

pri-mir-26a-1

chr3: 37969384-37969508 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	3	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.25 (61 / 476)	1.82 (155 / 476)
Cleavage homogeneity	1.14 (22 / 476)	1.18 (20 / 476)

Rank among
tested pri-miRs

