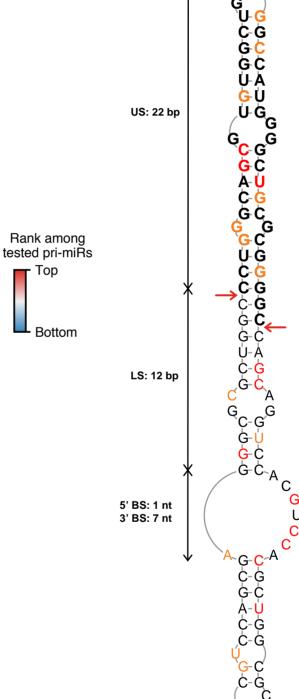
pri-mir-3187

chr19: 813559-813683 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.63 (315 / 476)	0.75 (415 / 476)
Cleavage homogeneity	0.71 (244 / 476)	0.56 (370 / 476)



Loop: 12 nt