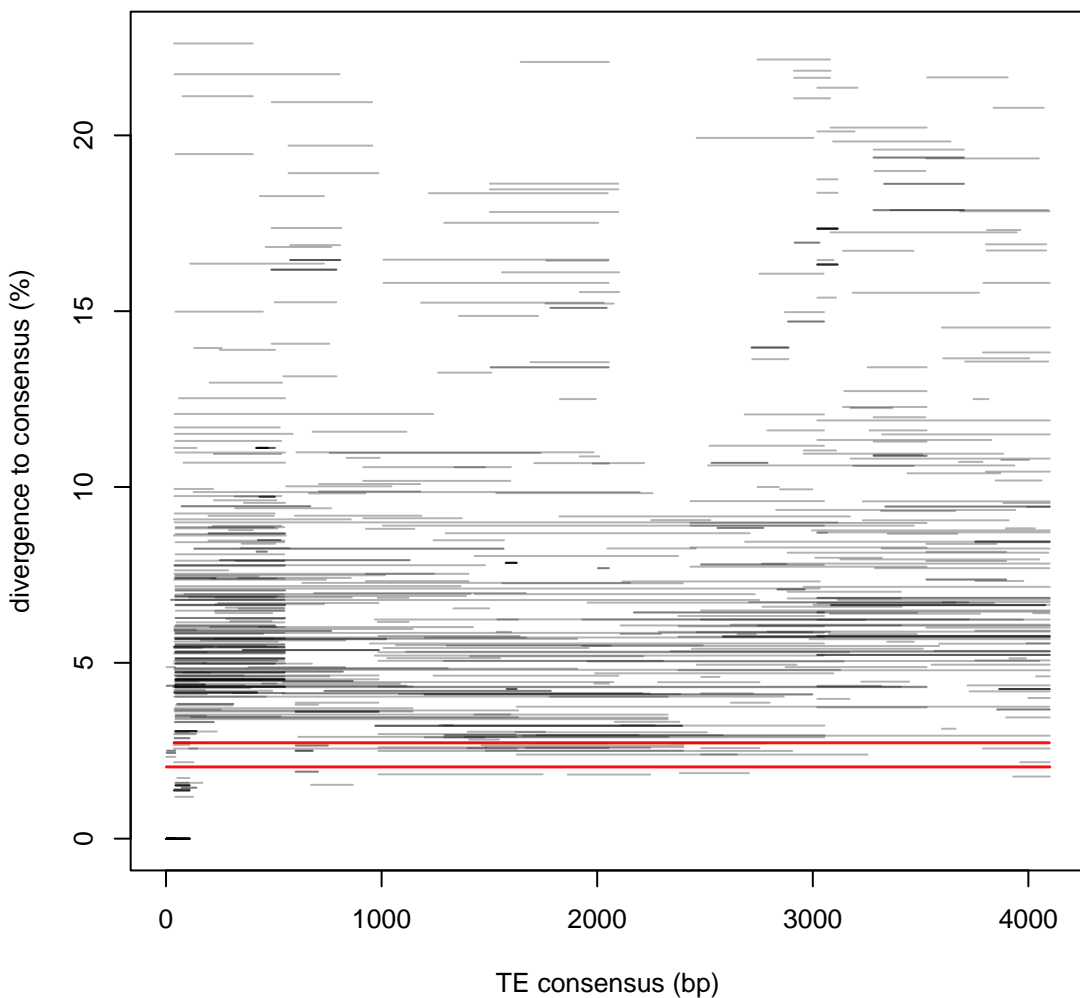
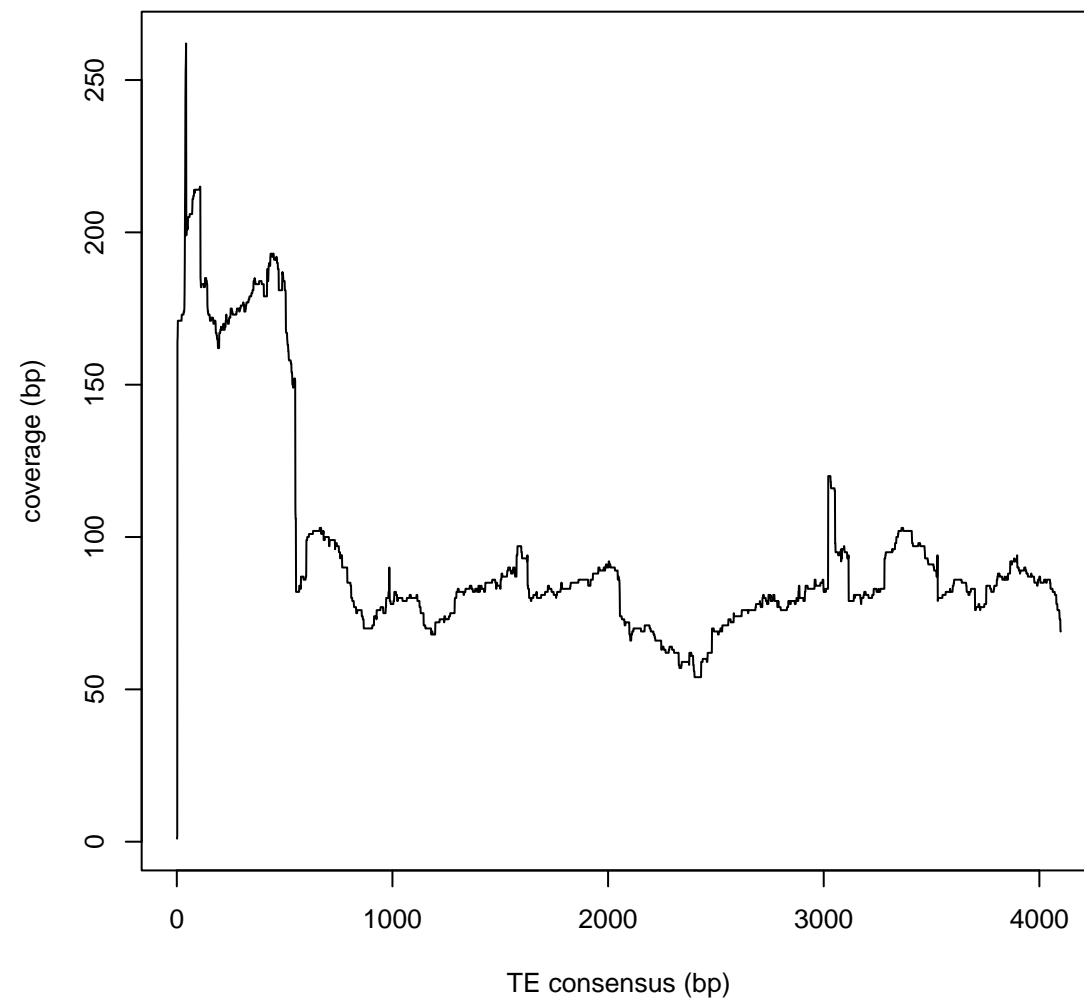


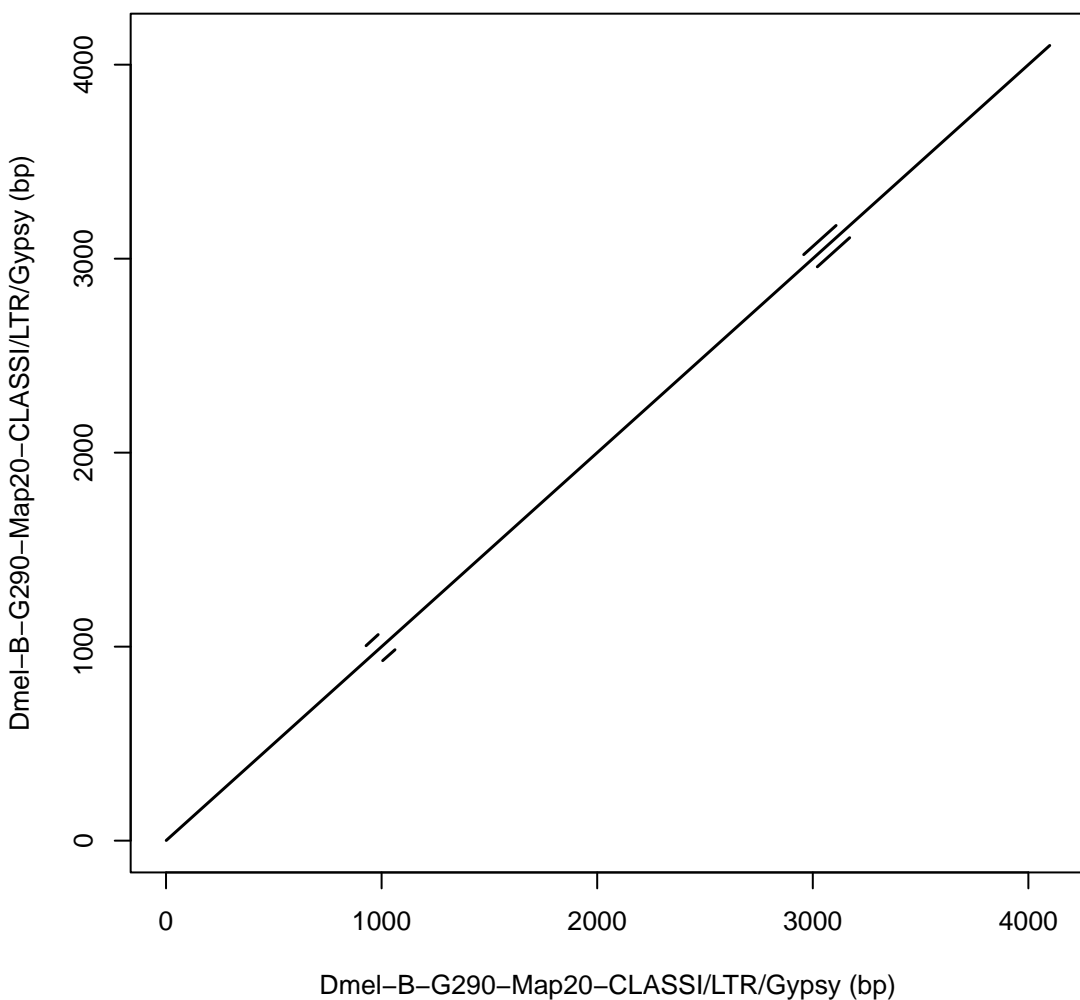
TE: Dmel-B-G290-Map20-CLASSI/LTR/Gypsy  
consensus size: 4099bp; fragments: 831; full length: 2 (>=3689.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

