

Figure 1 – figure supplement 1

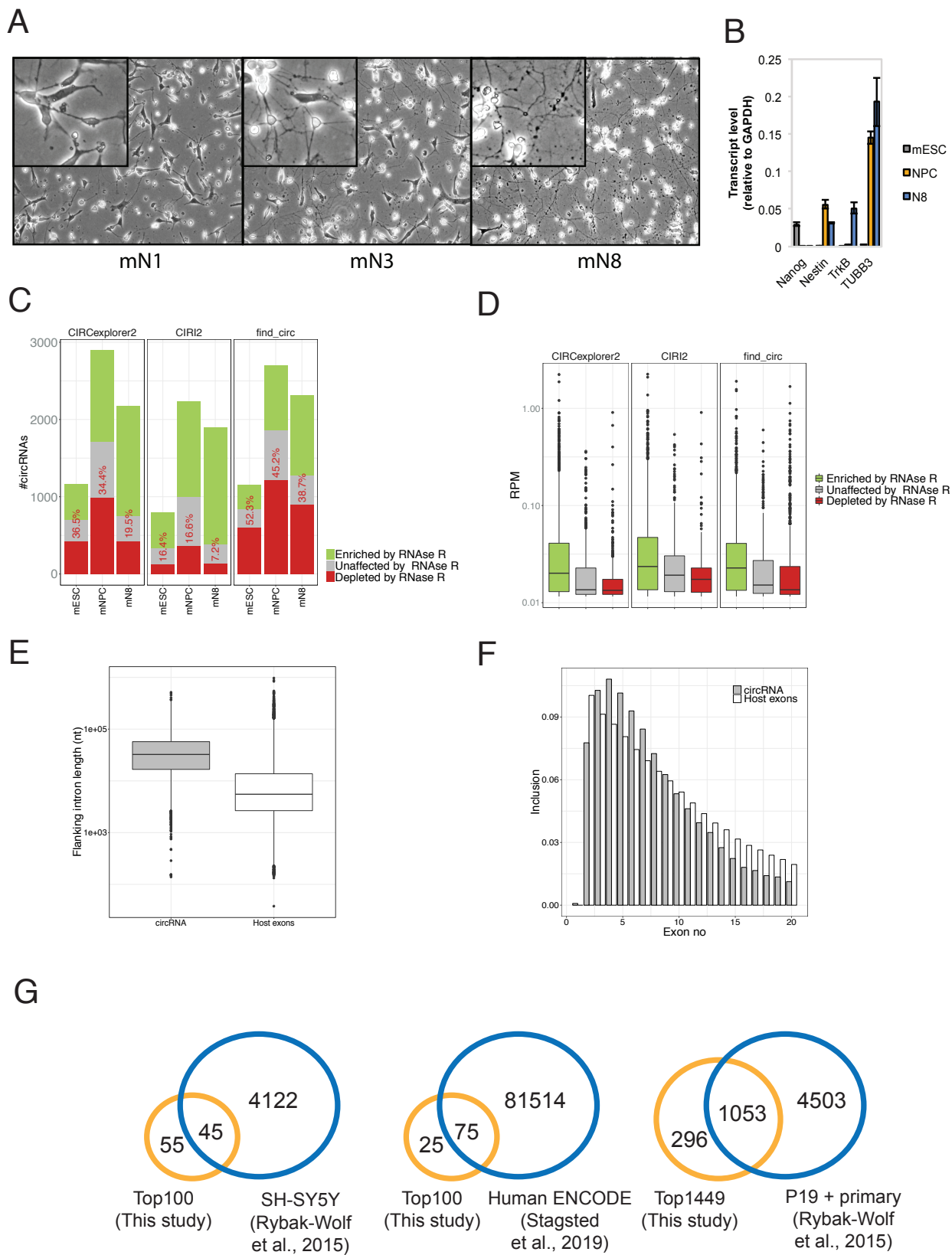


