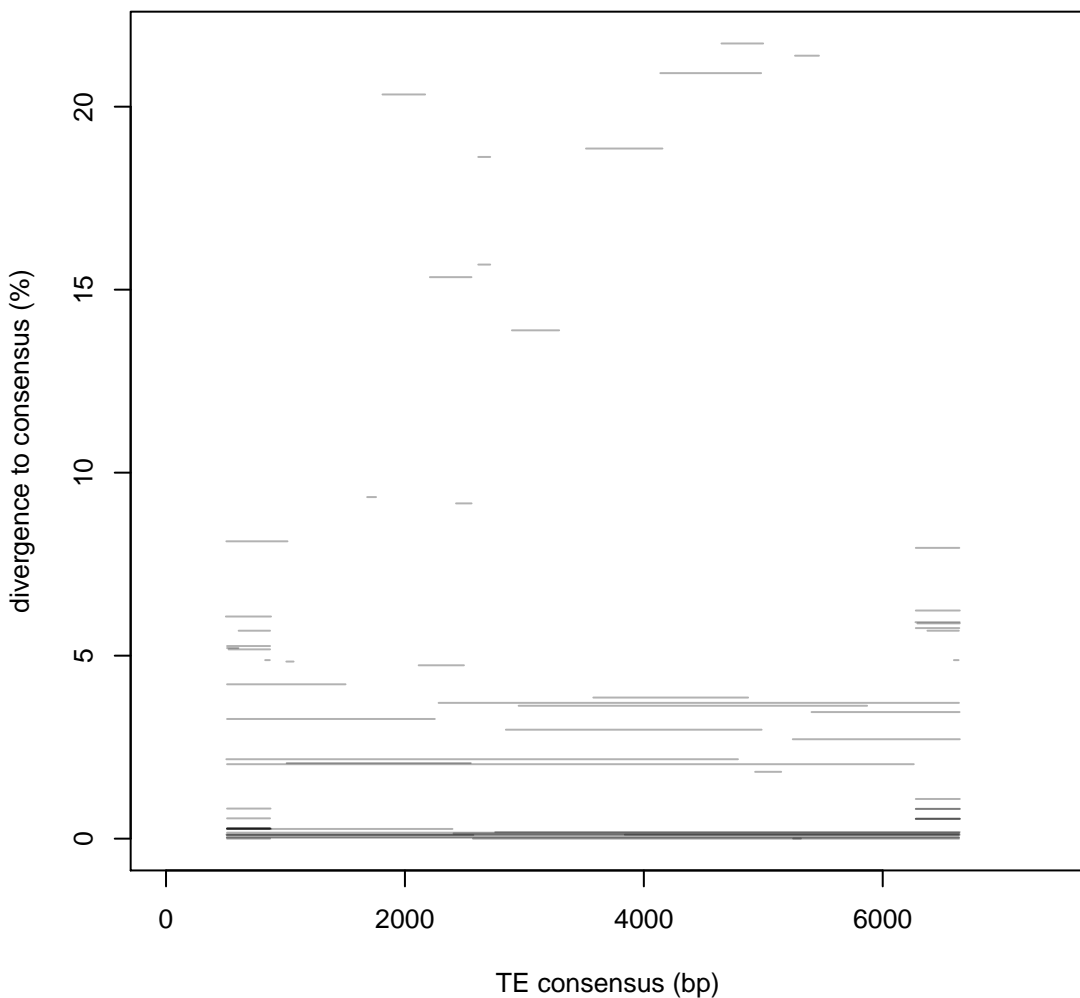
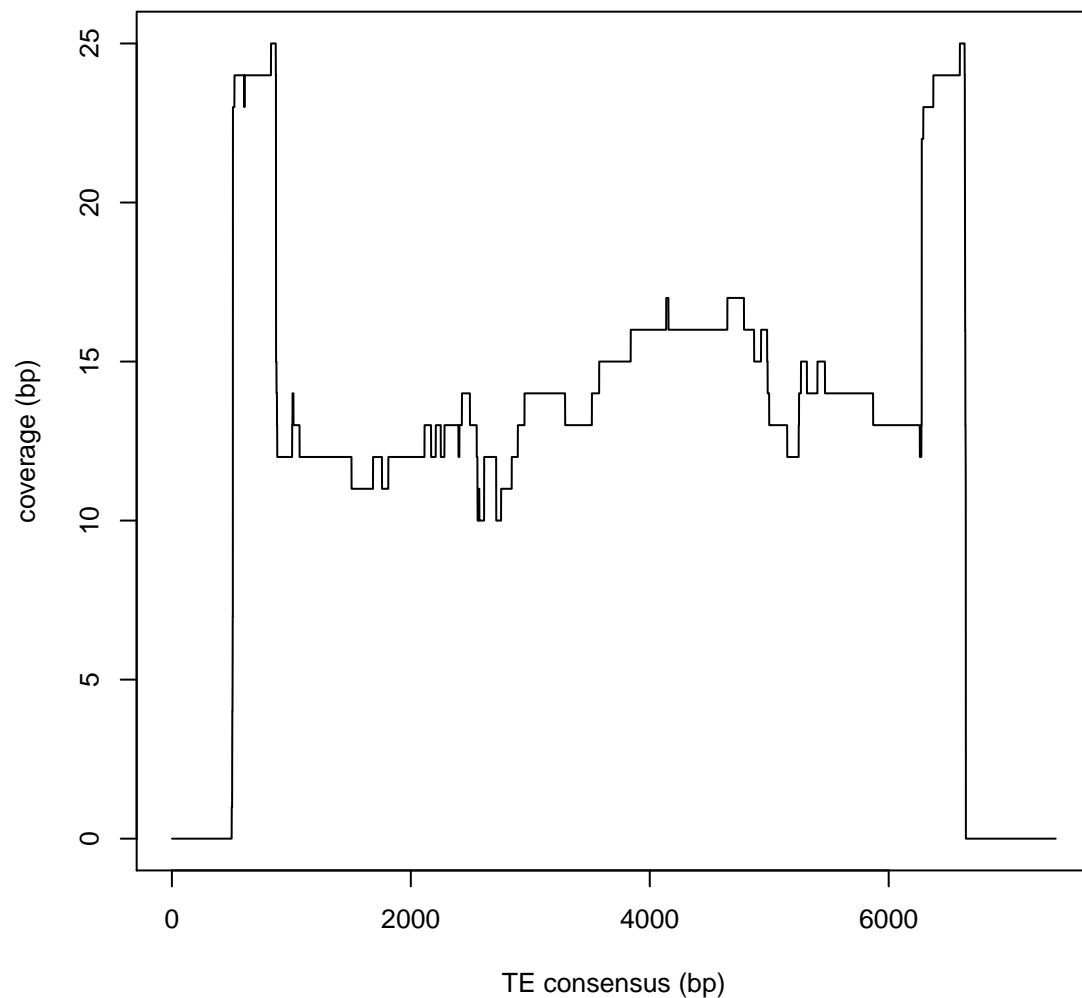


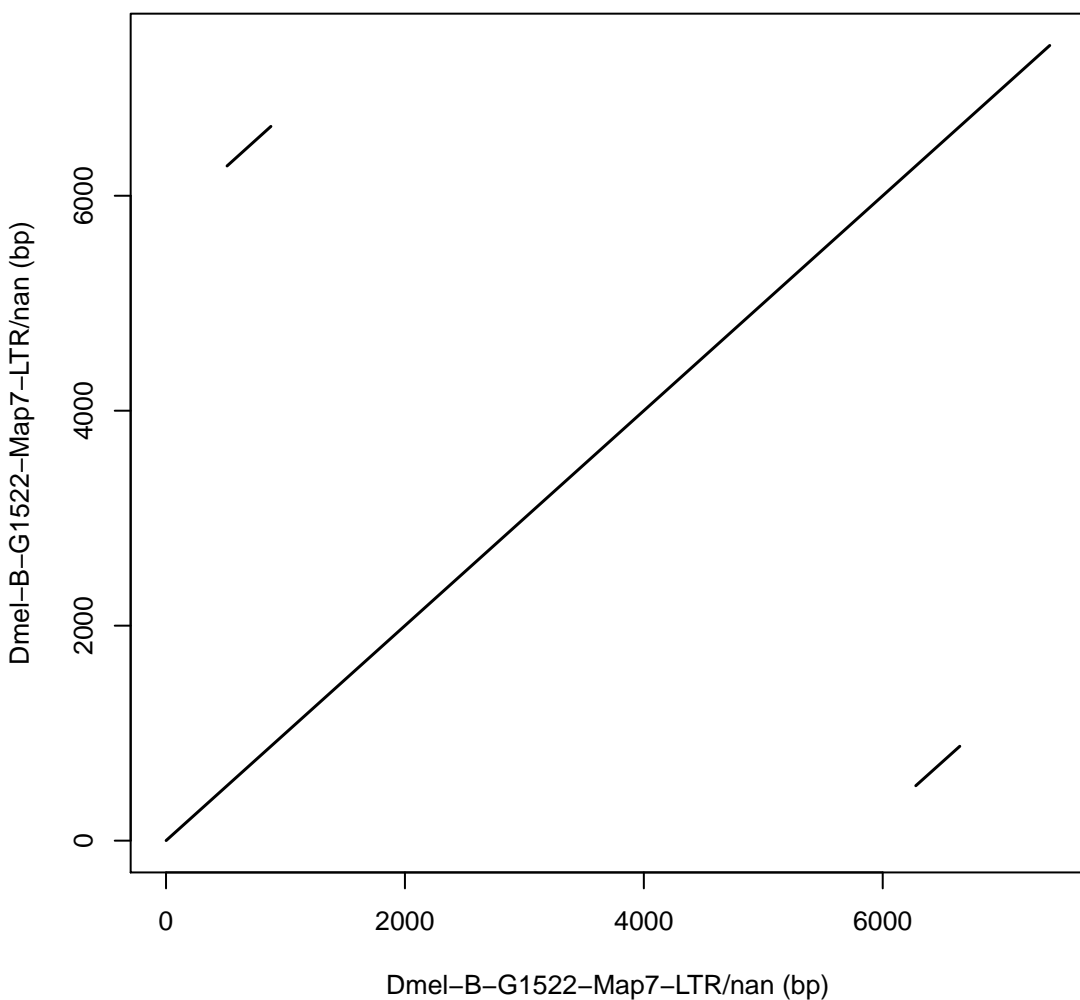
TE: Dmel-B-G1522-Map7-LTR/nan
consensus size: 7398bp; fragments: 65; full length: 0 (>=6658.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

