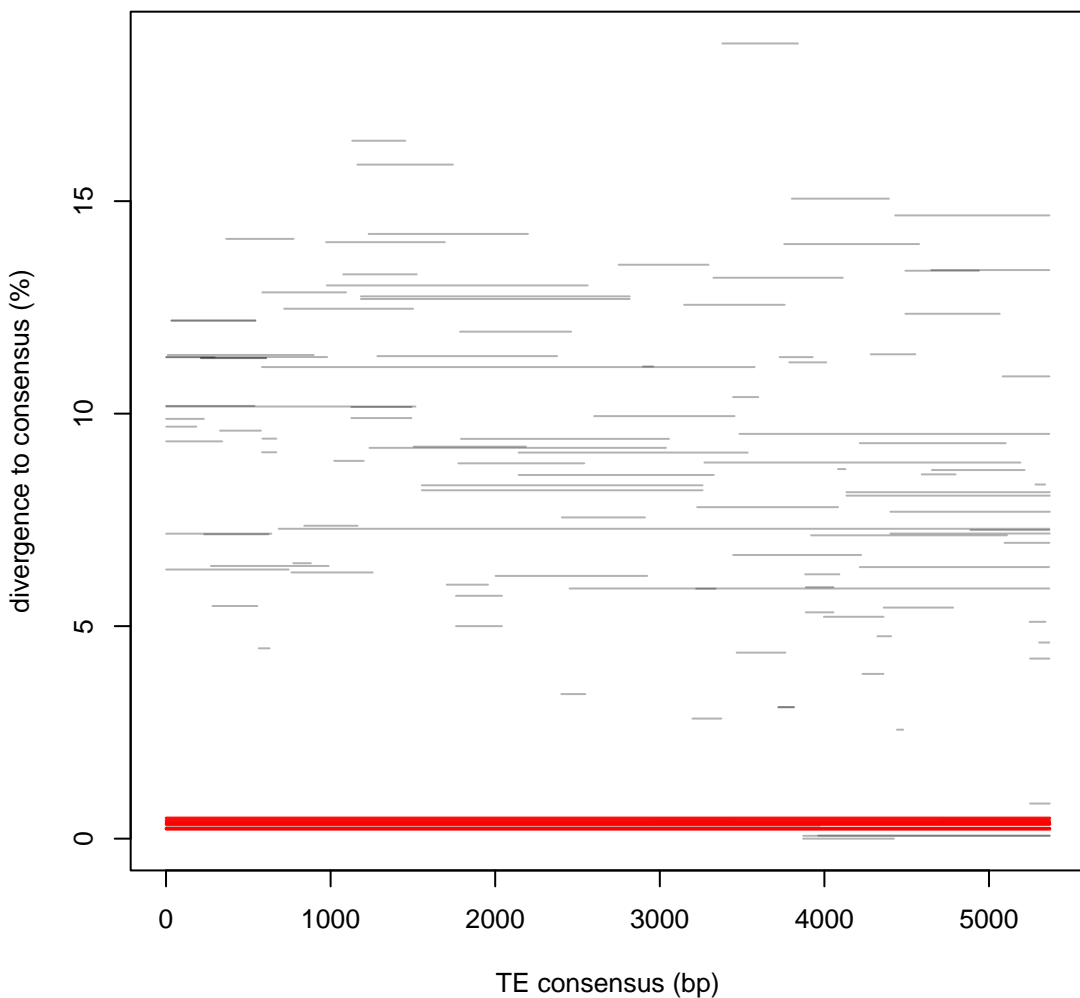
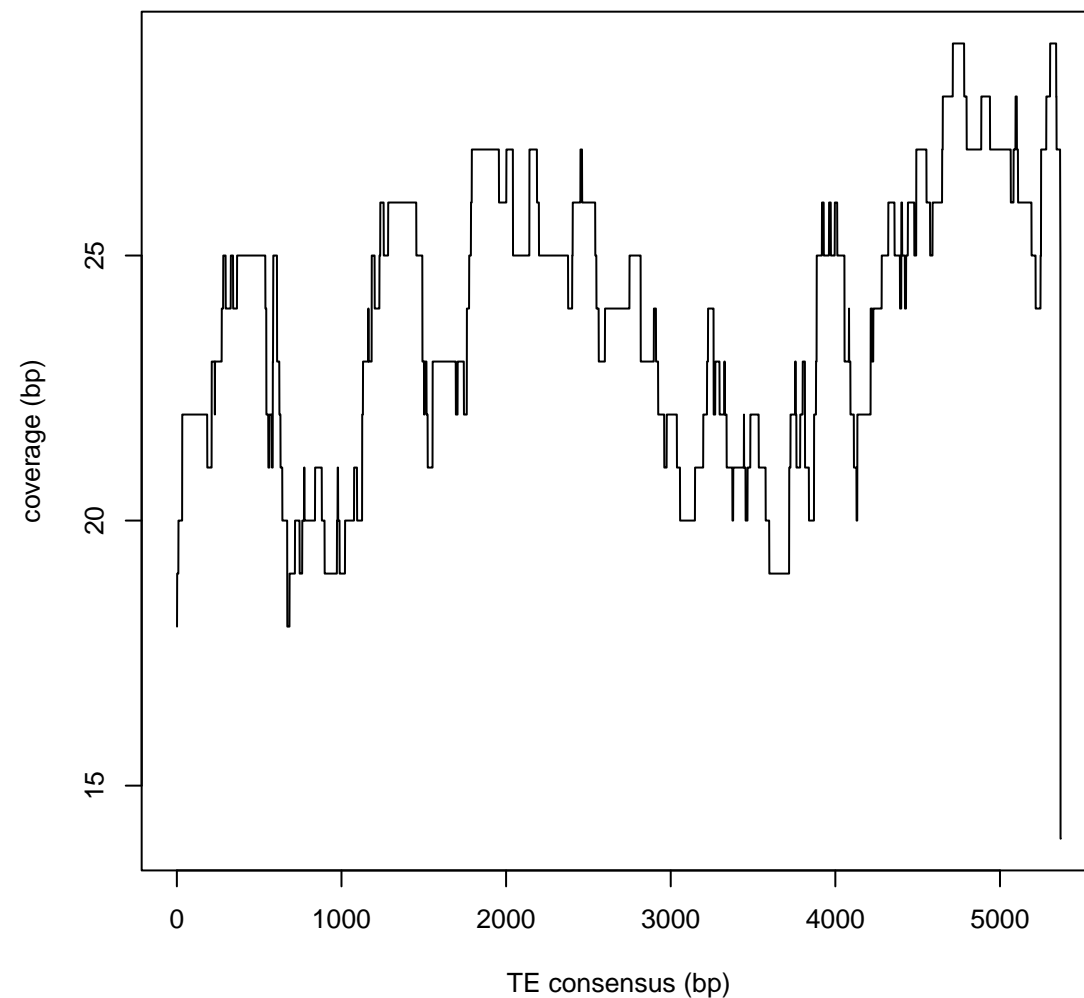


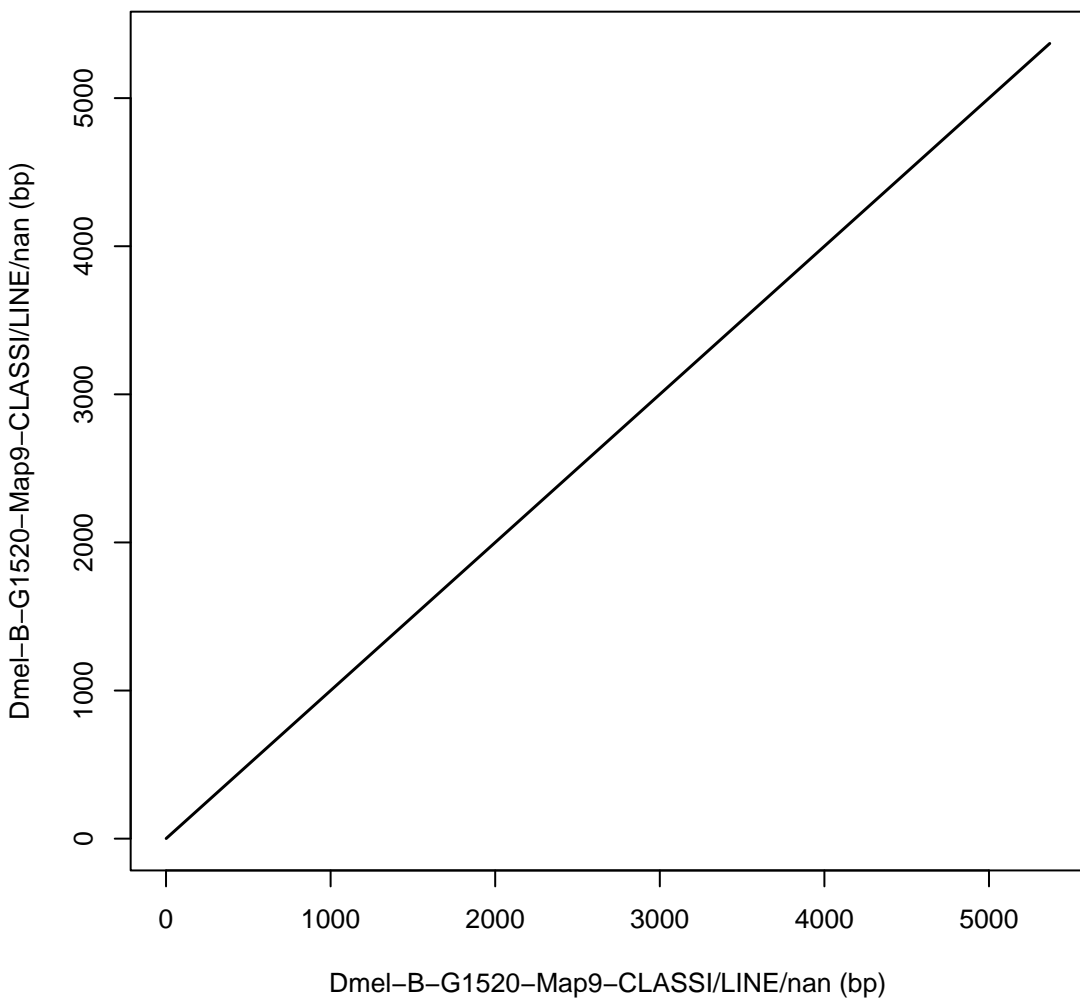
TE: Dmel-B-G1520-Map9-CLASSI/LINE/nan
consensus size: 5369bp; fragments: 123; full length: 8 (>=4832.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

