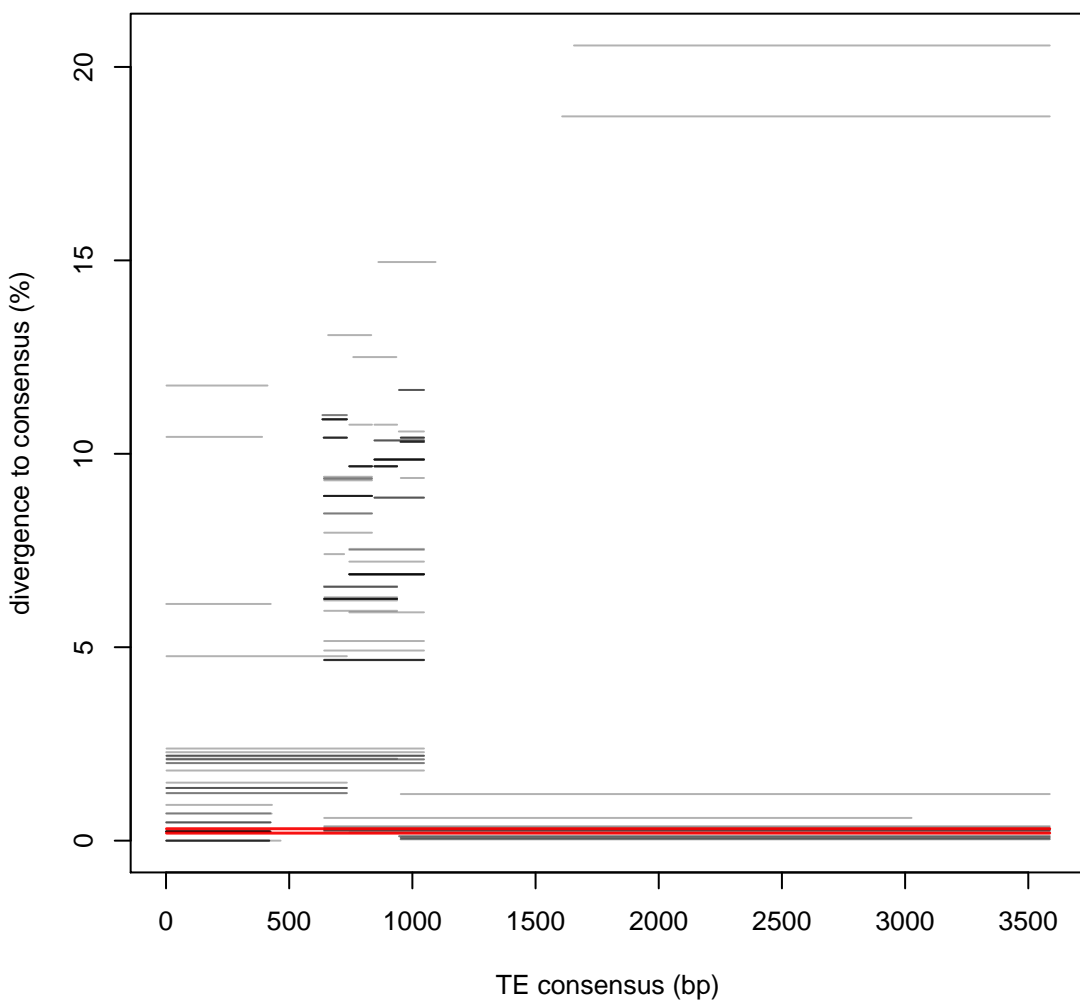
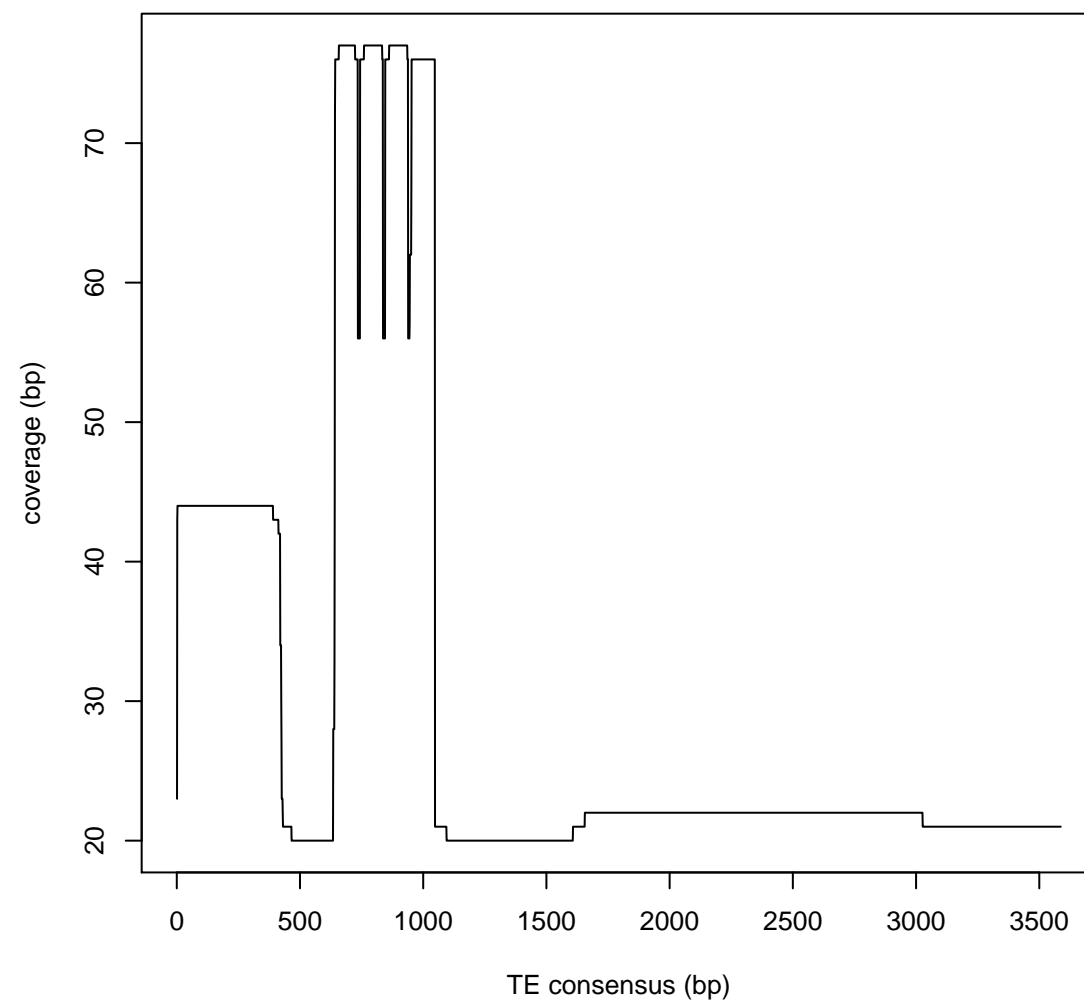


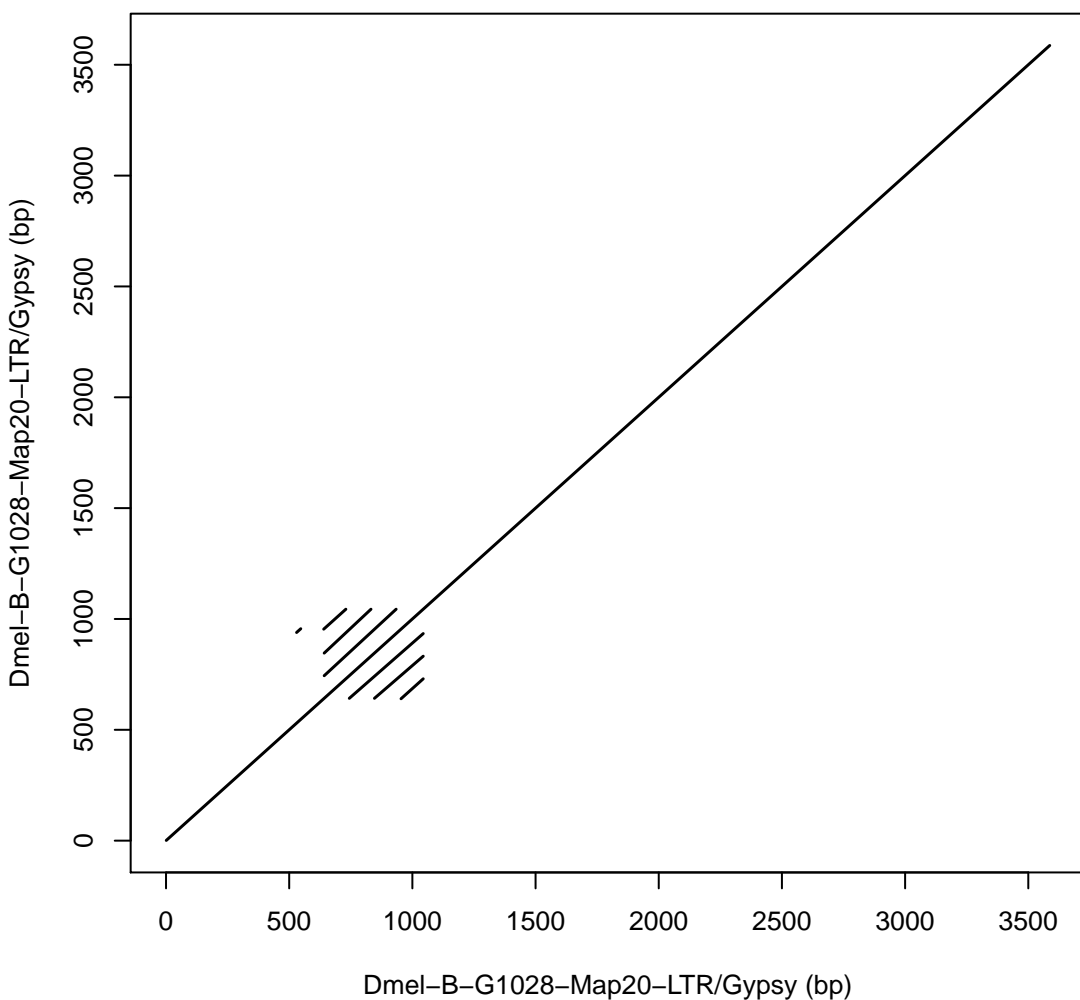
TE: Dmel-B-G1028-Map20-LTR/Gypsy
consensus size: 3587bp; fragments: 165; full length: 2 (>=3228.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

