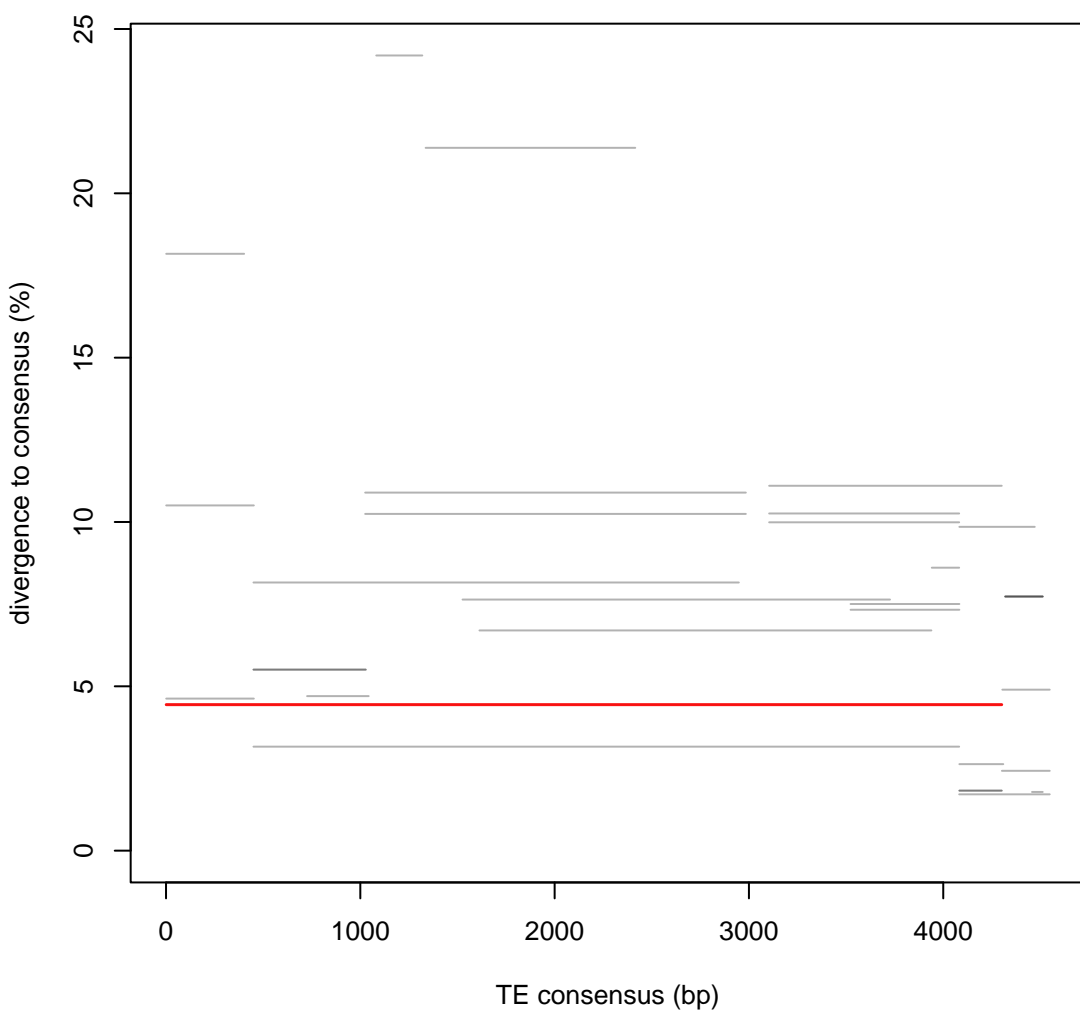
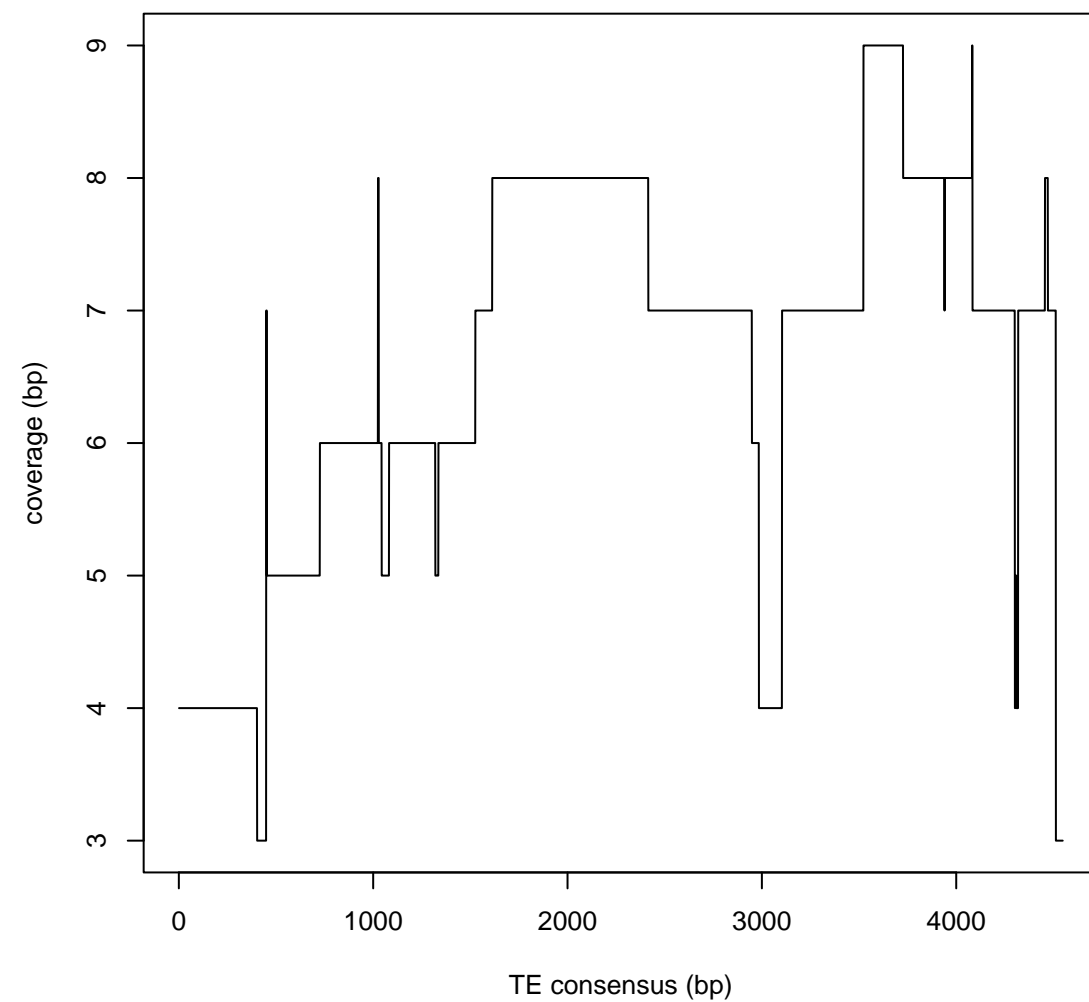


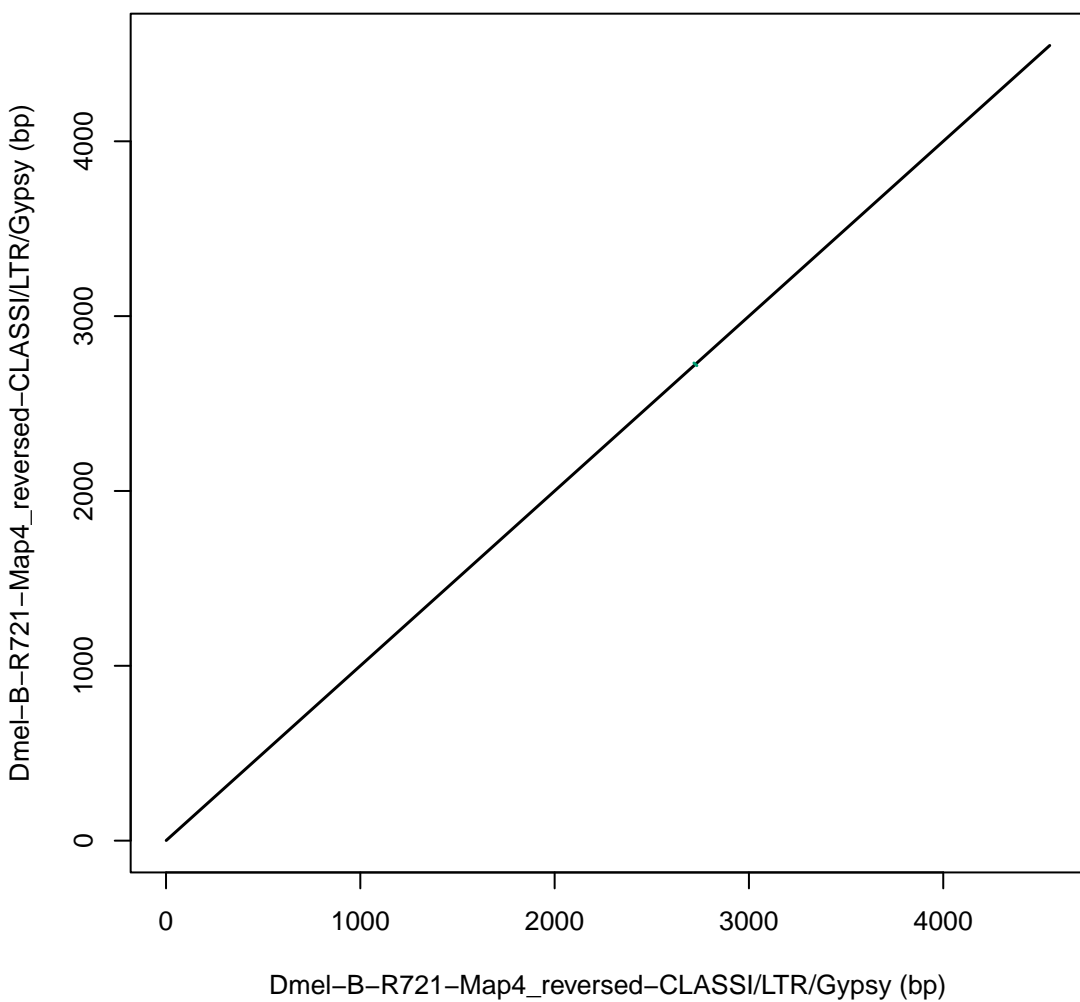
TE: Dmel-B-R721-Map4\_reversed-CLASSI/LTR/Gypsy  
consensus size: 4548bp; fragments: 32; full length: 1 (>=4093.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

