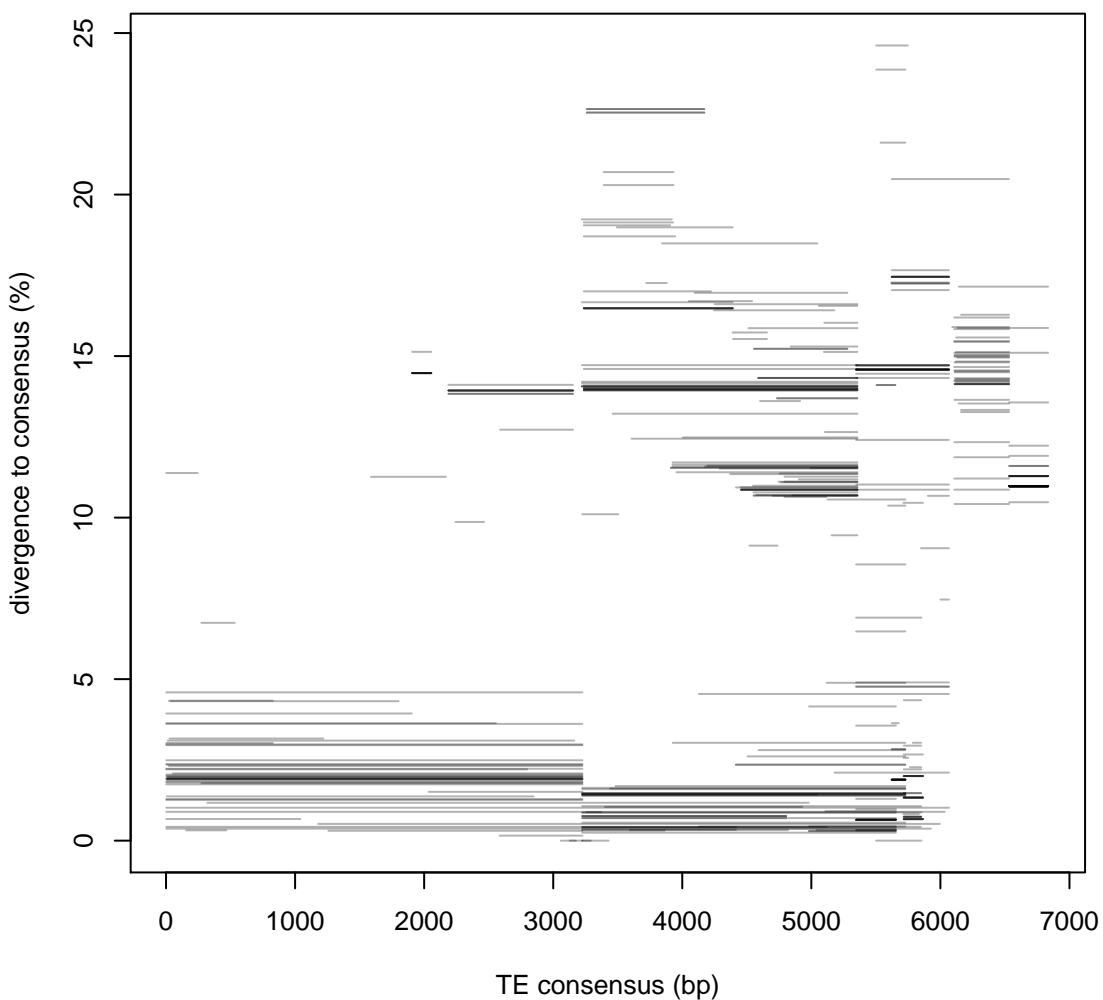
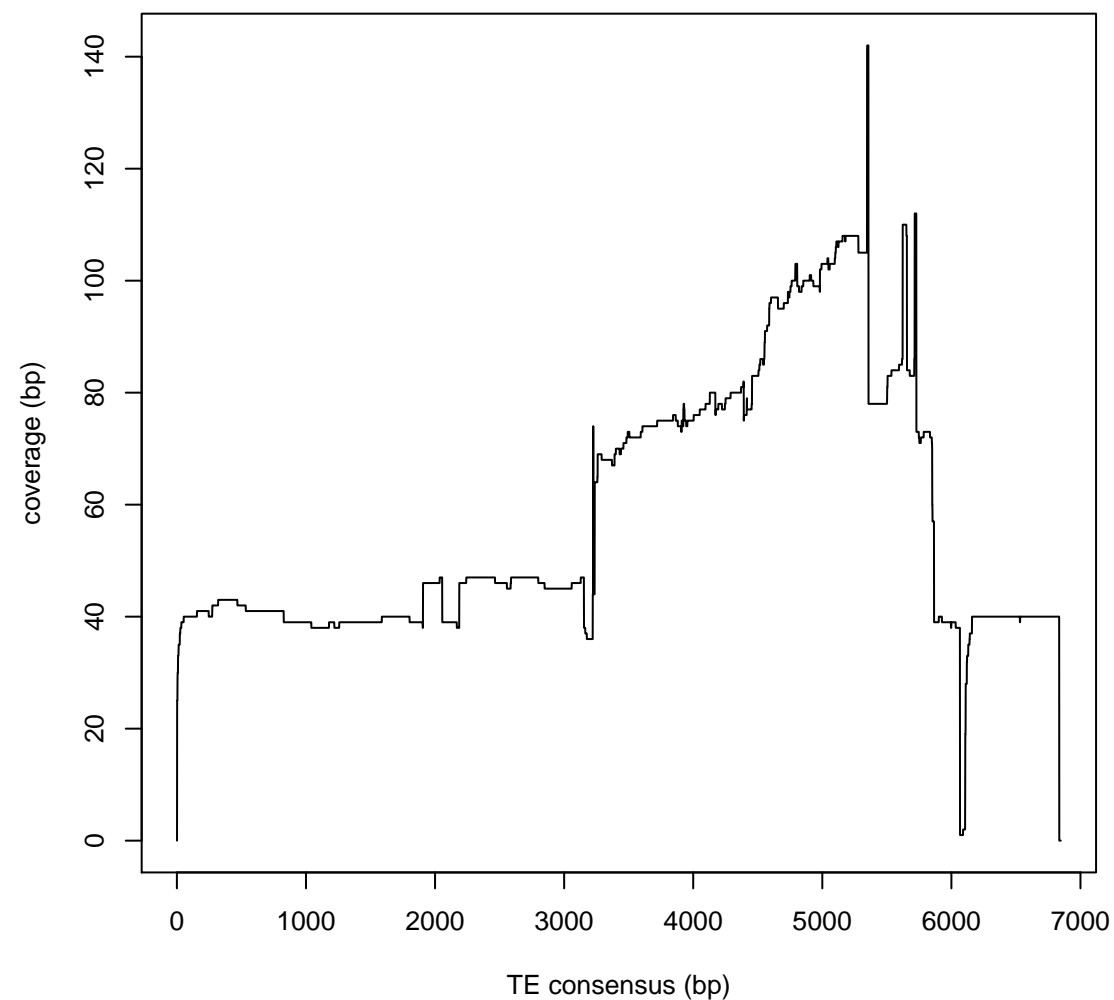


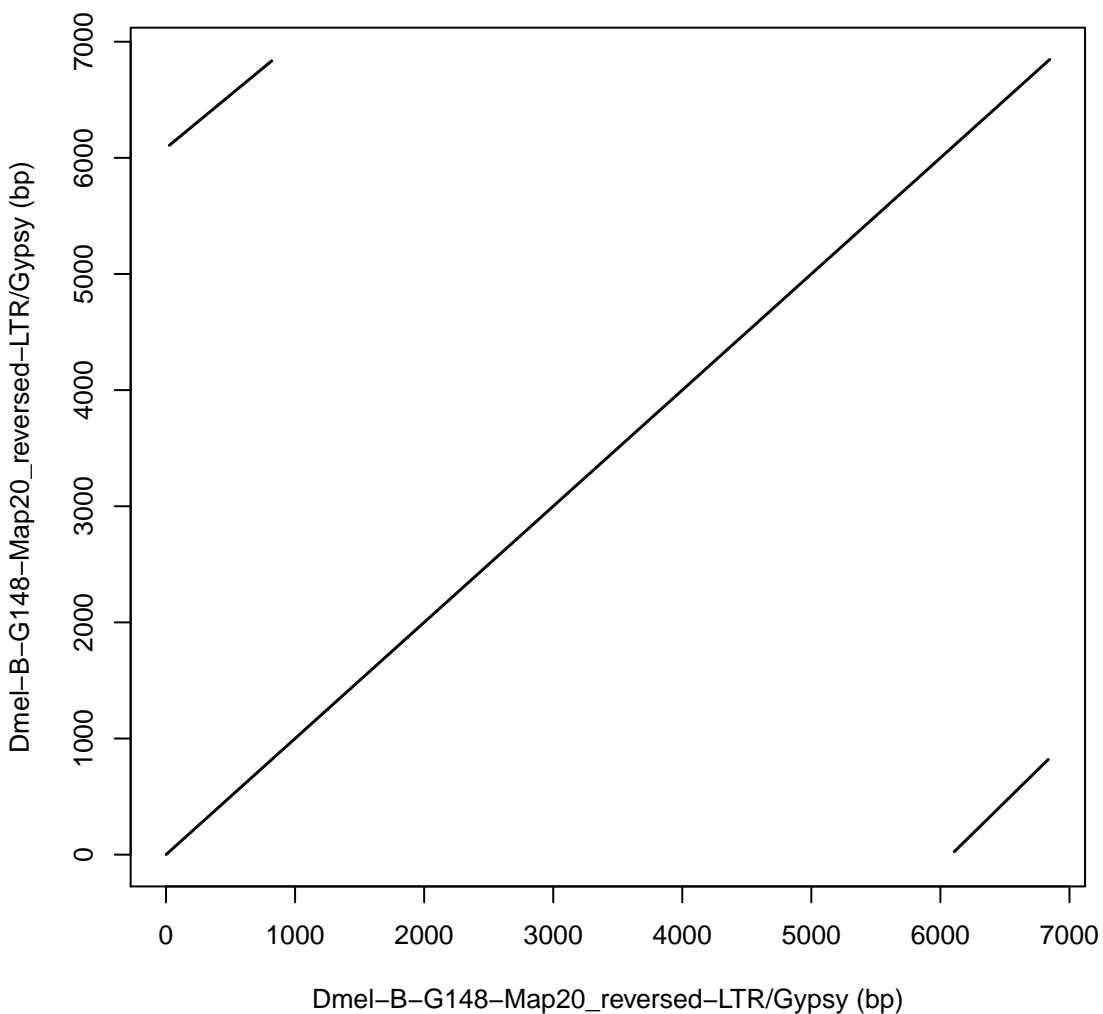
TE: Dmel-B-G148-Map20_reversed-LTR/Gypsy
consensus size: 6847bp; fragments: 393; full length: 0 (>=6162.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

