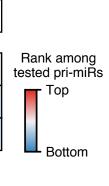
## pri-mir-371a

chr19: 53787650-53787774 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.79 (272 / 476)	1.48 (299 / 476)
Cleavage homogeneity	0.64 (262 / 476)	0.76 (301 / 476)



Loop: 11 nt

US: 23 bp

LS: 12 bp

5' BS: 2 nt 3' BS: 2 nt

