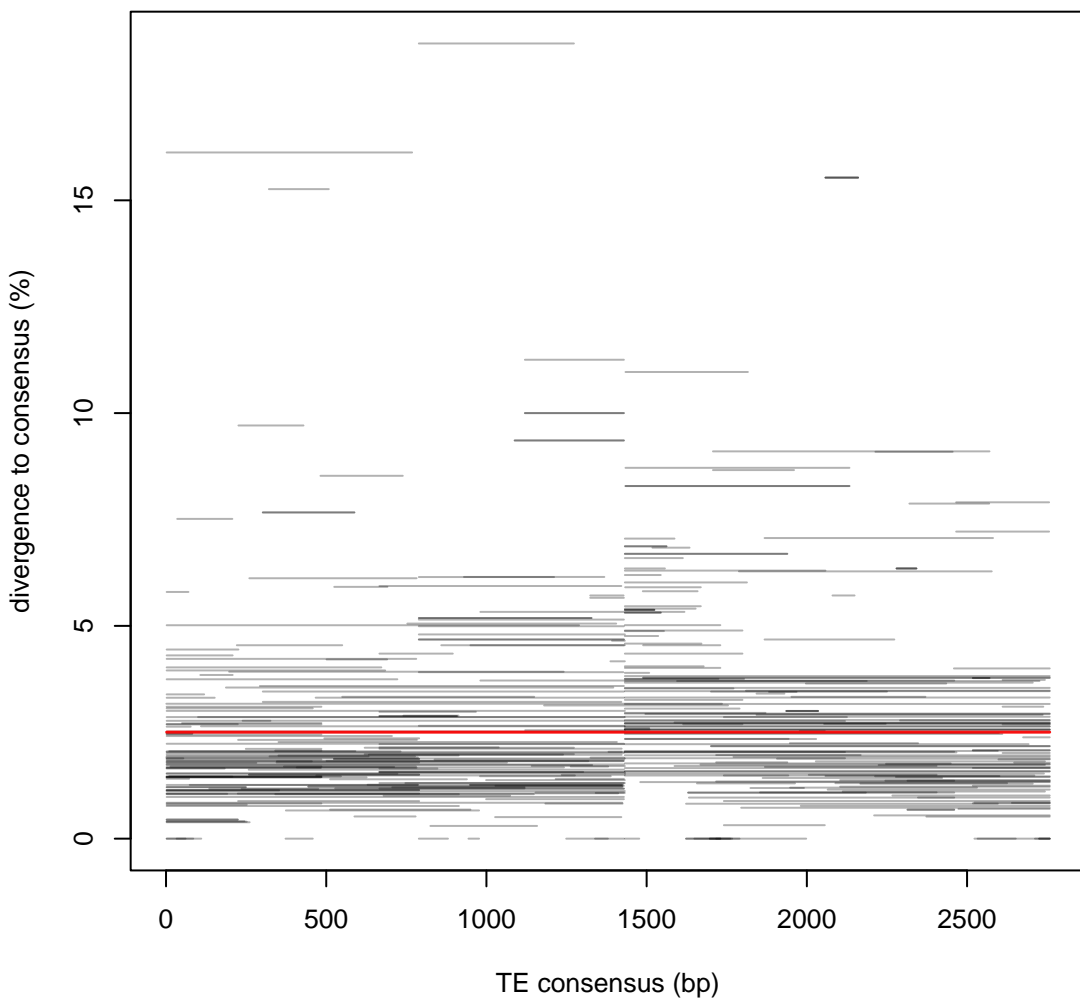
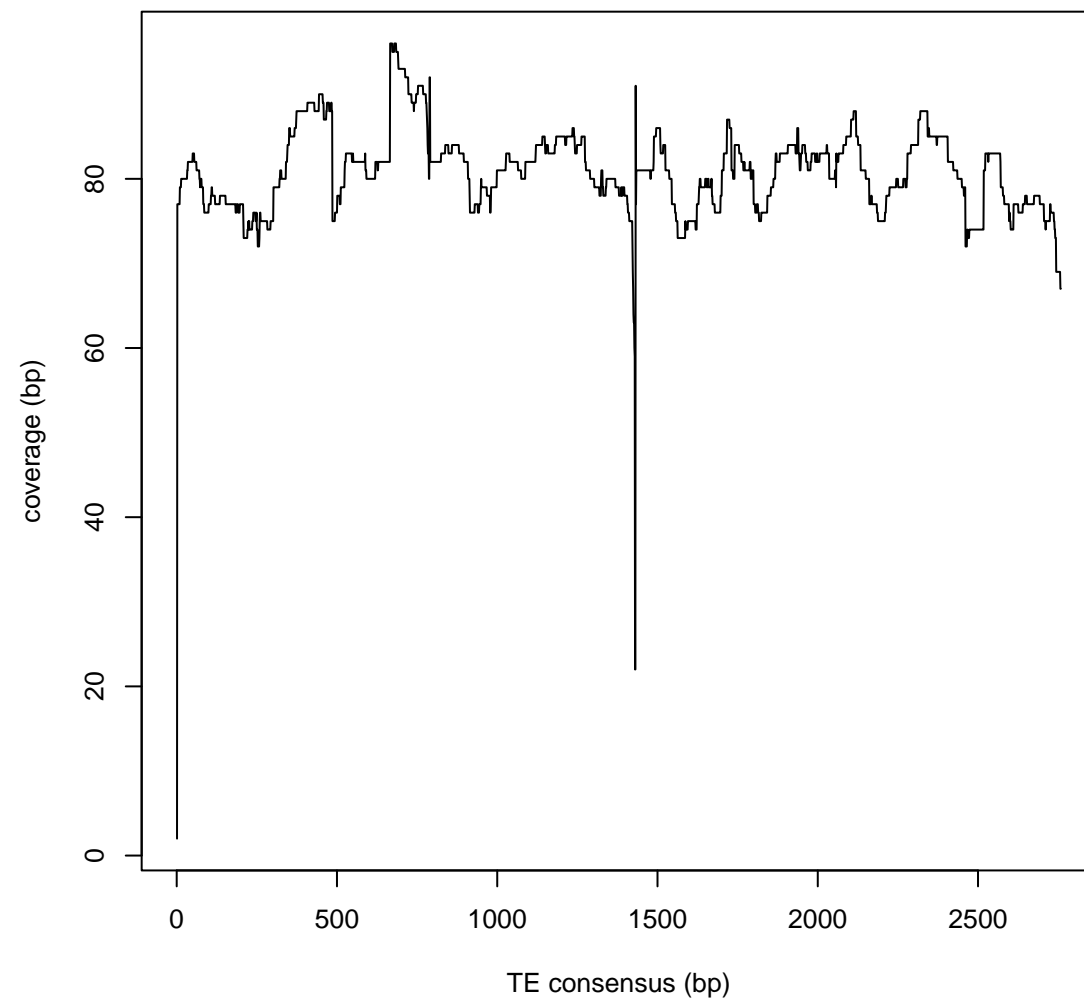


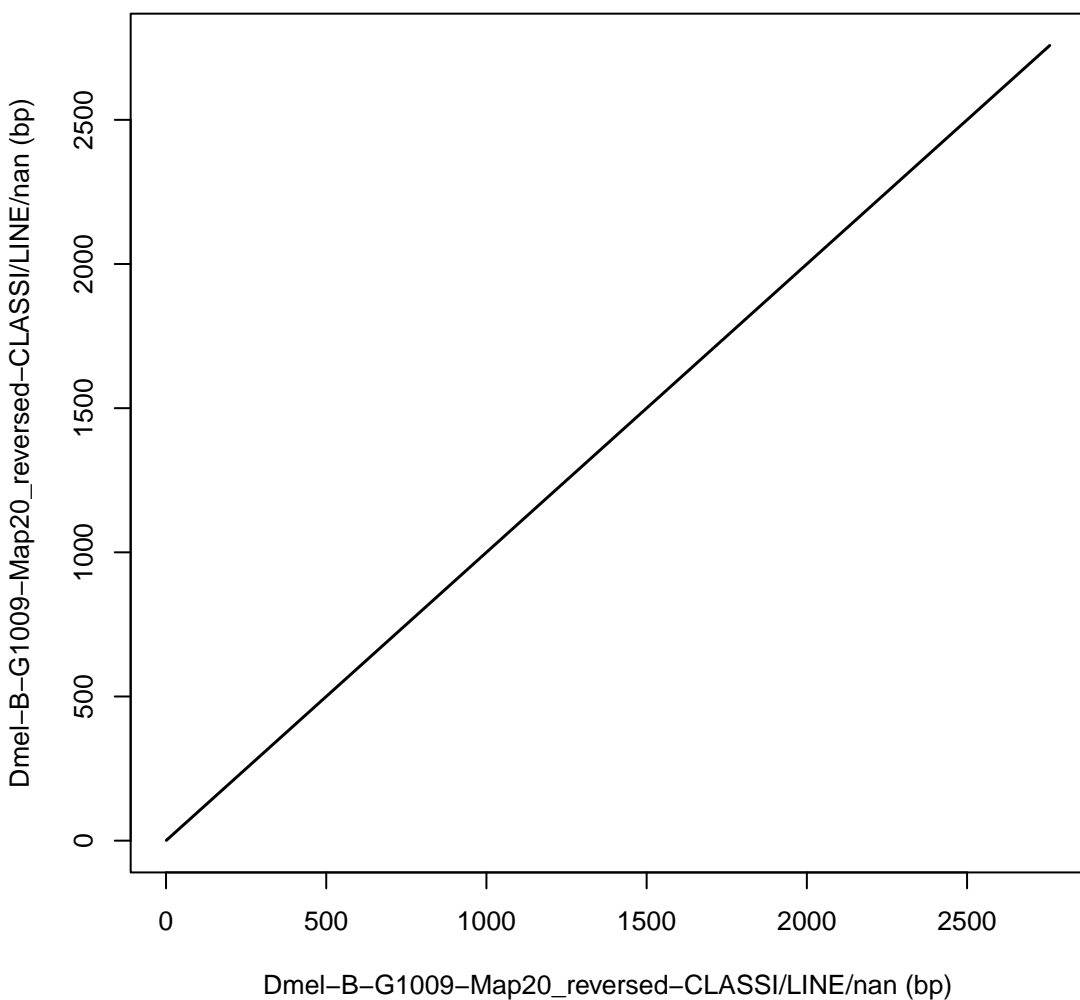
TE: Dmel-B-G1009-Map20_reversed-CLASSI/LINE/nan
consensus size: 2758bp; fragments: 398; full length: 1 (>=2482.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