Figure 2 – figure supplement 1

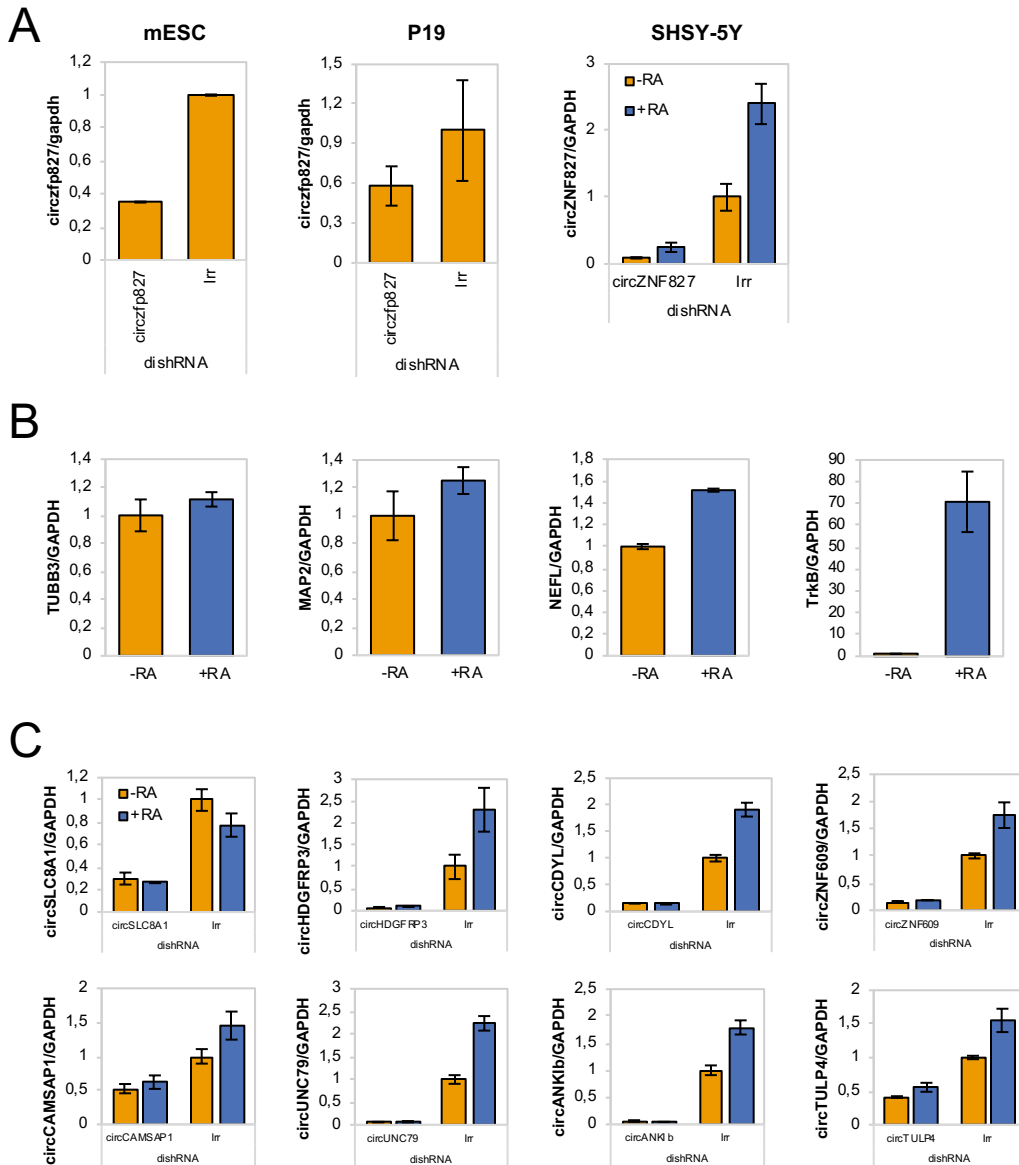
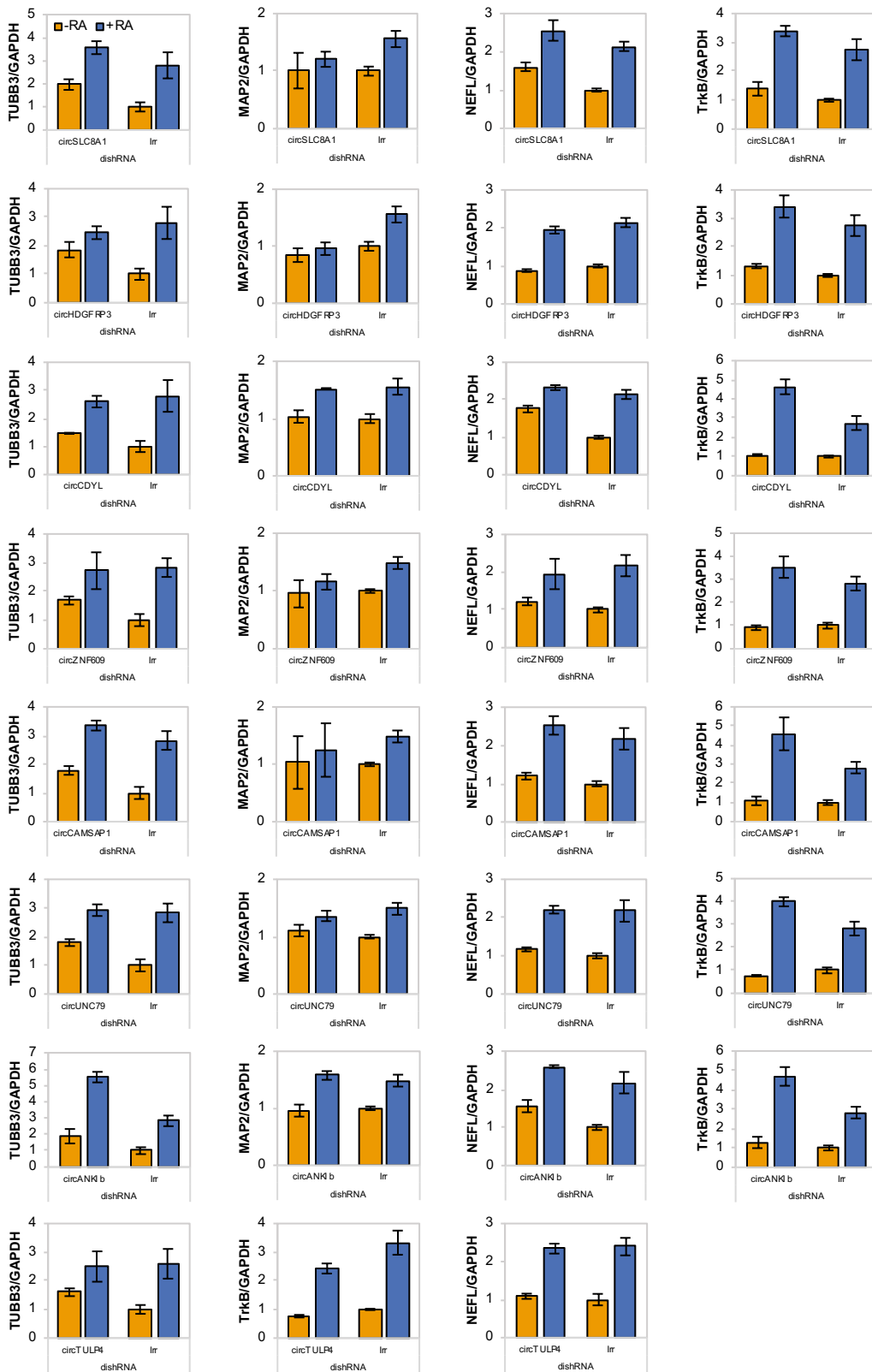


