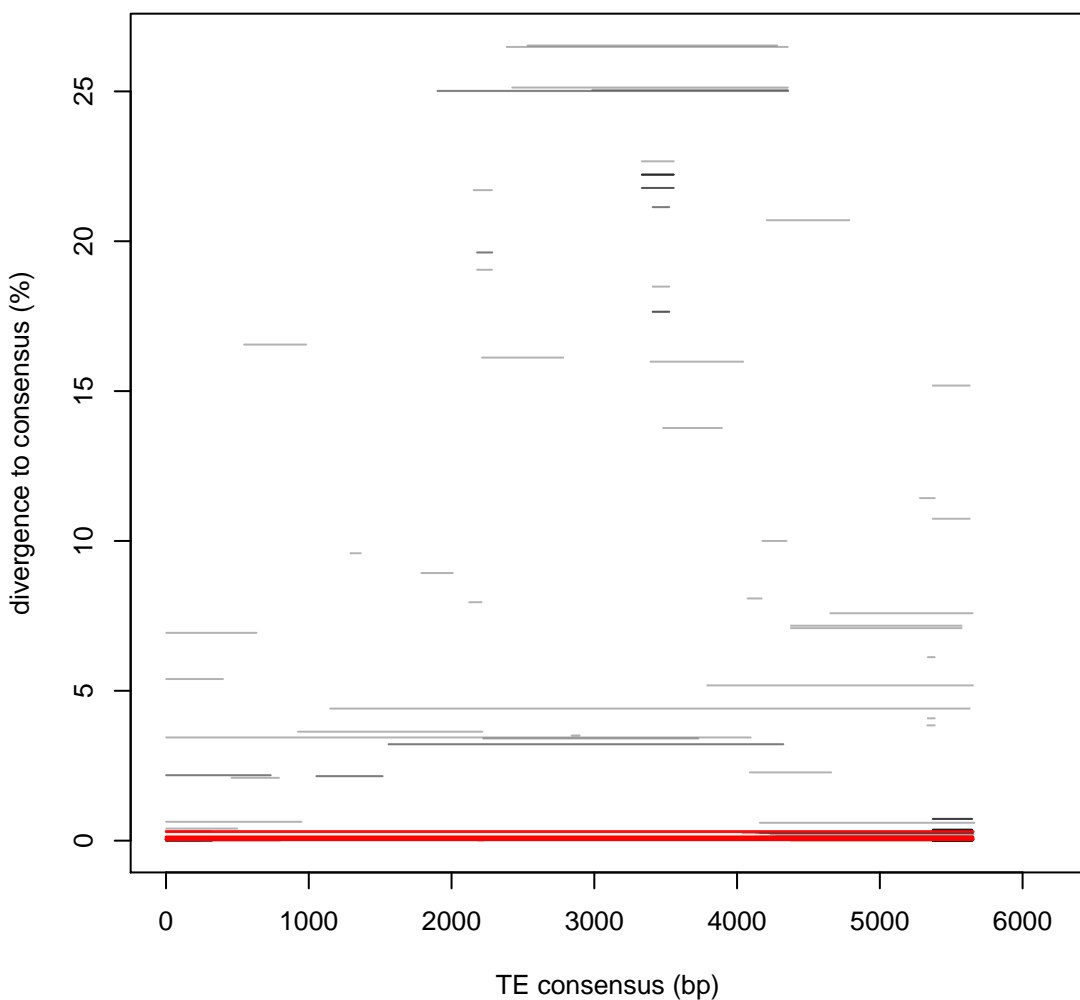
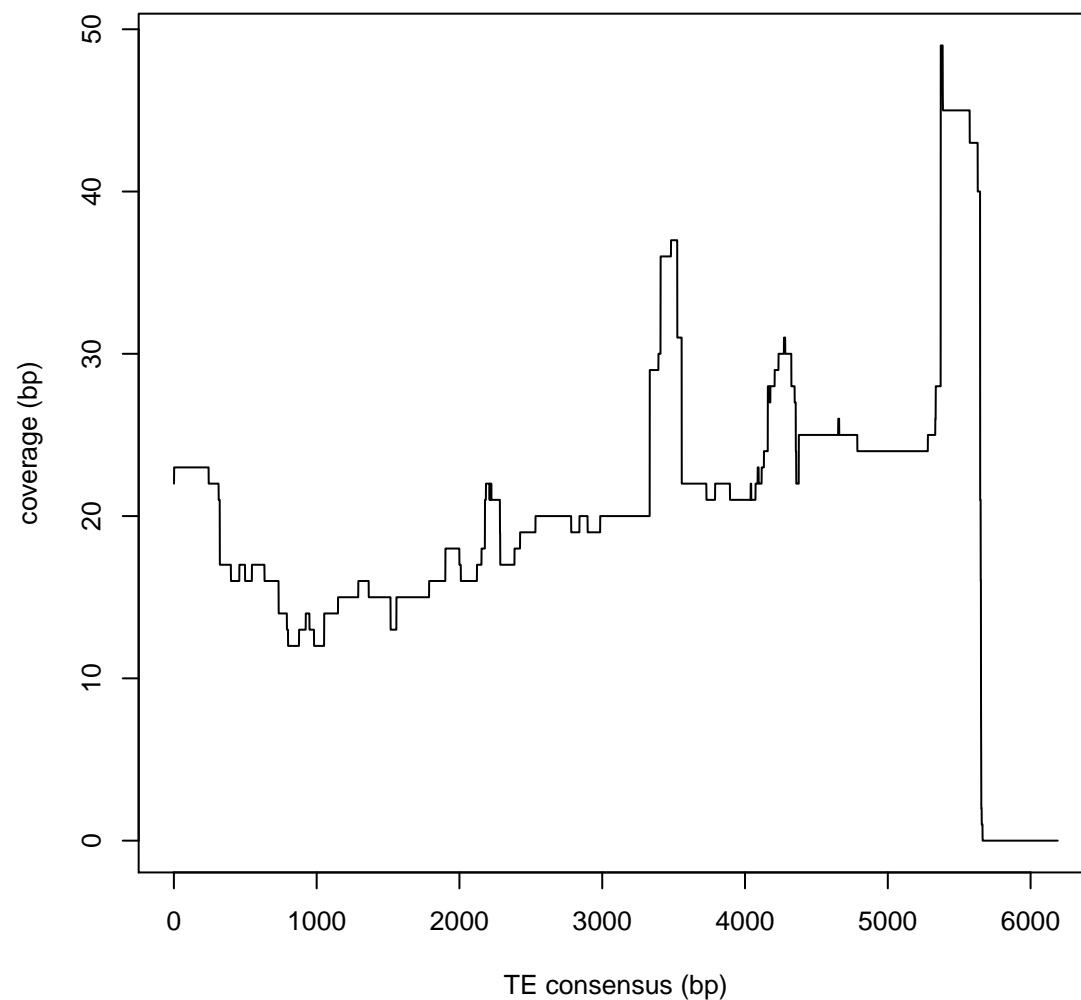


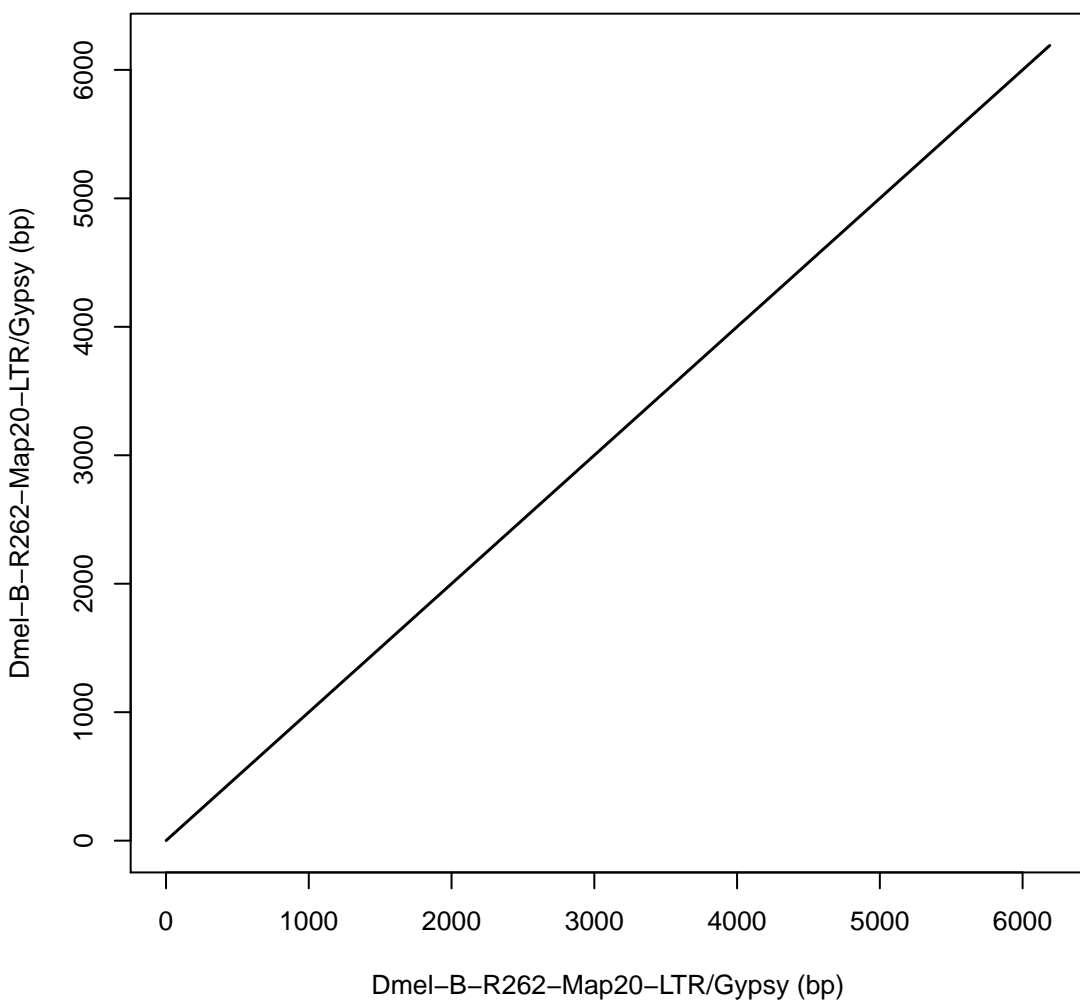
TE: Dmel-B-R262-Map20-LTR/Gypsy
consensus size: 6190bp; fragments: 109; full length: 8 (>=5571bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

