

## Supplementary Information

### **„Direct determination of lead in bones using slurry sampling high-resolution continuum source electrothermal atomic absorption spectrometry“**

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**Figure S1** Bone sample preparation and processing: (a) a rib sample with remnants of adherent tissue, (b) sample of the rib after cleaning with a ceramic knife, (c) bone specimens in sterile freeze-drying containers, (d) freeze dried bone sample, (e) a stainless steel vial (1" height x 1/2" diameter) with stainless steel ball pestle (1/4" diameter), (f) resulting sample after 5 min of grinding time, scale given by CZ coin of 23 mm diameter.



a



b



c



d

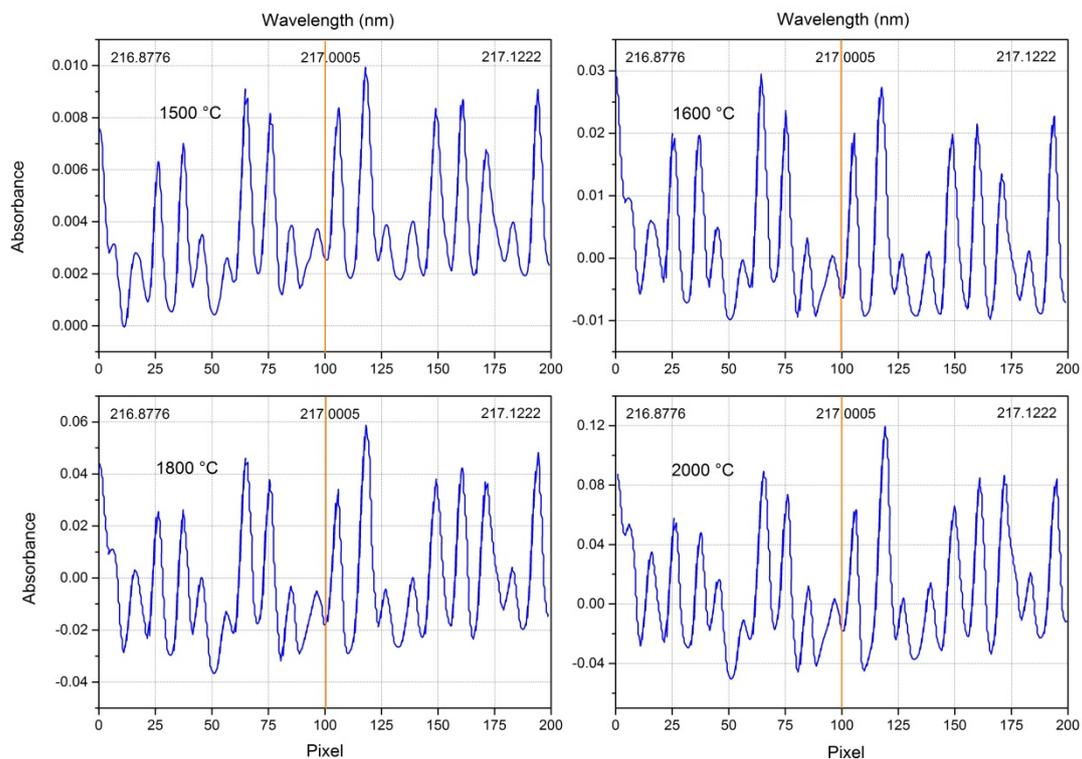


e

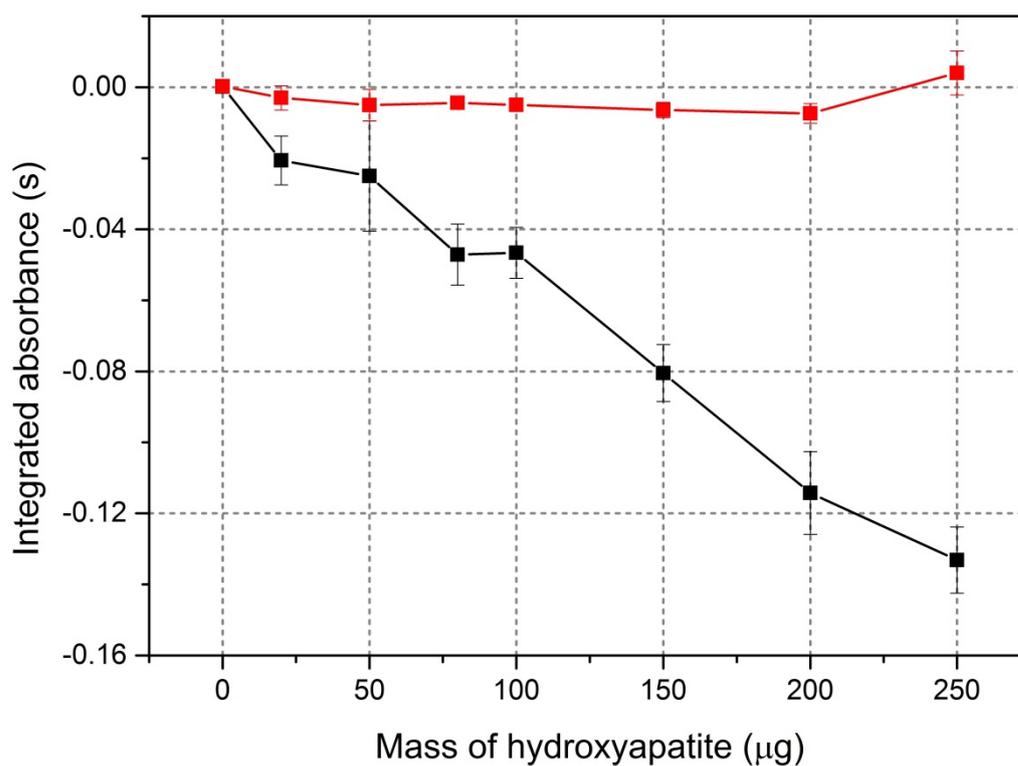


f

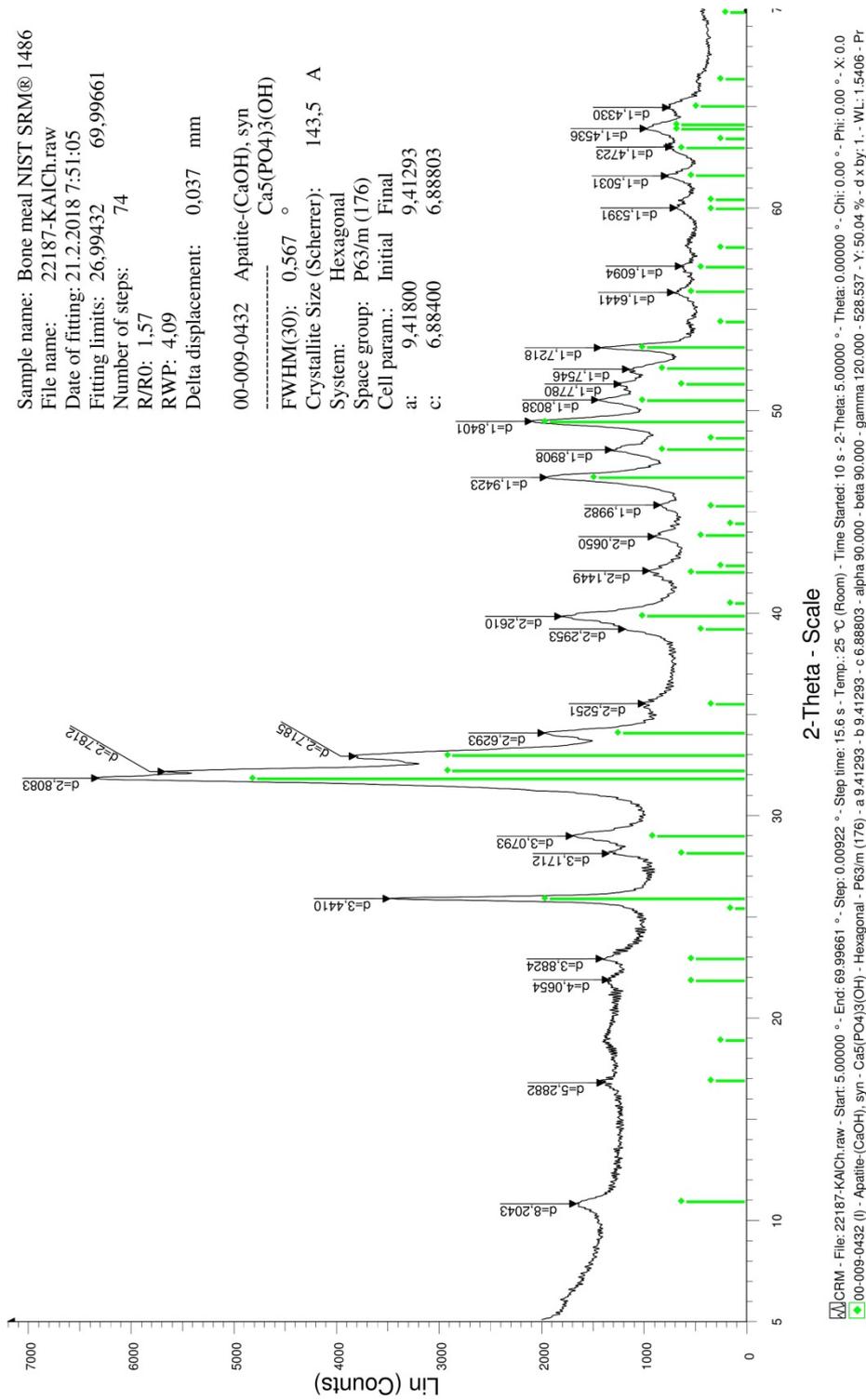
**Figure S2.** Absorbance spectra of PO recorded at different temperatures in the vicinity of the lead line at 217.001 nm



