

Candidate	<i>S. cerevisiae</i> homolog(s)	<i>H. sapiens</i> homolog
<i>ash2</i>	<i>BRE2</i>	ASH2L
** <i>cay1</i>	--	CACTIN
* <i>csx2</i>	--	ACAP1
* <i>ctr1</i>	--	CCDC174
** <i>cwf18</i>	--	CCDC12
** <i>cwf19</i>	--	CWF19L2/L1
<i>est1</i>	<i>EST1</i>	SMG6
<i>hba2</i>	<i>PDR5/10/12/15/18</i>	--
* <i>hri2</i>	--	EIF2AK2/3
<i>ies2</i>	<i>IES2</i>	INO80B
<i>imp1</i>	<i>SRP1</i>	KPNA1/5/6
<i>lem2</i>	<i>LEMD2</i>	SRC1, HEH2
<i>lsg1</i>	<i>CTK3</i>	--
<i>med20</i>	<i>SRB2</i>	MED20
<i>meu17</i>	<i>SGA1</i>	--
<i>mug154</i>	<i>NUR1</i>	--
<i>mug42</i>	--	--
<i>mug78</i>	<i>ATG13</i>	ATG13
<i>mug80</i>	<i>CLG1</i>	--
<i>not3</i>	<i>NOT3/5</i>	CNOT3
<i>nup170</i>	<i>NUP157, NUP170</i>	NUP155
* <i>nup37</i>	--	NUP37
<i>nxt3</i>	<i>BRE5</i>	G3BP1
<i>oma4</i>	<i>PMT4</i>	POMT1
<i>pds5</i>	<i>PDS5</i>	PDS5B
<i>pht1</i>	<i>HTZ1</i>	H2AFZ, H2AFV
<i>png2</i>	<i>YNG1/2, PHO23</i>	ING2/3/4/5
<i>pof4</i>	<i>EKA1</i>	TCEB3, TCEB3B
<i>ppk29</i>	<i>PRK1, ARK1, AKL1</i>	BMP2K, AAK1
** <i>ppn1</i>	--	PPP1R10
<i>prx</i>	--	--
** <i>pwi1</i>	--	SRRM1
* <i>red1</i>	--	ZFC3H1
<i>rhp51</i>	<i>RAD51</i>	RAD51
<i>rnh201</i>	<i>RNH201</i>	RNASEH2A
<i>rpb9</i>	<i>RPB9</i>	POLR21
<i>rpl2102</i>	<i>RPL21A,B</i>	RPL21
<i>rpl4301</i>	<i>RPL43A,B</i>	RPL37A

	Screen Hit	<i>S. cerevisiae</i> homolog(s)	<i>H. sapiens</i> homolog
*	<i>rtf2</i>	--	RTFDC1
*	<i>rtt109</i>	--	RTT109
*	<i>sdu1</i>	--	DES2
	<i>smd3</i>	SMD3	SNRPD3
	SPAC18G6.13	--	--
**	SPAC20H4.06c	--	GPATCH1
	SPAC22H10.11c	IFH1, CRF1	--
	SPBC1348.03	--	--
	SPBC28F2.08c	HRD3	SEL1L, SEL1L2, SEL1L3
*	SPBC713.05	--	WRD83
	SPCC18.13	TRM82	WDR4
	SPCC550.03c	SKI2	SKIV2L
	SPCC736.07c	BUD27	URI1
*	<i>sdj1</i>	--	PARK7
*	<i>spf31</i>	--	DNAJC8
	<i>ssu72</i>	SSU72	SSU72
	<i>swc2</i>	VPS72	VPS72
	<i>swi3</i>	Csm3	TIPIN
	<i>toc1</i>	--	--
	<i>trs130</i>	TRS130	TRAPPC10
	<i>ubr1</i>	UBR1, UBR2	UBR1, UBR2
	<i>vps71</i>	VPS71	ZNHIT1
	<i>yaf9</i>	YAF9	MLLT3, YEATS4

Table S4: Many candidates identified in our screen have no apparent *S. cerevisiae* homolog but have an apparent homolog in humans.

Listed here are the 61 candidates identified in our screen with their corresponding homolog(s) in *S. cerevisiae* and in humans. The red asterisks indicate the 17 genes that have no homolog in *S. cerevisiae* but do in humans. Of those 17 genes, the green asterisks indicate those genes that are known splicing factors. The blue asterisks indicate those genes that were predicted to be involved in splicing based on homology and have now been confirmed in this study.