TE: Dmel−B−G2605−Map3−CLASSI/LTR/Gypsy
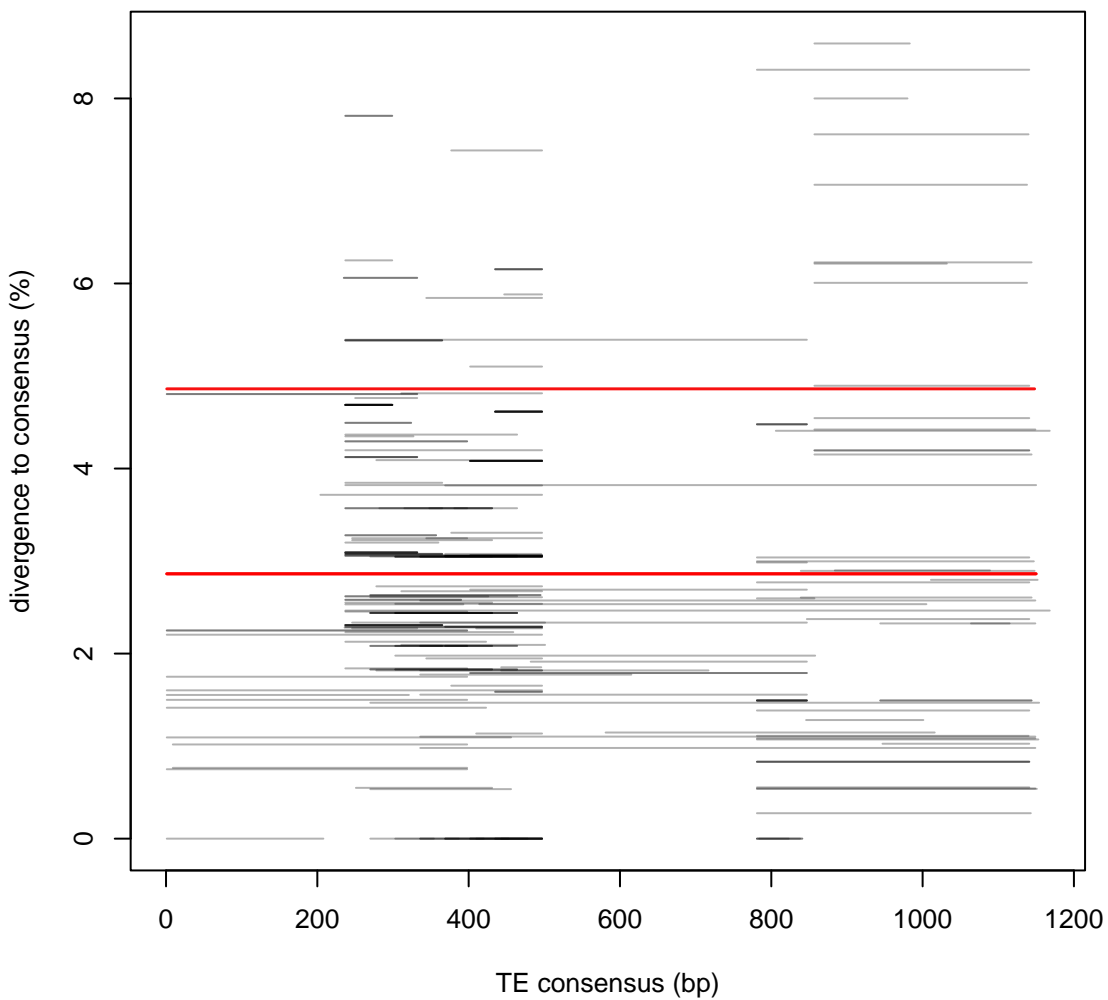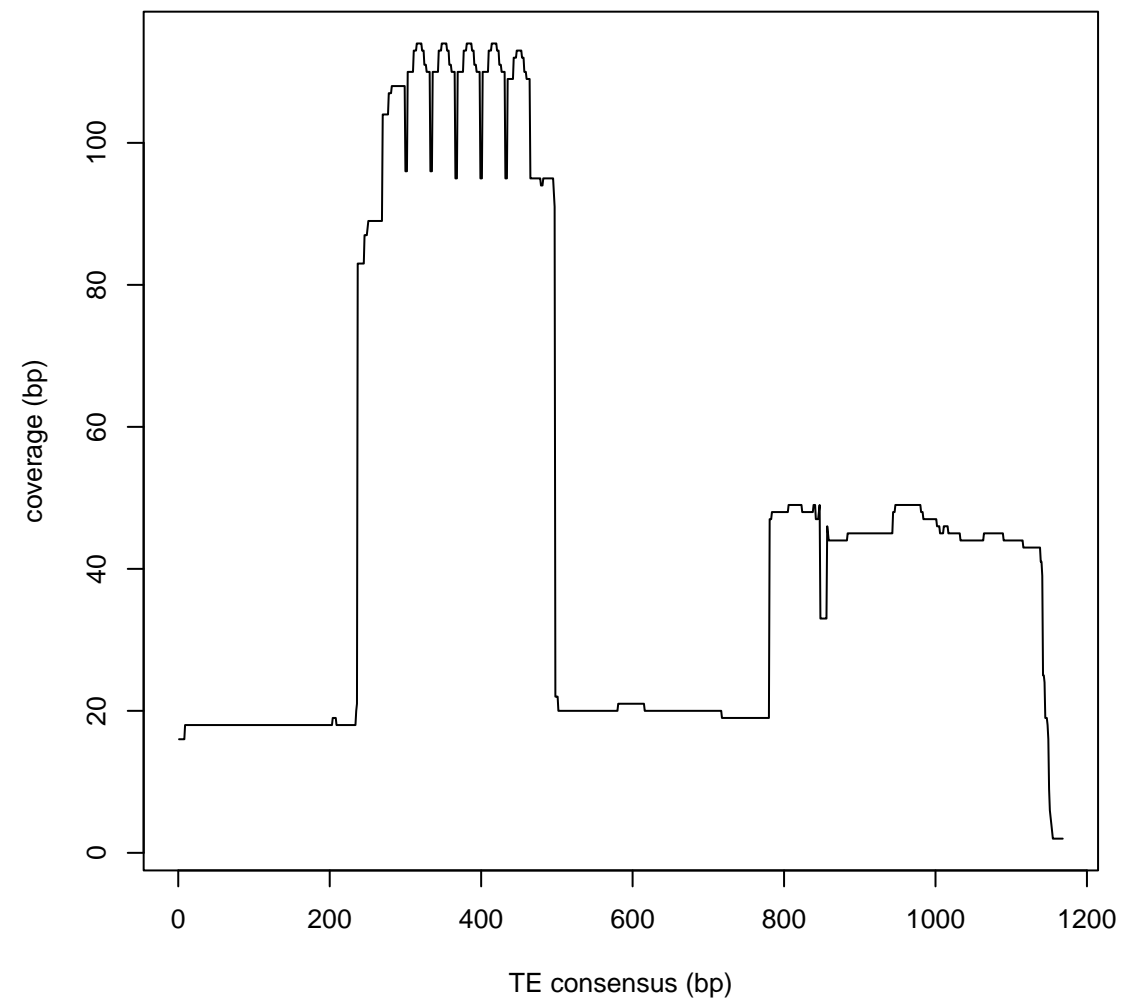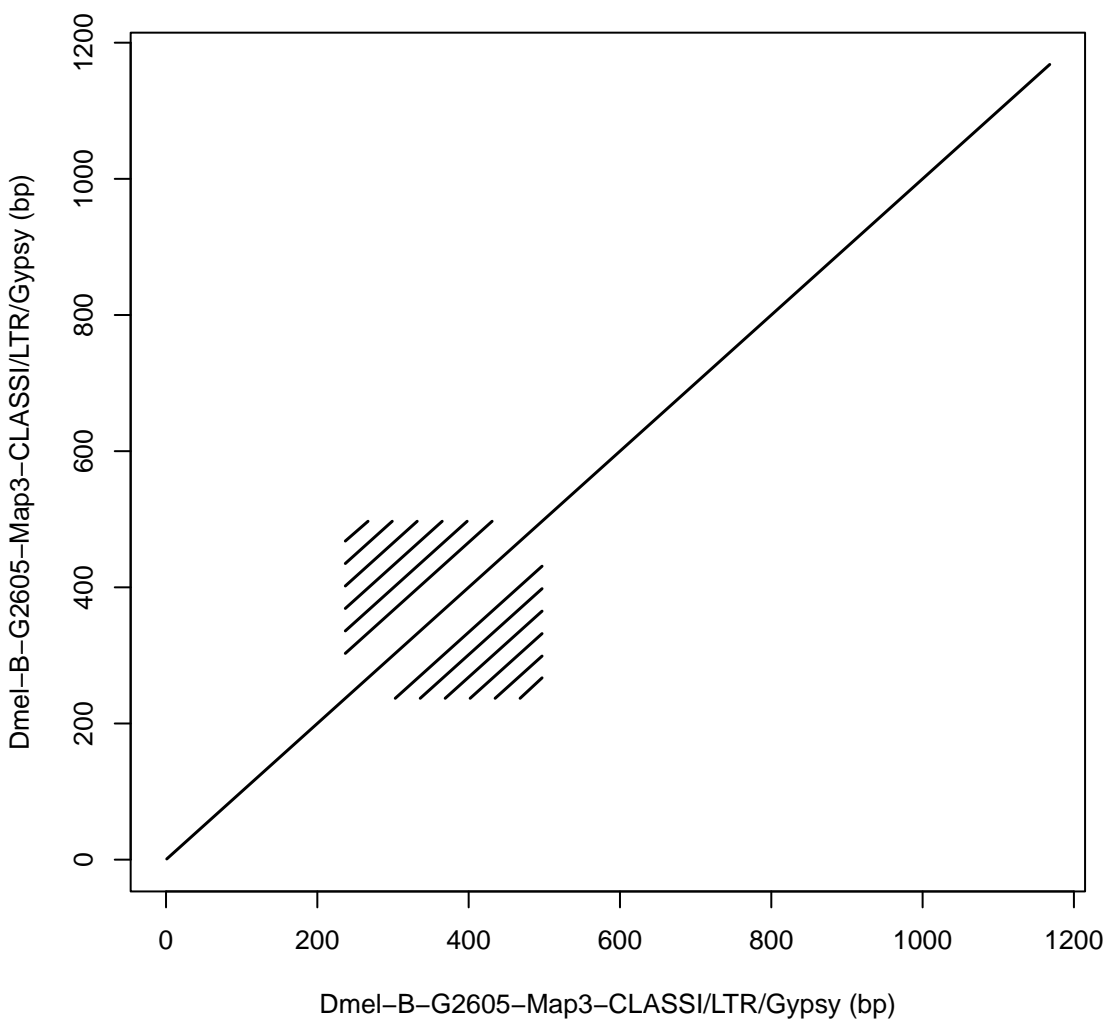**consensus size: 1168bp; fragments: 257; full length: 3 (>=1051.2bp)**

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected