TE: Dmel−B−R32−Map4−CLASSI/LINE/nan
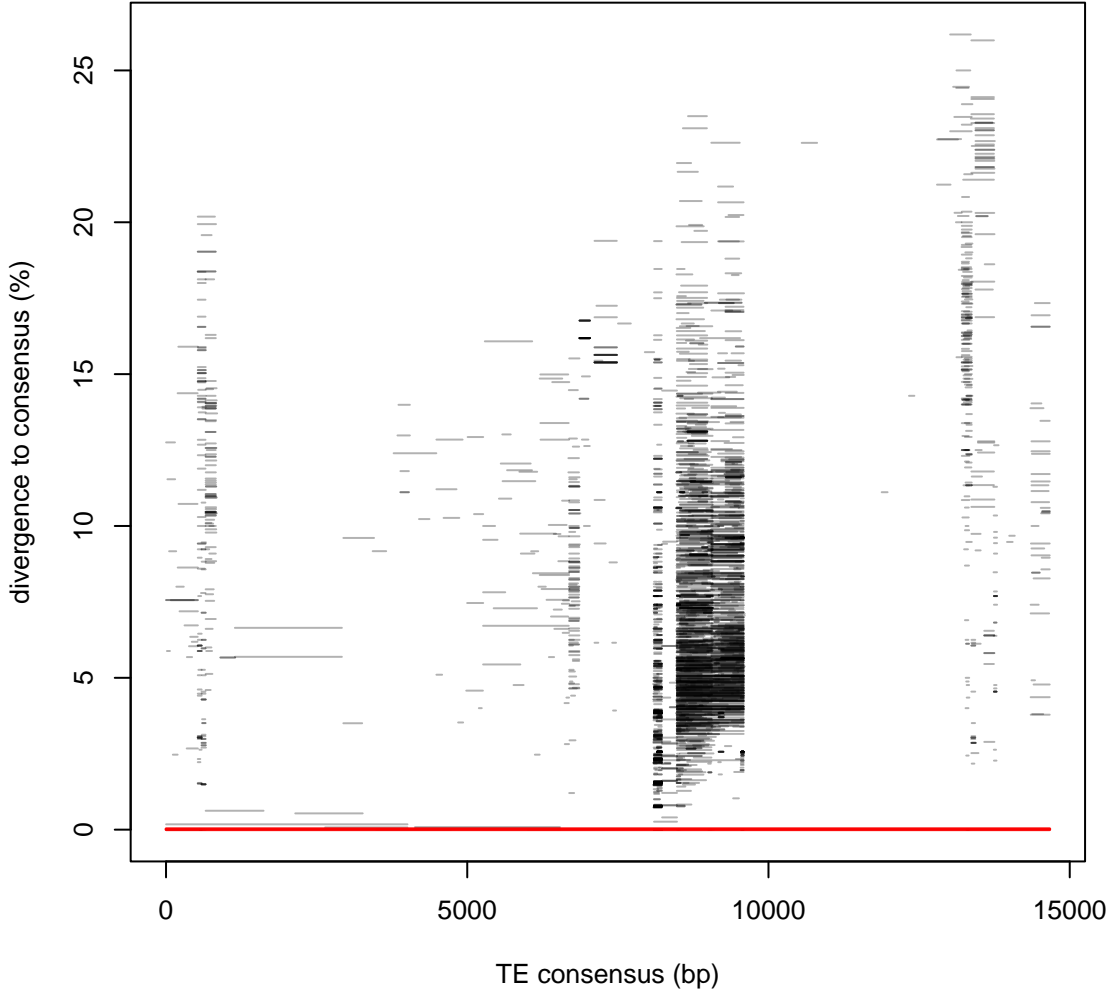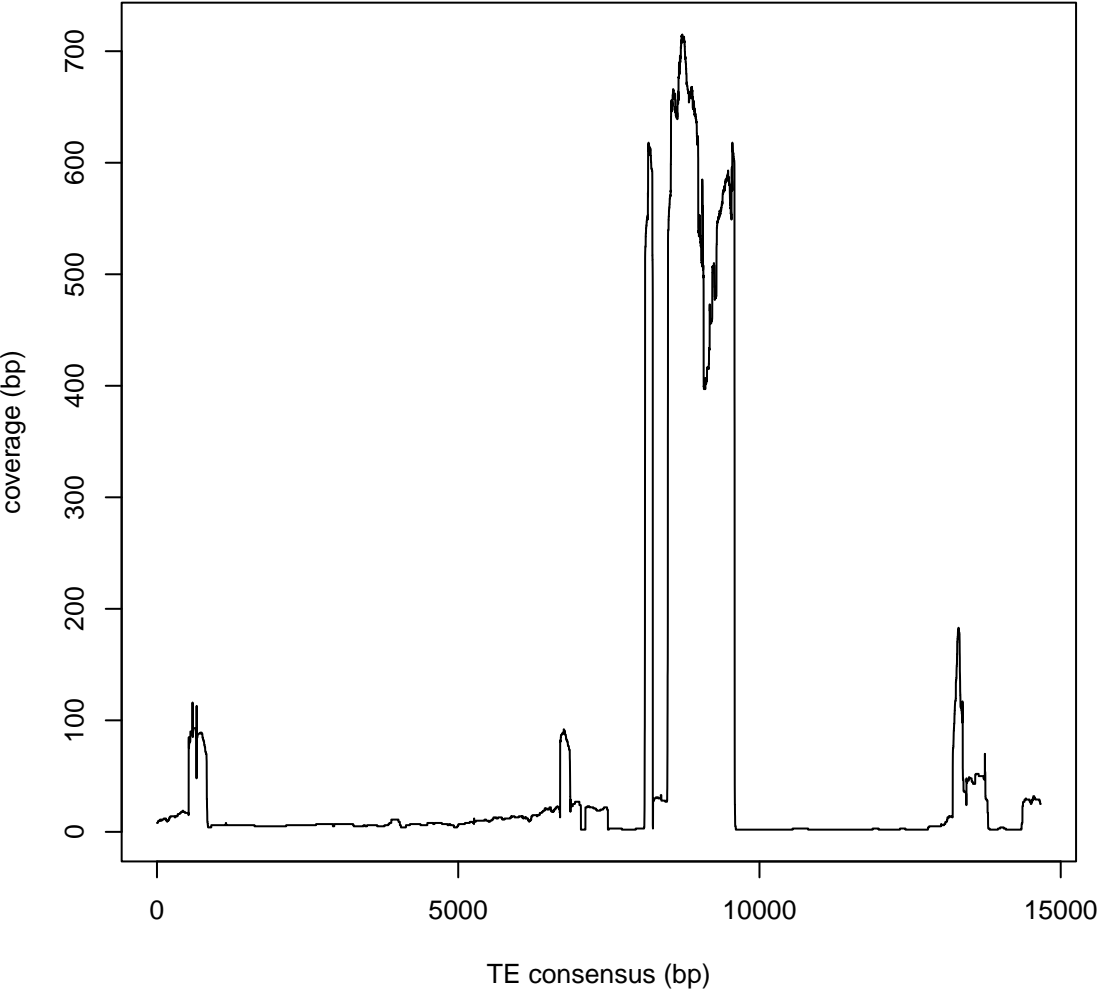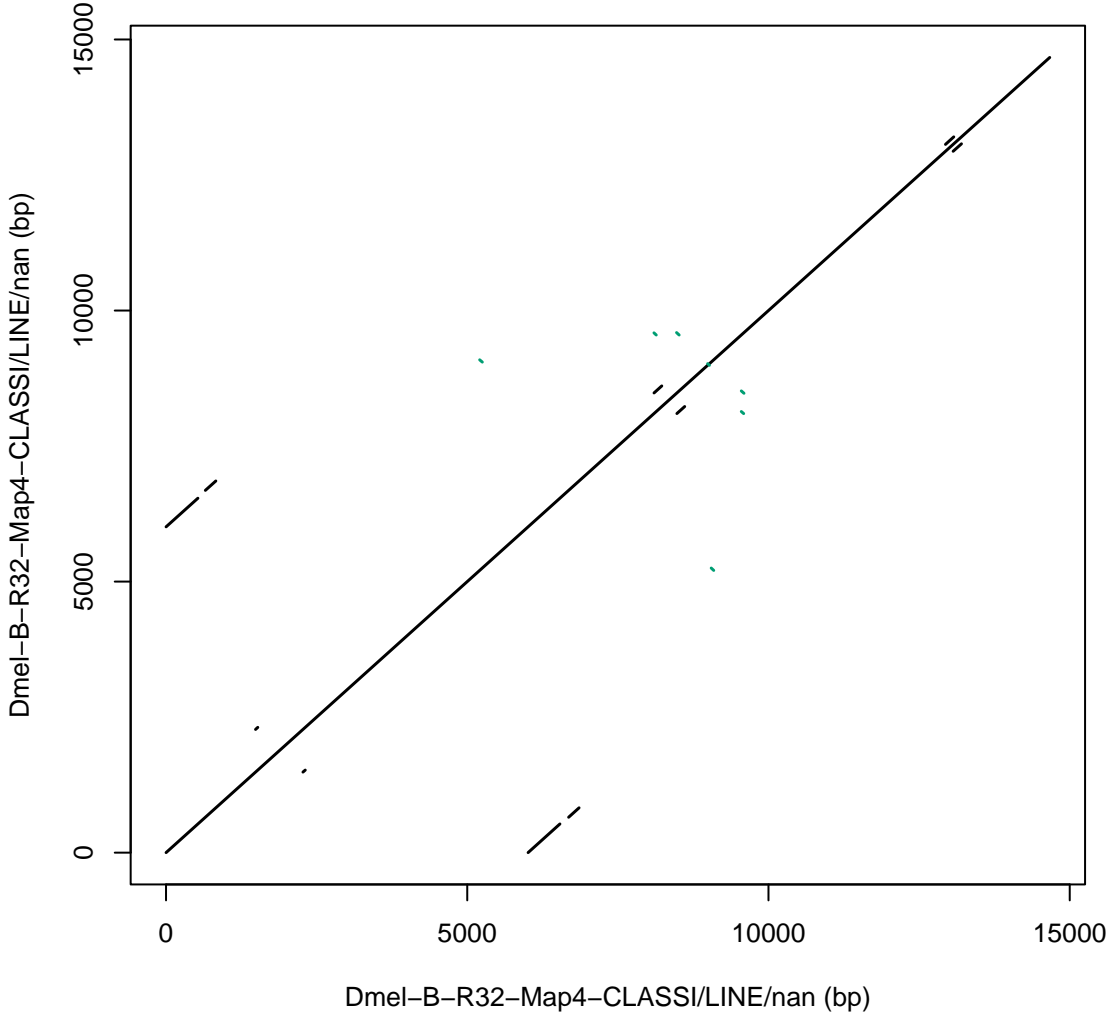consensus size: 14668bp; fragments: 3143; full length: 2 (>=13201.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

pol LINE/I−Jockey
Gypsy−6_DBp LTR/Gypsy
G3−DM_pol LINE/I−Jockey
BEL−6_DEI_pol_pol LTR/Pao
BEL1_SP−pol LTR/Pao
Copia−31_CCri LTR/Copia