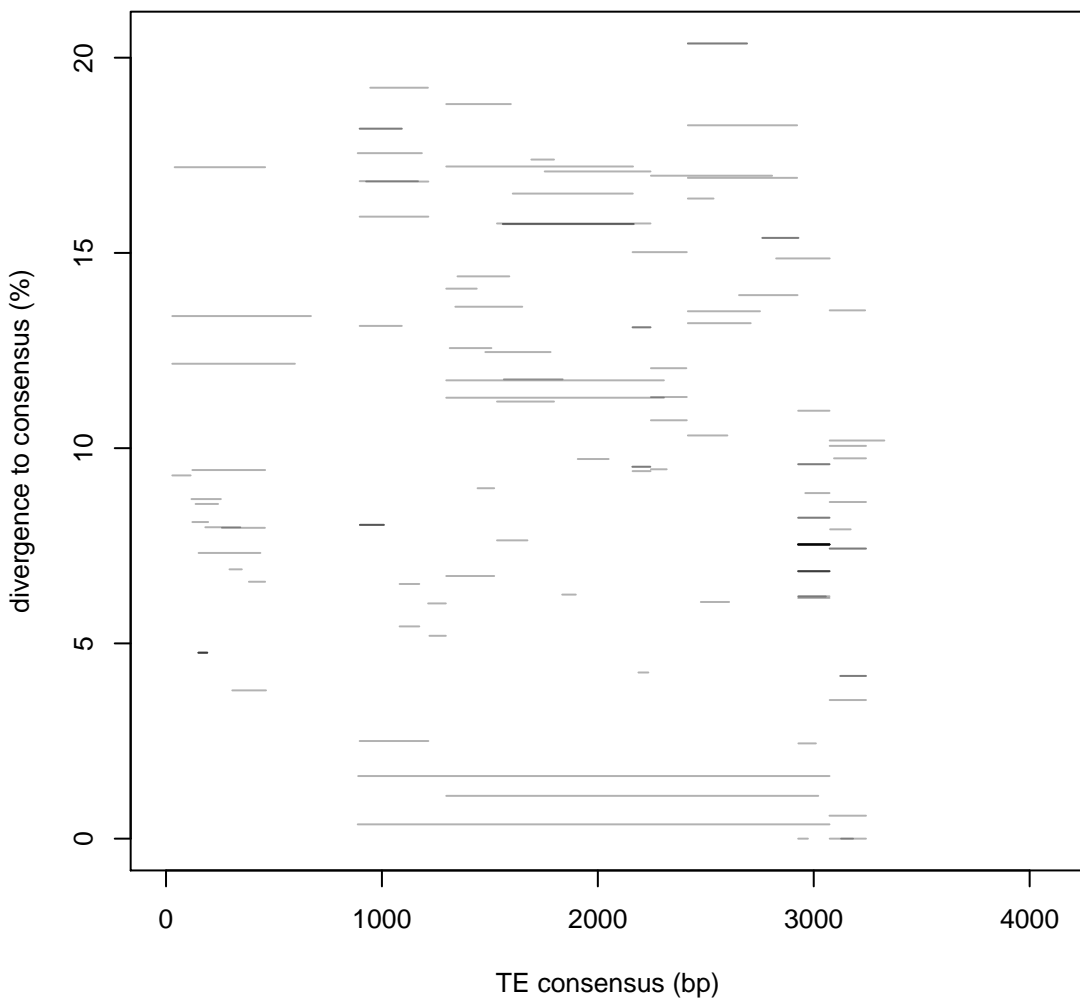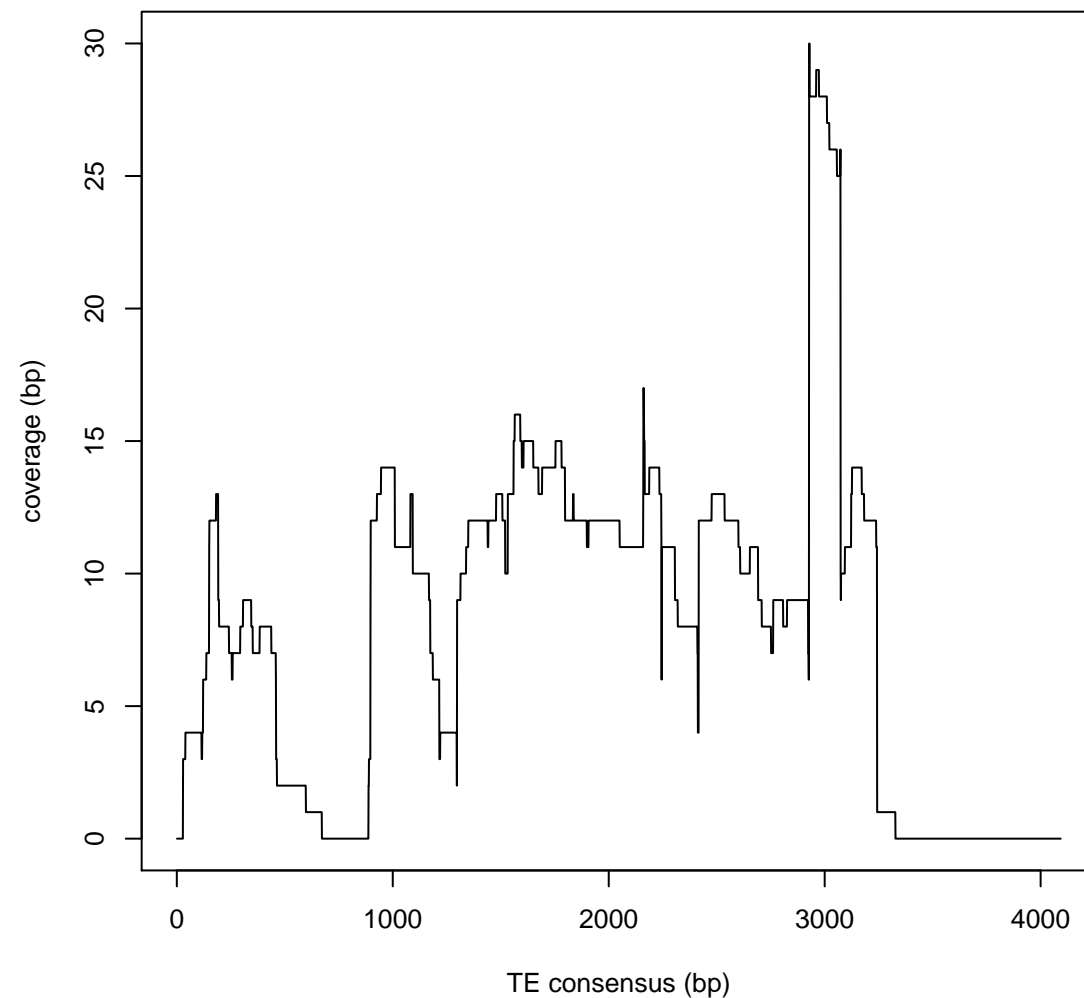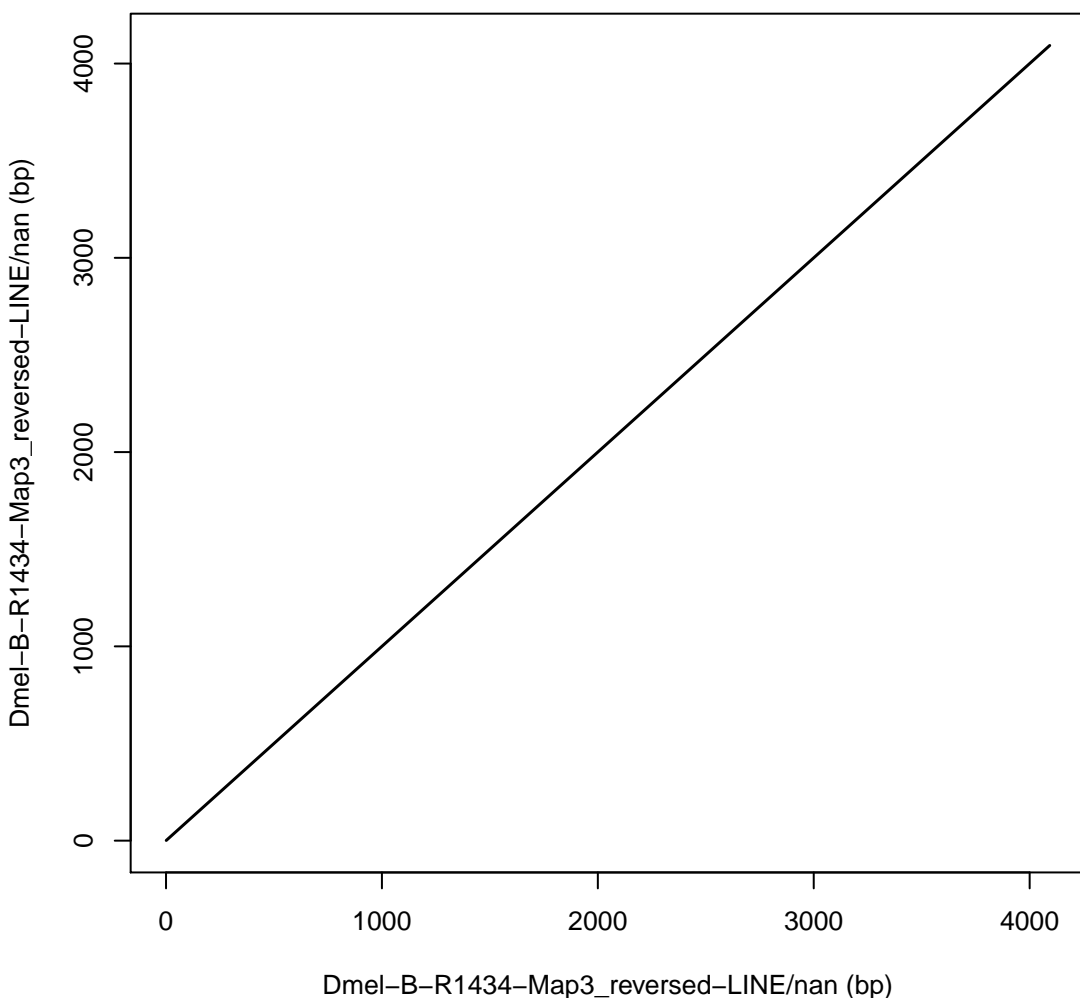**TE: DmeI−B−R1434−Map3_reversed−LINE/nan**
consensus size: 4093bp; fragments: 123; full length: 0 (>=3683.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

BURDOCK_pol LTR/Gypsy

BURDOCK_gag LTR/Gypsy

BURDOCK_gag LTR/G

divergence to consensus (%)

coverage (bp)

TE consensus (bp)

Dmel−B−R1434−Map3_reversed−LINE/nan (bp)

Dmel−B−R1434−Map3_reversed−LINE/nan (bp)