


pri-mir-1286

chr22: 20249109-20249233 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	13	
Apical loop size (nt)	6	Suboptimal
3' basal segment length (nt)	0	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.08 (414 / 476)	1.32 (349 / 476)
Cleavage homogeneity	0.03 (464 / 476)	0.64 (351 / 476)

Rank among
tested pri-miRs

Top

Bottom

