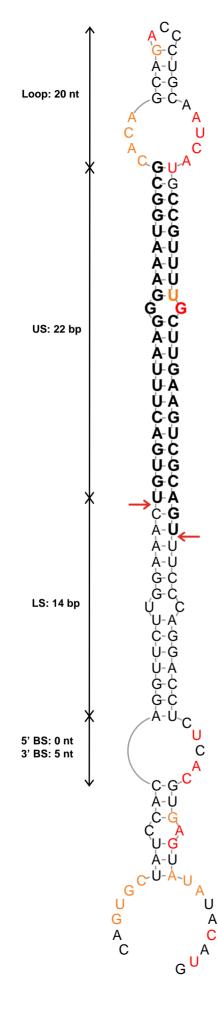
pri-mir-3164

chr11: 69083158-69083282 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	20	Optimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.99 (175 / 476)	1.71 (209 / 476)
Cleavage homogeneity	0.75 (214 / 476)	0.7 (329 / 476)



Rank among tested pri-miRs

Top

Bottom