

pri-mir-124-2

chr8: 64379142-64379266 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.31 (42 / 476)	2.06 (59 / 476)
Cleavage homogeneity	1.05 (61 / 476)	1.08 (74 / 476)

Rank among
tested pri-miRs

