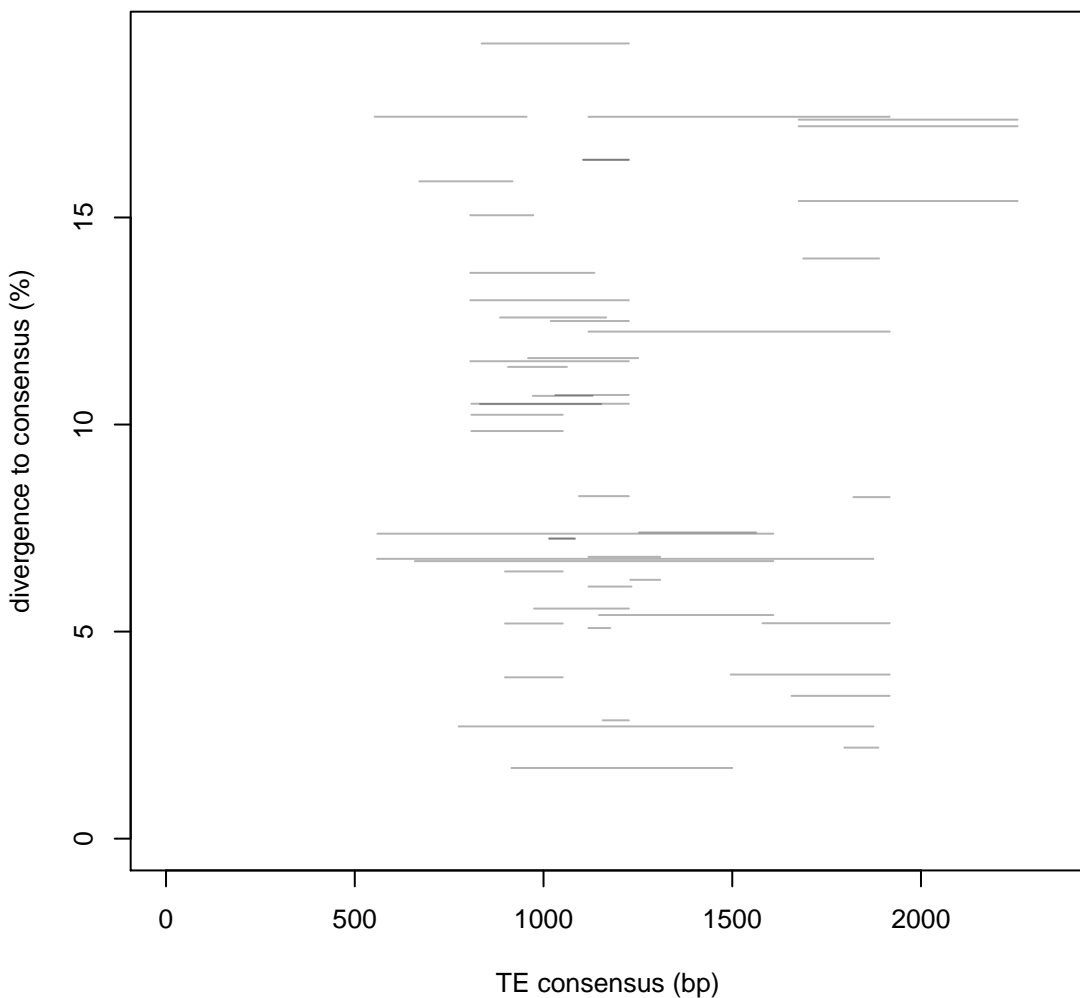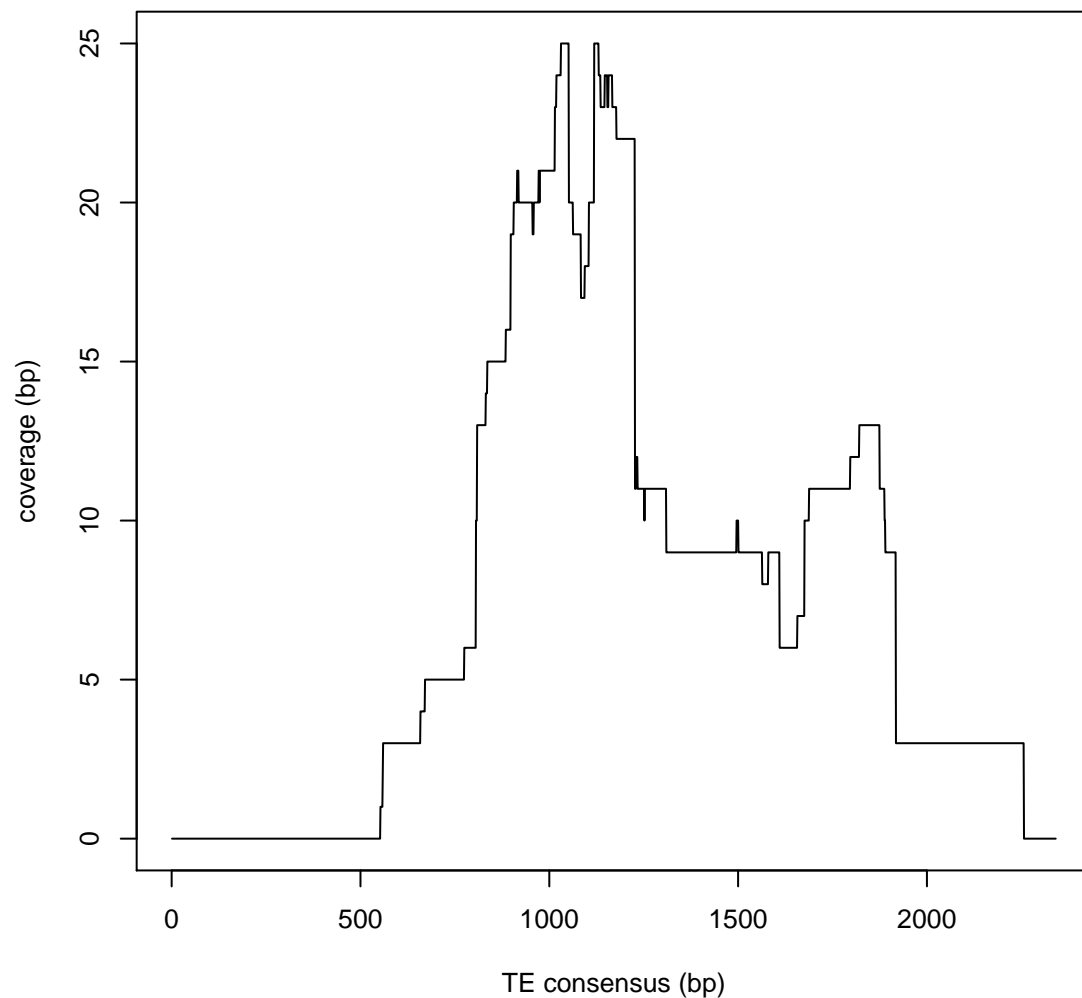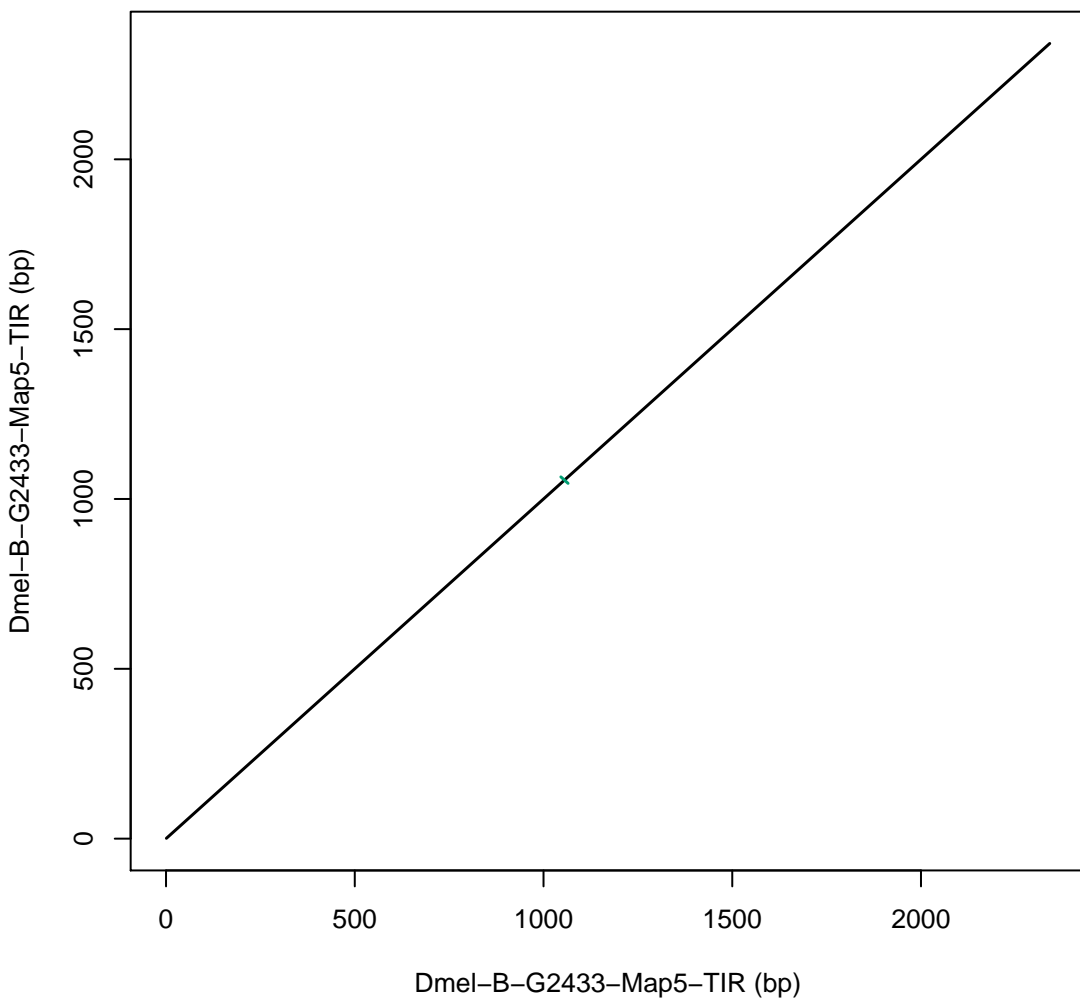# TE: Dmel−B−G2433−Map5−TIR
consensus size: 2341bp; fragments: 49; full length: 0 (>=2106.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected