## pri-mir-103a-1

chr5: 168560870-168560994 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	4	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	
Cleavage efficiency	1.06 (146 / 476)	2.04 (68 / 476)	
Cleavage homogeneity	1.06 (58 / 476)	1.17 (23 / 476)	



