

pri-mir-105-2

chrX: 152394387-152394511 (-)

| | Features | Optimality |
|------------------------------|---------------|------------|
| Stem length (bp) | 35 | Suboptimal |
| Lower stem length (bp) | 13 | Optimal |
| Mismatched pairs in the stem | 3 | |
| 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 |
|----------------------|---------------------|---------------------|
| Cleavage efficiency | 0.0 (476 / 476) | 0.02 (456 / 476) |
| Cleavage homogeneity | 0.05 (454 / 476) | 0.23 (436 / 476) |

Rank among tested pri-miRs

Top

Bottom

