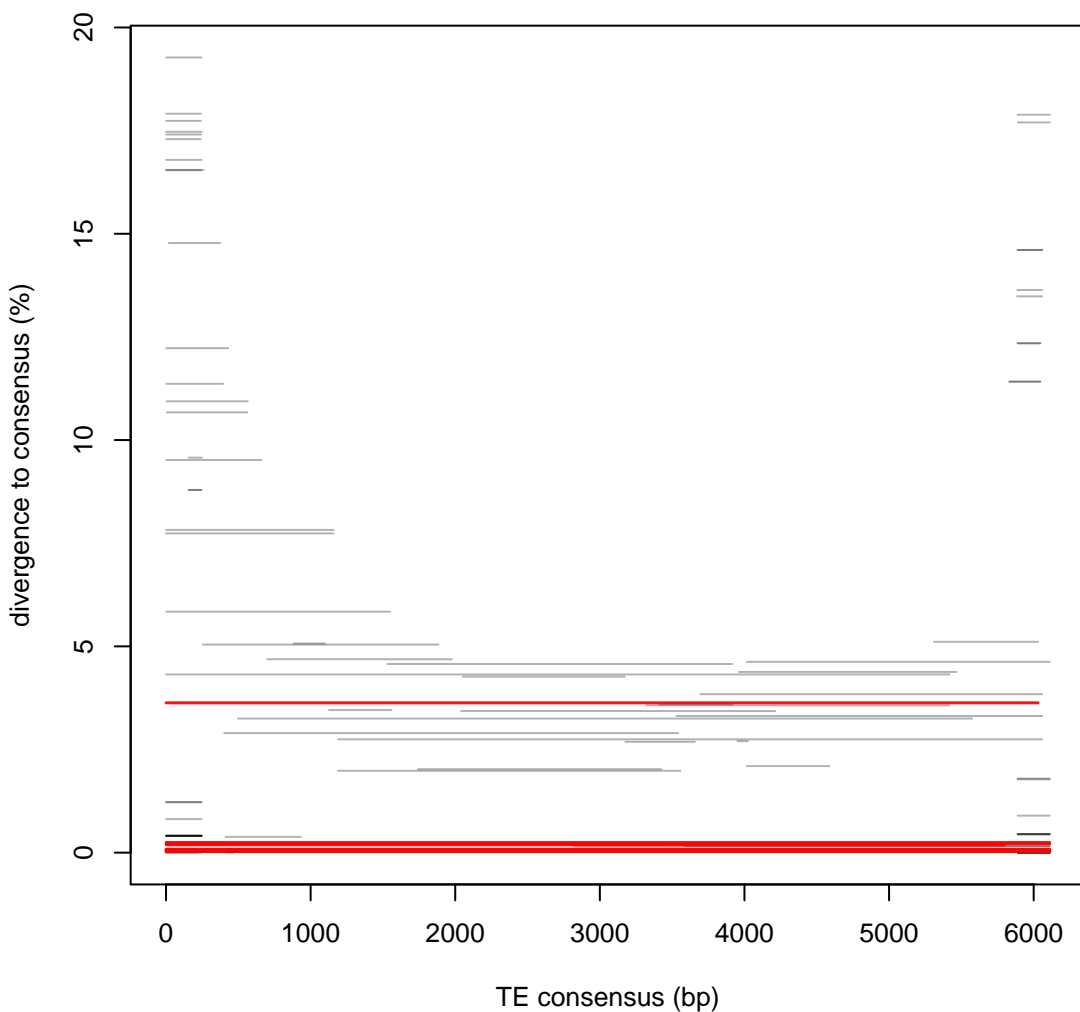
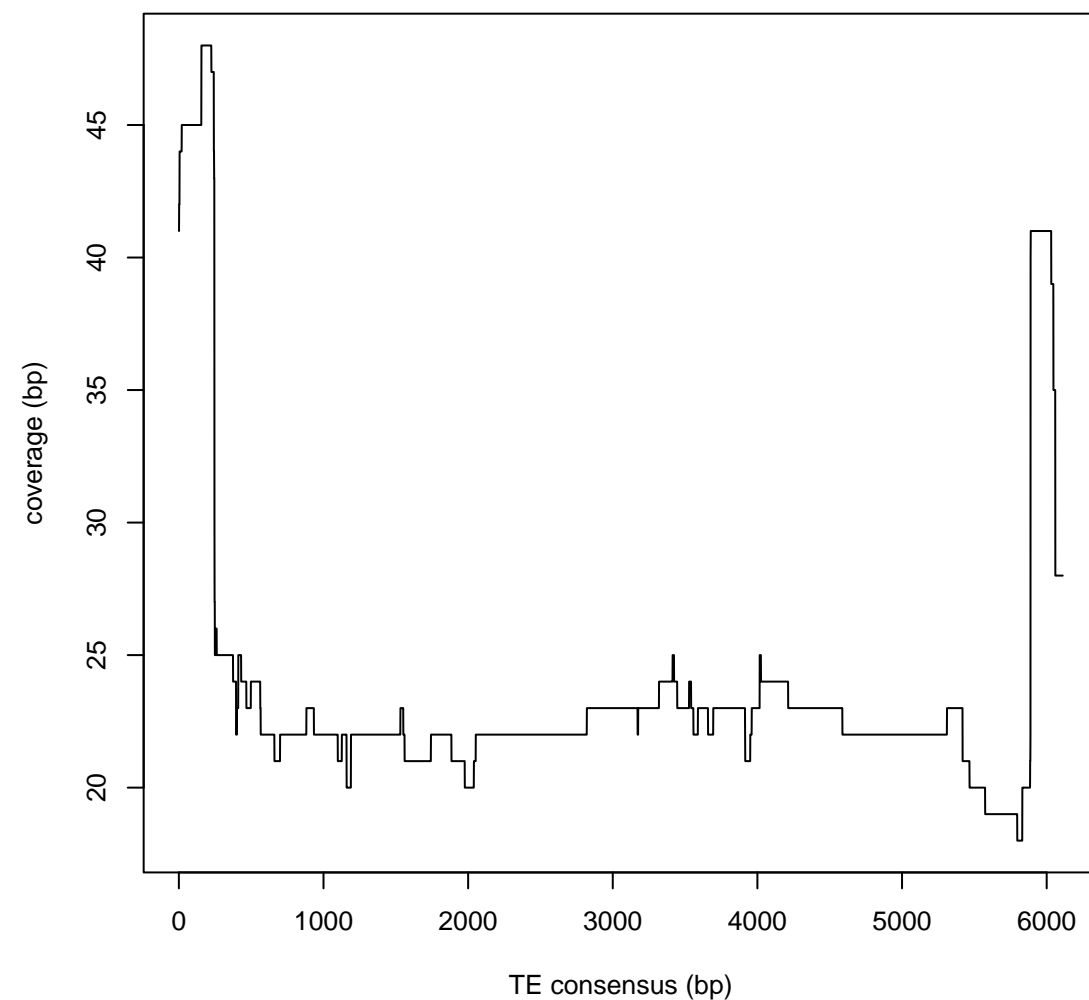


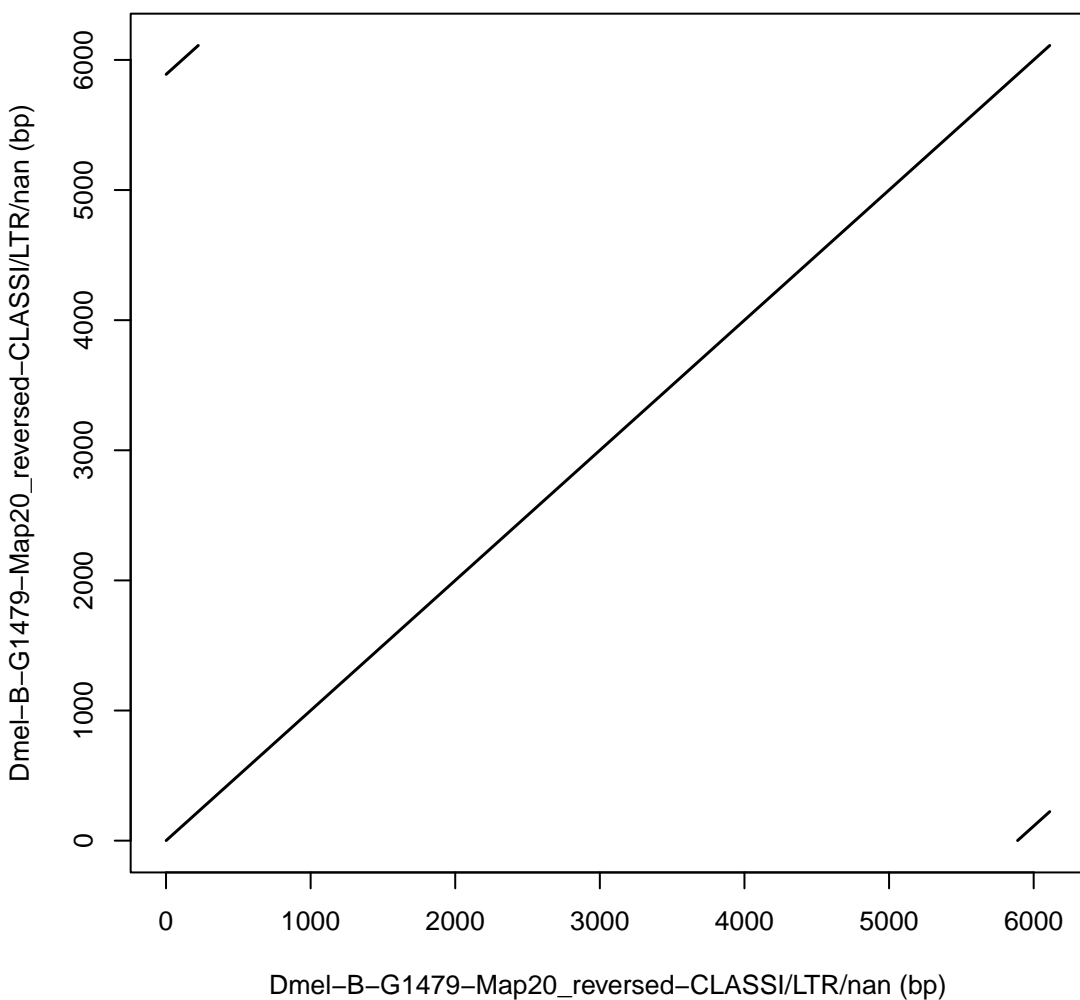
TE: Dmel-B-G1479-Map20_reversed-CLASSI/LTR/nan
consensus size: 6111bp; fragments: 97; full length: 12 (>=5499.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

