

Supporting Information

Similarities between *N*-acetylcysteine and Glutathione in Binding to Lead(II) Ions

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Table S1. Assignment of mass ions observed in ESI-MS spectra (positive-ion mode) for Pb(II) *N*-acetylcysteine solutions A, B and E ($C_{\text{Pb(II)}} = 10 \text{ mM}$; mole ratio $\text{H}_2\text{NAC}/\text{Pb(II)} = 2.1, 3.0$ and 10.0, respectively)

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Table S3. Structural parameters obtained from EXAFS least-squares curve-fitting for a Pb(II) GSH solution containing $C_{\text{Pb(II)}} = 10 \text{ mM}$, $C_{\text{GSH}} = 20 \text{ mM}$ (pH = 8.5) measured at RT, using different EXAFS fitting models.

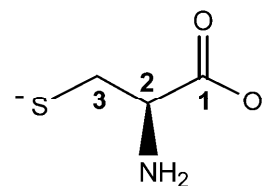
Table S1. Assignment of mass ions observed in ESI-MS spectra (positive-ion mode) for Pb(II) *N*-acetylcysteine solutions A, B and E ($C_{\text{Pb(II)}} = 10 \text{ mM}$; mole ratio $\text{H}_2\text{NAC}/\text{Pb(II)} = 2.1, 3.0$ and 10.0 , respectively) ^a

m/z (amu)	assignment	m/z (amu)	assignment
164.04	$[\text{H}_2\text{NAC} + \text{H}^+]^+$	555.01	$[\text{Na}^+ + \text{Pb}(\text{H}_2\text{NAC})_2 - 2\text{H}^+]^+$
186.01	$[\text{Na}^+ + \text{H}_2\text{NAC}]^+$	556.04	$[3\text{Na}^+ + 3(\text{H}_2\text{NAC}) - 2\text{H}^+]^+$
294.00	$[\text{Pb}(\text{C}_3\text{H}_4\text{NO}_2)]^+$	578.02	$[4\text{Na}^+ + 3(\text{H}_2\text{NAC}) - 3\text{H}^+]^+$
349.05	$[\text{Na}^+ + 2(\text{H}_2\text{NAC})]^+$	738.98	$[\text{Pb}_2(\text{H}_2\text{NAC})_2 - 3\text{H}^+]^+$
370.00	$[\text{Pb}(\text{H}_2\text{NAC}) - \text{H}^+]^+$	740.02	$[2\text{Na}^+ + \text{Pb}(\text{H}_2\text{NAC})_3 - 3\text{H}^+]^+$
371.03	$[2\text{Na}^+ + 2(\text{H}_2\text{NAC}) - \text{H}^+]^+$	741.05	$[4\text{Na}^+ + 4(\text{H}_2\text{NAC}) - 3\text{H}^+]^+$
393.01	$[3\text{Na}^+ + 2(\text{H}_2\text{NAC}) - 2\text{H}^+]^+$	763.04	$[5\text{Na}^+ + 4(\text{H}_2\text{NAC}) - 4\text{H}^+]^+$

^a H_2NAC ($\text{C}_5\text{H}_9\text{NO}_3\text{S}$); $m = 163.03$

Table S2. Comparison of differences in ^{13}C NMR chemical shifts ($\Delta\delta$) for lead(II) *N*-acetylcysteine solutions A – E ($C_{\text{Pb(II)}} = 10 \text{ mM}$) relative to those of a pure *N*-acetylcysteine solution (0.1 M ; $\text{pH} = 9.1$); see Figure 3

Solution	$\Delta\delta$ (^{13}C , ppm)	
	C_1	C_3
A	0.5	3.1
B	0.6	2.4
C	0.5	2.0
D	0.4	1.6
E	0.2	0.8



