



Figure S3. Depletion of *ESF2* has little or no impact on the protein and RNA associated with Utp18p, a protein component of the SSU processome (Dragon *et al.*, 2002). (A) Affinity purifications for two different isolates of a UTP18–TAP tet:*ESF2* strain, using a “low speed spin” protocol (Krogan *et al.*, 2004). (B) UTP “B” complex from Krogan *et al.*, 2004, showing strong similarity to the purifications in (A). (C) Scatter plot comparing RNAs co-purifying with UTP18–TAP protein detected using microarrays (Xing *et al.*, 2004). Ratio data from all 21,939 array spots is shown; any quantitative differences would appear as spots off the line $X = Y$. Enrichment for 5' ETS and U3 sequences is expected.