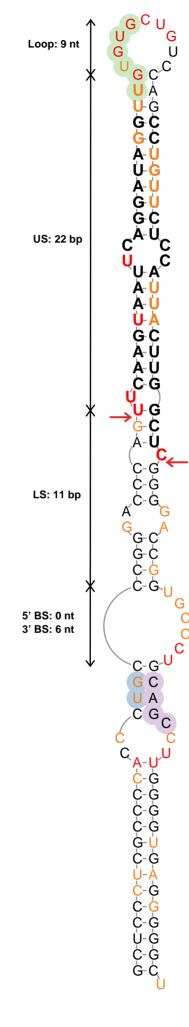
## pri-mir-26b

chr2: 218402626-218402750 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.2 (80 / 476)	1.94 (101 / 476)
Cleavage homogeneity	0.88 (154 / 476)	1.03 (126 / 476)



Rank among tested pri-miRs

Top

**Bottom**