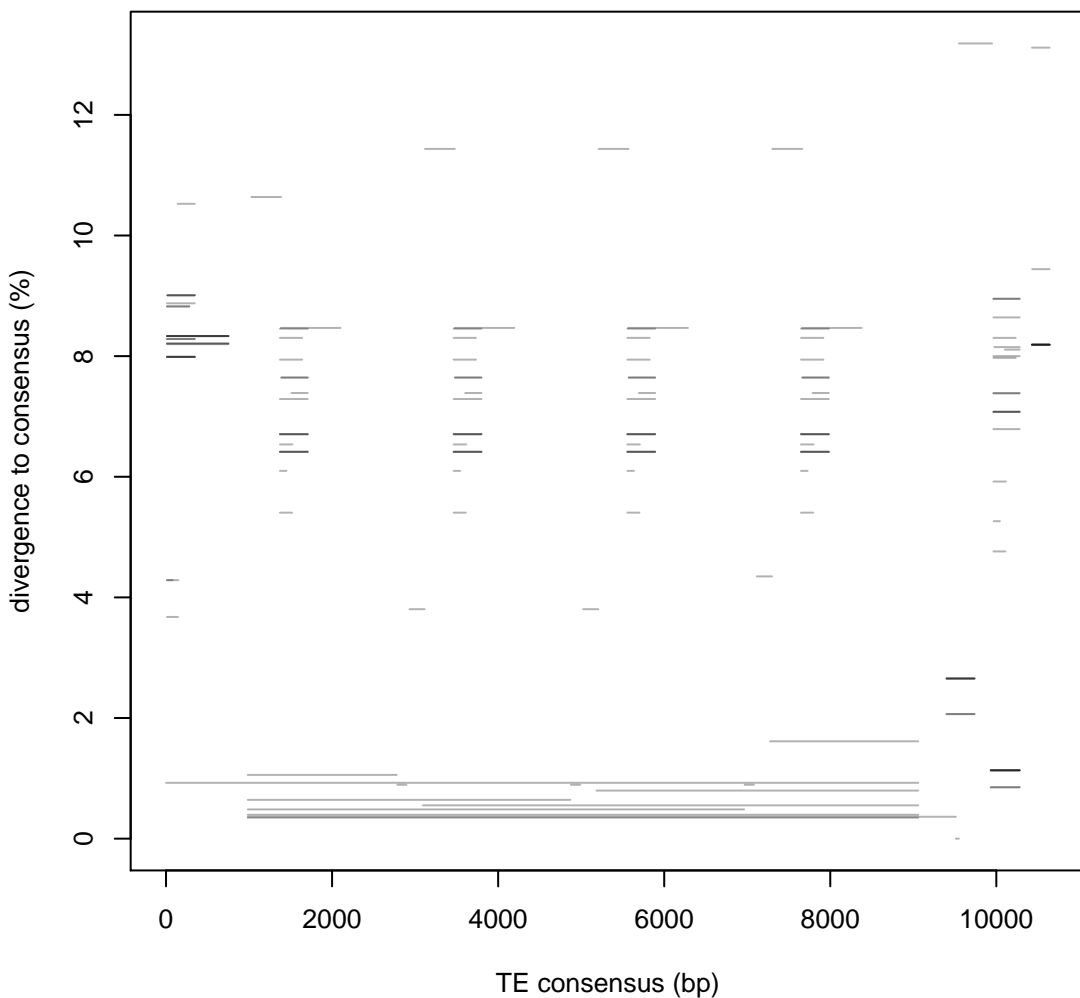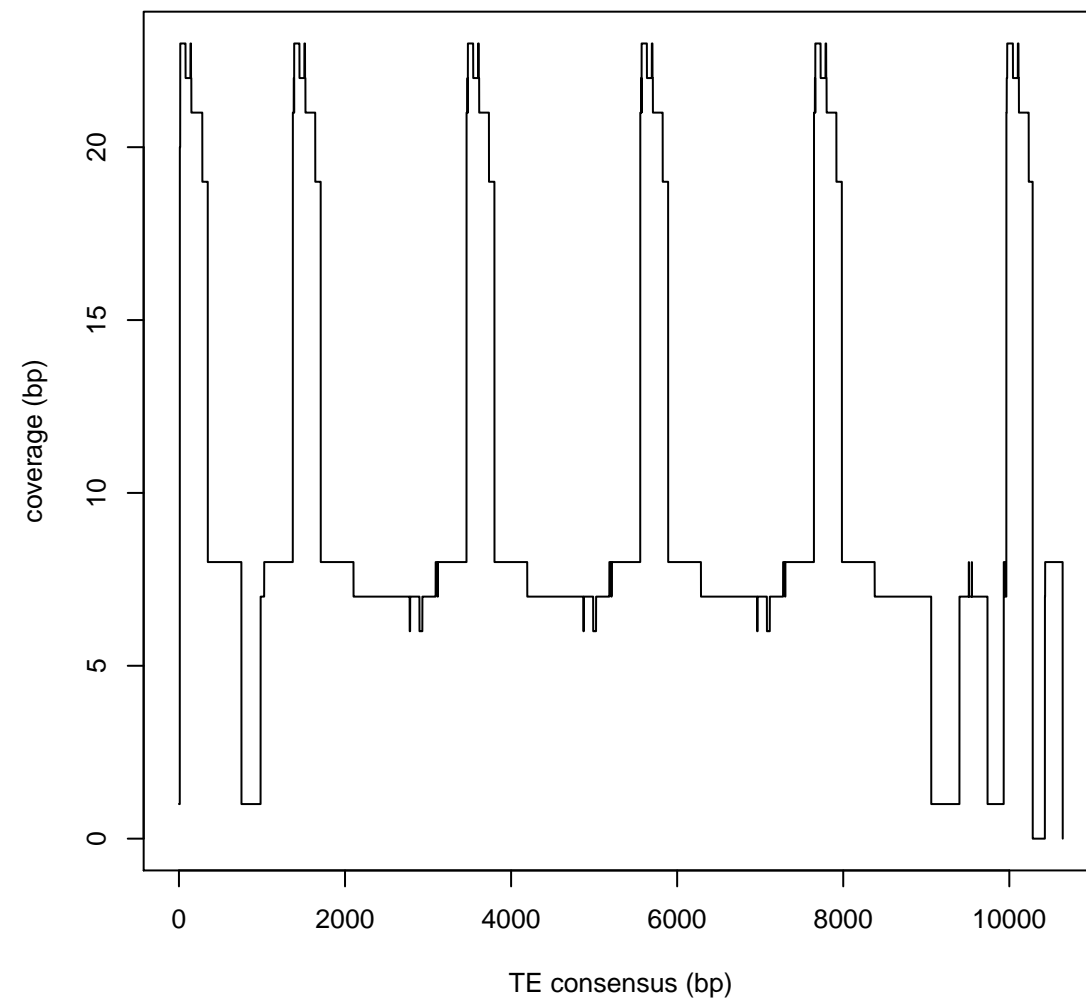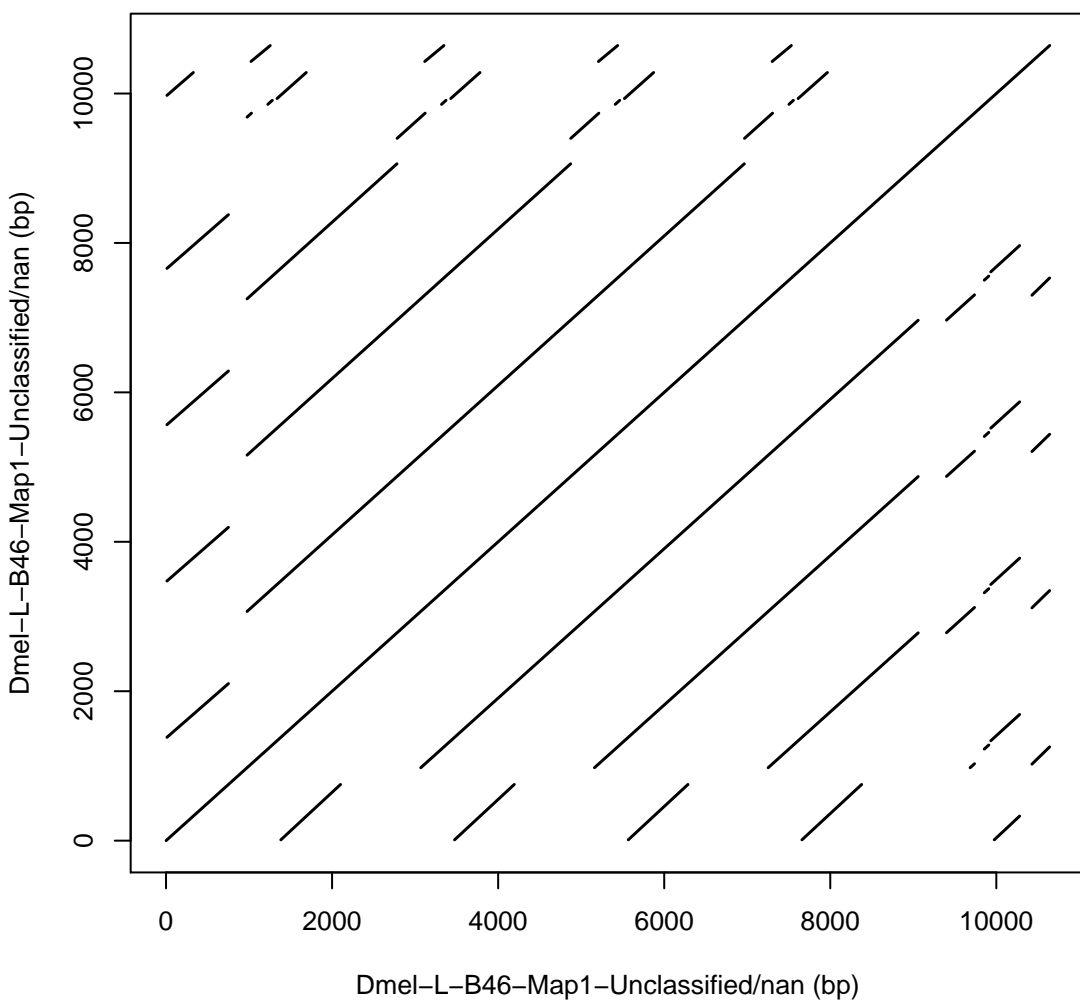**TE: Dmel−L−B46−Map1−Unclassified/nan**
**consensus size: 10643bp; fragments: 151; full length: 0 (>=9578.7bp)**

TE consensus (bp)

divergence to consensus (%)

**TE consensus genomic coverage**

coverage (bp)

TE consensus (bp)

**TE consensus self dotplot (blastn)**

Dmel−L−B46−Map1−Unclassified/nan (bp)

Dmel−L−B46−Map1−Unclassified/nan (bp)

**TE consensus structure and protein hits**

Dmel−L−B46−Map1−Unclassified/nan (bp)