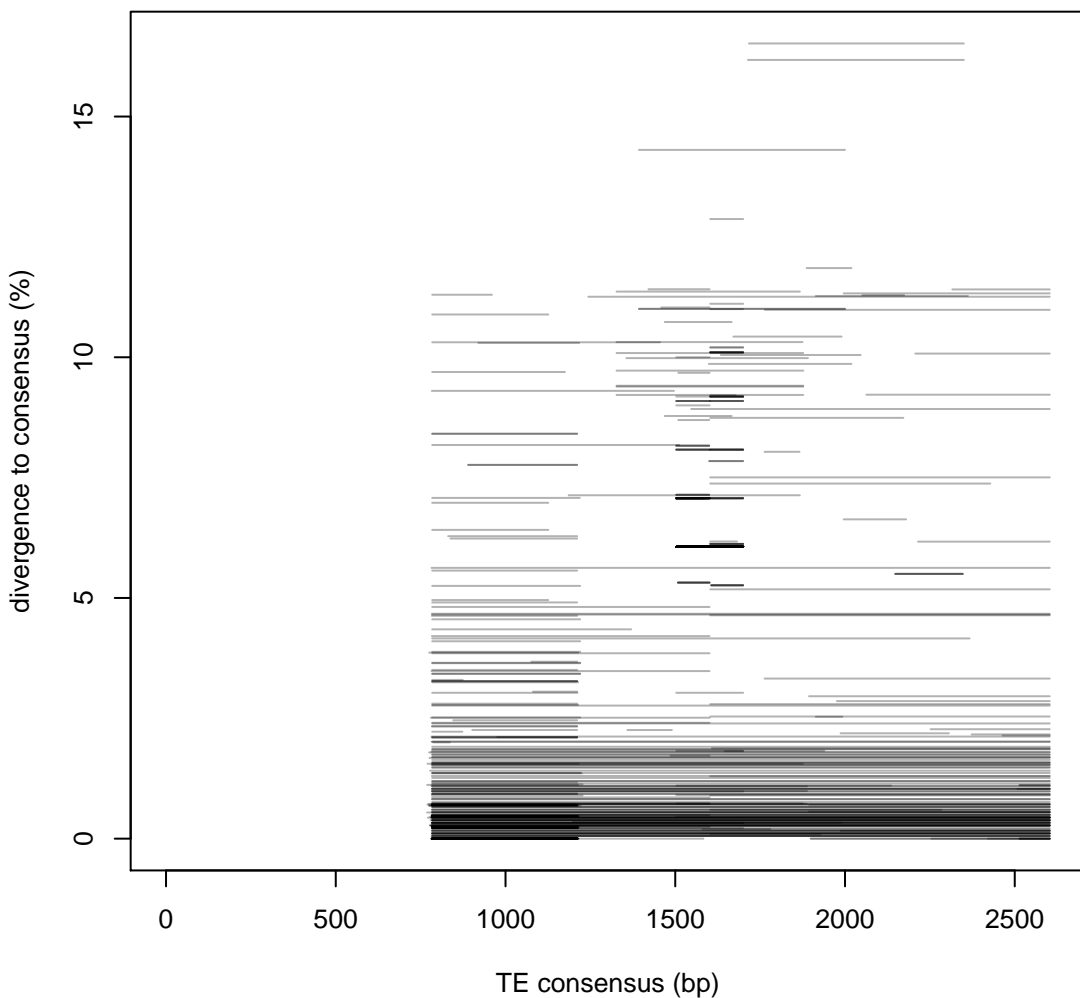
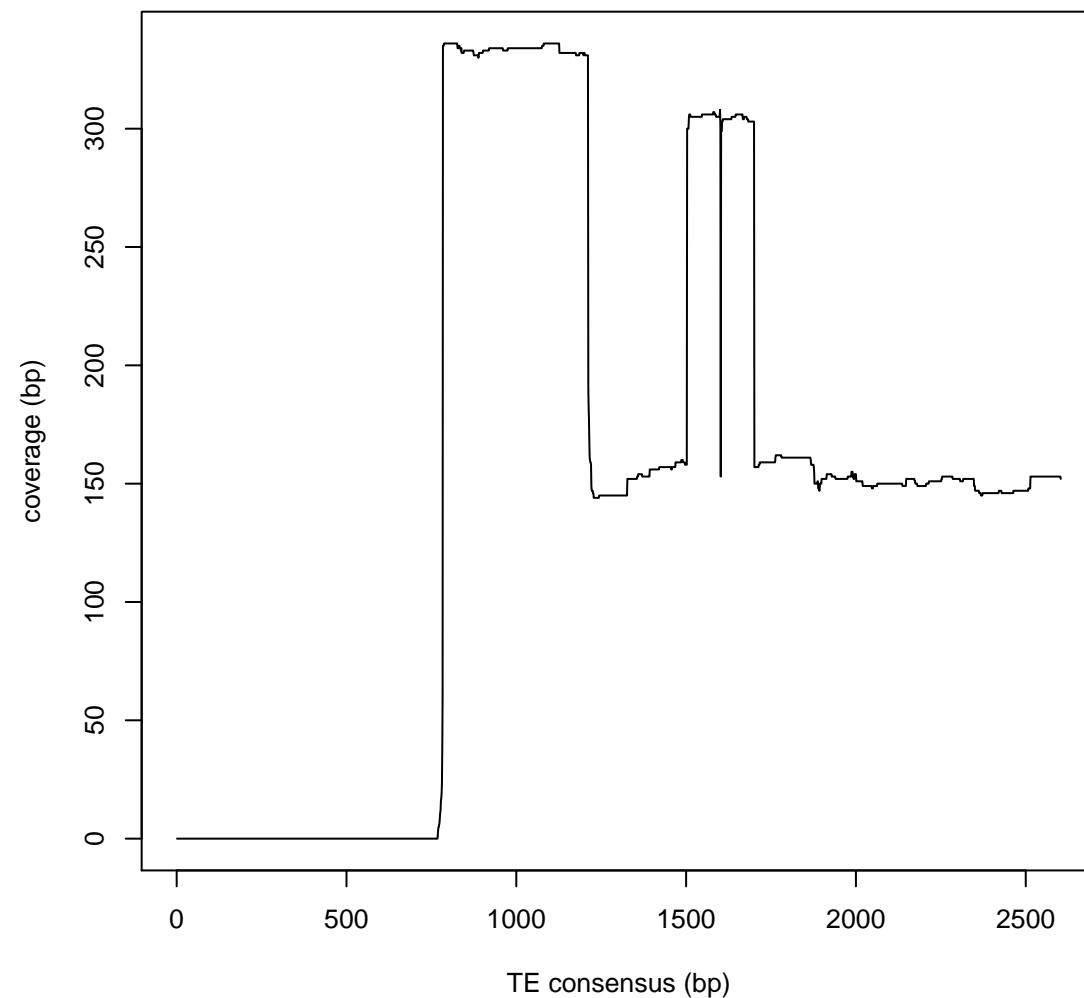


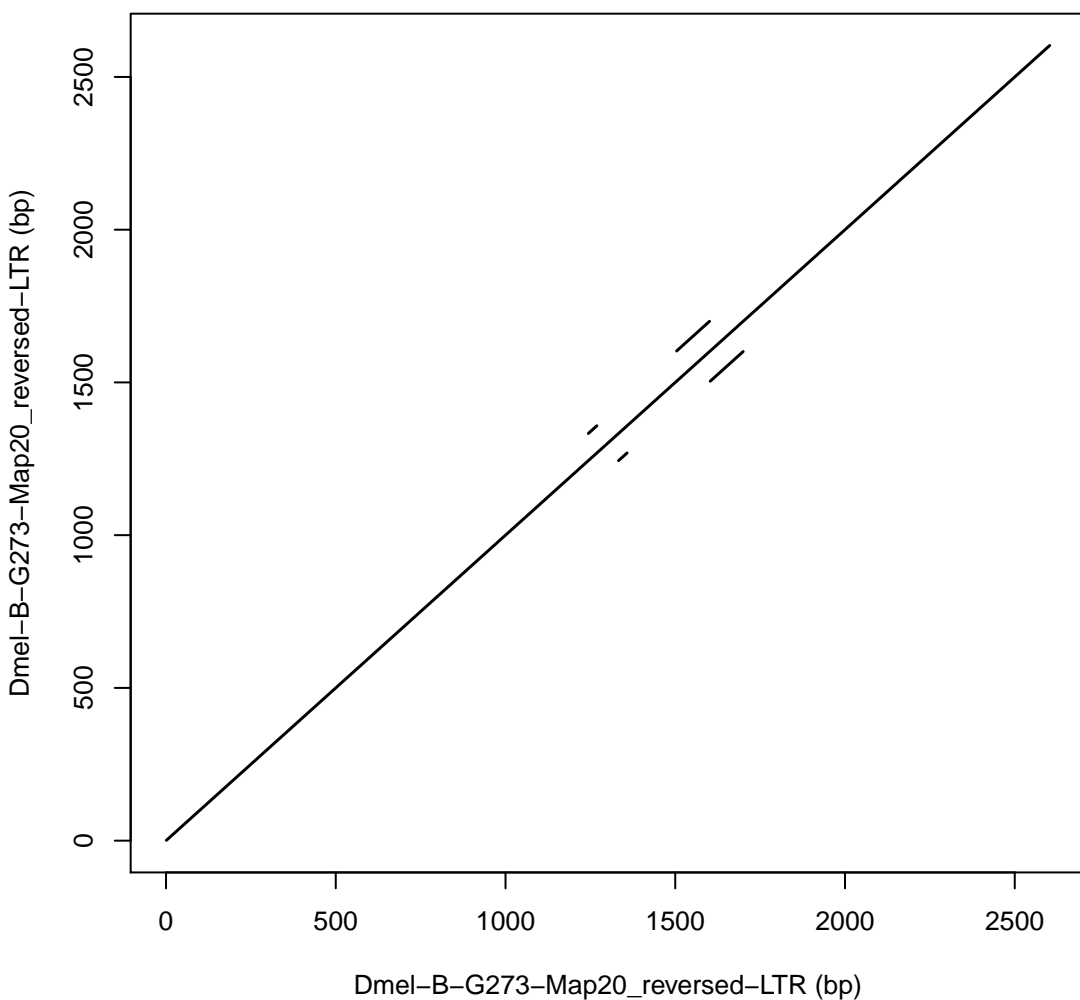
TE: Dmel-B-G273-Map20_reversed-LTR
consensus size: 2603bp; fragments: 710; full length: 0 (>=2342.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

