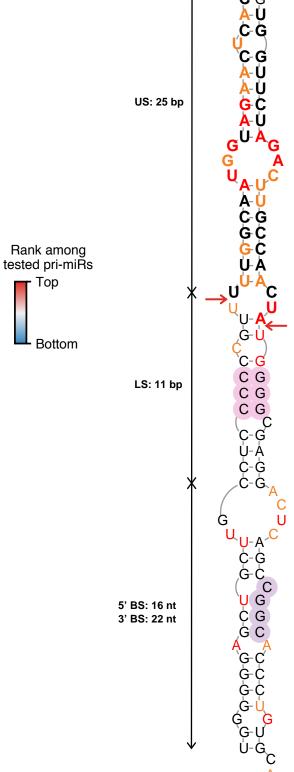
pri-mir-182

chr7: 129770374-129770498 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	4	
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.47 (13 / 476)	2.04 (64 / 476)
Cleavage homogeneity	1.02 (81 / 476)	1.15 (30 / 476)



Loop: 11 nt

U A' G Ų