TE: Dmel–B–R1189–Map3–Unclassified/nan
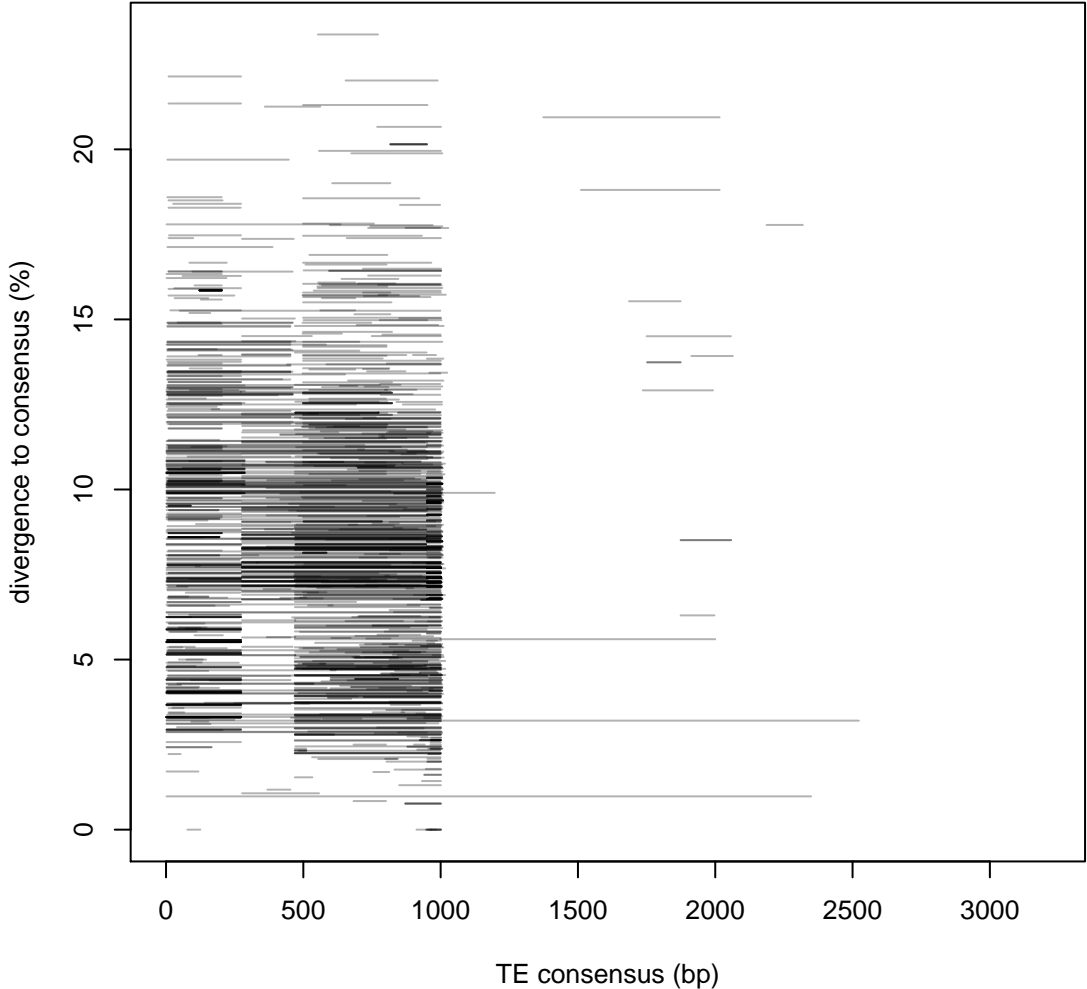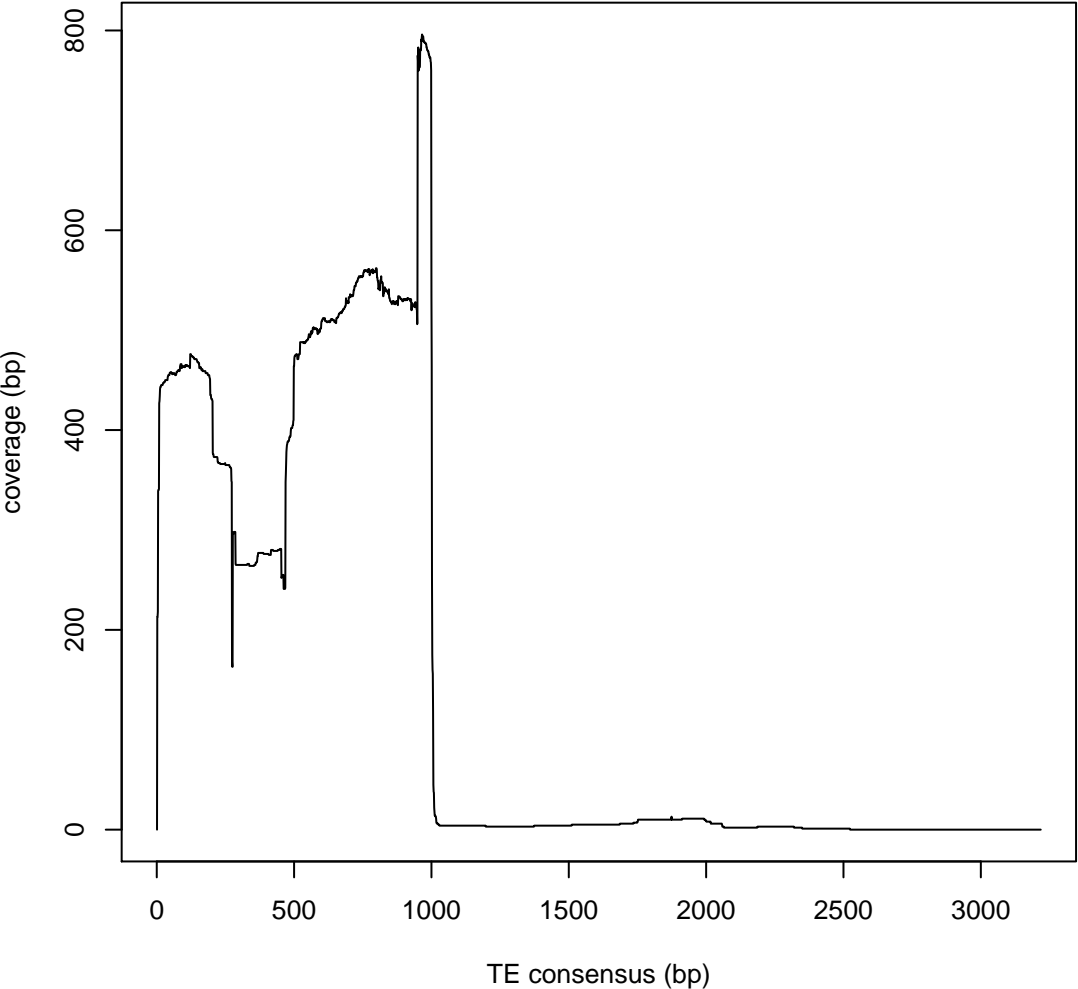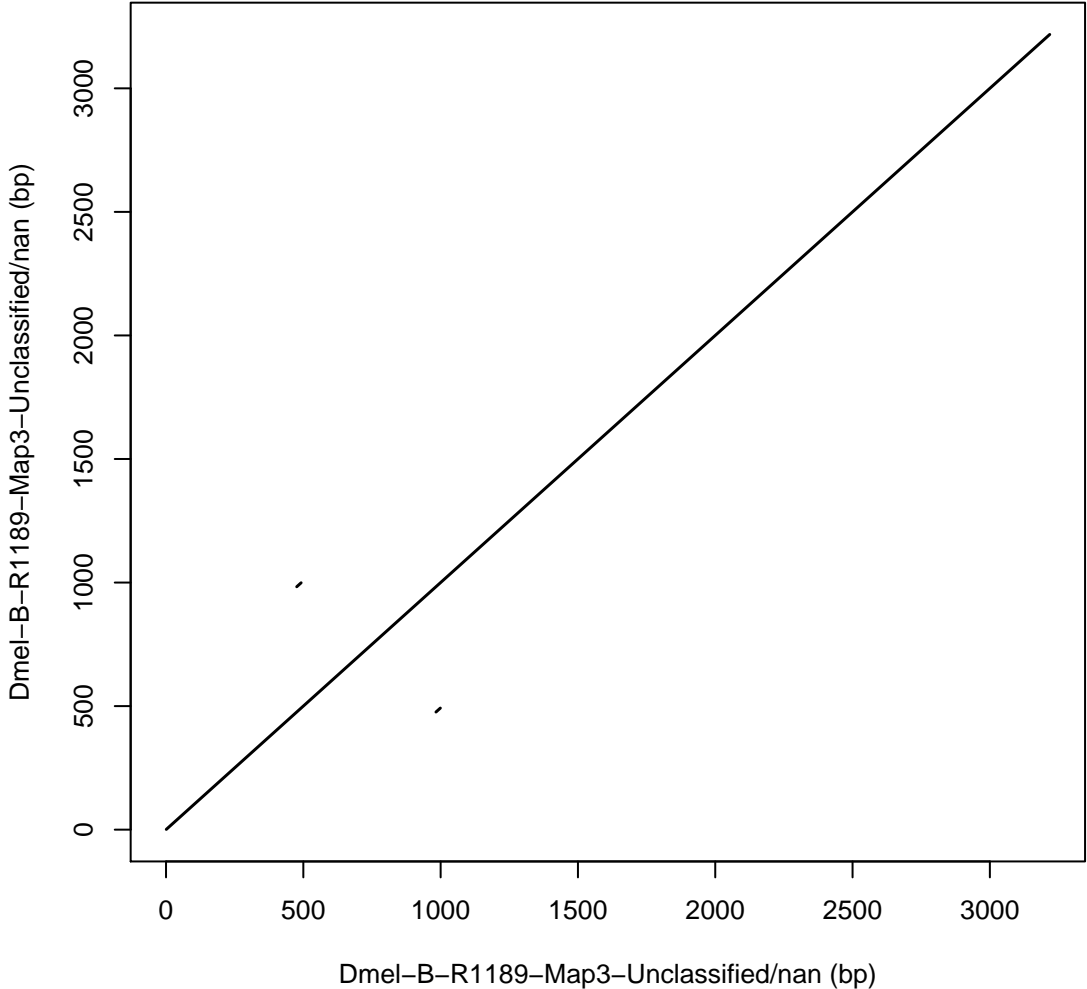### consensus size: 3218bp; fragments: 1532; full length: 0 (>=2896.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected