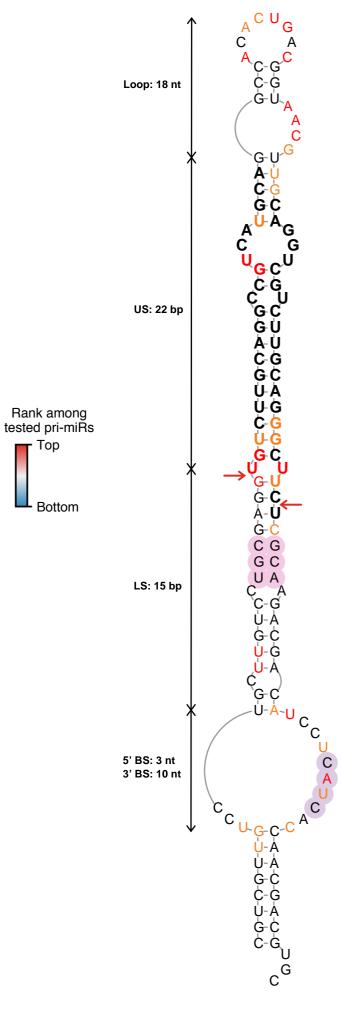
## pri-mir-431

chr14: 100880999-100881123 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	18	Optimal
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	0.93 (209 / 476)	1.35 (343 / 476)
Cleavage homogeneity	0.76 (210 / 476)	1.0 (152 / 476)



Top

**Bottom**