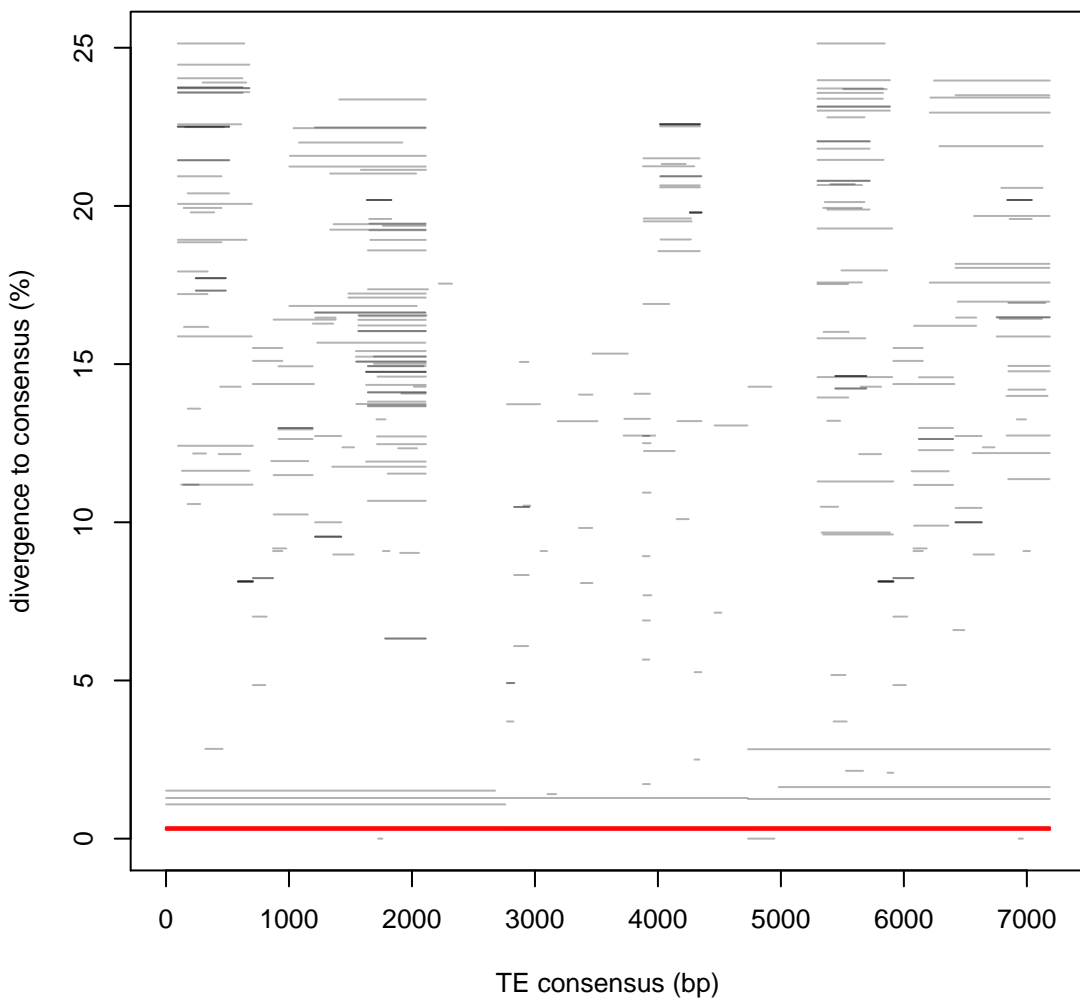
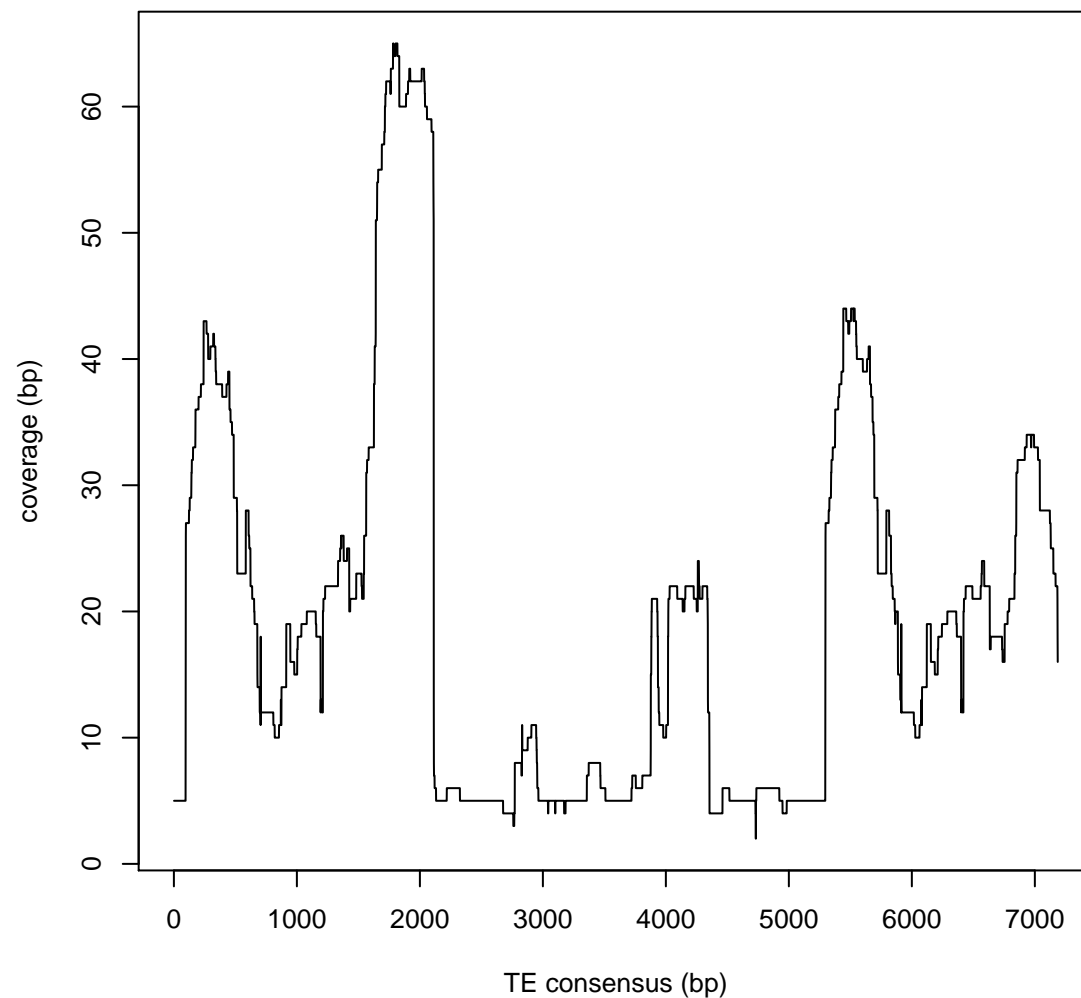


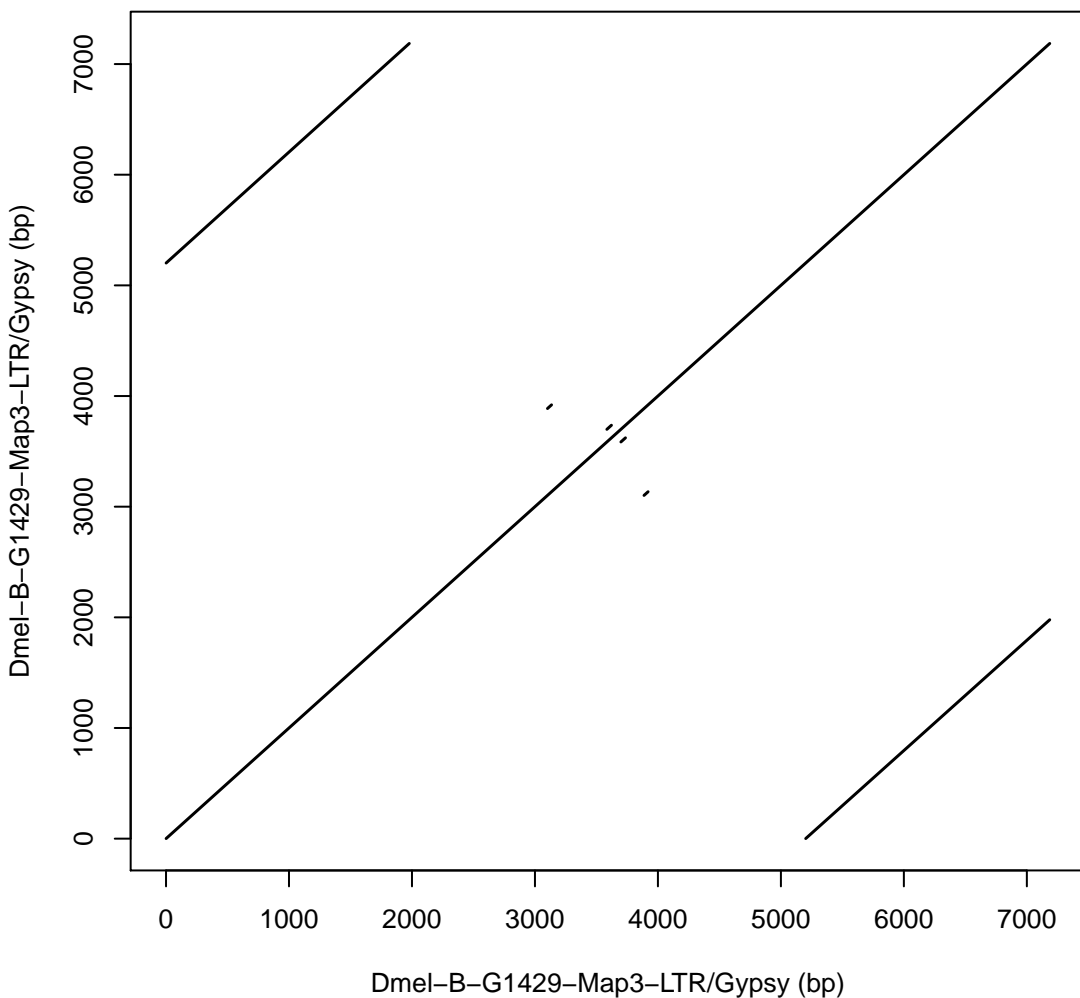
TE: Dmel-B-G1429-Map3-LTR/Gypsy
consensus size: 7186bp; fragments: 315; full length: 2 (>=6467.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

