TE: Dmel−B−G1614−Map3−LTR/Gypsy
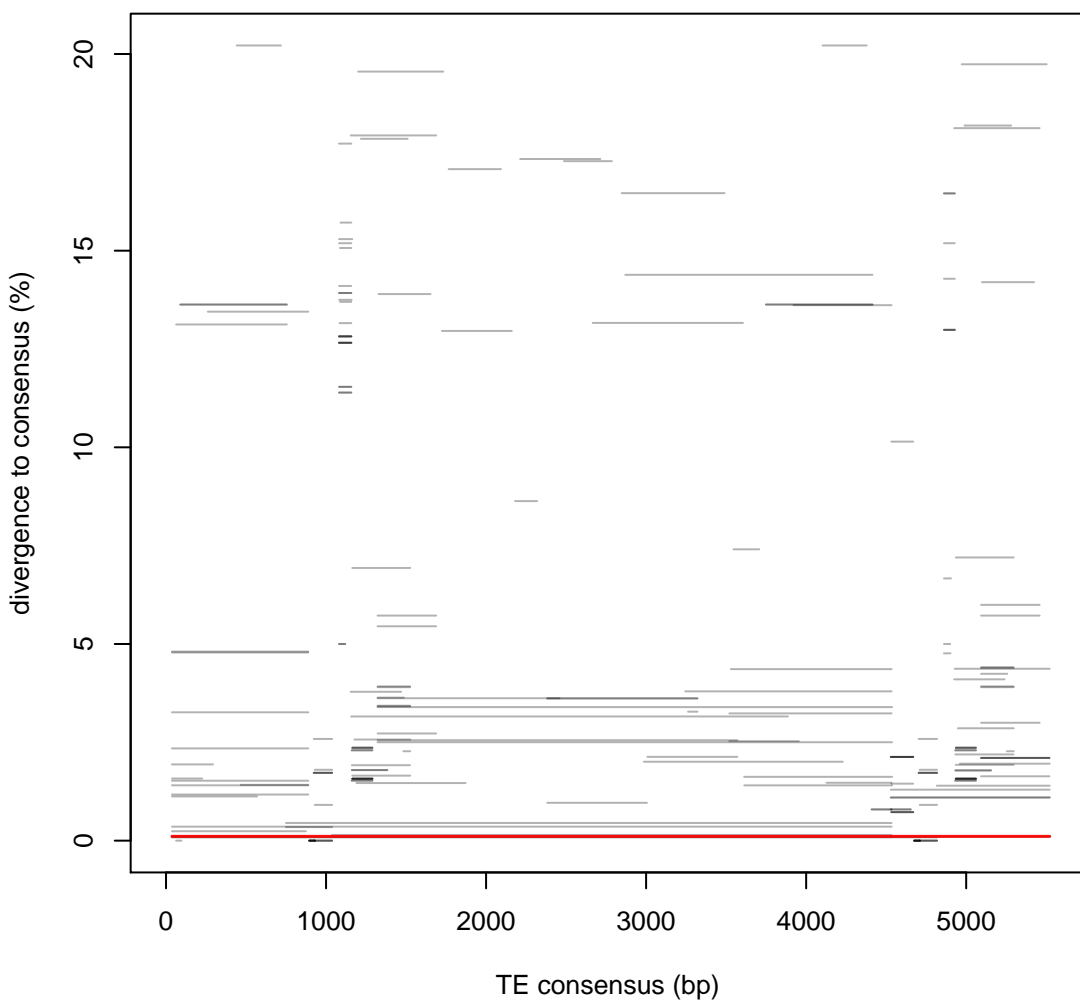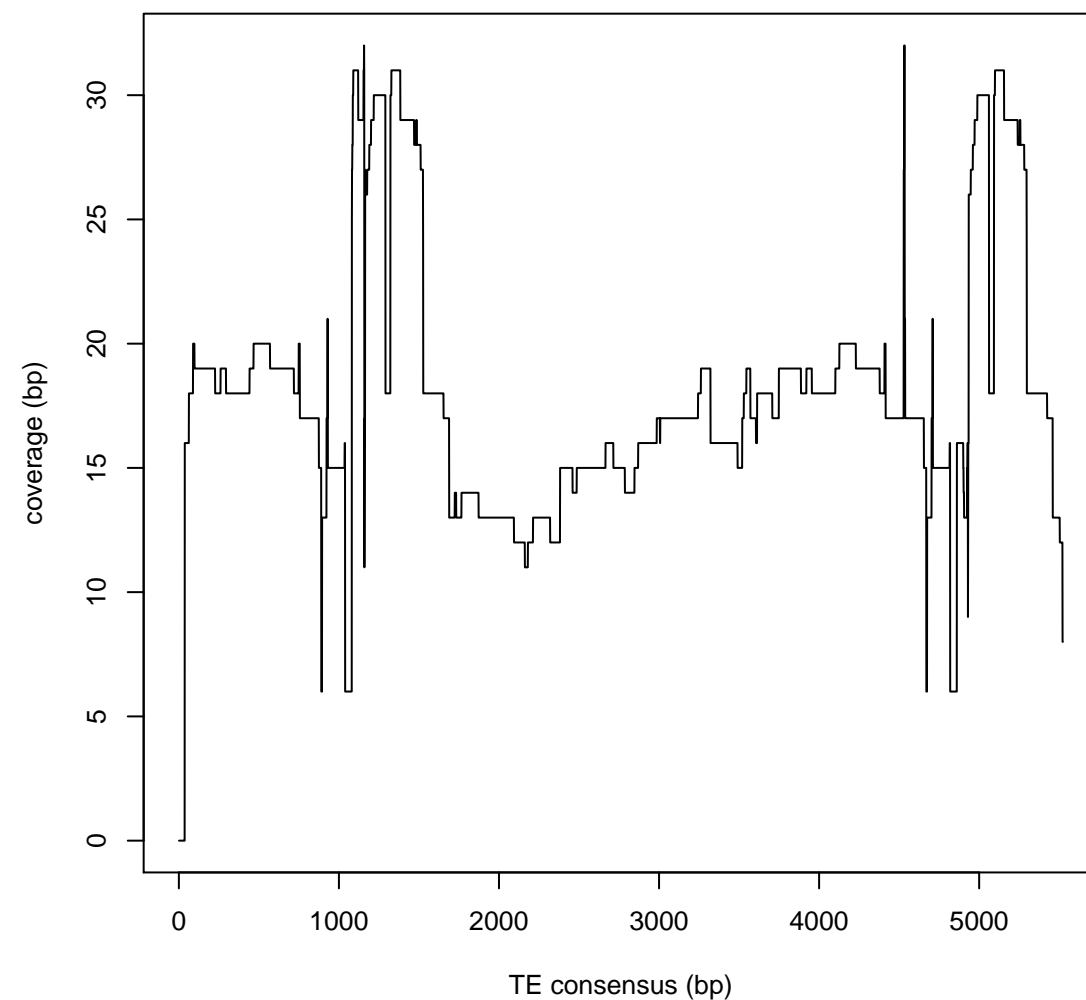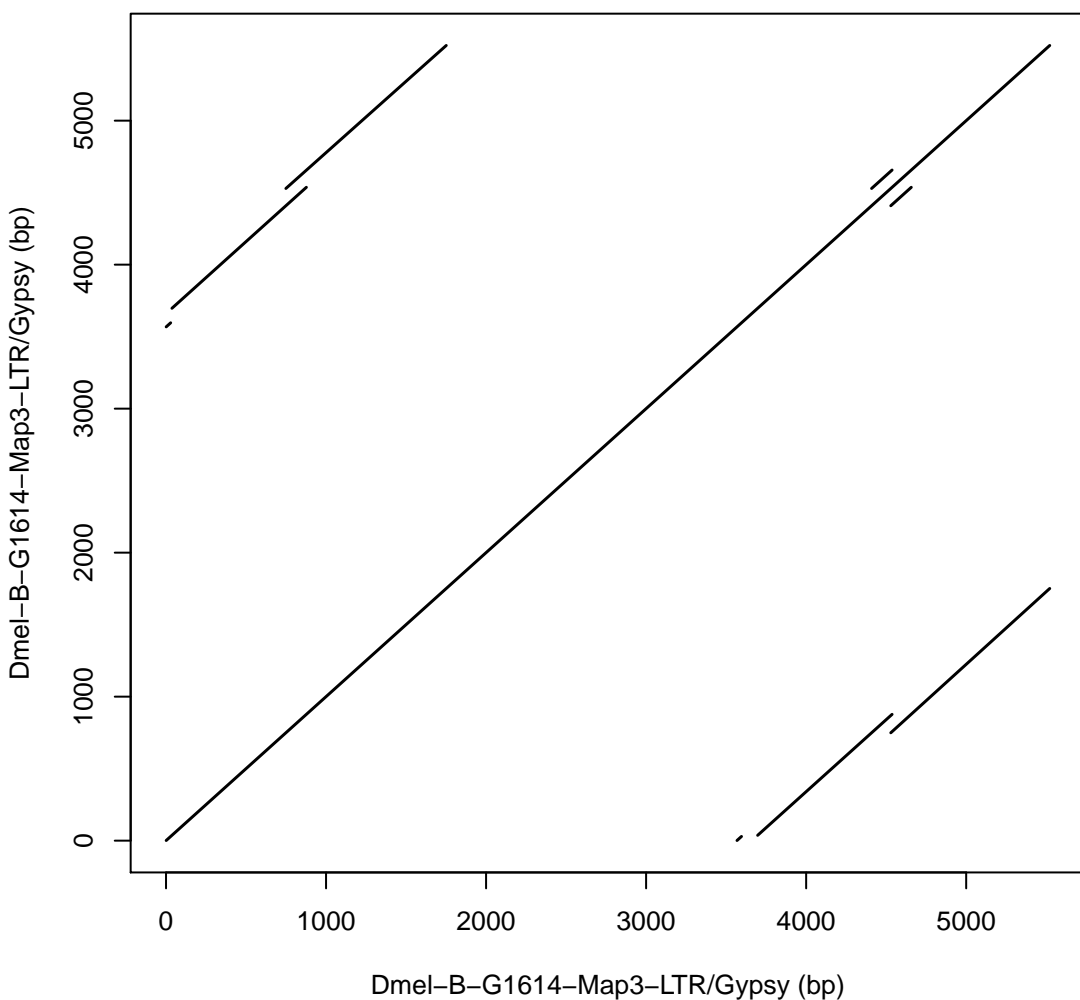consensus size: 5522bp; fragments: 213; full length: 1 (>=4969.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

y−30_DWil LTR/Gypsy

Gypsy−30_DWil LTR/Gypsy

Mariner−2_DBi_tp TIR/TcMar−Tc1

Gypsy−30_DWil LTR/Gypsy