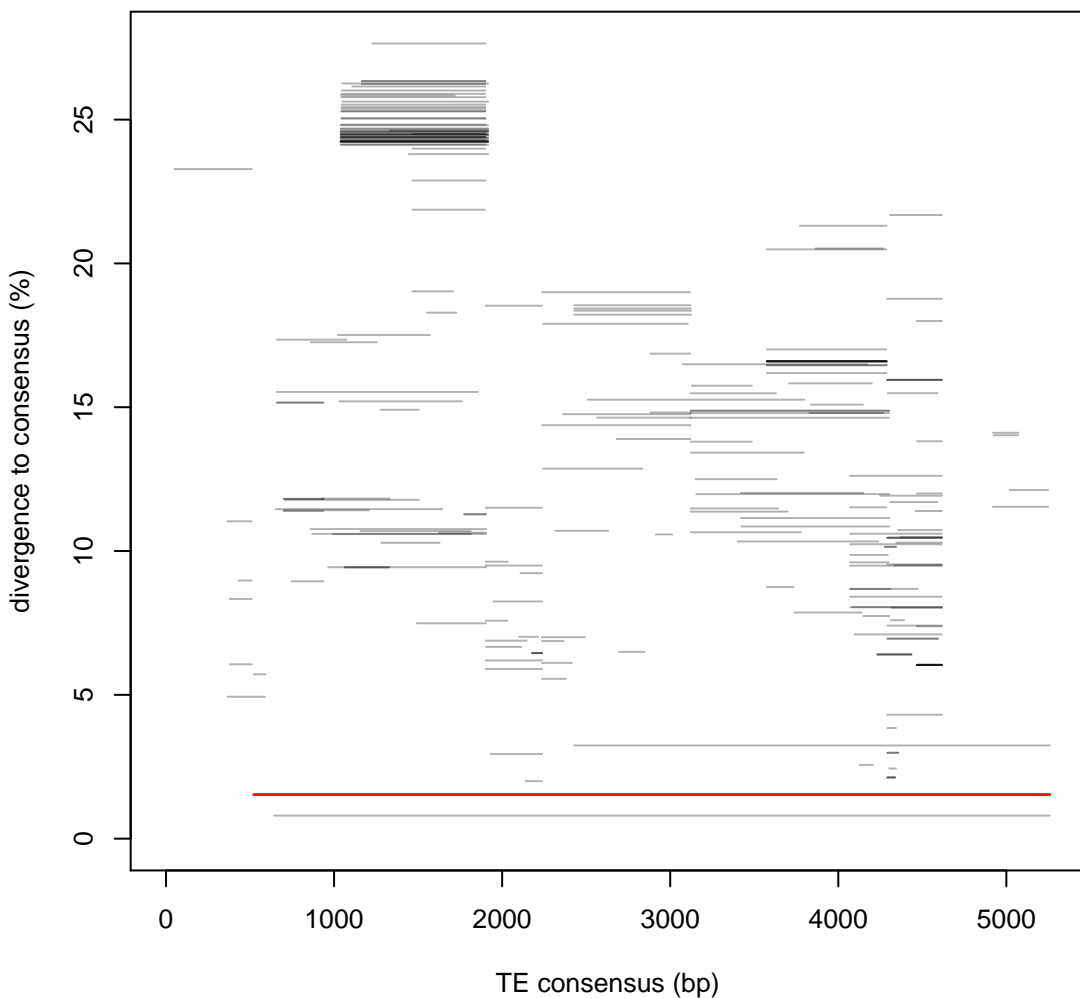
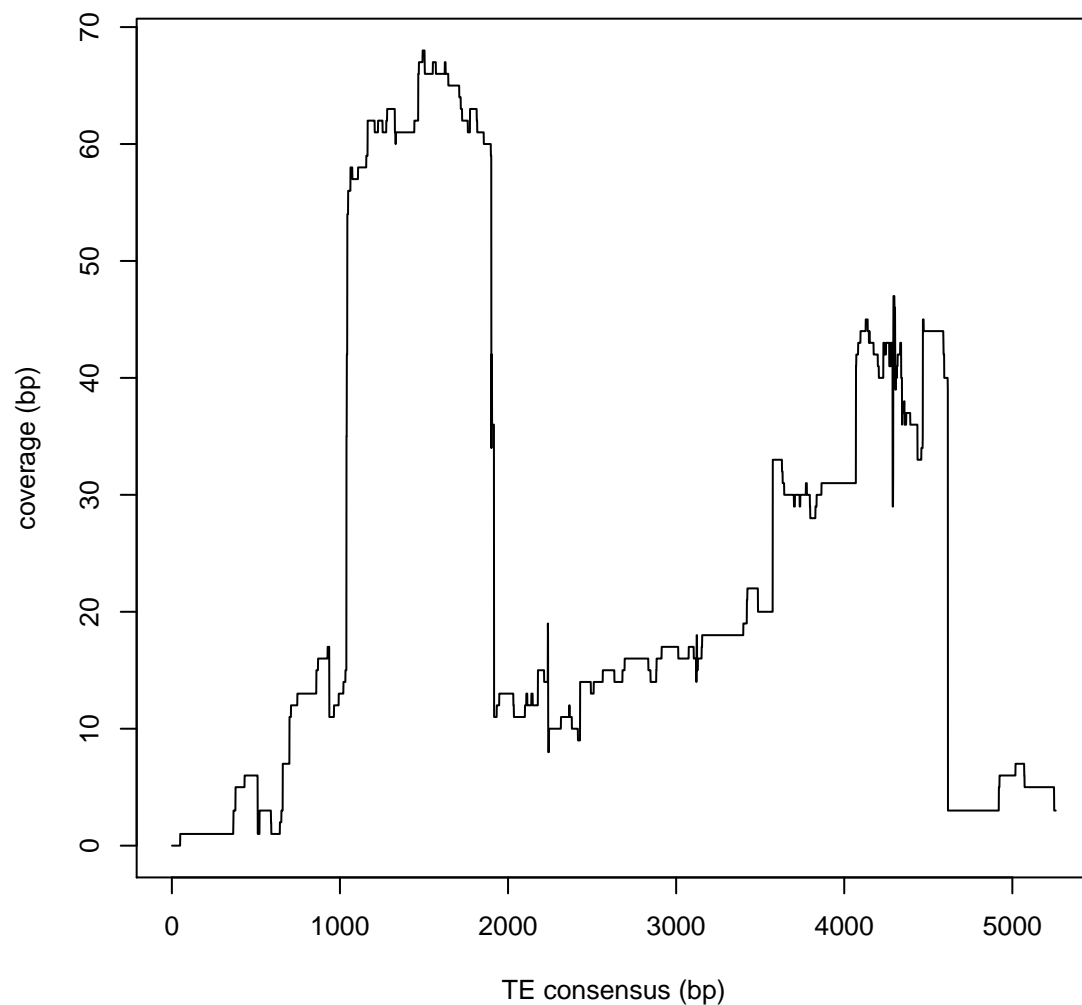


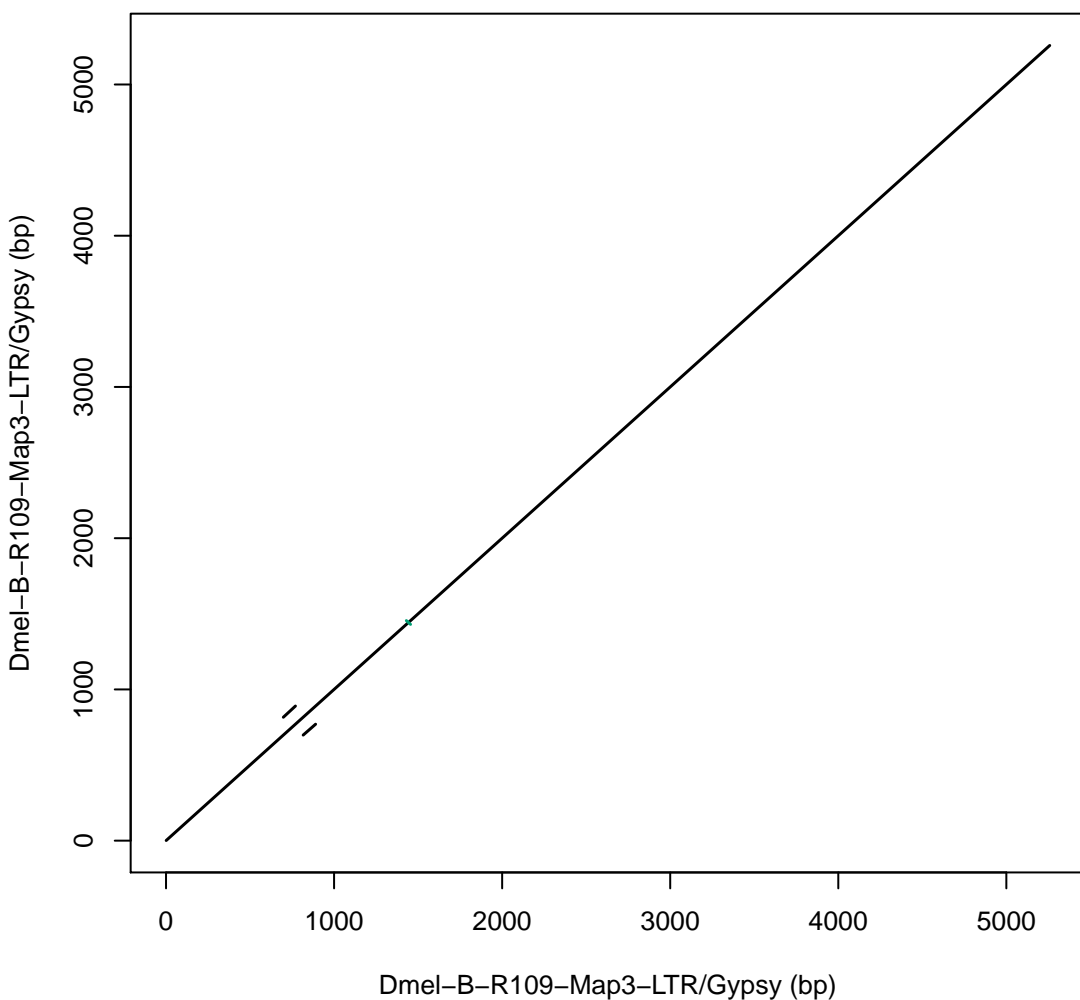
TE: Dmel-B-R109-Map3-LTR/Gypsy  
consensus size: 5258bp; fragments: 232; full length: 1 (>=4732.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

