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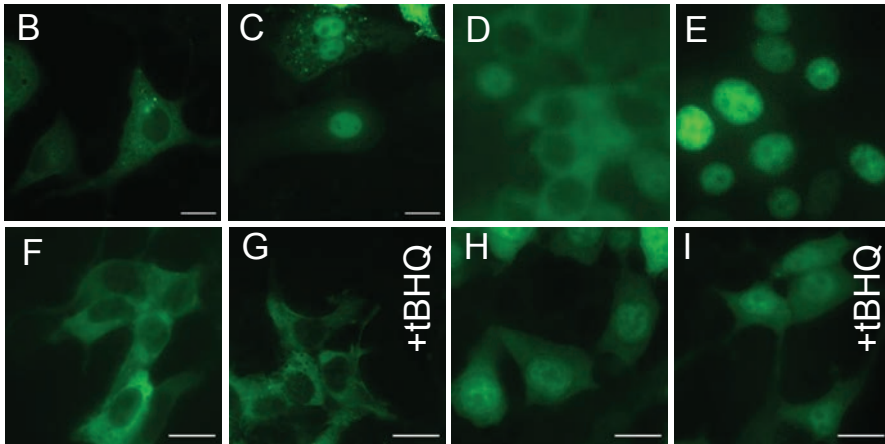
BLAST e-value Species Hit Description % Length

Invertebrates

3.4e-224	<i>C. brenneri</i>	CN:CN26609	CBN17984	90.4%
1.9e-217	<i>C. briggsae</i>	BP:CBP37043	CBR-WDR-23	93.3%
4.1e-86	<i>P. pacificus</i>	PP:PP43443	PPA-WDR-23	65.5%
7.8e-86	<i>B. malayi</i>	BM:BM31075	Bm3045, isoform e	73.0%
1.7e-82	<i>O. volvulus</i>	OV:OVP07517	OVOC1095	71.6%
1.4e-79	<i>C. japonica</i>	JA:JA42848	CJA27743	46.1%
9.1e-75	<i>C. remanei</i>	RP:RP41093	CRE-WDR-23	37.0%
1.1e-13	<i>D. melanogaster</i>	FLYBASE:CG9945	Flybase gene name is CG9945-PA	78.6%
4e-09	<i>S. cerevisiae</i>	SGD:YCR072C	WD-repeat protein involved in ribosome biogenesis; may interact with ribosomes; required for maturation and efficient intra-nuclear transport or pre-60S ribosomal subunits, localizes to the nucleolus	26.4%

Vertebrates

9.8e-83	<i>H. sapiens</i>	ENSEMBL:ENSP00000380146	Isoform 2 of DDB1- and CUL4-associated factor 11	65.1%
4.9e-82	<i>R. norvegicus</i>	SW:Q5M9G8	DDB1- and CUL4-associated factor 11	65.1%
4.9e-82	<i>B. taurus</i>	SW:Q5E9I8	DDB1- and CUL4-associated factor 11	64.9%
1e-81	<i>M. musculus</i>	SW:Q91VU6	DDB1- and CUL4-associated factor 11	71.8%



Lo and Curran  
Figure S1