

pri-mir-105-1

chrX: 152392194-152392318 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	10	Suboptimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.21 (394 / 476)	1.36 (341 / 476)	<div><div></div>Top</div> <div><div></div>Bottom</div>
Cleavage homogeneity	0.17 (405 / 476)	0.51 (387 / 476)	

