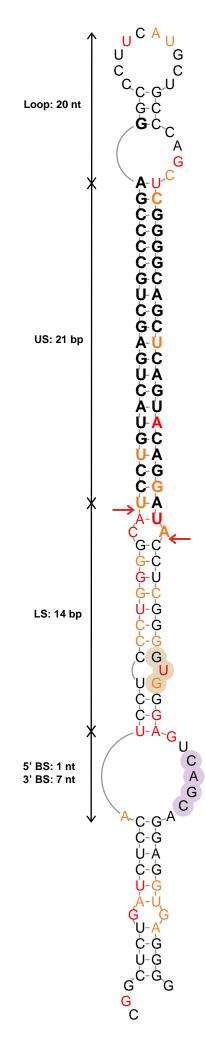
## pri-mir-486-1

chr8: 41660410-41660534 (-)

	Features	Optimality
Stem length (bp)	35	Optimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	20	Optimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.39 (24 / 476)	2.29 (10 / 476)
Cleavage homogeneity	0.6 (272 / 476)	0.76 (304 / 476)



Rank among tested pri-miRs

Top

**Bottom**