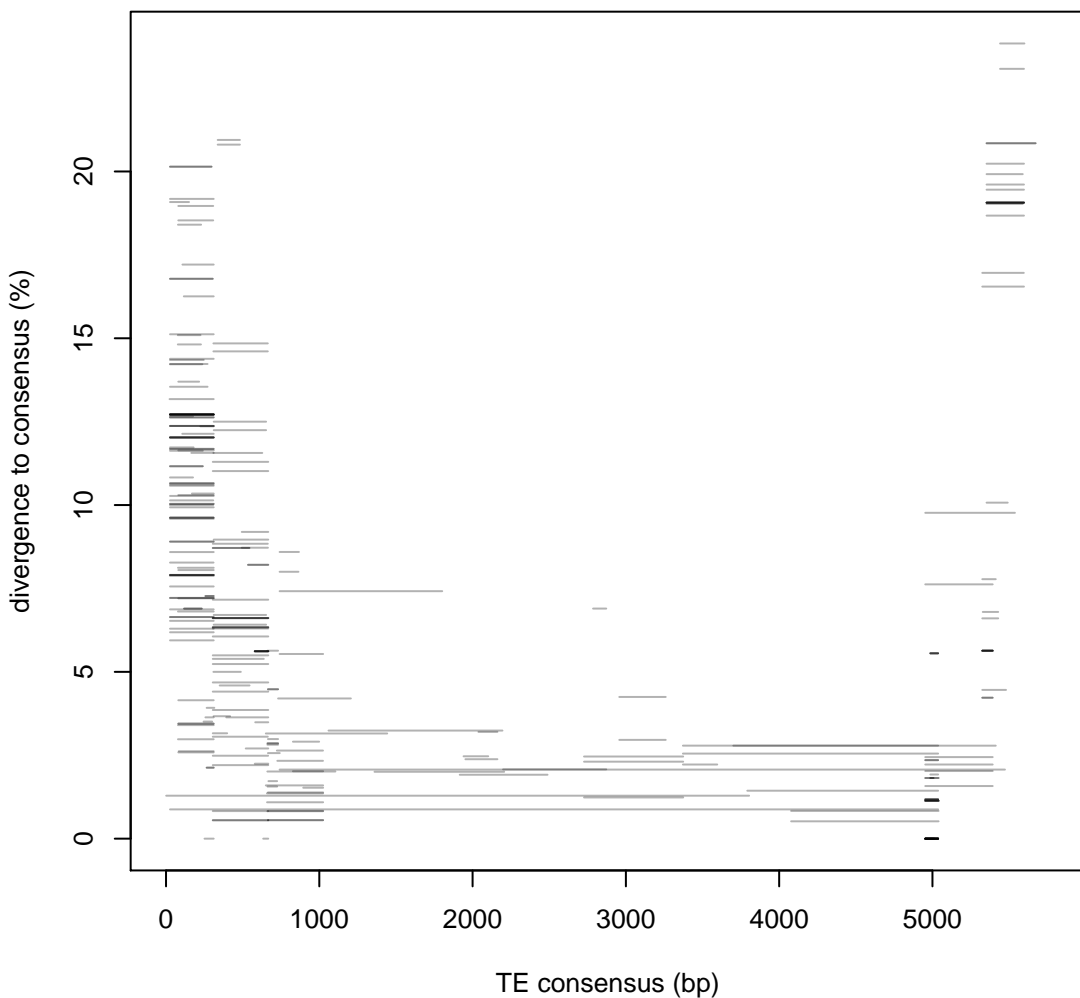
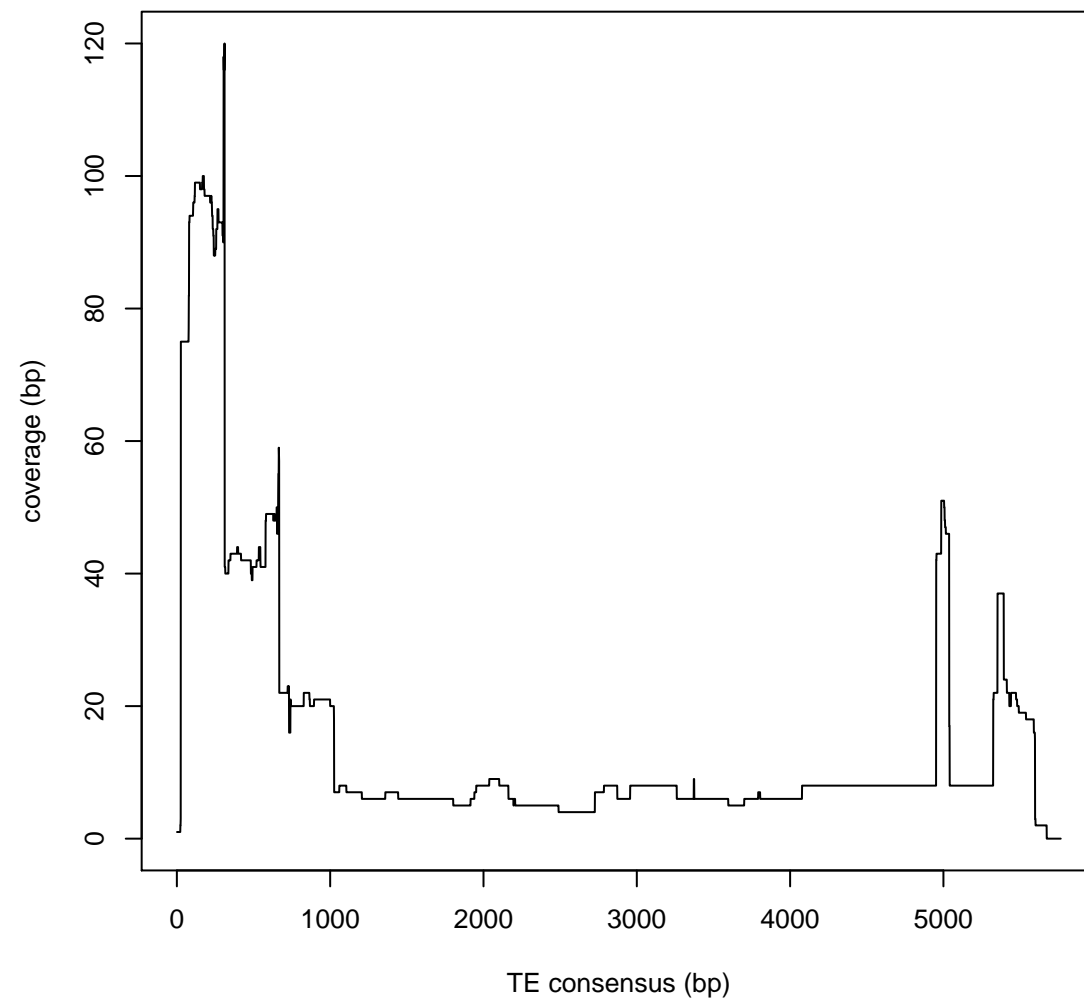


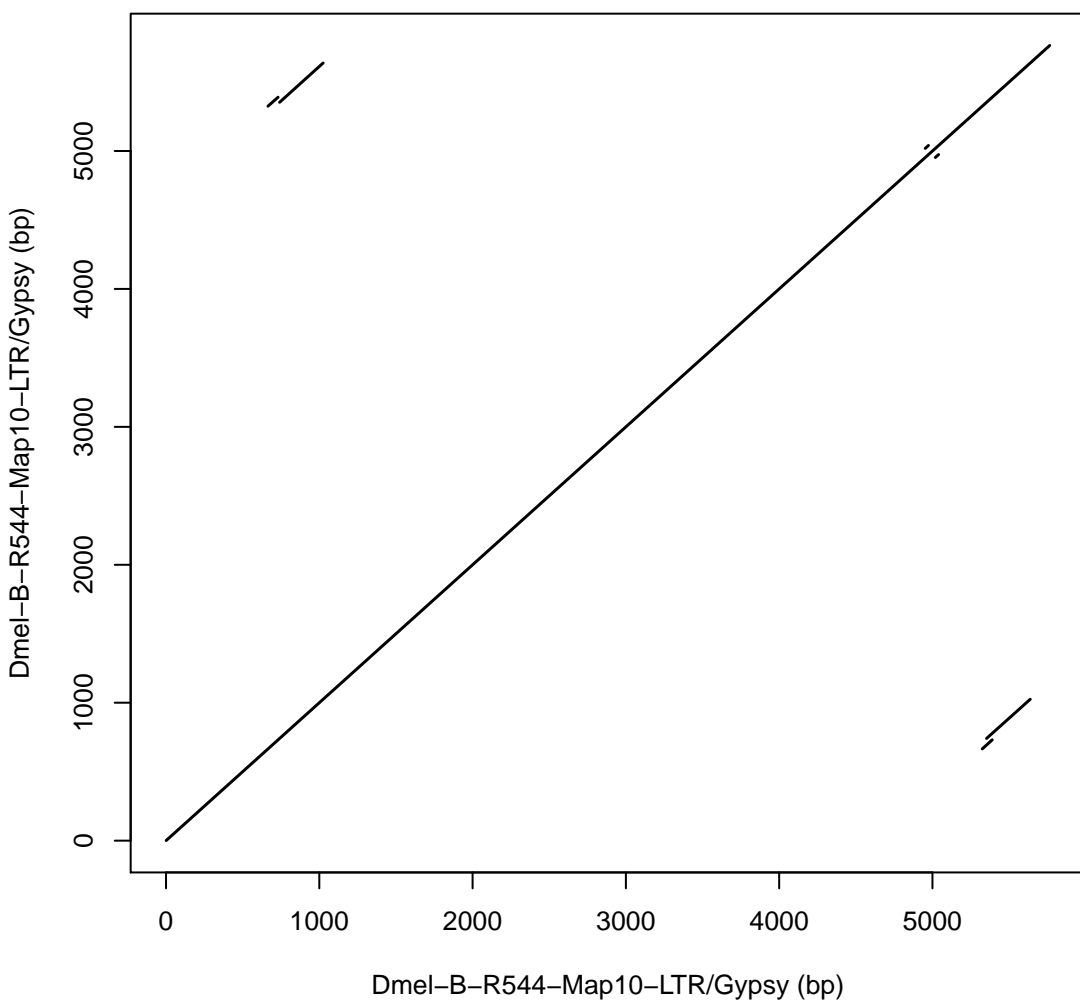
TE: Dmel-B-R544-Map10-LTR/Gypsy
consensus size: 5765bp; fragments: 291; full length: 0 (>=5188.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

