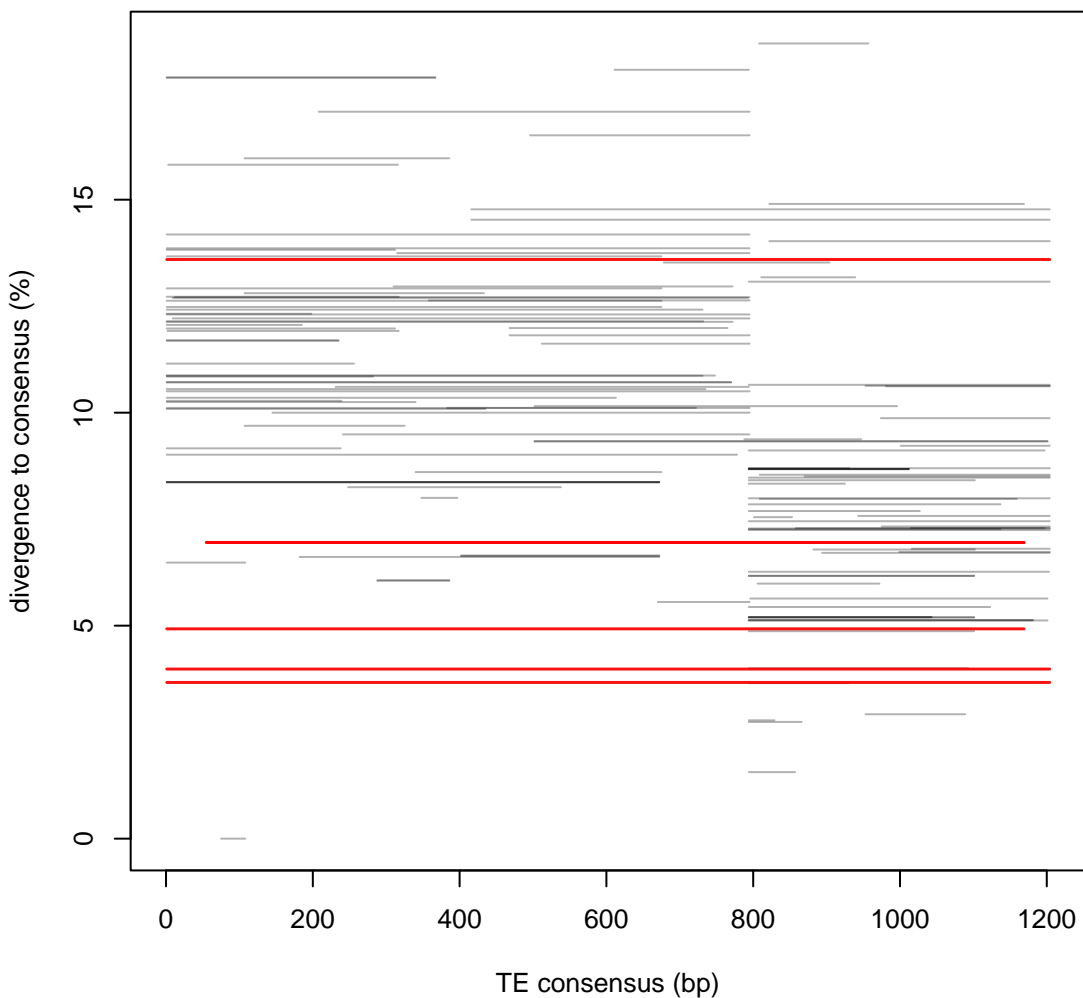
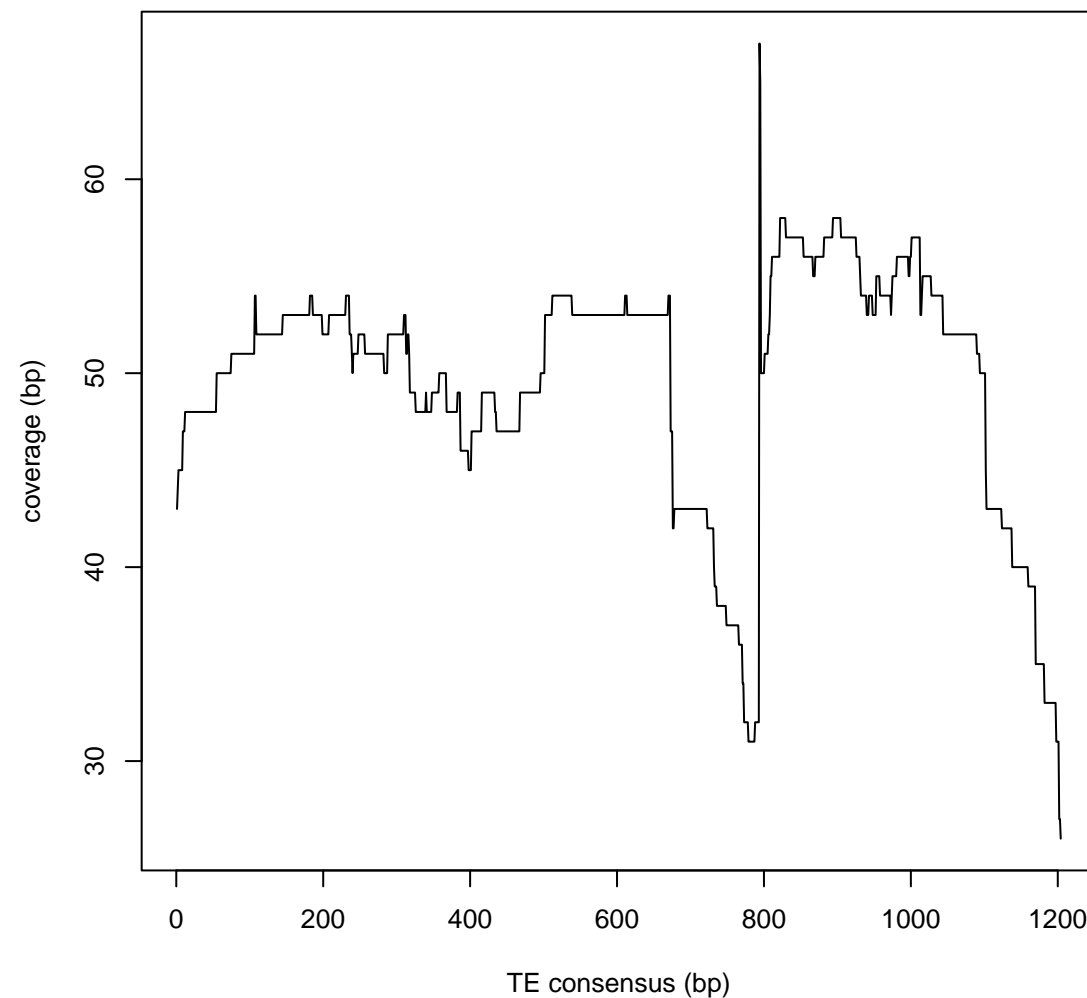


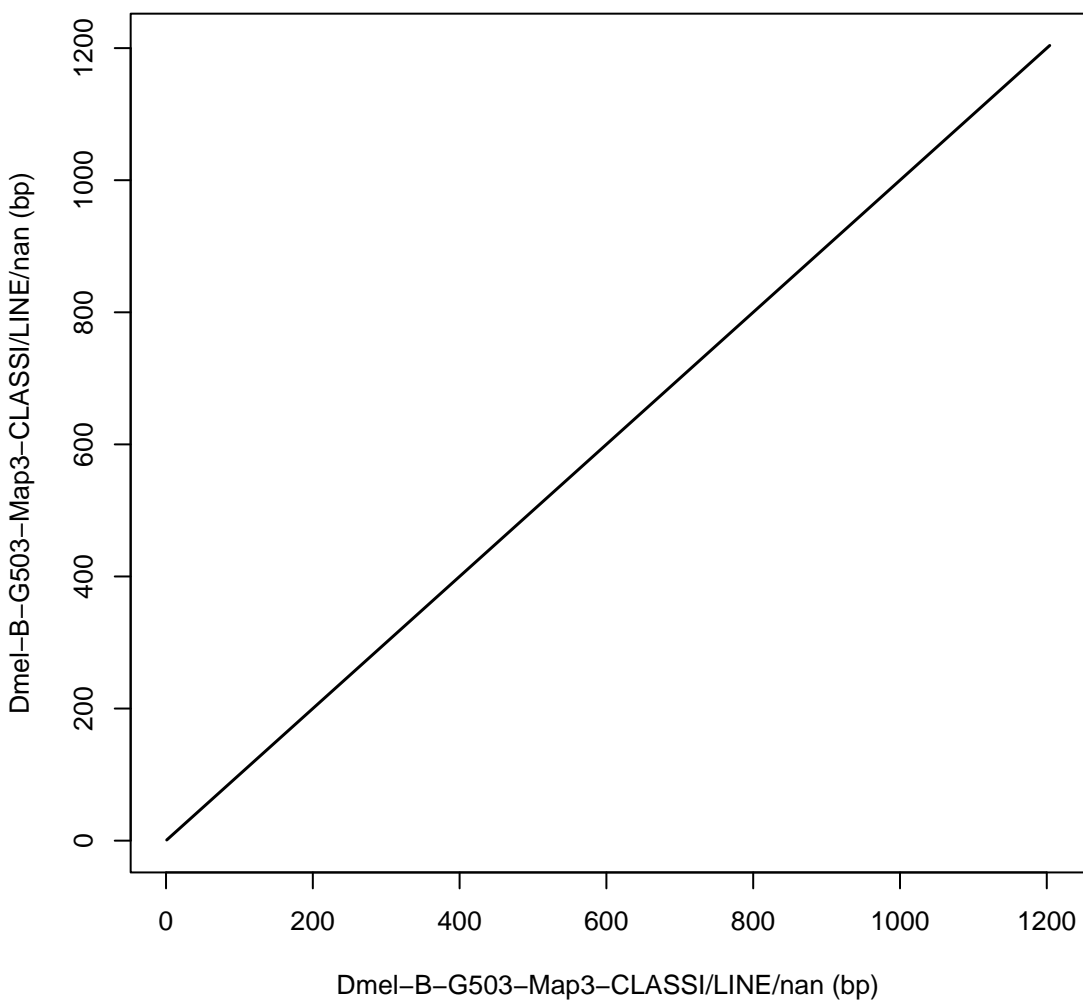
TE: Dmel-B-G503-Map3-CLASSI/LINE/nan
consensus size: 1204bp; fragments: 142; full length: 6 (≥ 1083.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

