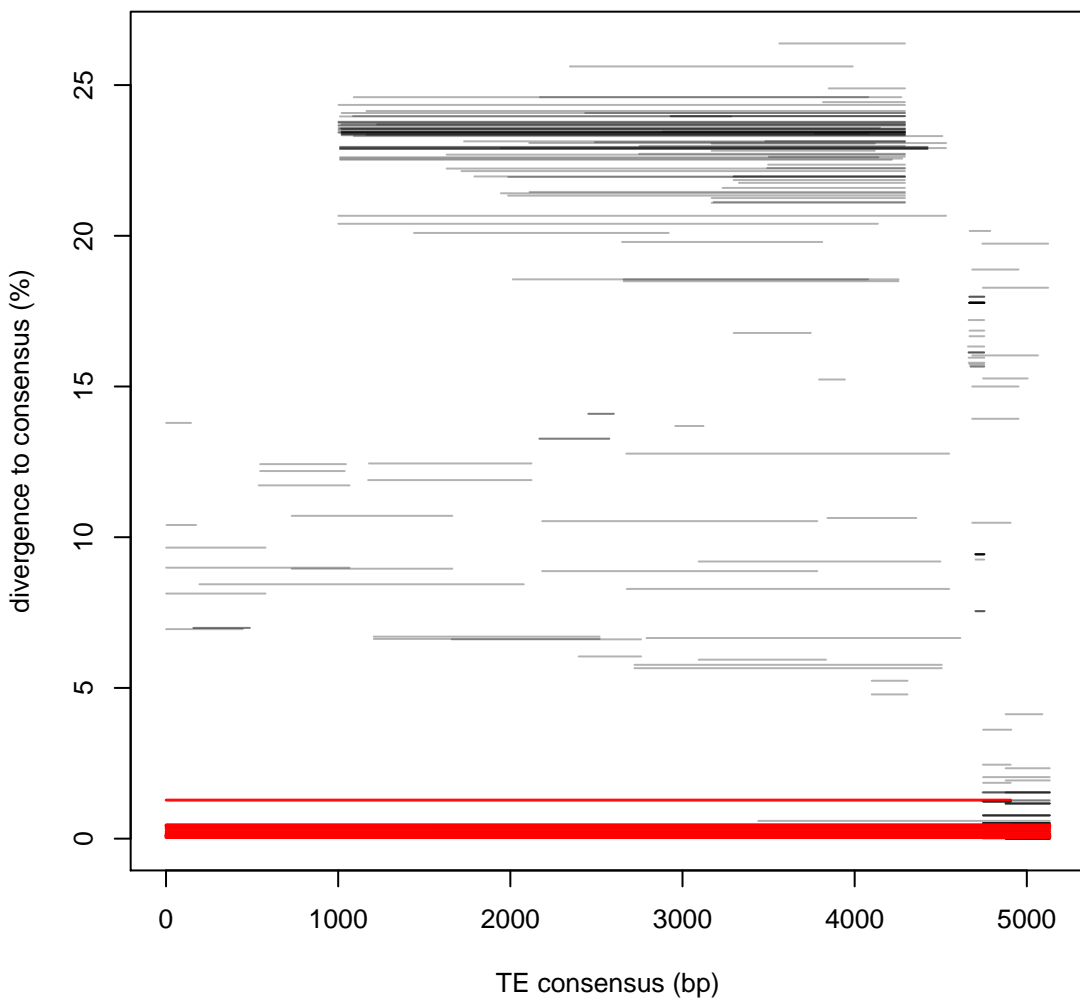
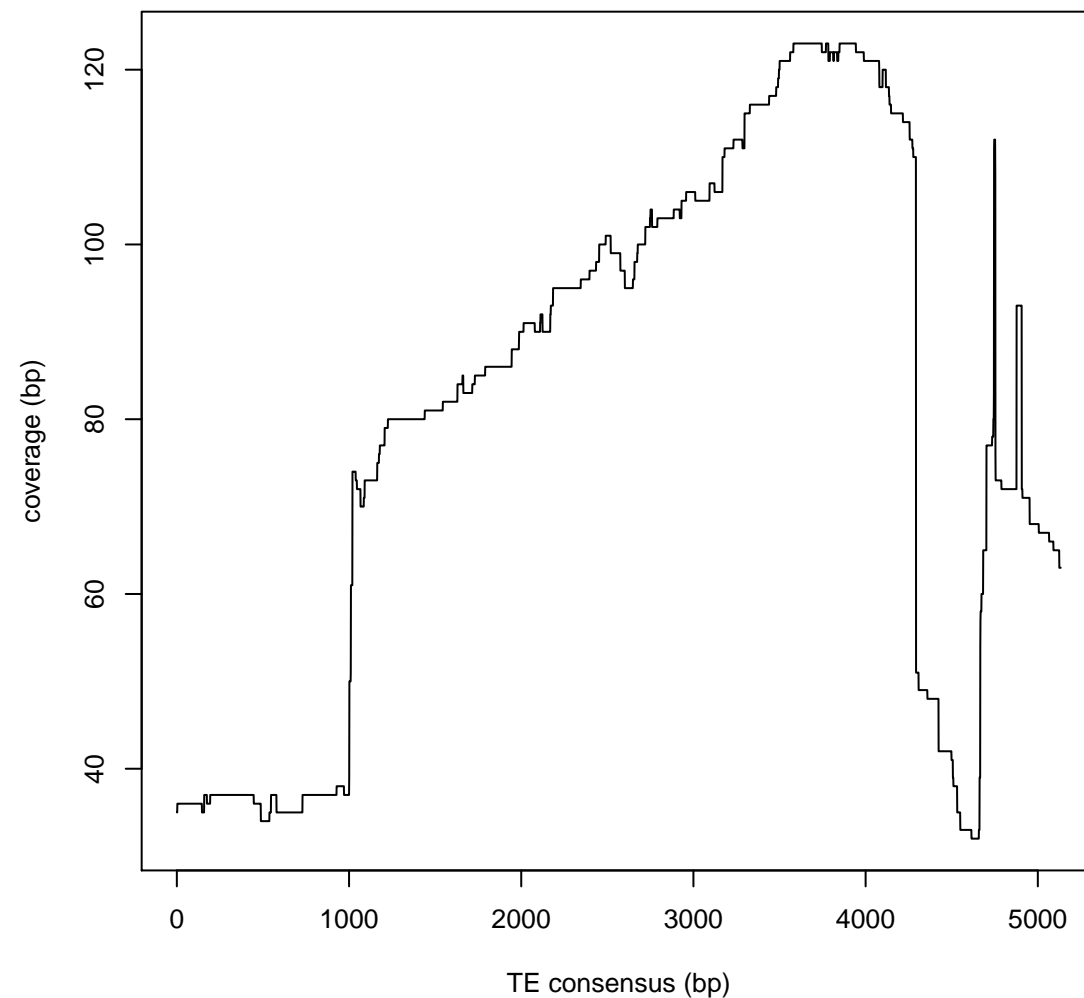


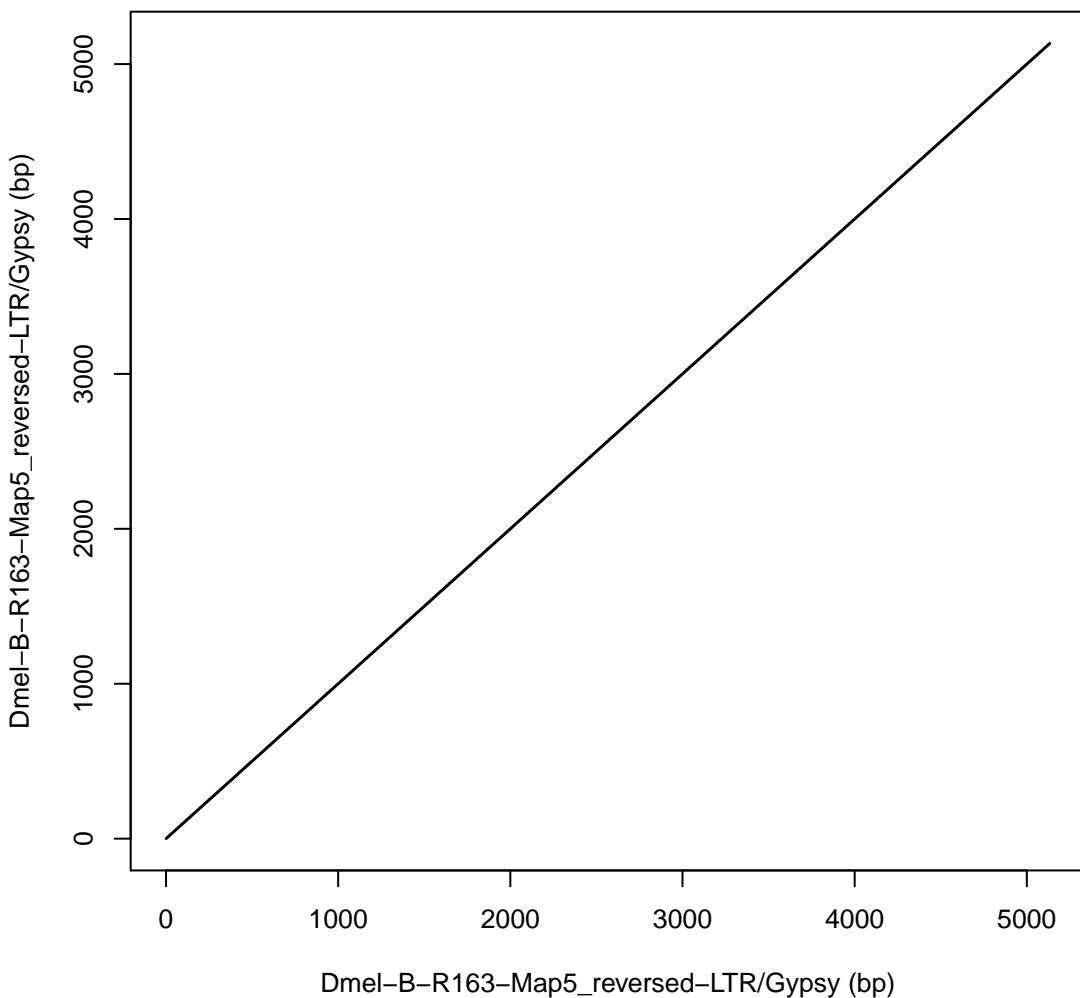
TE: Dmel-B-R163-Map5_reversed-LTR/Gypsy
consensus size: 5133bp; fragments: 260; full length: 28 (>=4619.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

