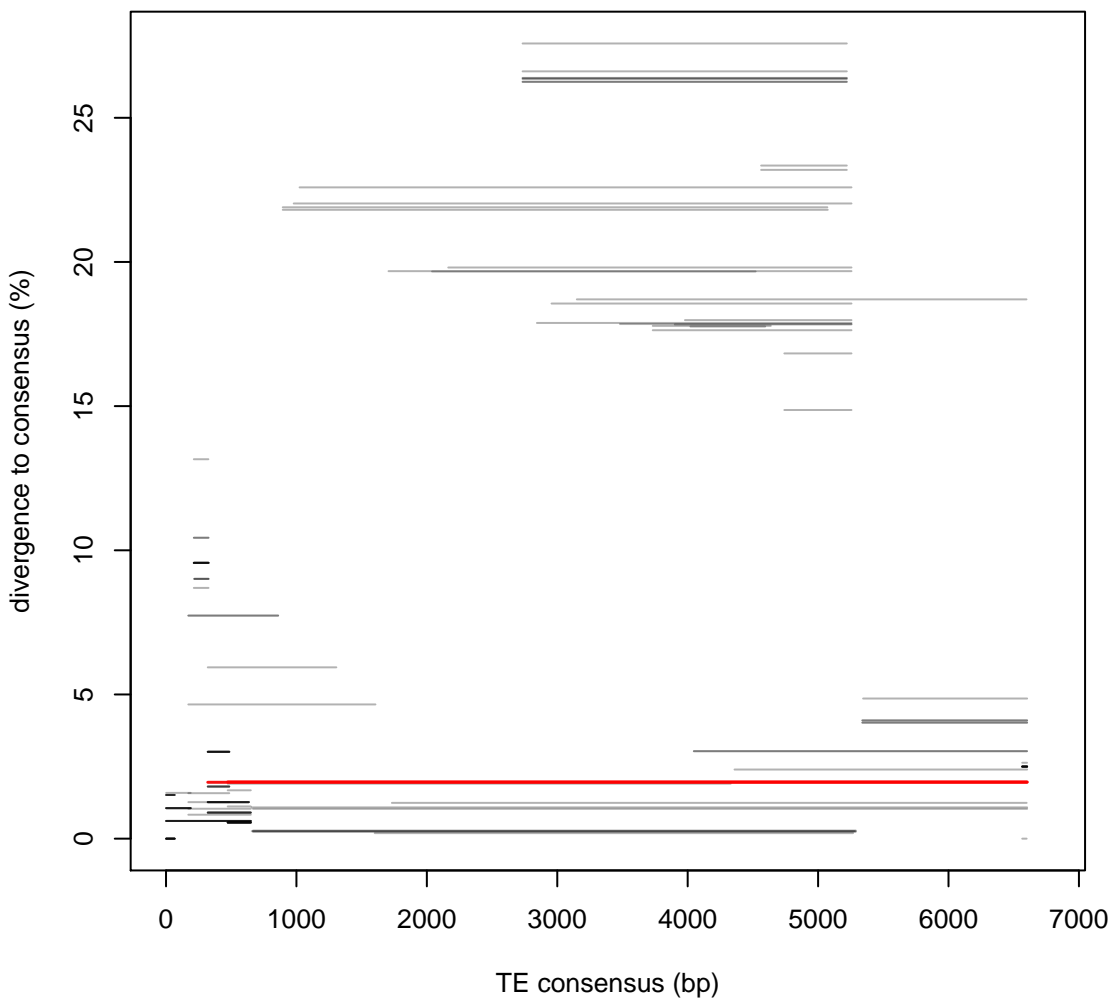
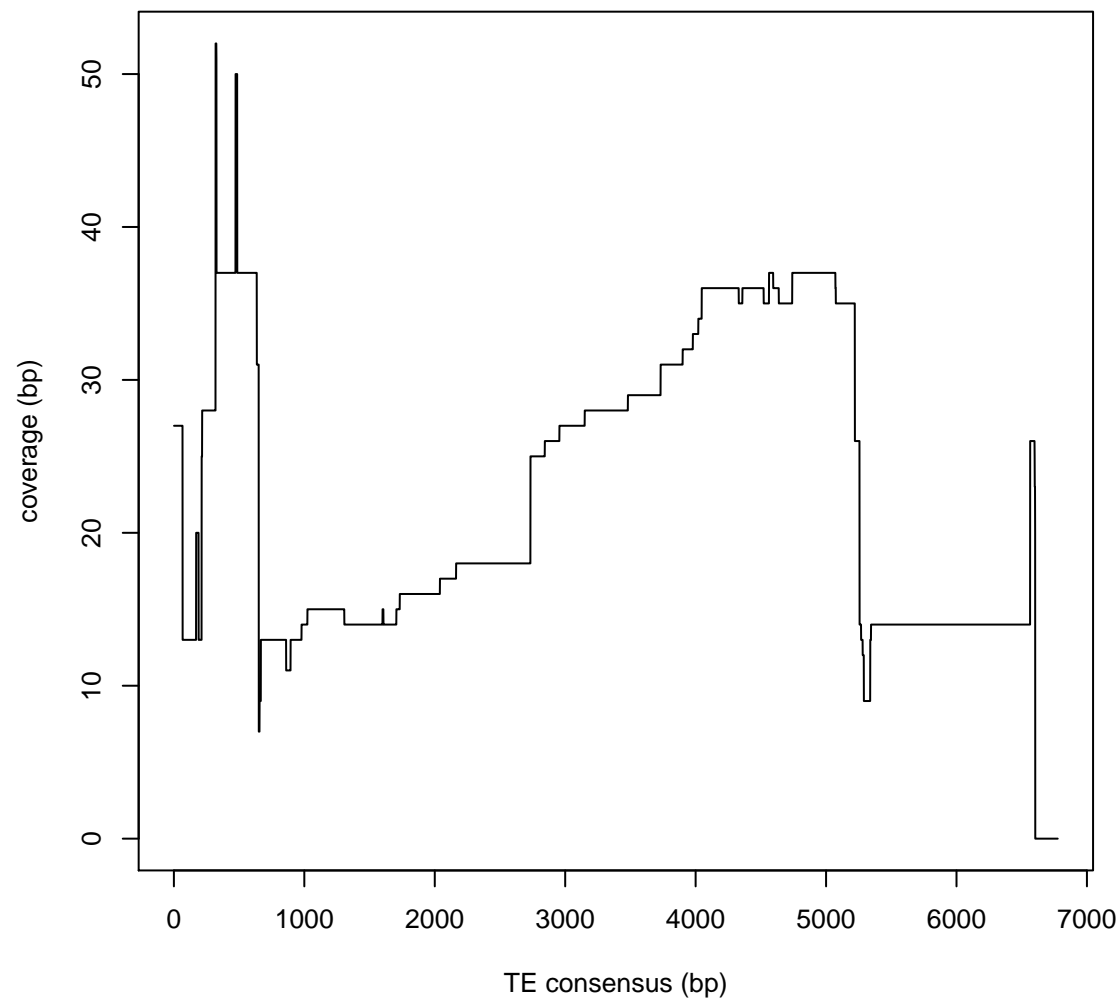


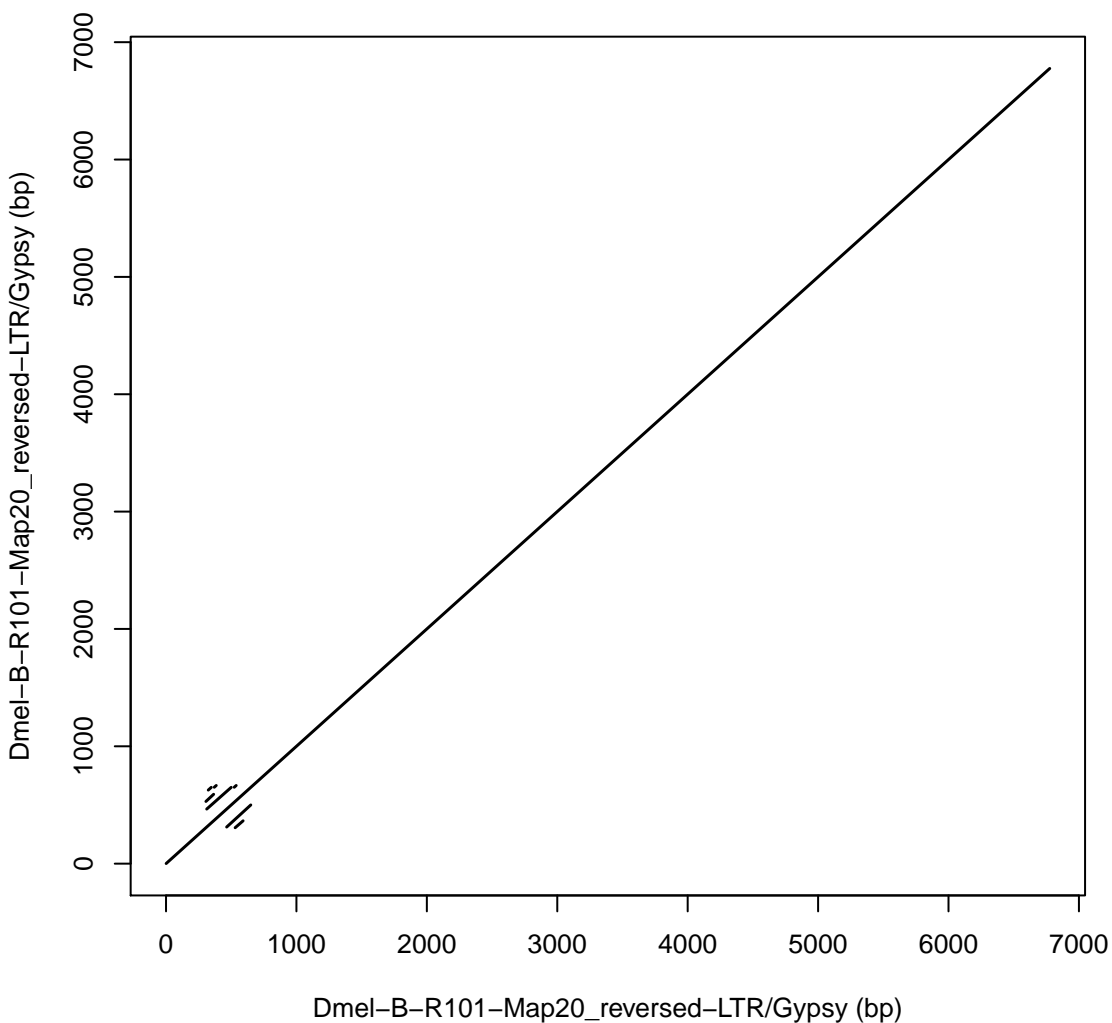
TE: Dmel-B-R101-Map20_reversed-LTR/Gypsy
consensus size: 6776bp; fragments: 141; full length: 2 (>=6098.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

