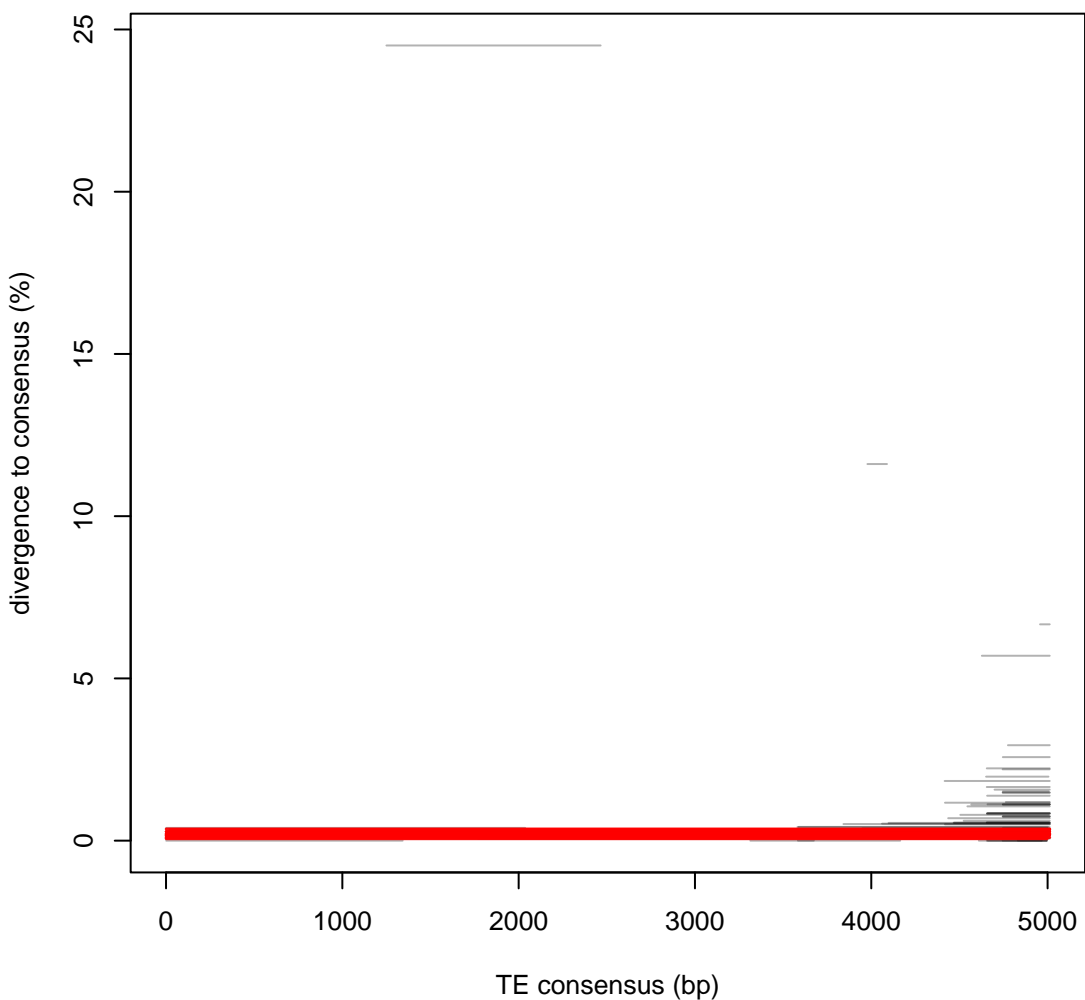
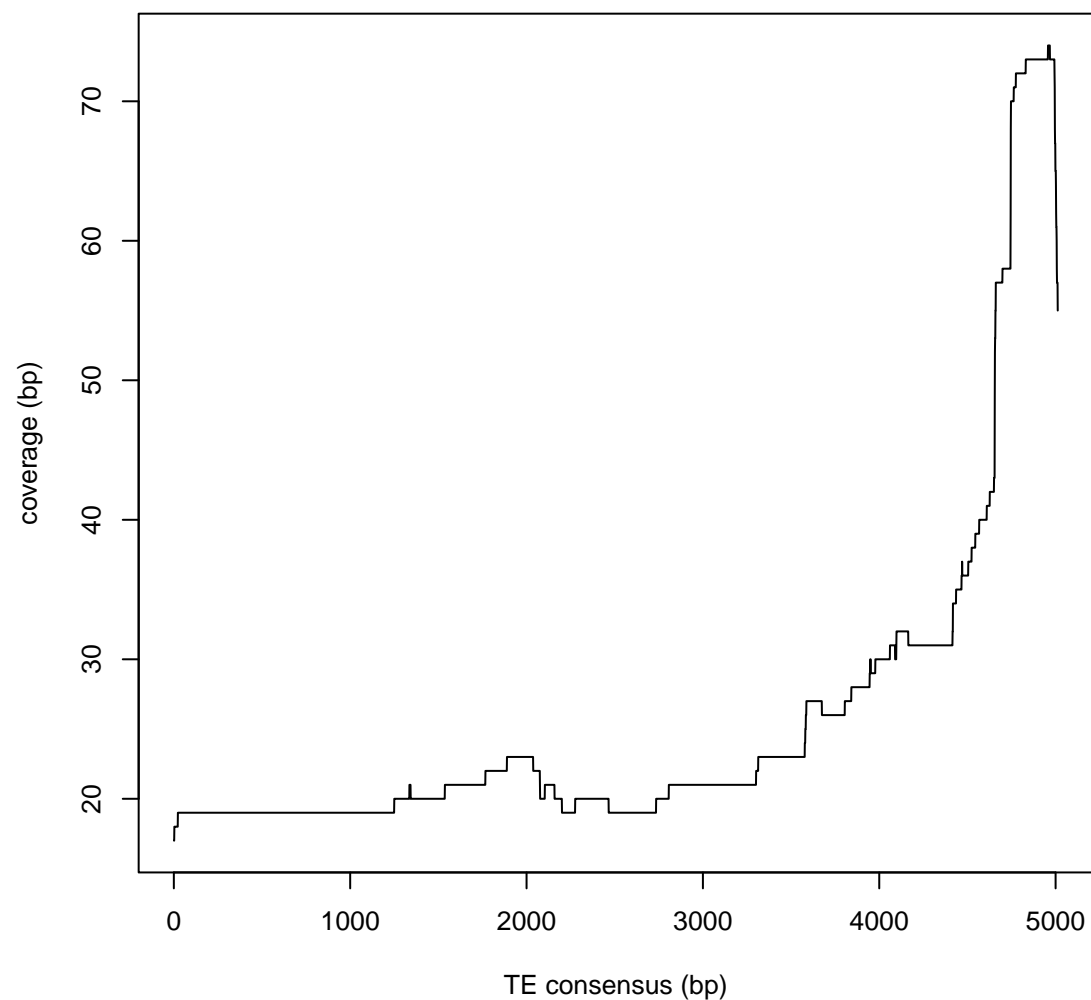


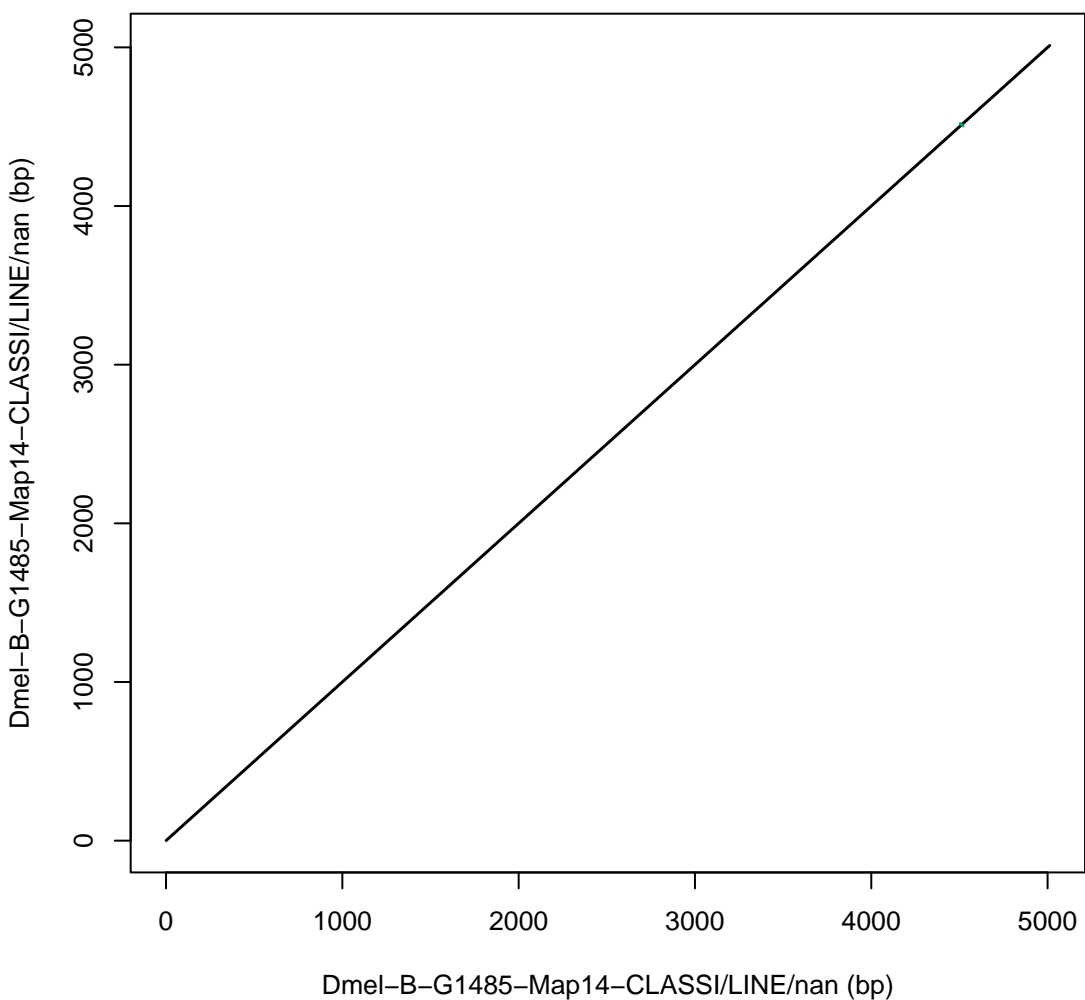
TE: Dmel-B-G1485-Map14-CLASSI/LINE/nan
consensus size: 5012bp; fragments: 86; full length: 12 (≥ 4510.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

