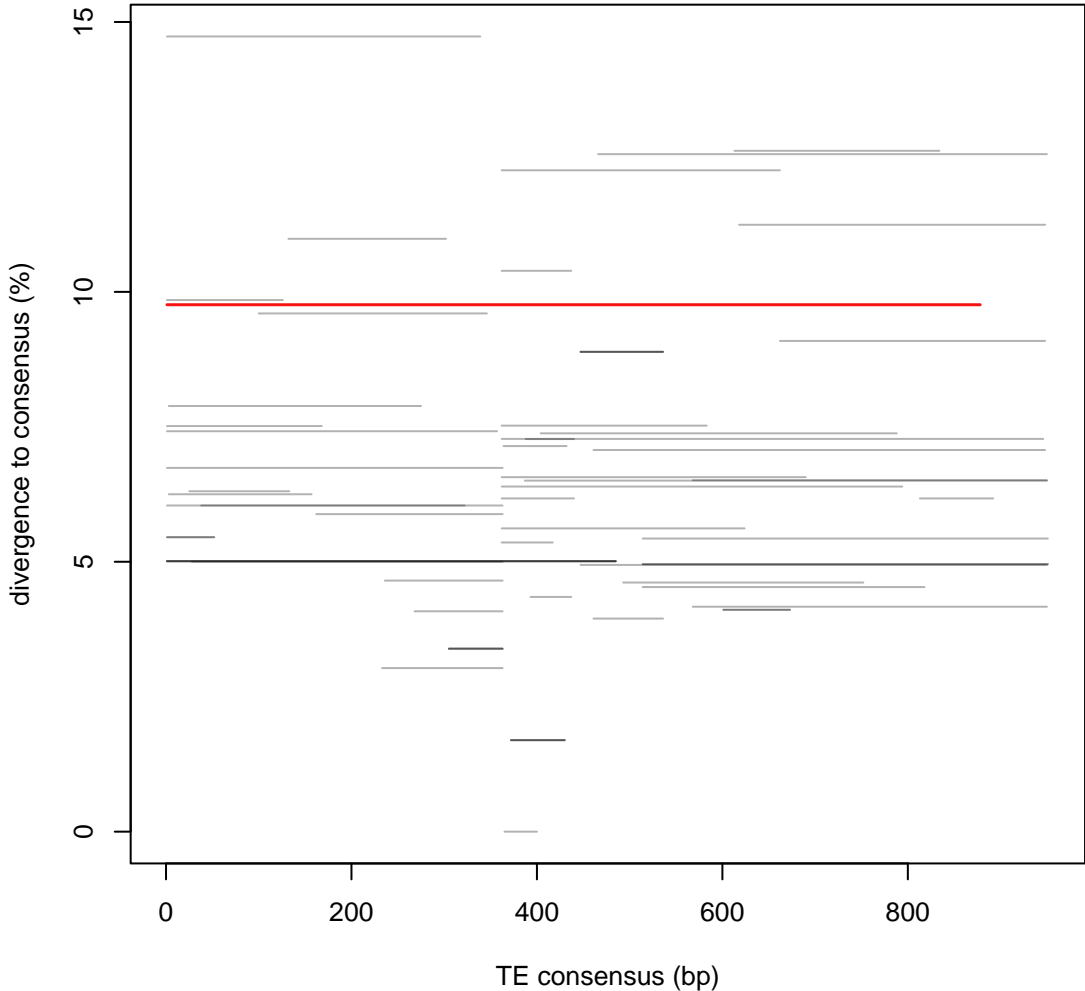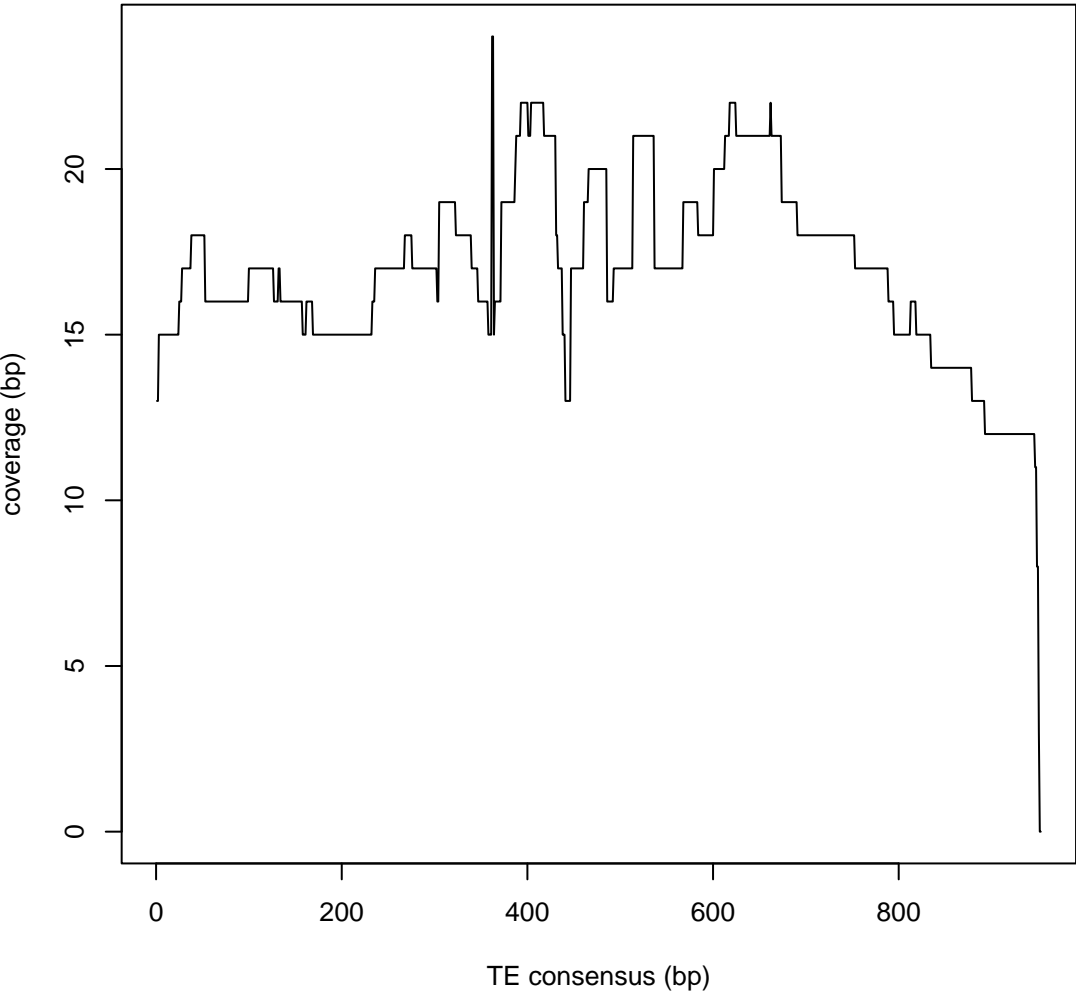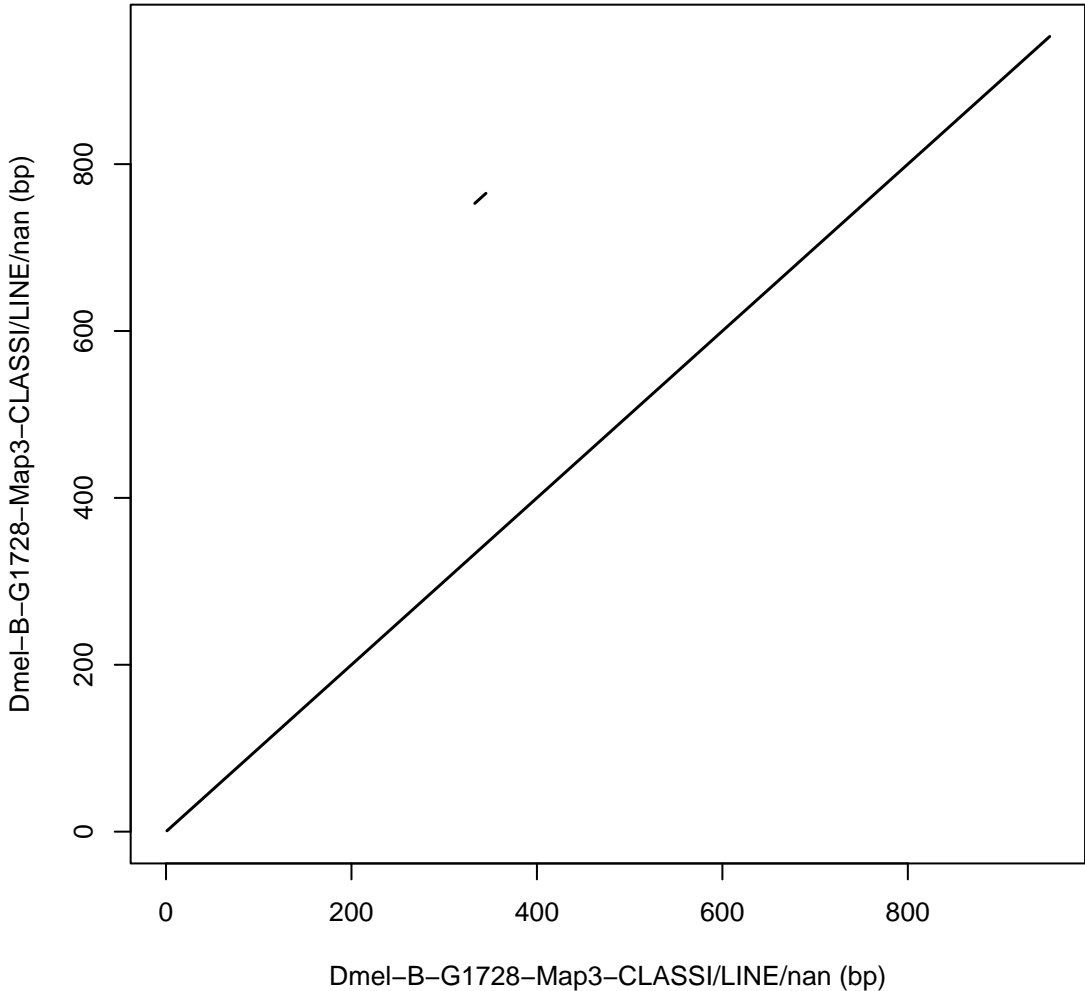## TE: Dmel−B−G1728−Map3−CLASSI/LINE/nan
consensus size: 953bp; fragments: 65; full length: 1 (>=857.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected