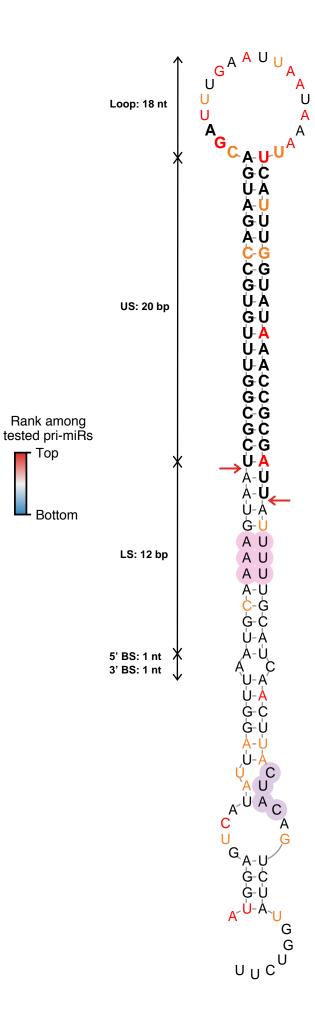
## pri-mir-579

chr5: 32394358-32394482 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	0	Optimal
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	18	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.11 (409 / 476)	1.7 (212 / 476)
Cleavage homogeneity	0.07 (449 / 476)	0.46 (405 / 476)



Top