pri-mir-143

chr5: 149428912-149429036 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	29	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.97 (186 / 476)	1.78 (175 / 476)
Cleavage homogeneity	0.89 (149 / 476)	1.14 (35 / 476)



