TE: Dmel−B−R1122−Map13−Helitron/nan
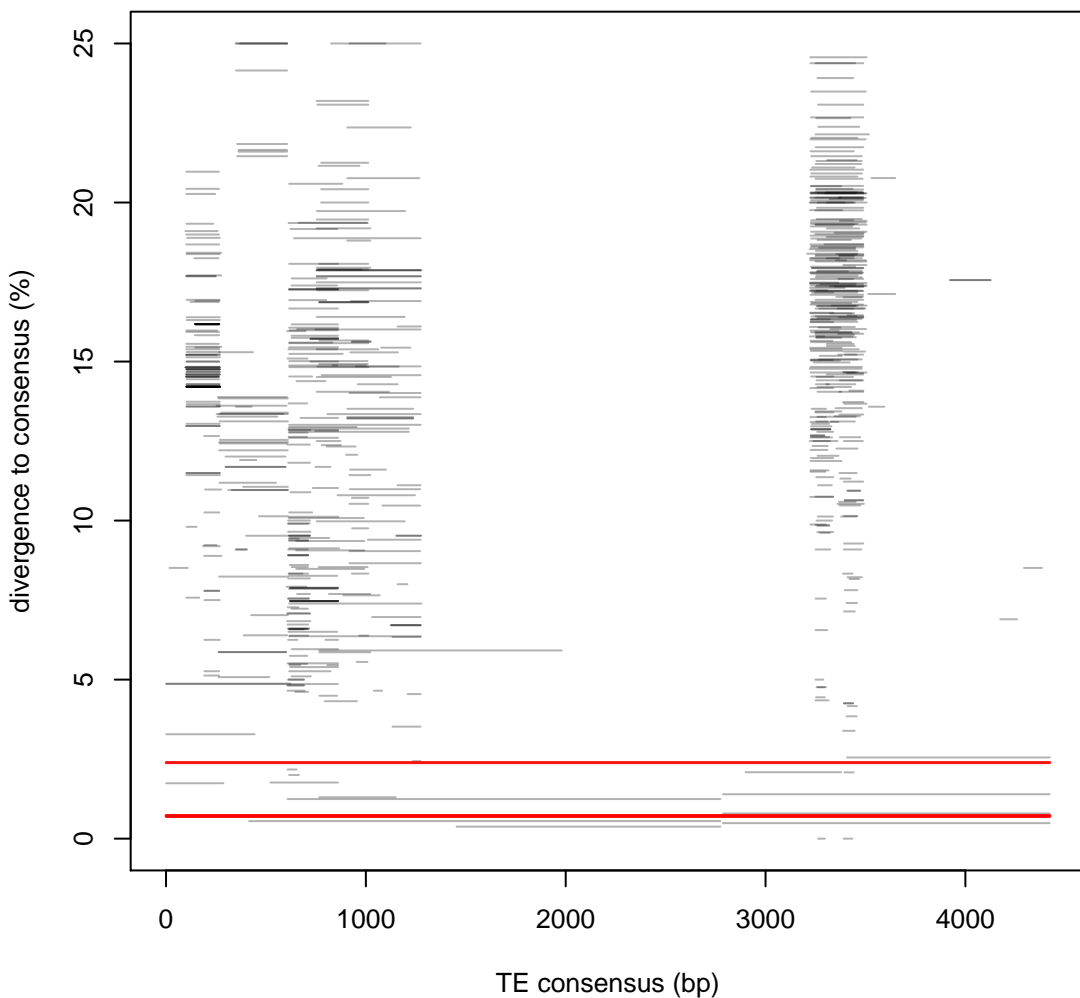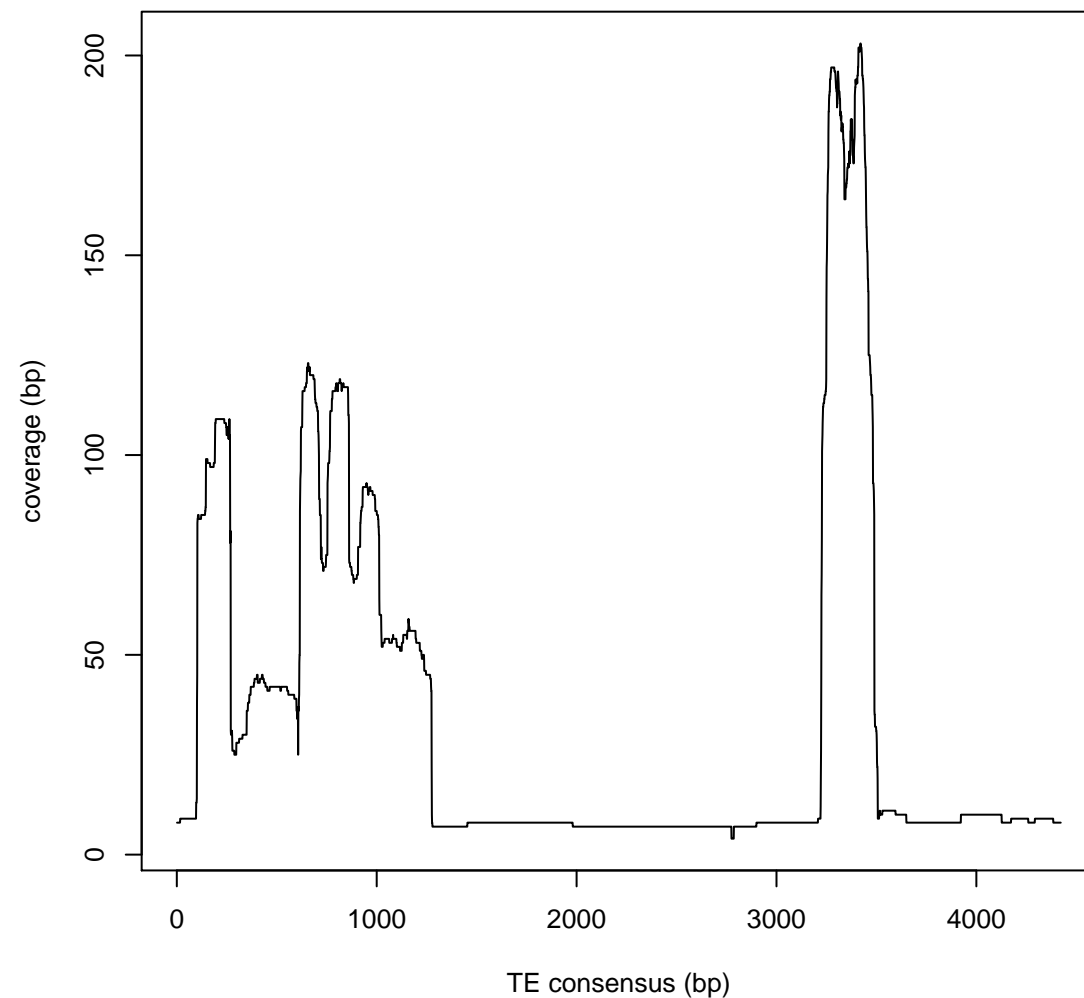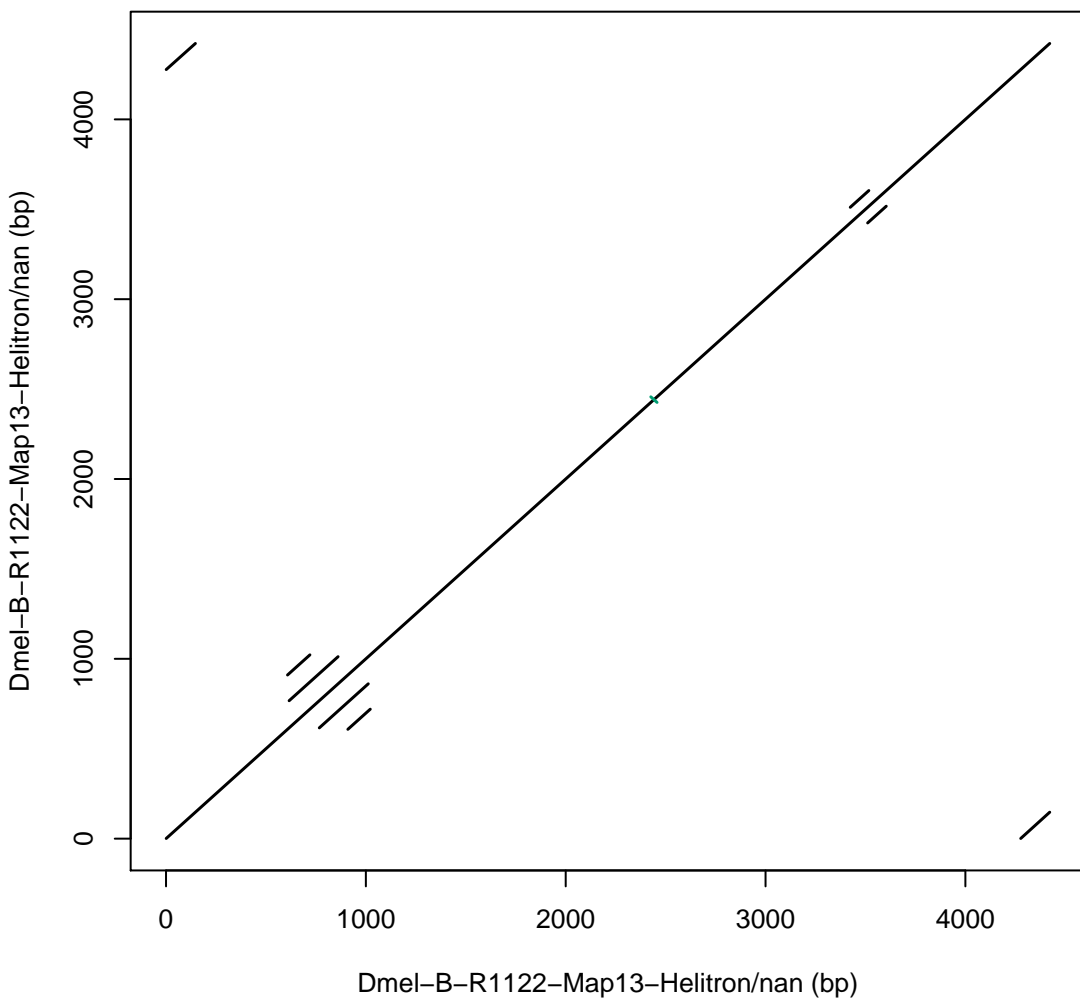consensus size: 4422bp; fragments: 659; full length: 4 (>=3979.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected