## pri-mir-146a

chr5: 160485341-160485465 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.97 (400 / 476)
Cleavage homogeneity	0.06 (451 / 476)	0.44 (411 / 476)



