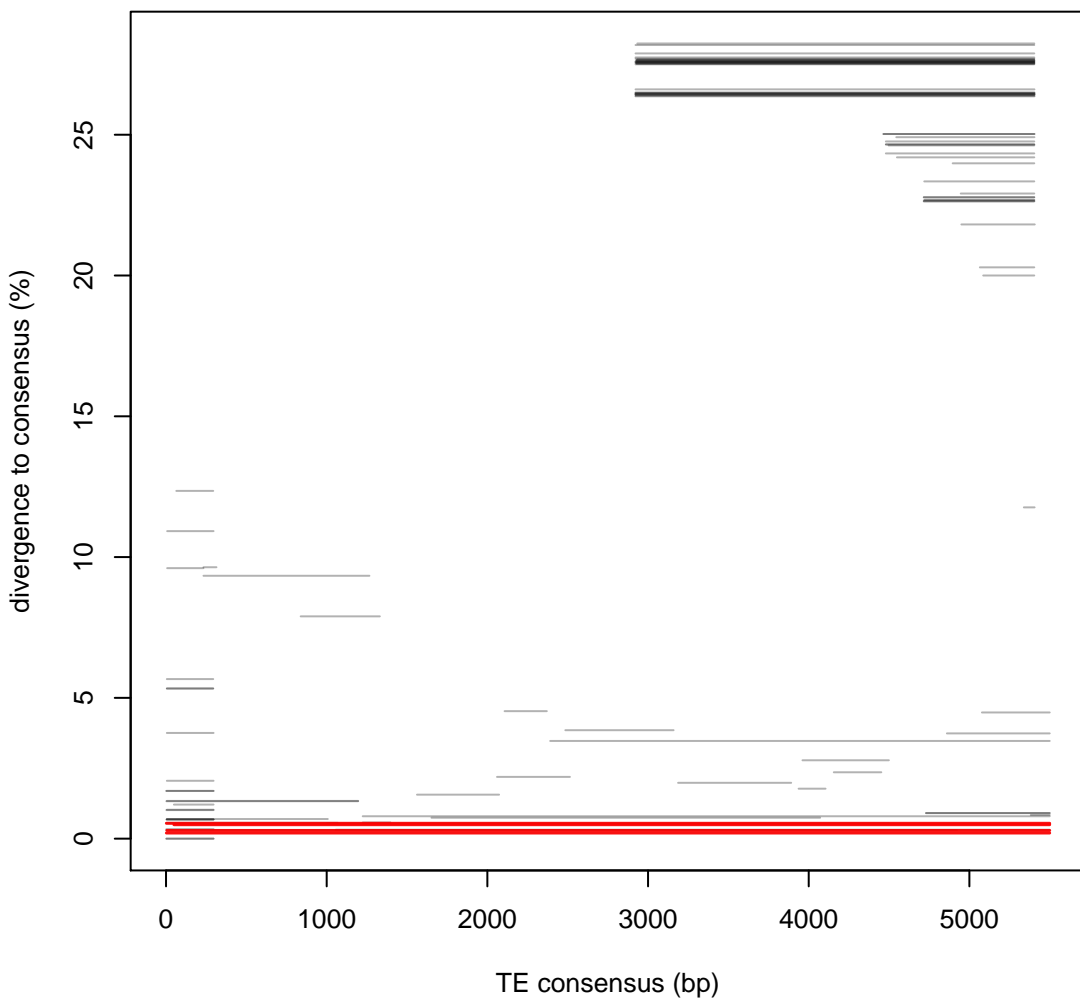
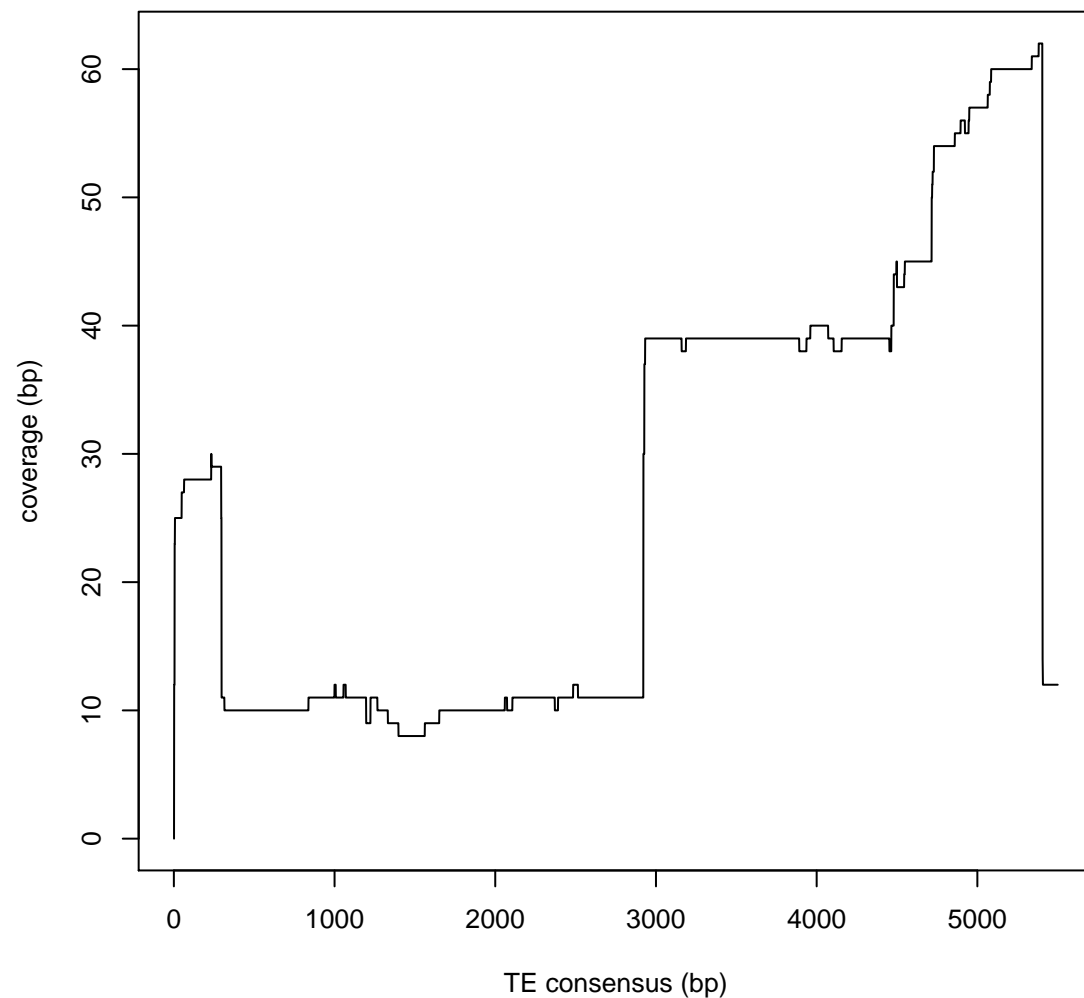


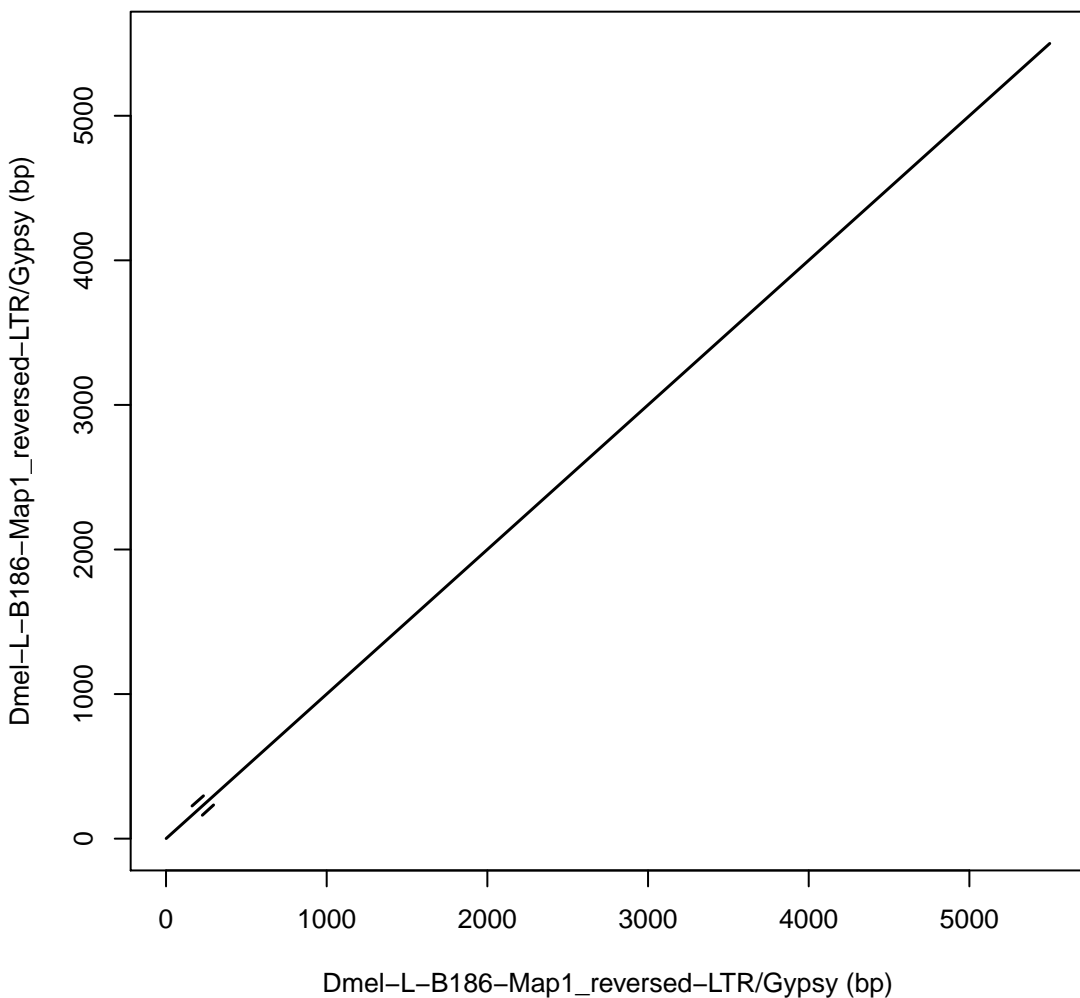
TE: Dmel-L-B186-Map1_reversed-LTR/Gypsy
consensus size: 5500bp; fragments: 100; full length: 4 (>=4950bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

