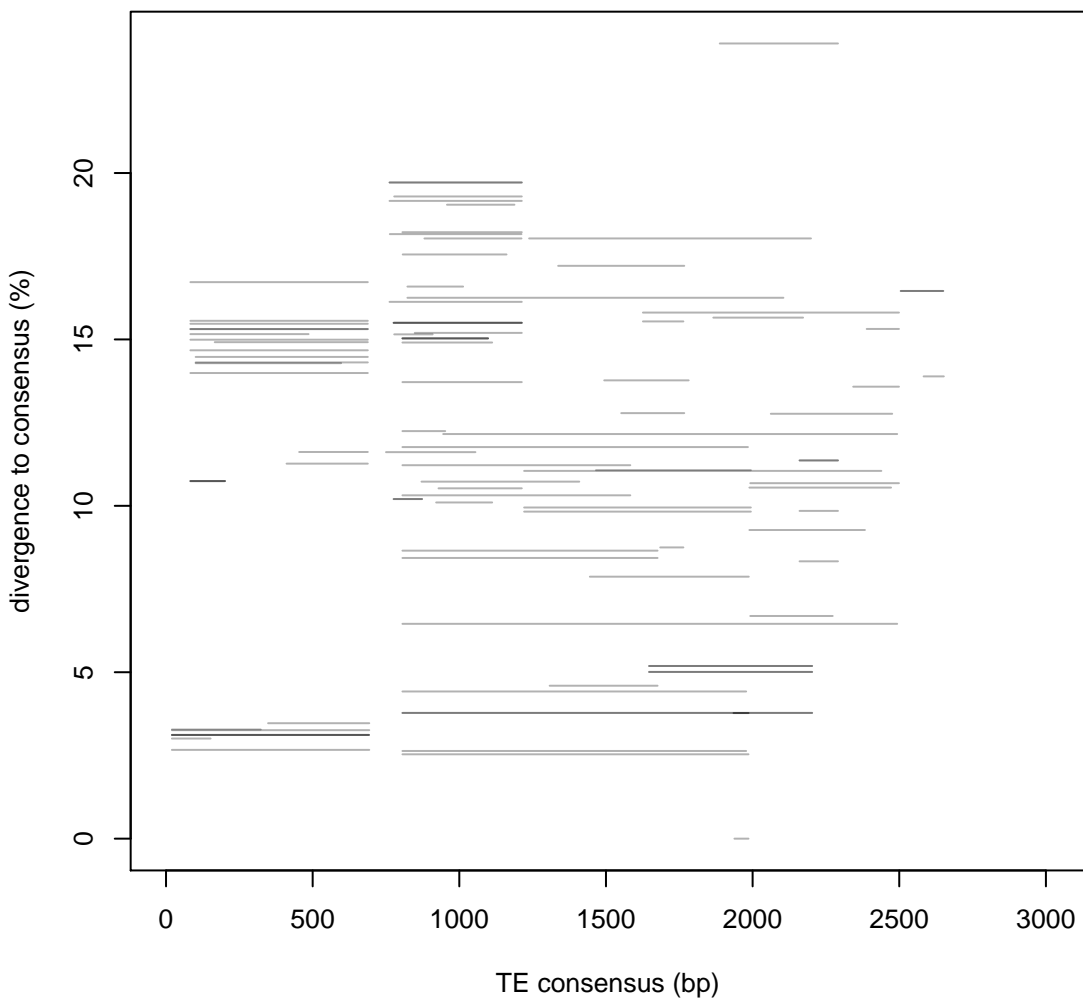
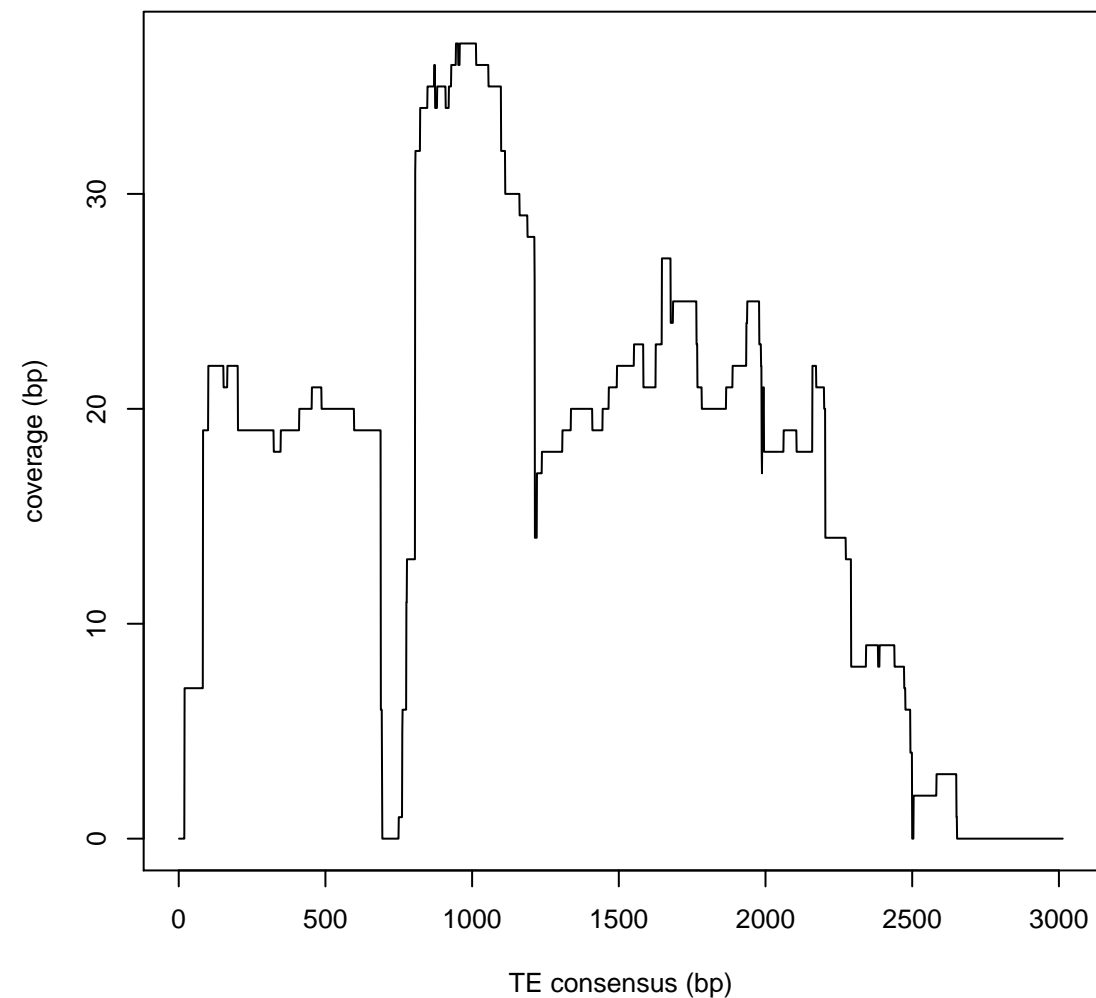


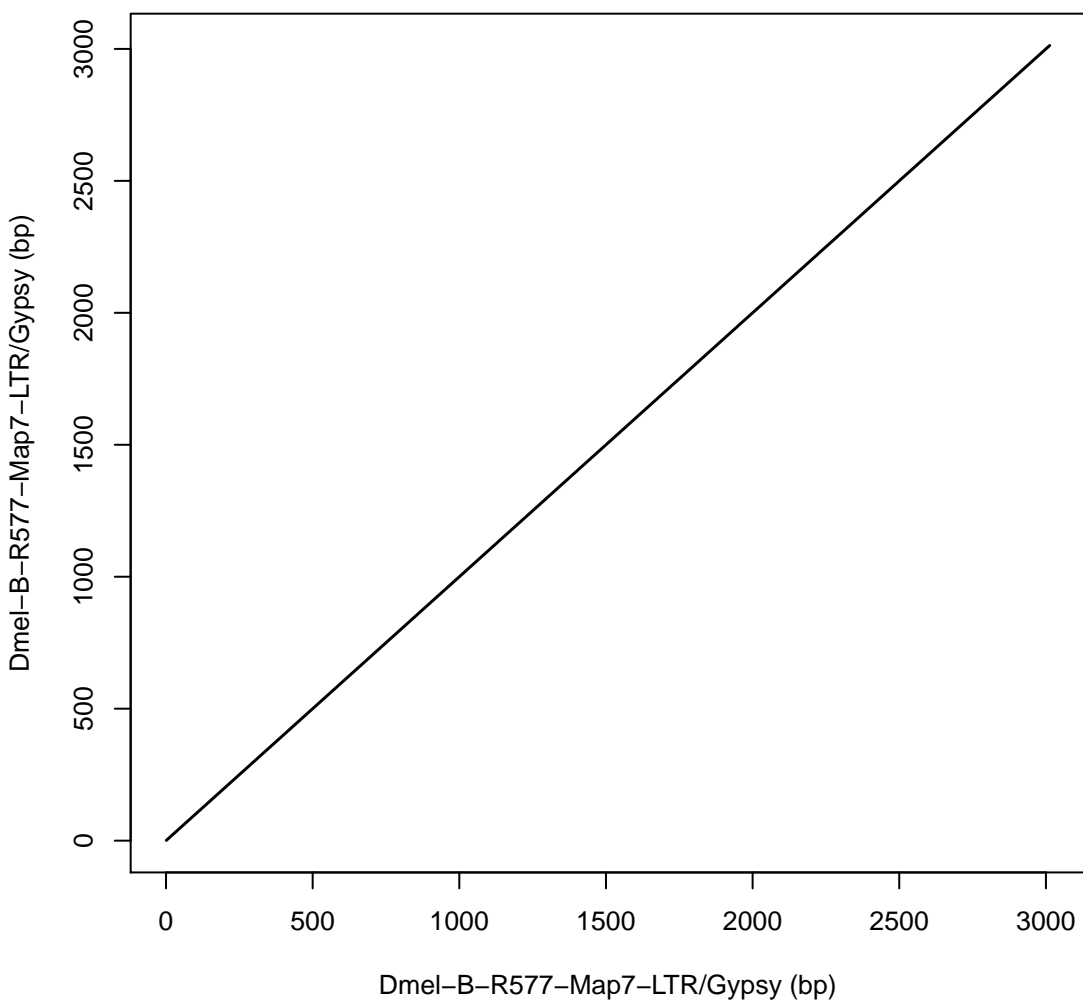
TE: Dmel-B-R577-Map7-LTR/Gypsy
consensus size: 3013bp; fragments: 103; full length: 0 (>=2711.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

