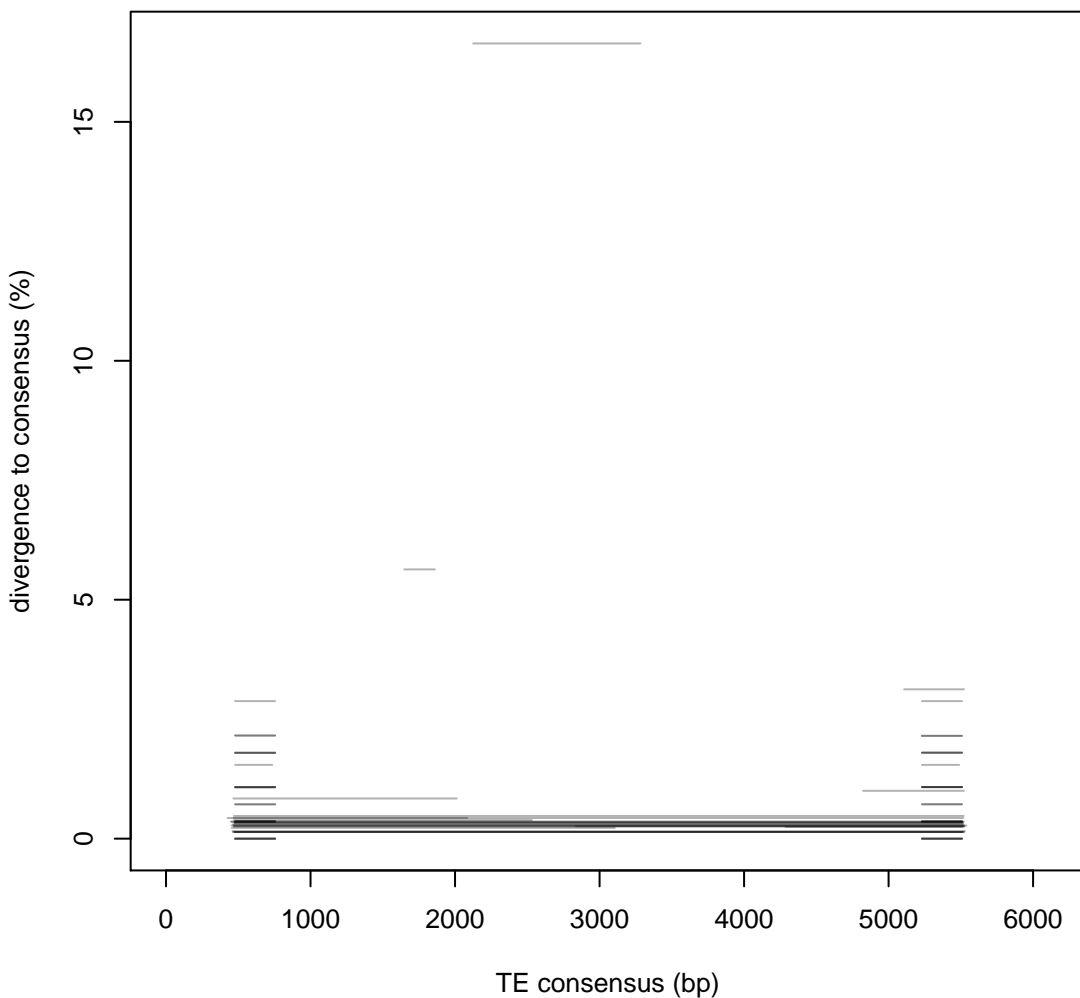
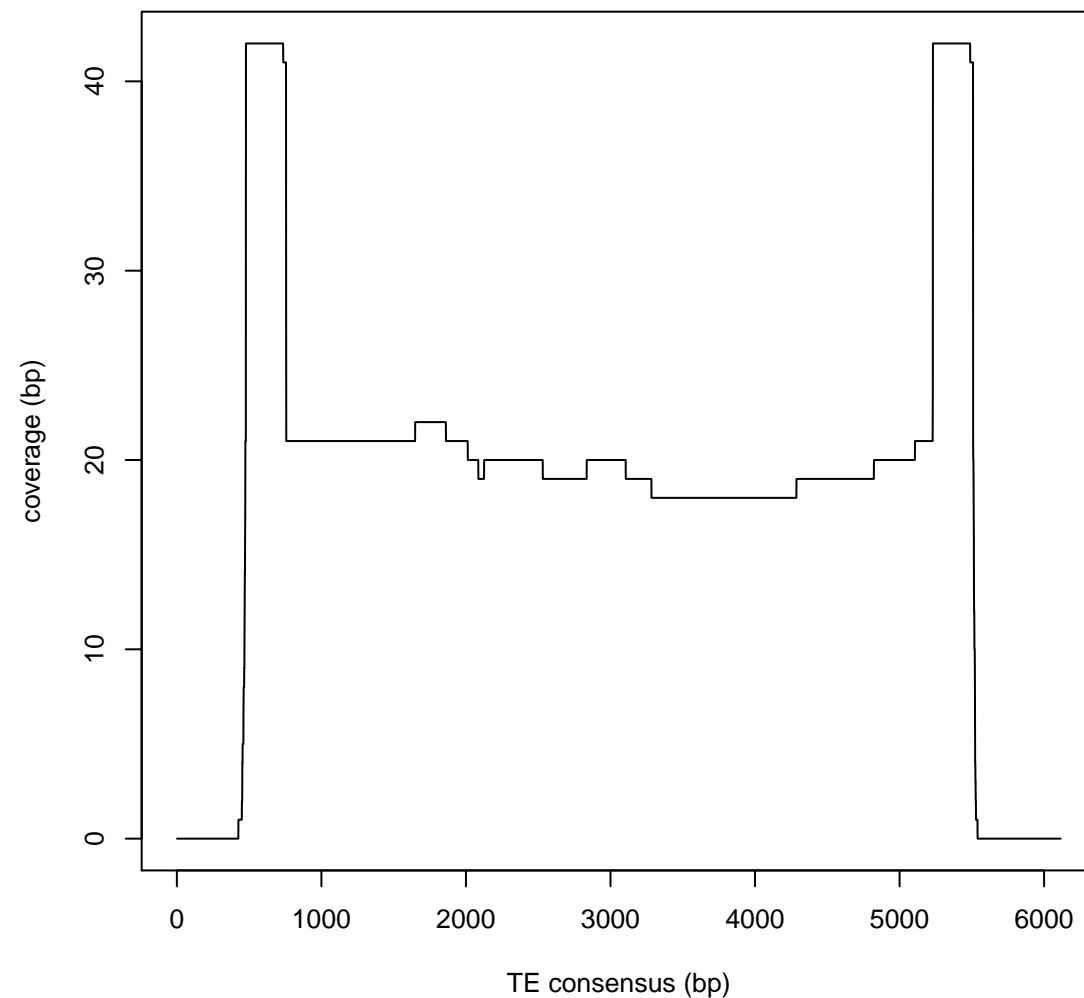


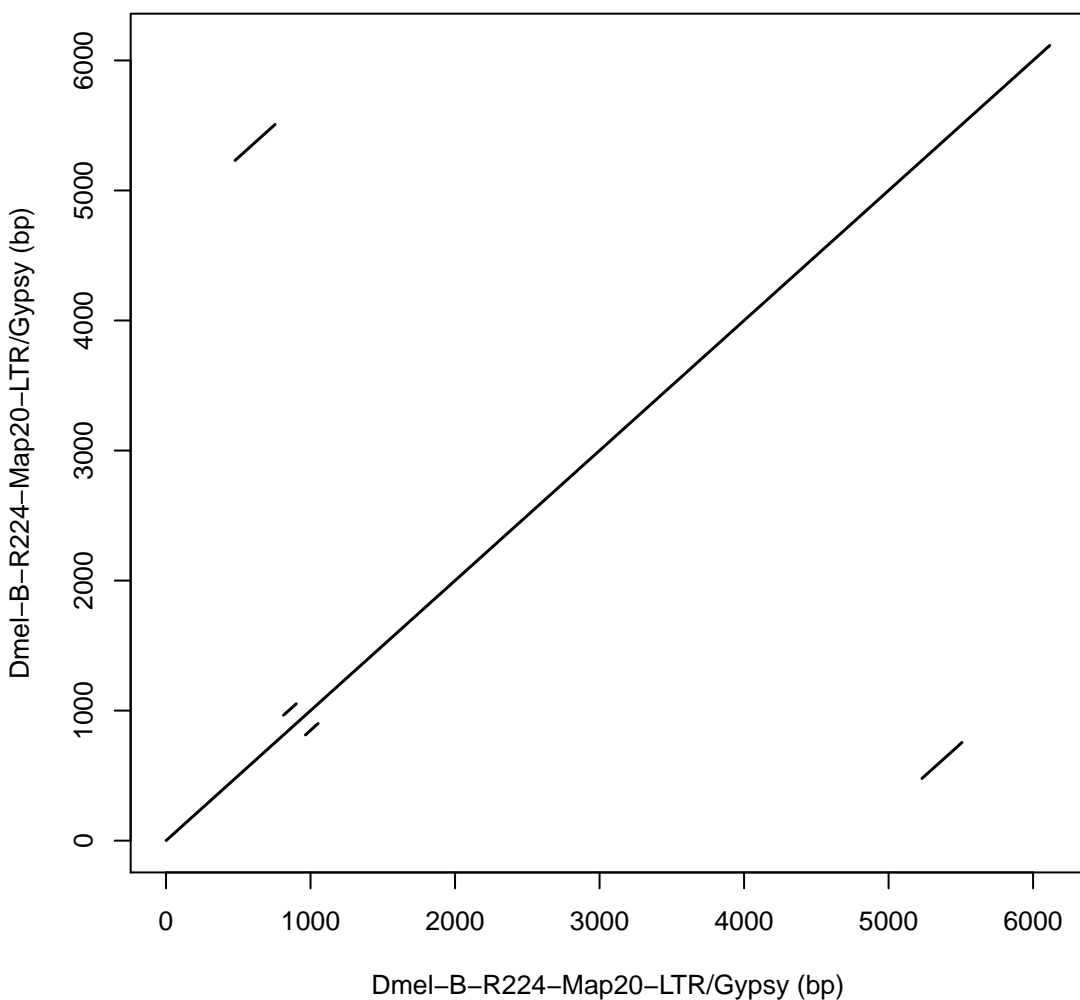
TE: Dmel-B-R224-Map20-LTR/Gypsy
consensus size: 6115bp; fragments: 69; full length: 0 (>=5503.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

