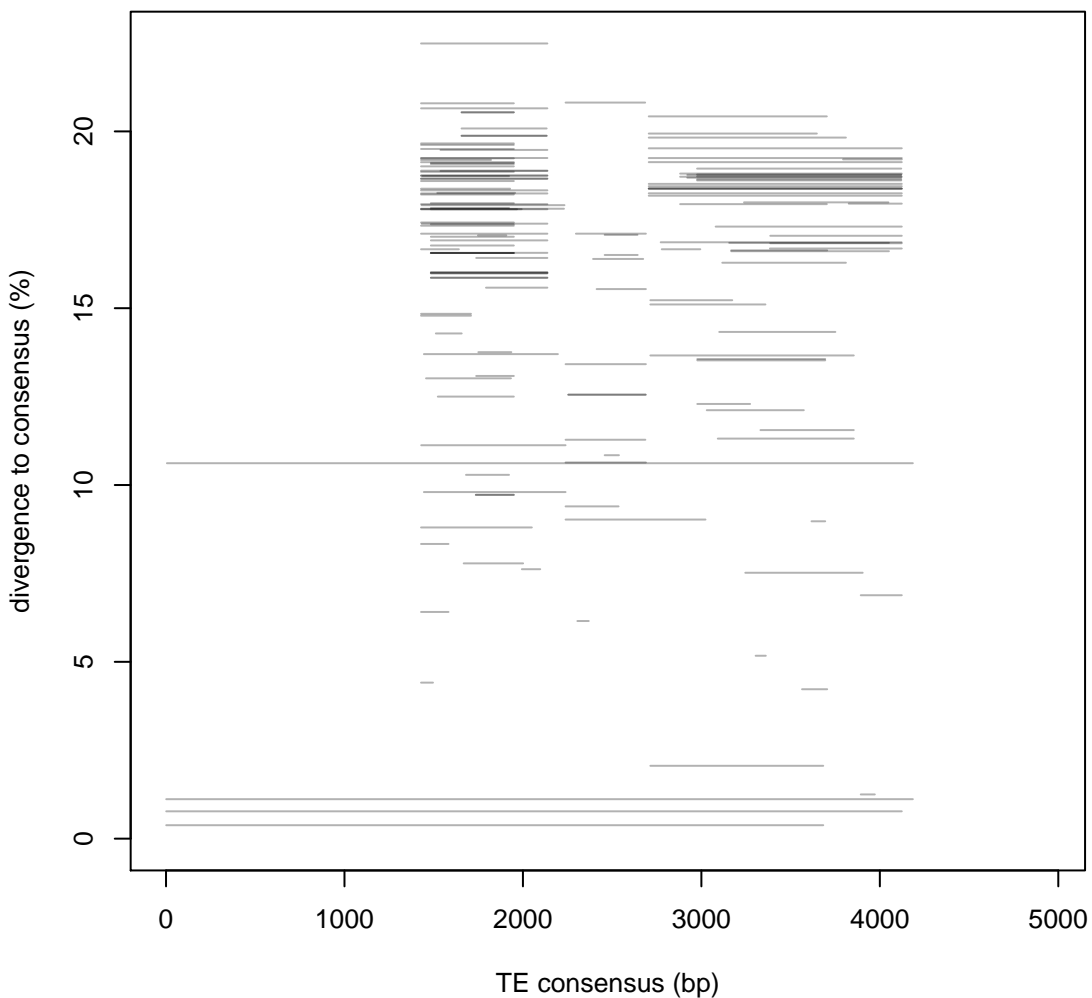
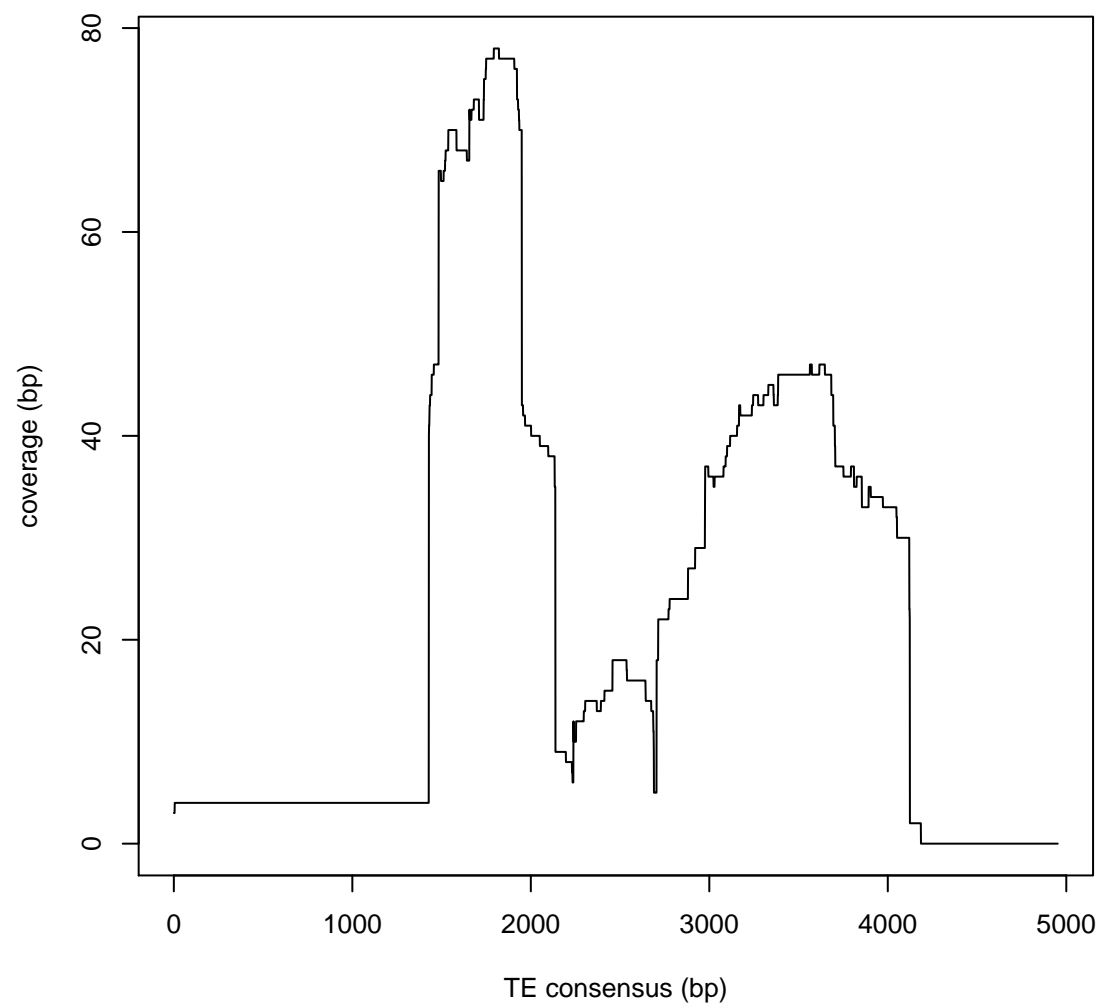


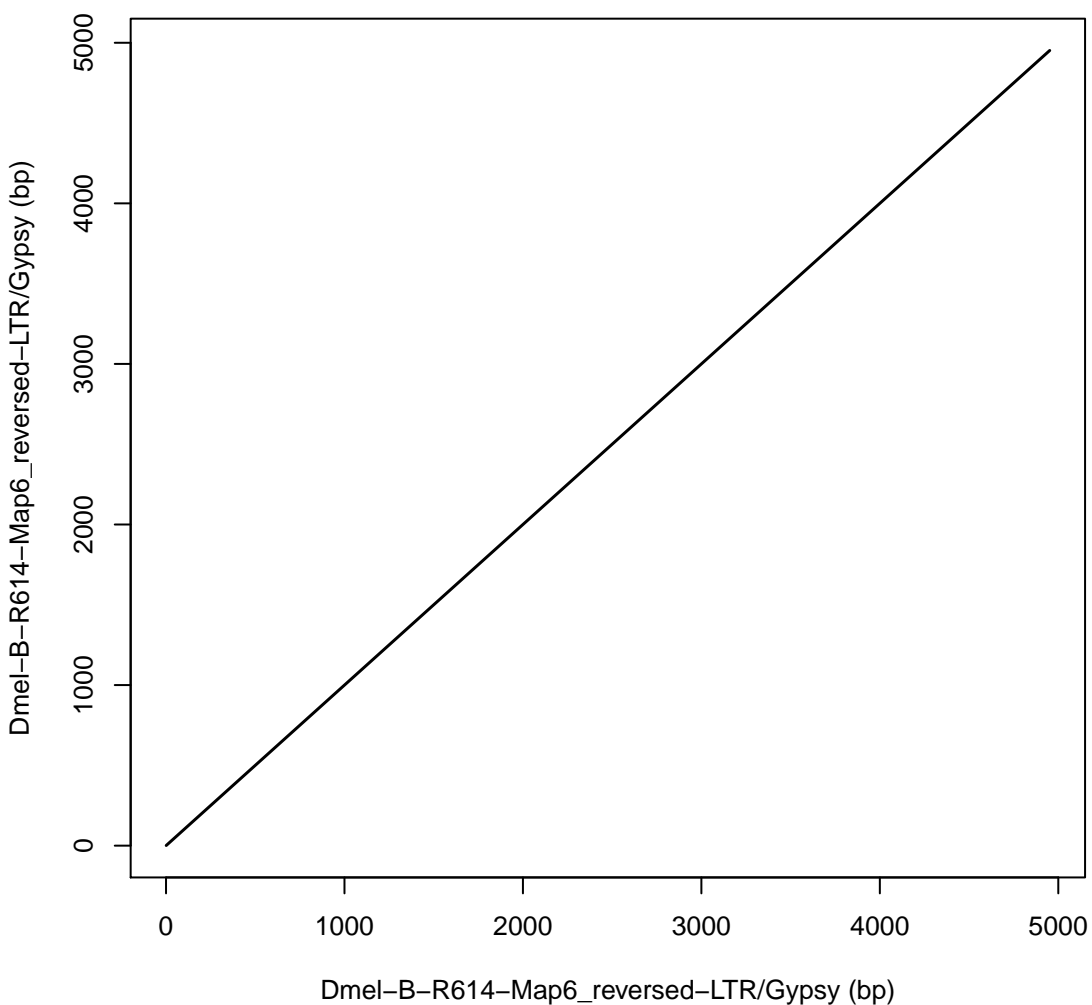
TE: Dmel-B-R614-Map6\_reversed-LTR/Gypsy  
consensus size: 4952bp; fragments: 154; full length: 0 (>=4456.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

