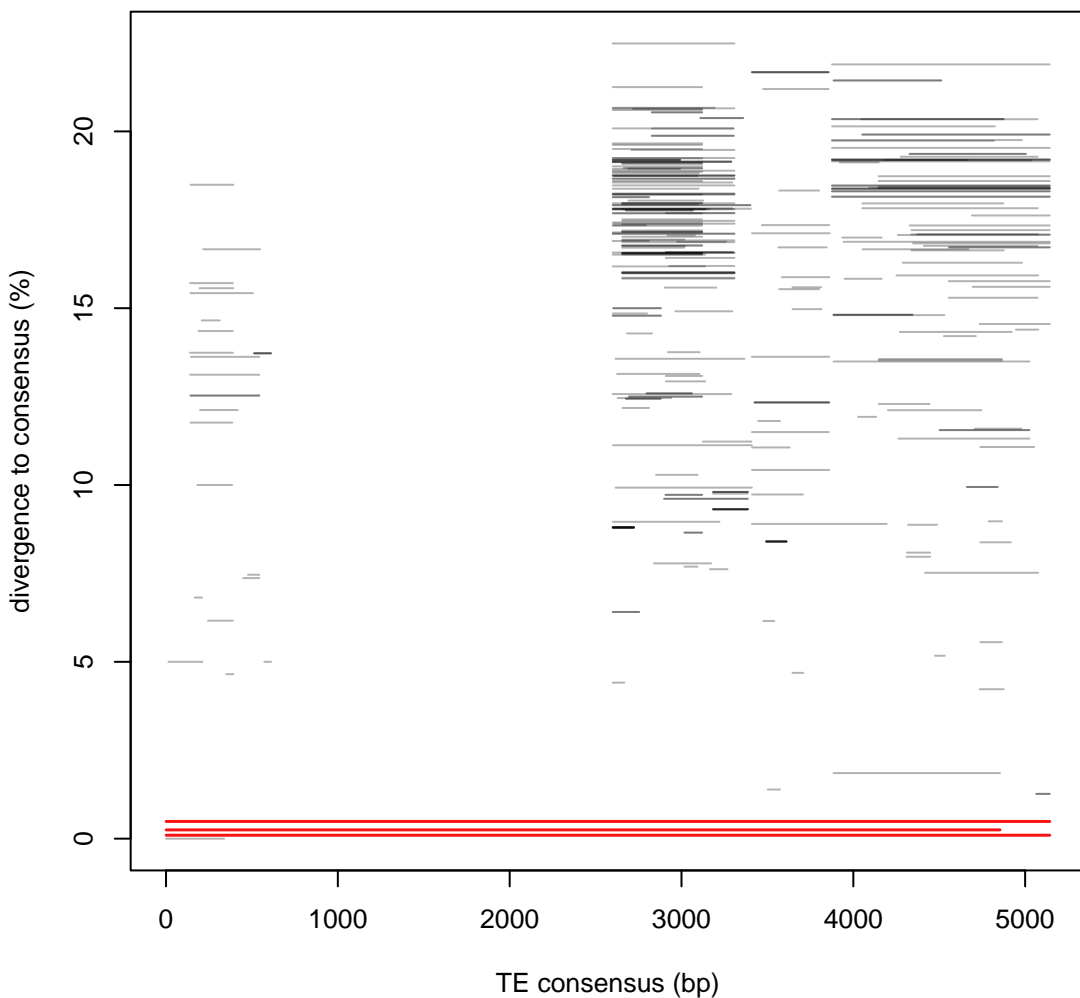
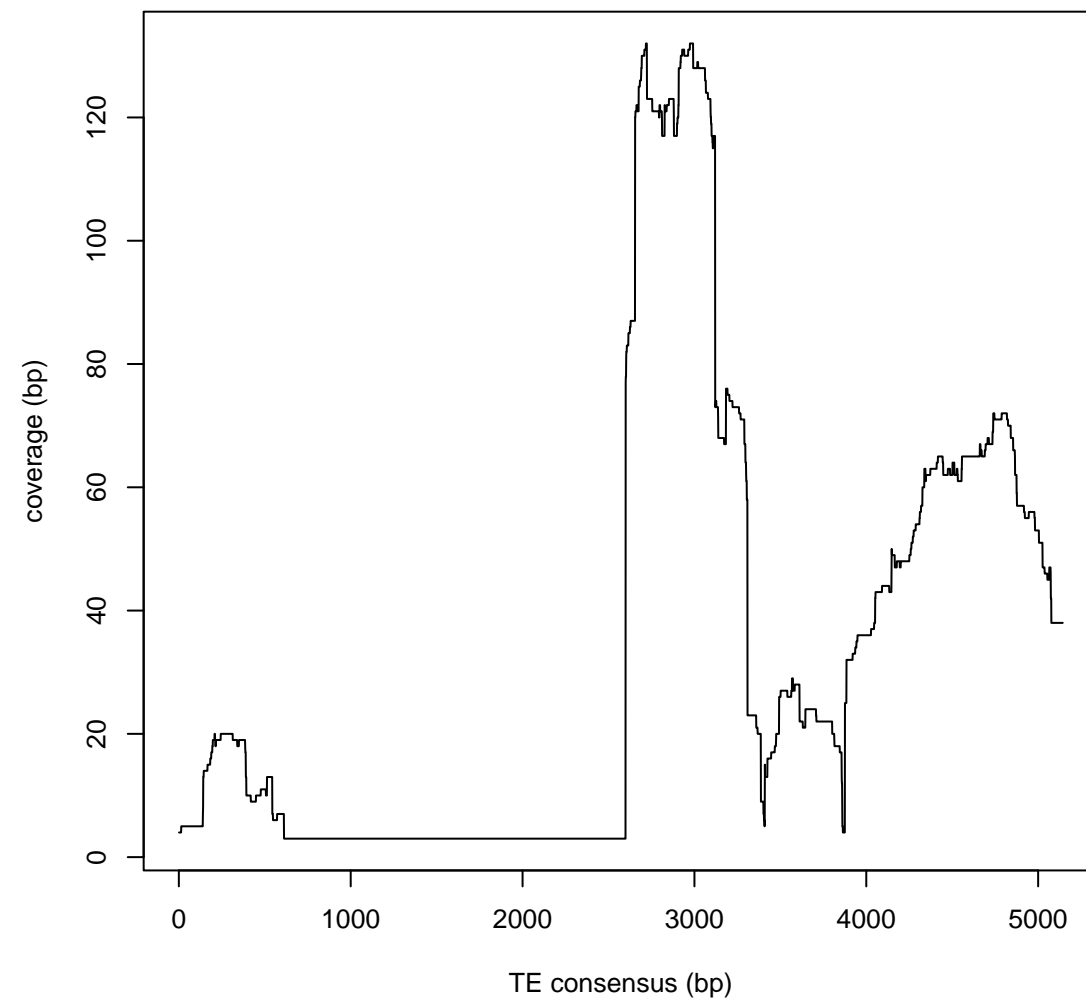


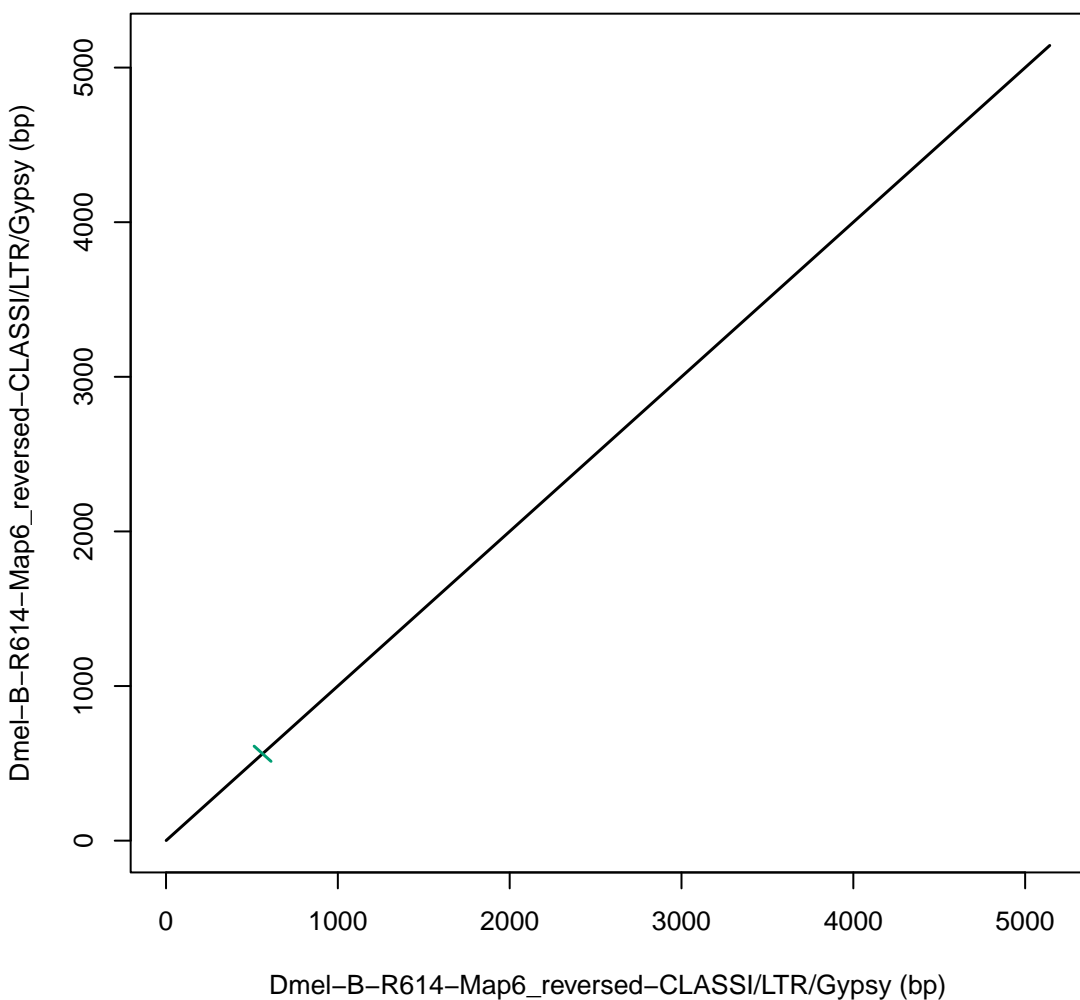
TE: Dmel-B-R614-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 5143bp; fragments: 321; full length: 3 (>=4628.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

