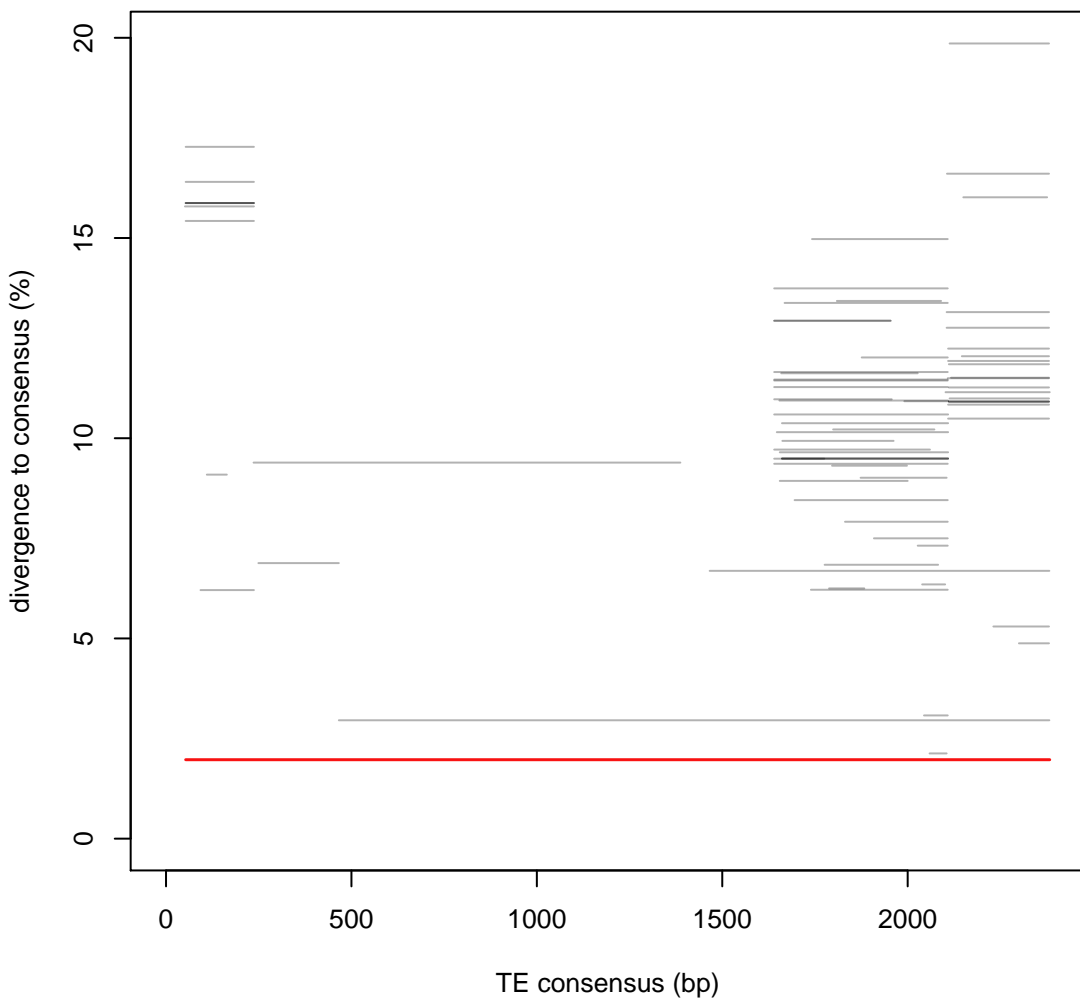
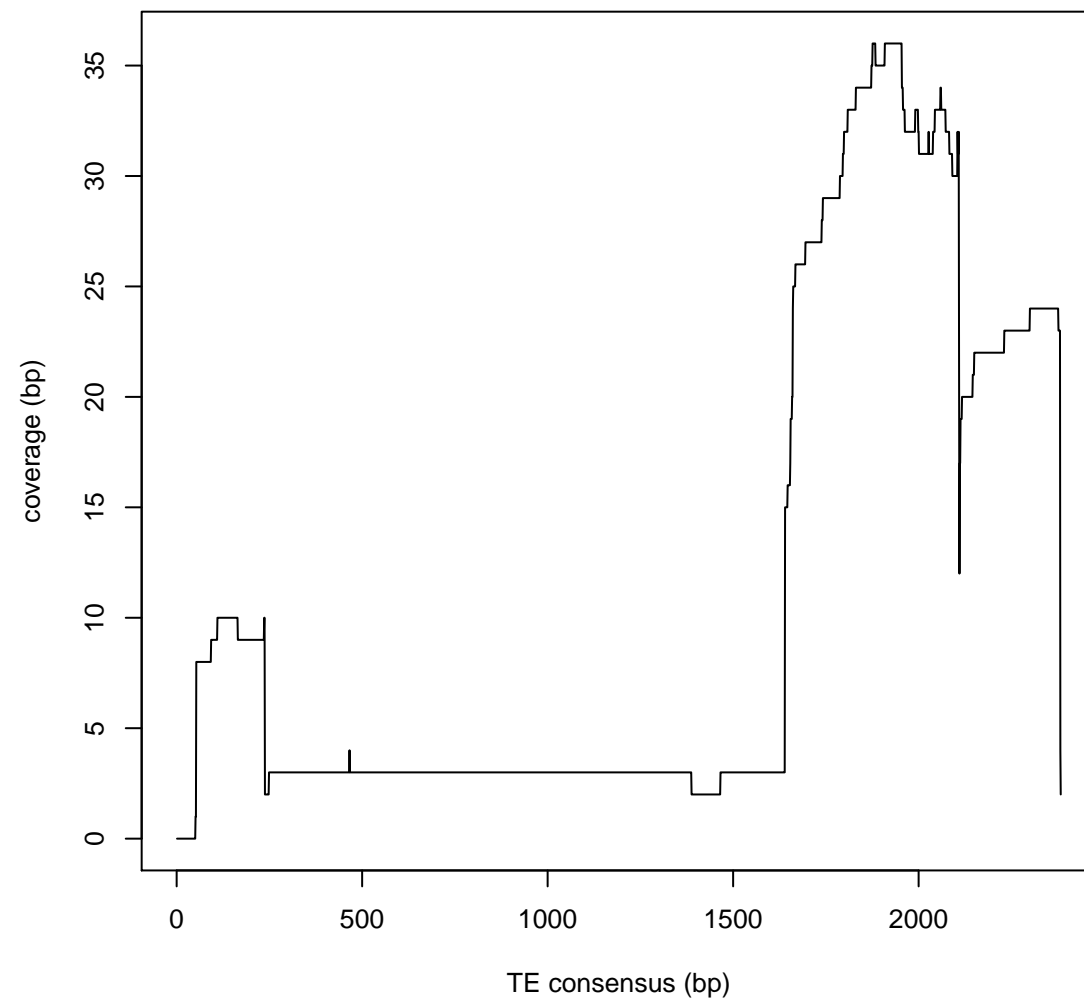


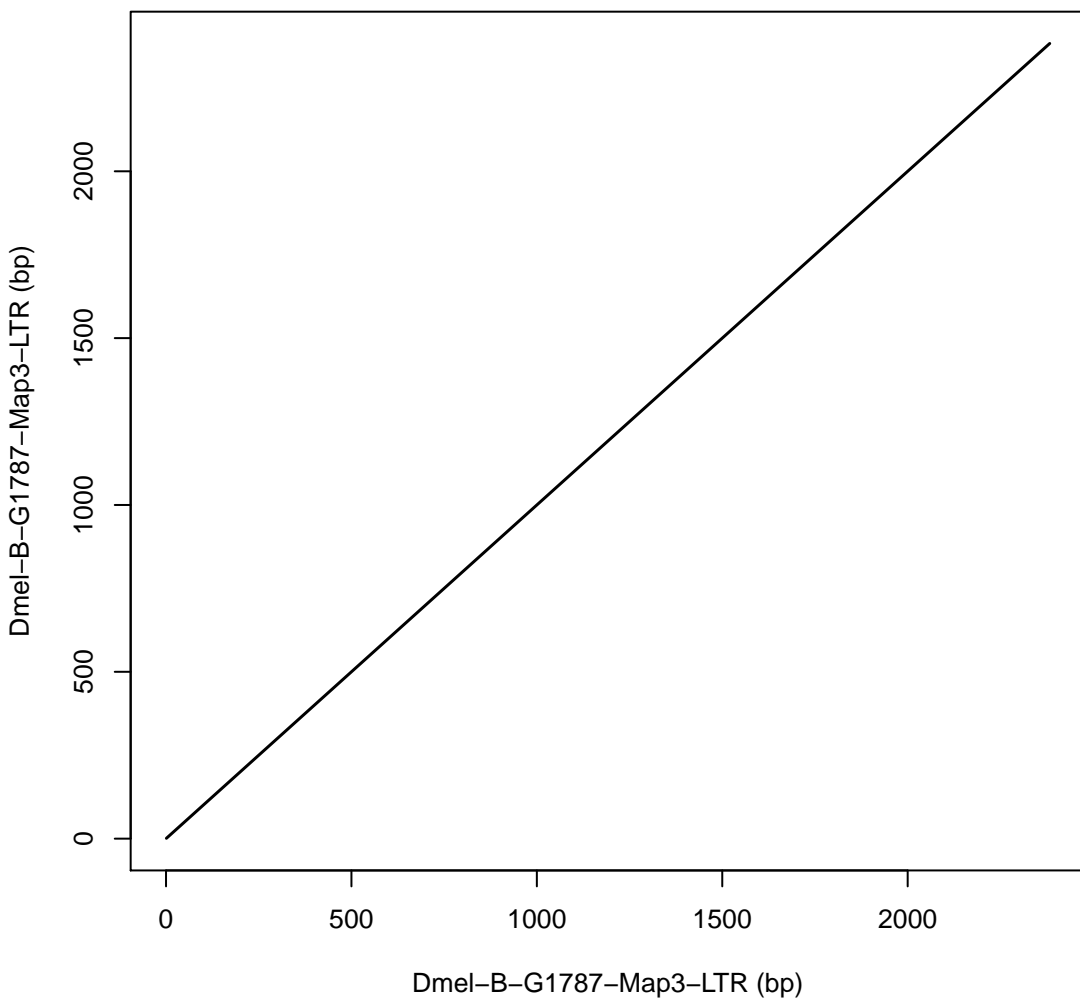
TE: Dmel-B-G1787-Map3-LTR
consensus size: 2383bp; fragments: 75; full length: 1 (>=2144.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