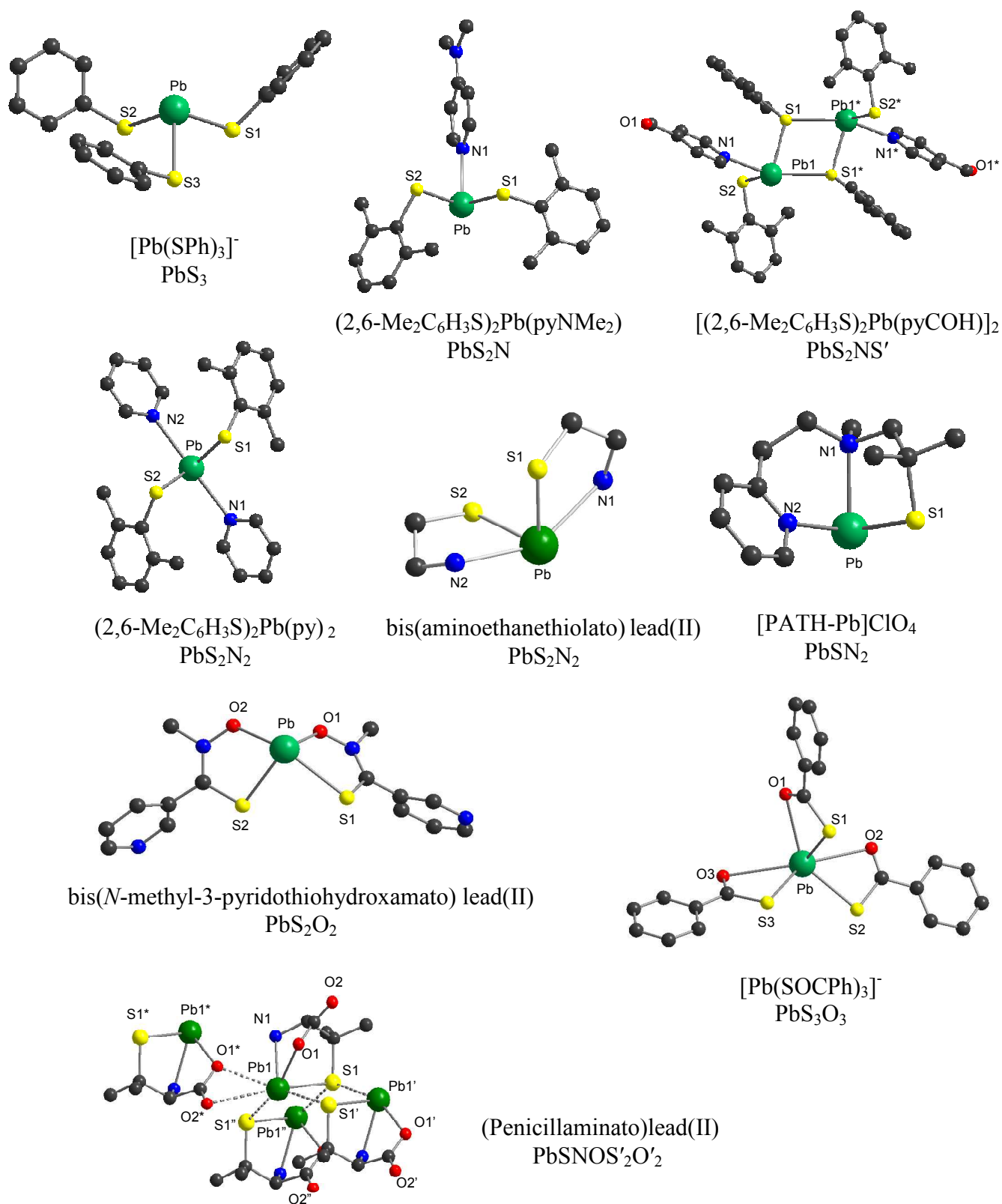


Figure S1. Structures of sulfur-bound Pb(II) complexes for which ²⁰⁷Pb NMR chemical shifts and references have been summarized in Table 2; H-atoms were removed for clarity.

