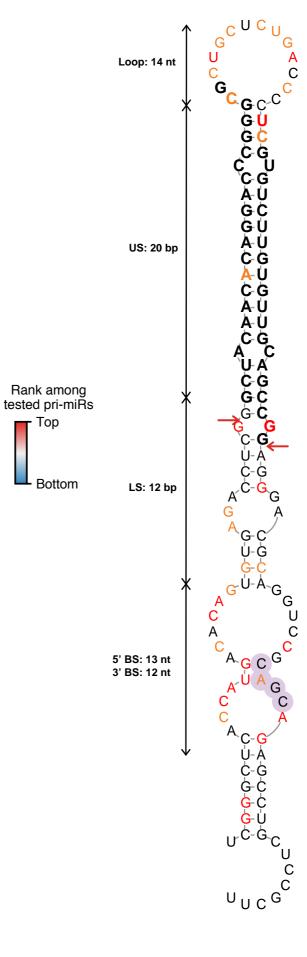
## pri-mir-187

chr18: 35904800-35904924 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	12	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.72 (294 / 476)	1.23 (375 / 476)
Cleavage homogeneity	0.41 (345 / 476)	0.93 (214 / 476)



Top

**Bottom**