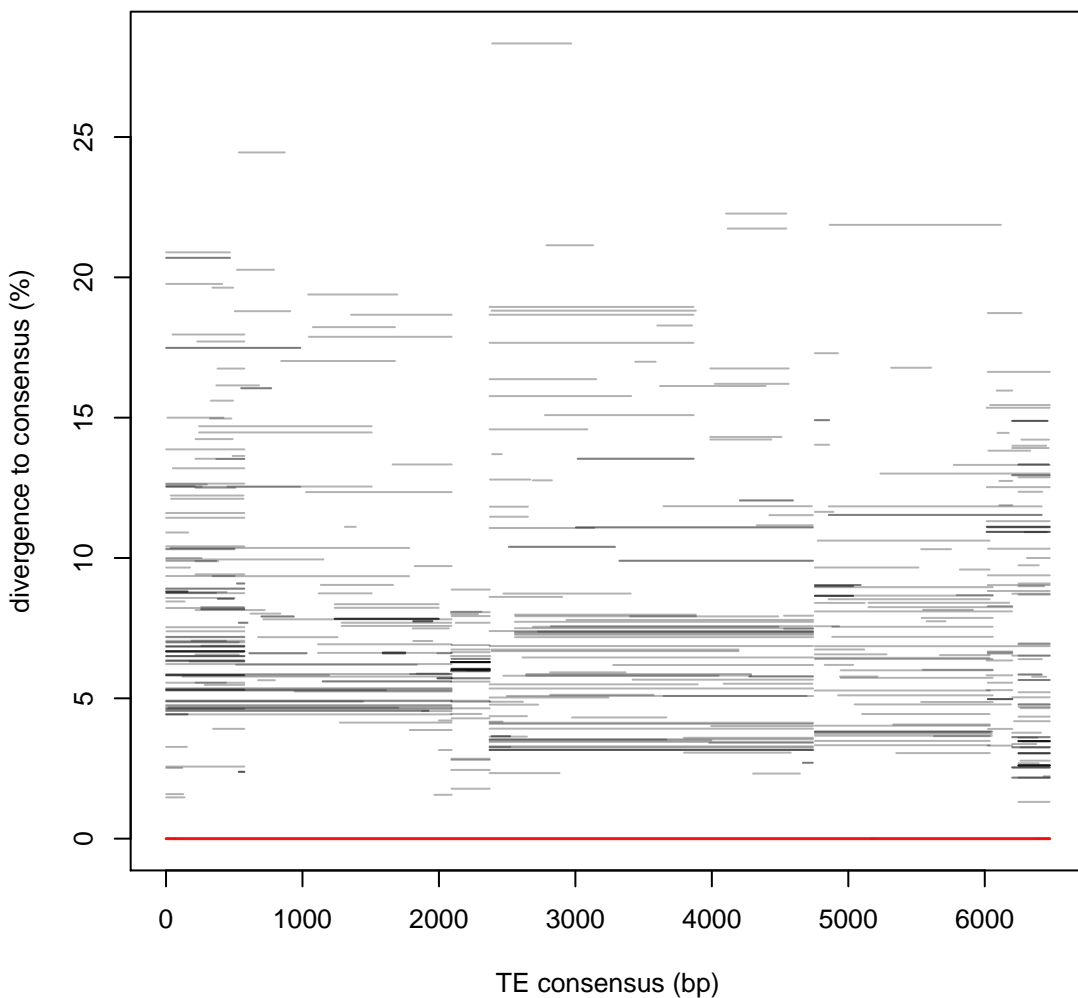
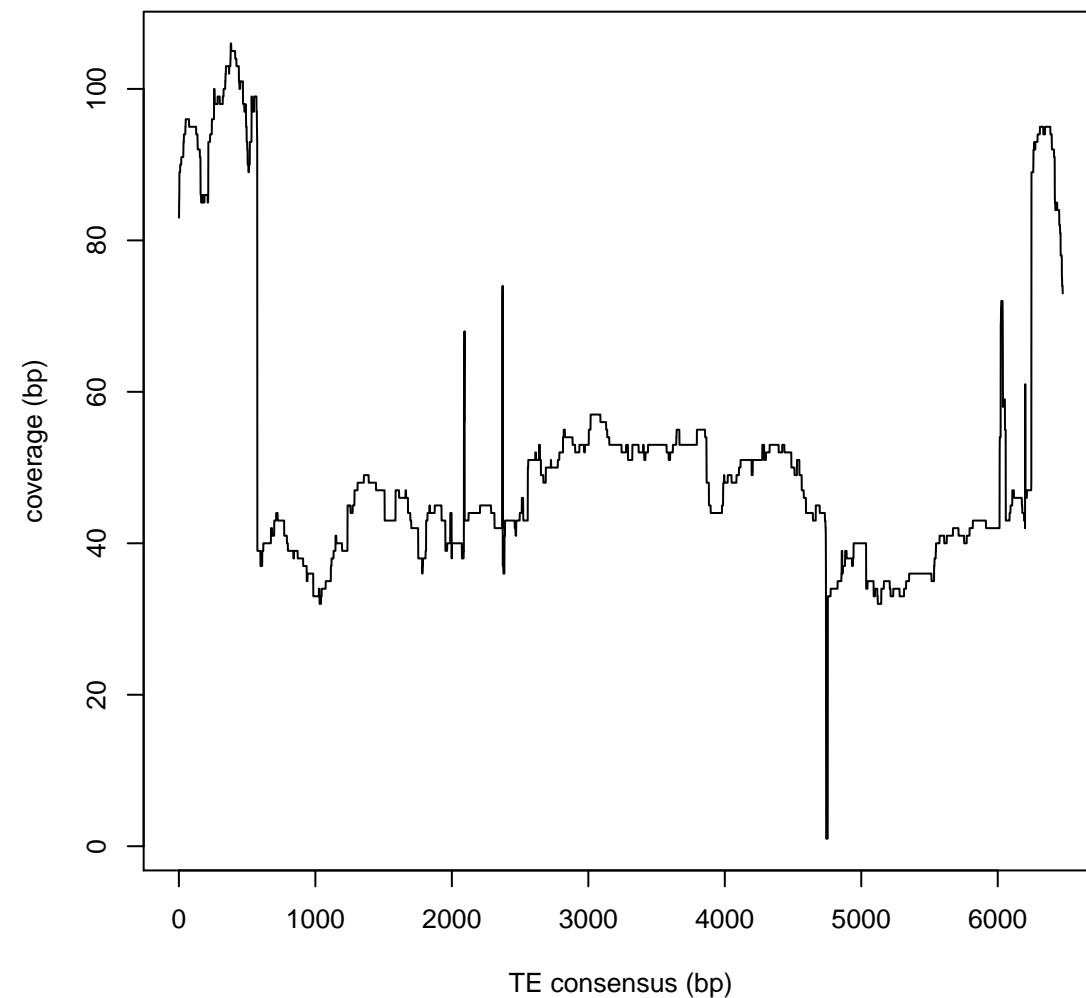


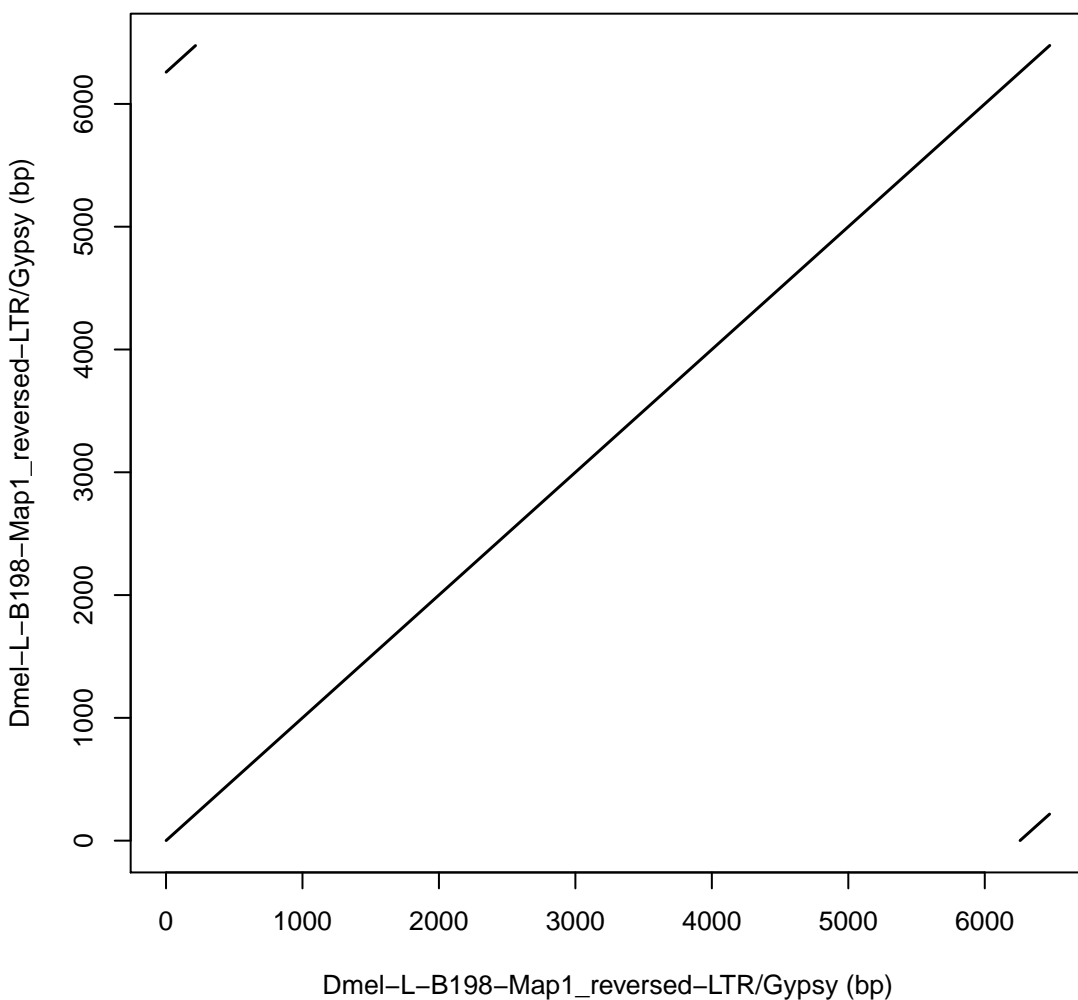
TE: Dmel-L-B198-Map1_reversed-LTR/Gypsy
consensus size: 6476bp; fragments: 515; full length: 1 (>=5828.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