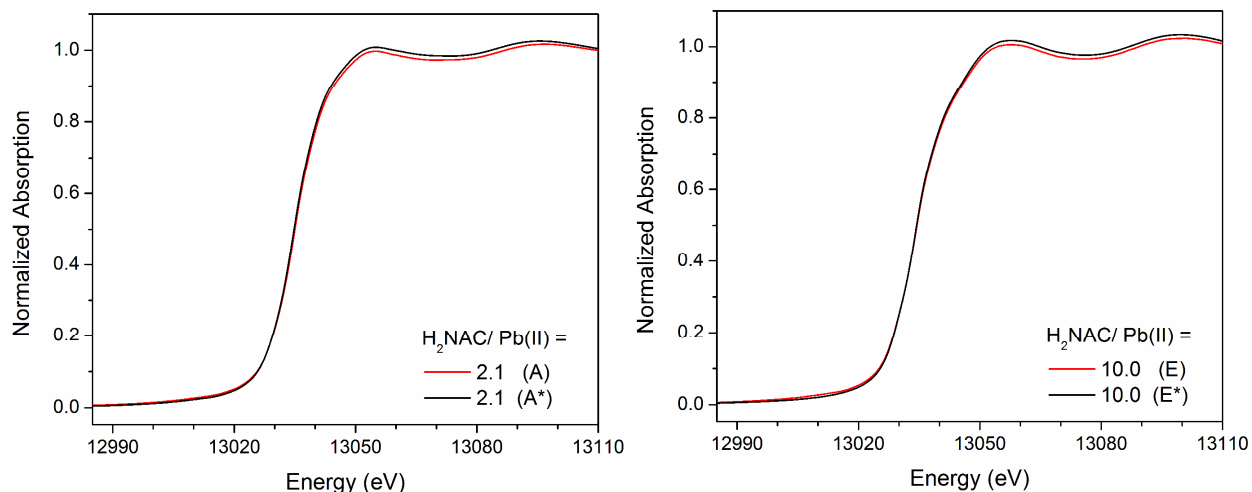


Figure S2. X-ray absorption near-edge features in the XANES spectra of Pb(II) *N*-acetylcysteine solutions, containing $C_{\text{Pb(II)}} = 10$ mM (red), 100 mM (black) and $\text{H}_2\text{NAC}/\text{Pb(II)}$ mole ratios 2.1 (pH = 9.4) and 10.0 (pH = 9.1), showing similar features for each set.

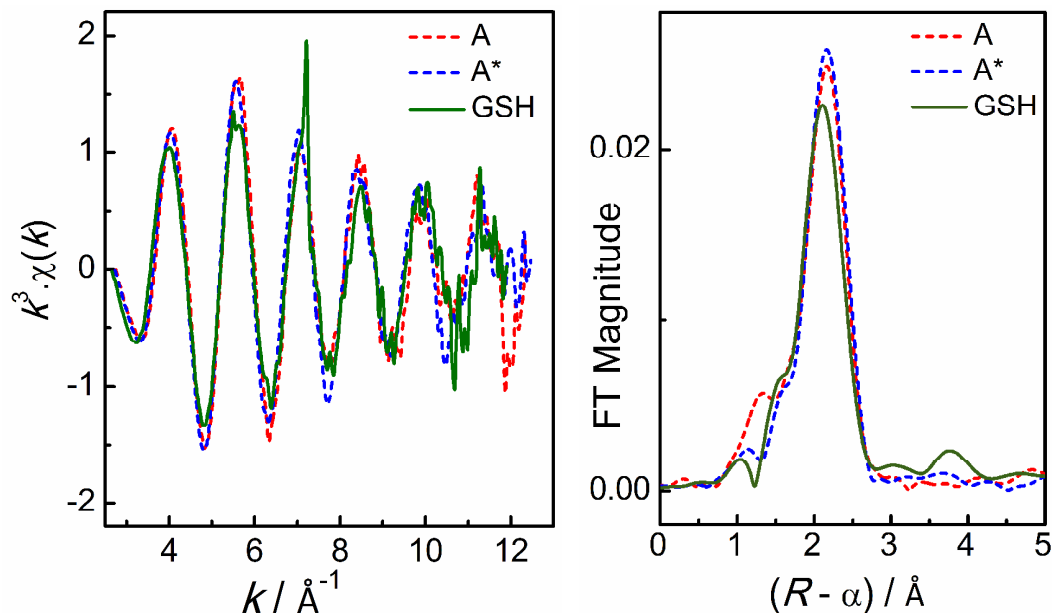


Figure S3. Pb L_{III} -edge k^3 -weighted EXAFS spectrum of a lead(II) glutathione solution ($C_{\text{Pb(II)}} = 10$ mM; $C_{\text{GSH}} = 20$ mM; pH = 8.5) measured in fluorescence mode (solution A in *Ref.* 26) compared with those of Pb(II) *N*-acetyl cysteine solutions A and A* (pH = 9.4), containing $\text{H}_2\text{NAC}/\text{Pb(II)}$ mole ratio 2.1, and their corresponding Fourier-transforms.

