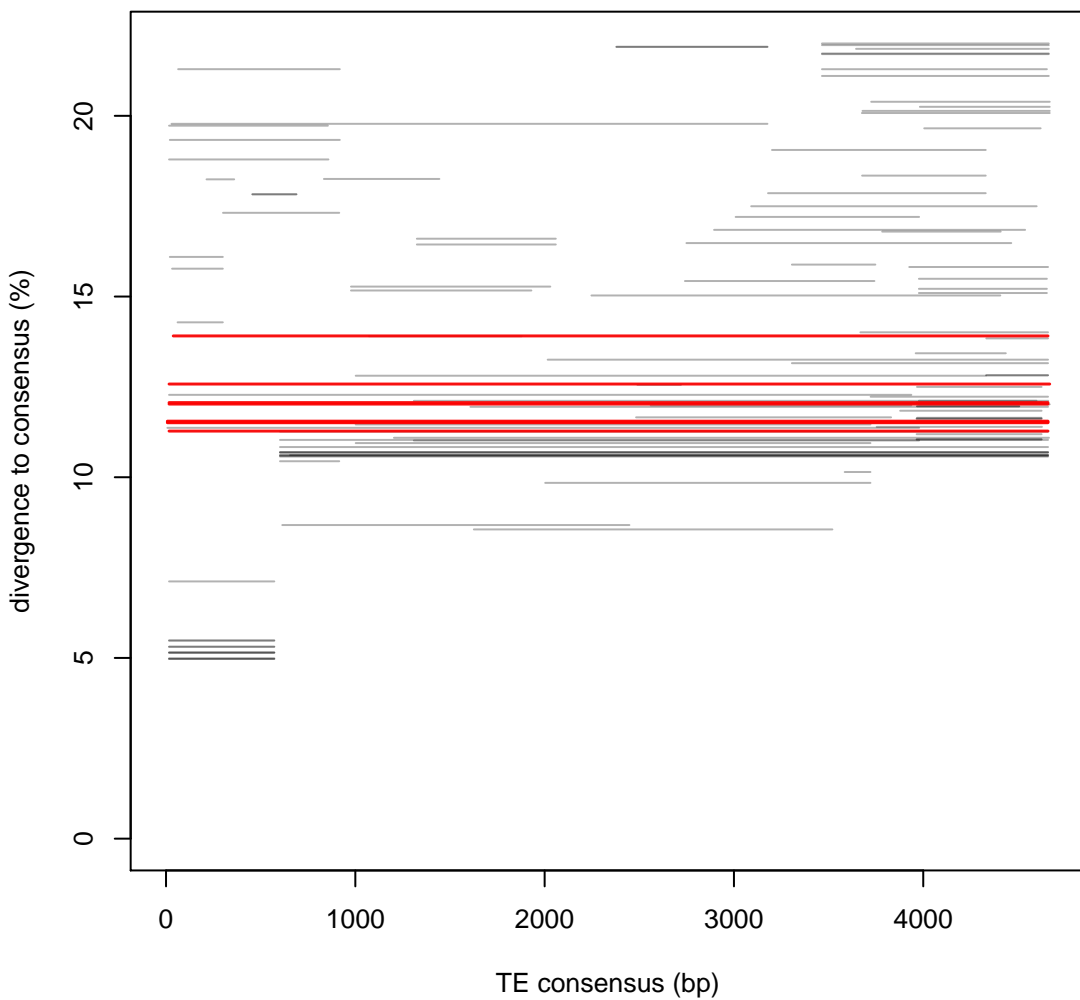
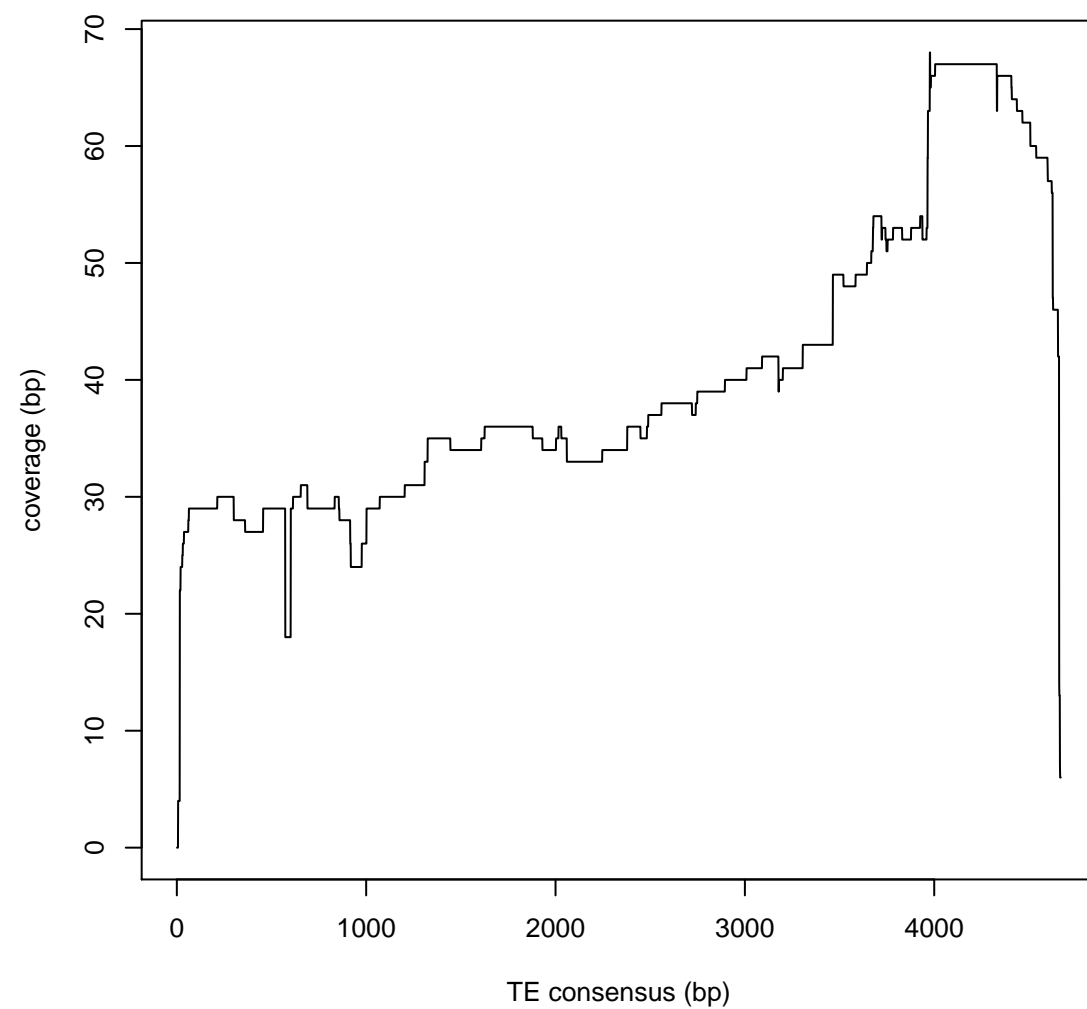


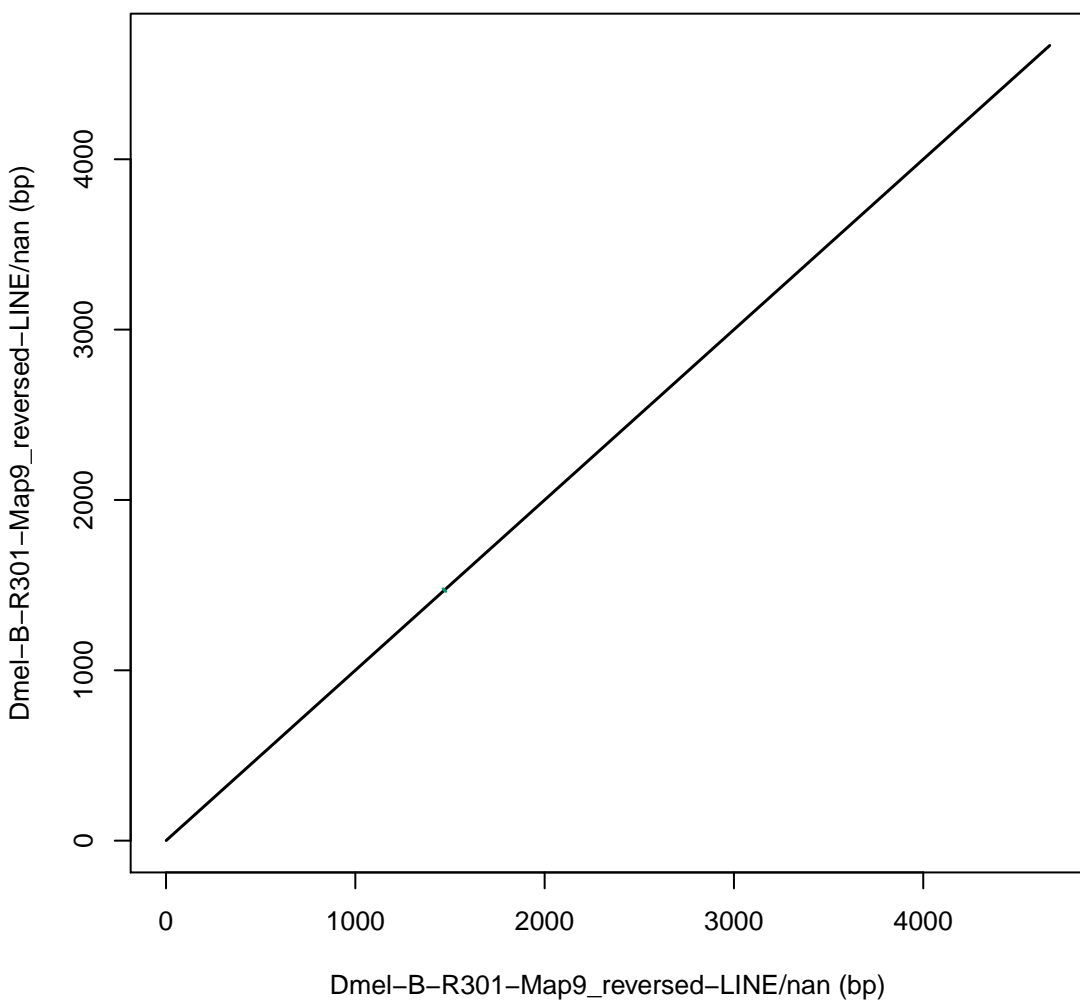
TE: Dmel-B-R301-Map9_reversed-LINE/nan
consensus size: 4668bp; fragments: 117; full length: 8 (>=4201.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

