

pri-mir-126

chr9: 136670586-136670710 (+)

	Features	Optimality
Stem length (bp)	26	Suboptimal
Lower stem length (bp)	7	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	19	Optimal
3' basal segment length (nt)	31	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.01 (166 / 476)	1.66 (234 / 476)
Cleavage homogeneity	0.67 (253 / 476)	0.84 (268 / 476)

