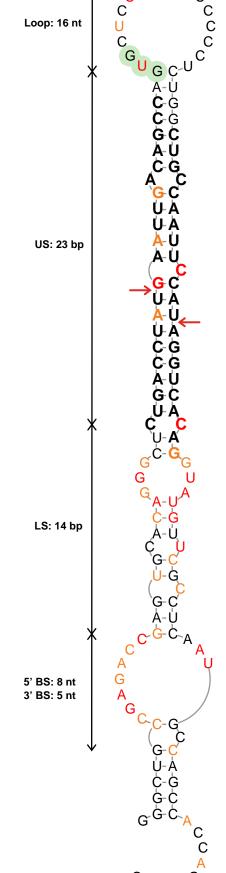
pri-mir-192

chr11: 64891127-64891251 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	2	
Apical loop size (nt)	16	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	0.46 (358 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.23 (391 / 476)	0.09 (453 / 476)



Rank among tested pri-miRs

Top

Bottom