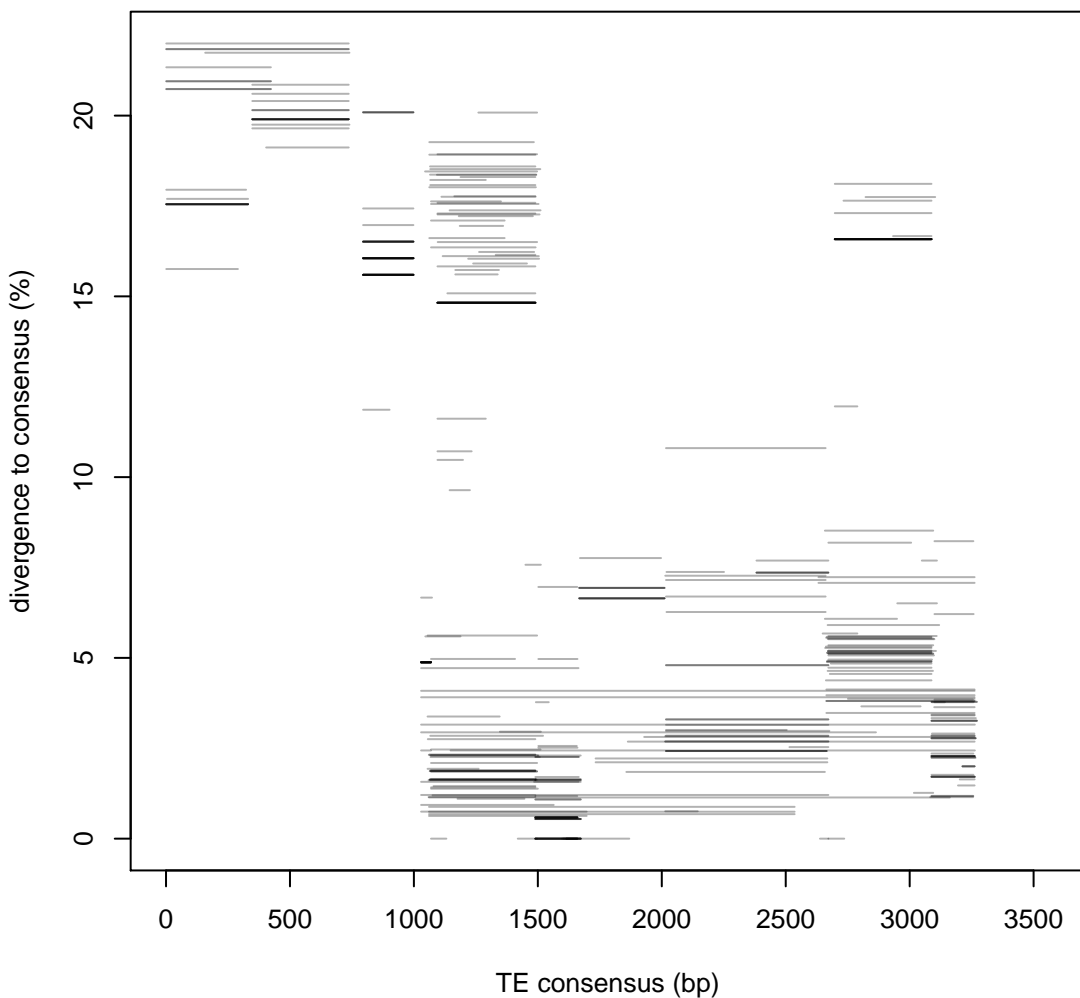
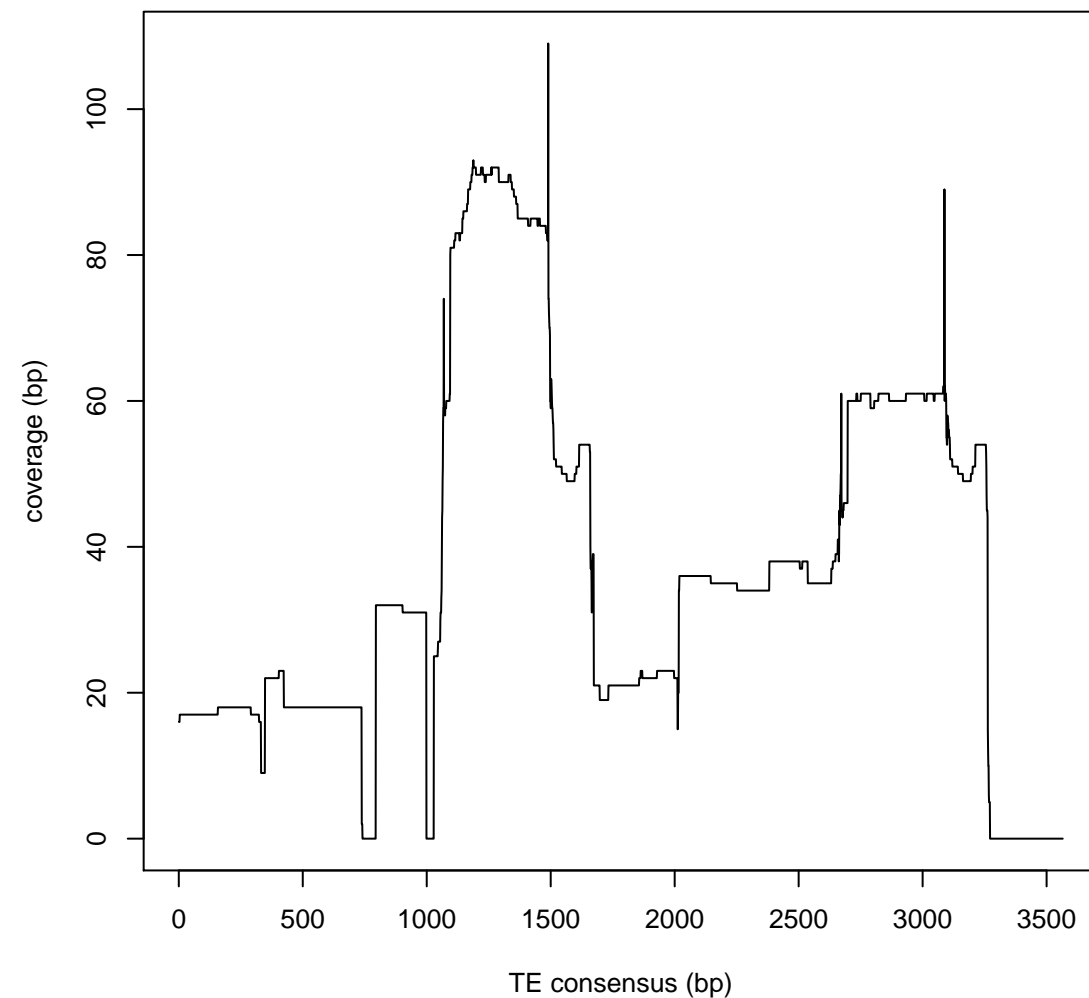


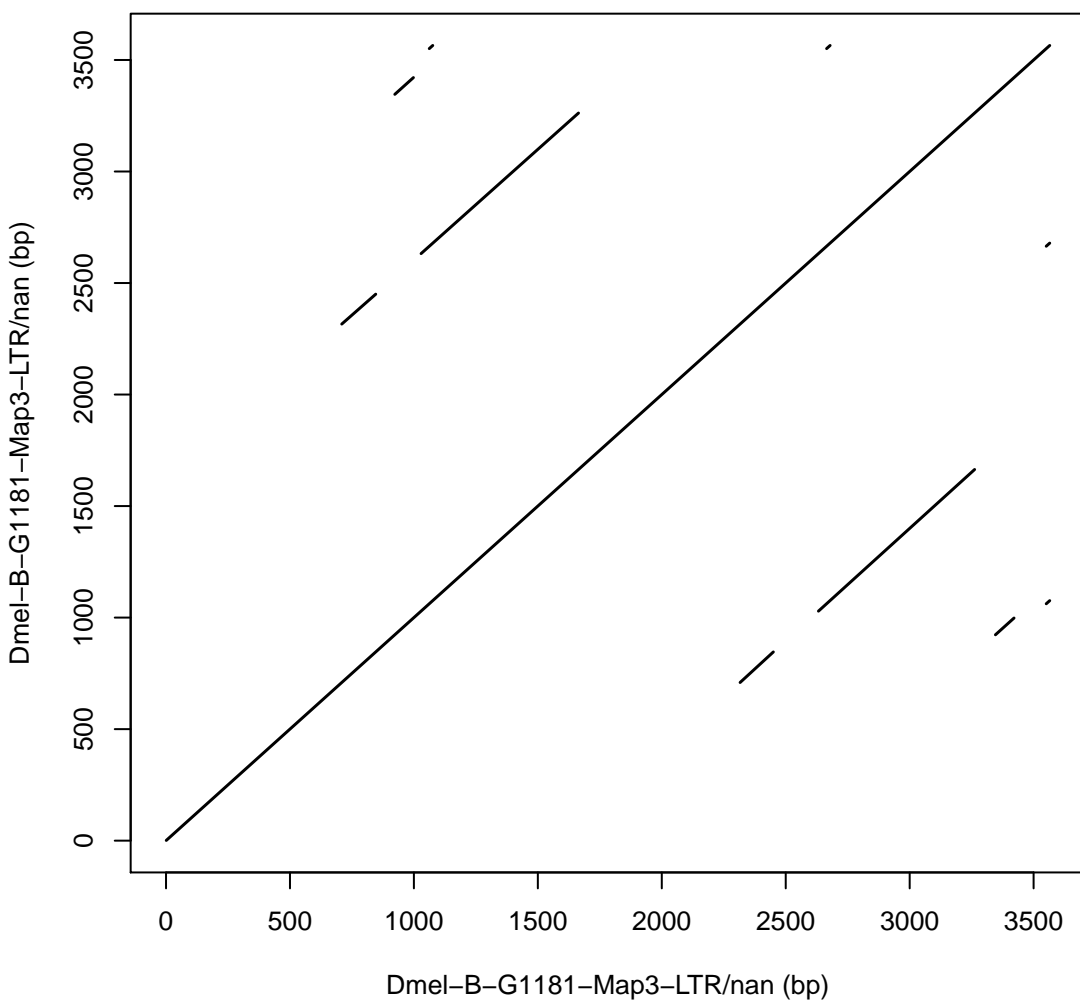
TE: Dmel-B-G1181-Map3-LTR/nan
consensus size: 3565bp; fragments: 364; full length: 0 (>=3208.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

