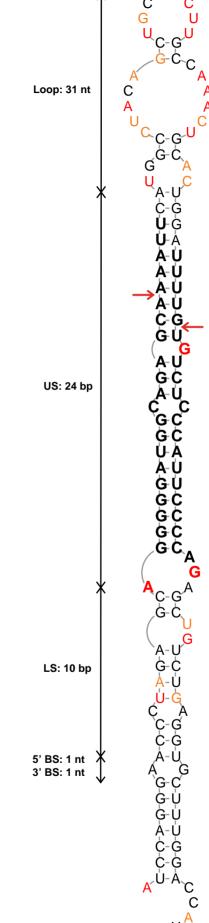
pri-mir-5010

chr17: 42514189-42514313 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	5	
Apical loop size (nt)	31	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.06 (418 / 476)	0.09 (448 / 476)
Cleavage homogeneity	0.17 (408 / 476)	0.11 (452 / 476)



Rank among tested pri-miRs

- Top

Bottom