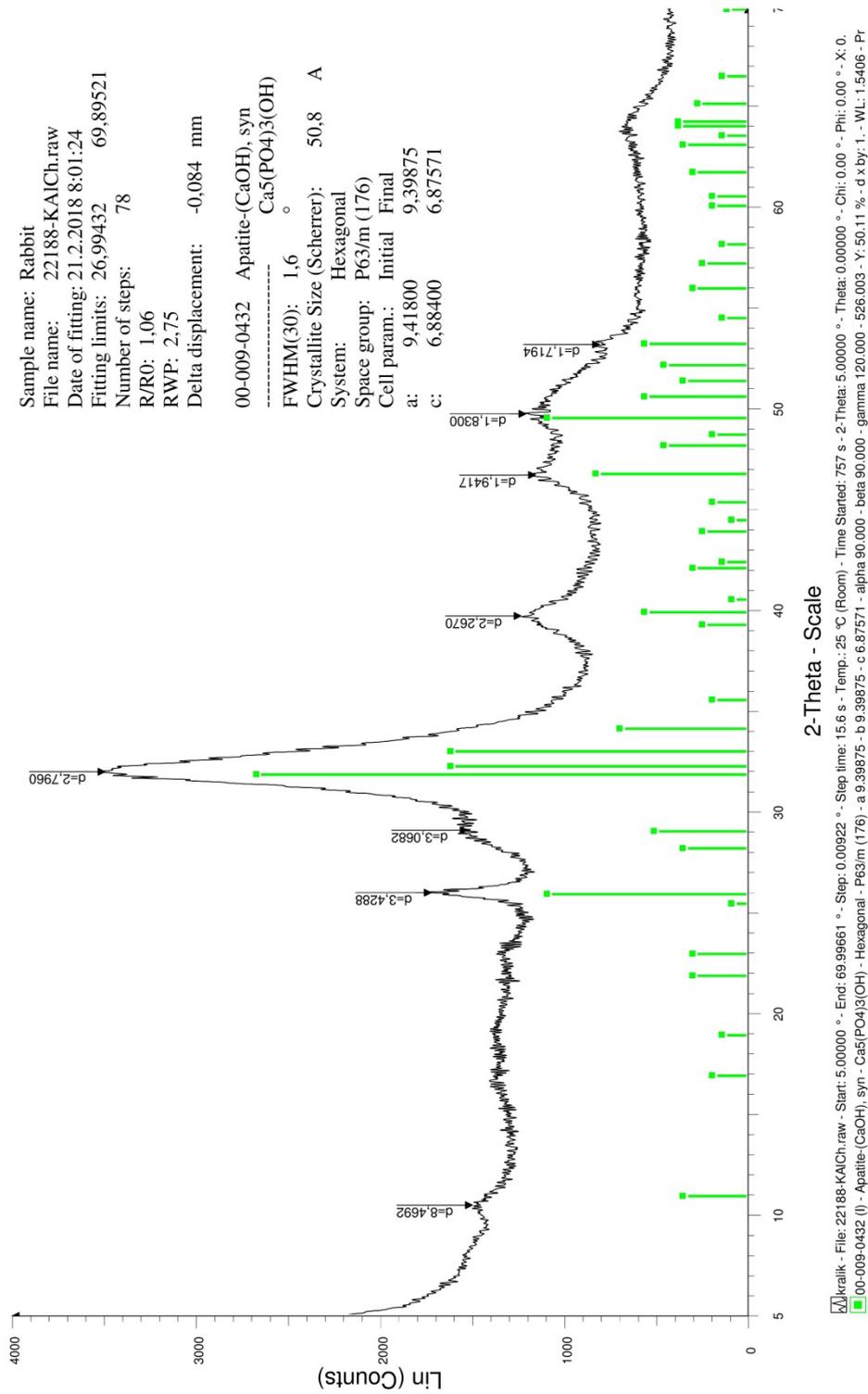
**Figure S3.** The influence of the amount of hydroxyapatite on the integrated absorbance determined at the Pb line 217.001 nm in the presence of 1  $\mu\text{g}$  Pd + 50  $\mu\text{g}$  citric acid for reagent blank samples without (black line) and with the use of automatic correction with the use of reference PO spectra and a least-square algorithm (red line). Pyrolysis and atomization temperature was 1000 and 2100  $^{\circ}\text{C}$ , respectively.



