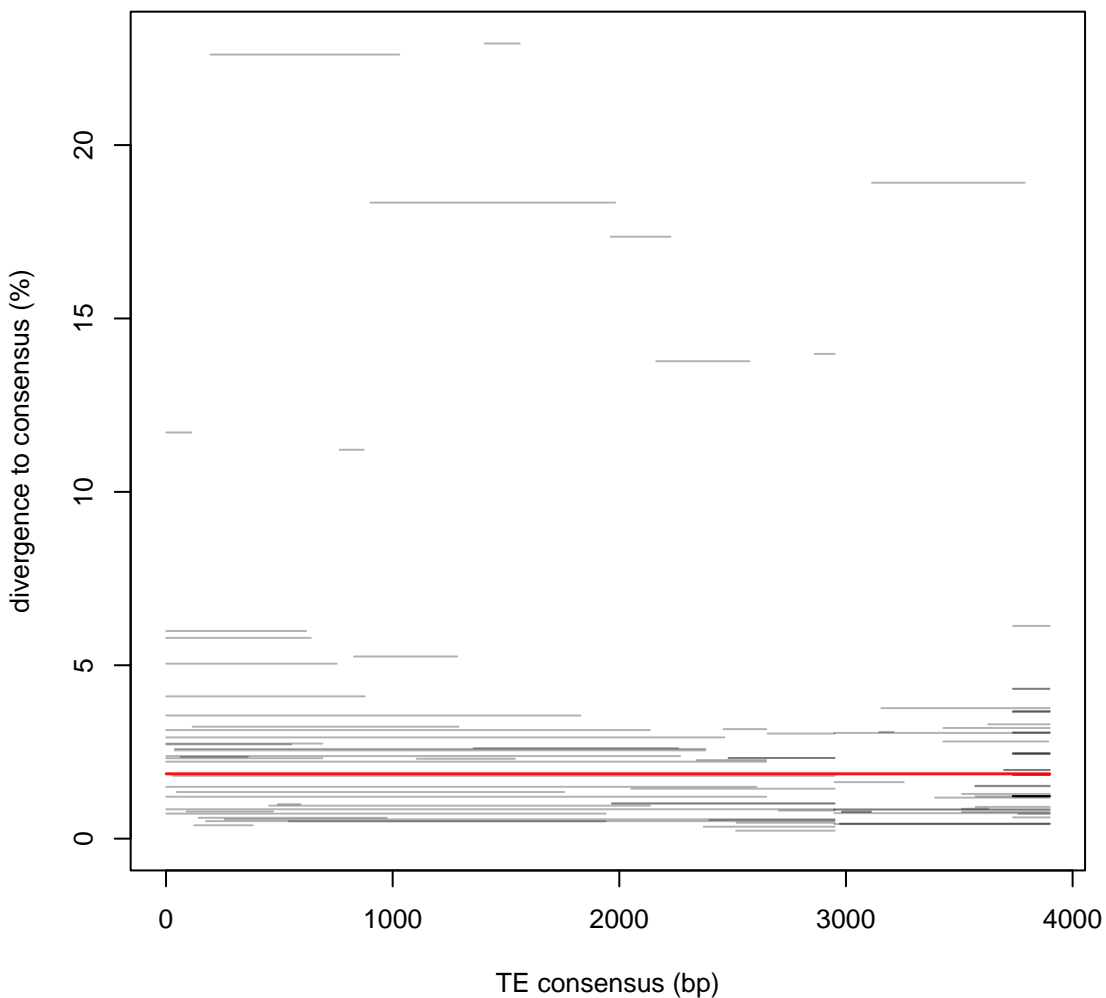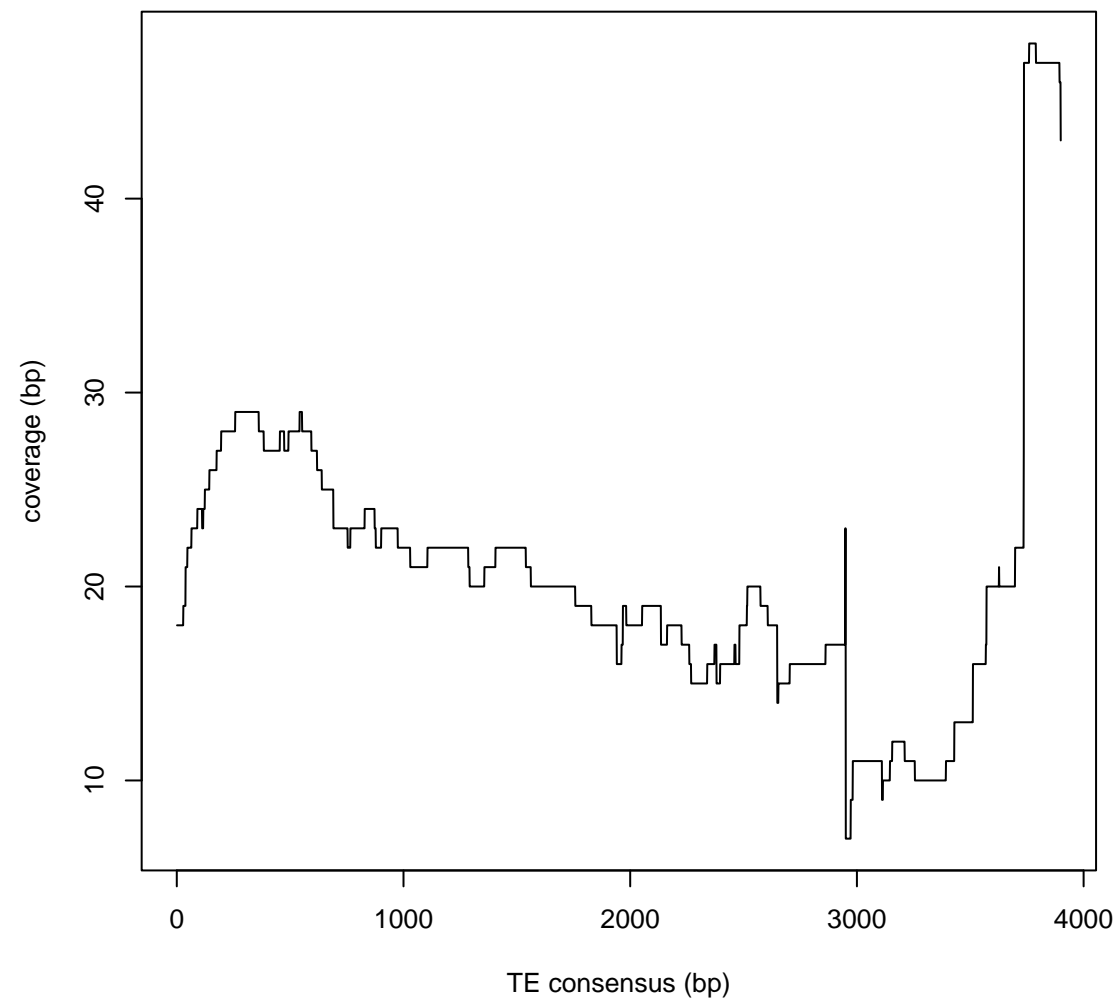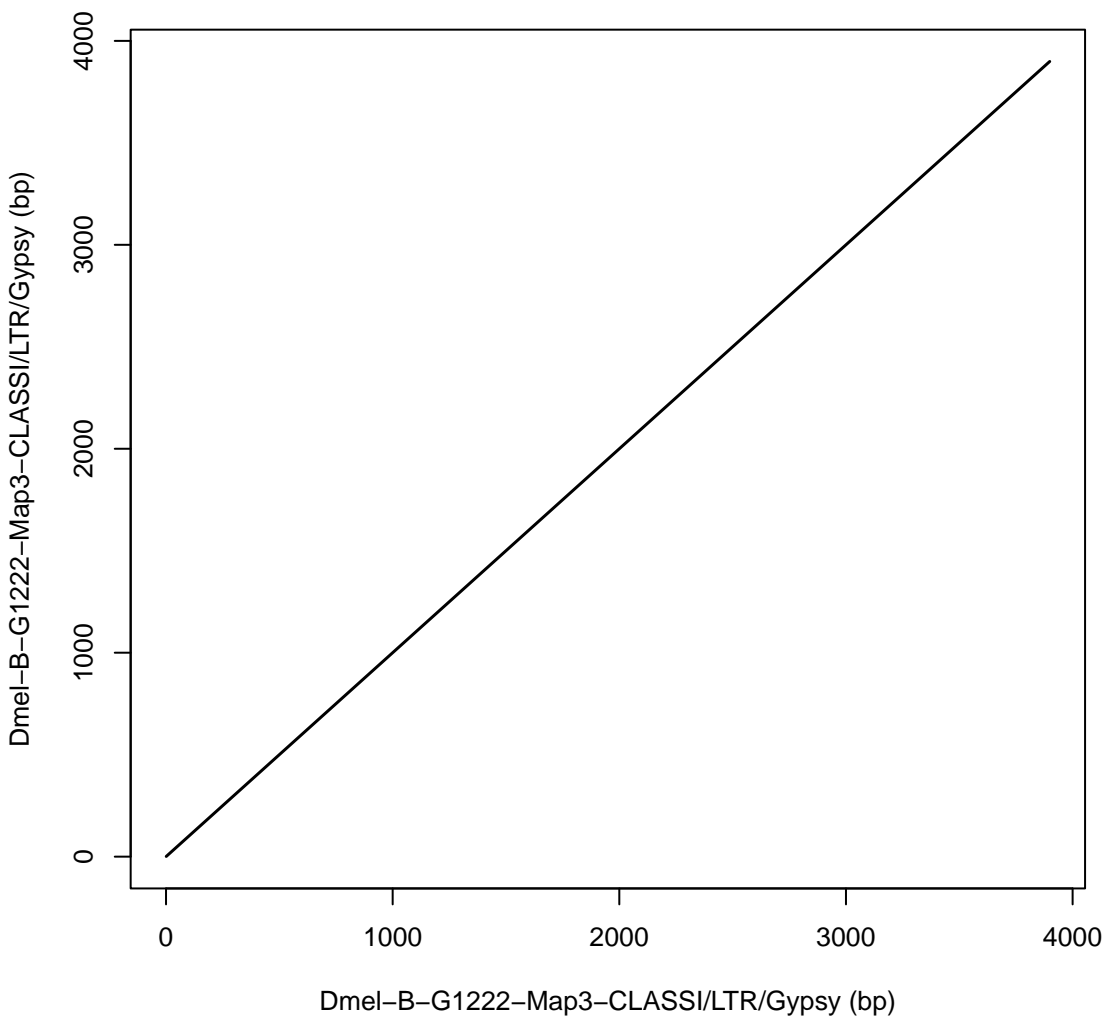**TE: Dmel−B−G1222−Map3−CLASSI/LTR/Gypsy**
consensus size: 3899bp; fragments: 107; full length: 1 (>=3509.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

sy−18_DAn_1p LTR/Gypsy

MuDR−1_DPu TIR/MULE−MuD

Bica_gagpol LTR/Gypsy