pri-mir-149

chr2: 240455986-240456110 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	4	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	2.21 (3 / 476)	2.15 (32 / 476)
Cleavage homogeneity	1.18 (6 / 476)	1.18 (19 / 476)

Rank among tested pri-miRs

Top

Bottom