**Figure S4a.** X-ray diffractogram of NIST SRM® 1486

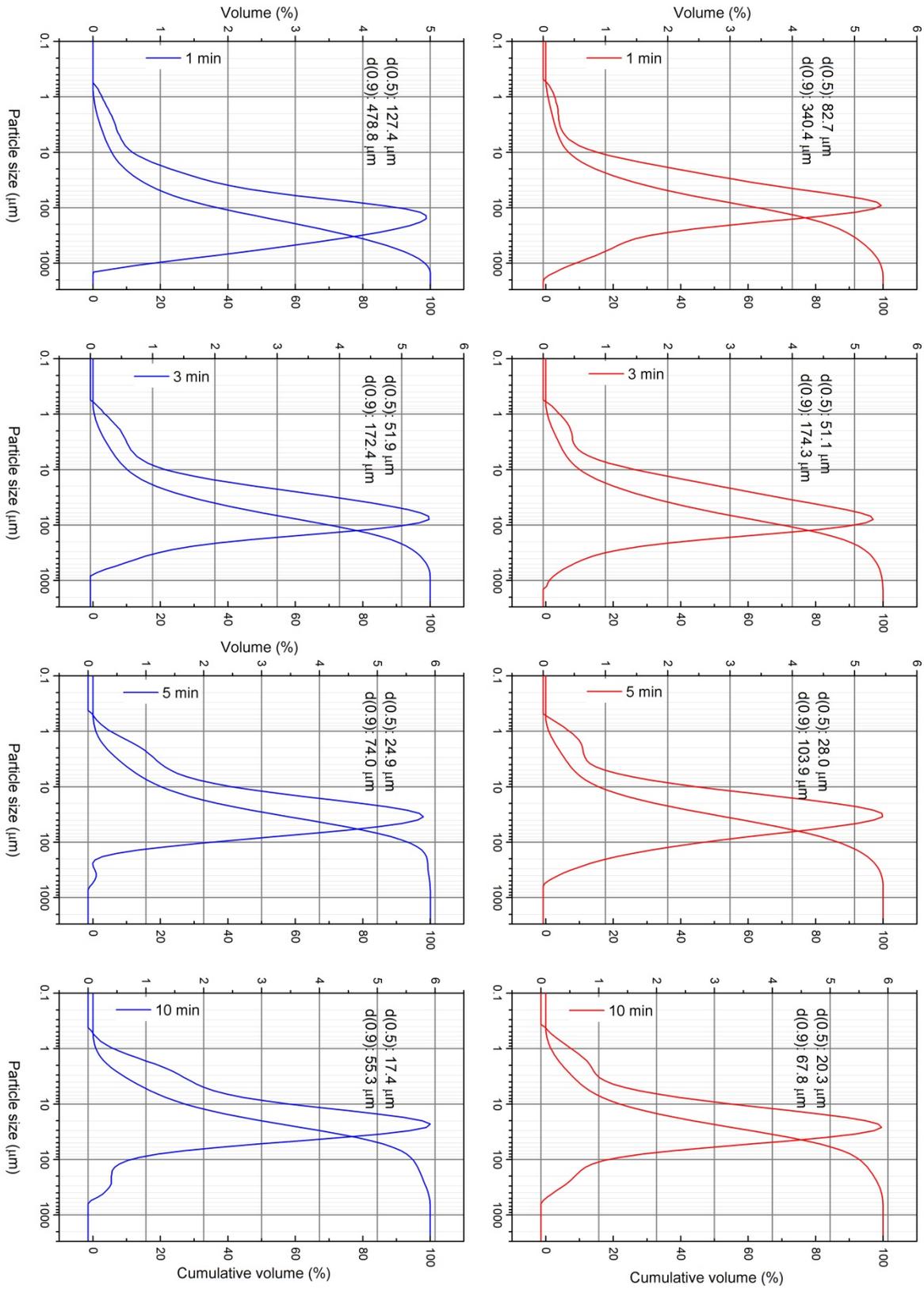


**Figure S4b.** X-ray diffractogram of rabbit rib bone

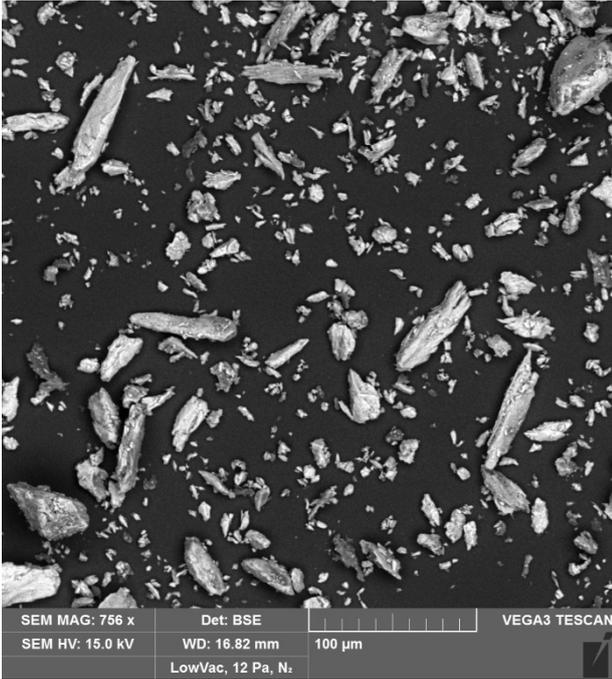




