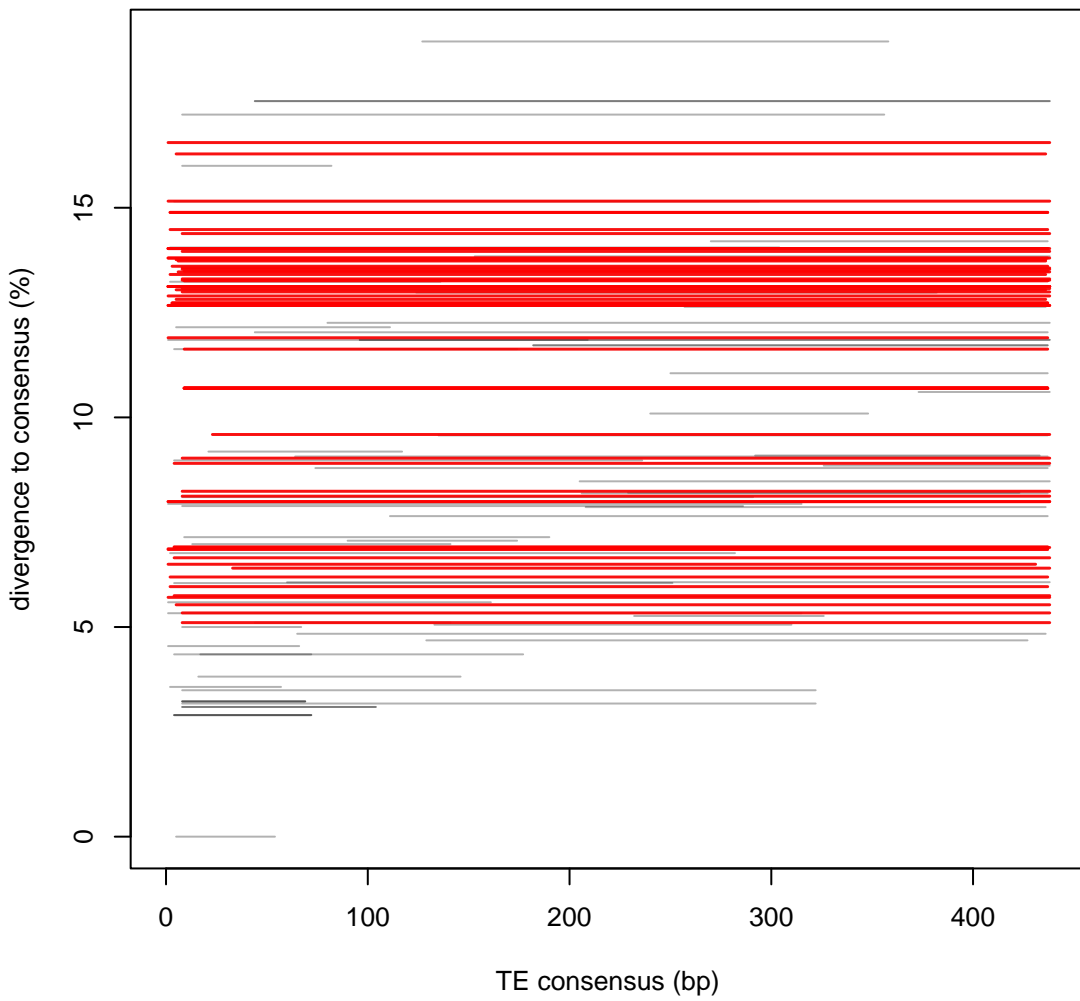
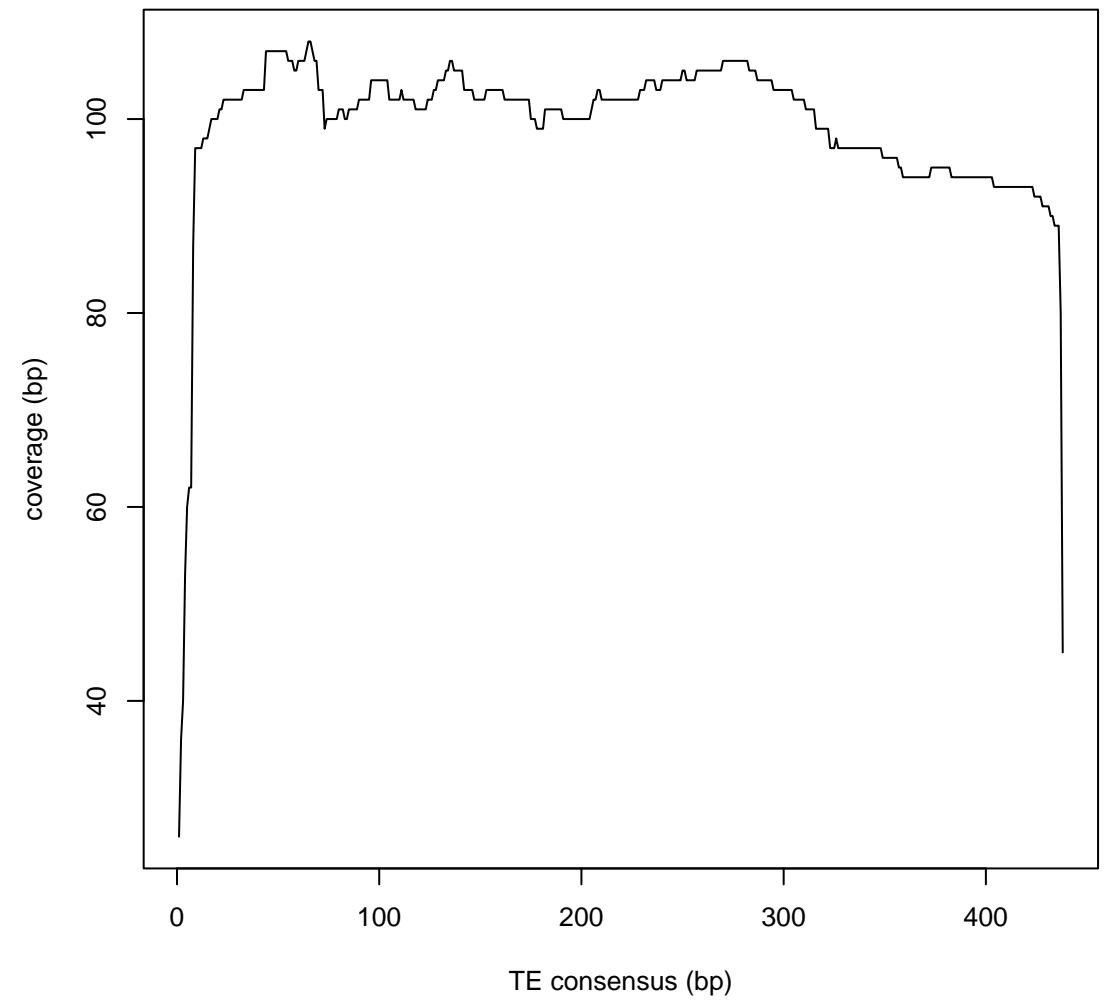


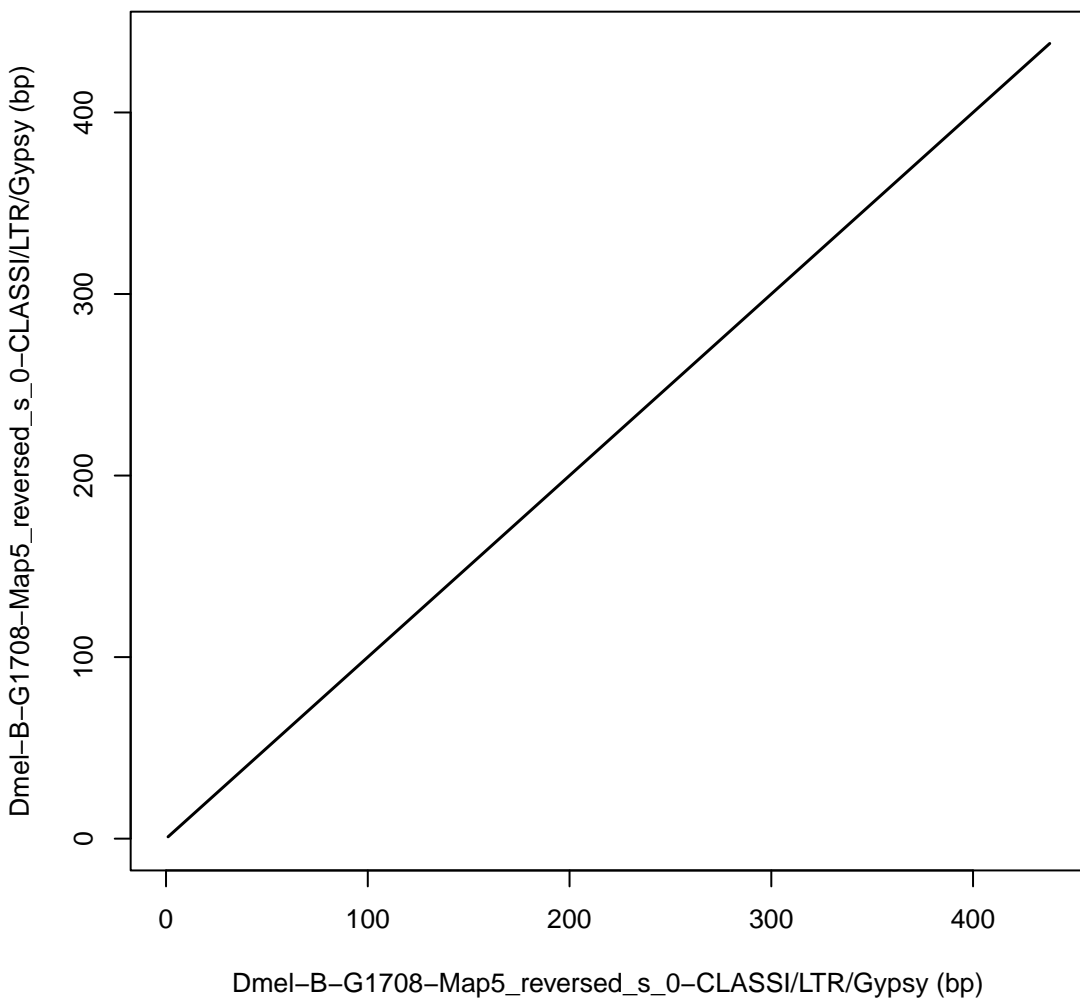
TE: Dmel-B-G1708-Map5_reversed_s_0-CLASSI/LTR/Gypsy
consensus size: 438bp; fragments: 137; full length: 67 (>=394.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

