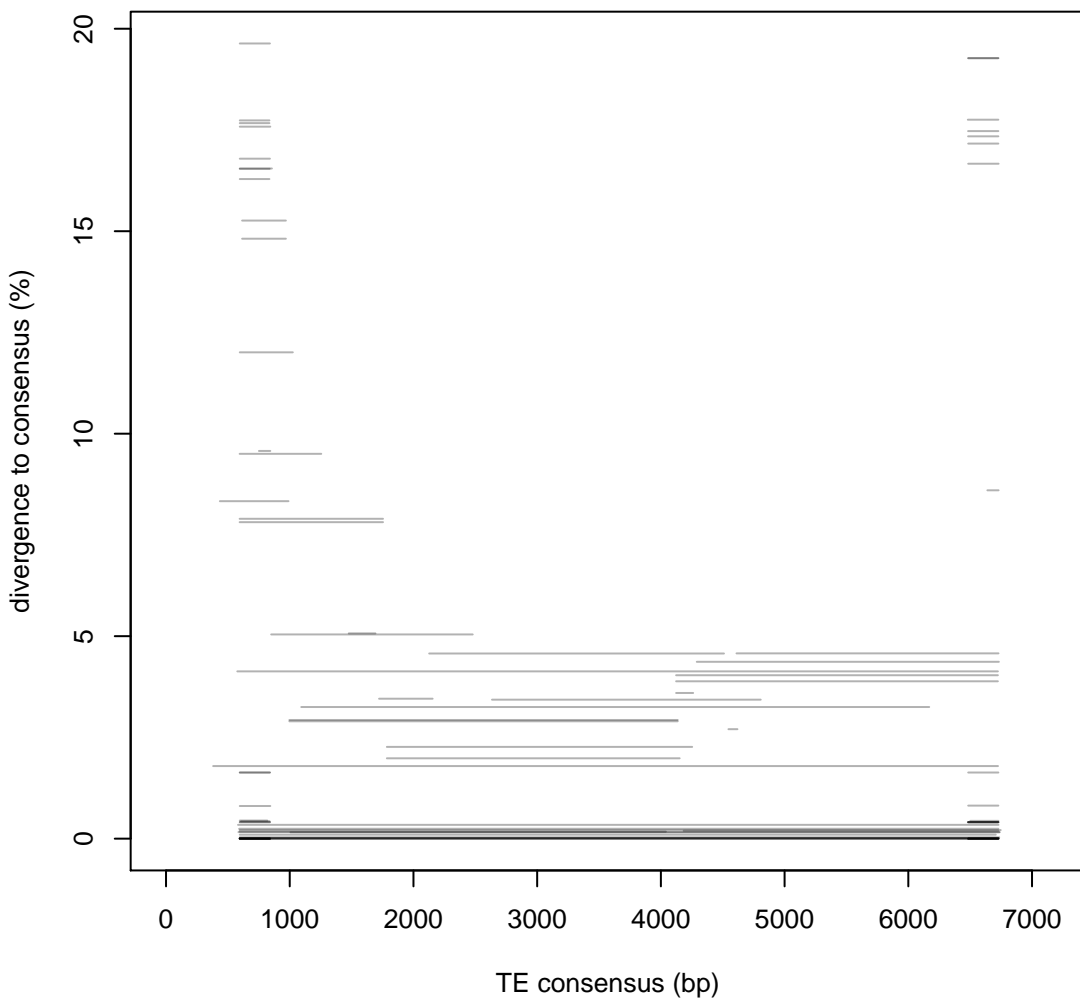
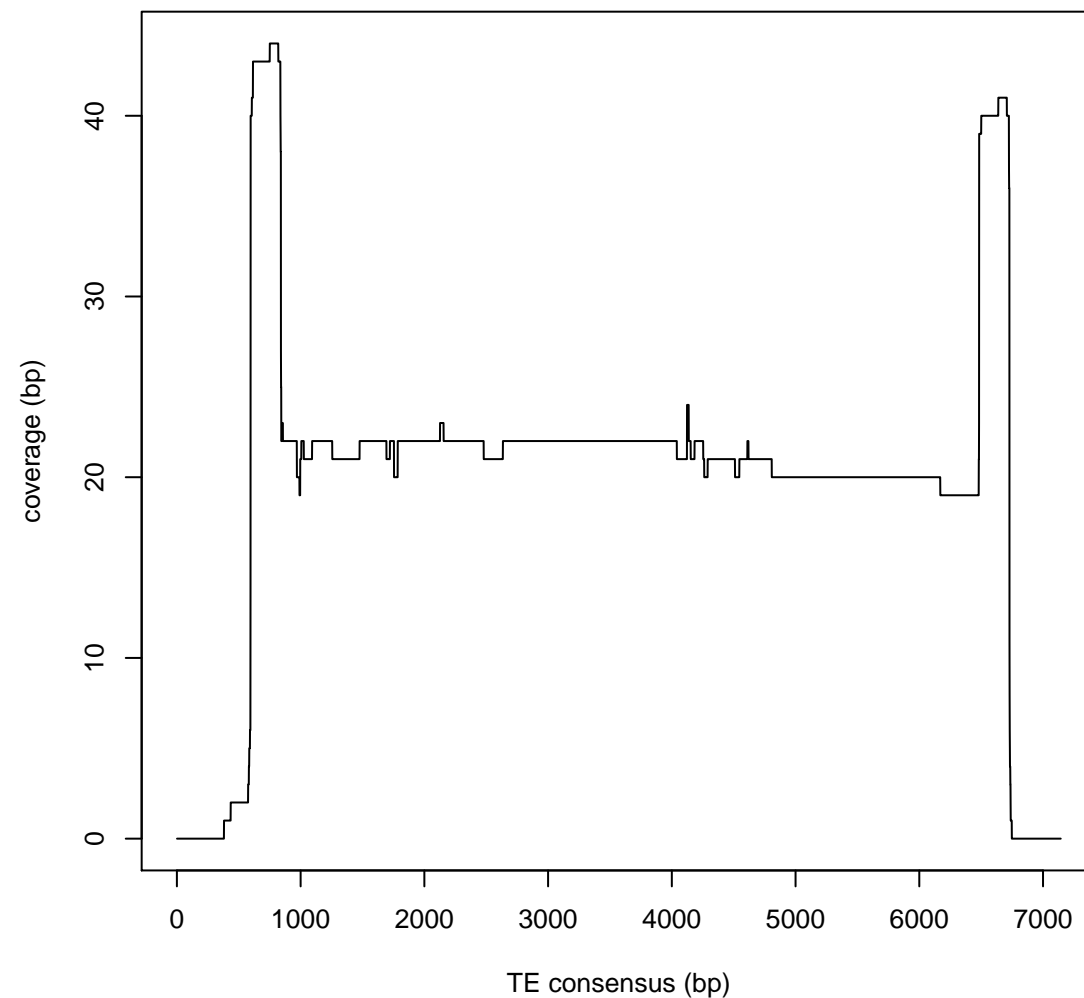


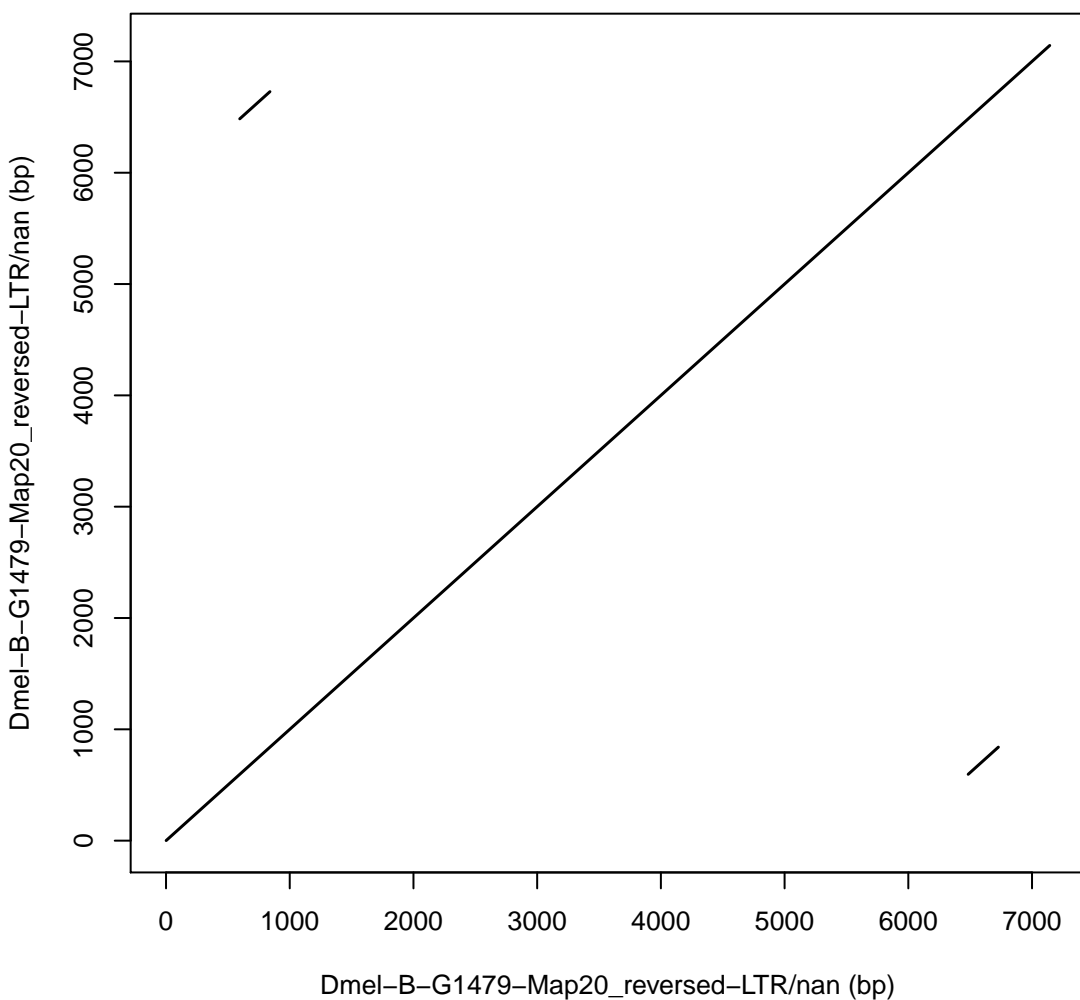
TE: Dmel-B-G1479-Map20_reversed-LTR/nan
consensus size: 7143bp; fragments: 84; full length: 0 (>=6428.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

