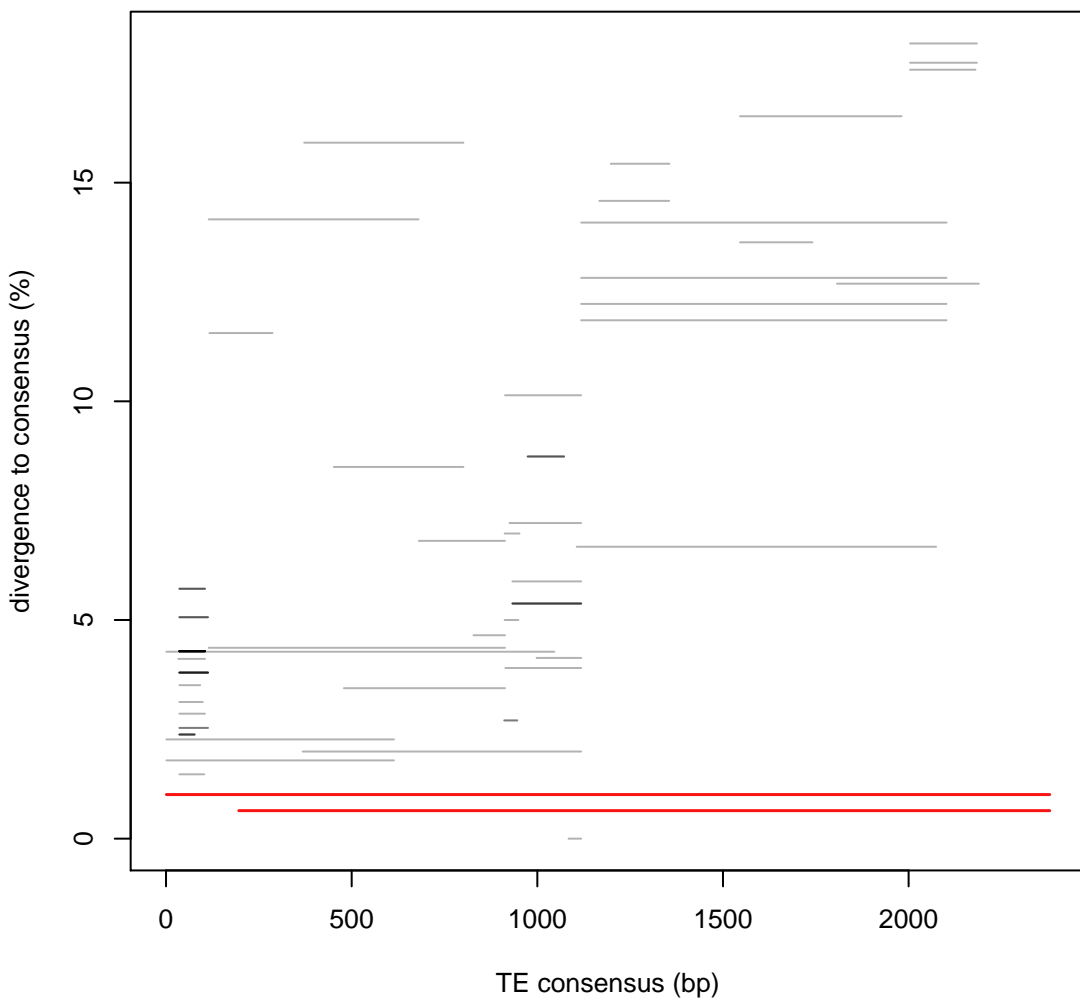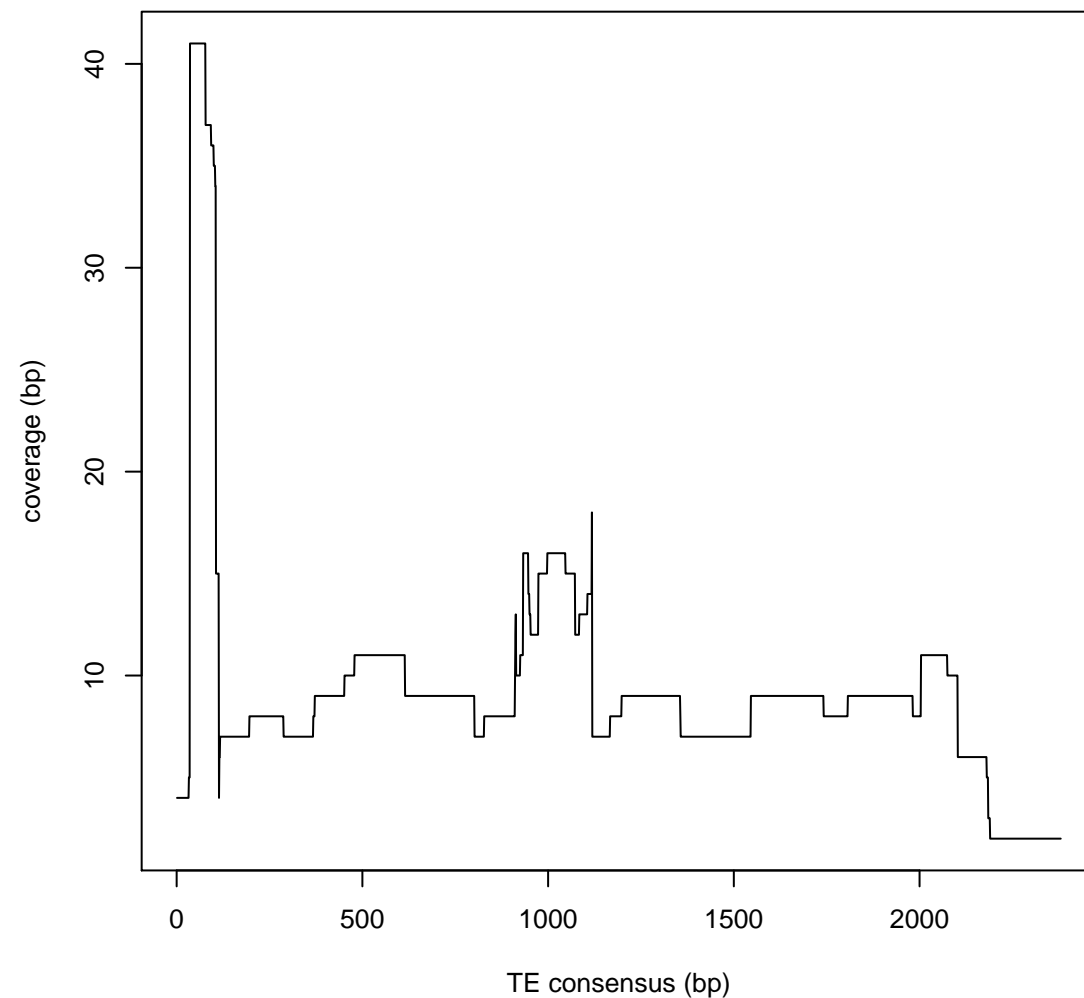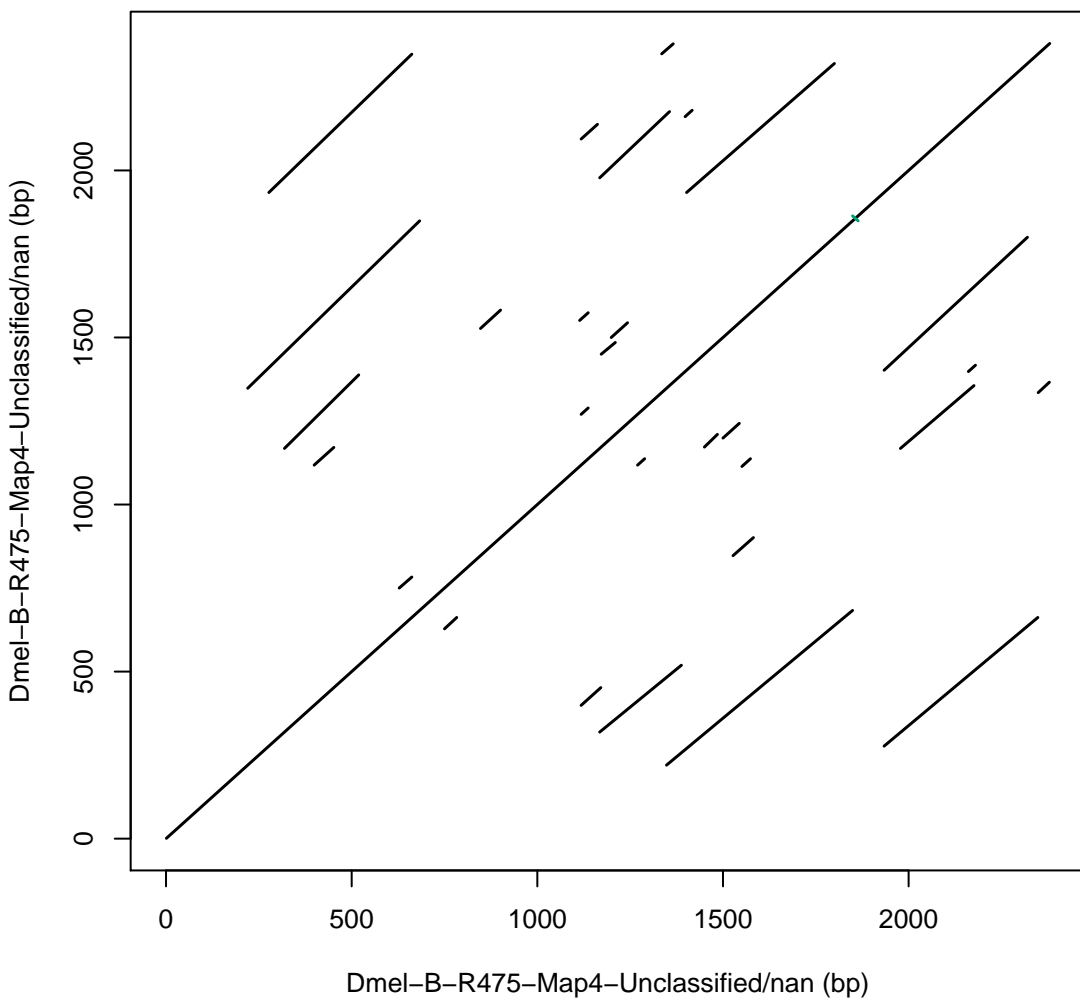**TE: Dmel-B-R475-Map4-Unclassified/nan**
consensus size: 2380bp; fragments: 81; full length: 2 (>=2142bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected