

REFERENCE

- Albuquerque, C.P.; Smolka, M.B.; Payne, S.H.; Bafna, V.; Eng, J. & Huilin Zhou (2008) A Multidimensional Chromatography Technology for In-depth Phosphoproteome Analysis. *Molecular and Cellular Proteomics*, Vol.7, No.7, (Jul 2008), pp.1389–1396
- Bantscheff, M.; Schirle, M.; Sweetman, G.; Rick, J. & Kuster B. (2007) Quantitative mass spectrometry in proteomics. a critical review. *Analytical Bioanalytical Chemistry*, Vol.389, No.4 (Oct 2007), pp.1017–1031.
- Chang, C.K., Wu, C.C., Wang, Y.S. & Chang, H.C. (2008) Selective extraction and enrichment of multiphosphorylated peptides using polyarginine-coated diamond nanoparticles. *Analytical Chemistry*, Vol.80, No.10 (Apr 2008), pp.3791–3797
- Domon, B. & Aebersold, R. (2006) Mass spectrometry and protein analysis. *Science*, Vol. 312, No.5771 (14 Apr 2006), pp.212–217

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