Figure 2 – figure supplement 2

A



B

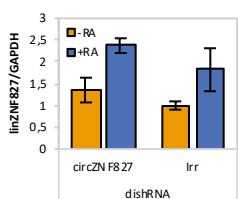
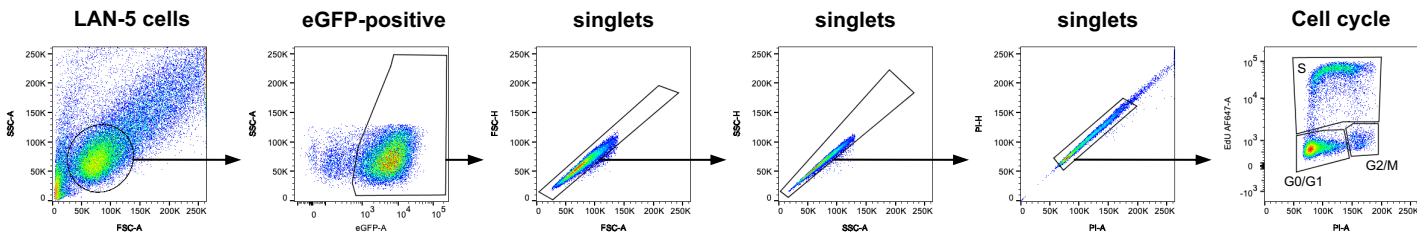


