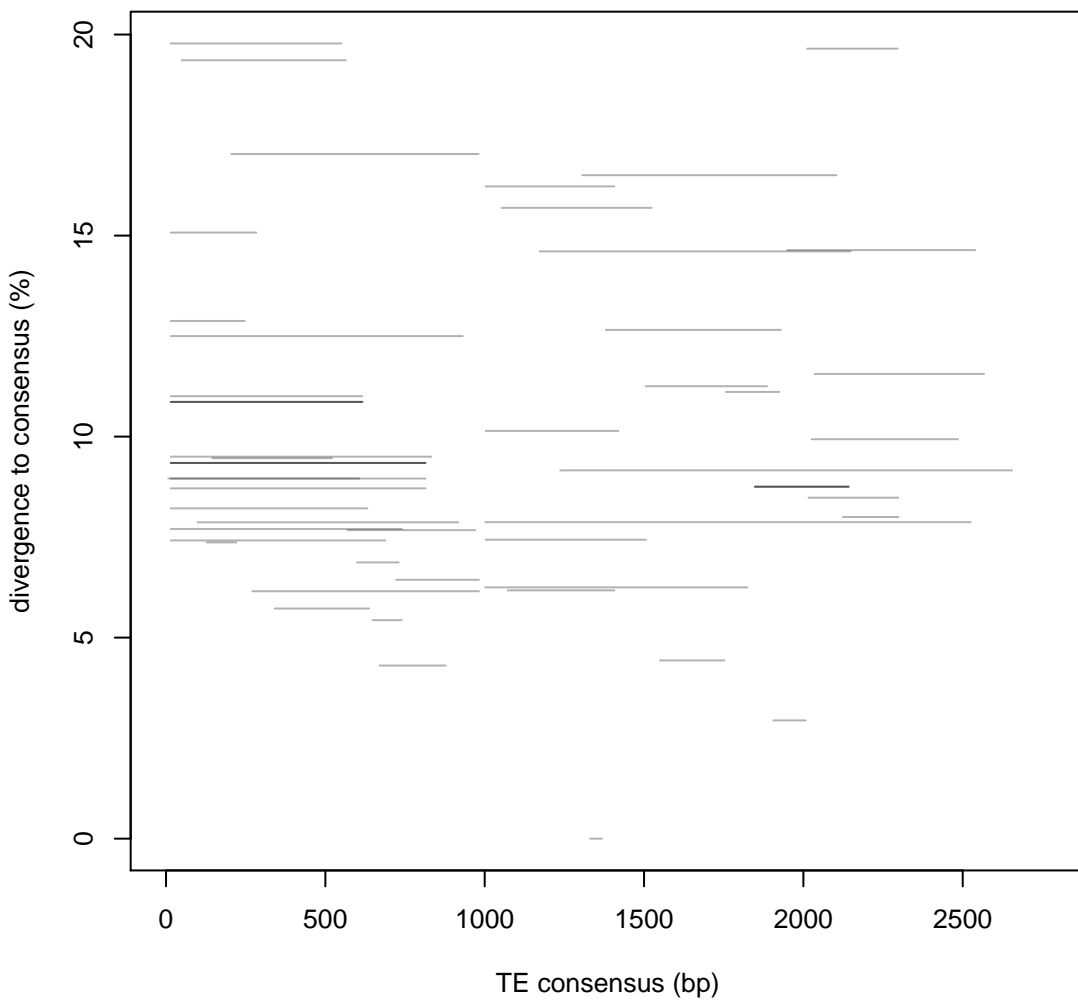
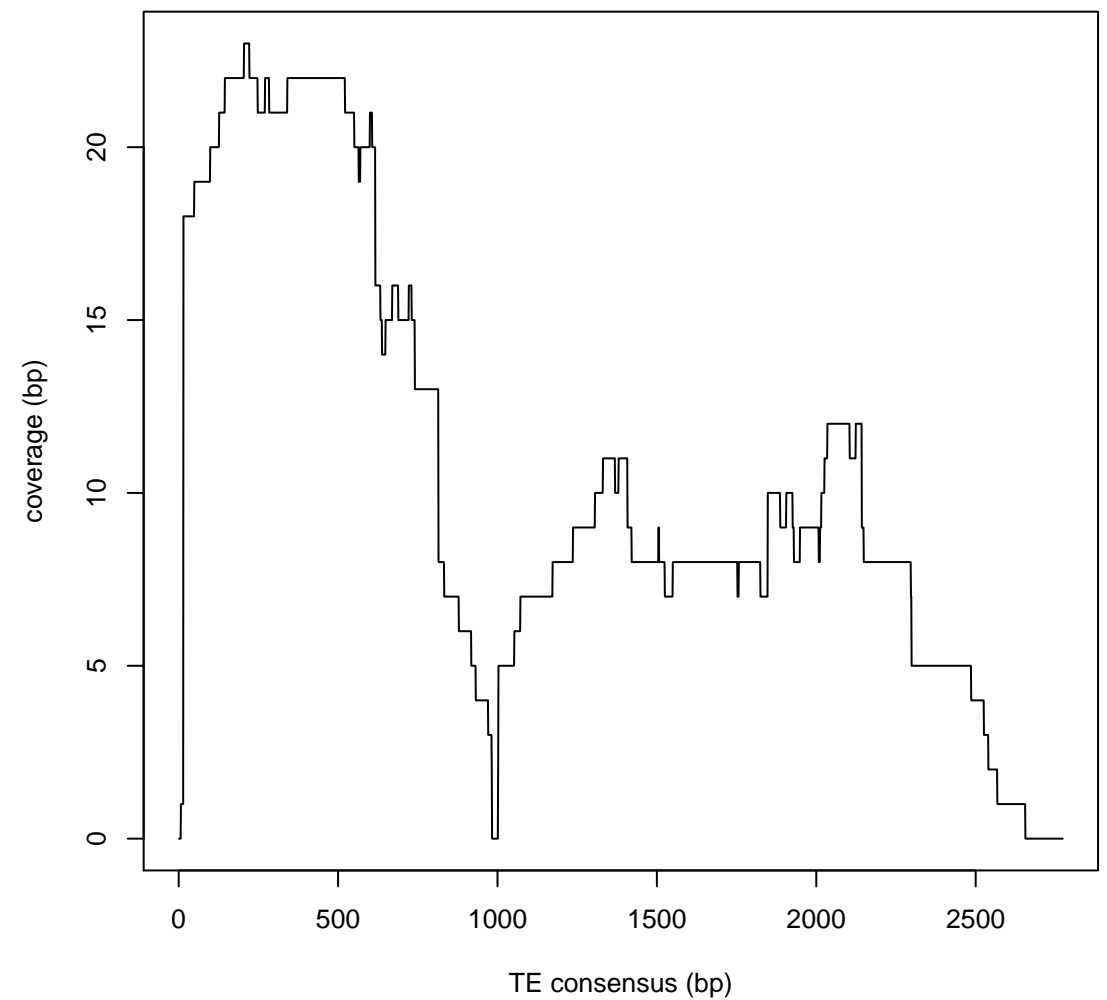


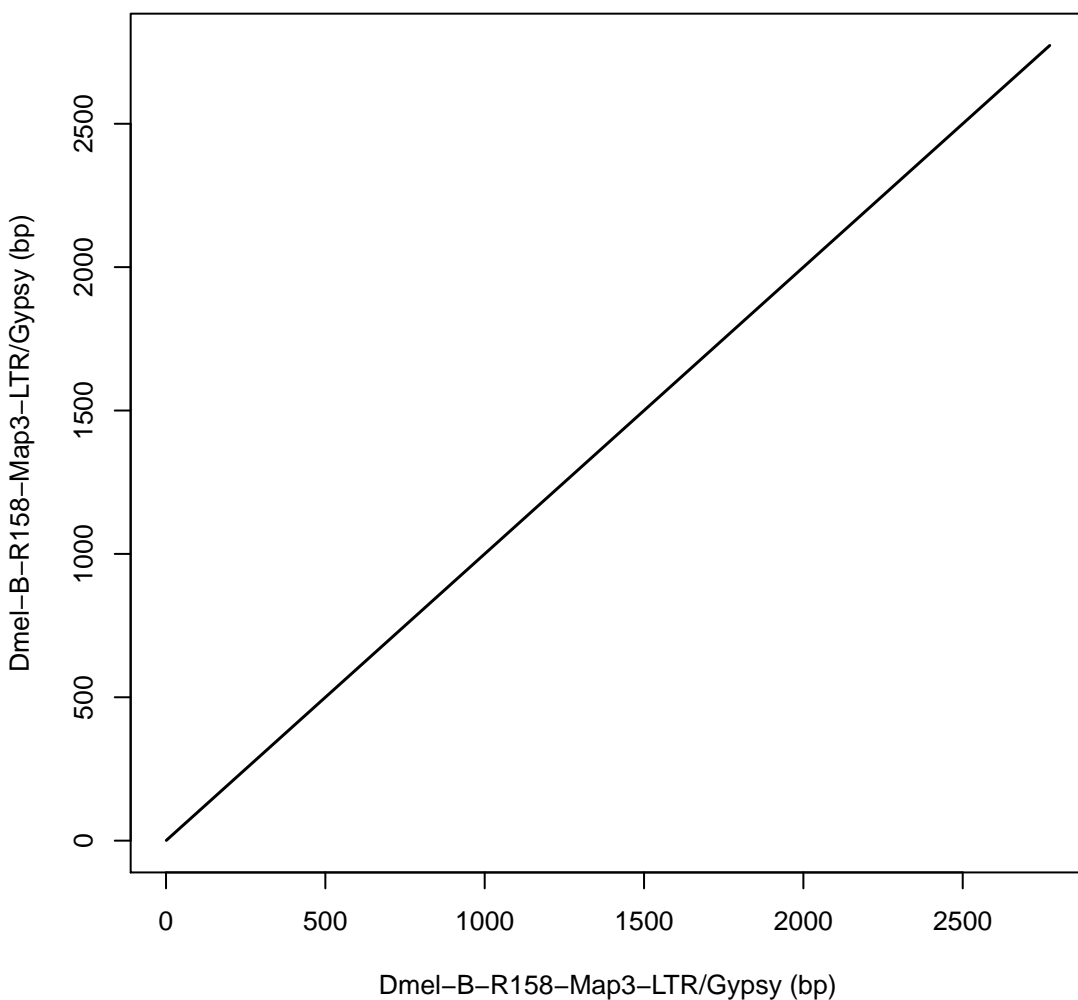
TE: Dmel-B-R158-Map3-LTR/Gypsy
consensus size: 2773bp; fragments: 55; full length: 0 (>=2495.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

