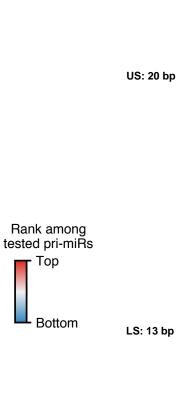
pri-mir-185

chr22: 20033121-20033245 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	19	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.09 (130 / 476)	2.1 (47 / 476)
Cleavage homogeneity	1.06 (57 / 476)	1.02 (129 / 476)



Loop: 13 nt

