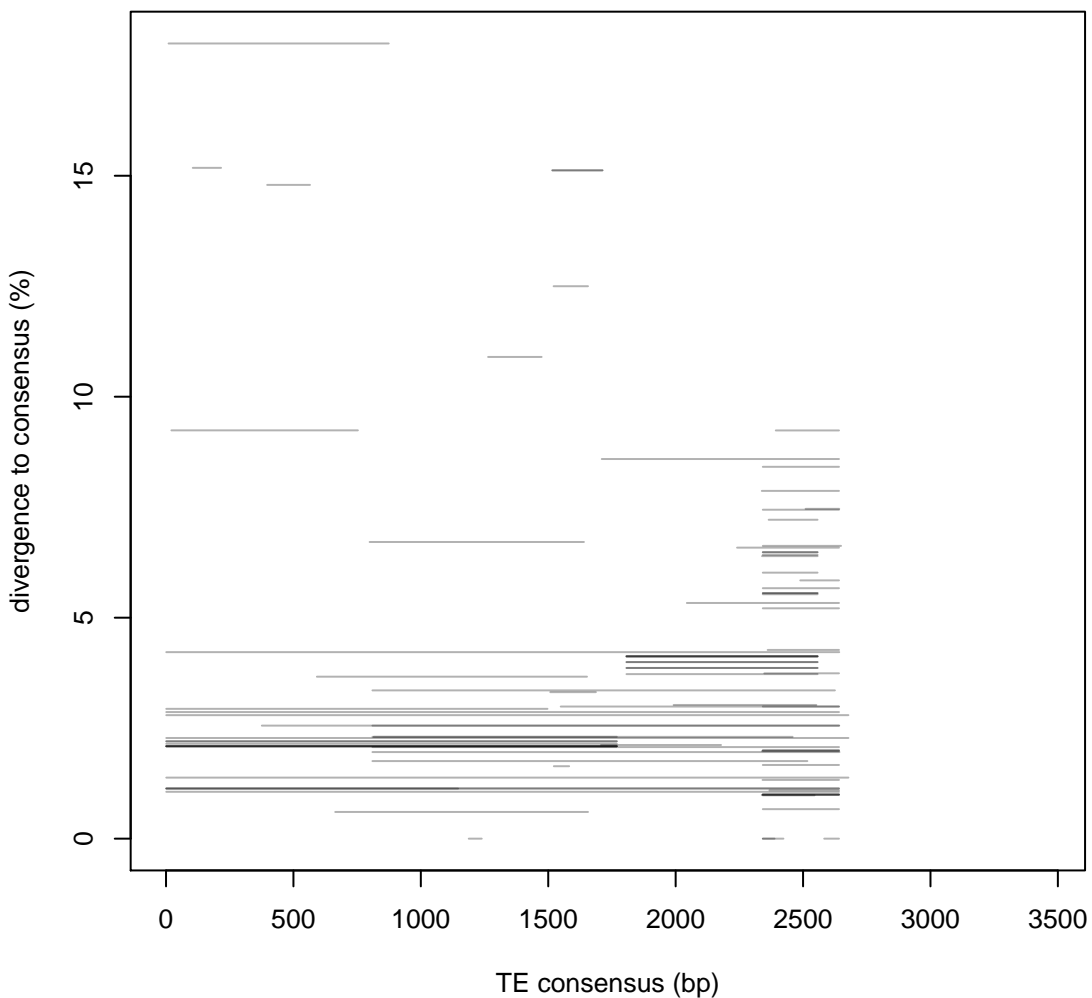
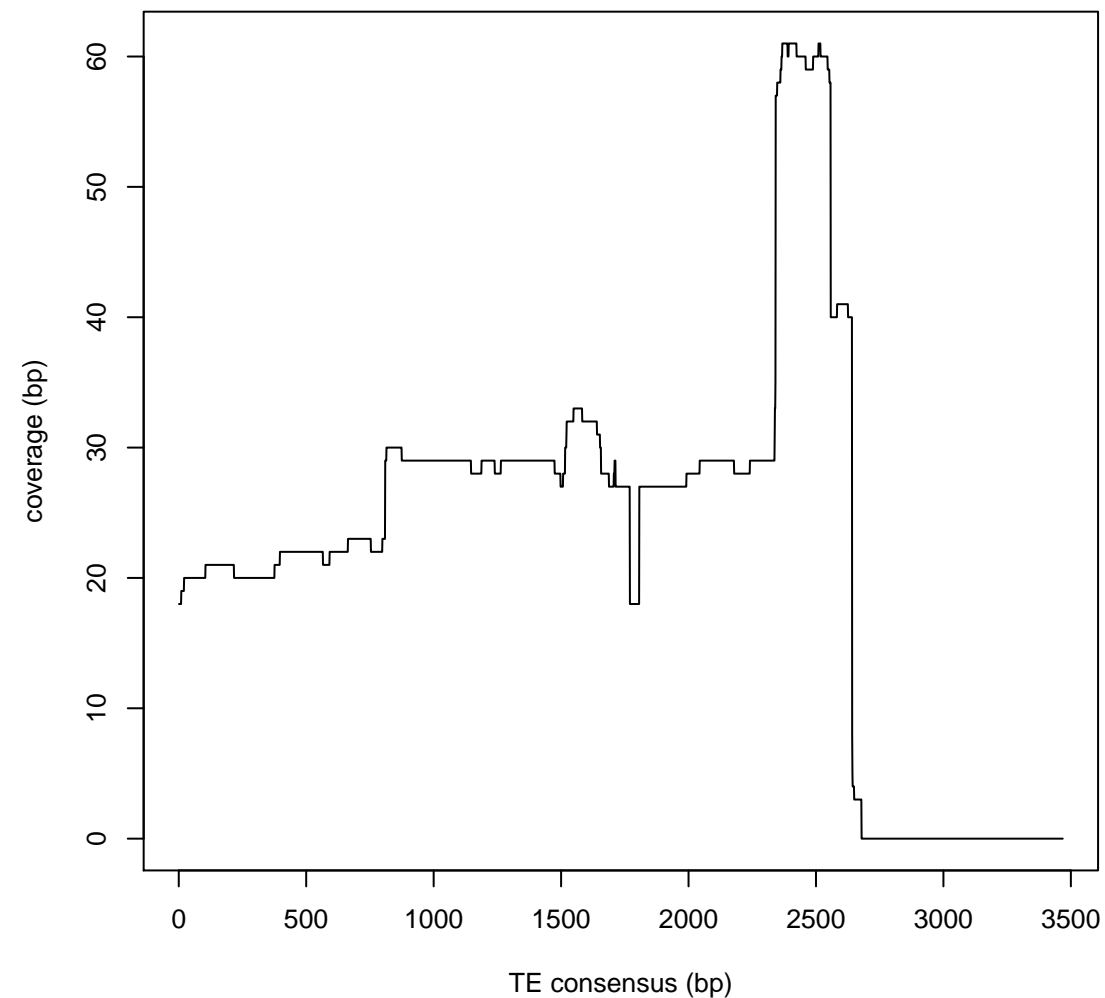


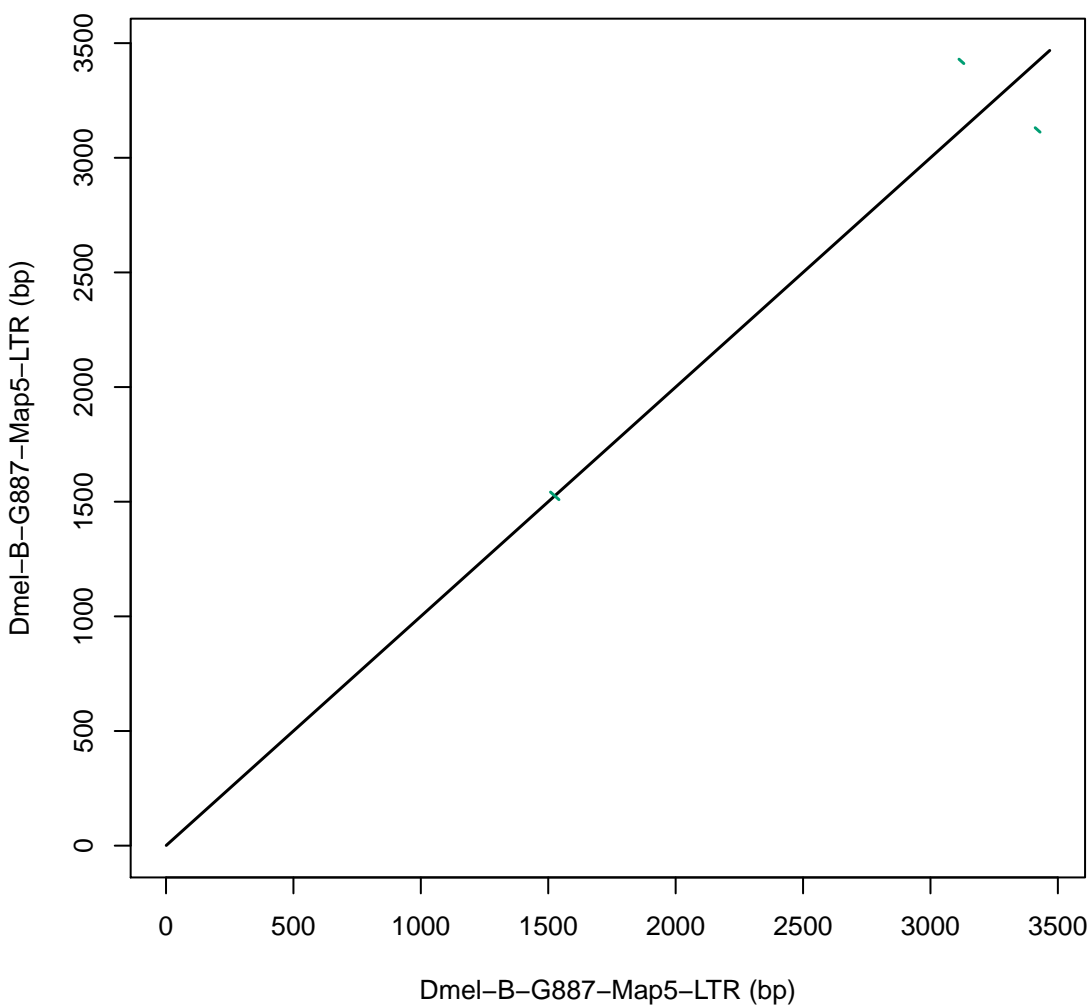
TE: Dmel-B-G887-Map5-LTR
consensus size: 3468bp; fragments: 91; full length: 0 (≥ 3121.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

