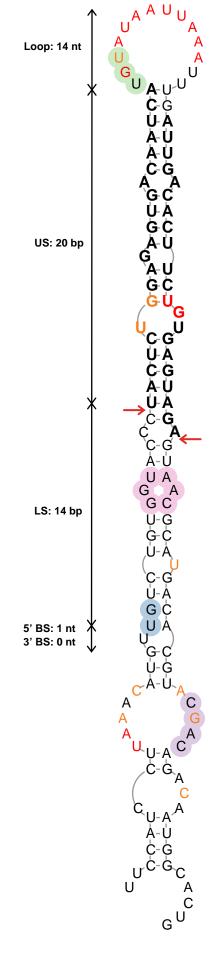
pri-mir-514a-1

chrX: 147279231-147279355 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	3	
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	0	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.3 (47 / 476)	1.81 (158 / 476)
Cleavage homogeneity	1.15 (19 / 476)	1.13 (45 / 476)



Rank among tested pri-miRs

Top

Bottom