

A**Overlap of Puf3 Target Sets (from IP data)**

	<i>N. crassa</i> (this study)*	<i>S. cerevisiae</i> (this study)	<i>S. cerevisiae</i> (Gerber <i>et al.</i>)
<i>N. crassa</i> (this study)*	(250)	83	48
<i>S. cerevisiae</i> (this study)	83	(392)	194
<i>S. cerevisiae</i> (Gerber <i>et al.</i>)	48	194	(220)

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	# RNAs	# with Motif Match	
		Non-3' UTR	3' UTR
<i>N. crassa</i> Puf3 Targets (this study)*	250	47 (19%)	83 (33%)
<i>S. cerevisiae</i> Puf3 Targets (this study)	392	99 (25%)	241 (61%)
<i>S. cerevisiae</i> Puf3 Targets (Gerber <i>et al.</i>)	220	50 (23%)	159 (73%)
All RNAs	6228	518 (8%)	710 (11%)

* - "targets" are RNAs bound by *N. crassa* Puf3 when ectopically expressed in *S. cerevisiae*