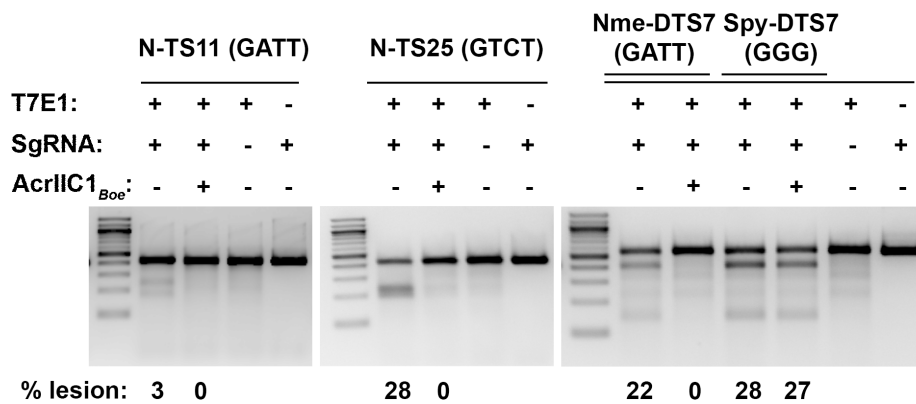
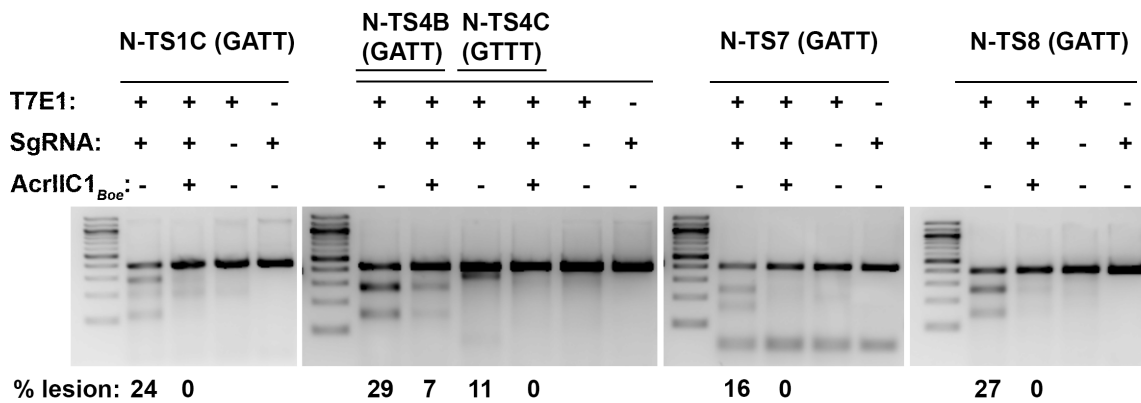


**A**



**B**

Site	Gene or locus	Spacer sequence	Target site
N-TS1C	SLC9A9	GUGGUCUGGGGUACAGCCUUGGCA	TACTTGGTCTGGGGTACAGCCTTGGC <u>ATCATGATT</u> TTG
N-TS4B	FLJ00328	GGACAGGAGUCGCCAGAGGCCGGU	GCAGGACAGGAGTGCAGAGAGGCCGGT <u>GGTGGATT</u> TCC
N-TS4C	FLJ00328	GGGGCUGGCUCCACGUCGCGCCGC	TGCGGGGCTGGCTCCACGTCGCGCCG <u>CGCGGTTT</u> GGG
N-TS7	LOC100505797	GAGGGAGAGAGGUGAGCGGAUGAA	GCAAAGGGAGAGAGGTGAGCGGATGAAGGGAGATTGGT
N-TS8	ESPN	GGACGCAAUCCAGAGGUGAUGGG	CGGCAGCAATTCCAGAGGTGATGGG <u>GAGTGATT</u> GTC
N-TS11	SMARCB1	GUUCCAGUUGGAAGGCCAGUGC	TAGATTCCAGTTGGGAAGGCCAGTGC <u>CTCCGATT</u> CCA
N-TS25	AC193513	GGUUUCUCAUCCUGUCUUCUGCCU	CCGCGTTTCTCATCTGTCTTCTGCTT <u>AGTGGATA</u> TGT
Nme-DTS3	ARHGEF9	GACUGAAGGCGAGGUCCGGGGCGG	GACTGAAGGCGAGGTCCGGGGCGG <u>AGGGGATT</u> GGG
Spy-DTS3	ARHGEF9	GAAGGCGAGGUCCGGGGCGG	GACTGAAGGCGAGGTCCGGGGCGG <u>AGGGGATT</u> GGG
Nme-DTS7	LSP1	GGCUGGCACCCUCAUGUACCCAG	GGCTGGCACCCCTCATGTACCCAG <u>GGGAGATT</u> CCA
Spy-DTS7	LSP1	GGCACCCUCAUGUACCCAG	GGCTGGCACCCCTCATGTACCCAG <u>GGGAGATT</u> CCA