

# Enterobactin Protonation and Iron Release: Structural Characterization of the Salicylate Coordination Shift in Ferric Enterobactin

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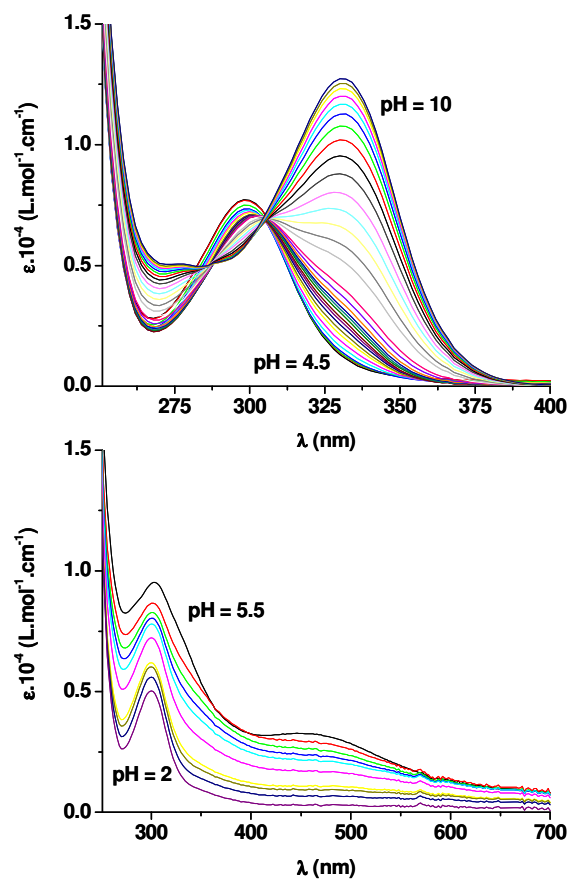
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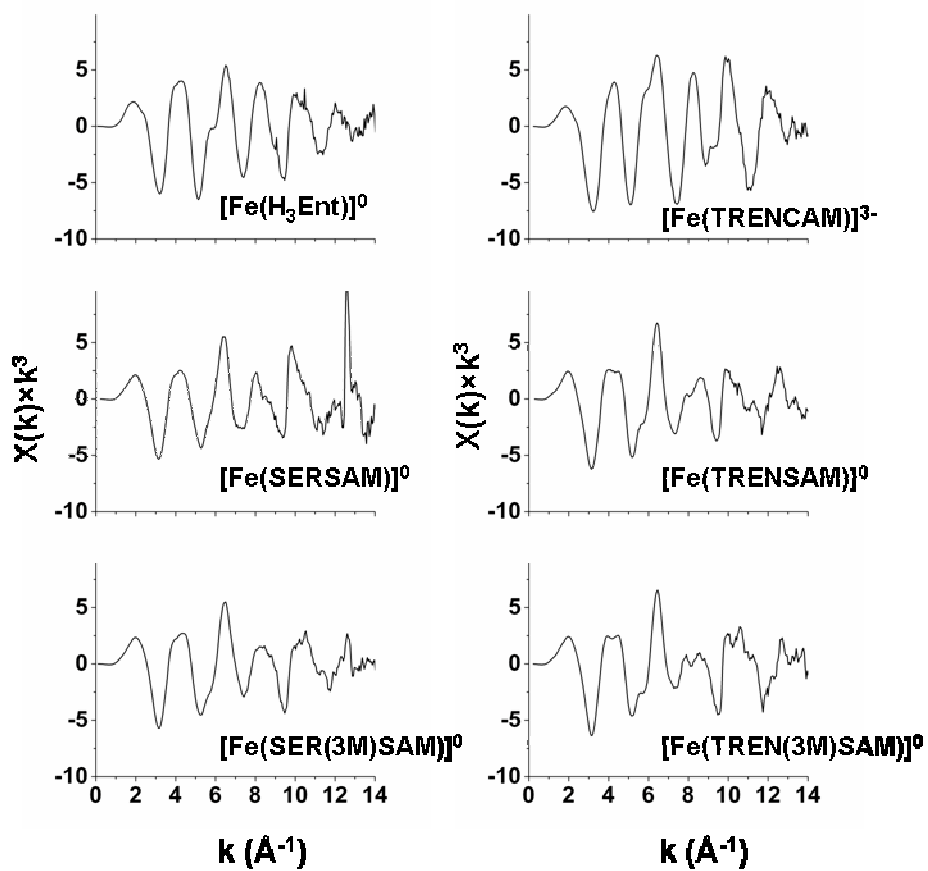
## Supporting Information

