Supporting Information for

Metal Affinity Capture Tandem Mass Spectrometry for the Selective Detection of

Phosphopeptides

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Figure S1. Collision-induced dissociation mass spectrum of protonated N_{α} , N_{α} -bis-(carboxymethyl)lysine (LysNTA + H)⁺ at m/z 263. CID spectrum obtained on a Bruker Esquire quadrupole ion trap.



Figure S2. Electrospray mass spectrum (Bruker Esquire ion trap) of a complex formed from 25 μ M LysNTA and 50 μ M GaCl₃ in 1:1 CH₃OH/H₂O buffered with 1% acetic acid. The position of the peak of (LysNTA + H)⁺ is denoted by an arrow. The Ga-containing ions show splitting due to the presence of ⁶⁹Ga and ⁷¹Ga isotopes.



Figure S3. Formation constants for ternary complexes of LysNTA, Ga^{III} and phosphoserine.



Figure S4. CID mass spectra of doubly charged LysNTA complexes with NQLLpTPLR and Ga^{III} (top) and Fe^{III} (bottom).



Figure S5. CID mass spectra of doubly charged LysNTA complexes with pYWQAFR and Ga^{III} (top) and Fe^{III} (bottom).



Figure S6. CID mass spectrum of a triply charged LysNTA- Ga^{III} complex with the α -case in tryptic peptide EKVNELSKDIGpSEpSTEDQAMEDIK.



Figure S7. CID mass spectrum of a triply charged LysNTA- Ga^{III} complex with the α -case in tryptic peptide QMEAEpSIpSpSEEIVPNSVEQK.



Figure S8. Neutral loss scan (49 Da difference) of phosphopeptides from trypsinolysis of 40 nM α-casein.



Figure S9. Parent scan (m/z 263) detection of LysNTA-Ga^{III} phosphopeptide complexes from trypsinolysis of 40 nM α -casein.