TE: Dmel−B−R1473−Map4−LTR
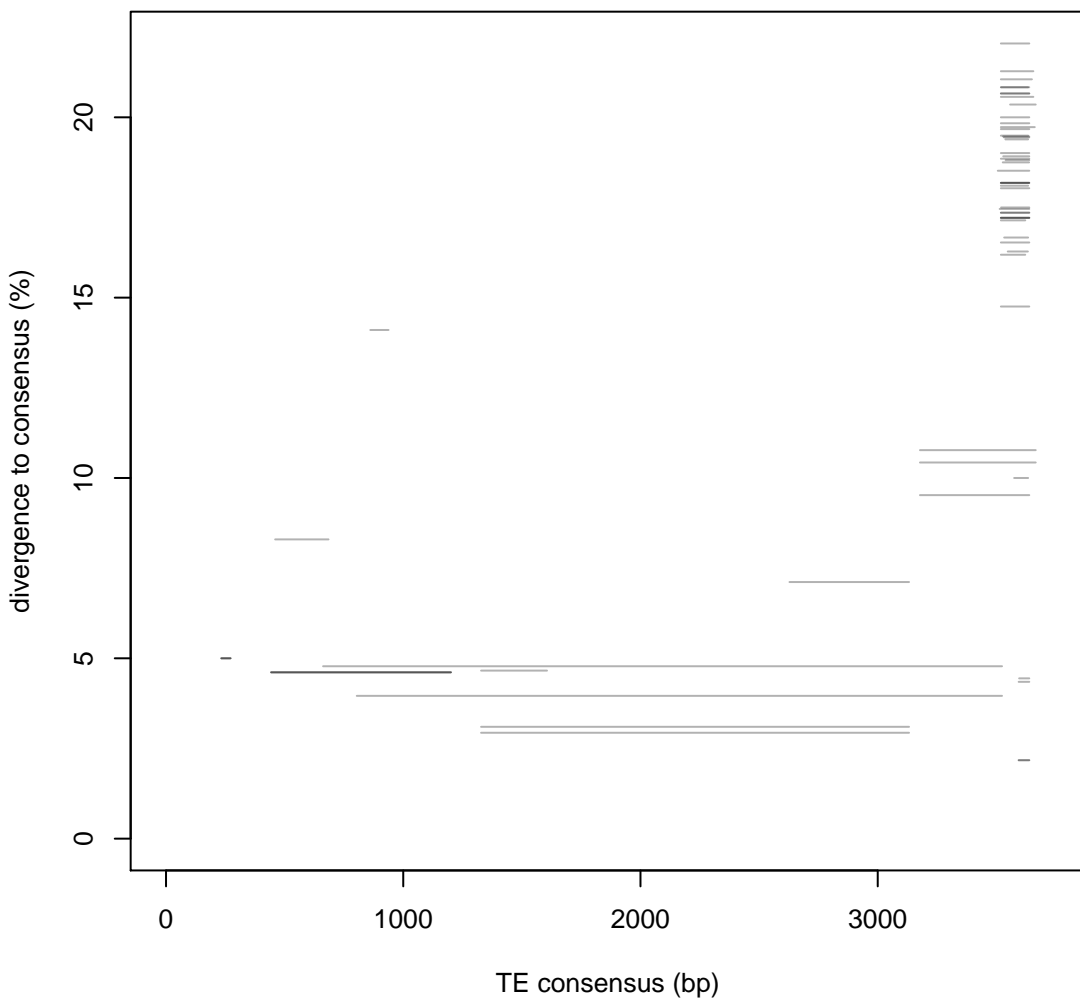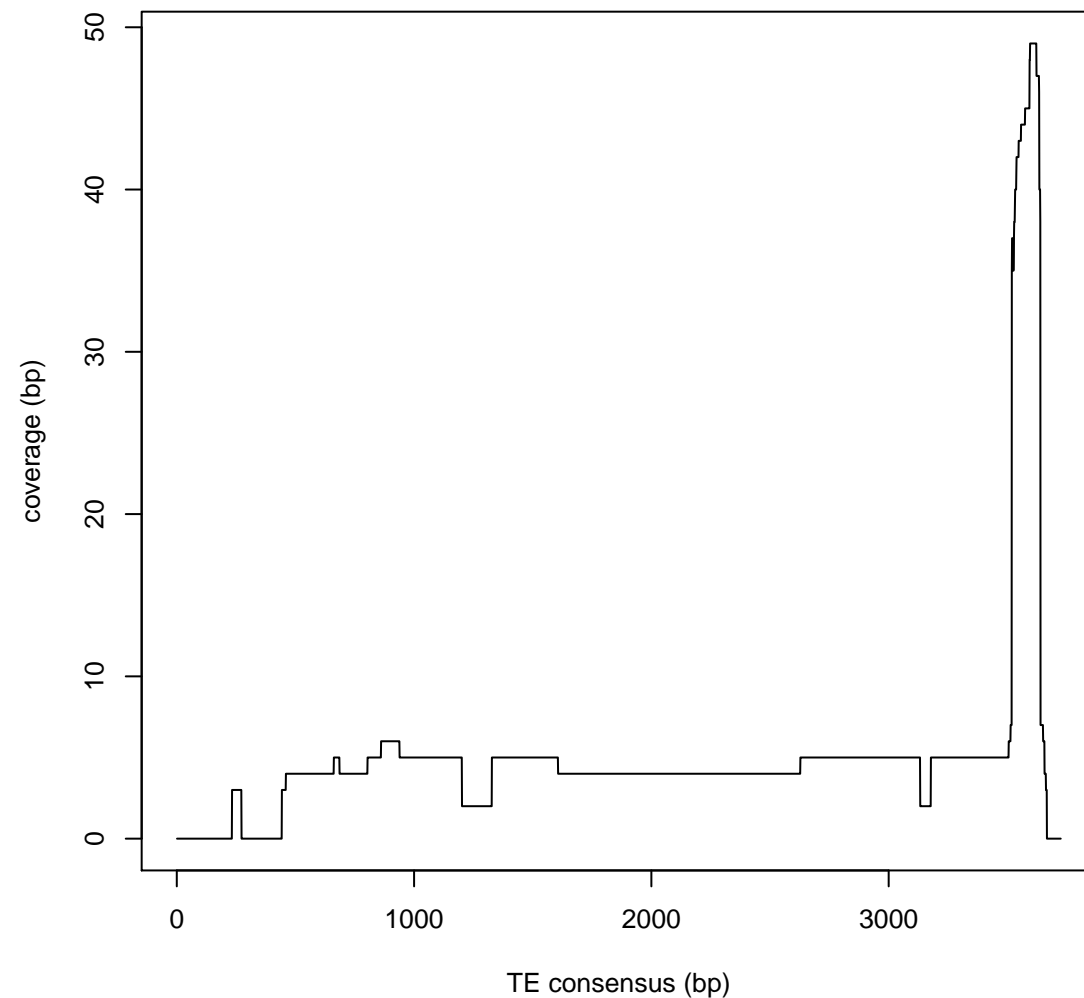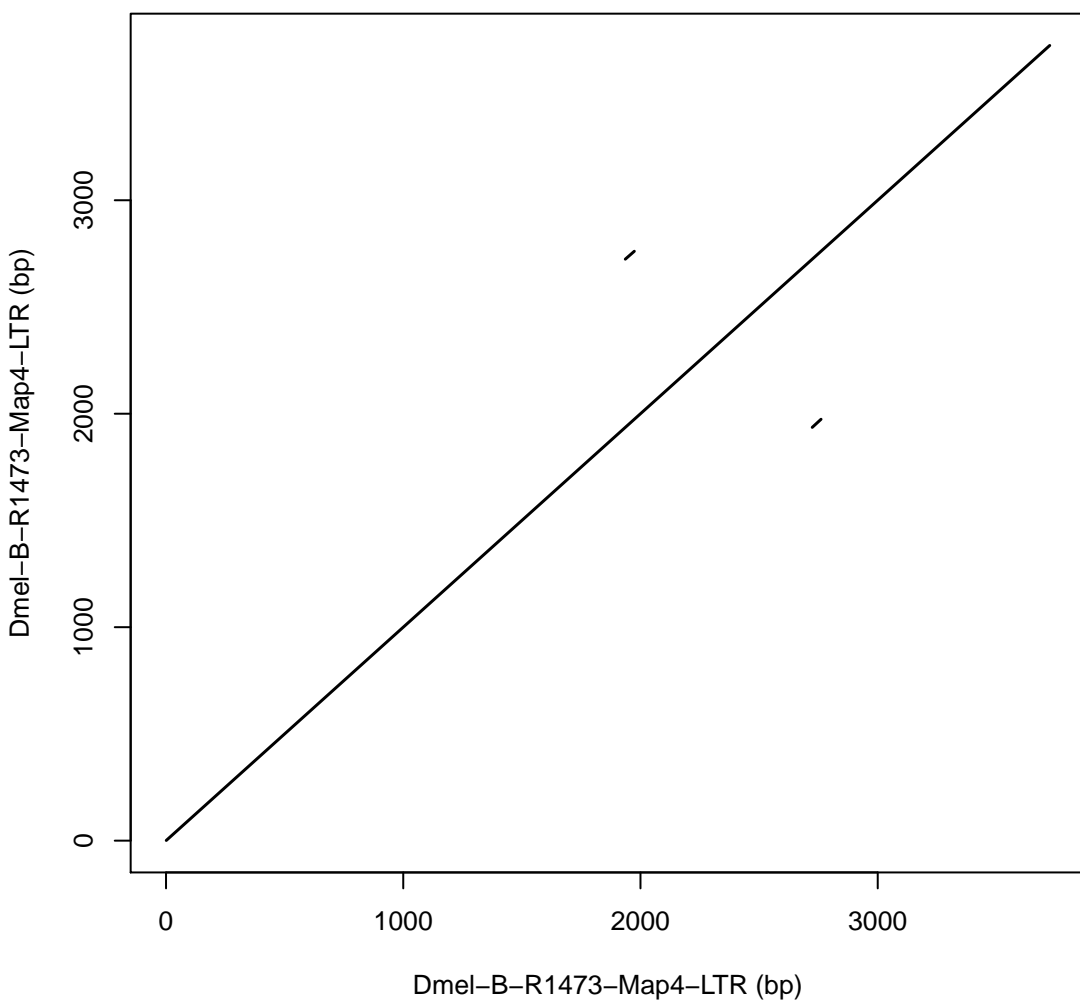consensus size: 3725bp; fragments: 63; full length: 0 (>=3352.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Copia12−NVi LTR/Copia