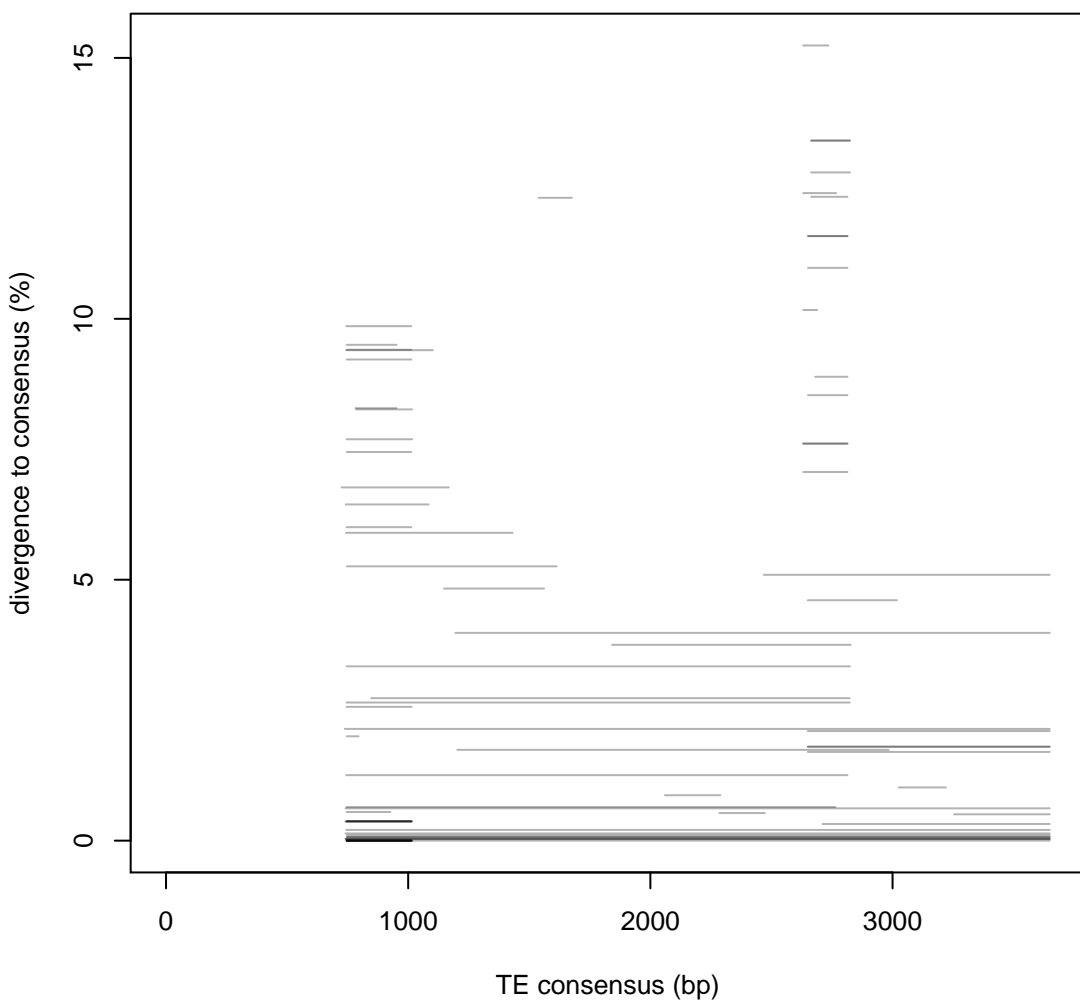
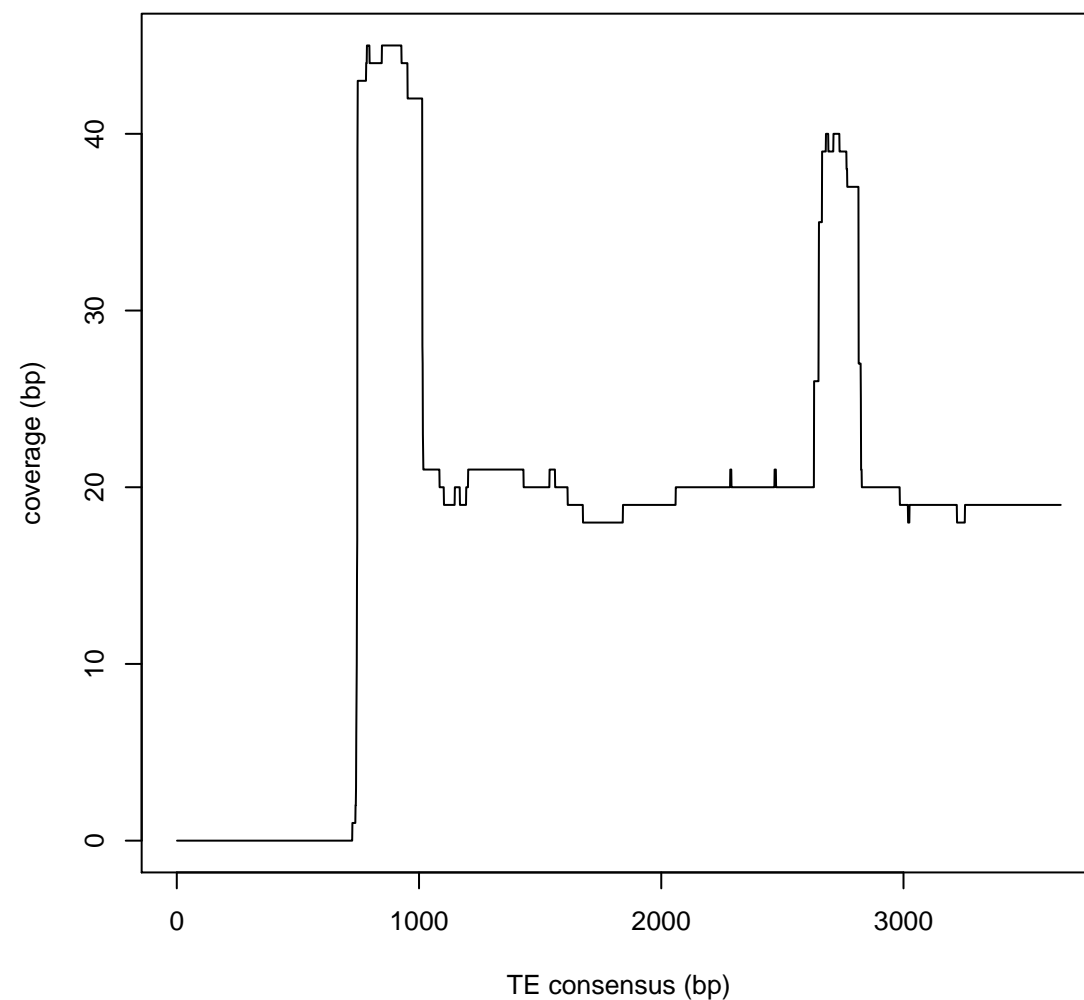


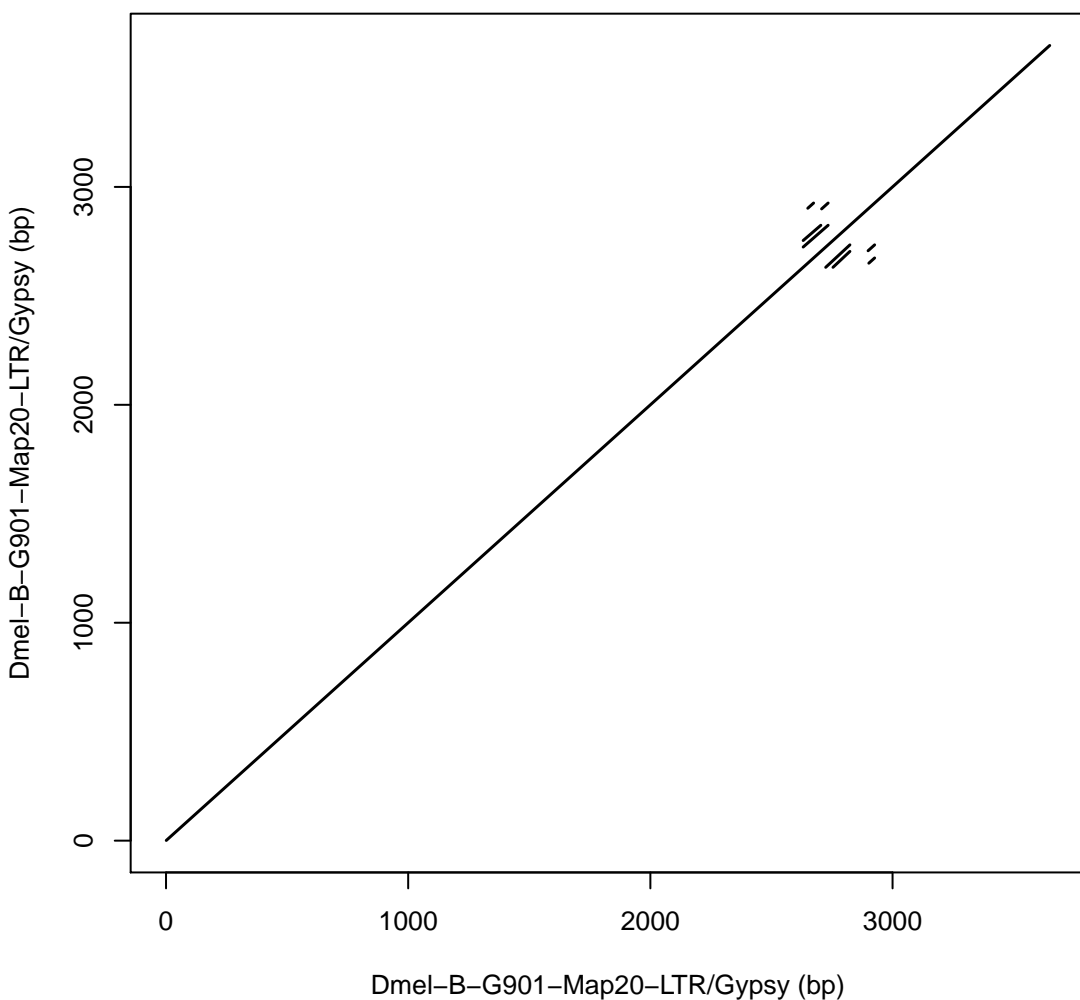
TE: Dmel-B-G901-Map20-LTR/Gypsy
consensus size: 3649bp; fragments: 77; full length: 0 (>=3284.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

