

Figure S1: Pairwise correlations of bait proteins based on pulled-down protein profiles.

Pearson correlation coefficients were computed for all pairs of bait proteins as well as the control fraction based on the pulled-down protein profiles. Positive correlations indicate that the bait proteins have similar pulled-down protein profiles. ApsA (DVU0847) – ApsB (DVU0846), RpoC (DVU2928) – RpoB (DVU2929) and RpoB (DVU2929) – Pnp (DVU0503) are all highly correlated with each other. The control fraction correlations provide information on baits with potential for false positives and in this study the maximum correlations between the baits and control are small (RoO (DVU3185) being the exception (see Methods)) after removing the prey identified as having false positive potential.