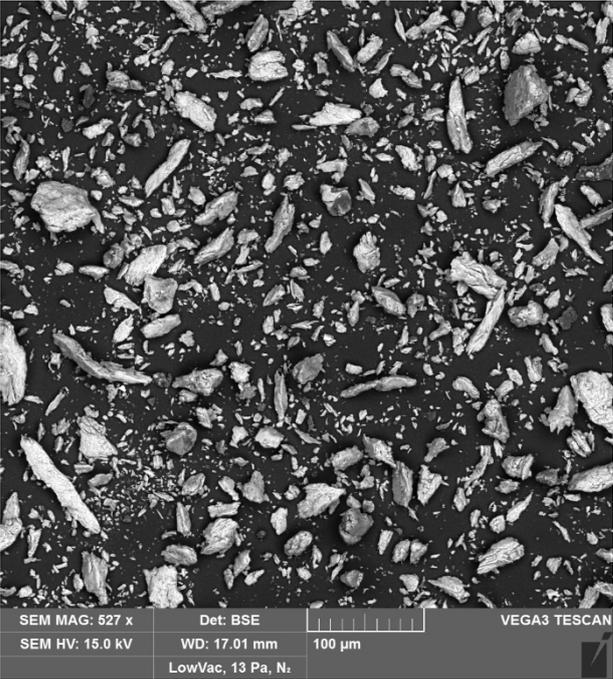
**Fig.S5** Effect of grinding time on the particle size distribution of rabbit (red line) and otter (blue line) bones



