

Supplementary Table 6 Transcripts upregulated in human cells depleted of Stau1 in three independently performed microarray analyses and that bind Stau1-HA₃ in three (white) or two (gray) independently performed microarray analyses

Transcript	Relative increase (microarray value)		Accession
	Stau1 depletion	Anti-HA IP	
<i>Homo sapiens</i> cDNA FLJ40697 fis clone THYMU2025406	2.5	16.6	AA398756
<i>Homo sapiens</i> transcribed sequence with moderate similarity to protein ref:NP_071431.1 (<i>H.sapiens</i>) cytokine receptor-like factor 2	2.3	9.2	AA824321
Interferon-induced protein with tetratricopeptide repeats 4 (IFIT4)	4.1	7.2	AI075407
Stathmin-like 3 (STMN3)	5.5	6.4	AL353715
Interferon alpha-inducible protein (clone IFI-15K) (G1P2)	3.2	4.1	NM_005101
Cyclin-dependent kinase 5 regulatory subunit 1 (p35) (CDK5R1)	2.1	3.8	AL567411
Decapping enzyme hDcp2 (DCP2)	2.7	2.6	AV715578
KIAA0143 protein (KIAA0143)	2.0	2.4	AW470003
Cyclin-E binding protein 1 (CEB1)	3.6	2.3	NM_016323
Rho GTPase activating protein 19 (ARHGAP19)	2.1	2.0	U79256
Guanine nucleotide binding protein (G protein) gamma 4 (GNG4)	4.8	4.3	NM_004485
Rho GTPase activating protein 19 (ARHGAP19)	2.1	2.1	AV717623
<i>Homo sapiens</i> cDNA FLJ37290 fis clone BRAMY2014469.	2.1	7.4	AW451135
Epiplakin 1 (EPPK1)	2.3	3.9	AL137725
Fibronectin 1 (FN1)	4.7	3.3	X02761
KIAA0931 protein (KIAA0931)	2.1	3.0	AB023148
<i>Homo sapiens</i> similar to KIAA0563-related gene (LOC376854) mRNA	3.4	2.6	BC040700
Likely ortholog of mouse Sds3 (SDS3)	2.1	2.6	AK026749
Thioredoxin reductase 3 (TXNRD3)	2.0	2.4	AI247566
KIAA0143 protein (KIAA0143)	2.3	2.3	AA805651