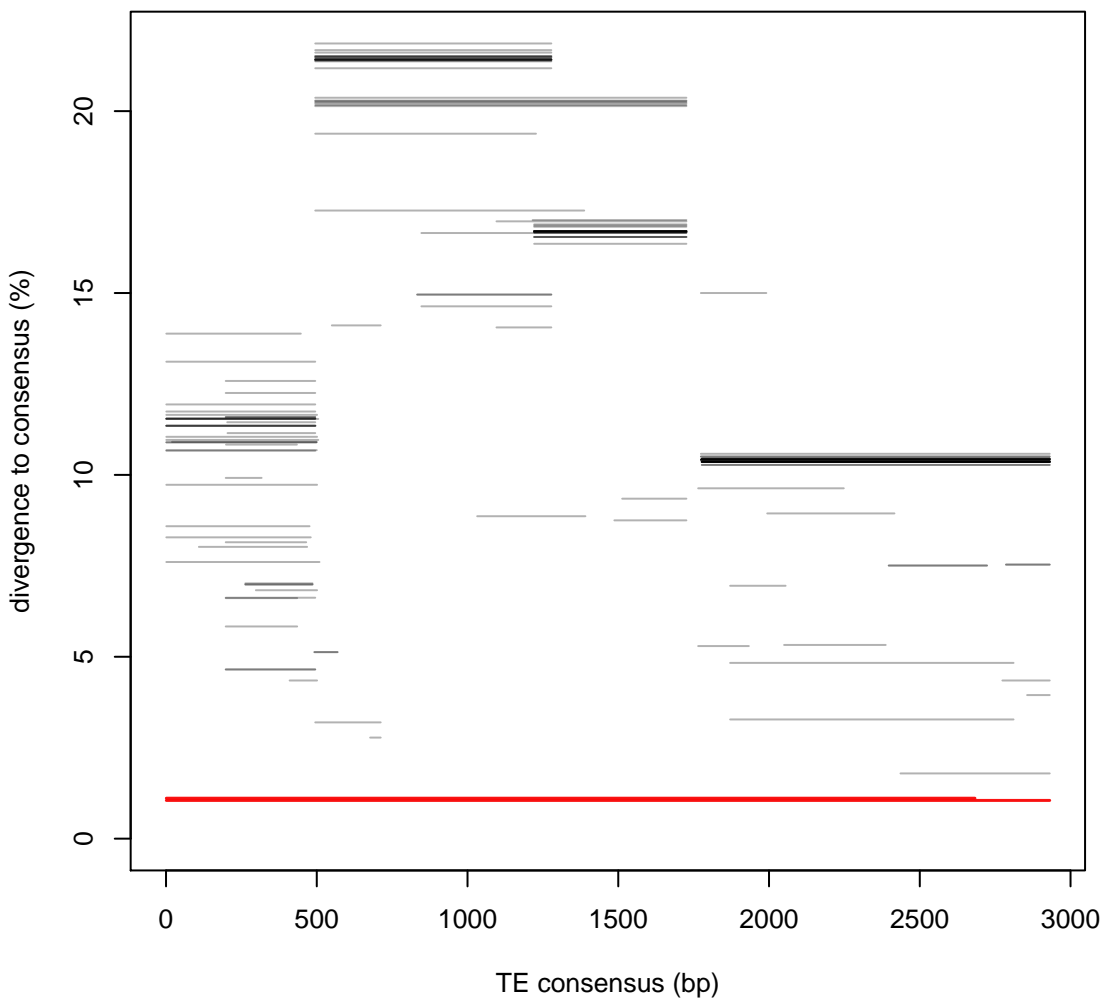
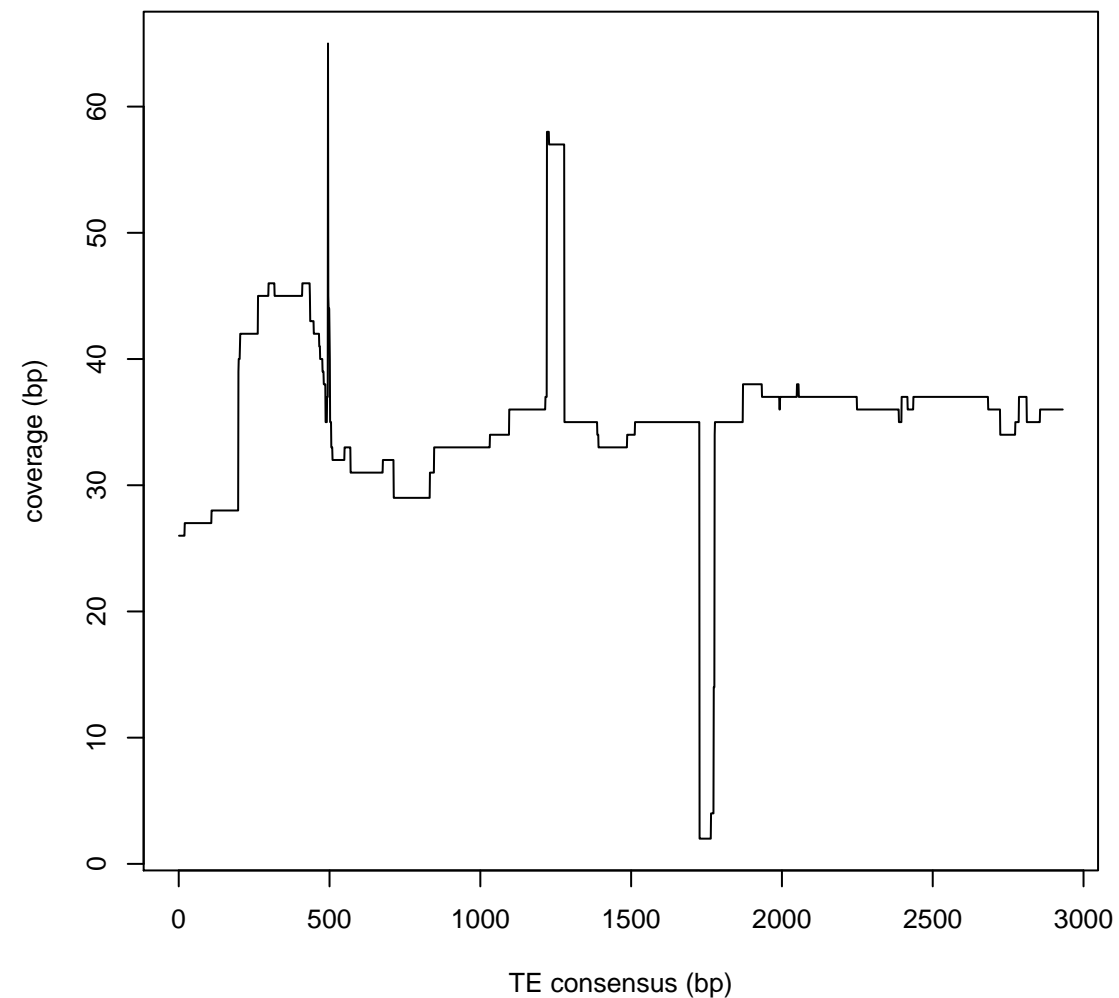


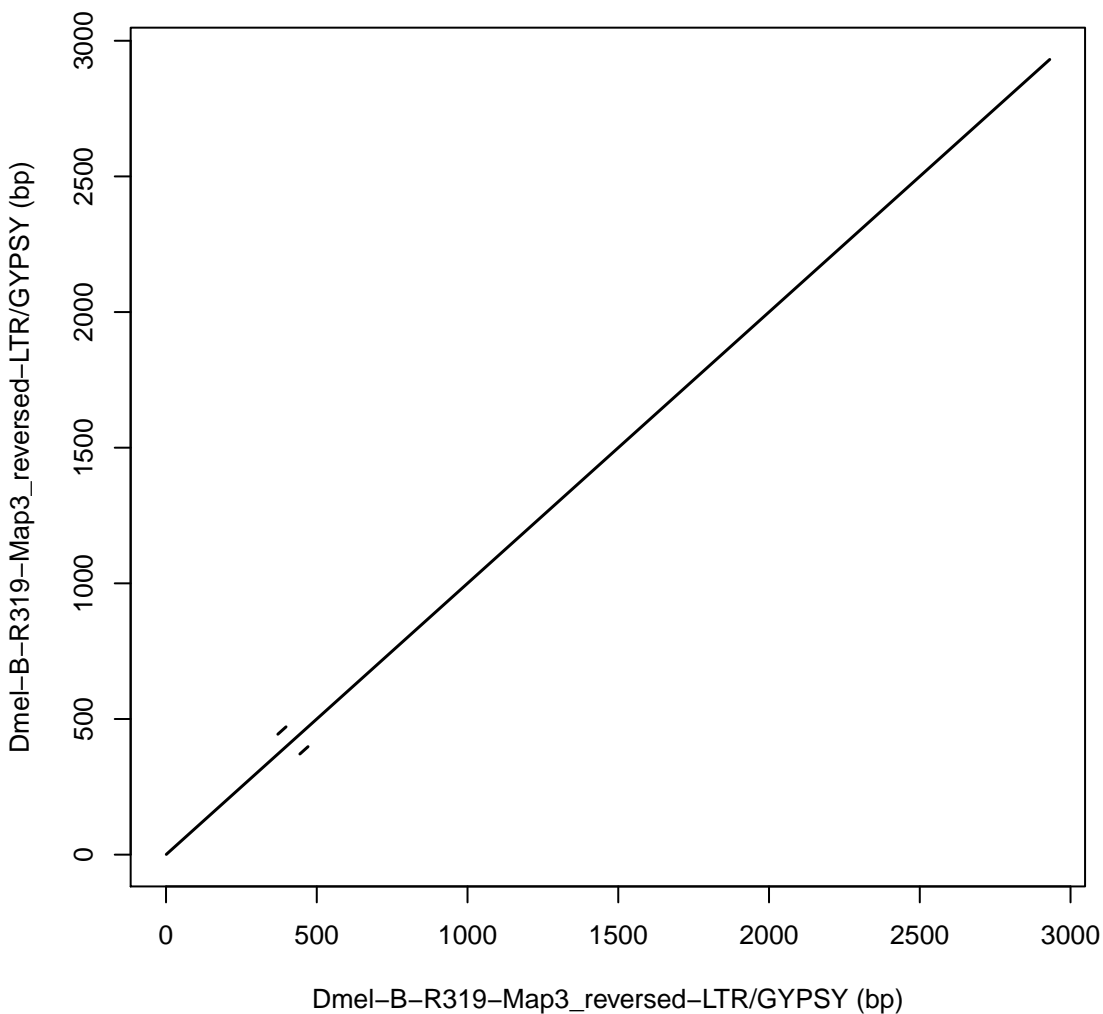
TE: Dmel-B-R319-Map3\_reversed-LTR/Gypsy  
consensus size: 2931bp; fragments: 155; full length: 2 (>=2637.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

