

pri-mir-1283-1

chr19: 53688465-53688589 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	23	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.61 (323 / 476)	1.11 (389 / 476)	<div><div></div>Top</div> <div><div></div>Bottom</div>
Cleavage homogeneity	0.82 (180 / 476)	1.01 (146 / 476)	

