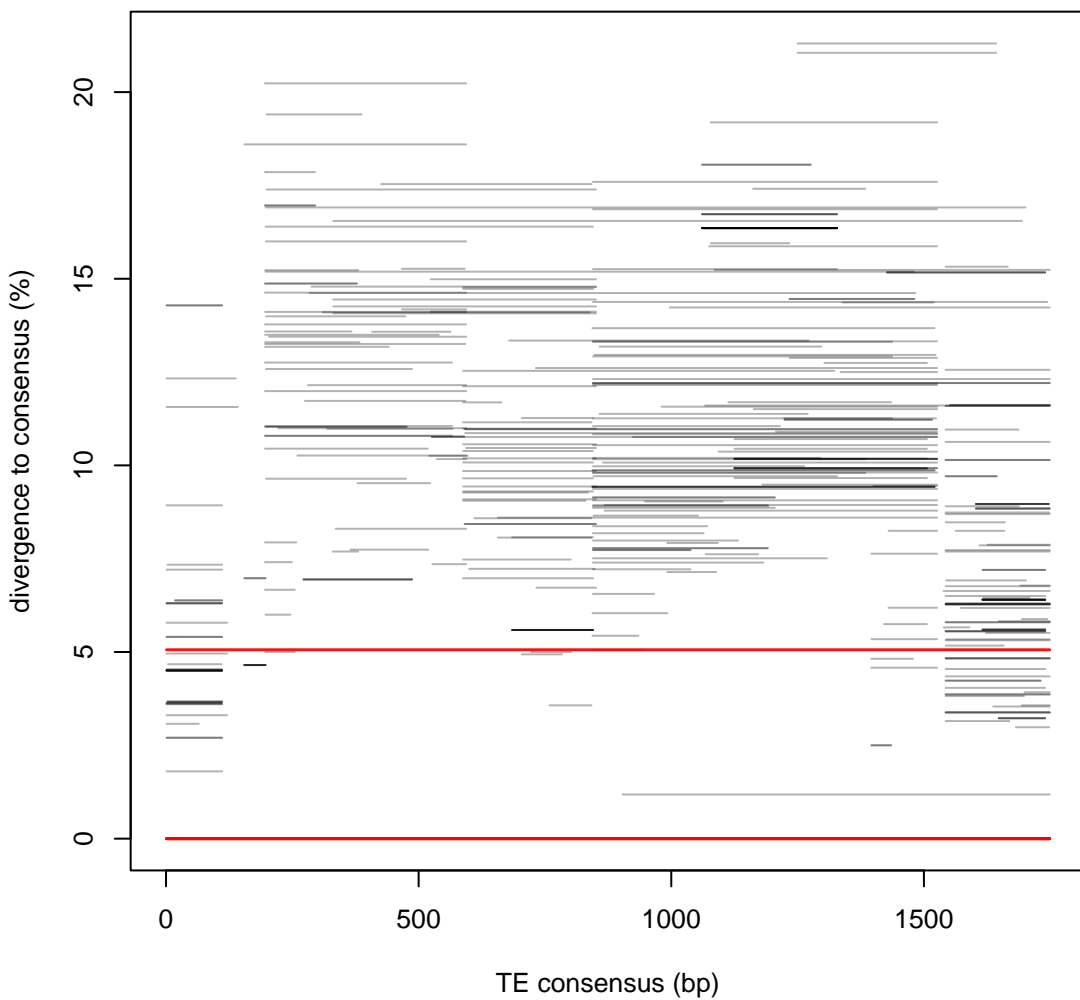
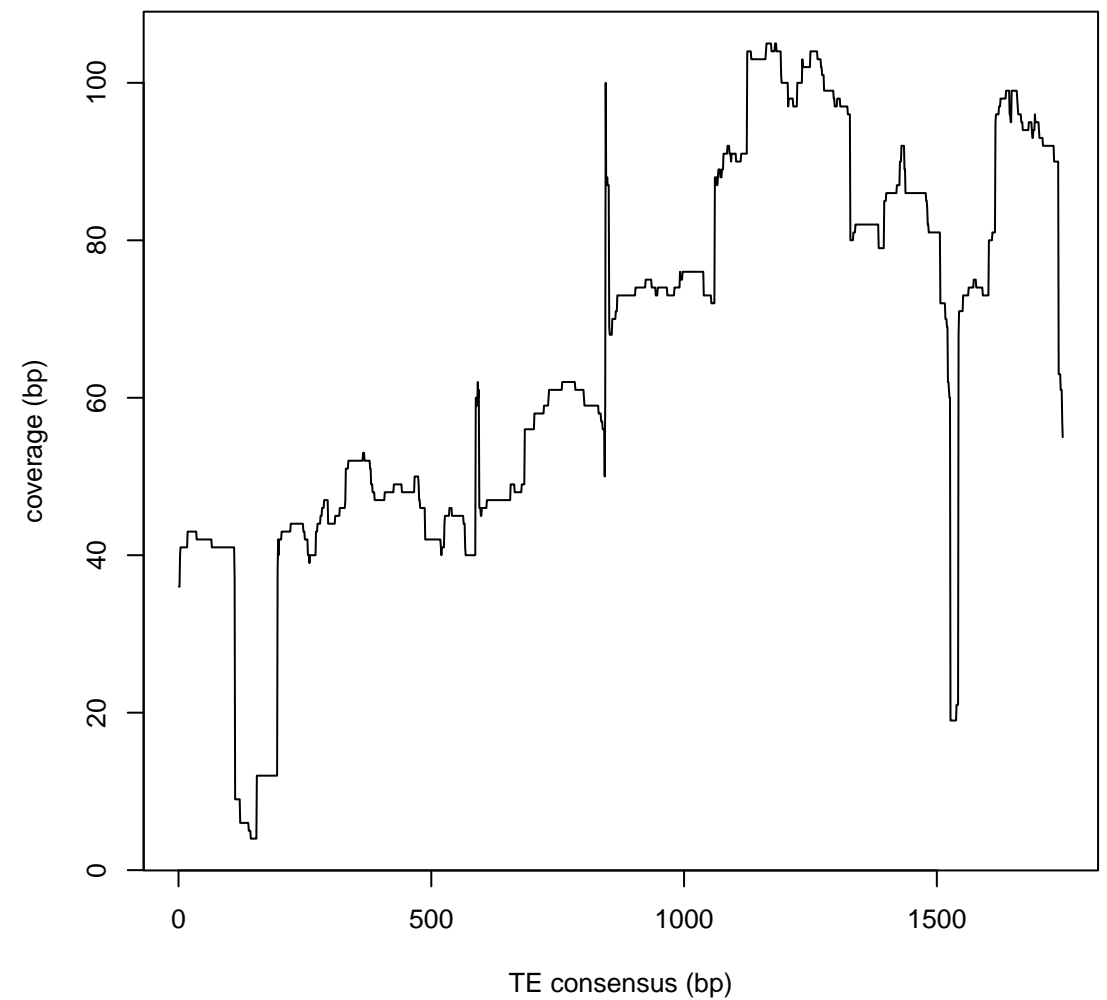


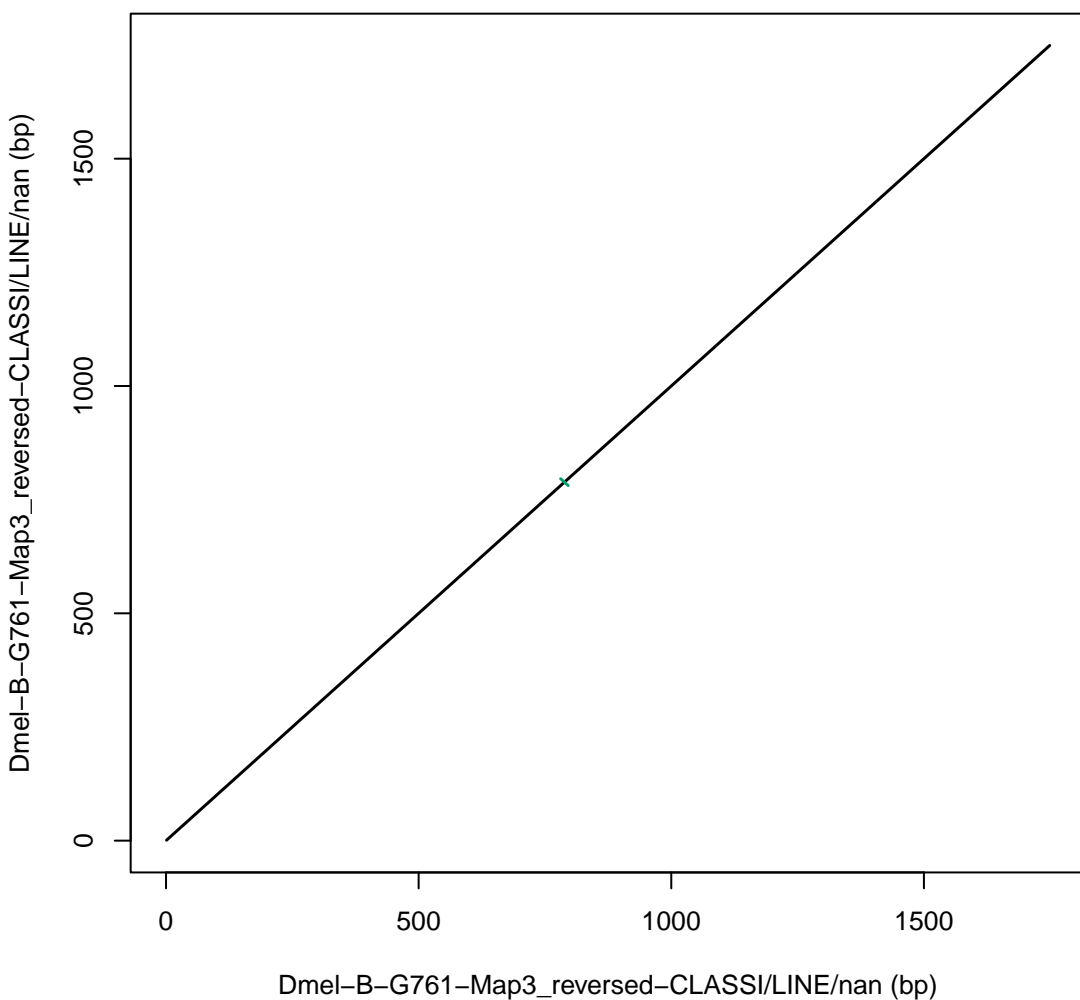
TE: Dmel-B-G761-Map3_reversed-CLASSI/LINE/nan
consensus size: 1749bp; fragments: 386; full length: 4 (≥ 1574.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