**Fig.S6** SEM images of (A) rabbit and (B) river otter bone after 3 min of grinding

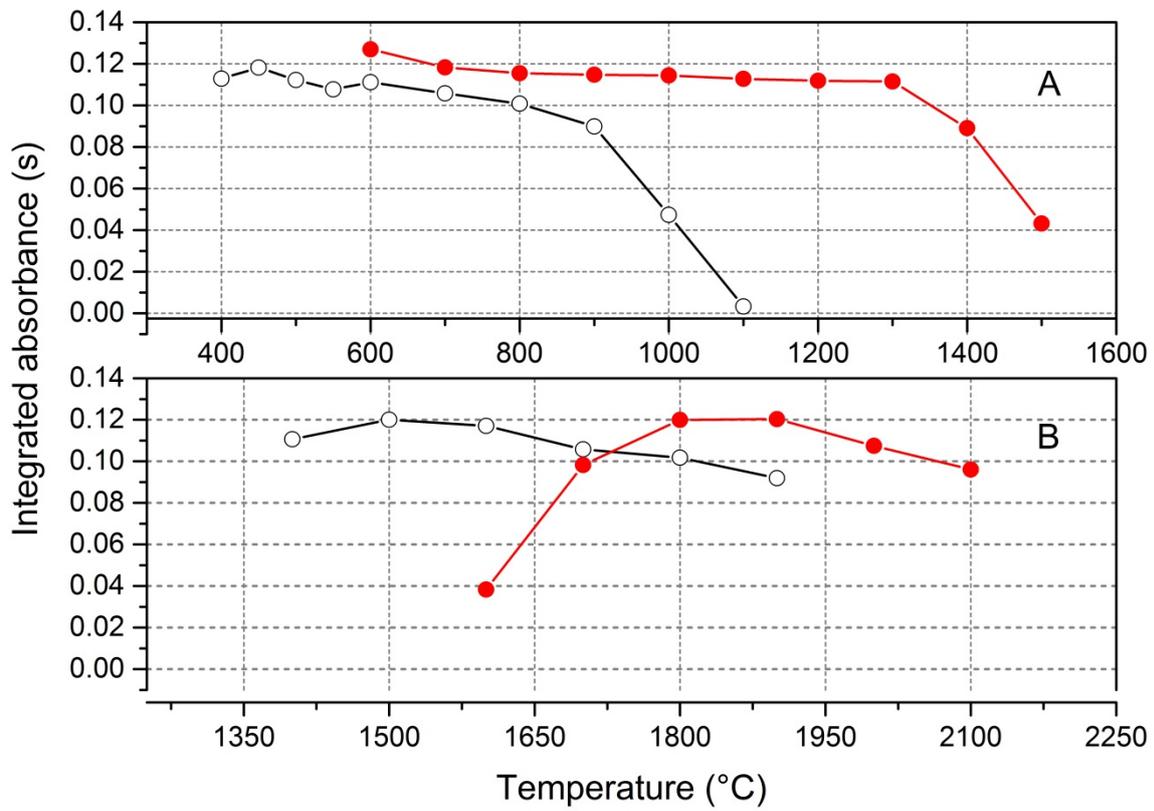


A



B

**Figure S7.** The effect of pyrolysis (A) and atomization (B) temperature on the integrated absorbance of 200 pg Pb in aqueous standard without (○) and with 1 μg Pd + 50 μg citric acid (●) chemical modifier.



**Table S1.** Design and experimental results<sup>a</sup> for analysis of Pb in NIST SRM 1486 Bone Meal

Exp.	Replic.	Glycerol (%)	HNO <sub>3</sub> (%)	Sonication (min)	Particles (μm)	Tp (°C)	Ta (°C)	Modifier (μL)	Integrated absorbance
2	1	20.0	0.0	2.0	54	900	2100	10	0.0409
11 (C)	1	10.0	2.5	6.0	160	1150	1800	6	0.1297
23	3	0.0	0.0	2.0	315	1400	2100	2	0.0040
26	3	20.0	5.0	2.0	315	900	1500	2	0.1476
13	2	20.0	0.0	2.0	54	900	2100	10	0.0367
16	2	0.0	0.0	10.0	315	900	1500	10	0.0158
17	2	20.0	0.0	10.0	54	1400	1500	2	0.0040
22 (C)	2	10.0	2.5	6.0	160	1150	1800	6	0.1340
1	1	0.0	0.0	2.0	315	1400	2100	2	0.0055
12	2	0.0	0.0	2.0	315	1400	2100	2	0.0040
28	3	20.0	0.0	10.0	54	1400	1500	2	0.0030
