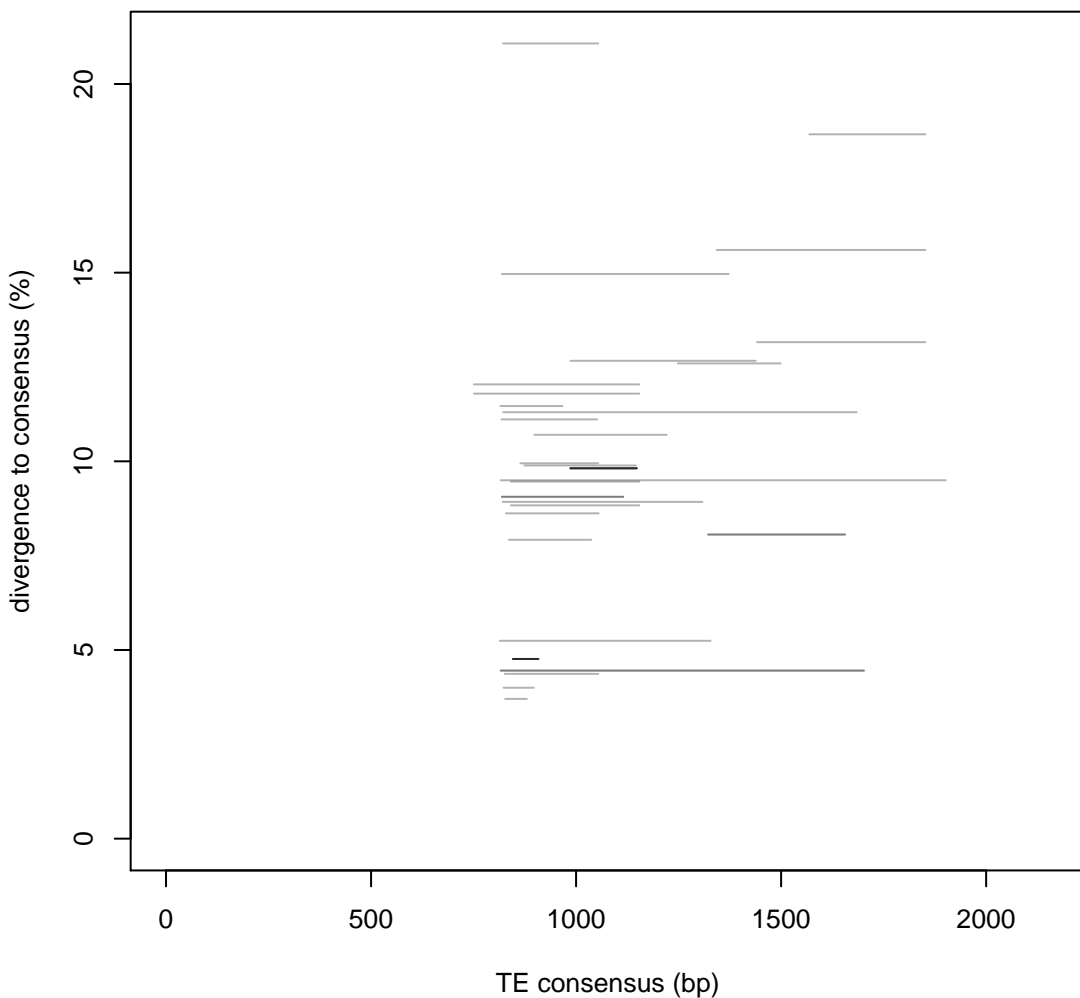
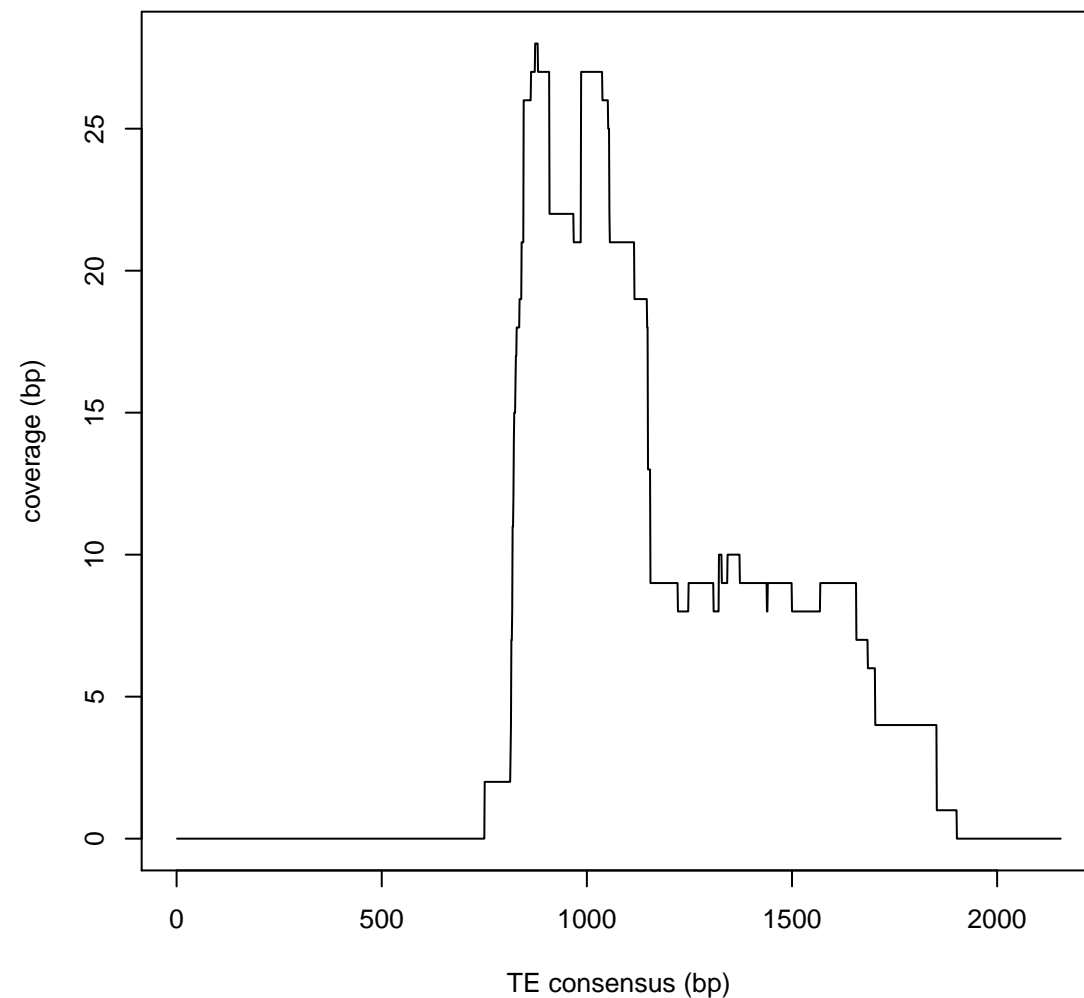


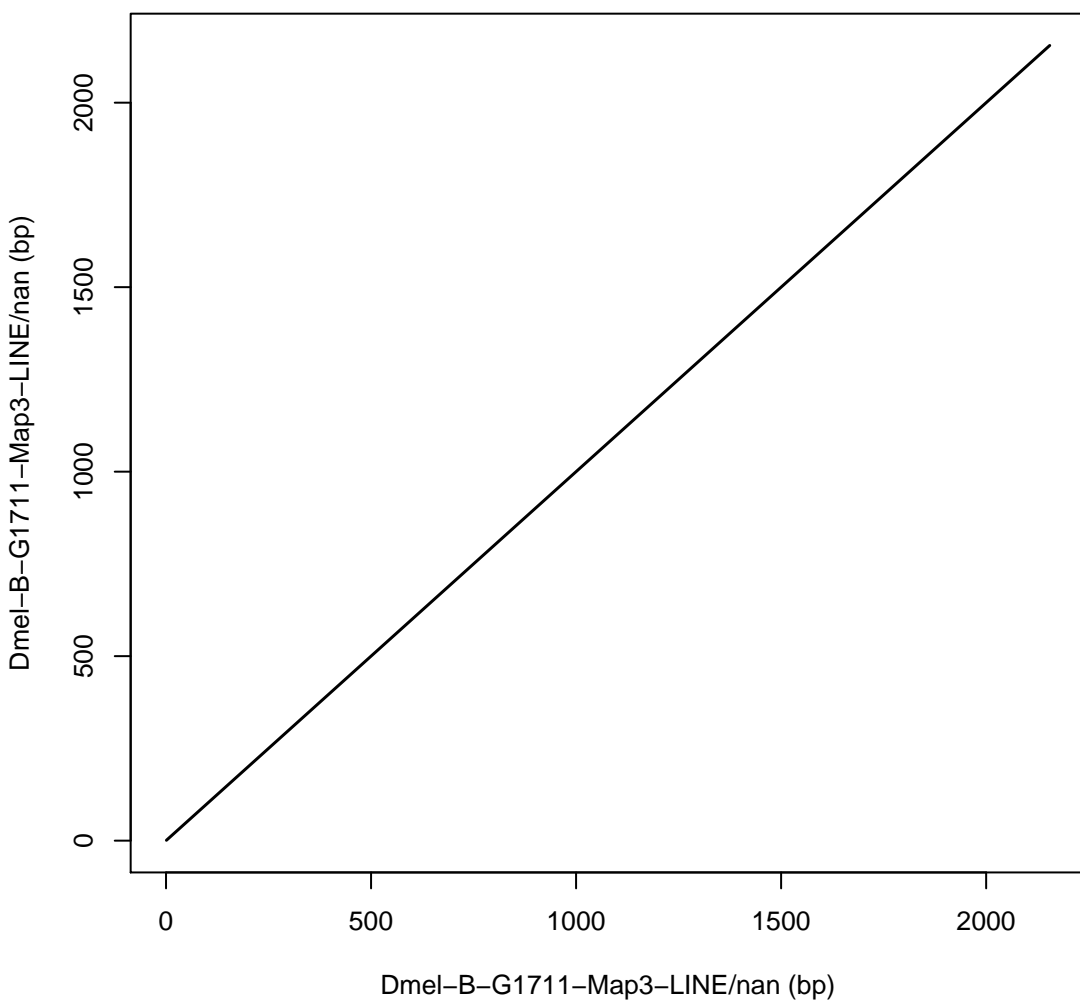
TE: Dmel-B-G1711-Map3-LINE/nan
consensus size: 2155bp; fragments: 41; full length: 0 (≥ 1939.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

