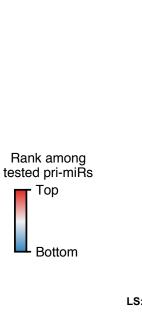
pri-mir-362

chrX: 50008938-50009062 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	5	
Apical loop size (nt)	5	Suboptimal
3' basal segment length (nt)	13	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.24 (66 / 476)	2.21 (22 / 476)
Cleavage homogeneity	0.8 (191 / 476)	0.85 (262 / 476)



Loop: 5 nt

