TE: Dmel−L−B431−Map1−LARD/nan
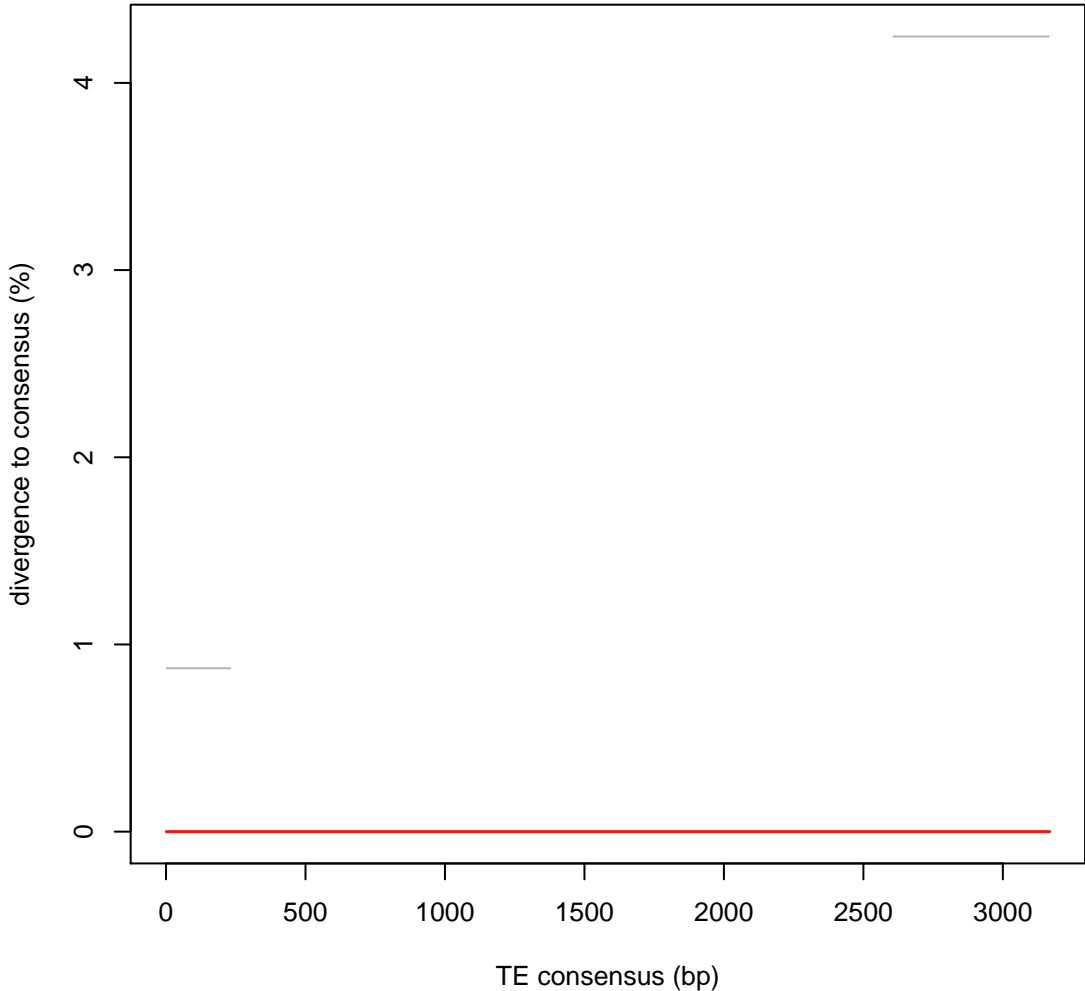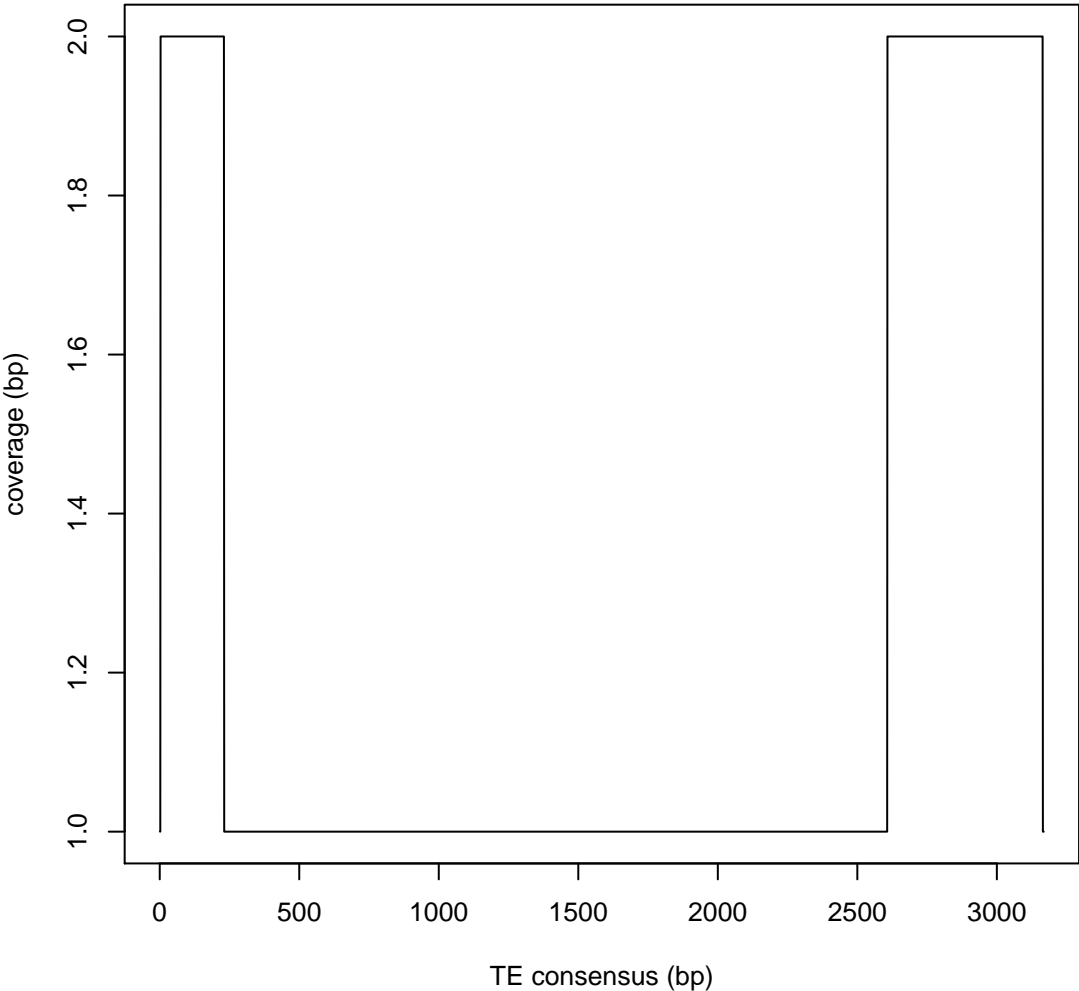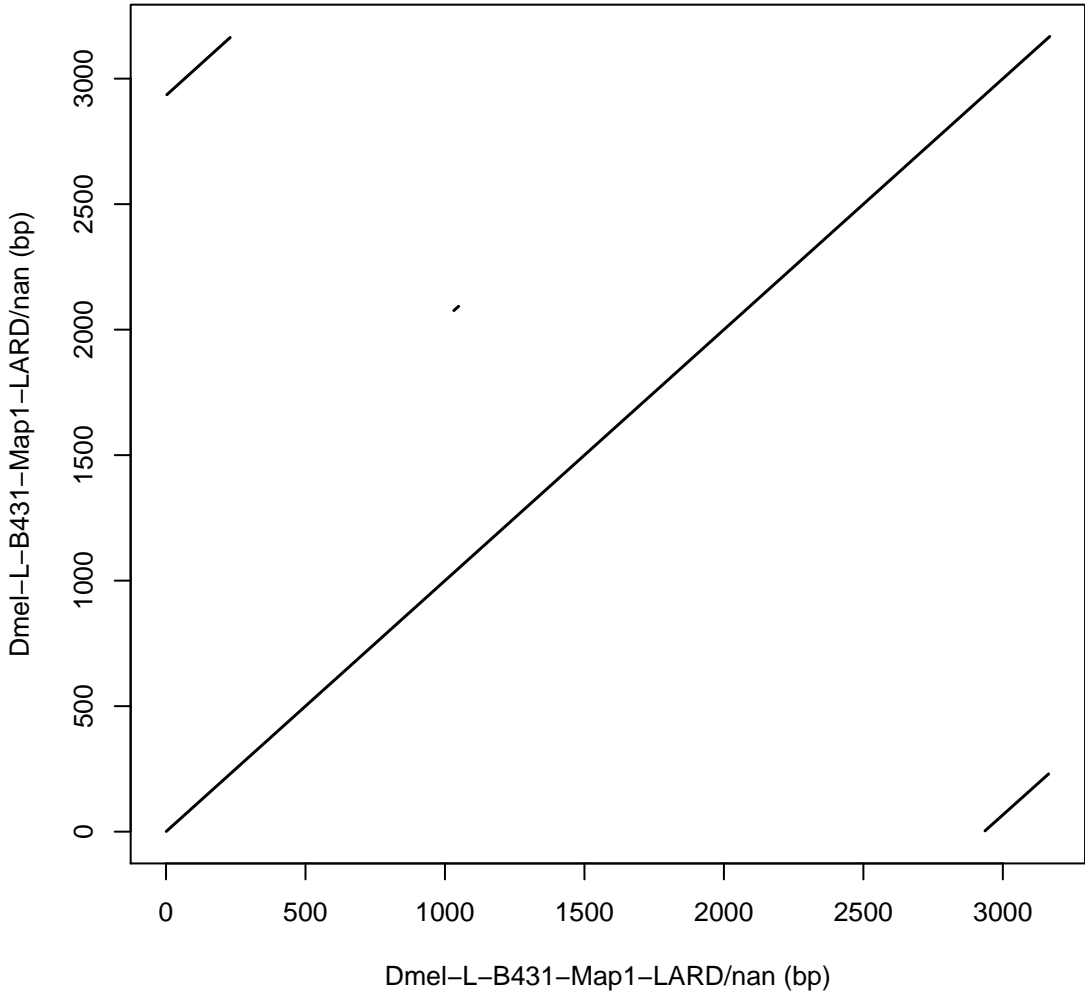consensus size: 3168bp; fragments: 3; full length: 1 (>=2851.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected