Figure S6. A comprehensive database of the yeast interactome (http://tap.med.utoronto.ca/). Querying of a protein name (1) leads to an interaction view summarizing the results of all purifications involving the queried protein, as well as links to YPD, SGD, MIPS and BioGRID databases. Links from this table lead to details of the LC/MS/MS (2) or MALDI-TOF (6 and 7) experiments. From the LC/MS/MS page, mass spectrum search algorithm scores (2 and 3) and spectral images (5) can be retrieved, in addition to results from SIMS site-specific modification searches (4). Similarly, the link to the MALDI-TOF data allows for retrieval of search algorithm scores (7), silver-stained gels (6) and spectral images (8).