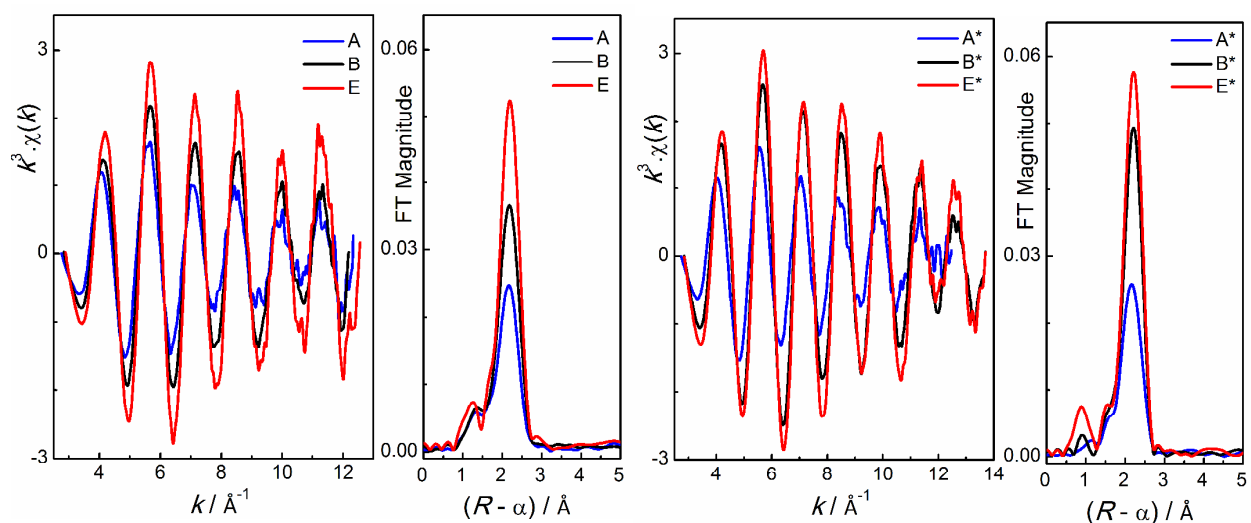


Figure S4. Comparison between Pb L_{III}-edge k^3 -weighted EXAFS spectra and corresponding Fourier-transforms for Pb(II) *N*-acetyl cysteine solutions, containing $C_{\text{Pb(II)}} = 10$ mM (*left*), 100 mM (*right*), and H₂NAC/Pb(II) mole ratios 2.1 (A, A*, pH = 9.4), 3.0 (B, B*, pH = 9.1) and 10.0 (E, E*, pH = 9.1).

Table S3. Structural parameters obtained from EXAFS least-squares curve-fitting for a Pb(II) GSH solution containing $C_{\text{Pb(II)}} = 10$ mM, $C_{\text{GSH}} = 20$ mM (pH = 8.5; solution A in *Ref.* 26) measured at RT, using different EXAFS fitting models.^{a,b}

Model	Pb-S			Pb-N			Pb...Pb			R
	N	R (Å)	σ^2 (Å ²)	N	R (Å)	σ^2 (Å ²)	N	R (Å)	σ^2 (Å ²)	
I	2.7	2.63	0.0107							28.5
II	$2f$	2.66	0.0122	$1f$	2.48	0.0036				29.8
III ^c	$2f$	2.65	0.0156	$2f$	2.51	0.0056				29.4 ^c
IV	$2f$	2.62	0.0080				$1f$	3.84	0.015	27.9
	$1f$	2.76	0.0117							
V	1.9	2.62	0.0076				$1f$	3.84	0.015	27.9
	$1f$	2.75	0.0109							
VI	$1.9f$	2.62	0.0076				$1f$	3.84	0.015	27.9
	0.9	2.75	0.0100							

^a Refined N accurate within ± 20 %; $S_0^2 = 0.9$, f fixed; R = residual. ^b Estimated error limits for R is ± 0.04 Å and for $\sigma^2 \pm 0.002$ Å²; ^c the model shown in Figure 7 and Table 5 of *Ref.* 26