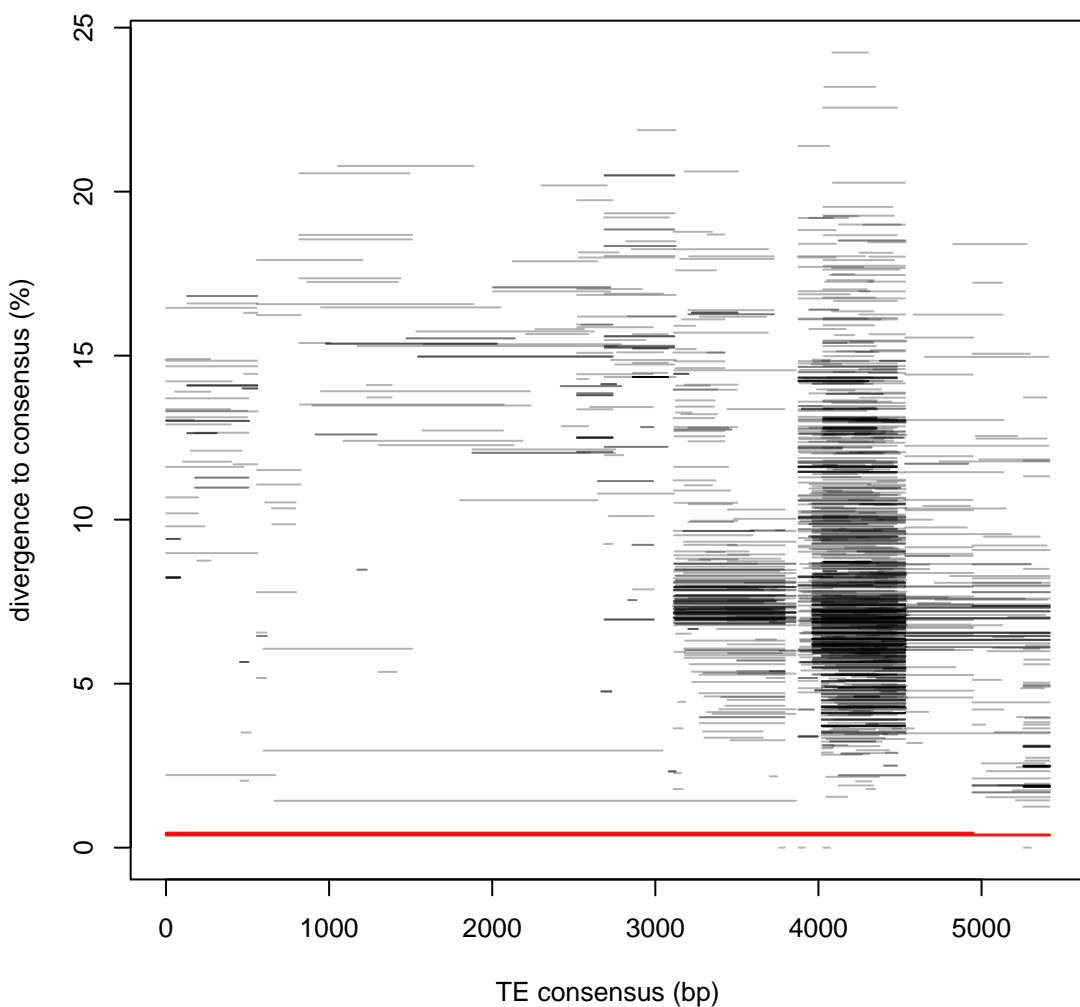
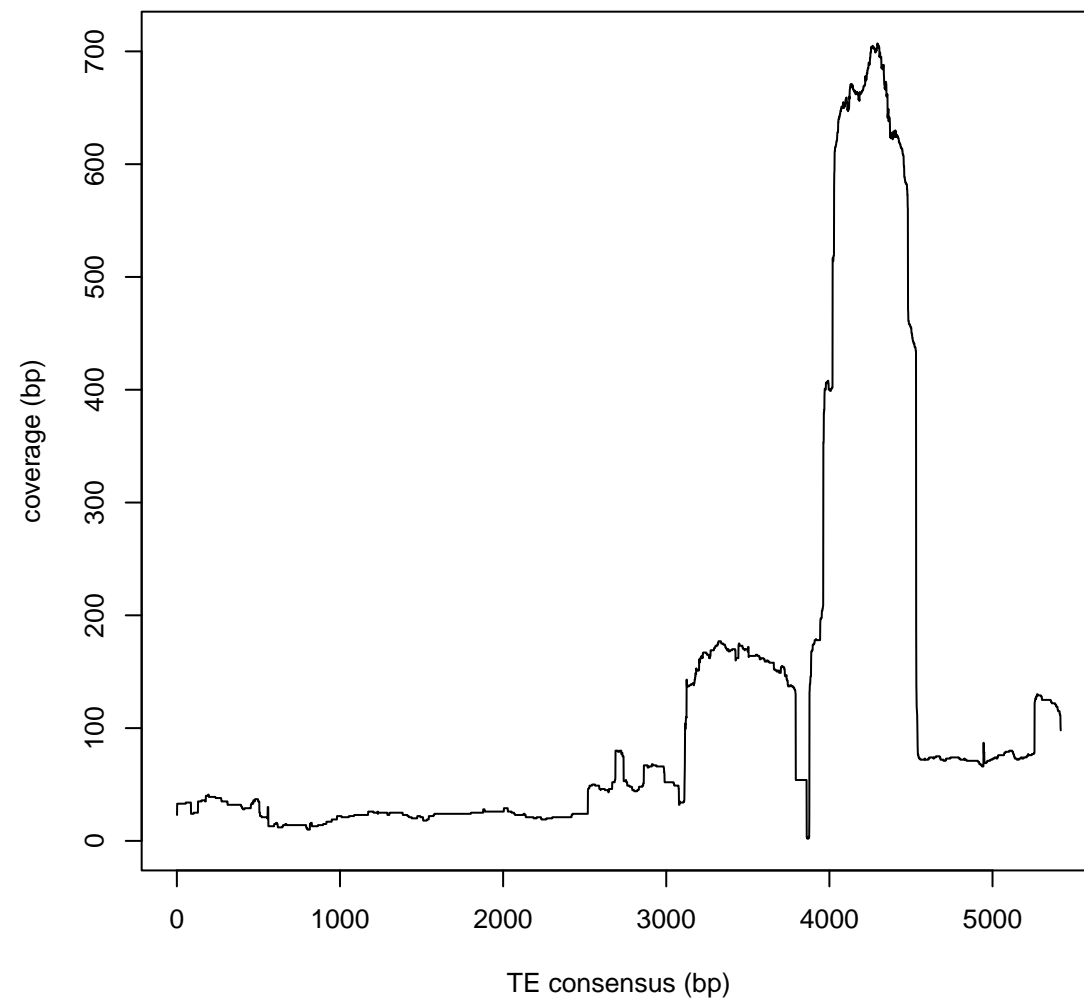


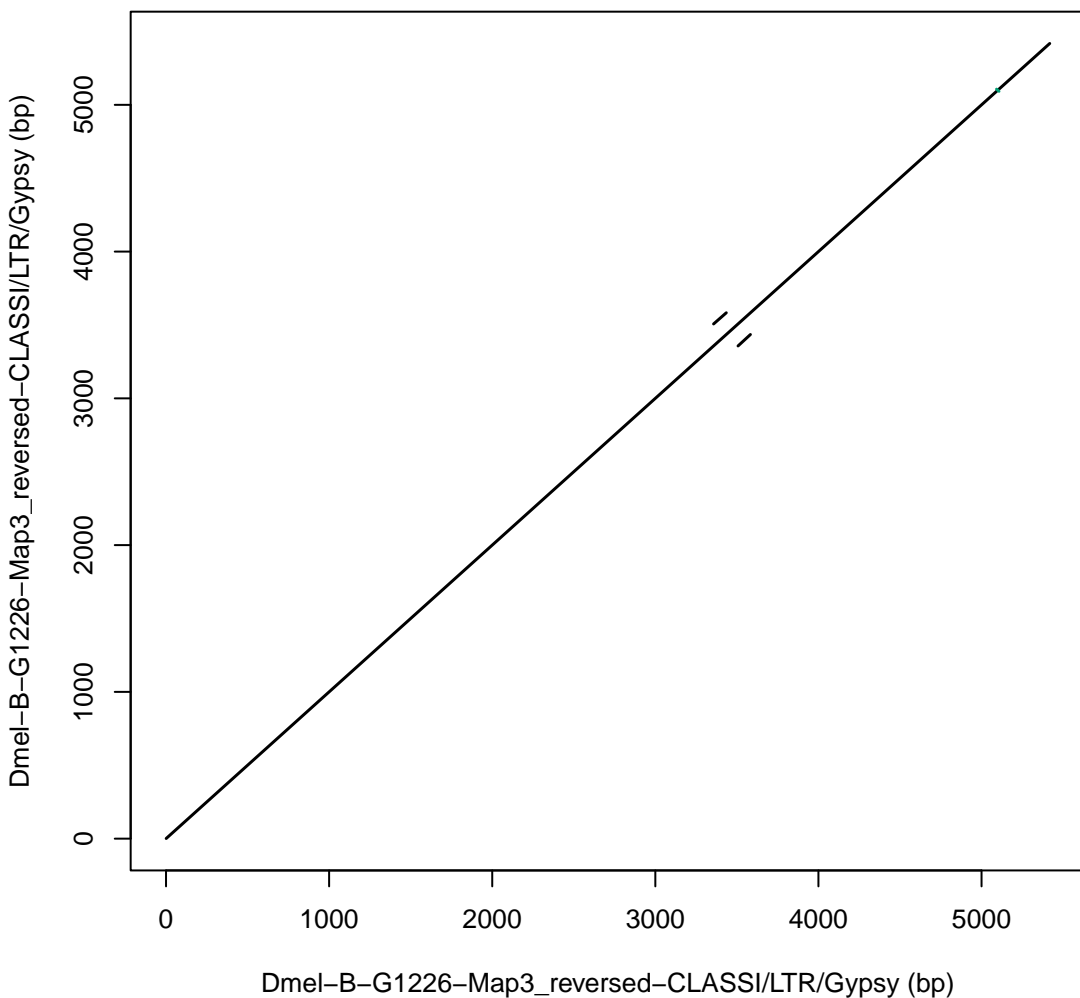
TE: Dmel-B-G1226-Map3\_reversed-CLASSI/LTR/Gypsy  
consensus size: 5418bp; fragments: 1529; full length: 2 (>=4876.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

