## **Supporting Information**

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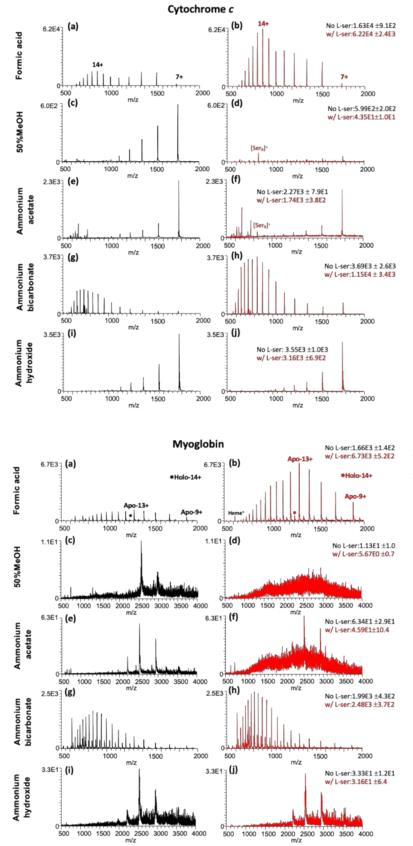


Figure S1. Representative DESI-MS spectra of nativelv deposited and myoglobin cytochrome С analyzed with five different desorption solvent systems with 100 µM L-serine (shown in red) and without it (shown in black) arranged by increasing desorption solvent pH. (a), (b) 0.1% formic acid in 50% MeOH. (c), (d) 50% MeOH. (e), (f) 200 mM ammonium acetate in 50% MeOH. (g), (h) 200 mM ammonium bicarbonate in 50% MeOH. (i), (j) ammonium hydroxide 1% in 50%MeOH. The intensities reported are the average intensity of the highest observed charge state (HICS) of the protein in each solvent system with L-serine (in red) and without it (in black). Statistically significant improvement in signal intensity were observed for cytochrome c (b) and (h) and myoglobin (b) (p-value<0.001).

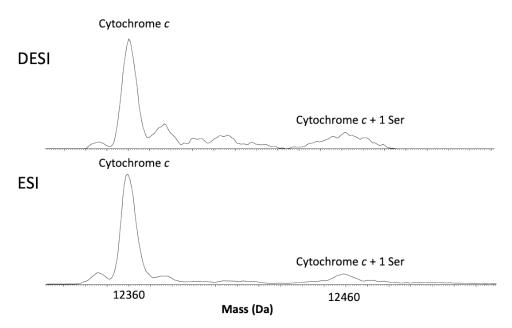
	рН	Average fold change Cyt c HICS Intensity	Average fold change Cyt c Integrated Intensity	Average fold change apo-Myo HICS Intensity	Average fold change Apo-myo Integrated Intensity	Average fold change holo-Myo HICS Intensity	Average fold change Holo-myo Integrated Intensity
Formic acid	2.99	3.8±0.1	3.1±0.4	$4.3{\pm}0.3$	$\textbf{3.7}{\pm}~\textbf{0.2}$	$3.5{\pm}0.3$	3.31±0.3
50% MeOH	5.75	$0.1{\pm}0.02$	0.4±0.02	N/A	N/A	$0.5{\pm}0.05$	N/A
Ammonium acetate	7.32	0.8± 0.2	0.7±0.1	N/A	N/A	0.7±0.2	N/A
Ammonium bicarbonate	7.97	3.1± 0.2	3.1±0.3	1.4± 0.2	1.2± 0.2	1.3± 0.2	$1.3\pm0.2$
Ammonium hydroxide	10.02	0.9± 0.2	0.8±0.2	N/A	N/A	0.9± 0.2	N/A

**Table S1-** Improvement in signal intensity for each protein with addition of L-serine to the five different solvent systems.

Cytochrome c							
Solvent	Intensity Fold Change	≈pH	Protein Net Charge (pl 10.3)	Serine Population	pH-pka₁ (2.21)	pH-pka₂ (9.15)	
Formic acid	3.1	3	+	60-70% zwitterion 30-40% positive	0.79		
50% MeOH	0.8	6	+	100% zwitterion	3.79		
Ammonium acetate	0.8	7	+	99% zwitterion 1% negative		-2.15	
Ammonium bicarbonate	3.1	8	+	92-94% zwitterion 6-8% negative		-1.15	
Ammonium hydroxide	0.9	10	-	10-20% zwitterion 80-90% negative		0.85	

**Table S2.** Approximate net charge on protein and serine in different solvent systems.

Myoglobin							
Solvent	Intensity Fold Change	≈pH	Protein Net Charge (pl 7.4)	Serine Population	pH-pka₁ (2.21)	pH-pka₂ (9.15)	
Formic acid	3.7	3	+	60-70% zwitterion 30-40% positive	0.79		
50% MeOH	0.5	6	+	100% zwitterion	3.79		
Ammonium acetate	0.7	7	-	99% zwitterion 1% negative		-2.15	
Ammonium bicarbonate	1.2	8	-	92-94% zwitterion 6-8% negative		-1.15	
Ammonium hydroxide	0.9	10	-	10-20% zwitterion 80-90% negative		0.85	



**Figure S2.** Representative deconvoluted spectra of cytochrome *c* shows presence of L-serine adducts on cytochrome *c* peaks that were detected at low temperature (70°C) in DESI and ESI with 0.1% formic acid in 50% MeOH with 100  $\mu$ M L-serine as desorption solution and working solution respectively.