrlicka-Vavrusova, J. Sramkova, T. Cernohorsky, M.
603 Bednarikova and L. Pilarova, *Microchim Acta*, 2011, 173, 173-181.
- 604 56. L. Husakova, I. Urbanova, J. Sramkova, T. Cernohorsky, A. Krejcova, M.
605 Bednarikova, E. Frydova, I. Nedelkova and L. Pilarova, *Food Chem*, 2011, 129, 1287-
606 1296.

- 607 57. G. E. P. Box, W. G. Hunter and J. S. Hunter, *Statistics for experimenters : an*
608 *introduction to design, data analysis, and model building*, Wiley, New York, 1978.
609 58. D. C. Montgomery, *Design and analysis of experiments*, Wiley, New York, 3rd edn.,
610 1991.
611 59. M. A. Bezerra, R. E. Santelli, E. P. Oliveira, L. S. Villar and L. A. Escaleira, *Talanta*,
612 2008, 76, 965-977.
613 60. D. L. G. Borges, A. F. da Silva, B. Welz, A. J. Curtius and U. Heitmann, *J Anal Atom*
614 *Spectrom*, 2006, 21, 763-769.
615 61. D. R. Lide, *CRC handbook of chemistry and physics : a ready-reference book of*
616 *chemical and physical data*, CRC, Boca Raton, Fla. ; London, 90th edn., 2009.
617 62. C. Santos, F. Alava-Moreno, I. Lavilla and C. Bendicho, *J Anal Atom Spectrom*, 2000,
618 15, 987-994.
619 63. I. Urbanova, L. Husakova and J. Sramkova, *Environ Monit Assess*, 2013, 185, 3327-
620 3337.
621 64. L. Husakova, J. Sramkova, T. Cernohorsky and I. Urbanova-Dolezalova, *Talanta*,
622 2009, 77, 1504-1509.
623 65. L. Husakova, T. Cernohorsky, J. Sramkova, K. Hubackova and I. Dolezalova, *Anal*
624 *Chim Acta*, 2008, 614, 38-45.
625 66. J. F. Rego, A. Virgilio, J. A. Nobrega and J. A. G. Neto, *Talanta*, 2012, 100, 21-26.
626 67. Z. Arslan and J. F. Tyson, *Microchem J*, 2007, 86, 227-234.
627 68. L. Kubaszewski, A. Ziola-Frankowska, M. Frankowski, P. Rogala, Z. Gasik, J.
628 Kaczmarczyk, A. Nowakowski, M. Dabrowski, W. Labedz, G. Miekisiak and R.
629 Gasik, *J Orthop Surg Res*, 2014, 9.
630 69. G. A. Drasch and J. Ott, *Sci Total Environ*, 1988, 68, 61-69.
631 70. M. Meloun, M. Forina and J. Militký, *Chemometrics for analytical chemistry*, Ellis
632 Horwood, New York, 1992.
633
634