

# mRNA stability measurements

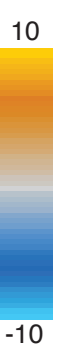
Stable transcripts

Unstable transcripts



Over-representation

Under-representation



name	location	motif	structure	MI	frequency	z-score	robustness	conservation
sRSM9	DN		(((((.....))))))	0.006	0.010	8.278	7/10	1
sRSM10	UP		(((((.....))))))	0.006	0.008	7.994	7/10	0.9969
sRSM7	DN		(((((.....))))))	0.007	0.008	11.800	10/10	1
sRSM11	UP		(((((.....))))).)	0.006	0.004	9.335	10/10	0.759
sRSM8	DN		..(((.....))))))	0.017	0.164	35.564	10/10	1
sRSM12	DN		(((((.....))))))	0.007	0.010	10.206	10/10	1
sRSM13	DN		(((((.....))))))	0.006	0.007	8.768	10/10	0.9996
sRSM14	DN		(((((.....))))))	0.006	0.005	8.380	8/10	0.9967
sRSM6	UP		.(((.....))))))	0.009	0.227	15.789	10/10	1
sRSM15	DN		.(((.....))))))	0.007	0.021	10.957	9/10	1
sRSM16	DN		(((((.....))))).)	0.006	0.005	8.197	10/10	0.9992
sRSM17	DN		(((((.....))))))	0.006	0.003	9.101	9/10	0.9092
sRSM4	DN		(((((.....))))))	0.012	0.131	23.753	10/10	1
sRSM5	UP		(((((.....))))))	0.015	0.156	29.670	10/10	1
sRSM1	DN		(((((.....)))))..)	0.054	0.431	122.220	10/10	1
sRSM19	DN		(((((.....))))))	0.008	0.063	13.767	10/10	1
sRSM2	DN		.(((.....))))).)	0.032	0.350	68.985	10/10	1
sRSM20	DN		(((((.....))))).)	0.005	0.004	7.357	7/10	0
sRSM3	DN		(((((.....))))).)	0.034	0.419	77.138	10/10	1
sRSM21	DN		.(((.....))))))	0.005	0.003	8.781	9/10	1
sRSM22	DN		(((((.....))))))	0.005	0.003	7.626	8/10	1