pri-mir-9-1

chr1: 156420320-156420444 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	8	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.91 (222 / 476)	1.82 (156 / 476)
Cleavage homogeneity	1.01 (89 / 476)	1.08 (73 / 476)

Rank among tested pri-miRs

Top

Bottom