


pri-mir-155

chr21: 25573954-25574078 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	4	
Apical loop size (nt)	13	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	0.96 (192 / 476)	1.32 (351 / 476)
Cleavage homogeneity	1.0 (92 / 476)	1.03 (123 / 476)

Rank among
tested pri-miRs

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