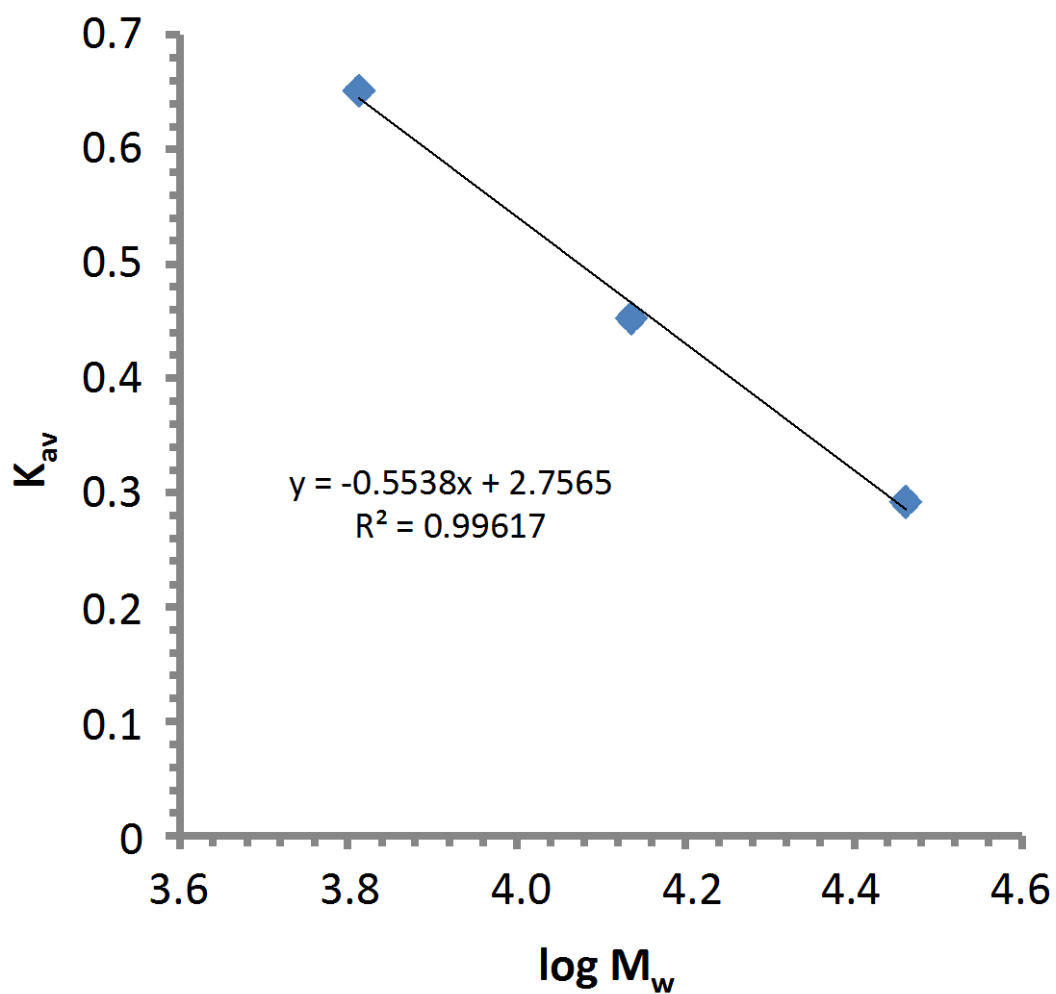*SUPPORTING INFORMATION*

Spectroscopic and Metal Binding Properties of a *De Novo* Metalloprotein Binding a Tetrazinc Cluster

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Figure S1. Calibration curve with standard proteins: carbonic anhydrase, ribonuclease A and aprotinin.

Table S1. Calibration curve parameters.

	K_{av}	$\log M_w$
Aprotinin	0.65	3.8
Ribonuclease A	0.45	4.1
Carbonic Anhydrase	0.29	4.5