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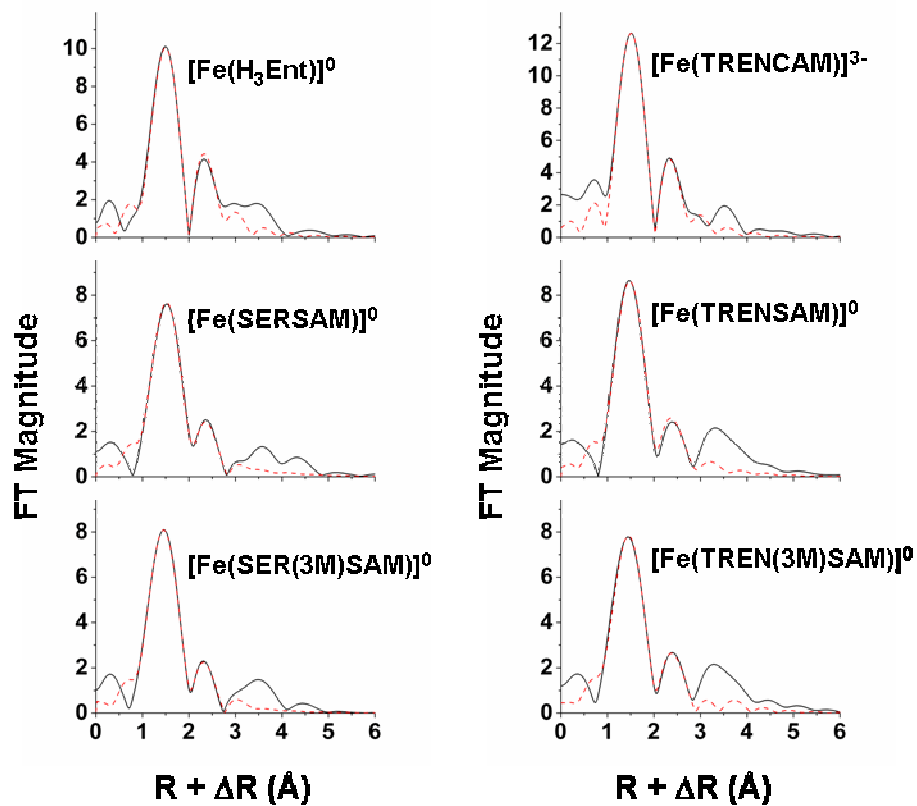
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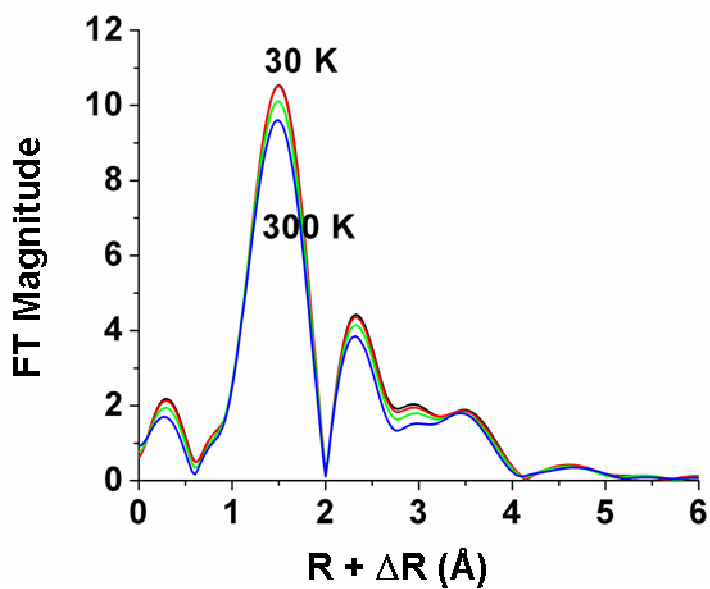
**Figure S1.** Spectrophotometric titrations of SERSAM by KOH (top) and  $[\text{Fe}^{\text{III}}(\text{SERSAM})]^0$  by HCl (bottom) in water.  $I = 0.1$  (KCl),  $T = 25.0$  °C,  $l = 1$  cm.



