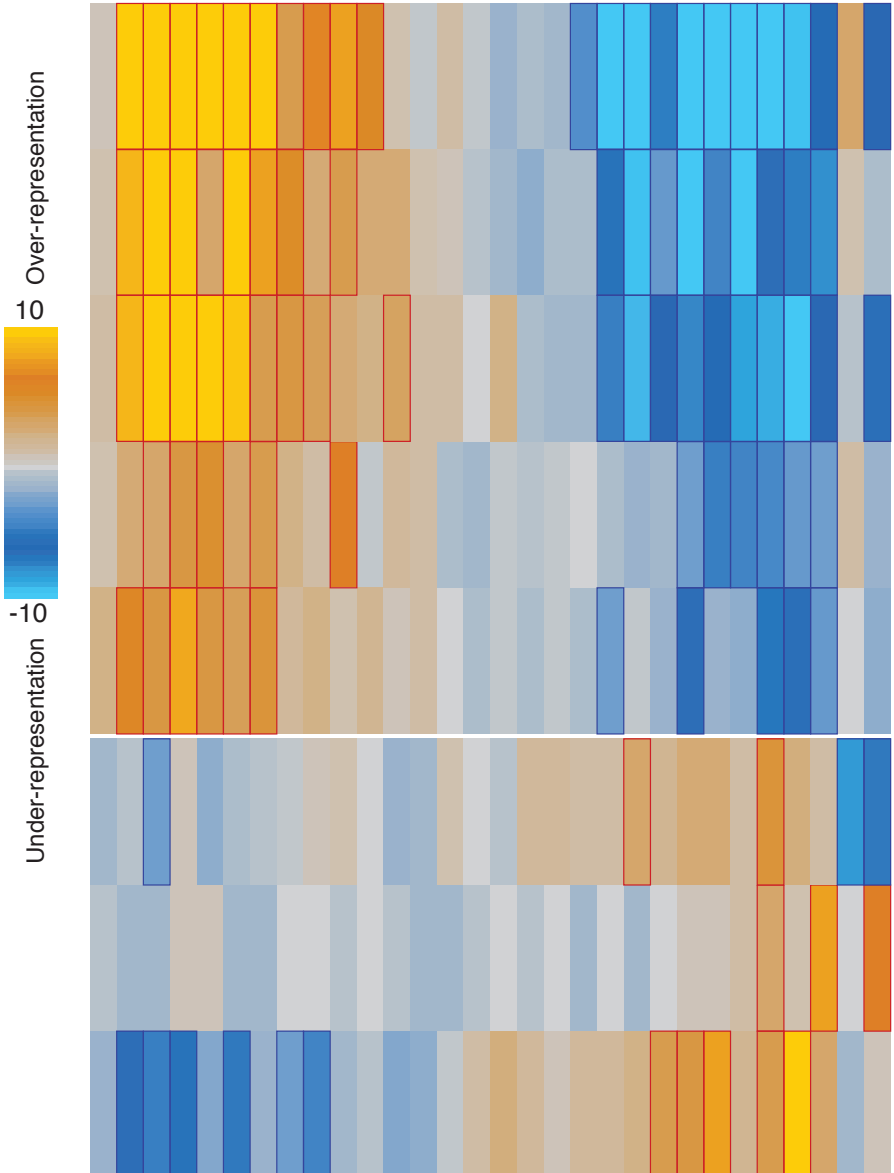
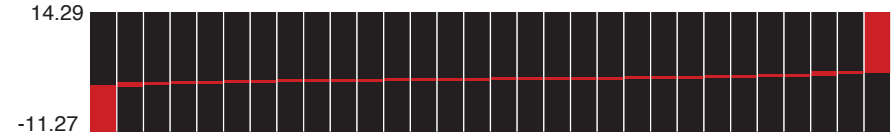


mRNA stability measurements

Stable transcripts

Unstable transcripts



name	location	motif	MI	frequency	z-score
sRSM1	DN		0.054	0.43	122
sRSM2	DN		0.032	0.35	69
sRSM3	DN		0.034	0.42	77
sRSM4	DN		0.012	0.13	24
sRSM5	UP		0.015	0.16	30
sRSM6	UP		0.009	0.23	16
sRSM7	DN		0.007	0.01	12
sRSM8	DN		0.017	0.16	36

