

pri-mir-1908

chr11: 61815137-61815261 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	8	
Apical loop size (nt)	15	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)	0.0 (476 / 476)
Cleavage homogeneity	0.08 (443 / 476)	0.06 (457 / 476)

