

pri-mir-136

chr14: 100884684-100884808 (+)

	Features	Optimality
Stem length (bp)	38	Suboptimal
Lower stem length (bp)	16	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.25 (59 / 476)	2.17 (29 / 476)
Cleavage homogeneity	0.49 (313 / 476)	0.31 (428 / 476)

Rank among
tested pri-miRs

 Top

- Bottom