Figure 2 - figure supplement 3

A



B

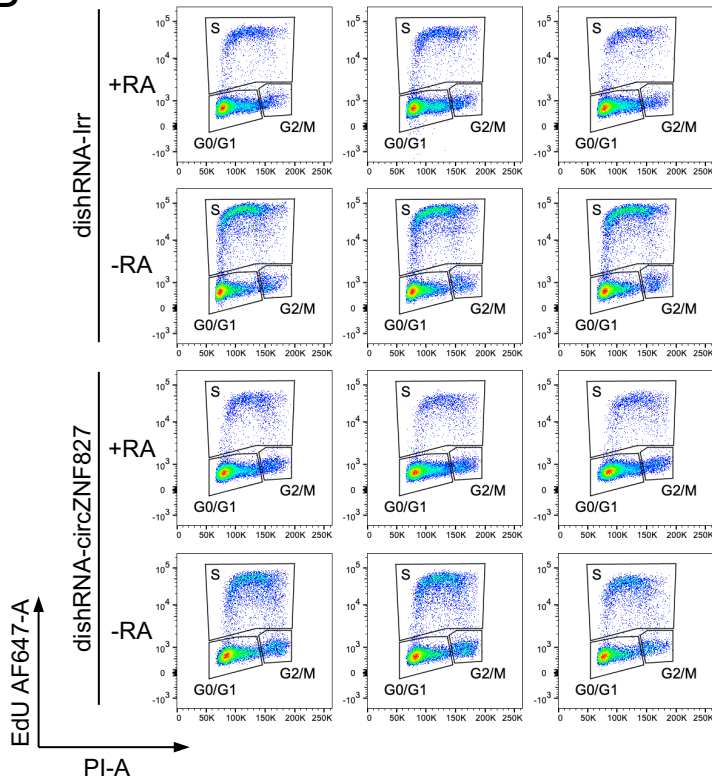


Figure 3 - figure supplement 1

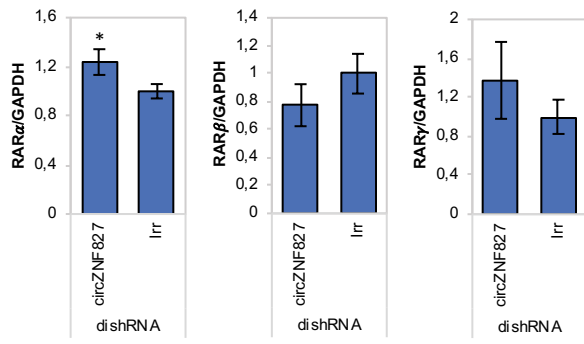


Figure 4 - figure supplement 1

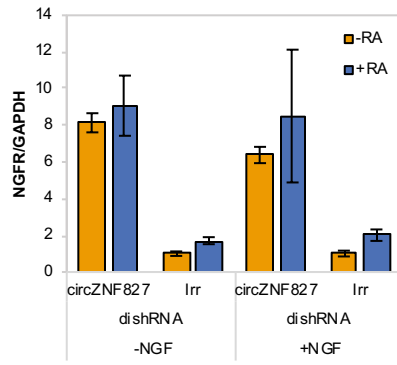
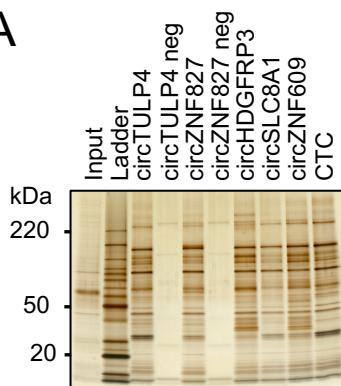


Figure 5 – figure supplement 1

A



C

hsa_circZNF827:
 BSJ_CGGAAAGCTACAAGGAAACACAGATGGTGAAGATTAAAGAGGAACCCATGGAGGTT
 GACATCCAGGACTCCCATGTCTCGATATCACCCAGCCGGAATGTTGGCTACAGCATTTA
 ATCGGGCGAGAGAAAACCGAACCCCTTACAGAAGATGCCAGAGGGCAGAGTACCCCCAGAG
 AGAAACCTCTTCAGTCAGGATATCTCTGTGAAAATGGCTCCGAGCTCCTCTTTCACTG
 TCAGAAAAAGTGAGCAAAGAGCACAATCATACAAAAGAAAACACCATCCGGA**CCACGACC**
AGCCCTTTCTTTTCAGA**AGACACATT**TAGACAATCACCATTCACCTCCAATTCAAAGAA
 CTGCTGCCAGTGACTCCGTGCTGCACGGAGAATATCAGCTCCAG_BSJ

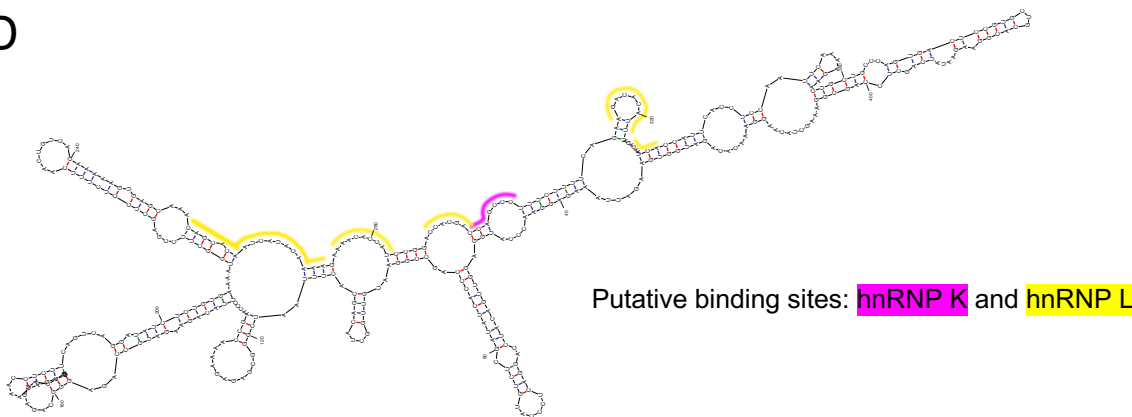
Putative binding sites: **hnRNP K** and **hnRNP L**

