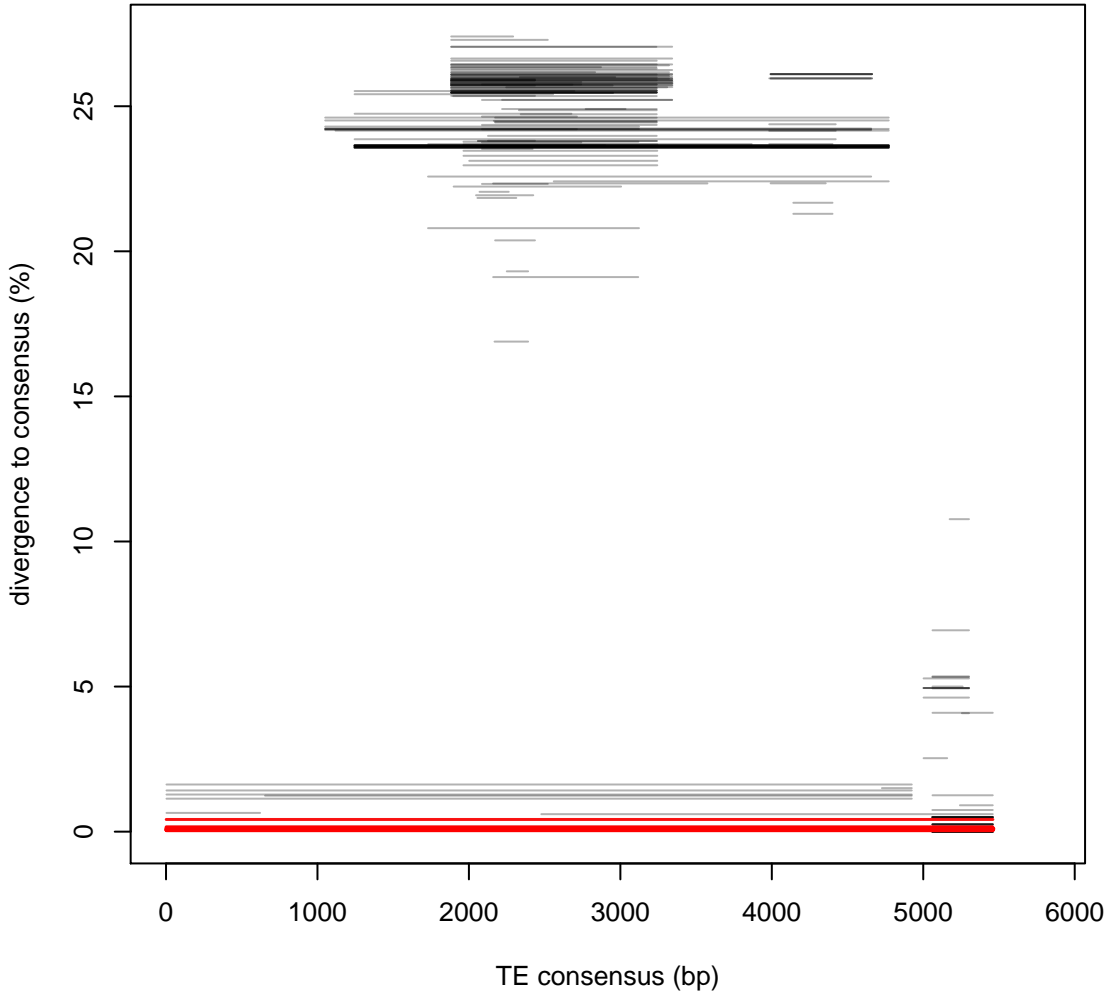
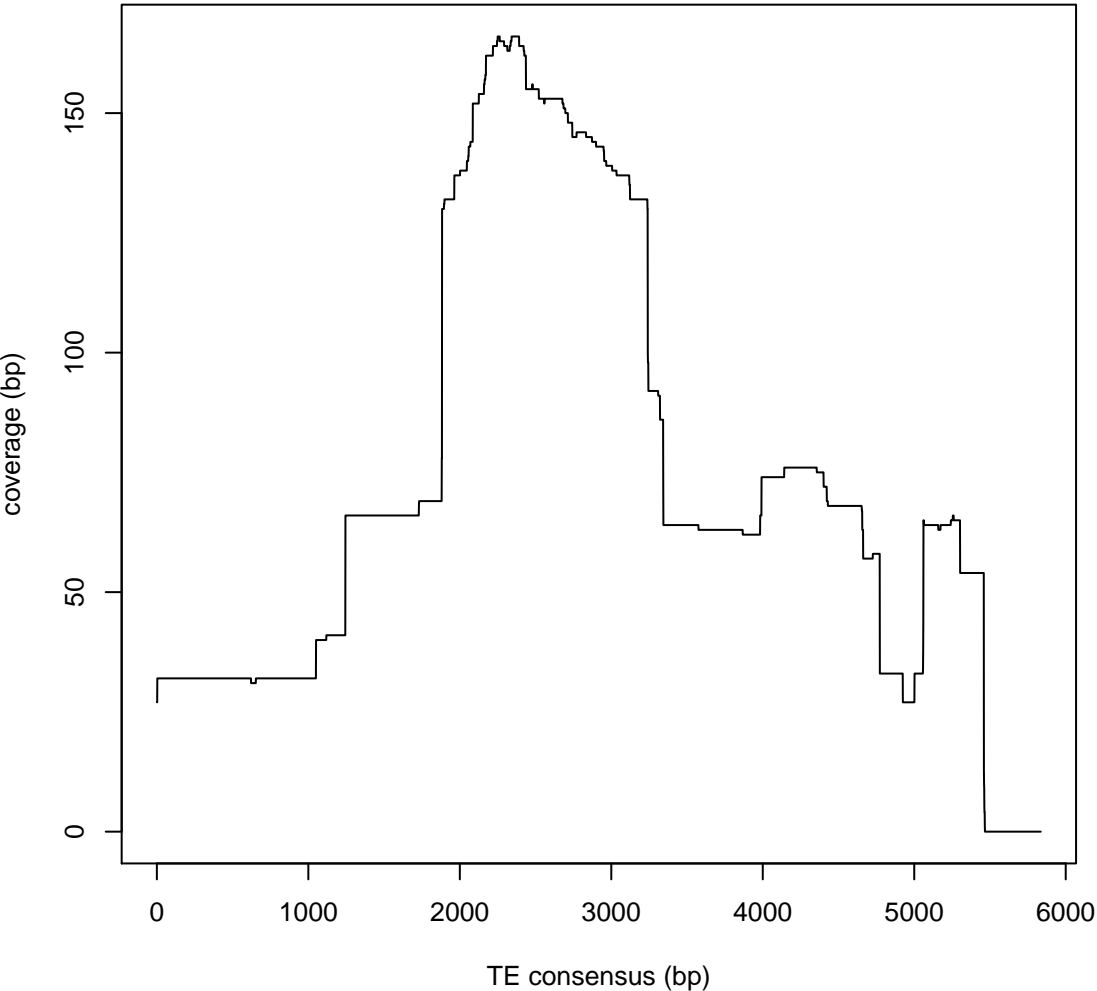


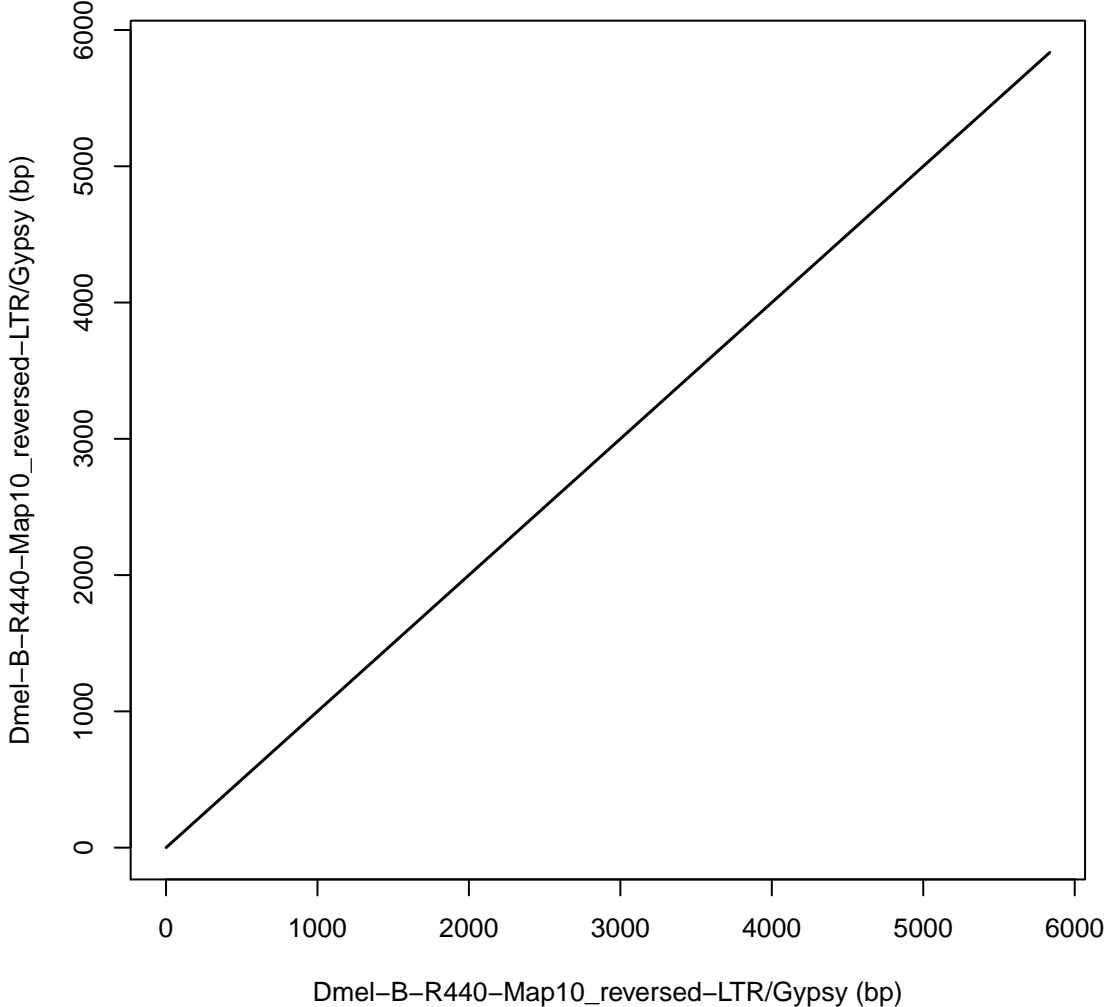
TE: Dmel-B-R440-Map10_reversed-LTR/Gypsy
consensus size: 5835bp; fragments: 229; full length: 25 (>=5251.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

