

pri-mir-23a

chr19: 13836558-13836682 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	
Apical loop size (nt)	10	Suboptimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.39 (29 / 476)	2.28 (11 / 476)
Cleavage homogeneity	0.83 (179 / 476)	0.93 (213 / 476)

