S2 Table. Yeast two-hybrid screening of human placenta library

Results summary listing the protein partners identified (gene name), the number of independent clones and the global PBS ^a. Ordering is performed based on global PBS and then on number of clones. Afadin is indicated in red.

| Gene name | Number of independent clones | Global PBS a |
|-----------|------------------------------|--------------|
| RPS8 | 26 | A |
| RBM5 | 14 | A |
| TSHZ1 | 12 | A |
| RPL5 | 9 | A |
| AEBP1 | 7 | A |
| NAB2 | 6 | A |
| THAP3 | 6 | A |
| NAB1 | 4 | A |
| DNMT3A | 5 | В |
| SAFB | 5 | В |
| ZNF653 | 5 | В |
| EXOSC3 | 4 | В |
| RPL7A | 4 | В |
| AF6 | 3 | В |
| FLJ22329 | 3 | В |
| SP3 | 3 | В |
| ZNF653 | 5 | В |
| ZNF6 | 3 | C |
| CDC27 | 2 | C |
| EPS8L2 | 2 | C |
| PHF23 | 2 | C |
| ZNF283 | 2 | C |

^aGlobal Predicted Biological Score (PBS). Confidence score of each interaction. A: very high confidence interaction; B: high confidence in the interaction; and C: good confidence in the interaction.