

| Bait Protein | Enriched mRNAs (SDs) | Function of the bait protein |
|--------------|---|--|
| Apc13-TAP | | Anaphase-promoting complex subunit |
| Ark1-myc | | Aurora-B kinase |
| Arp10-TAP | | Dynactin subunit |
| Arp1-TAP | | Dynactin subunit |
| Arp2-TAP | <i>arp8</i> (3.8, 4.1, 2.9) <i>arp9</i> (2.8, 2.9, 2.4) | Member of Arp2/3 complex; involved in actin polymerization |
| Arp3-TAP | <i>arp3</i> (3.3, 2.5) | Member of Arp2/3 complex; involved in actin polymerization |
| Arp42-TAP | <i>snf21</i> (4.4, 4.7) <i>snf22</i> (3.0, 3.6) | SWI/SNF and RSC complexes; chromatin remodelling |
| Alp5-TAP | <i>alp5</i> (4.6, 6.6) | NuA4, Ino80 and Swr1 complexes; chromatin remodelling |
| Arp5-TAP | <i>arp5</i> (5.8, 2.2) | Ino80 complex; chromatin remodelling |
| Arp6-TAP | <i>alp5</i> (4.2, 4.2, 3.8) <i>arp6</i> (4.3, 2.4, 3.8) | SWR1 complex; chromatin remodelling |
| Arp8-TAP | <i>ino80</i> (4.7, 2.3) | Ino80 complex; chromatin remodelling |
| Arp9-TAP | <i>snf21</i> (4.8, 4.9) <i>snf22</i> (2.7, 3.8) | SWI/SNF and RSC complexes; chromatin remodelling |
| Atf1-myc | <i>atf1</i> (5.4, 4.3, 4.5) <i>pcr1</i> (3.9, 2.7, 2.0) | Transcription factor; stress responses |
| Cdc11-TAP | <i>cdc11</i> (4.4) | SIN complex scaffold protein |
| Cdc2-TAP | <i>rum1</i> (5.8±2.1, n=7) <i>cdc18</i> (3.1±0.5, n=6) | Cyclin-dependent protein kinase |
| Cut17-myc | | Survivin protein |
| Cut7-TAP | <i>cut7</i> (5.3) | Spindle pole body motor |
| Ekc1-myc | | Protein phosphatase regulatory subunit |
| Klp3-TAP | | Kinesin-like protein |
| Klp8-TAP | <i>klp8</i> (5.8) | Kinesin-like protein |
| Lid1-TAP | | Anaphase-promoting complex subunit |
| Rpt2-myc | <i>ubp6</i> (5.0, 4.9) <i>rhp23</i> (3.9, 3.7) | 19S proteasome regulatory subunit |
| Rpn12-myc | <i>ecm29</i> (3.8, 3.6) <i>rpn1301</i> (2.2, 2.9) <i>rpn1302</i> (3.0, 2.6) | 19S proteasome regulatory subunit |
| Nup211-TAP | <i>nup211</i> (4.1, 11.2) | Nucleoporin |
| Pcr1-myc | <i>pcr1</i> (5.9, 4.8, 4.1) | Transcription factor; stress responses |
| Rif1-myc | | Telomere length regulator |
| Mnh1-TAP | <i>SPBC19C7.01/mni1</i> (4.4, 5.3) | Mago nashi homolog |
| Sty1-myc | <i>pyp2</i> (5.4±0.8, n=8) <i>cip2</i> (4.9±1.4, n=6) | MAP kinase; stress responses |
| Swo1-TAP | | Hsp90 chaperone |
| Tea2-TAP | <i>tip1</i> (5.2±1.5, n=7) | Kinesin-like protein |
| Tip1-TAP | <i>tip1</i> (7.7) | CLIP-170 family proteins |