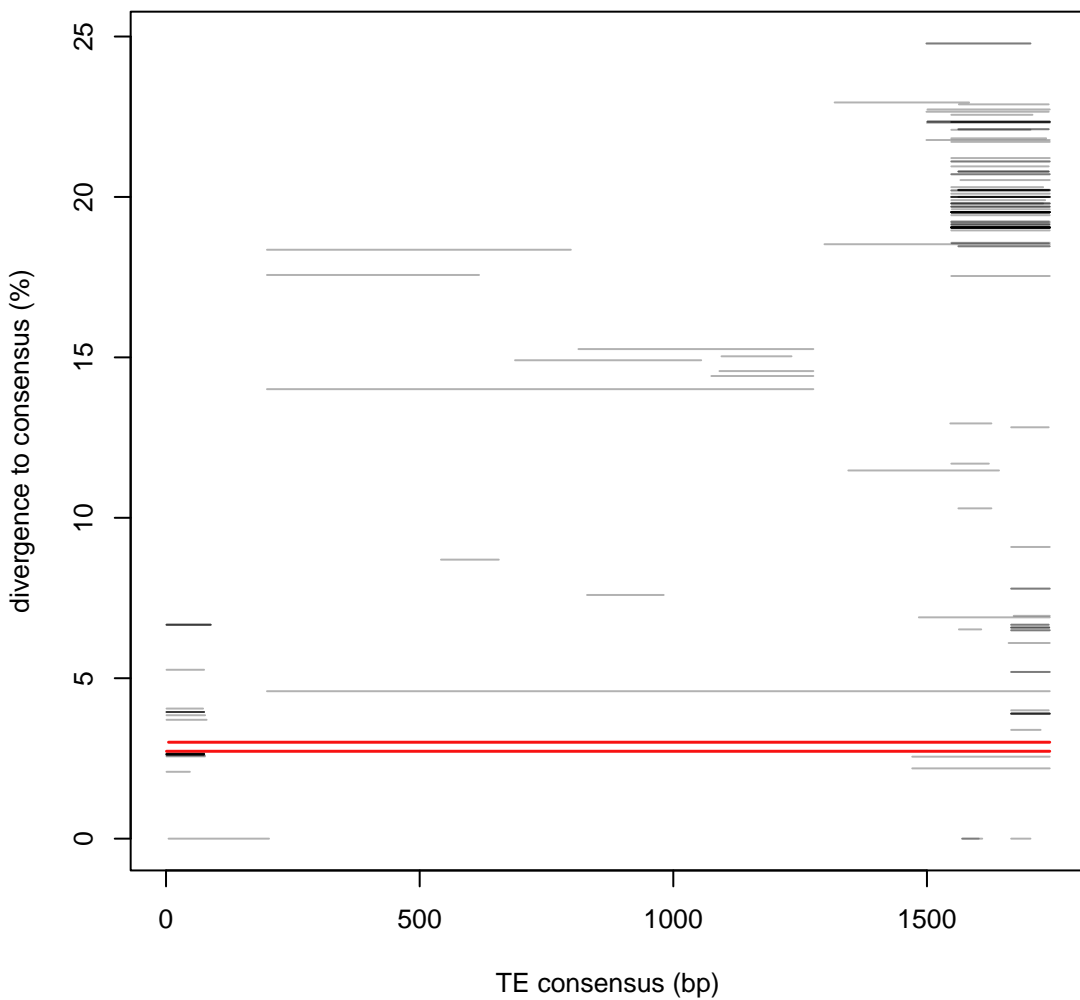
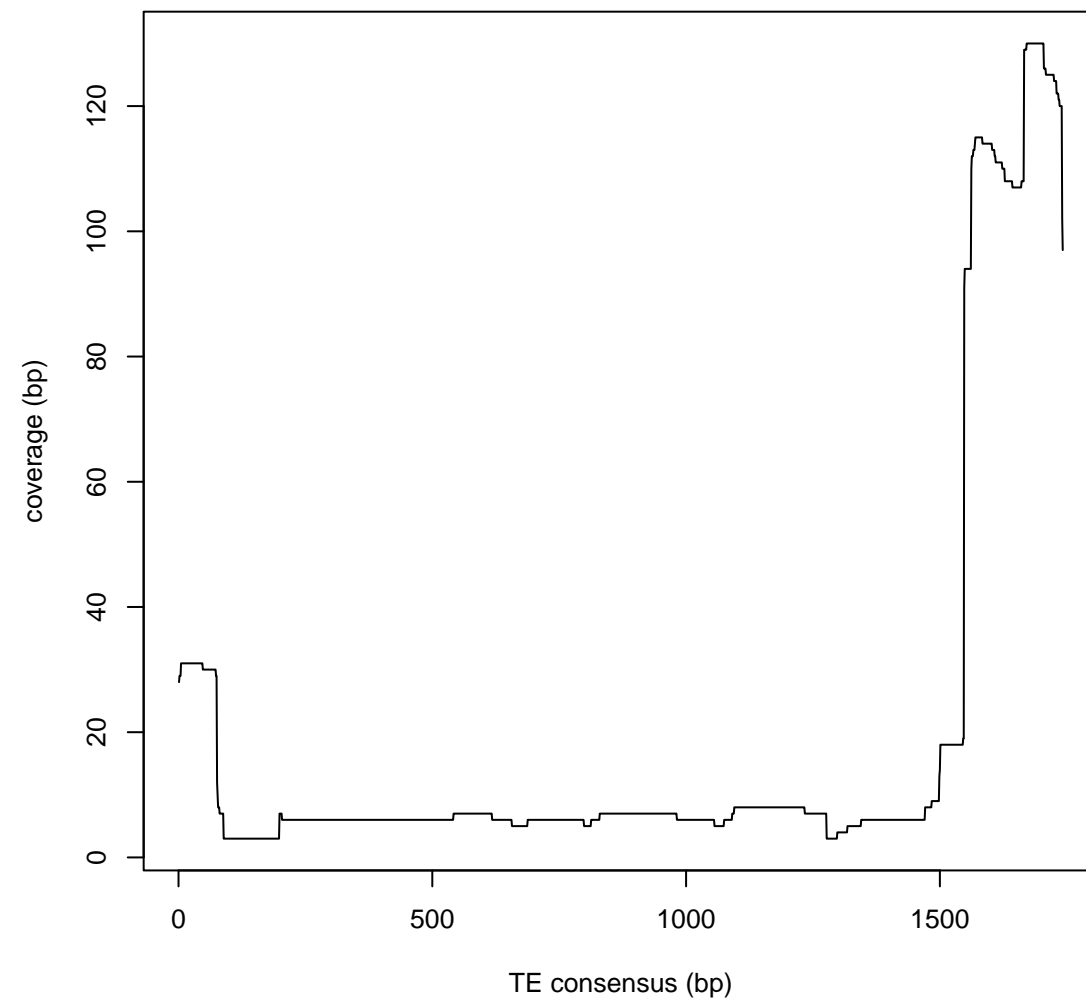


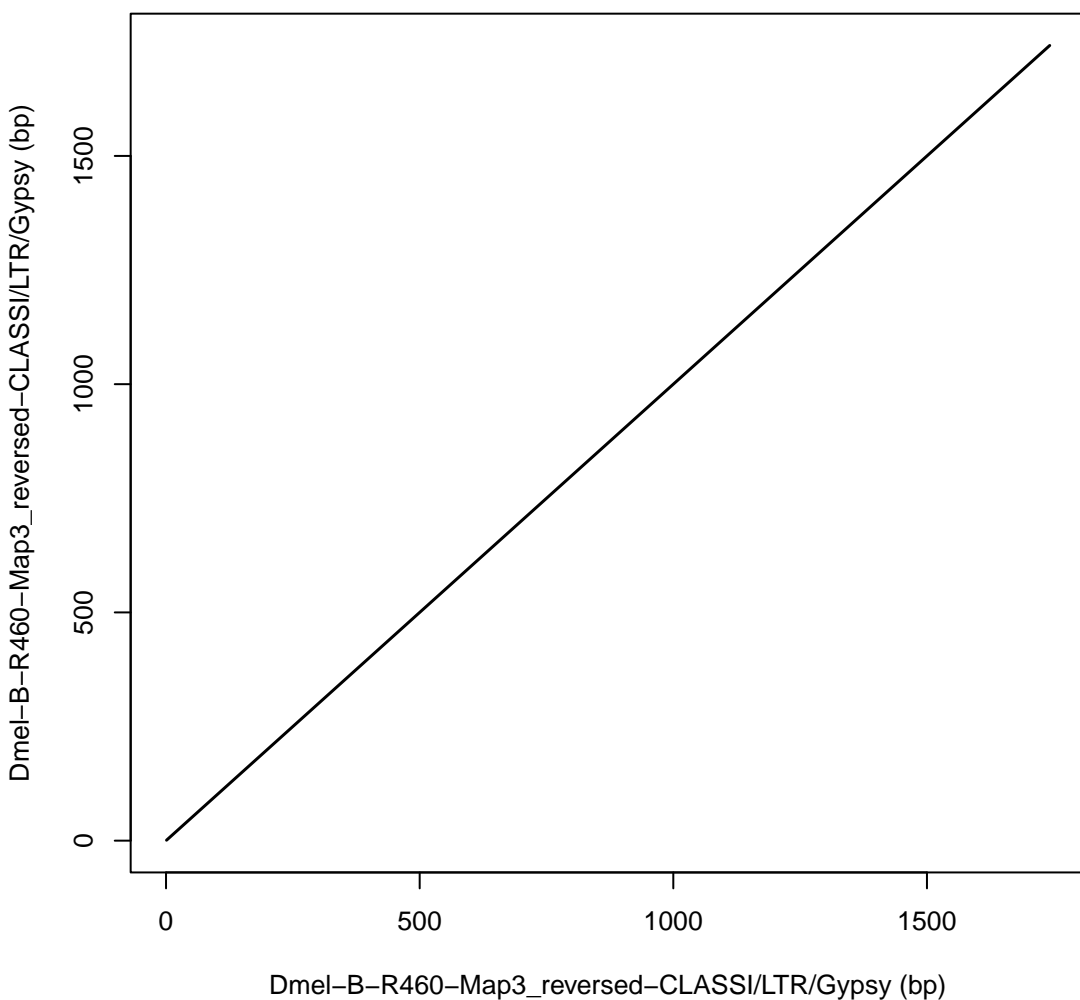
TE: Dmel-B-R460-Map3\_reversed-CLASSI/LTR/Gypsy  
consensus size: 1742bp; fragments: 177; full length: 2 ( $\geq 1567.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