32 (C)	3	10.0	2.5	6.0	160	1150	1800	6	0.1316
20 (C)	2	10.0	2.5	6.0	160	1150	1800	6	0.1325
8	1	20.0	5.0	10.0	315	1400	2100	10	0.0584
33 (C)	3	10.0	2.5	6.0	160	1150	1800	6	0.1203
14	2	0.0	5.0	2.0	54	1400	1500	10	0.0305
19	2	20.0	5.0	10.0	315	1400	2100	10	0.0466
30	3	20.0	5.0	10.0	315	1400	2100	10	0.0472
27	3	0.0	0.0	10.0	315	900	1500	10	0.0457
4	1	20.0	5.0	2.0	315	900	1500	2	0.1303
21 (C)	2	10.0	2.5	6.0	160	1150	1800	6	0.1065
9 (C)	1	10.0	2.5	6.0	160	1150	1800	6	0.1191
25	3	0.0	5.0	2.0	54	1400	1500	10	0.0225
31 (C)	3	10.0	2.5	6.0	160	1150	1800	6	0.1237
6	1	20.0	0.0	10.0	54	1400	1500	2	0.0017
3	1	0.0	5.0	2.0	54	1400	1500	10	0.0300
7	1	0.0	5.0	10.0	54	900	2100	2	0.1288
15	2	20.0	5.0	2.0	315	900	1500	2	0.1233
29	3	0.0	5.0	10.0	54	900	2100	2	0.1262
24	3	20.0	0.0	2.0	54	900	2100	10	0.0400
18	2	0.0	5.0	10.0	54	900	2100	2	0.1217
5	1	0.0	0.0	10.0	315	900	1500	10	0.0261
10 (C)	1	10.0	2.5	6.0	160	1150	1800	6	0.1271

<sup>a</sup> Results are presented as integrated absorbances evaluated as the means of 3 consecutive analytical runs.

