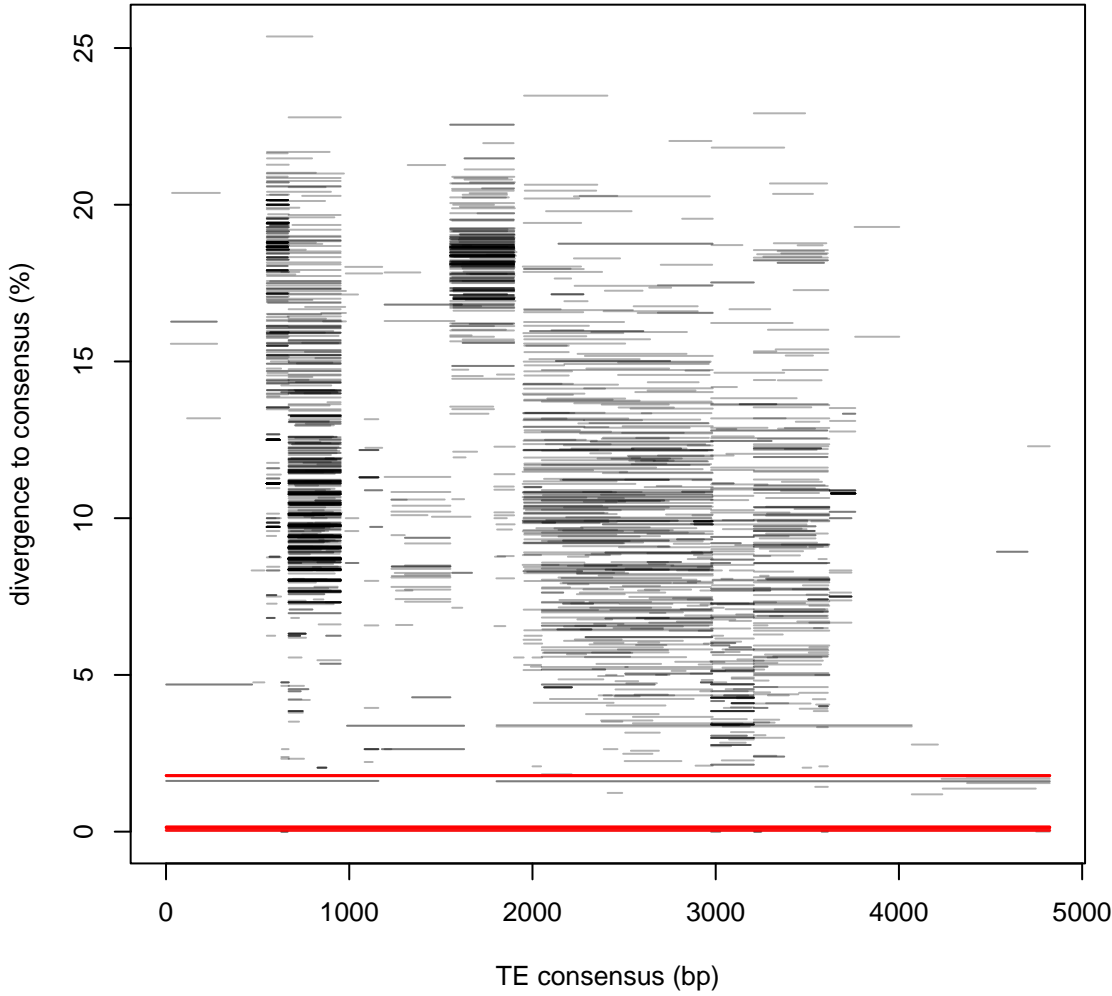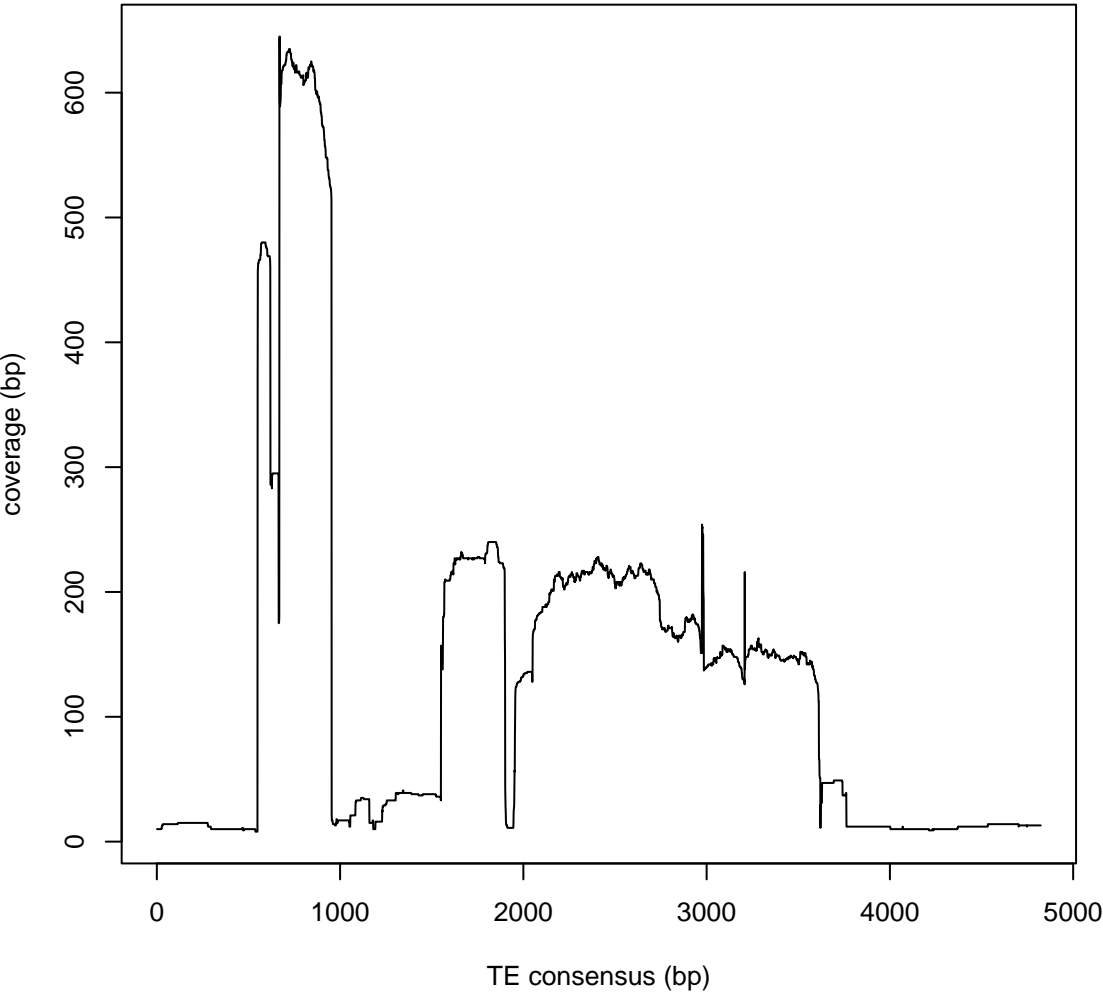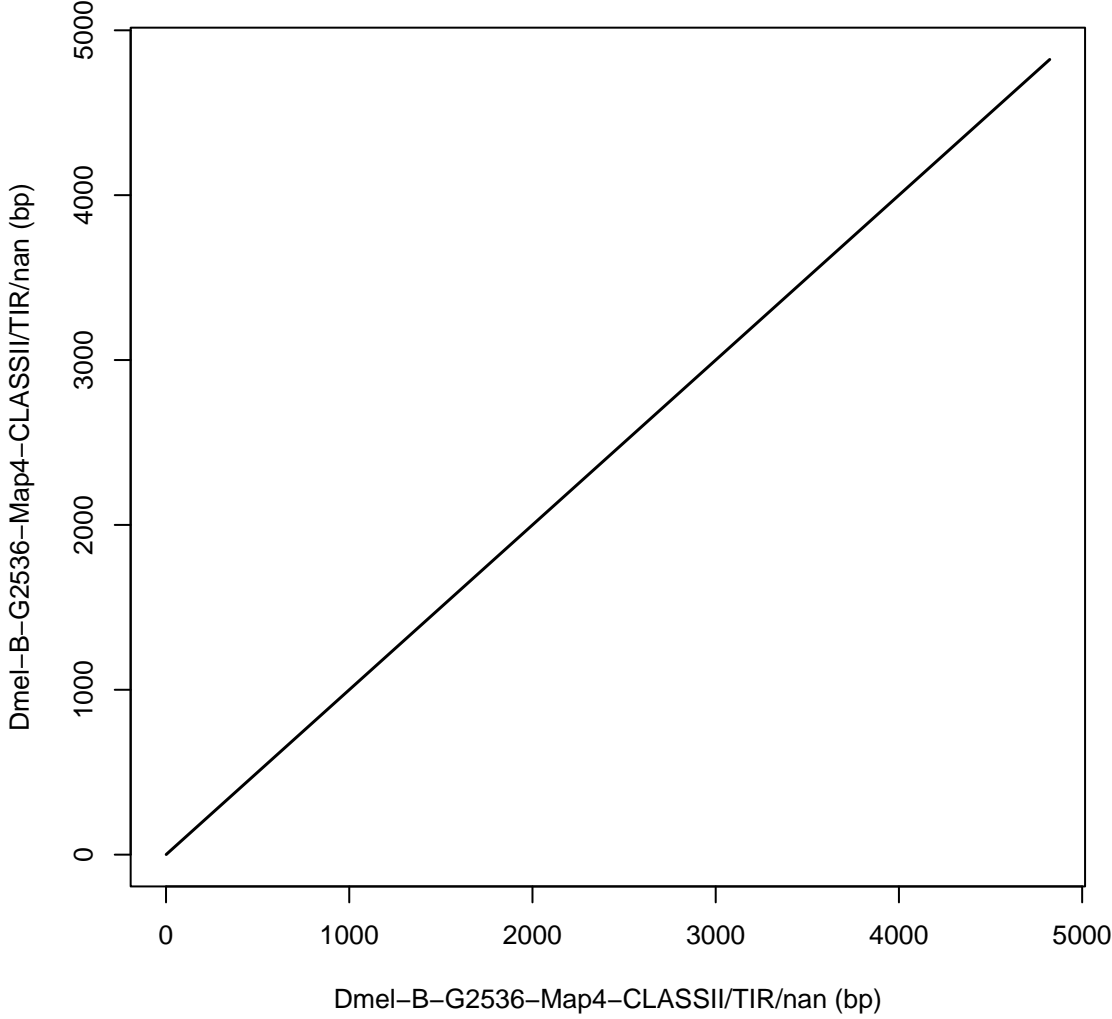**TE: Dmel−B−G2536−Map4−CLASSII/TIR/nan**
consensus size: 4823bp; fragments: 2272; full length: 6 (>=4340.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

CR1−8_AMi_pol LINE/CR1