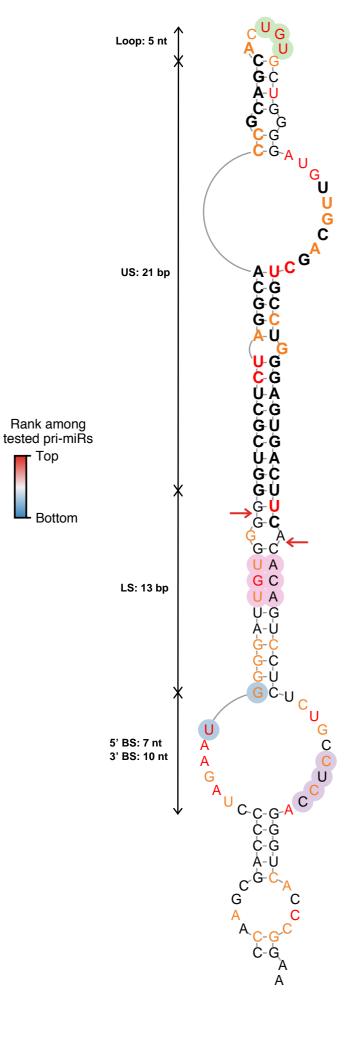
pri-mir-1301

chr2: 25328617-25328741 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	11	
Apical loop size (nt)	5	Suboptimal
3' basal segment length (nt)	10	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.95 (193 / 476)	1.64 (251 / 476)
Cleavage homogeneity	0.41 (347 / 476)	0.47 (400 / 476)



Top

Bottom