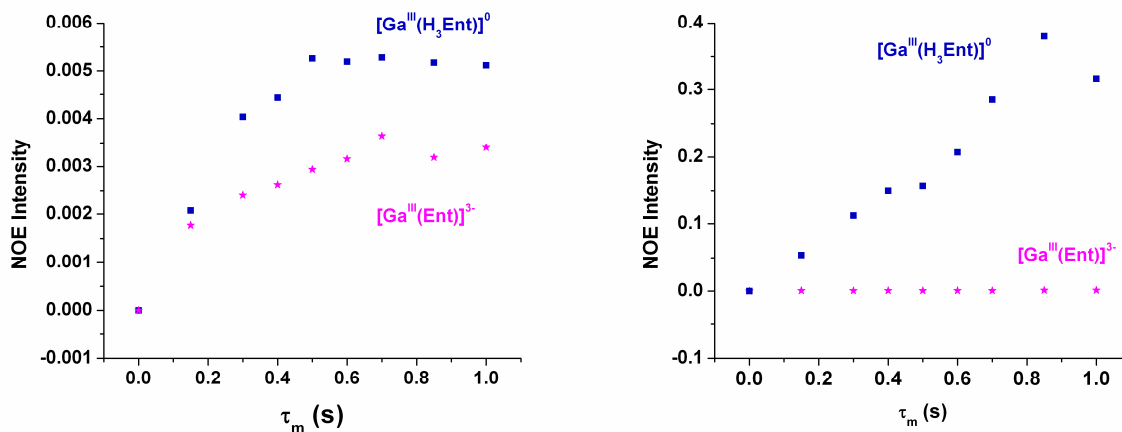
**Figure S2.** k-space spectra of the studied ferric complexes at 200 K.



**Figure S3.** Fourier transforms (solid lines) and R-space fits (dashed lines) of the studied ferric complexes at 200 K.



**Figure S4.** Fourier transforms temperature dependence for  $[\text{Fe}^{\text{III}}(\text{H}_3\text{Ent})]^0$ .



**Figure S5.** NOE intensities build up for the cross peaks between the amide proton and its two closest protons: the seryl  $\beta$ -proton ( $H_b$ , left) and the benzoyl *ortho*-proton ( $H_a$ , right).

Complex	$N_{idp}$	$N_{vp}$	$R$ (30K)	$R$ (100K)	$R$ (200K)	$R$ (300K)
$[Fe^{III}(Ent)]^{3-}$	18	6	6.0 %	6.3 %	6.1 %	6.4 %
$[Fe^{III}(H_3ent)]^0$	18	6	12.1 %	11.9 %	11.5 %	10.7 %
$[Fe^{III}(SERSAM)]^0$	18	6	5.4 %	3.9 %	3.4 %	4.7 %
$[Fe^{III}(SER(3M)SAM)]^0$	18	6	5.1 %	5.3 %	5.4 %	5.8 %
$[Fe^{III}(TRENCAM)]^{3-}$	18	6	10.1 %	10.2 %	10.5 %	10.0 %
$[Fe^{III}(TRENSAM)]^0$	18	4	8.2 %	8.3 %	8.6 %	8.2 %
$[Fe^{III}(TREN(3M)SAM)]^0$	18	6	6.3 %	5.8 %	5.8 %	5.4 %

**Table S1.** Fitting Parameters Used for EXAFS Spectral Analysis.<sup>a</sup>

<sup>a</sup>Number of independent parameters ( $N_{idp}$ ), variable parameters ( $N_{vp}$ ) and  $R$  factors for each studied complex. The  $R$  factors are averaged from at least two independent scans.

Complex	$\Delta H$ (hartrees)	$\Delta H$ (kcal/mol)	Ga-O <sub>ortho</sub> distance (Å)	Ga-O <sub>meta</sub> distance (Å)	Ga-O <sub>carbonyl</sub> distance (Å)	Average Ga-O length (Å)
[Ga <sup>III</sup> (Ent)] <sup>3-</sup>	-2450.775	-3905555	2.034	1.953	6.211	1.993
			2.029	1.953	6.211	
			2.033	1.956	6.200	
cat-[Ga <sup>III</sup> (H <sub>3</sub> Ent)] <sup>0</sup>	-2452.536	-3908361	1.835	2.184	5.947	2.010
			1.835	2.185	5.957	
			1.834	2.186	5.945	
sal-[Ga <sup>III</sup> (H <sub>3</sub> Ent)] <sup>0</sup>	-2452.551	-3908385	1.890	4.444	2.009	1.951
			1.893	4.443	2.013	
			1.893	4.440	2.010	

**Table S2.** Calculated Heats of Formation, Atomic Distances and Bond Lengths for Different Binding Modes of Gallic Enterobactin.

Complex	NH-H <sub>b</sub> distance (Å)	Average NH-H <sub>b</sub> distance (Å)	NH-H <sub>d</sub> distance (Å)	Average NH-H <sub>d</sub> distance (Å)
[Ga <sup>III</sup> (Ent)] <sup>3-</sup>	3.071	3.063	4.456	4.456
	3.052		4.458	
	3.067		4.456	
cat-[Ga <sup>III</sup> (H <sub>3</sub> Ent)] <sup>0</sup>	3.440	3.438	4.448	4.449
	3.438		4.450	
	3.435		4.448	
sal-[Ga <sup>III</sup> (H <sub>3</sub> Ent)] <sup>0</sup>	2.971	2.972	1.956	1.963
	2.972		1.963	
	2.973		1.969	

**Table S3.** Calculated Proton Distances for Different Binding Modes of Gallic Enterobactin.