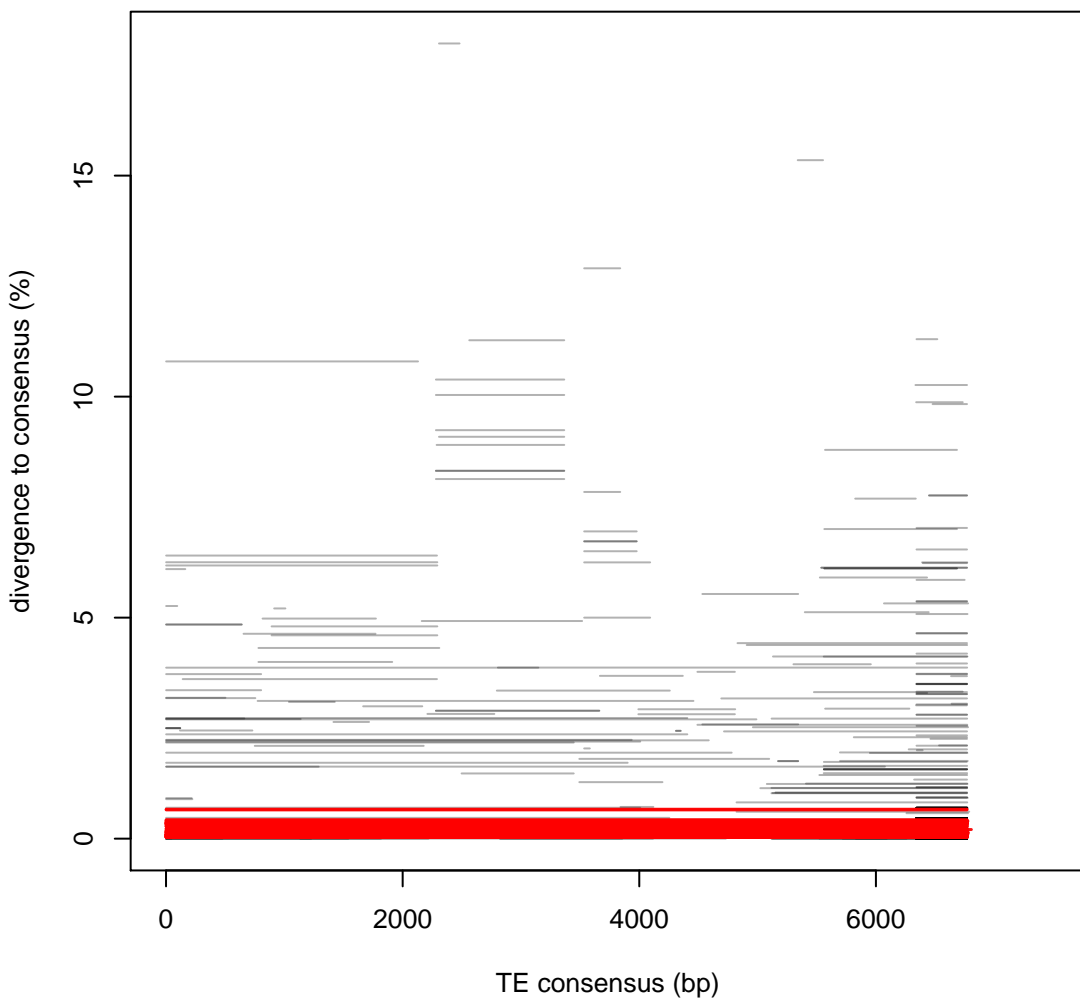
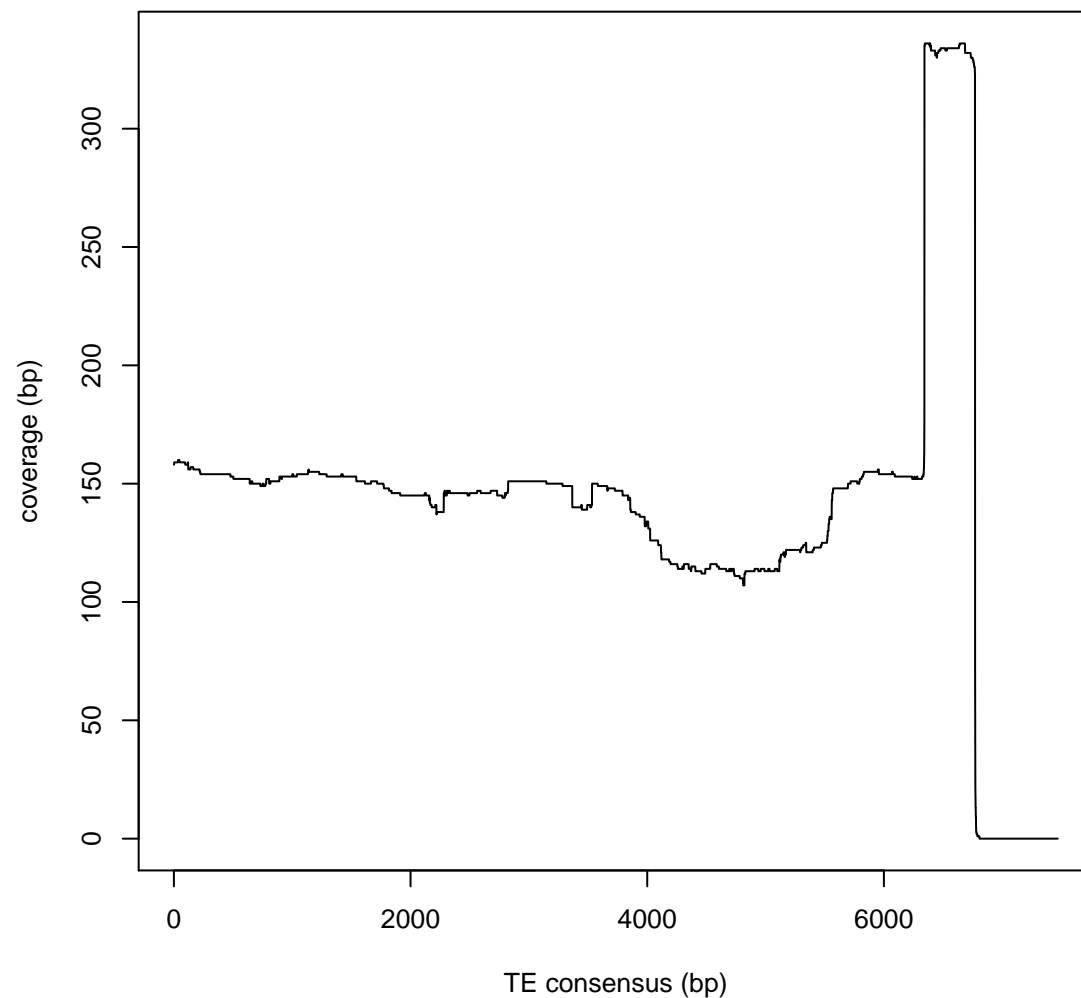


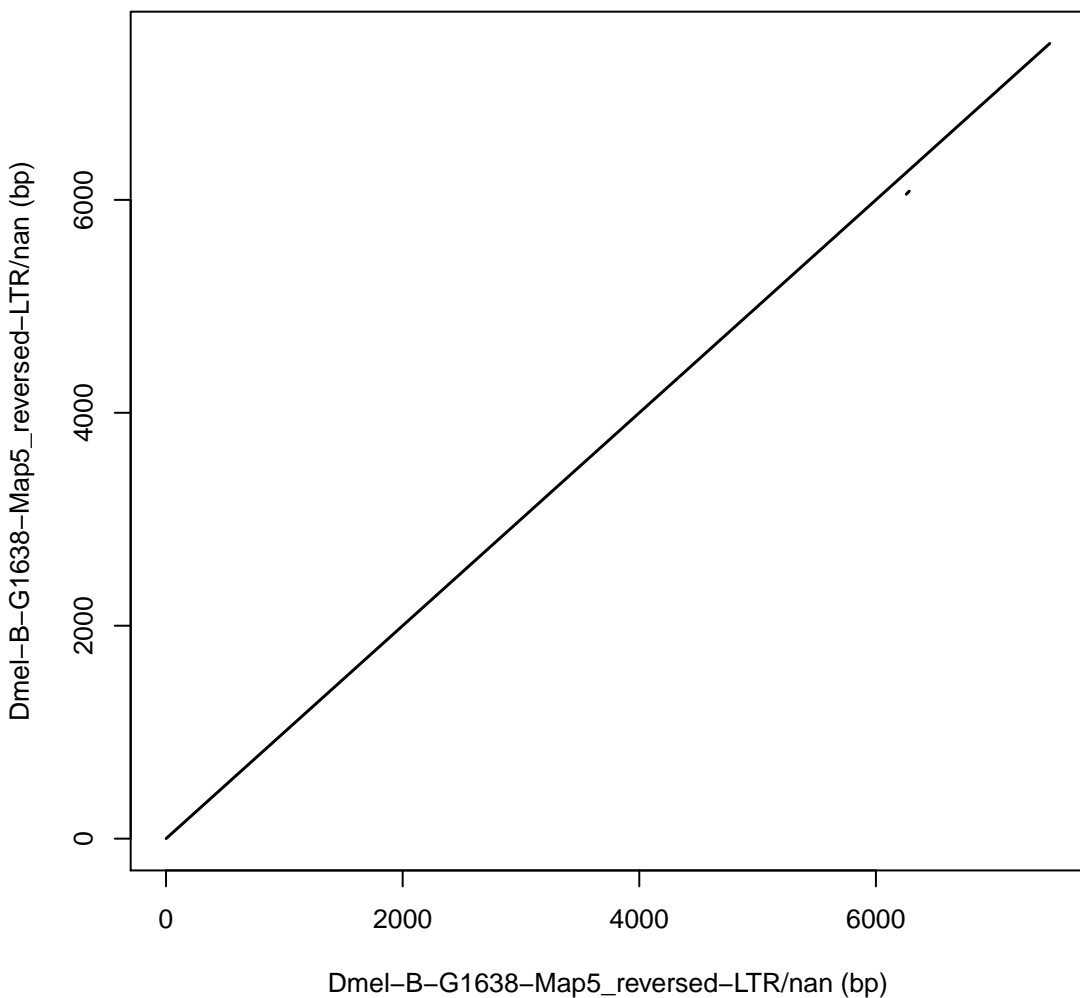
TE: Dmel-B-G1638-Map5_reversed-LTR/nan
consensus size: 7469bp; fragments: 496; full length: 78 (>=6722.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

