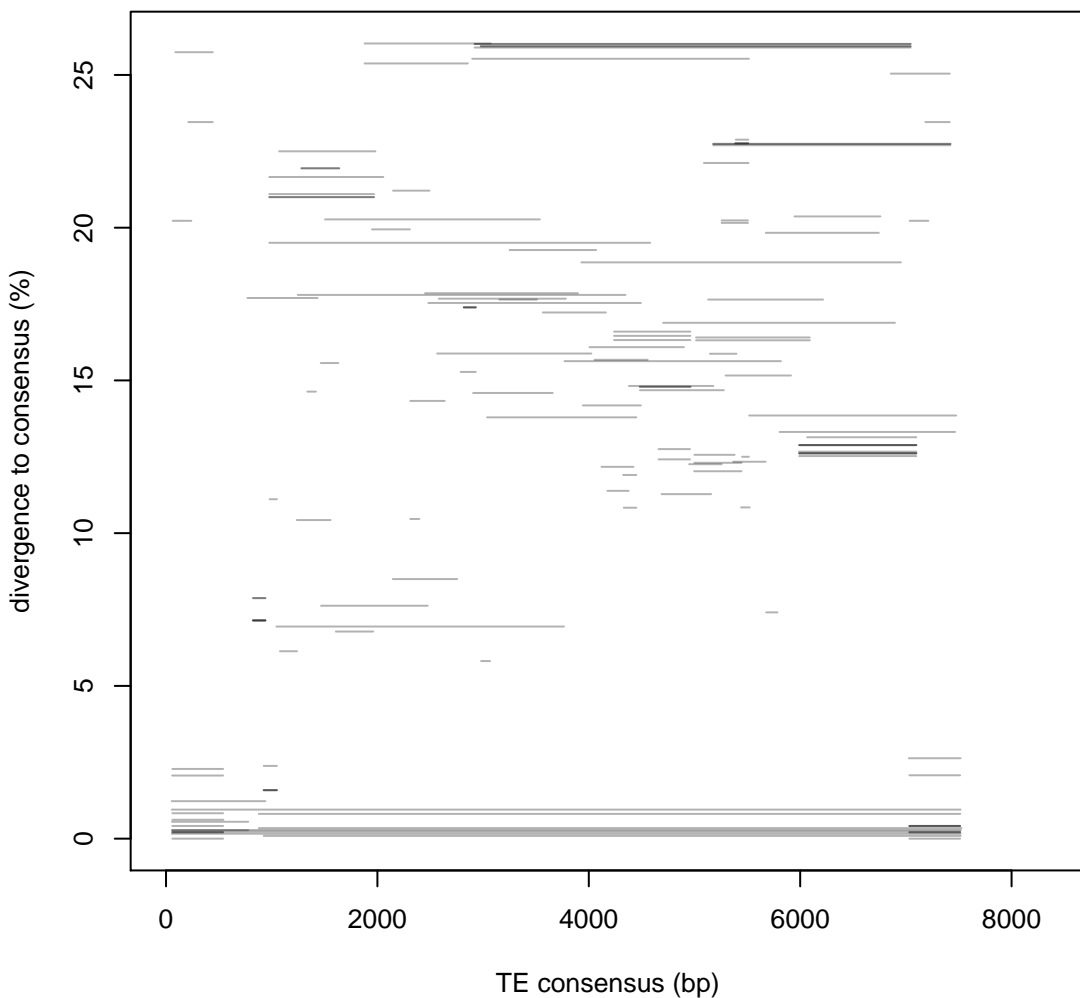
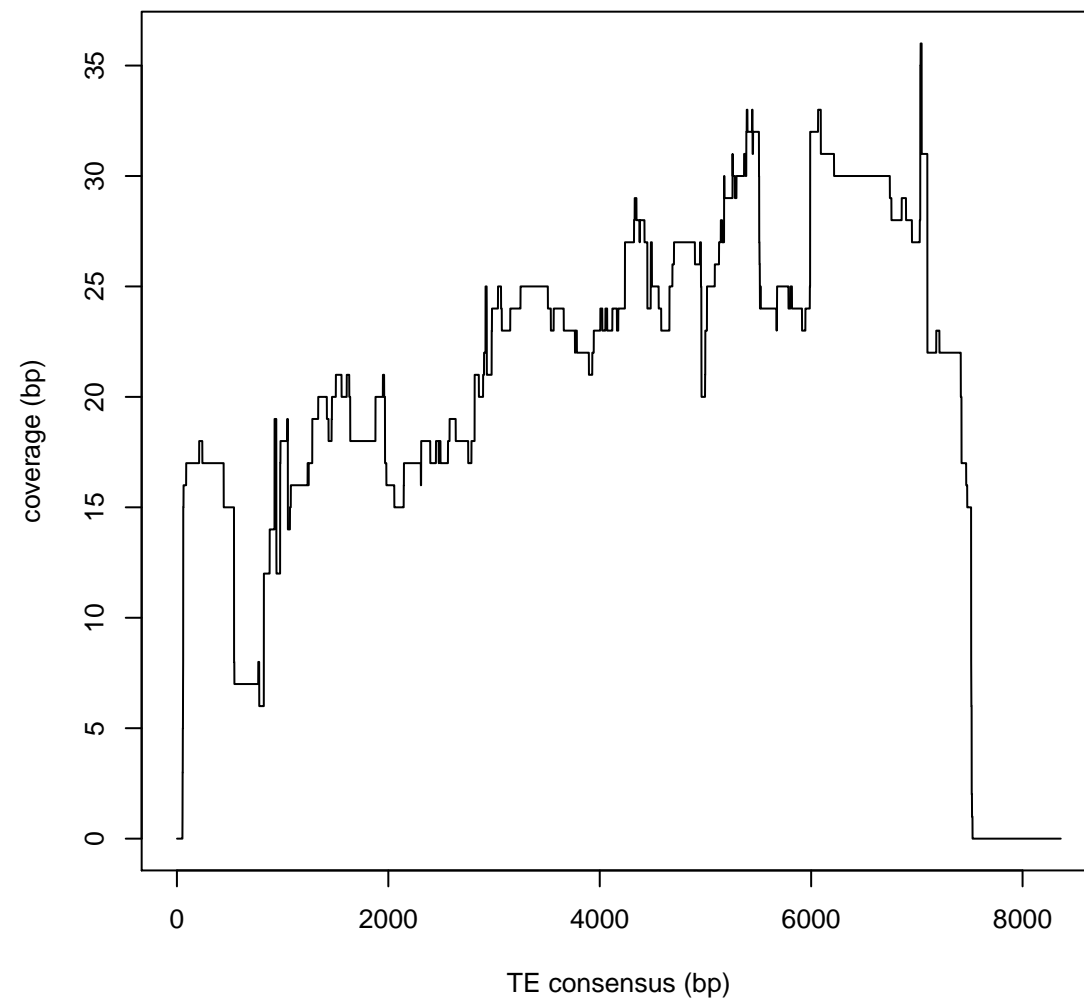


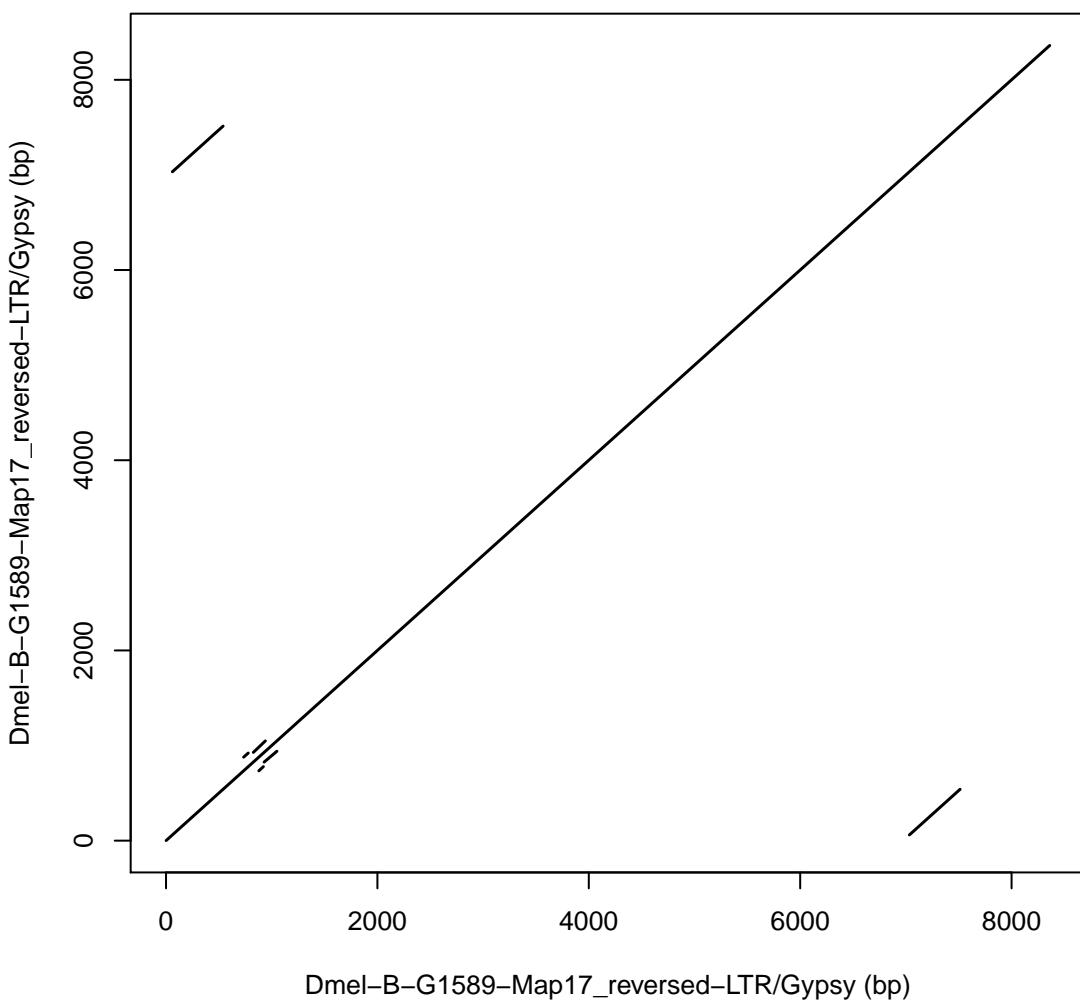
TE: Dmel-B-G1589-Map17_reversed-LTR/Gypsy
consensus size: 8361bp; fragments: 143; full length: 0 (>=7524.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

