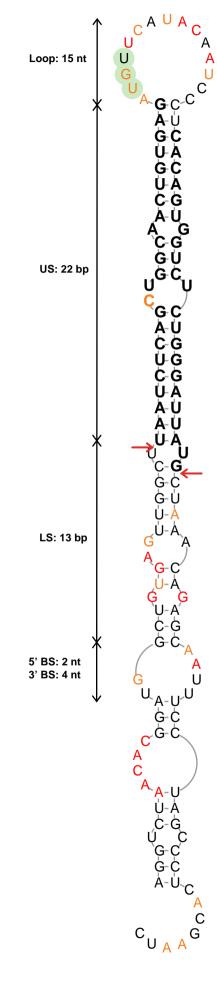
pri-mir-216a

chr2: 55988947-55989071 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.13 (117 / 476)	1.64 (247 / 476)
Cleavage homogeneity	0.93 (121 / 476)	0.9 (228 / 476)



Rank among tested pri-miRs

Top

Bottom