## pri-mir-128-1

chr2: 135665379-135665503 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	4	Suboptimal
3' basal segment length (nt)	26	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3
	Cleavage efficiency	0.86 (246 / 476)	2.1 (43 / 476)
	Cleavage homogeneity	0.35 (367 / 476)	0.75 (309 / 476)



