

pri-mir-1185-1

chr14: 101042961-101043085 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	10	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	1.11 (121 / 476)	1.66 (229 / 476)	<div><div></div>Top</div> <div><div></div>Bottom</div>
Cleavage homogeneity	0.93 (128 / 476)	1.08 (76 / 476)	