Tp and Ta: pyrolysis and atomization temperature, respectively.

**Table S2.** Determination of Ca, P, Na and Mg in NIST SRM 1486 Bone Meal, rabbit and otter ribs by EDX

Constituent element	Mass fraction (%)		
	SRM	Rabbit	River otter
Ca	25.82 ± 0.66	25.02 ± 0.64	29.08 ± 0.83
P	12.26 ± 0.41	10.17 ± 0.34	9.97 ± 0.38
Na	0.49 ± 0.06	0.54 ± 0.06	0.74 ± 0.07
Mg	0.42 ± 0.05	0.46 ± 0.05	0.27 ± 0.04

Certified<sup>a</sup> or information<sup>b</sup> mass fraction values (Dry-Mass Basis) for SRM 1486: Ca<sup>a</sup>: 26.58 ± 0.24 %, Mg<sup>a</sup>: 0.466 ± 0.017 %, P<sup>a</sup>: 12.30 ± 0.19 %, Na<sup>b</sup>: 0.5 %

**Table S3.** Analysis of variance (ANOVA) for Response Surface Quadratic Model

Source	DF	Adj SS	Adj MS	F-value	p-value <sup>a</sup>
Model	8	0.089235	0.011154	172.63	0.000
Linear	4	0.053645	0.013411	207.56	0.000
x <sub>1</sub> (Glycerol)	1	0.000589	0.000589	9.12	0.006
x <sub>2</sub> (HNO <sub>3</sub> )	1	0.025722	0.025722	398.09	0.000
x <sub>3</sub> (Tp)	1	0.021943	0.021943	339.61	0.000
x <sub>4</sub> (Modifier)	1	0.005391	0.005391	83.43	0.000
Square	1	0.035127	0.035127	543.64	0.000
x <sub>1</sub> (Glycerol) × x <sub>1</sub> (Glycerol)	1	0.035127	0.035127	543.64	0.000
2-Way Interaction <sup>b</sup>	3	0.000463	0.000154	2.39	0.094
x <sub>1</sub> (Glycerol) × x <sub>2</sub> (HNO <sub>3</sub> )	1	0.000196	0.000196	3.03	0.095
x <sub>1</sub> (Glycerol) × x <sub>3</sub> (Tp)	1	0.000004	0.000004	0.06	0.804
x <sub>1</sub> (Glycerol) × x <sub>4</sub> (Modifier)	1	0.000263	0.000263	4.08	0.055
Error	24	0.001551	0.000065		
Total	32	0.090786			

Model Summary: R<sup>2</sup> = 98.3 %; R<sup>2</sup>(adj) = 97.7 %; R<sup>2</sup>(pred) = 96.8 %, where: R<sup>2</sup> is the percentage of variation in the response that is explained by the model, R<sup>2</sup>(adj) is the percentage of the variation in the response that is explained by the model, adjusted for the number of predictors in the model relative to the number of observations, R<sup>2</sup>(pred) provides a measure of how well the regression model predicts the response outside the range of data sample used for the regression.

<sup>a</sup> Parameter is statistically insignificant at significance level 0.05 when its p-value is above 0.05

<sup>b</sup> The following terms cannot be estimated and were removed: x<sub>2</sub> × x<sub>2</sub> (HNO<sub>3</sub> × HNO<sub>3</sub>); x<sub>3</sub> × x<sub>3</sub> (Tp × Tp); x<sub>4</sub> × x<sub>4</sub> (Modifier × Modifier); x<sub>2</sub> × x<sub>3</sub> (HNO<sub>3</sub> × Tp); x<sub>2</sub> × x<sub>4</sub> (HNO<sub>3</sub> × Modifier); x<sub>3</sub> × x<sub>4</sub> (Tp × Modifier).