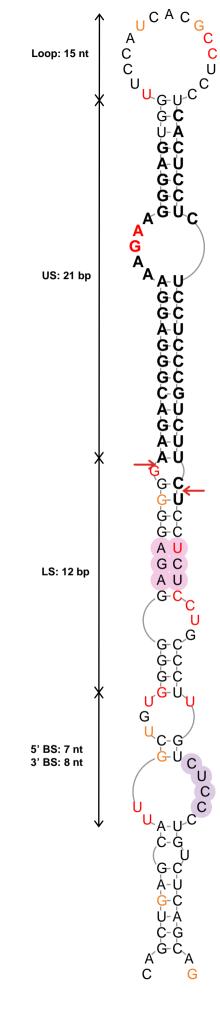
pri-mir-483

chr11: 2134108-2134232 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	8	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	8	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.3 (45 / 476)	1.96 (94 / 476)
Cleavage homogeneity	0.86 (163 / 476)	0.77 (298 / 476)



Rank among tested pri-miRs

Top

Bottom