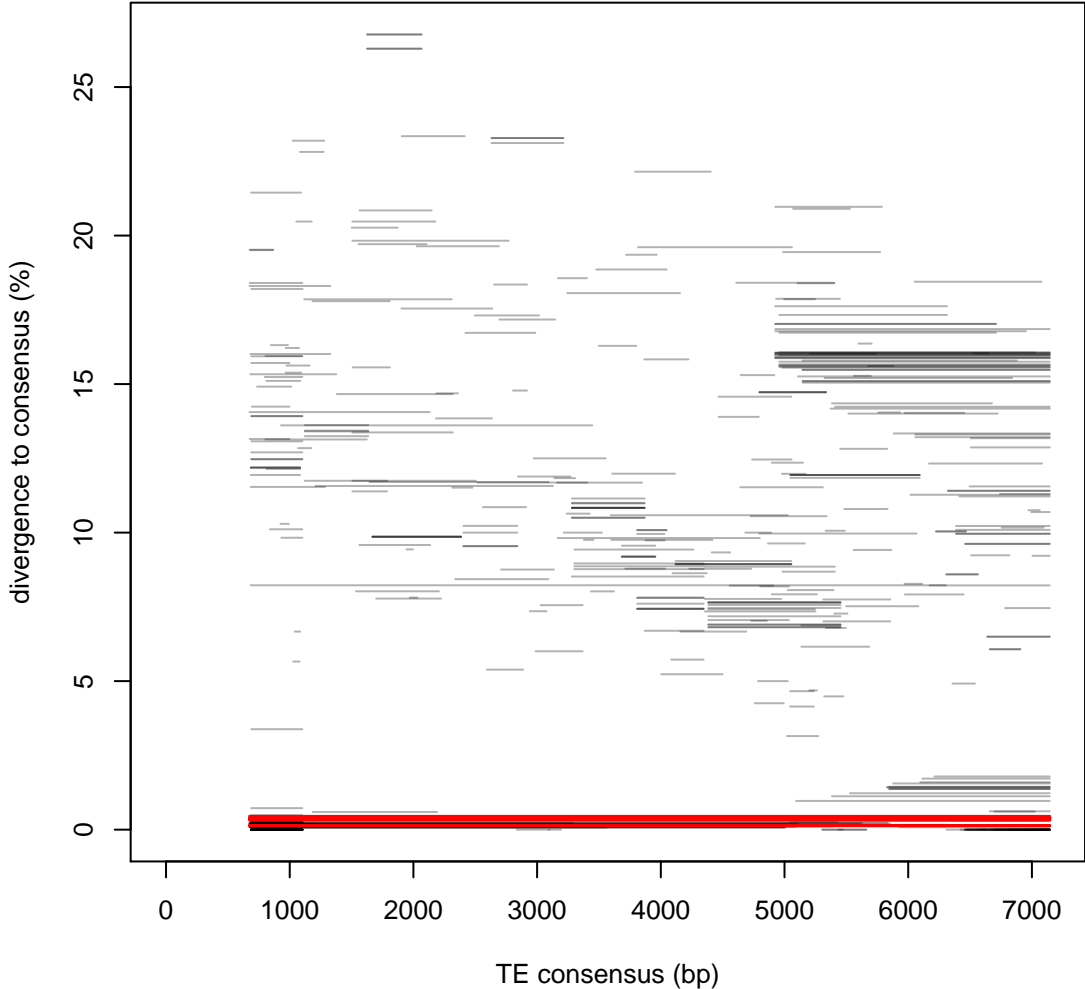
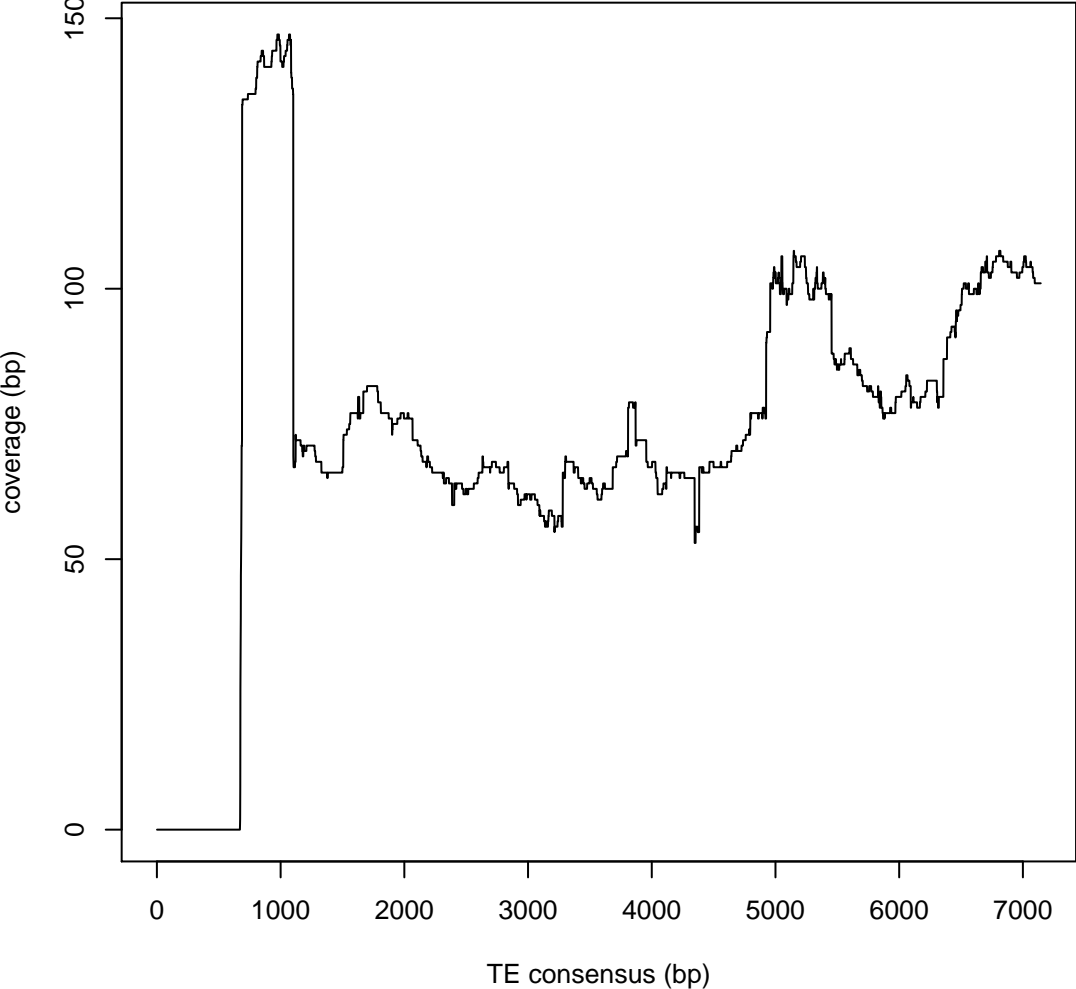


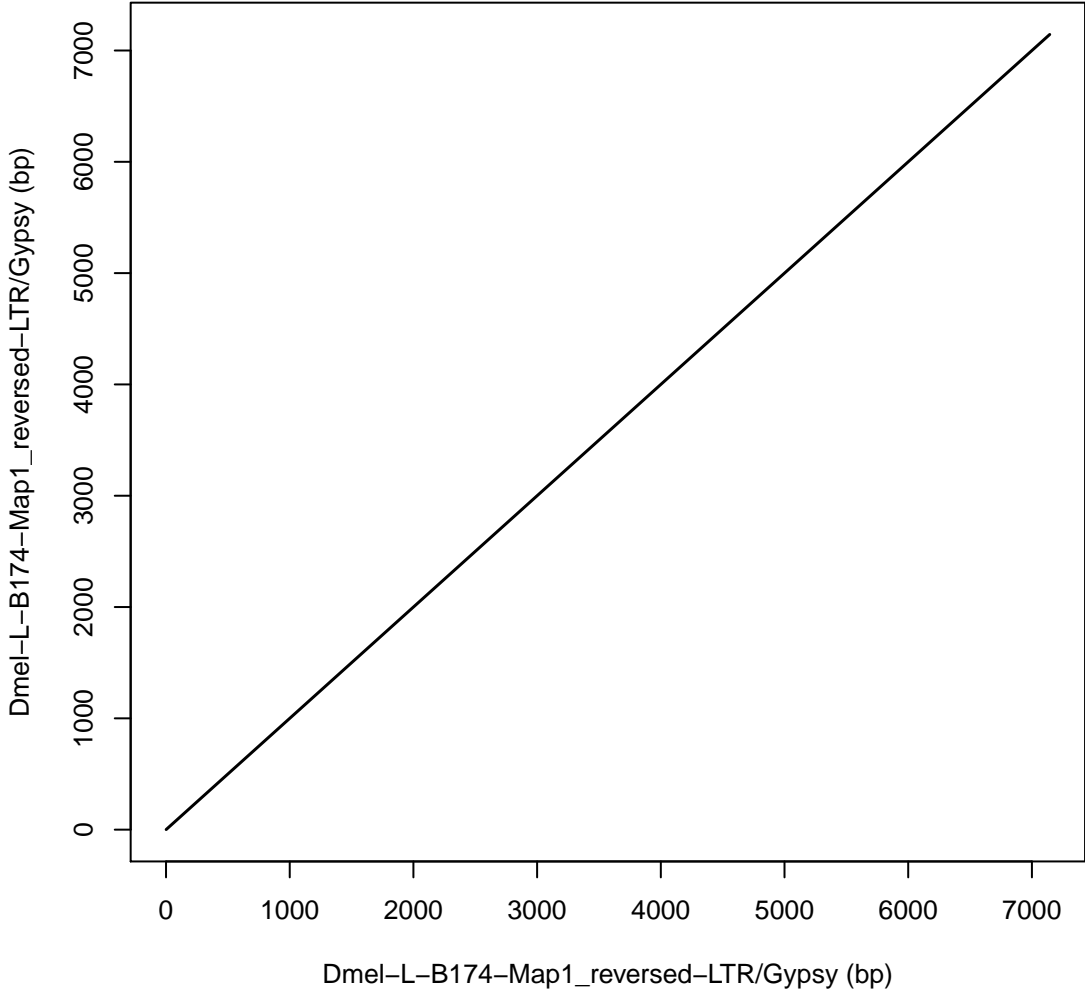
TE: Dmel-L-B174-Map1_reversed-LTR/Gypsy
consensus size: 7144bp; fragments: 467; full length: 16 (>=6429.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

