

pri-mir-181b-2

chr9: 124693695-124693819 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	4	
Apical loop size (nt)	7	Suboptimal
3' basal segment length (nt)	15	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.15 (444 / 476)
Cleavage homogeneity	0.12 (418 / 476)	0.28 (432 / 476)

