

pri-mir-222

chrX: 45747001-45747125 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	16	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	7	
Apical loop size (nt)	23	Optimal
3' basal segment length (nt)	18	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.75 (286 / 476)	1.61 (264 / 476)
Cleavage homogeneity	0.55 (286 / 476)	1.08 (72 / 476)

Rank among tested pri-miRs

Top

Bottom

