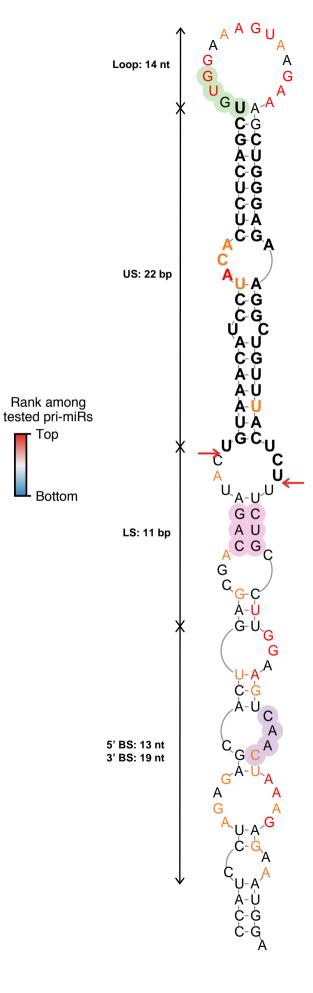
## pri-mir-30c-2

chr6: 71376931-71377055 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	4	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	19	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.7 (298 / 476)	1.02 (397 / 476)
Cleavage homogeneity	0.47 (317 / 476)	0.74 (312 / 476)



- Top

Bottom