


chr17: 1713880-1714004 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	14	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	0.73 (292 / 476)	1.33 (347 / 476)	
Cleavage homogeneity	0.69 (248 / 476)	0.86 (251 / 476)	

