

**pri-mir-337**

chr14: 100874485-100874609 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	3	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	0	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.7 (419 / 476)
Cleavage homogeneity	0.03 (462 / 476)	0.55 (373 / 476)

