


chr8: 91205464-91205588 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.07 (416 / 476)	1.97 (89 / 476)	 Top Bottom
Cleavage homogeneity	0.34 (368 / 476)	0.84 (267 / 476)	

