TE: Dmel−B−P15_s_1_s_1−CLASSI/LTR/TRIM
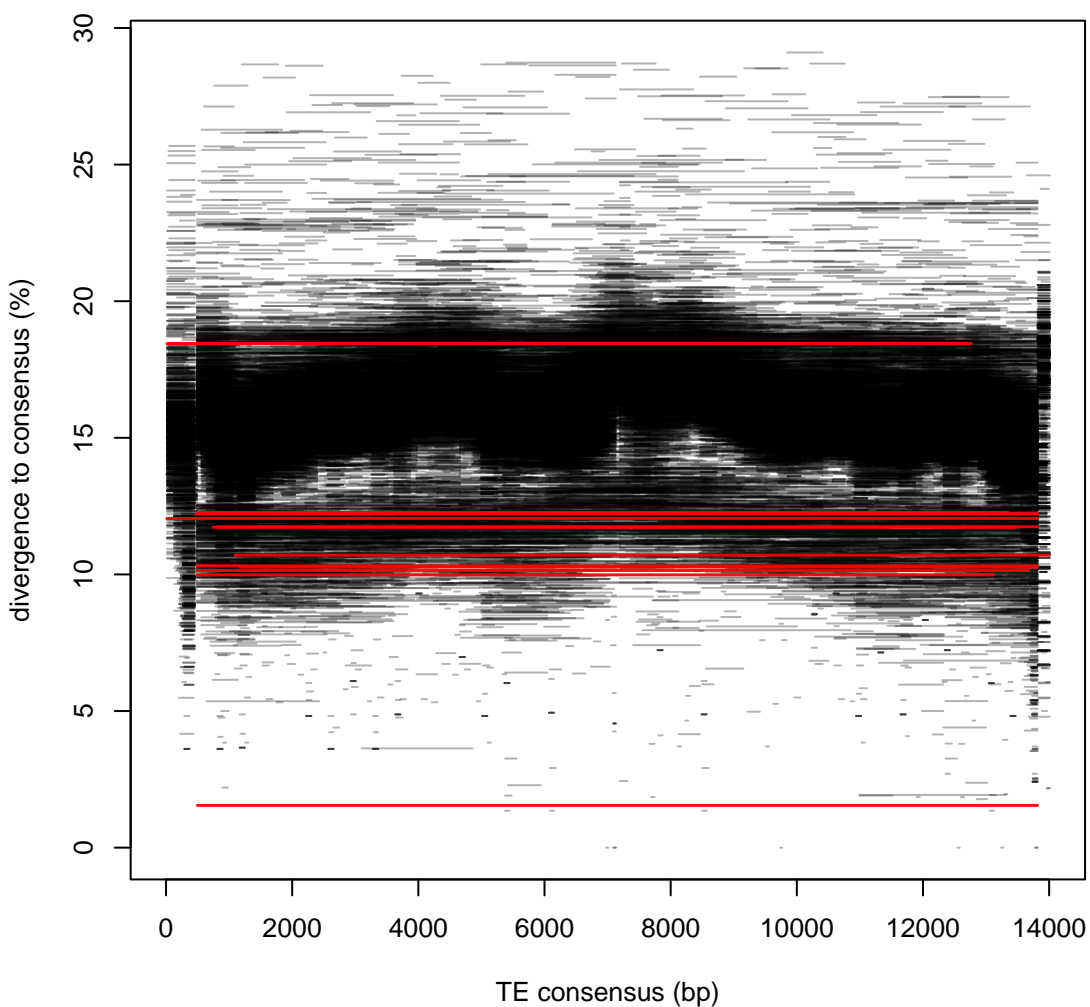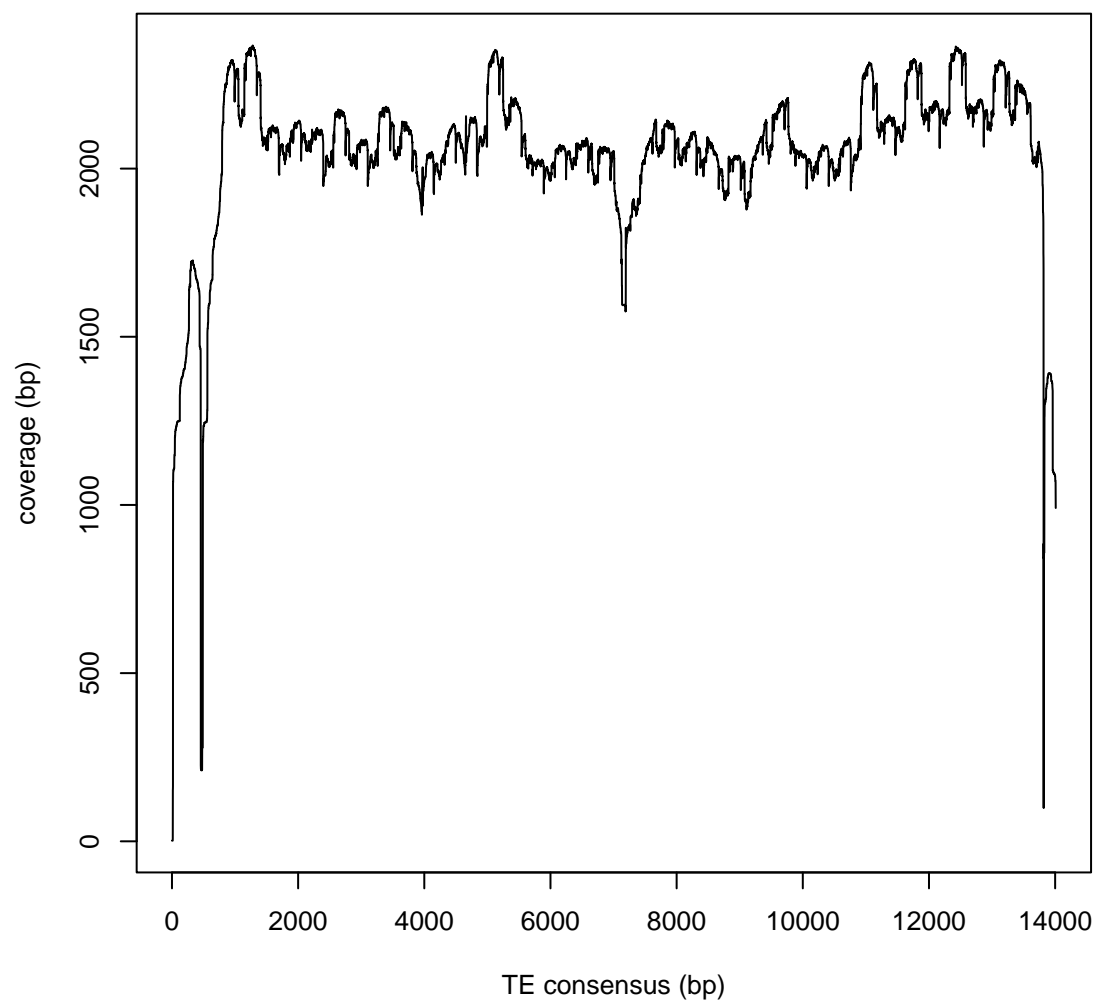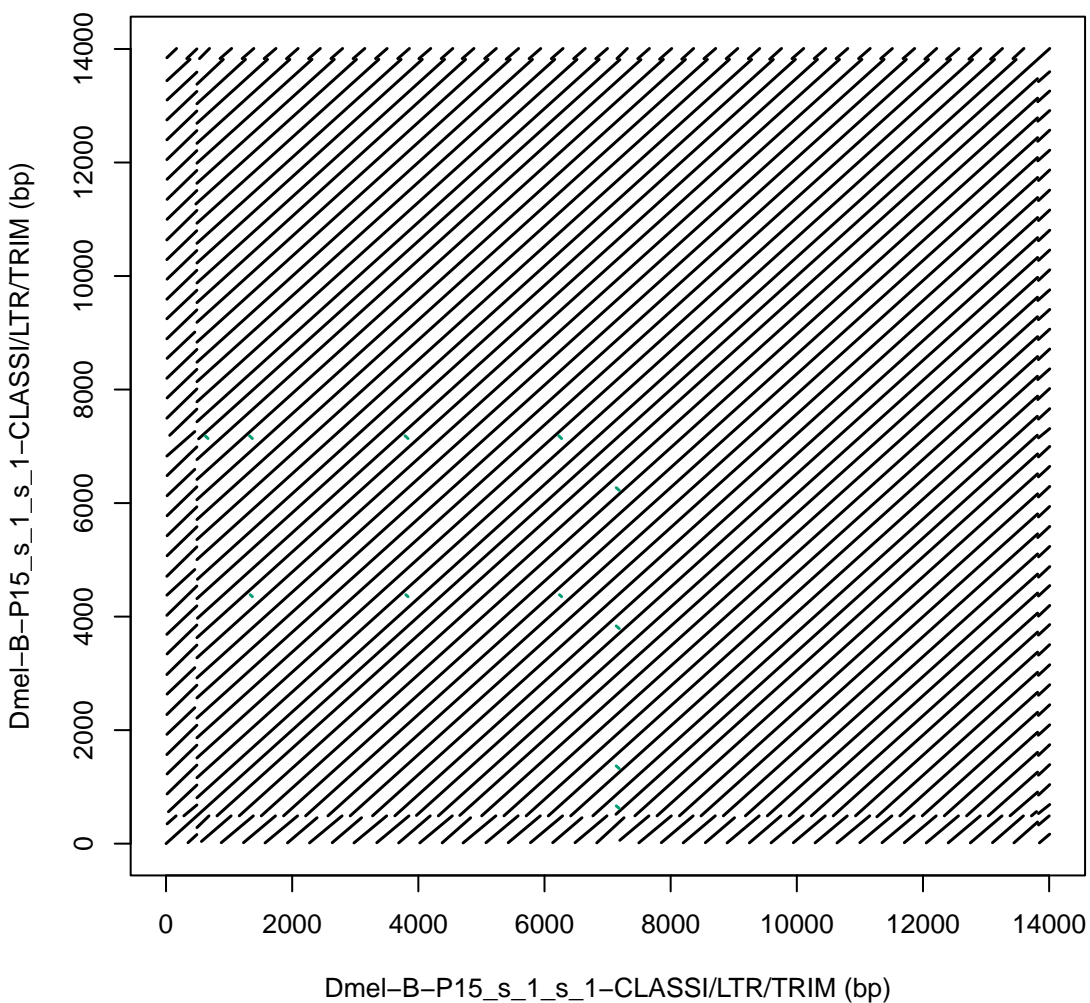consensus size: 14009bp; fragments: 28877; full length: 13 (>=12608.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no off >400 bp detected