TE: Dmel−L−B489−Map1−TRIM/nan
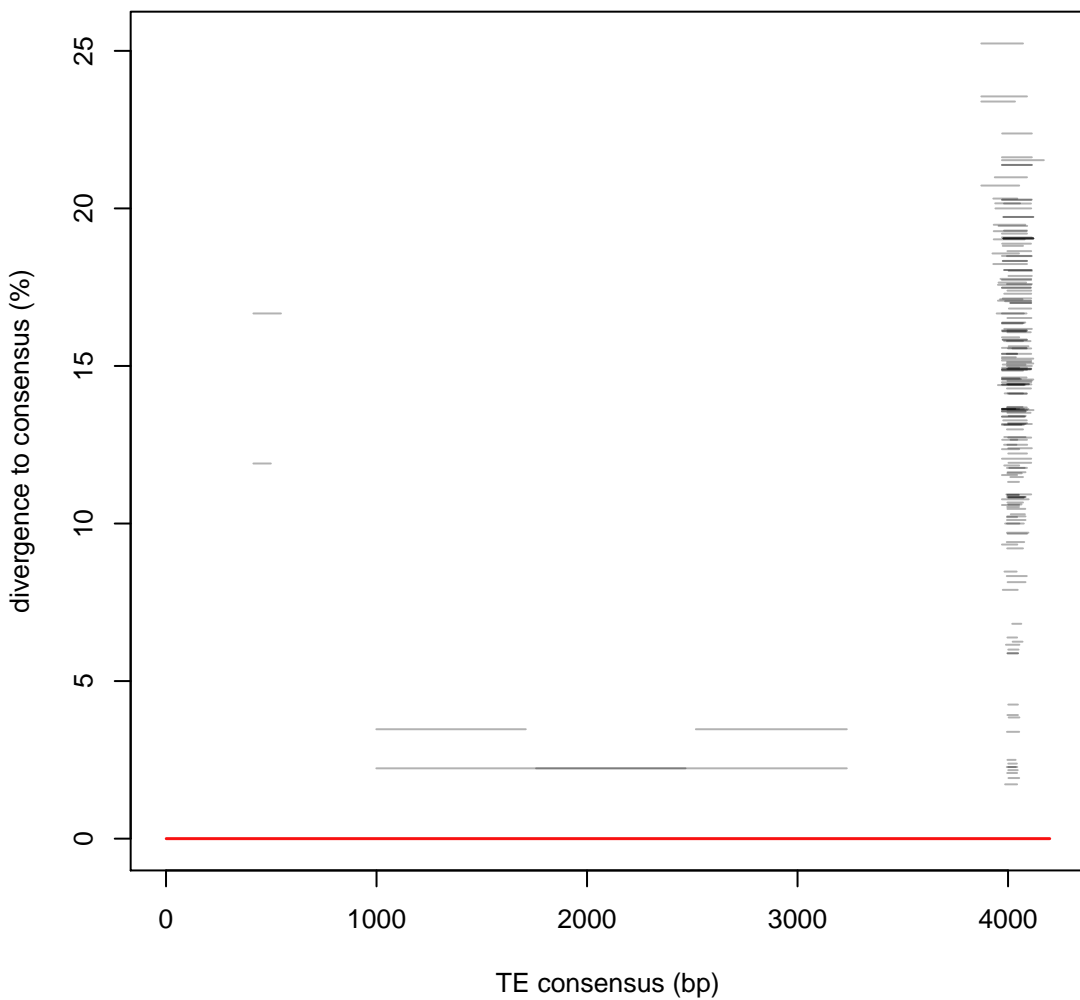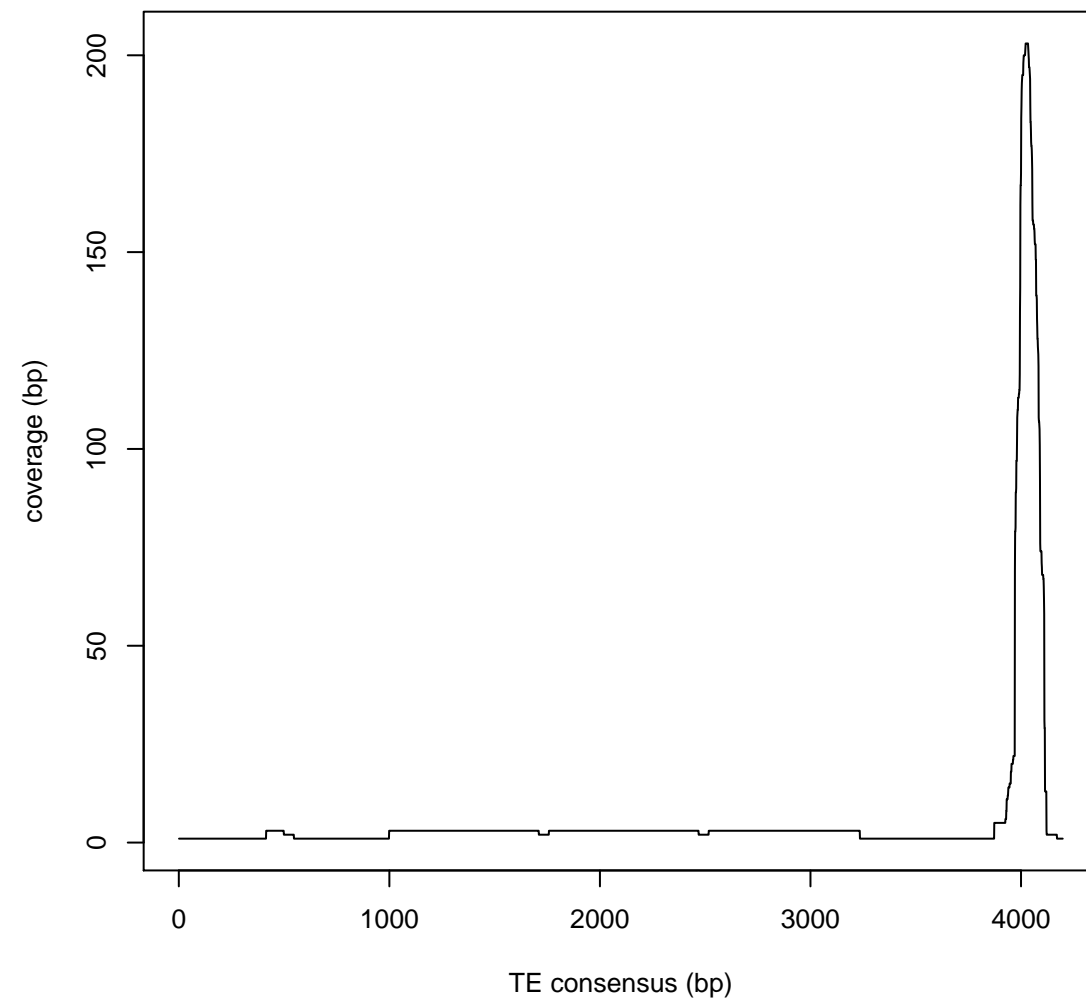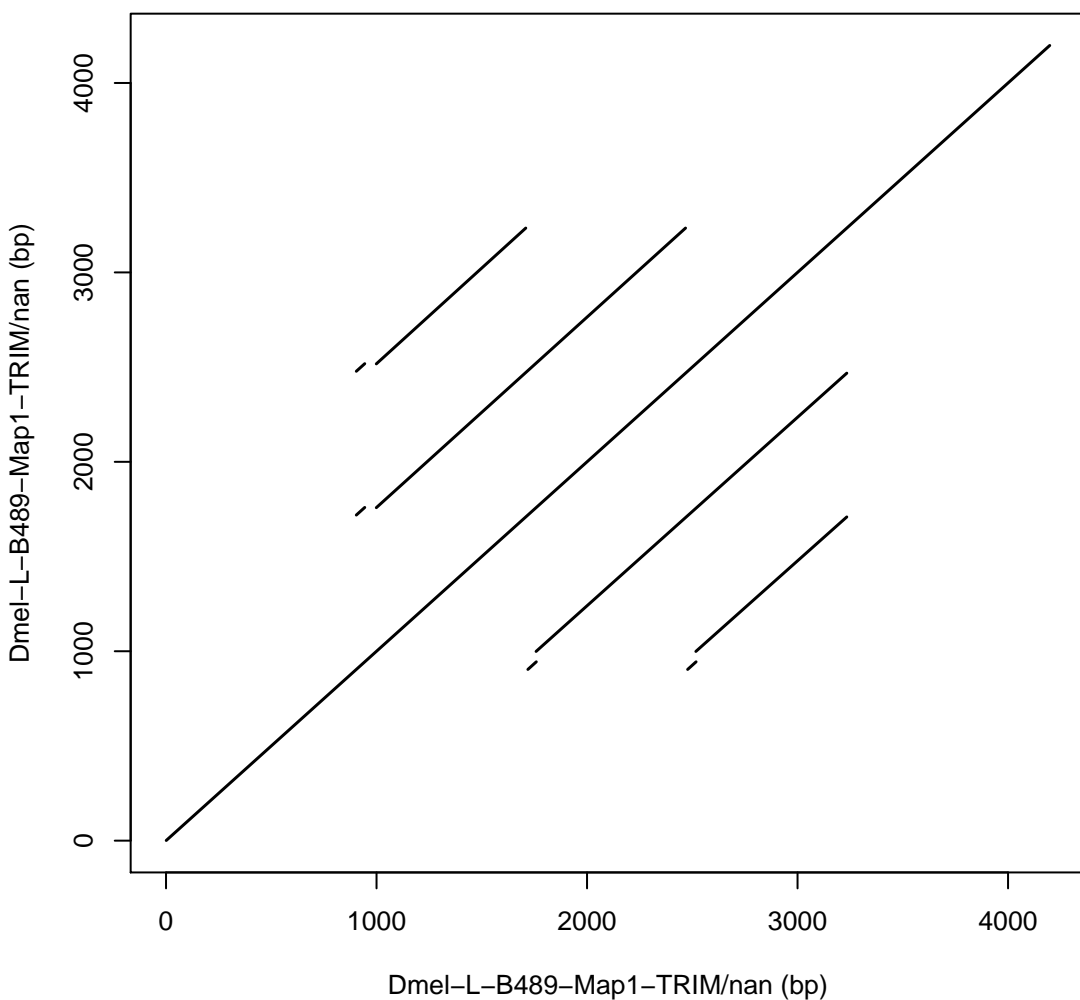### consensus size: 4198bp; fragments: 209; full length: 1 (>=3778.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected