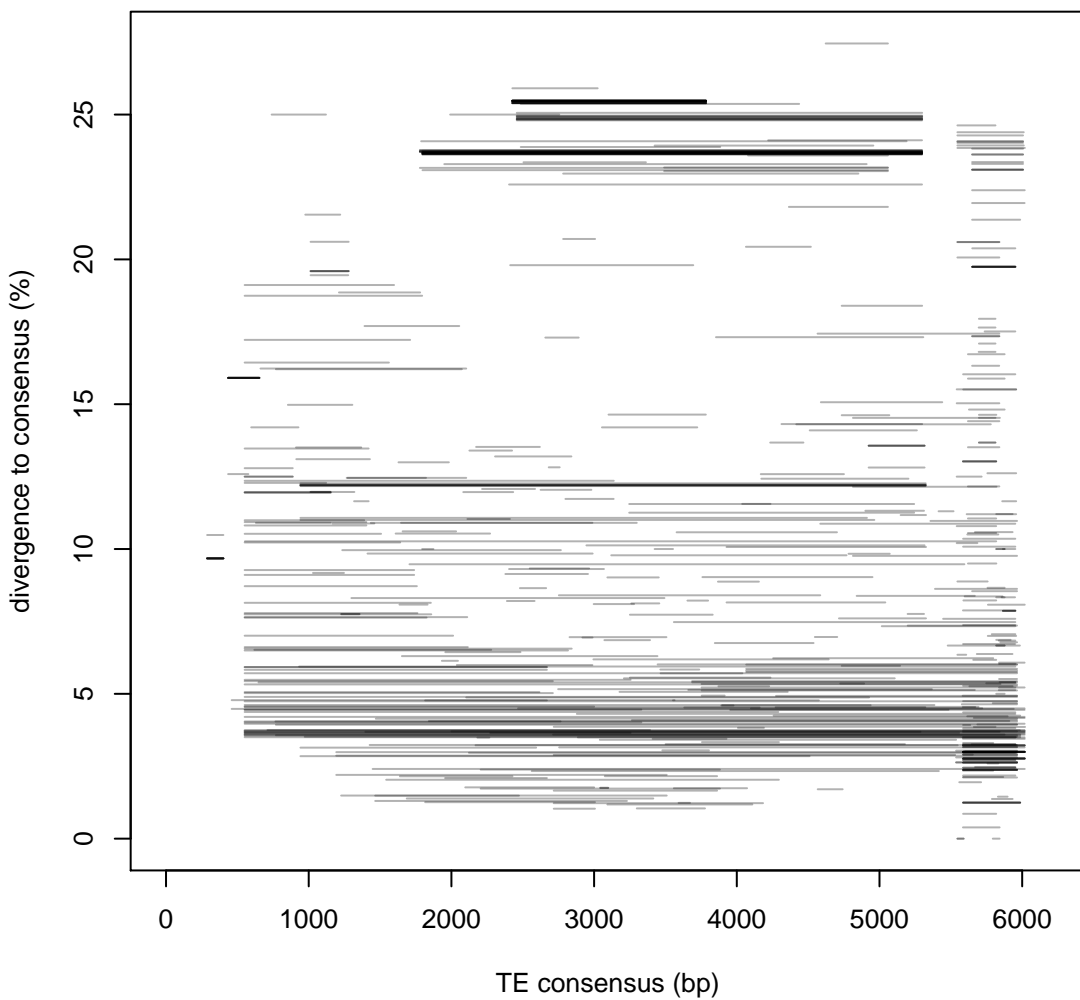
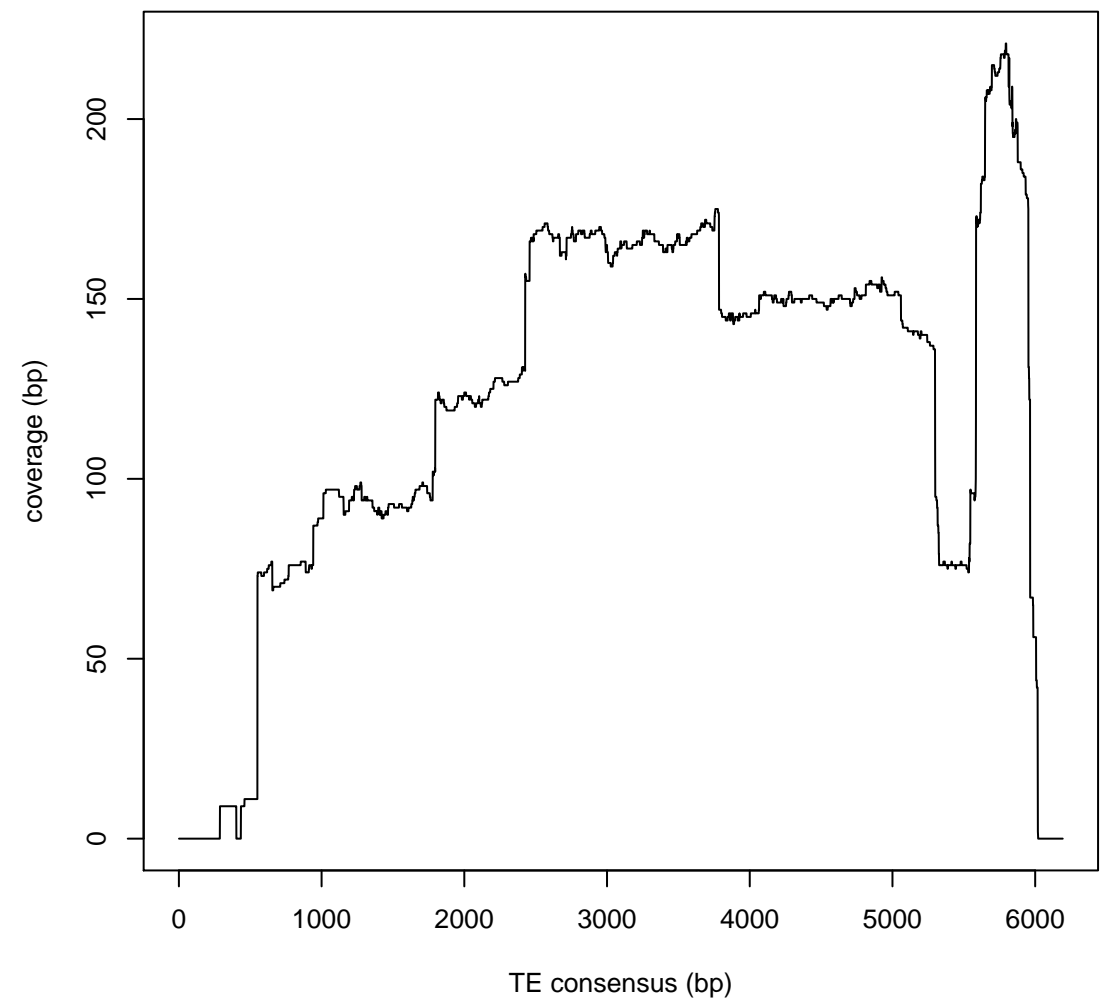


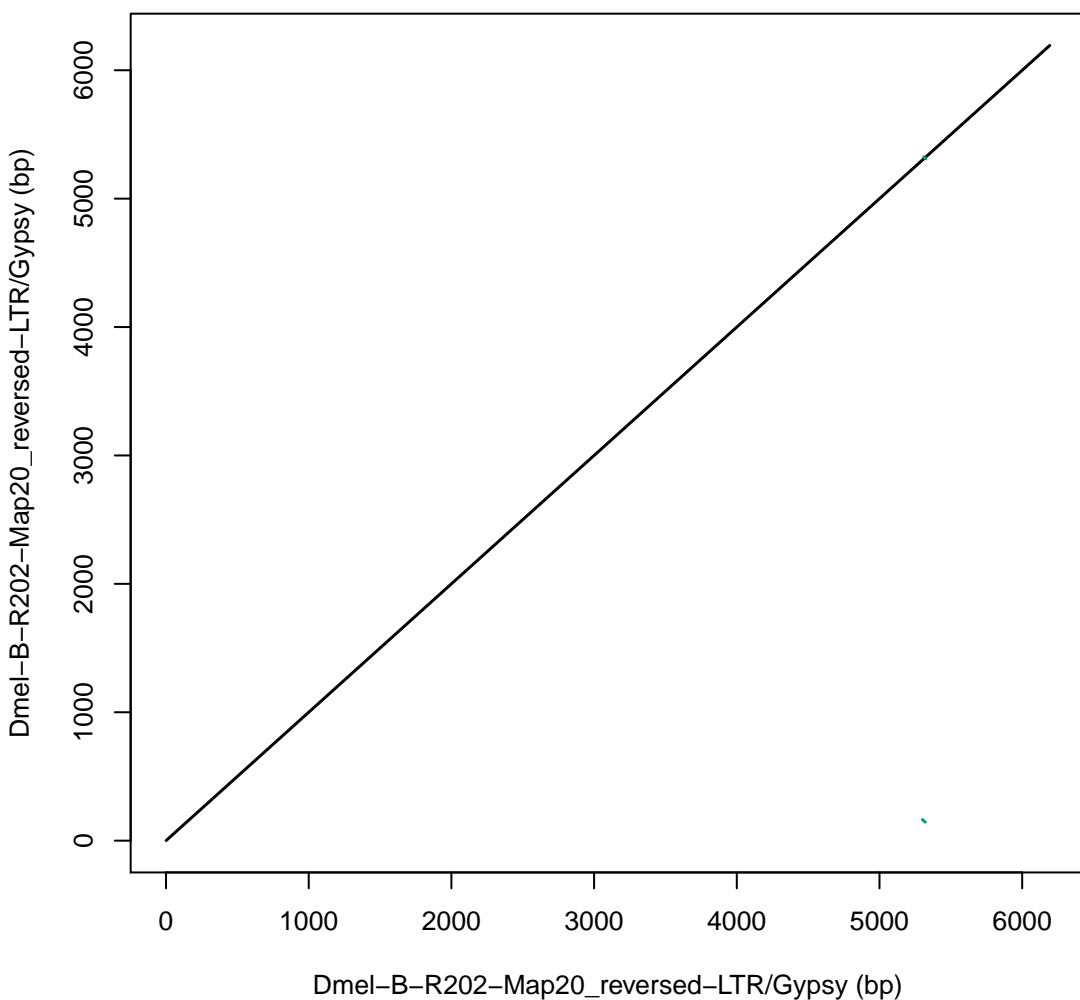
TE: Dmel-B-R202-Map20_reversed-LTR/Gypsy
consensus size: 6193bp; fragments: 573; full length: 0 (>=5573.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

