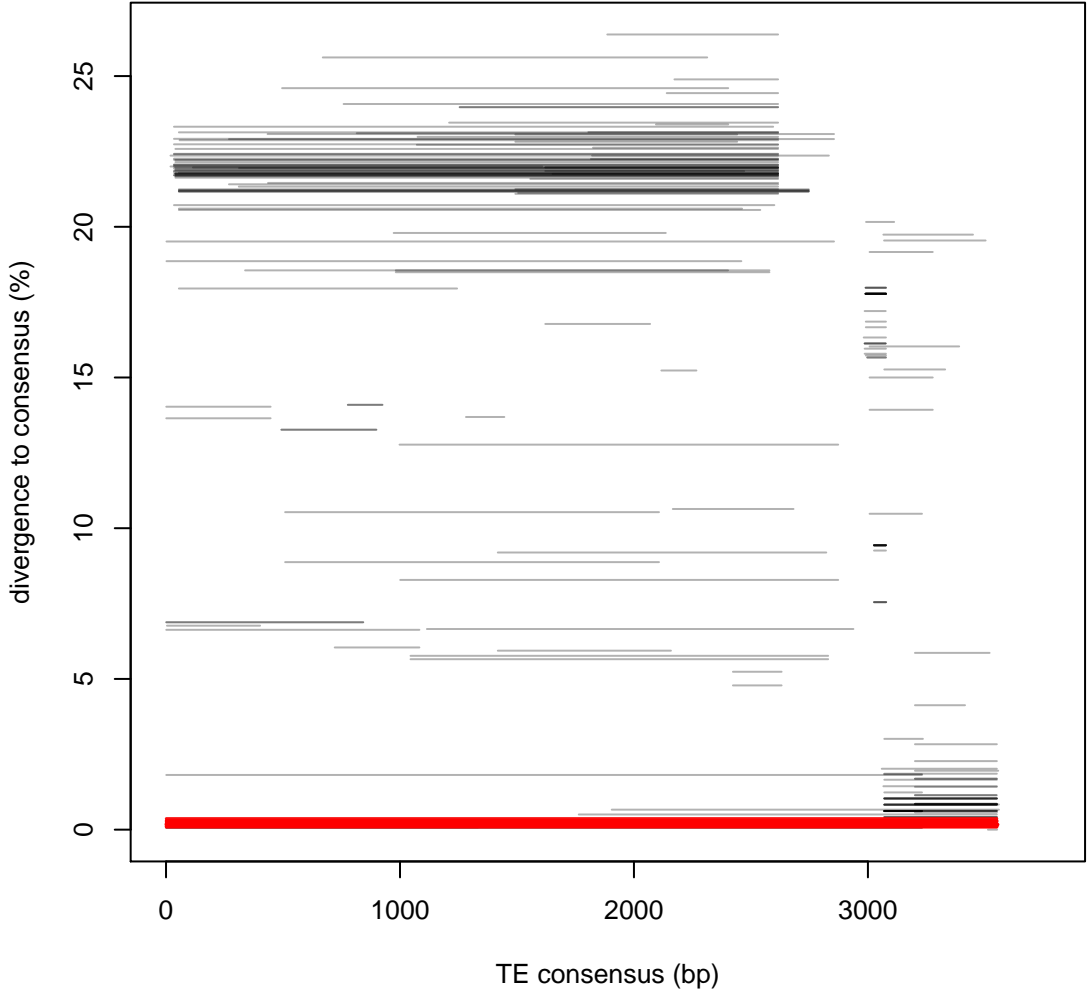
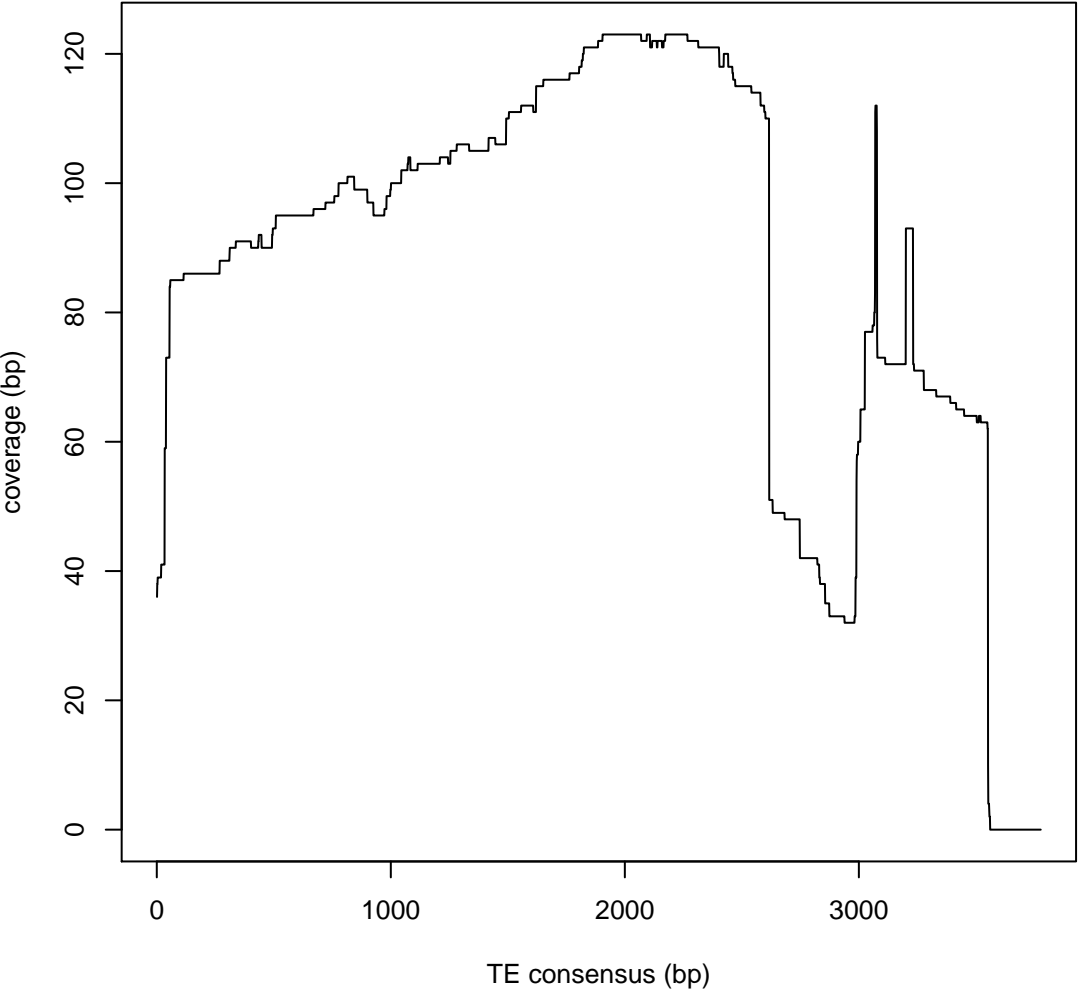


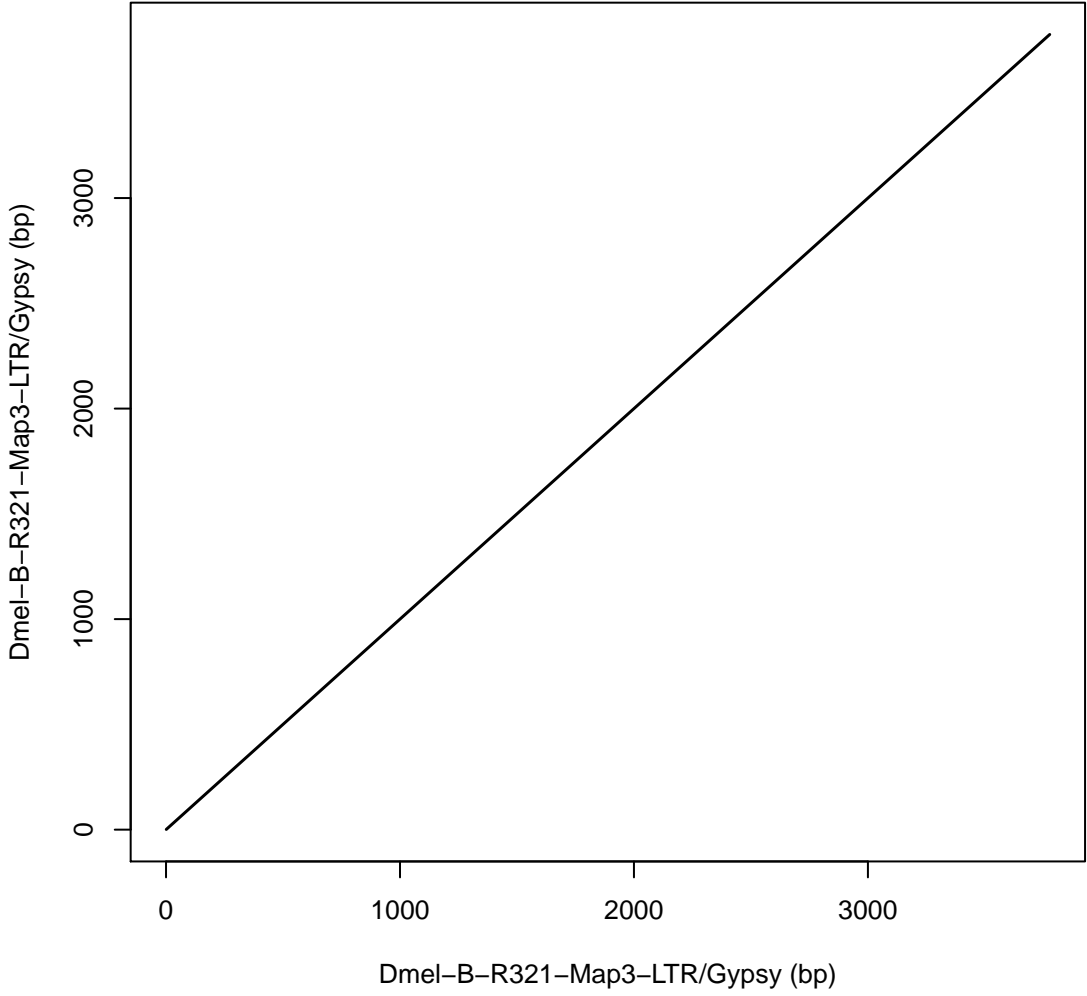
TE: Dmel-B-R321-Map3-LTR/Gypsy  
consensus size: 3777bp; fragments: 247; full length: 19 (>=3399.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

