

pri-mir-3163

chr11: 66934405-66934529 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	1.1 (391 / 476)
Cleavage homogeneity	0.29 (380 / 476)	0.48 (397 / 476)

