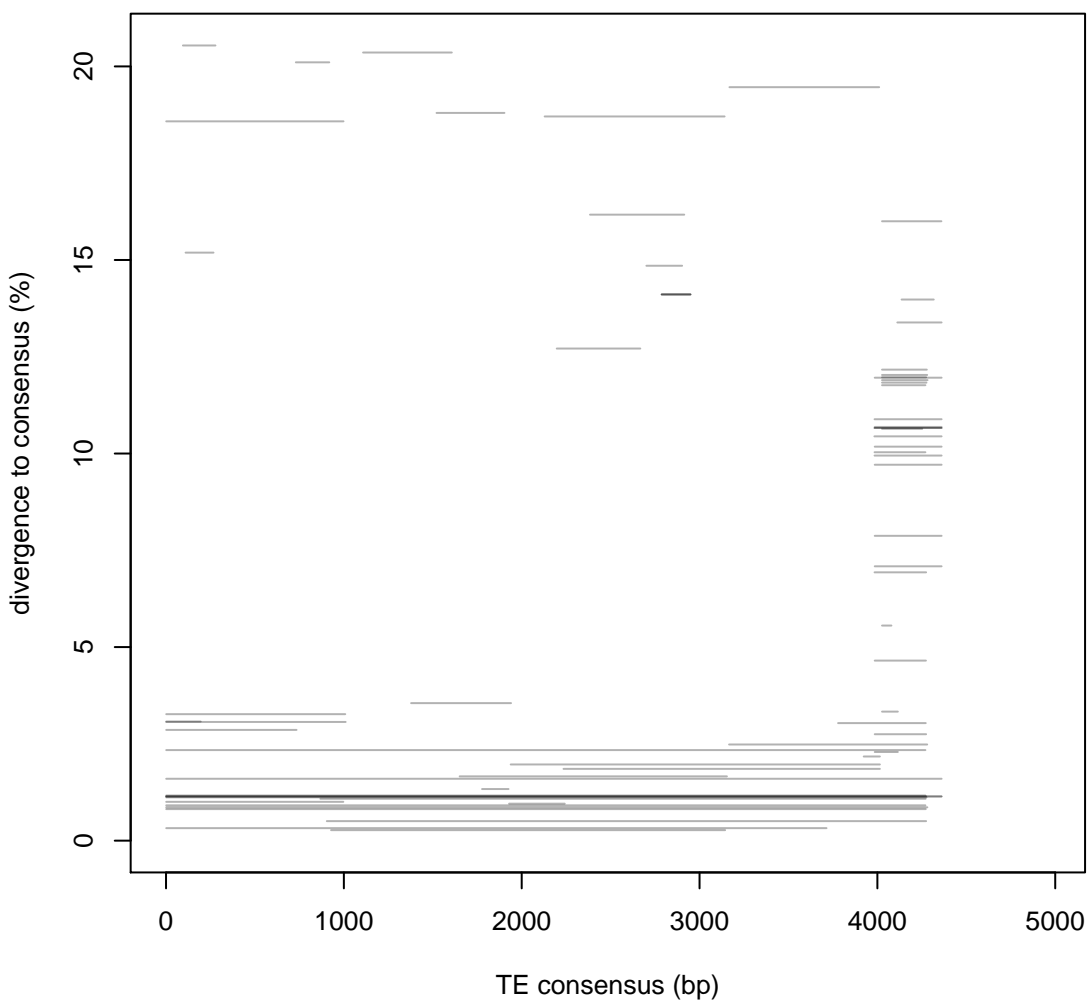
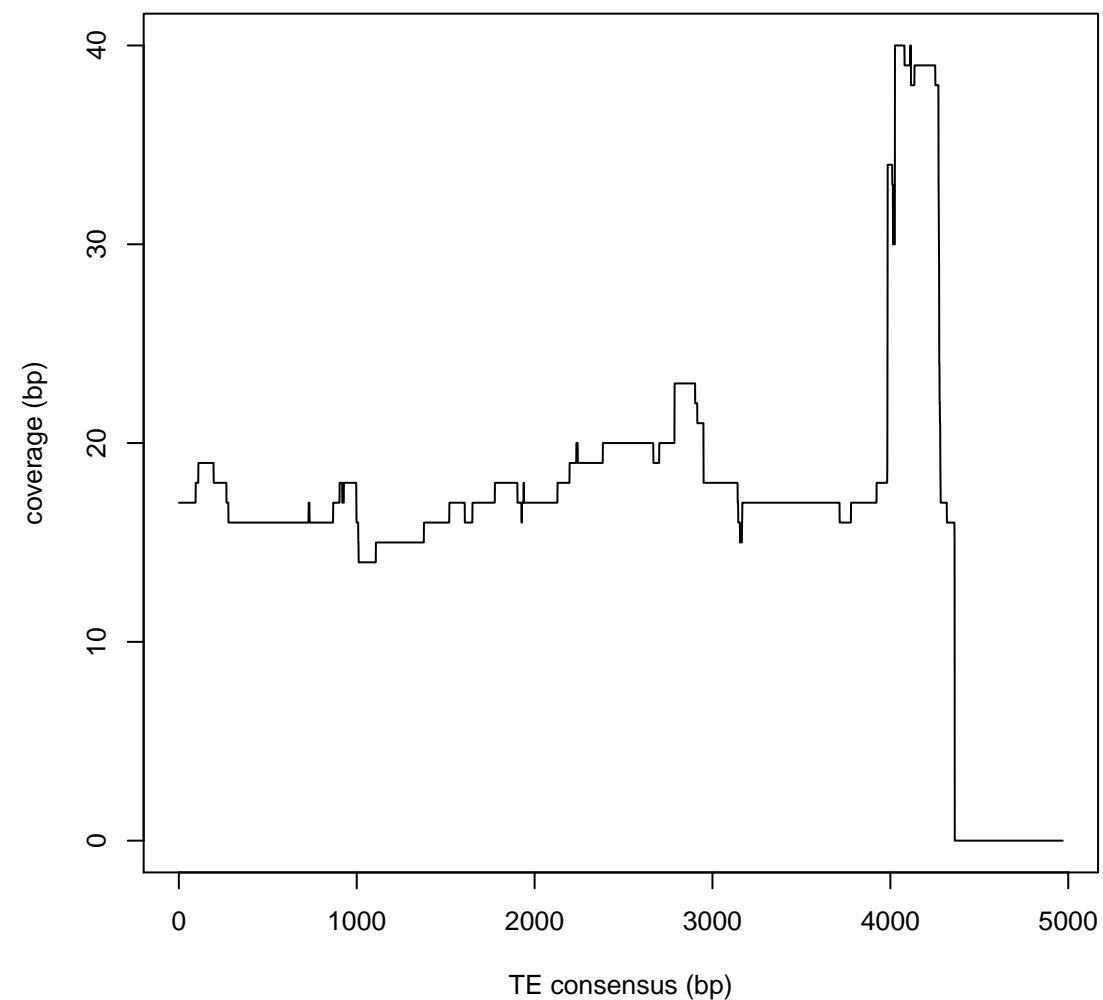


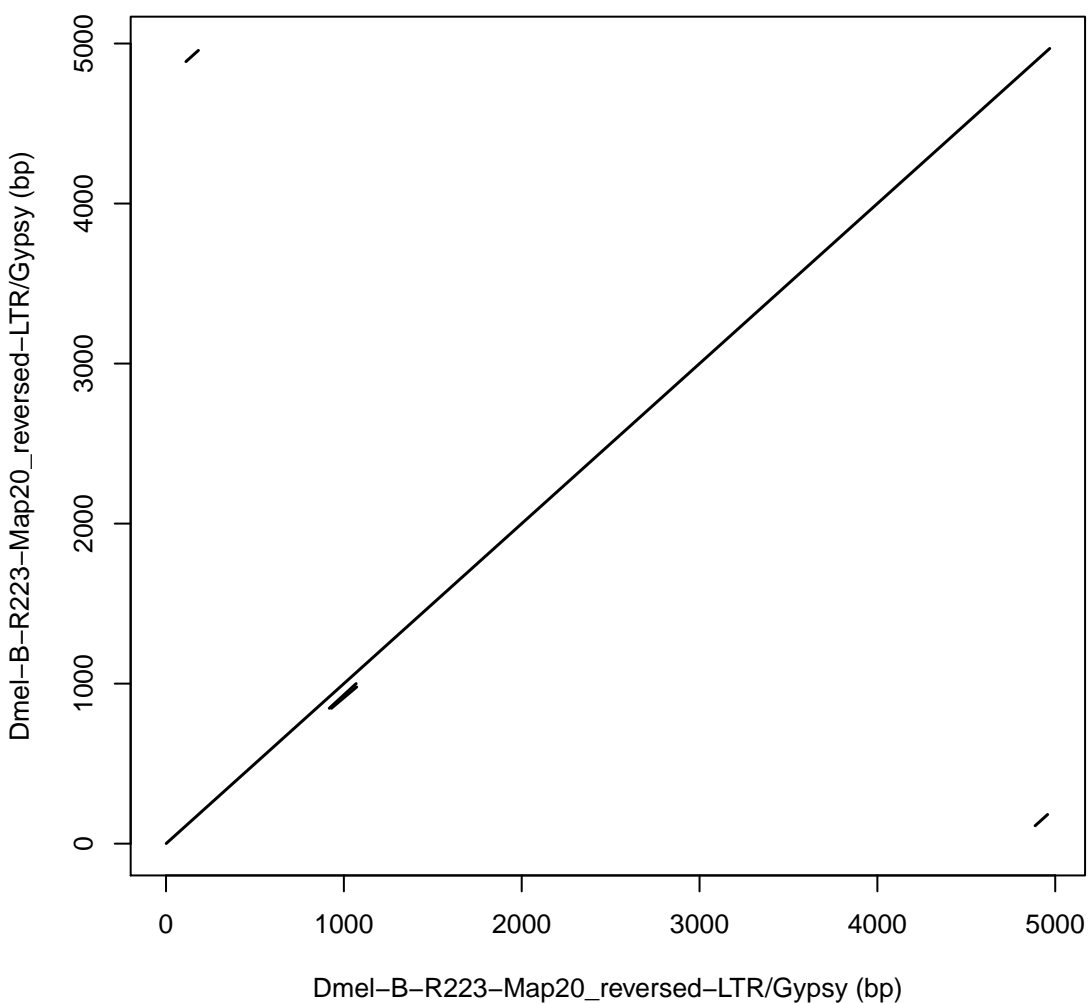
TE: Dmel-B-R223-Map20_reversed-LTR/Gypsy
consensus size: 4969bp; fragments: 70; full length: 0 (>=4472.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

