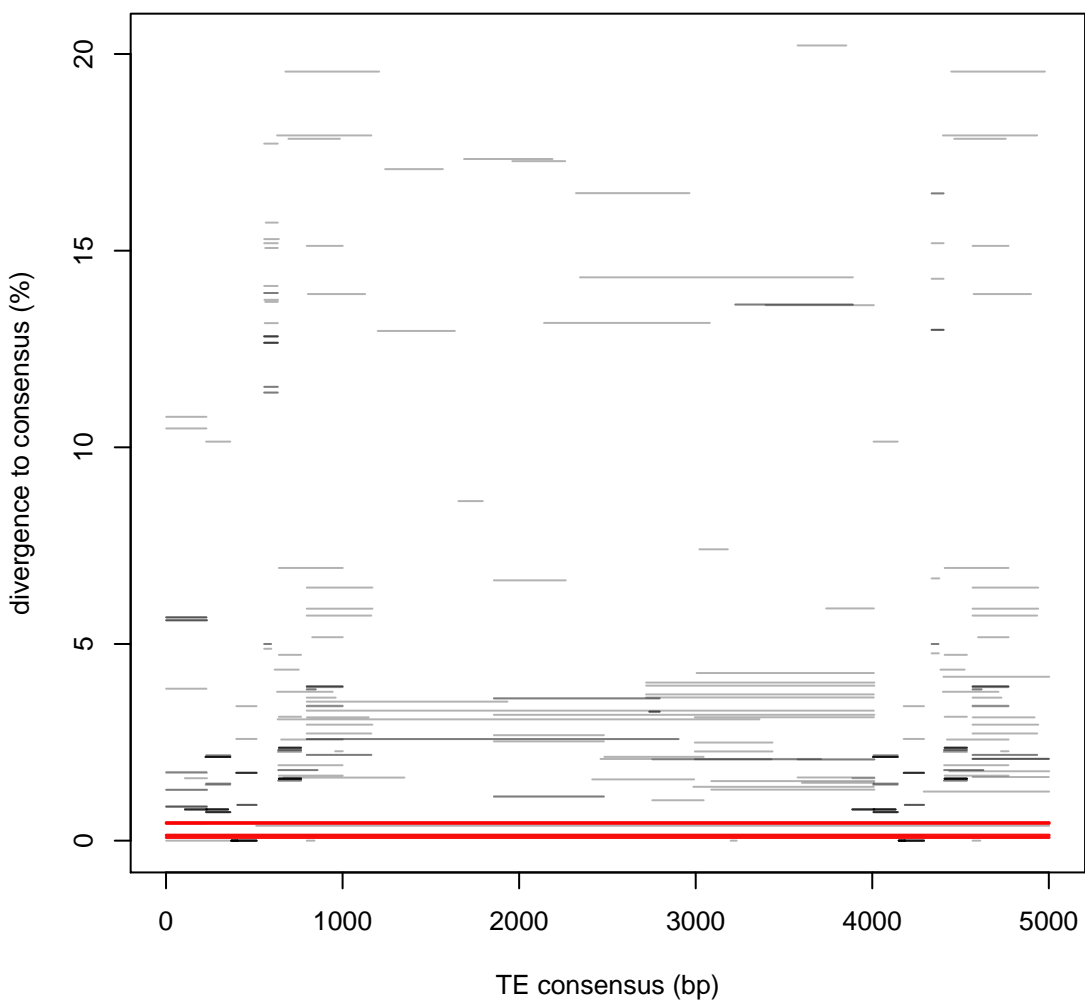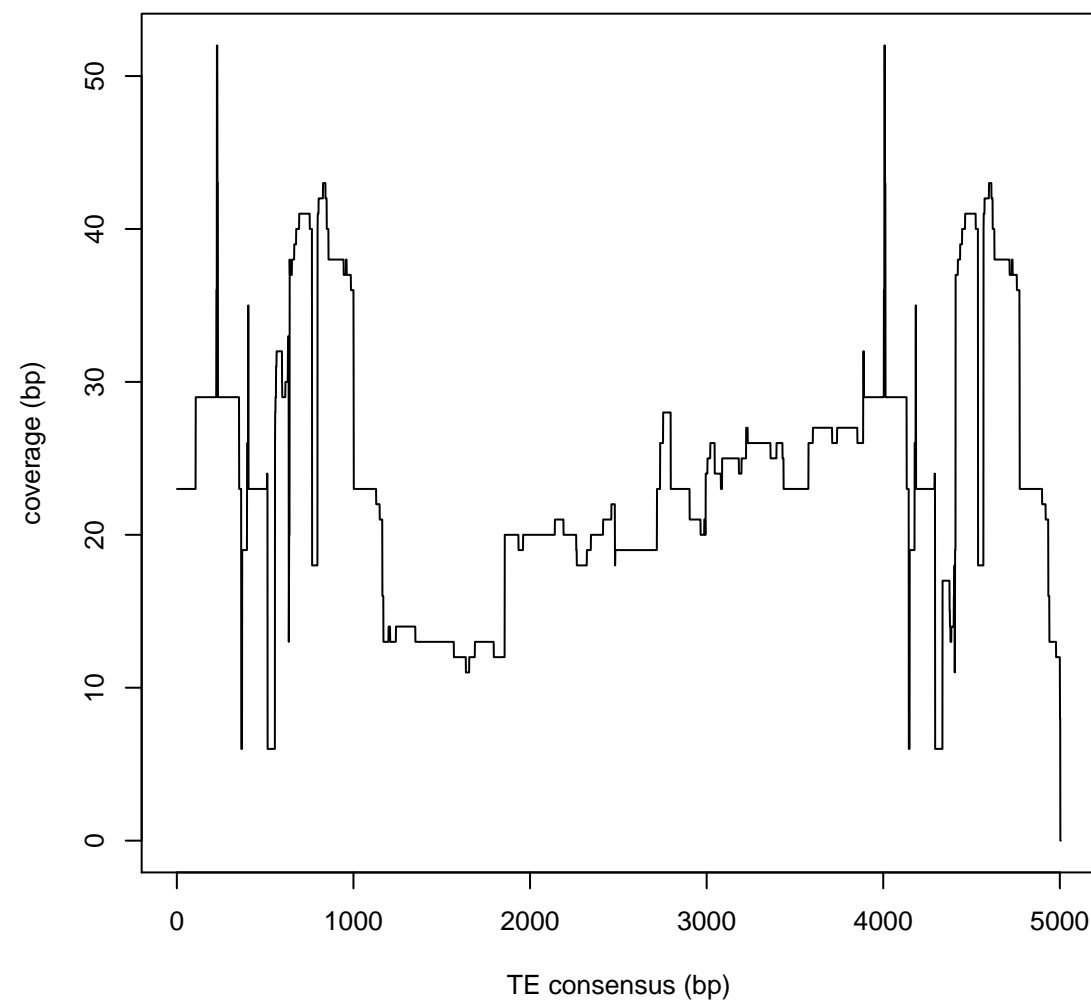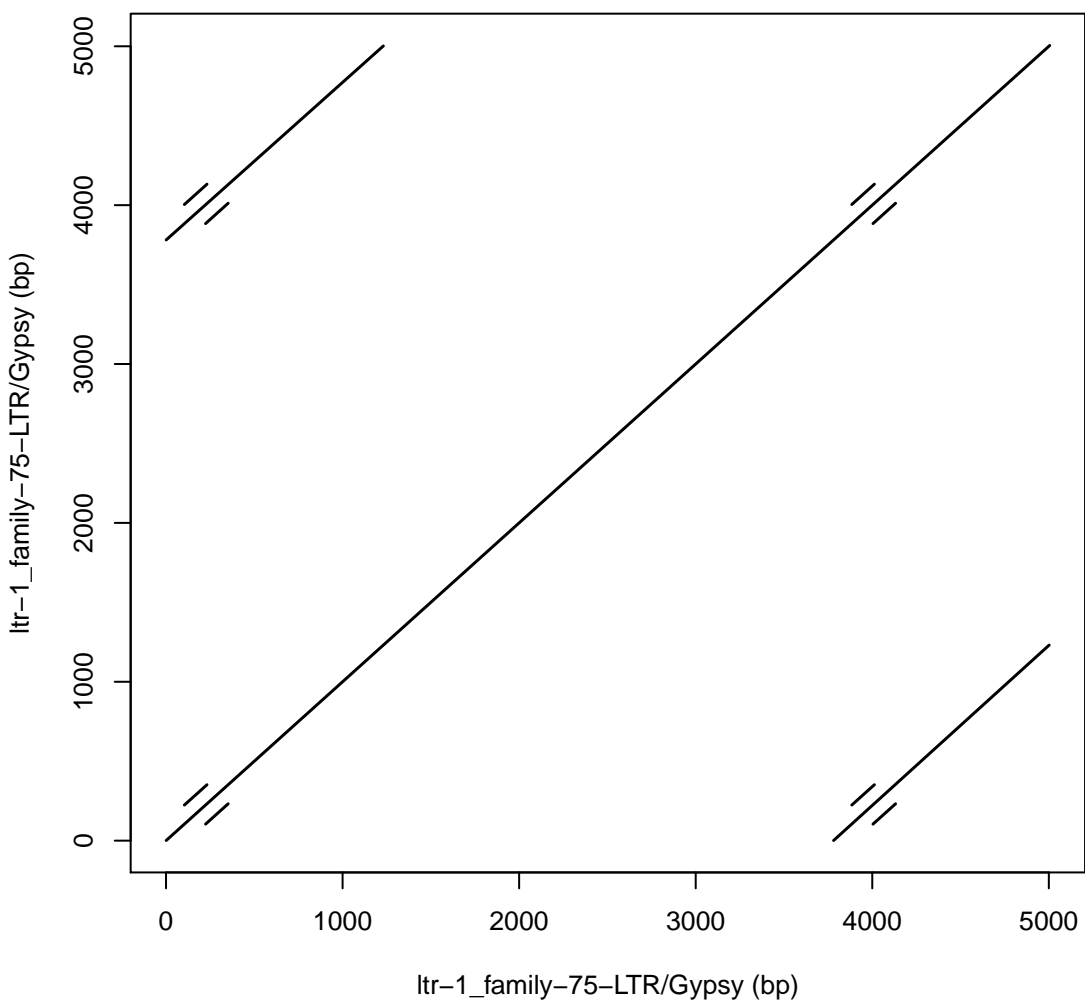**TE: ltr-1_family-75-LTR/Gypsy**
consensus size: 5005bp; fragments: 348; full length: 4 (>=4504.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

DBi_tp TIR/TcMar-Tc1
Gypsy-30_DWil LTR/Gypsy
Mariner-2_DBi_tp TIR/TcMar-Tc1
Gypsy-30_DWil LTR/Gypsy