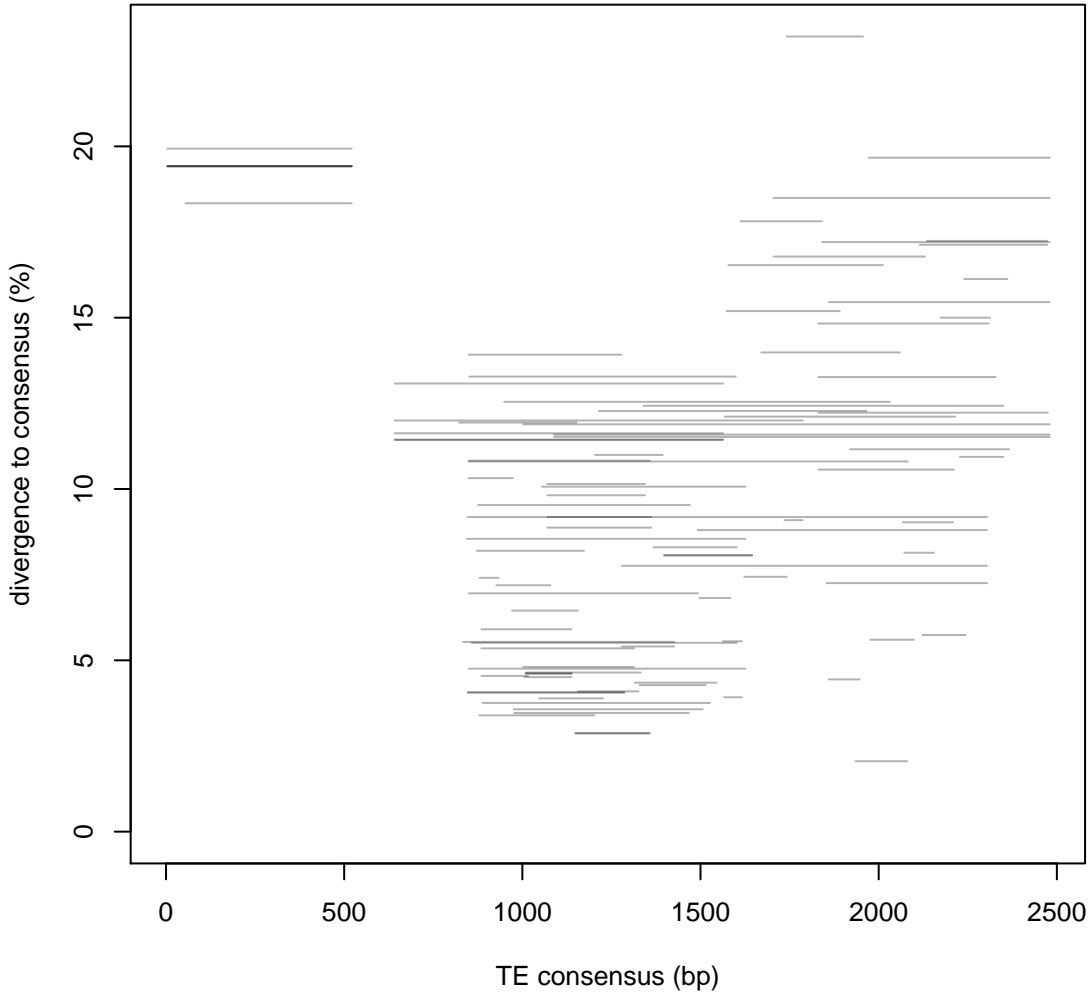
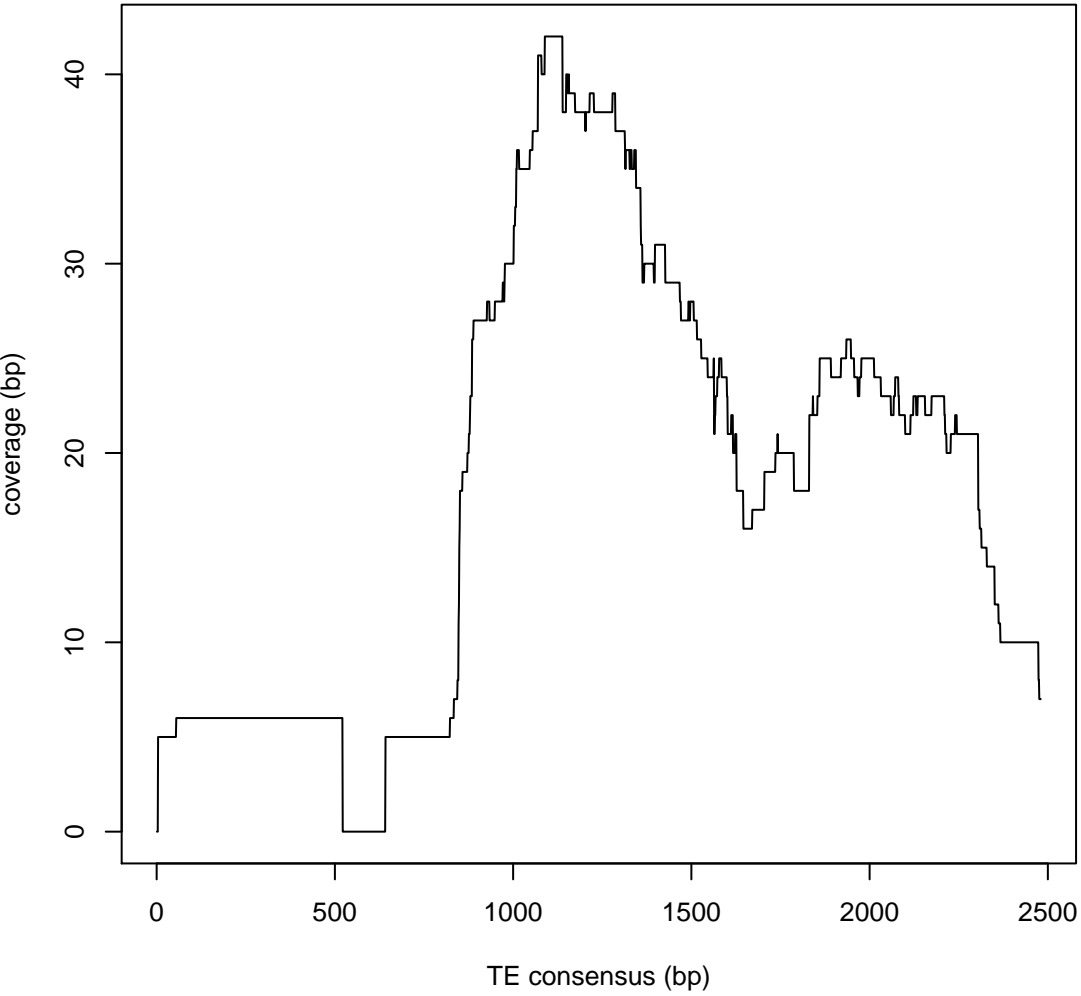


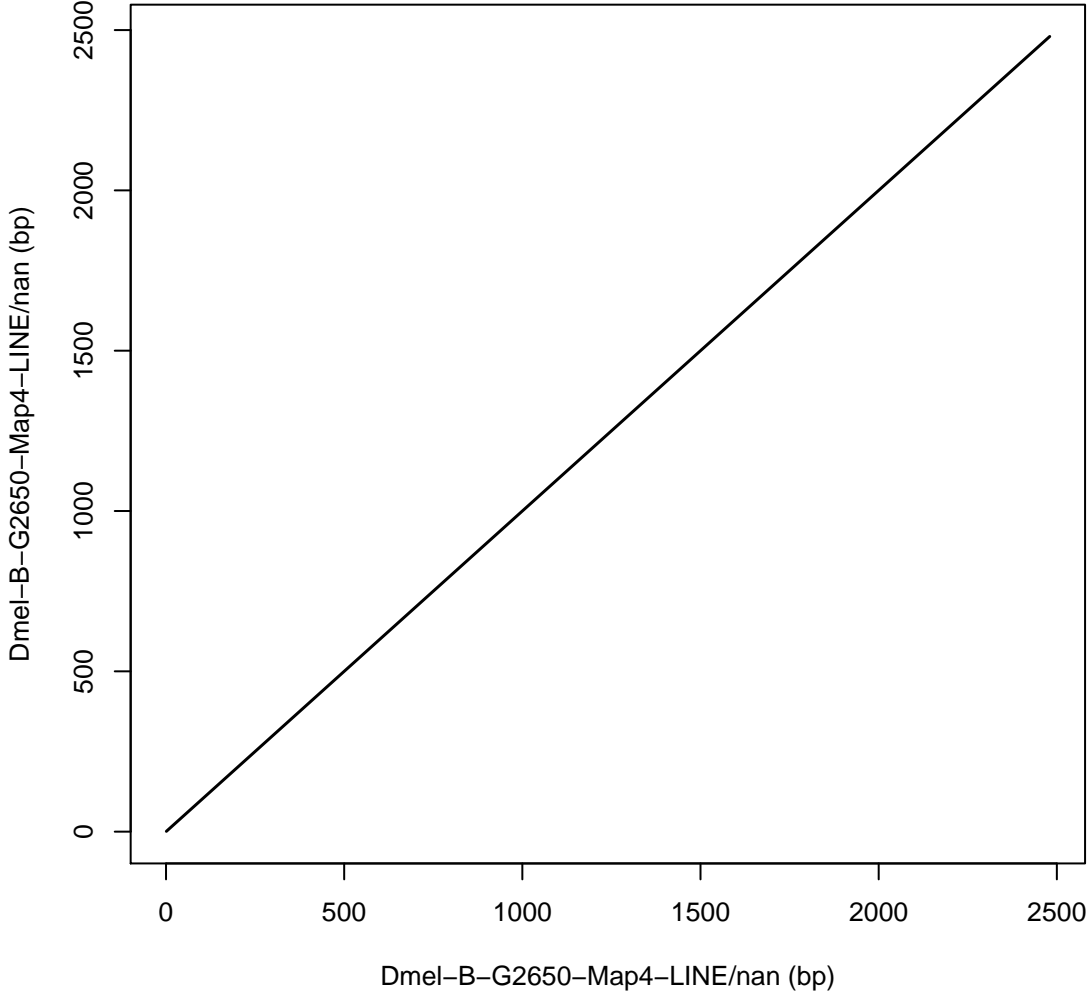
TE: Dmel-B-G2650-Map4-LINE/nan
consensus size: 2480bp; fragments: 99; full length: 0 (>=2232bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

