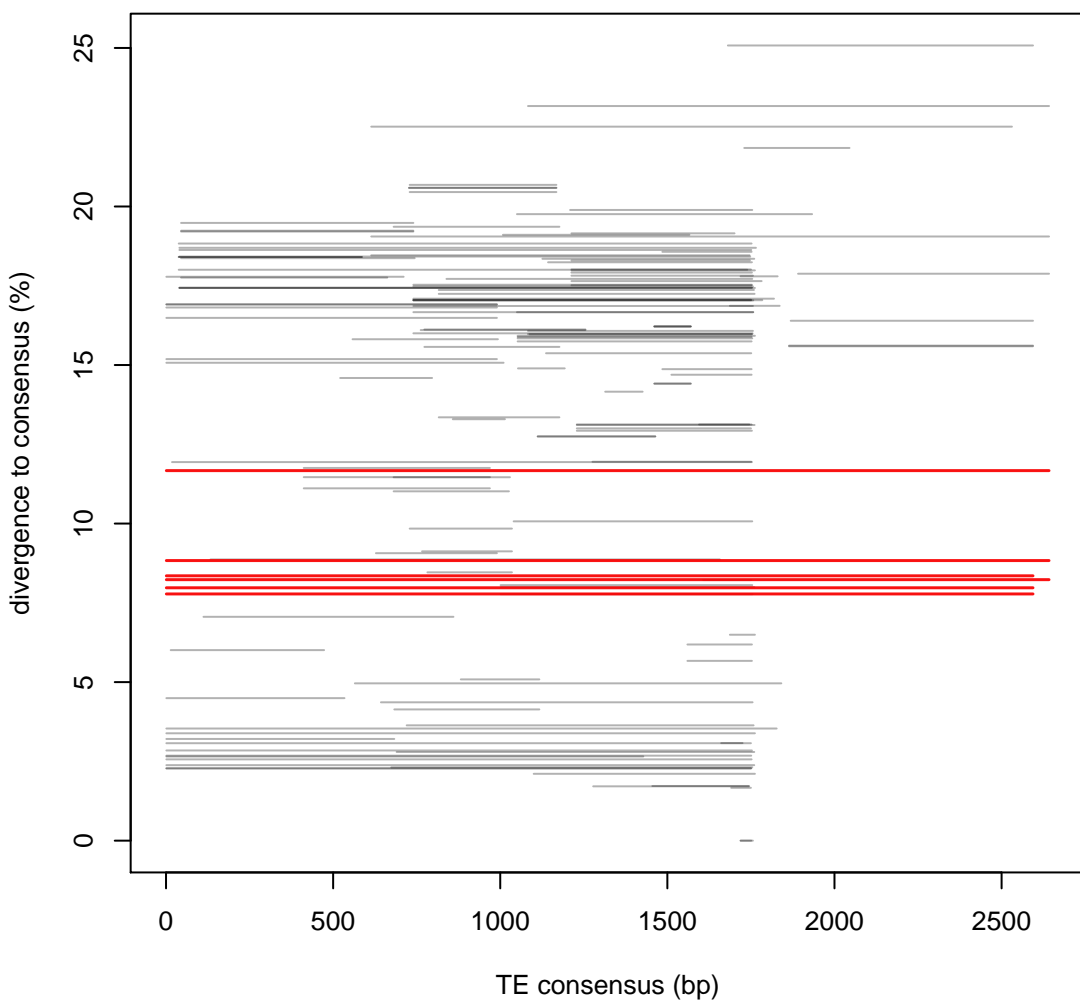
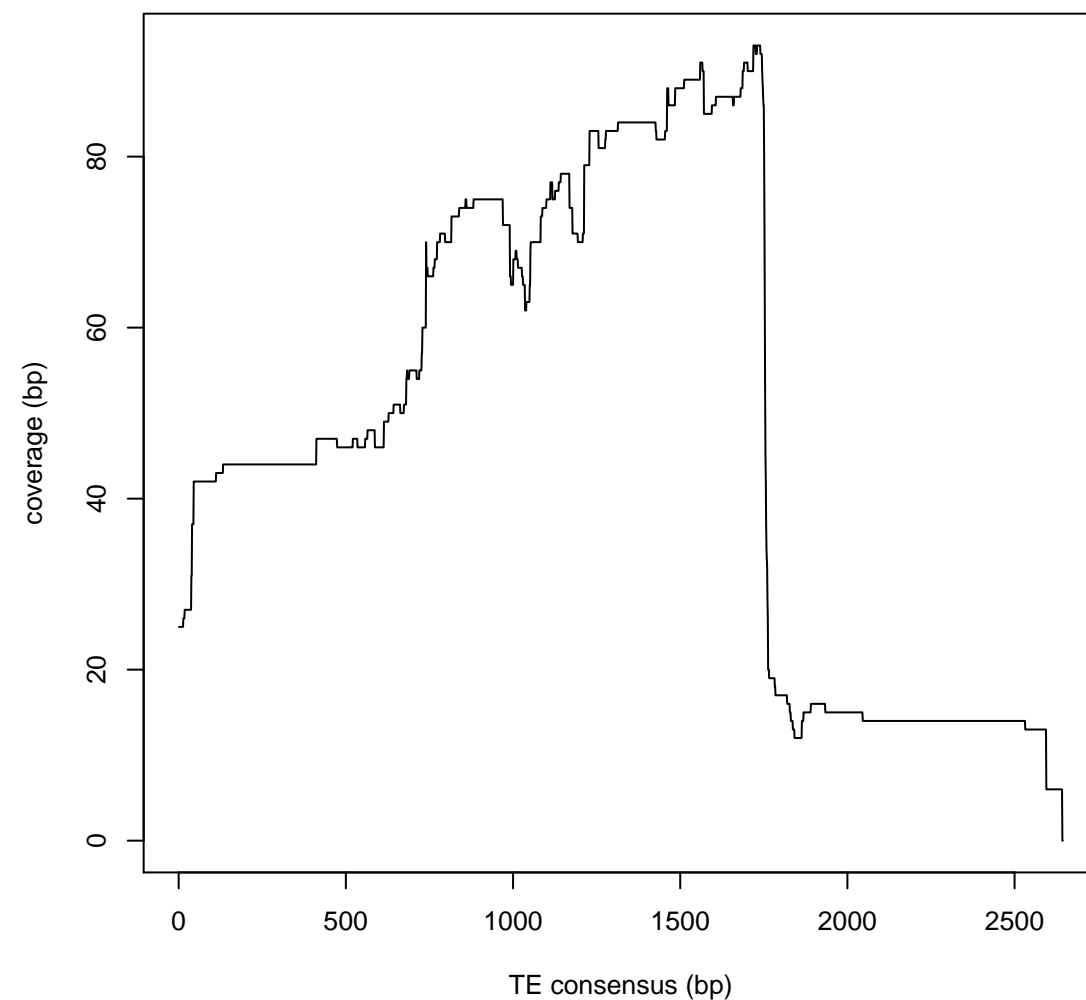


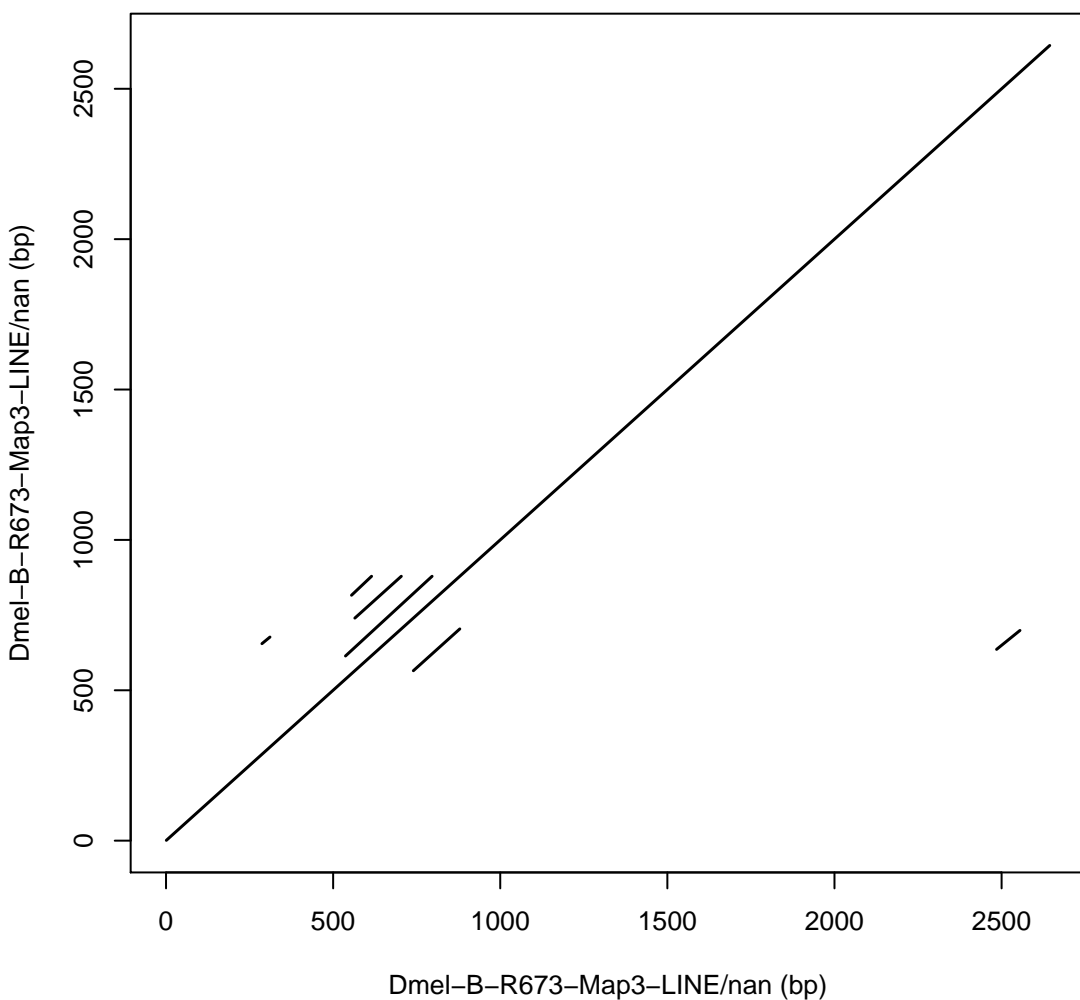
TE: Dmel-B-R673-Map3-LINE/nan
consensus size: 2644bp; fragments: 152; full length: 6 (≥ 2379.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

