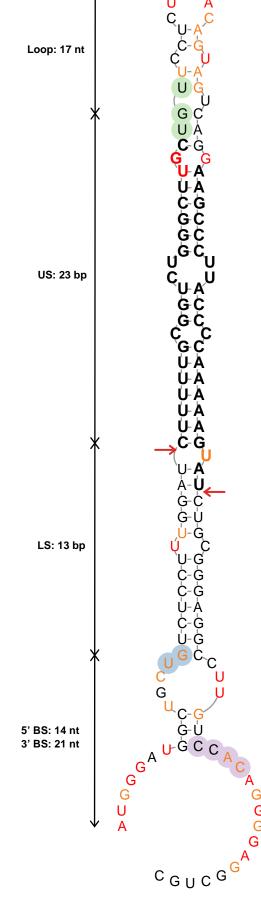
## pri-mir-129-1

chr7: 128207849-128207973 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	21	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.06 (147 / 476)	1.88 (131 / 476)
Cleavage homogeneity	0.68 (251 / 476)	1.02 (128 / 476)



Rank among tested pri-miRs

Top

**Bottom**