

pri-mir-1-1

chr20: 62554283-62554407 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	1.36 (30 / 476)	2.13 (35 / 476)	<div><div></div><div>Top</div><div></div><div>Bottom</div></div>
Cleavage homogeneity	1.04 (64 / 476)	1.13 (43 / 476)	

