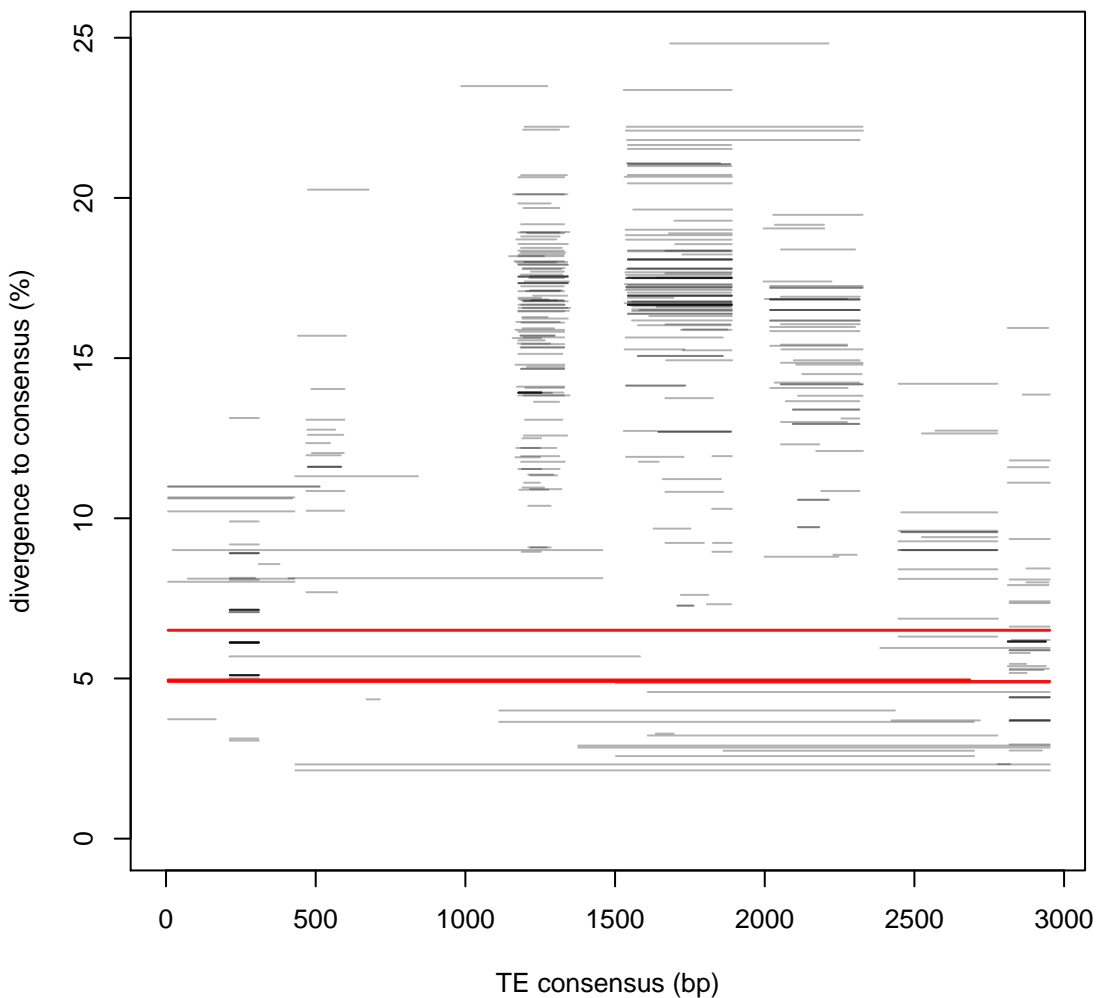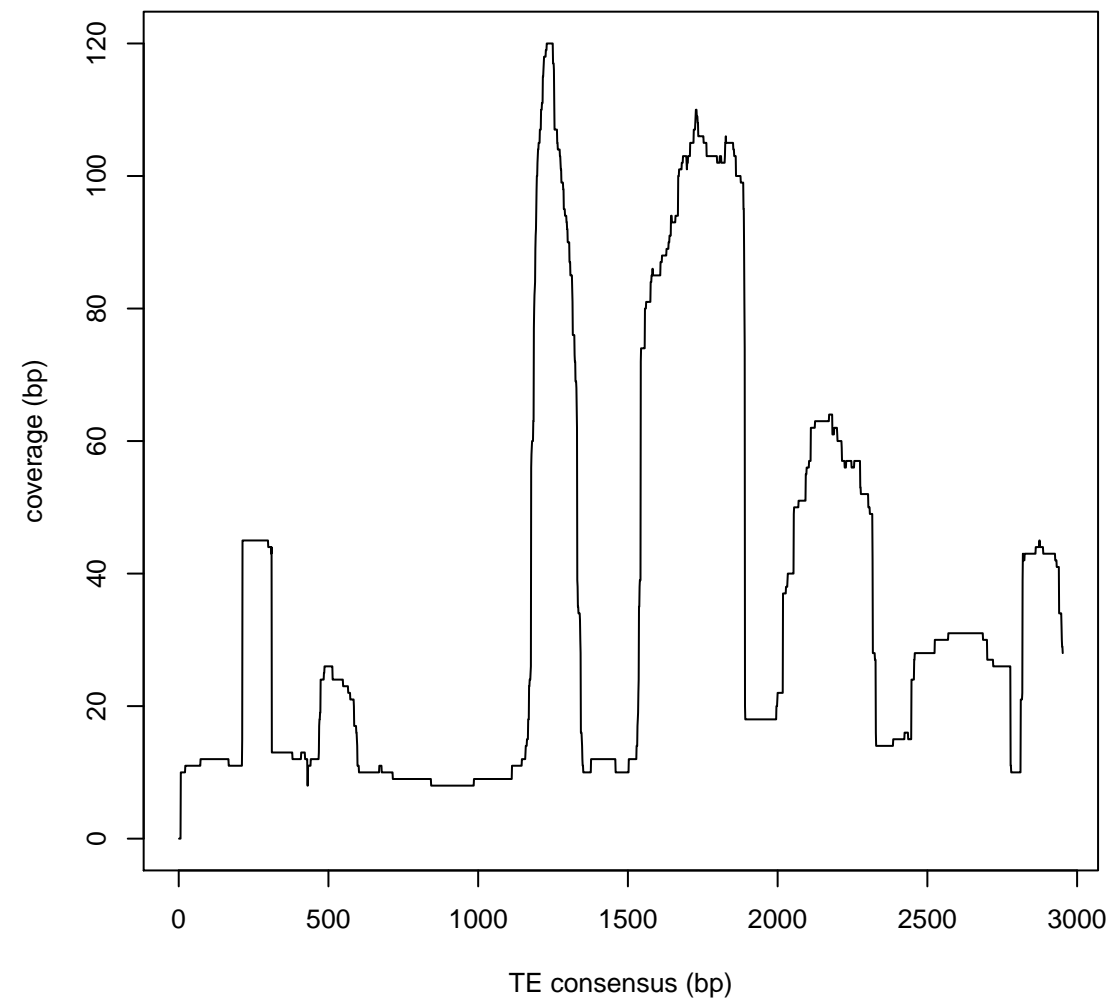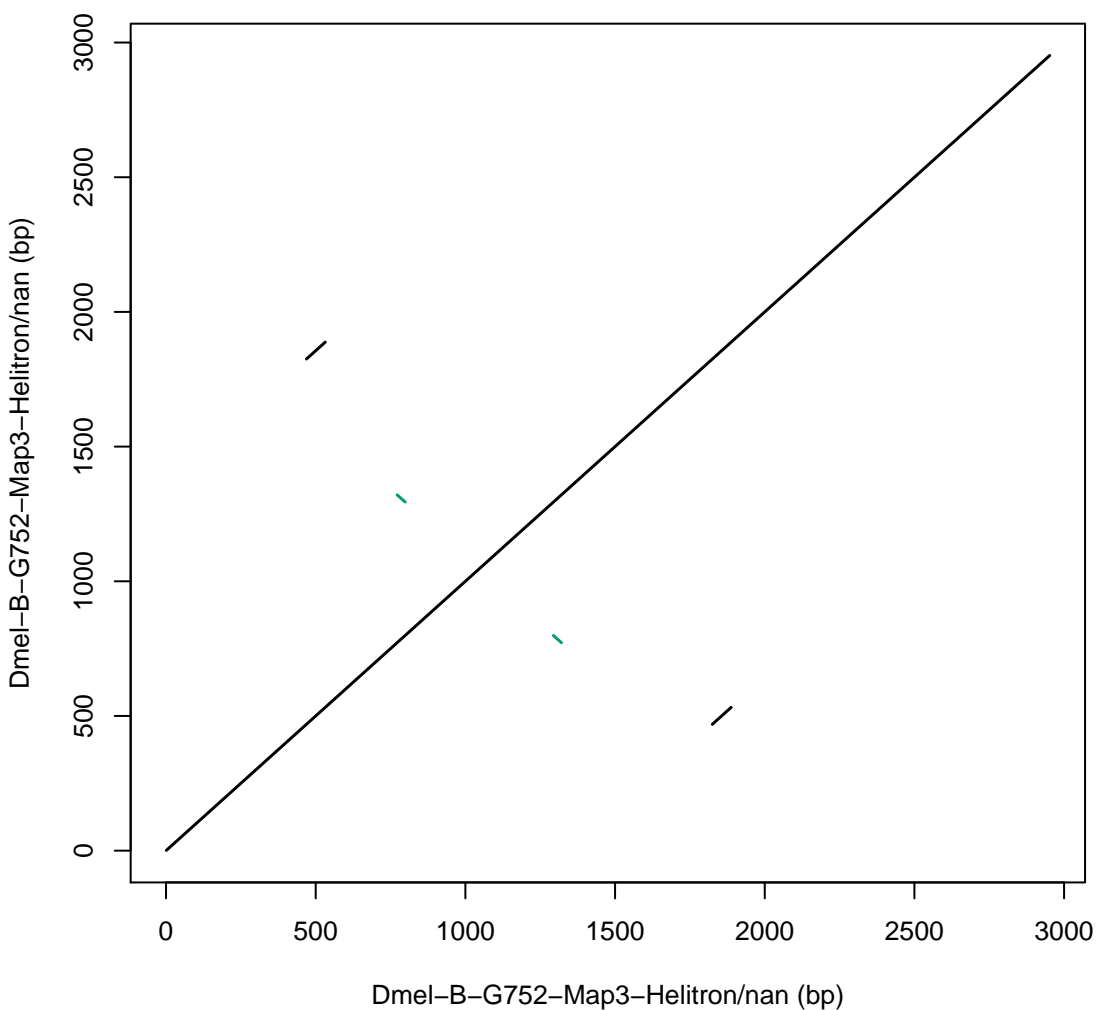**TE: Dmel-B-G752-Map3-Helitron/nan**
consensus size: 2952bp; fragments: 397; full length: 3 (>=2656.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected