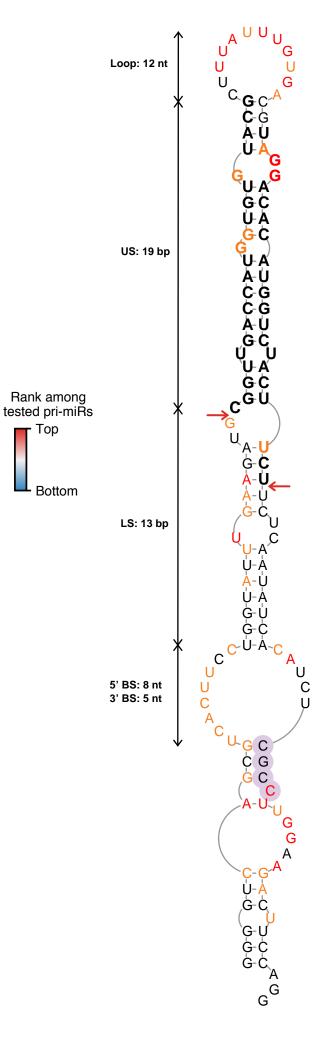
pri-mir-1197

chr14: 101025552-101025676 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	6	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.29 (384 / 476)	0.31 (436 / 476)
Cleavage homogeneity	0.32 (374 / 476)	0.54 (377 / 476)



Top

Bottom