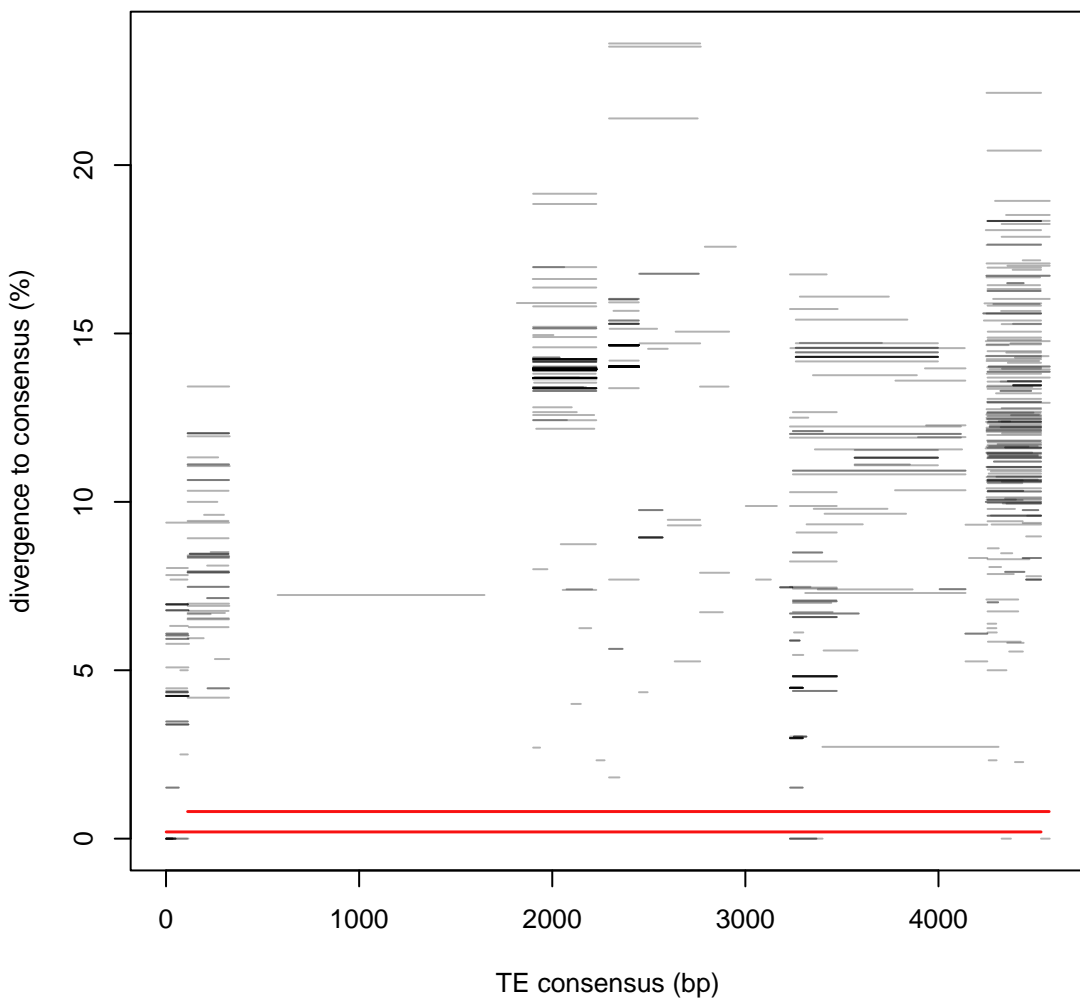
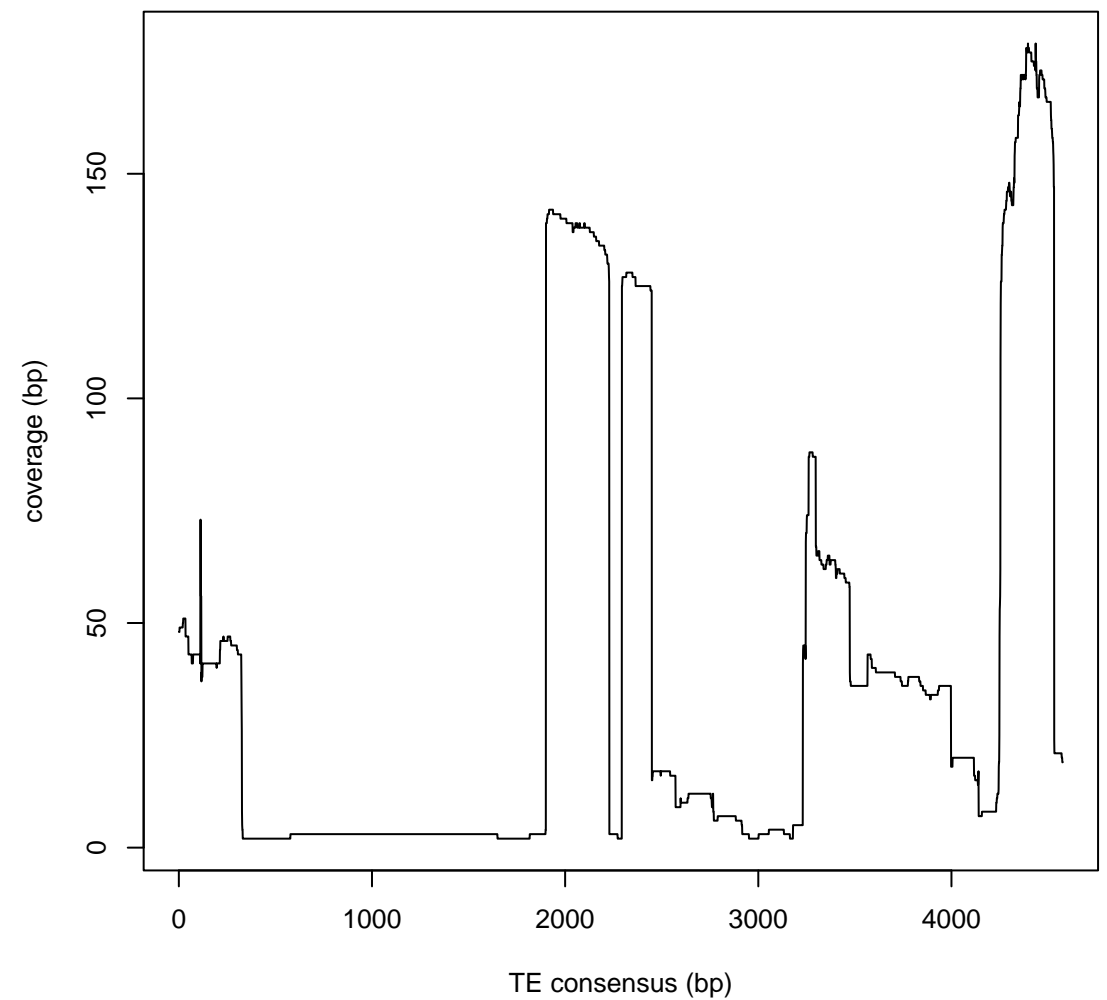


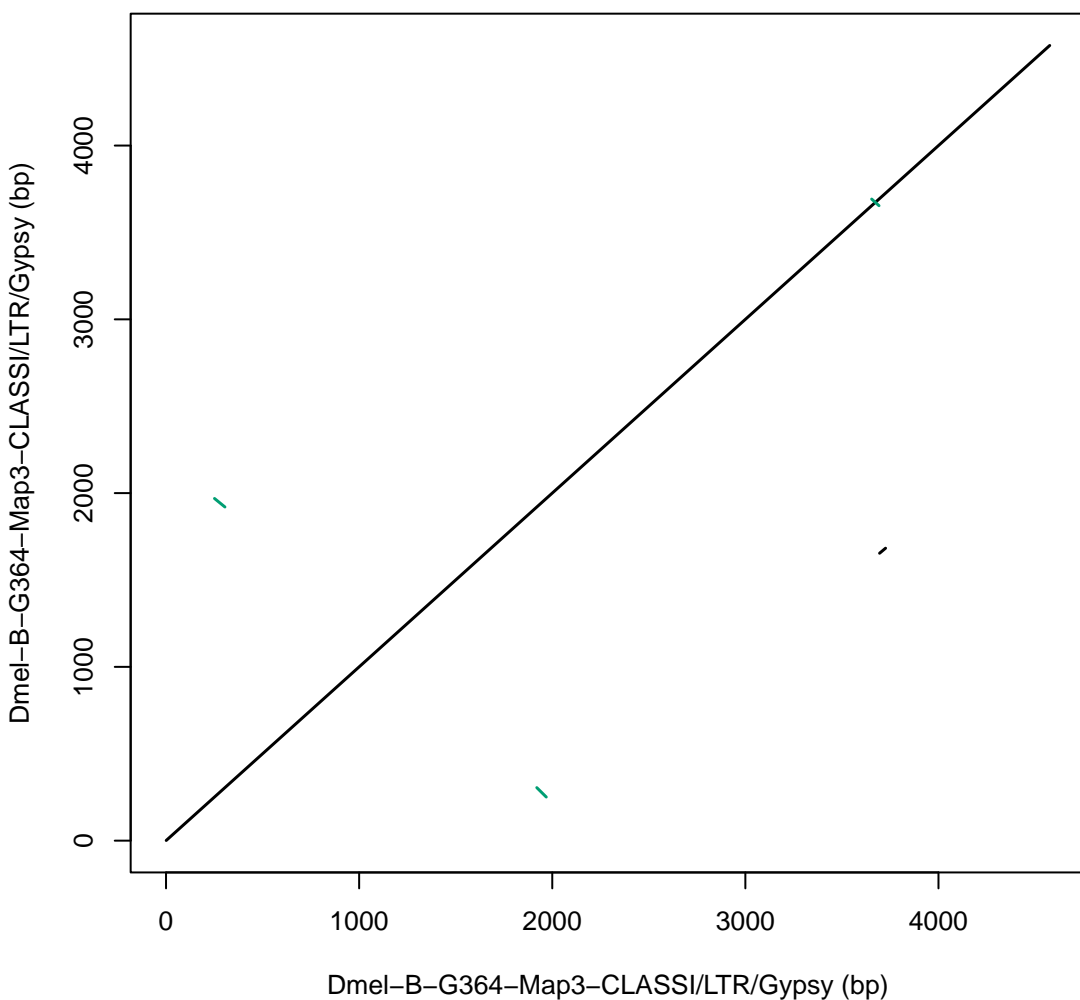
TE: Dmel-B-G364-Map3-CLASSI/LTR/Gypsy
consensus size: 4576bp; fragments: 718; full length: 2 (>=4118.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

