TE: Dmel−B−R578−Map6−CLASSI/LTR/Gypsy
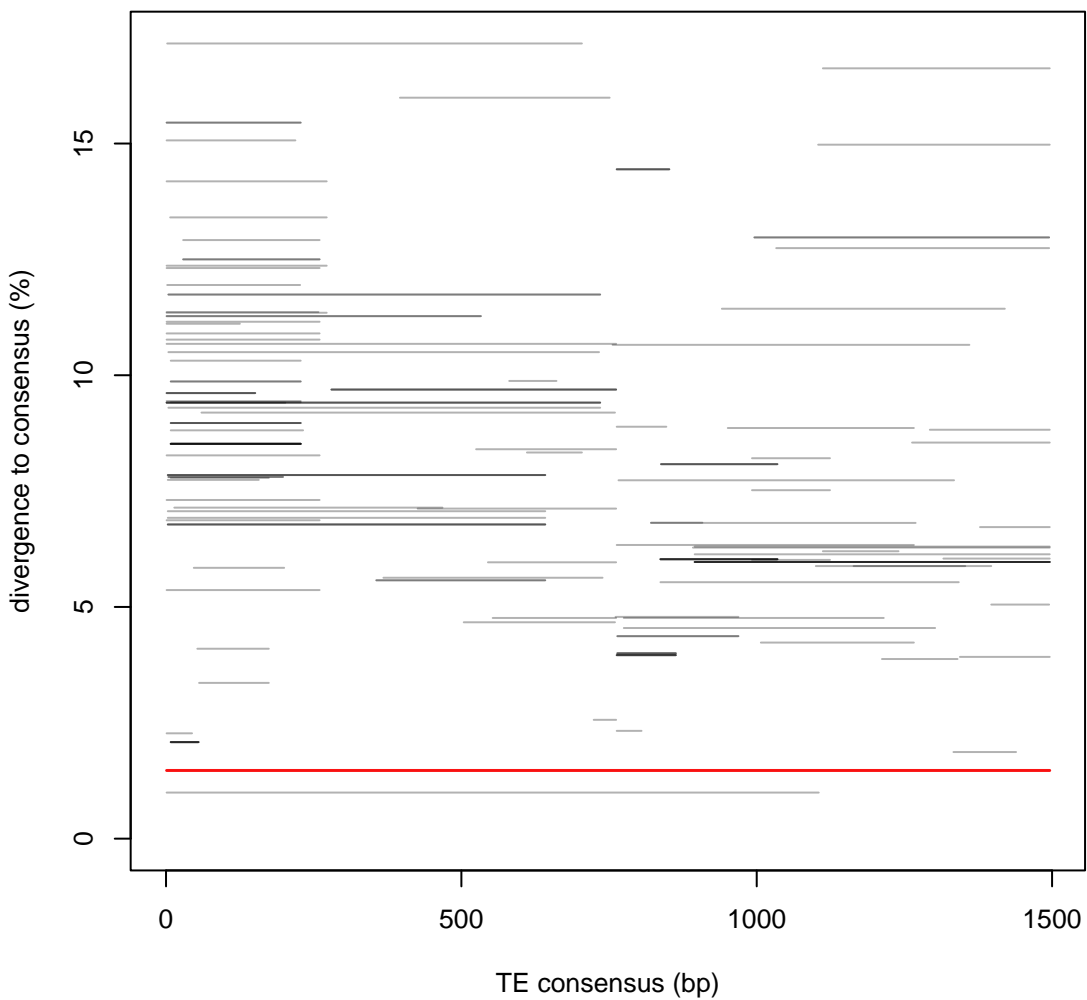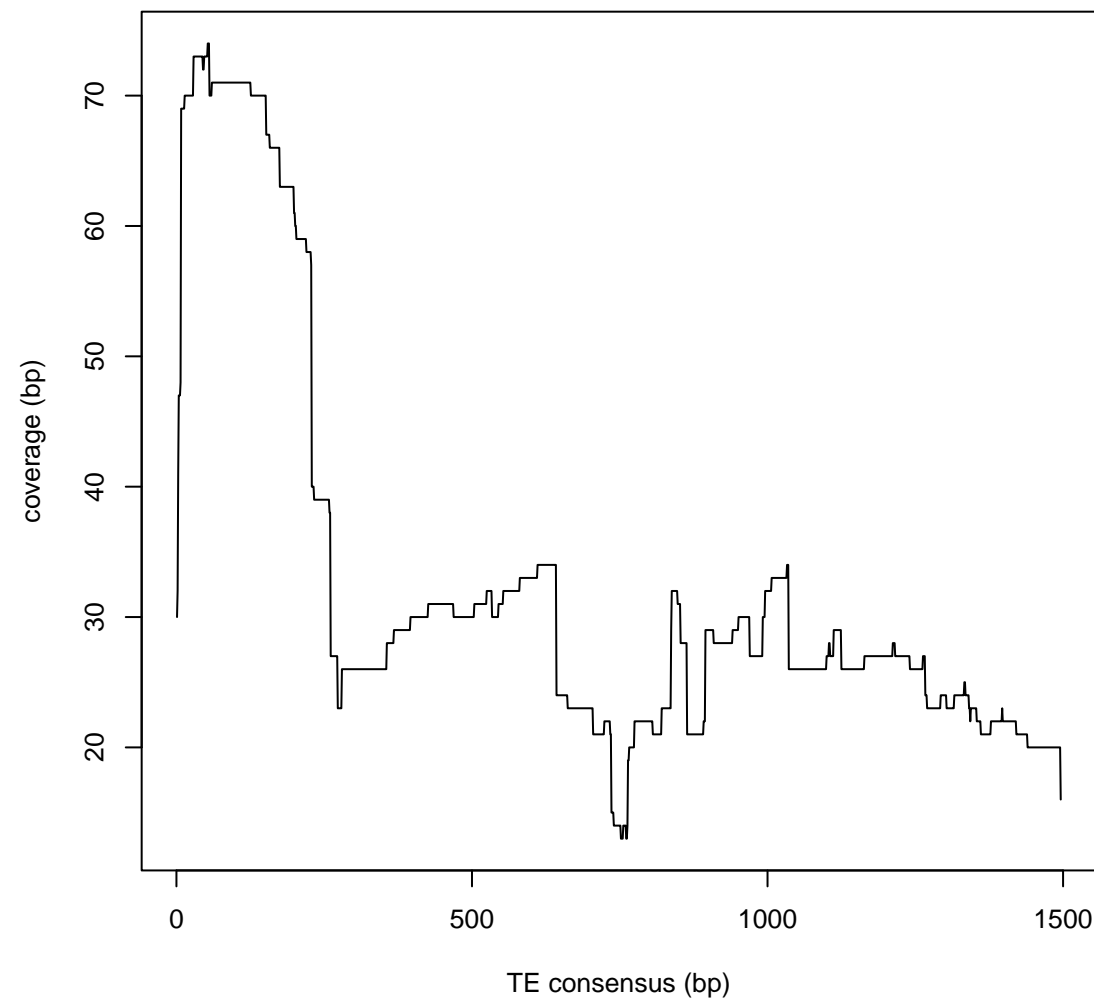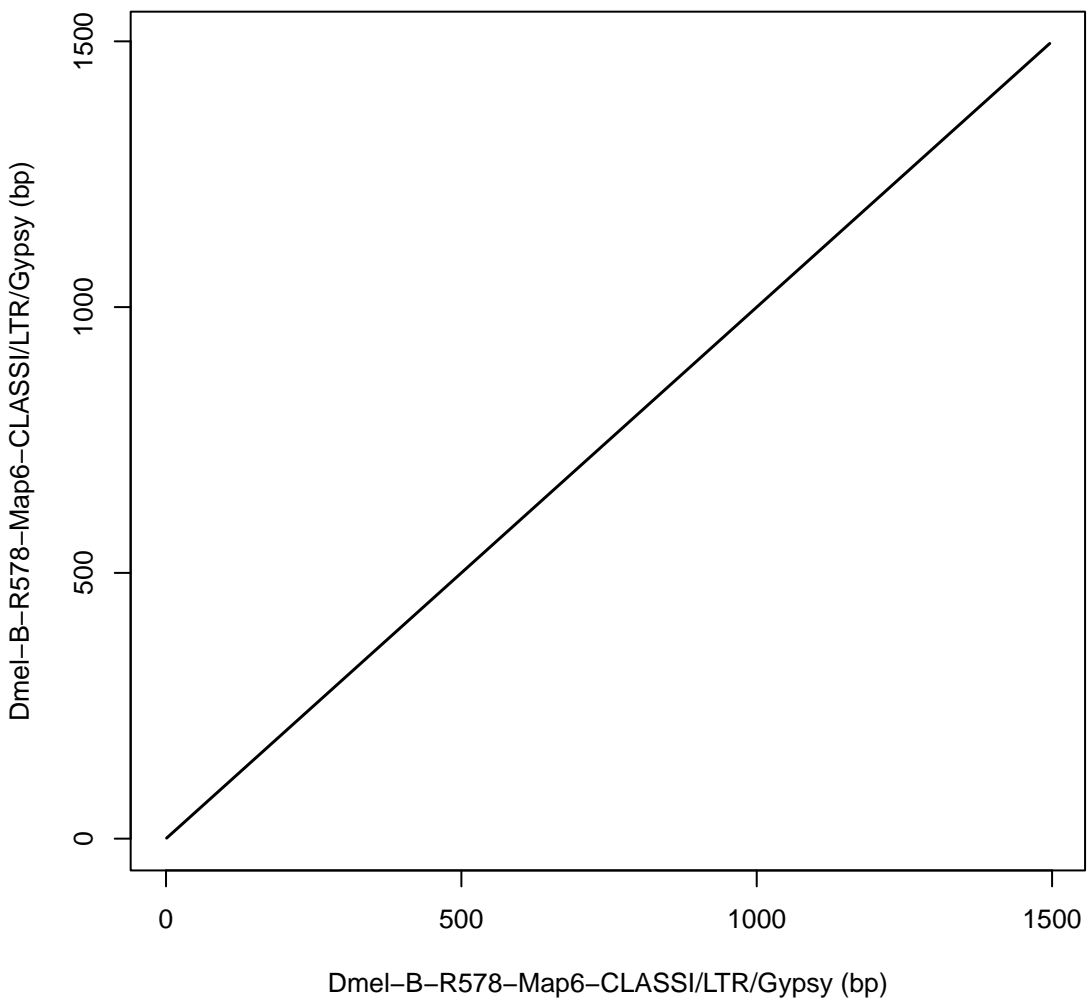**consensus size: 1496bp; fragments: 153; full length: 1 (>=1346.4bp)**

TE consensus genomic coverage

divergence to consensus (%)

TE consensus (bp)

coverage (bp)

TE consensus (bp)

TE consensus self dotplot (blastn)

Dmel−B−R578−Map6−CLASSI/LTR/Gypsy (bp)

Dmel−B−R578−Map6−CLASSI/LTR/Gypsy (bp)

TE consensus structure and protein hits

QUASIMODO_gag LTR/Gypsy

Dmel−B−R578−Map6−CLASSI/LTR/Gypsy (bp)