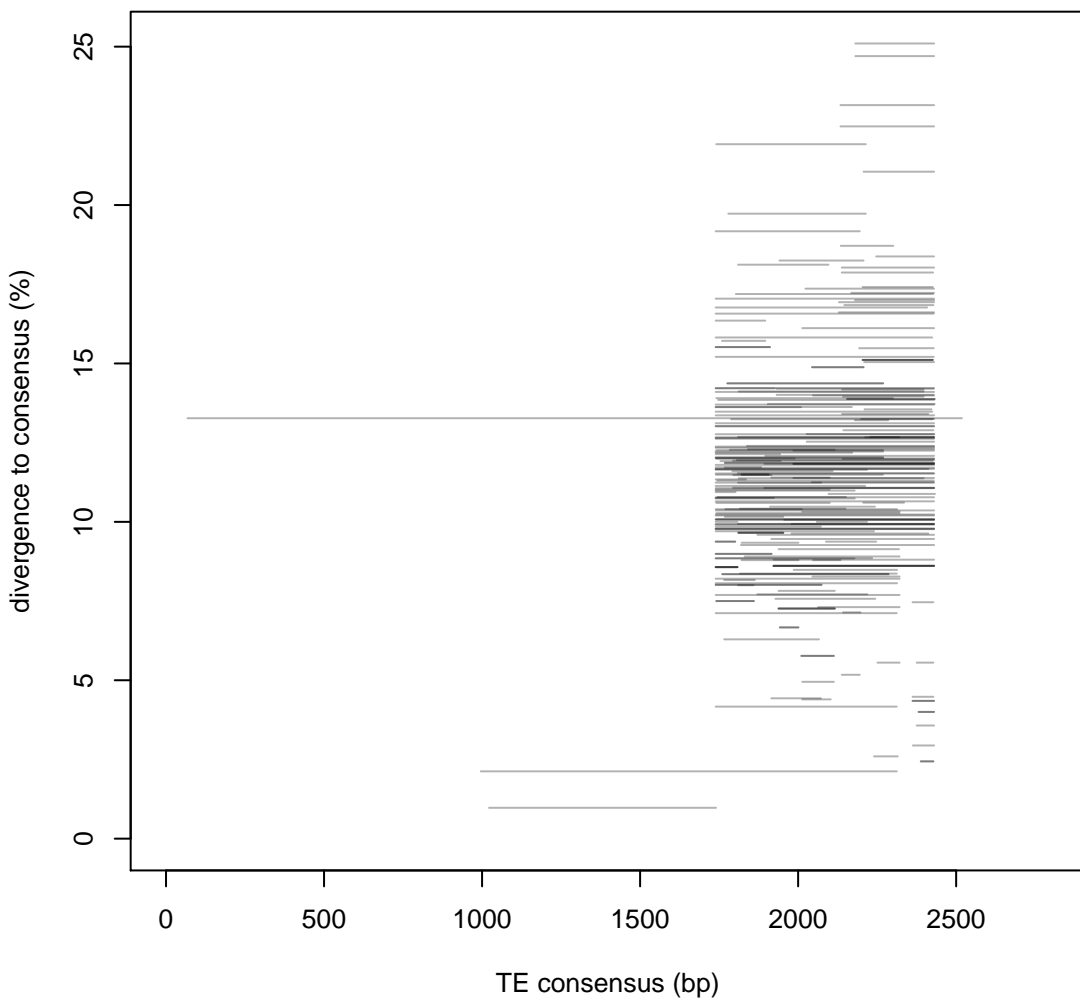
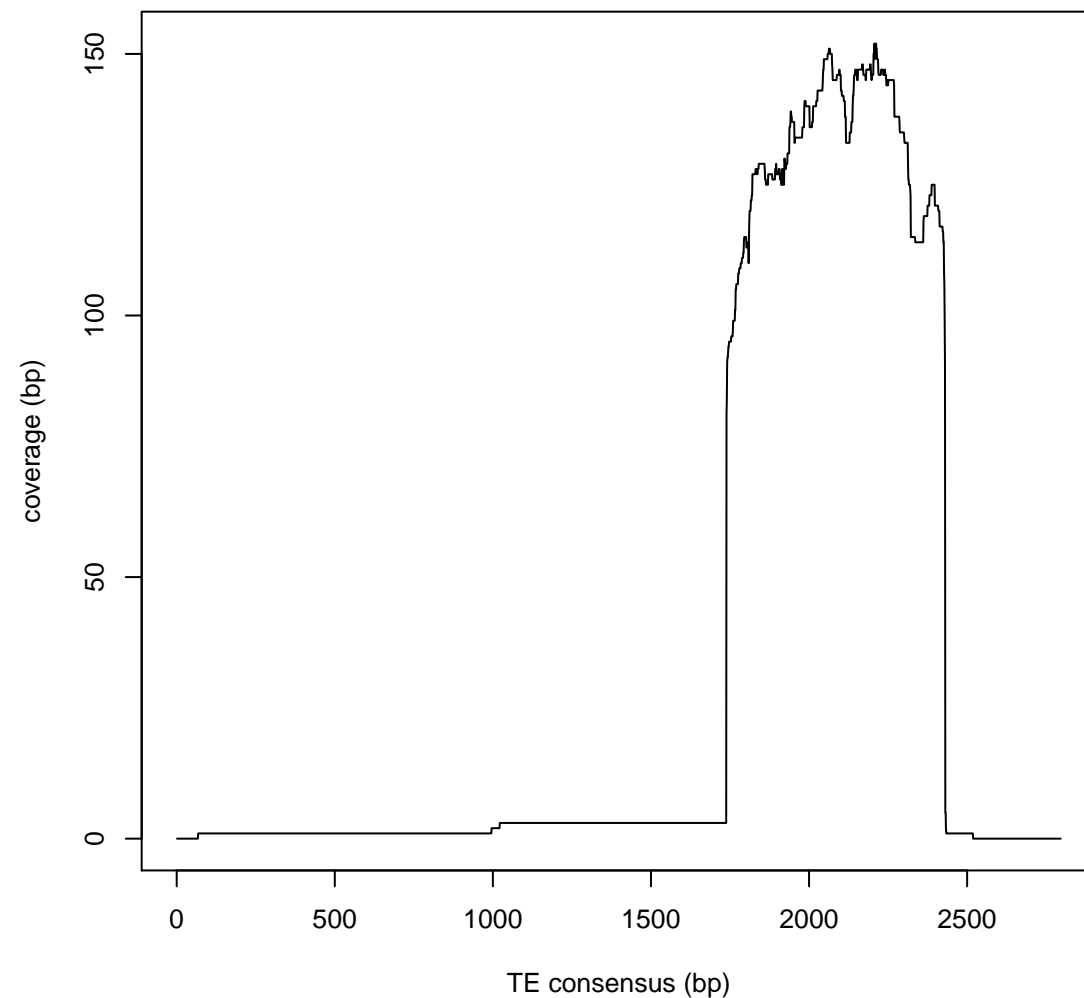


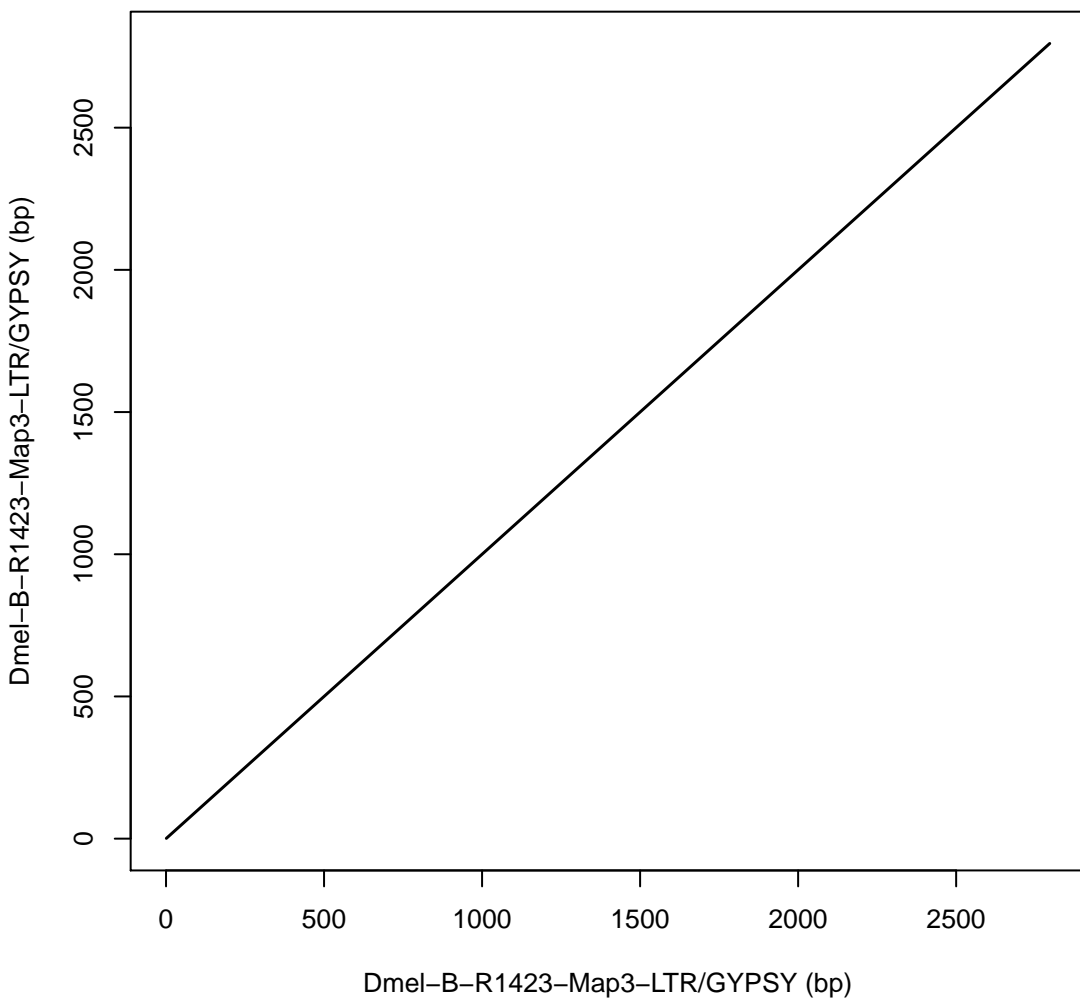
TE: Dmel-B-R1423-Map3-LTR/Gypsy  
consensus size: 2796bp; fragments: 240; full length: 0 (>=2516.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