B

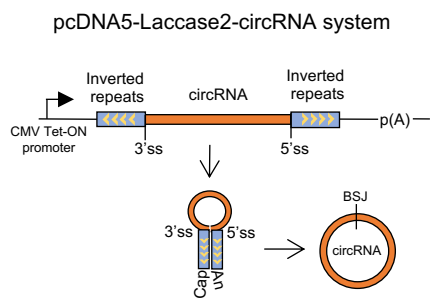
Protein: HNRNPK(Hs/Mm)					
Position	Motif	Occurrence	Z-score	P-value	
295	ccawmcc	aaagaaaaaccacccgaccacga ccagccc uuuuuuucagaagacacauuuga	1.739	4.10e-02	
Protein: HNRNPL(Hs/Mm)					
Position	Motif	Occurrence	Z-score	P-value	
257	acacrav	aacugucagaaaaagugagcaaaag gcacaa ucauacaaaaagaaacacccauccgga	2.887	1.94e-03	
259	amayama	cugucagaaaaagugagcaaaag gcacaa ucauacaaaaagaaacacccauccgga	2.813	2.45e-03	
263	amayama	cagaaaaagugagcaaaagcaca ucauca aaagaaaaacacccauccgga	2.987	1.41e-03	
265	acacrav	gaaaaagugagcaaaagcaca ucauca aaagaaaaacacccauccgga	3.394	3.44e-04	
267	acacrav	aaaagugagcaaaagcaca ucauca aaagaaaaacacccauccgga	3.465	2.65e-04	
275	amayama	gcaaaagcacaaucauacaaaa aaacac auccggaaccagaccgcccuuu	3.093	9.91e-04	
277	acacrav	aaagcacaaucauacaaaa aaacac auccggaaccagaccgcccuuu	3.563	1.83e-04	
289	acacrav	cauacaaaaagaaacacccauccgga ccacga ccgcccuuuucagaagacaca	2.972	1.48e-03	
314	amayama	ccagcaccgcccuuuucaga agacac uuuagacaaucacauaccucca	2.480	6.57e-03	
316	amayama	acgaccgcccuuuucaga agacac uuuagacaaucacauaccucca	2.133	1.65e-02	
326	amayama	cuuuuuucagaagacacauuag acaauc acauuaccuccaaaucaaaagac	2.160	1.54e-02	

RBPmap precisions, <http://rbpmap.technion.ac.il/>

D



E



F

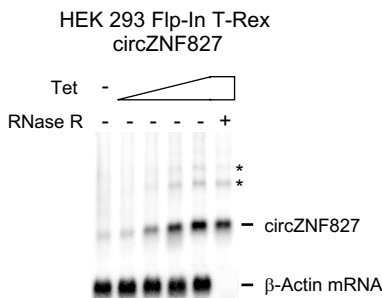
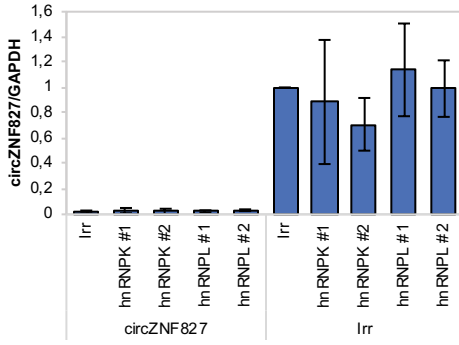
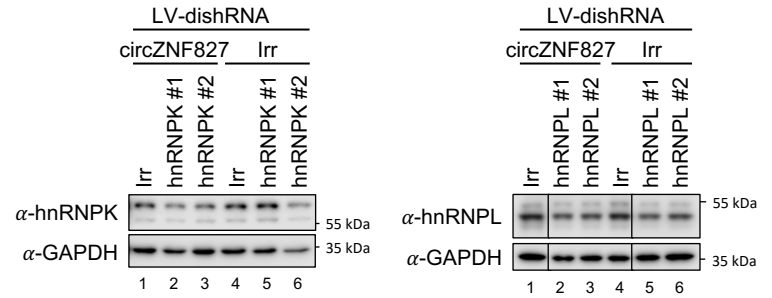


Figure 6 – figure supplement 1

A



B



C

