

Supplementary Table S3. sRSEs in the 3'UTR of GNG7

Index	Sequence	Length	3UTR Start	3UTR End	Stem loop	GC ratio
1	UGUCUCUGUCUCAGGCA	17	59	75	((((.....))))	52.94%
2	GGCCUCUCAGUUUGGGCC	19	238	256	((((.....))))	68.42%
3	GAUGGUUGAACCCGUC	16	583	598	((((.....))))	56.25%
4	GAGGCCGGGAGAUGCCUU	18	1424	1441	((((.....))))	66.67%
5	AACCCGUGCCGAGAGGGUU	19	1744	1762	((((.....))))	63.16%
6	GCGUCCCCACACGC	15	2116	2130	(((.....)))	80.00%
7	GCUGCUGCAGAGACGCAGC	19	2401	2419	((((.....))))	68.42%
8	GAAGGCCGUUUCUCCUUC	18	2967	2984	((((.....))))	55.56%
9	CUGGGCACCUUCCCGG	16	3087	3102	((((.....))))	75.00%
10	UGCCCAAGUUCUGGGGCA	18	3376	3393	((((.....))))	61.11%
11	GGUGGAGGUGGAGCCACC	18	3451	3468	((((.....))))	72.22%