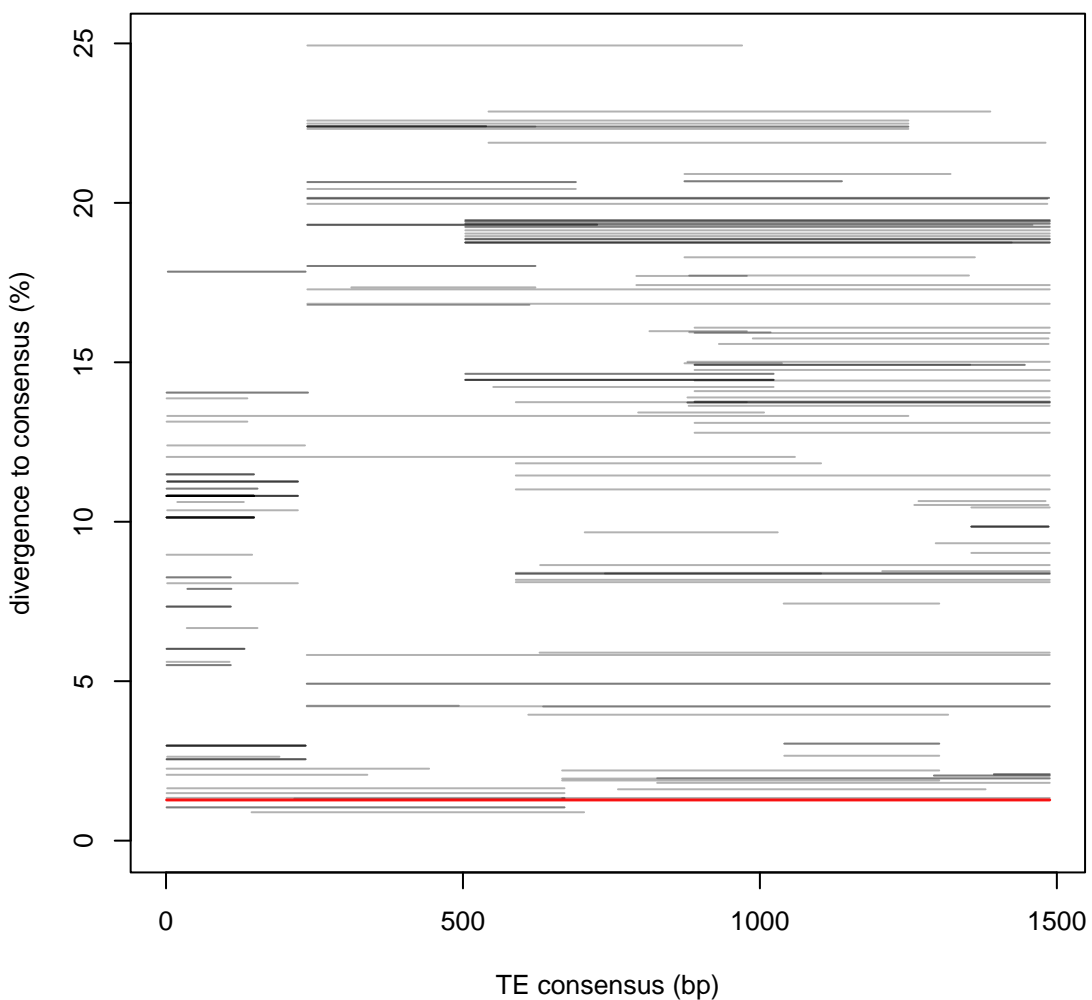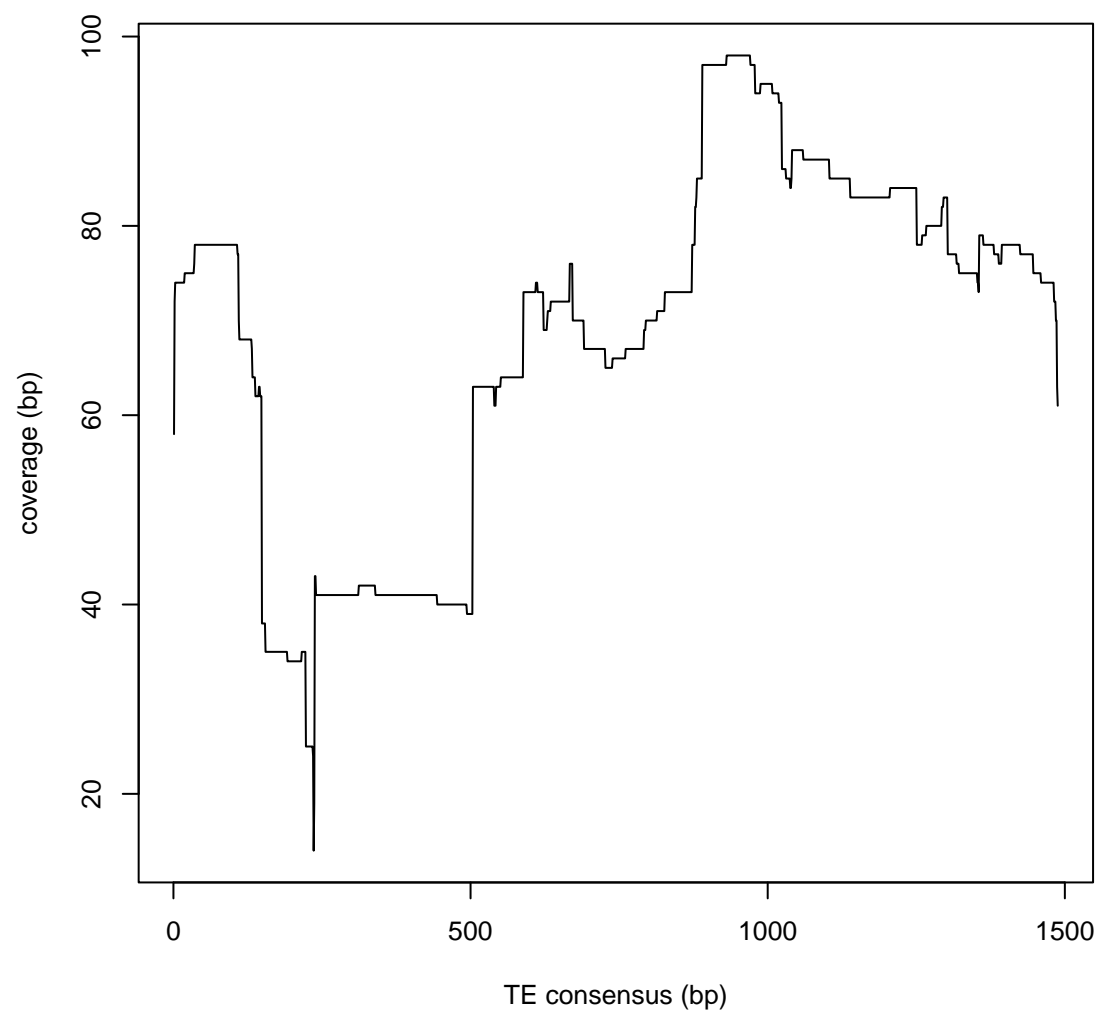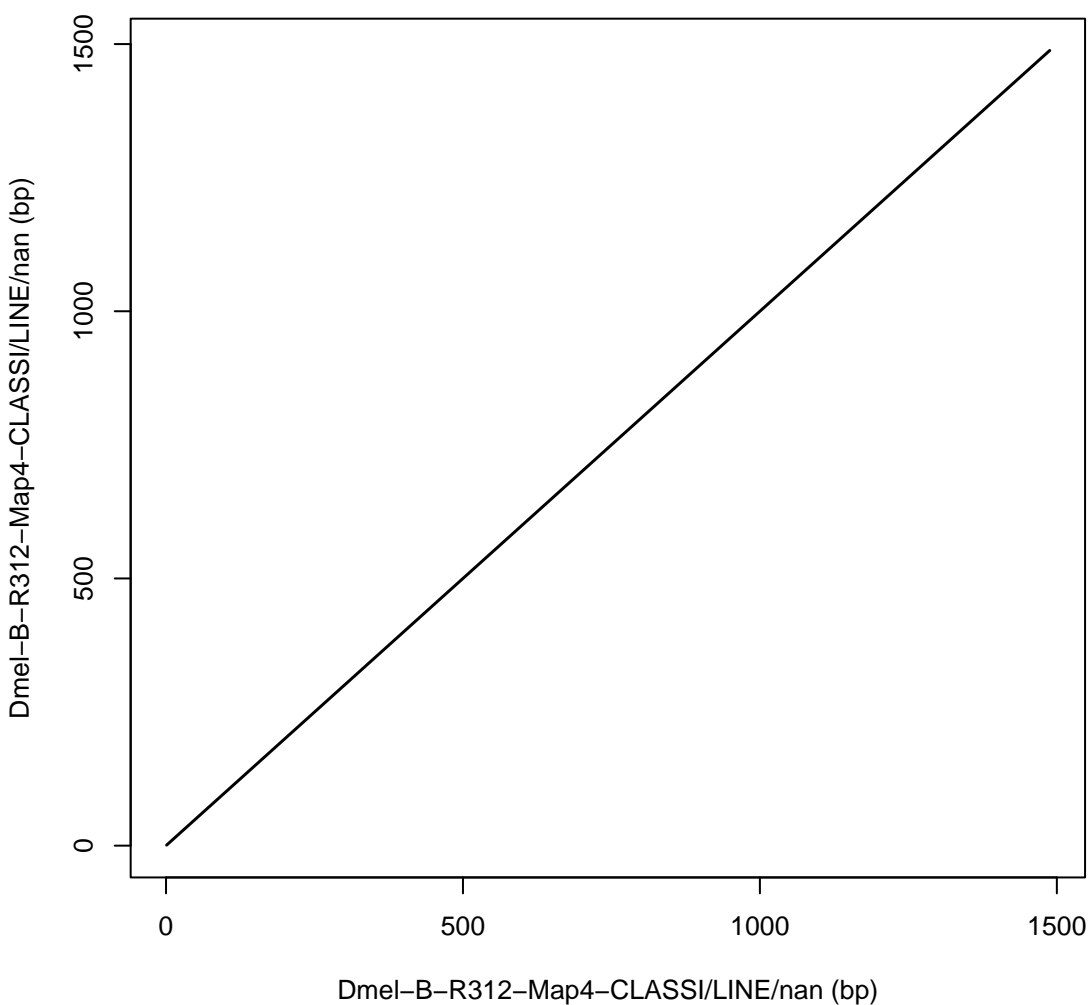TE: Dmel−B−R312−Map4−CLASSI/LINE/nan
consensus size: 1488bp; fragments: 207; full length: 1 (>=1339.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected