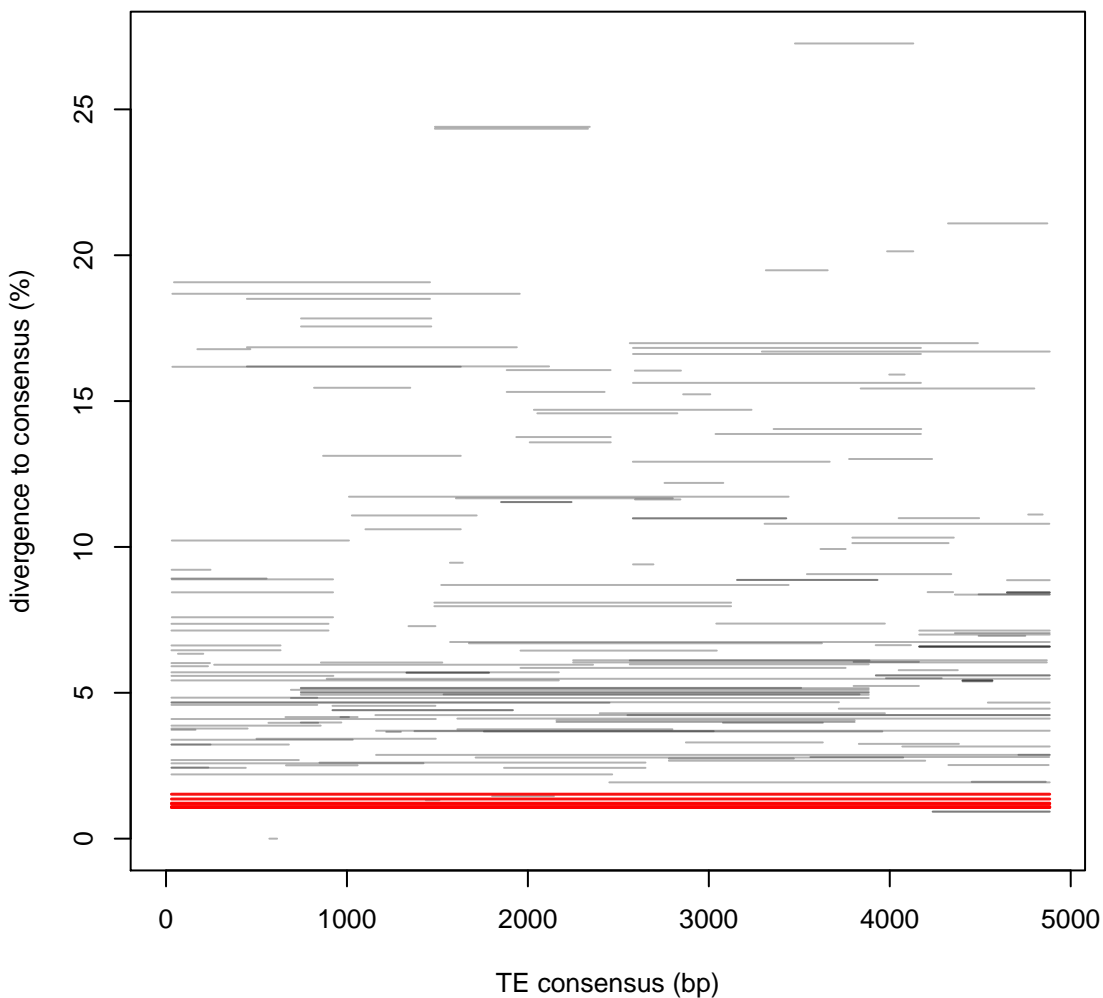
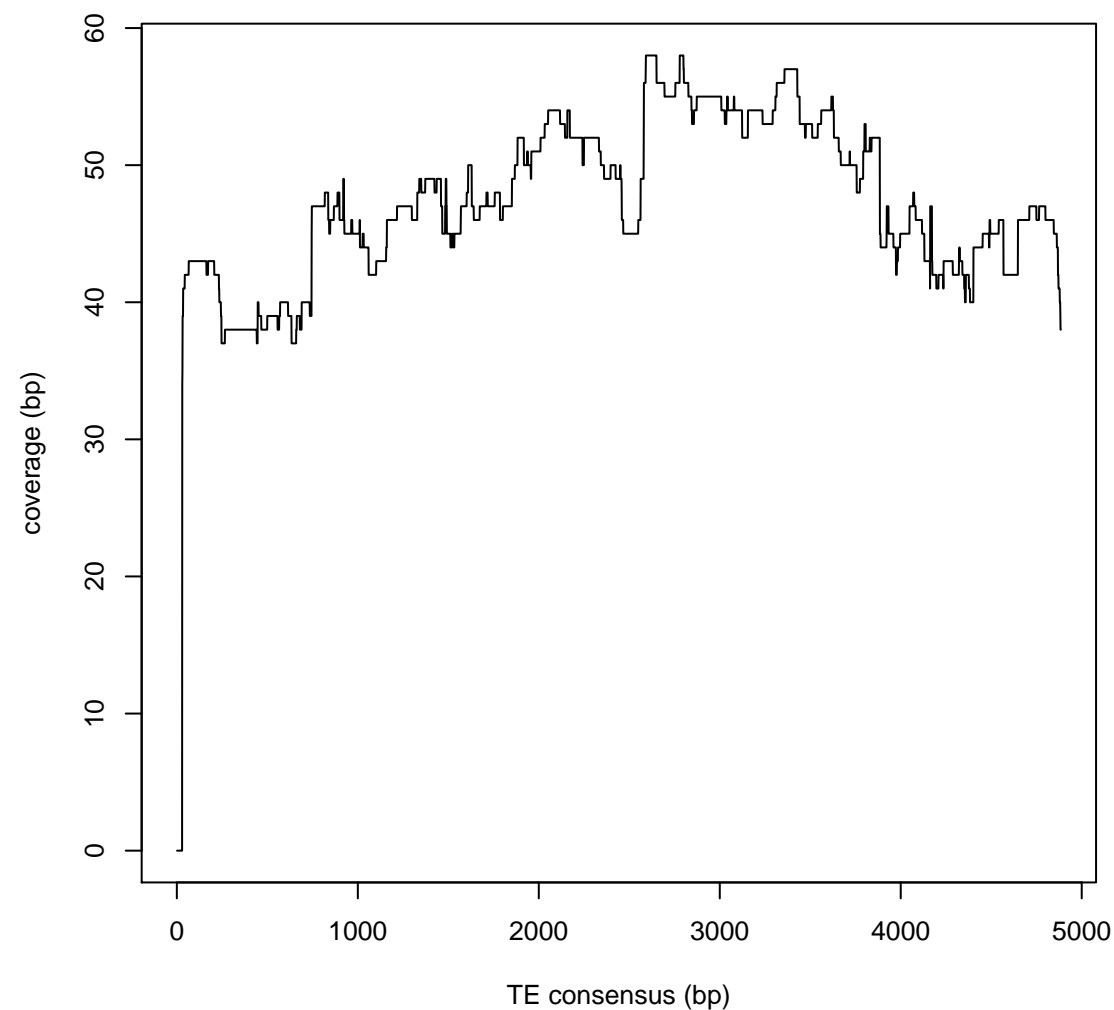


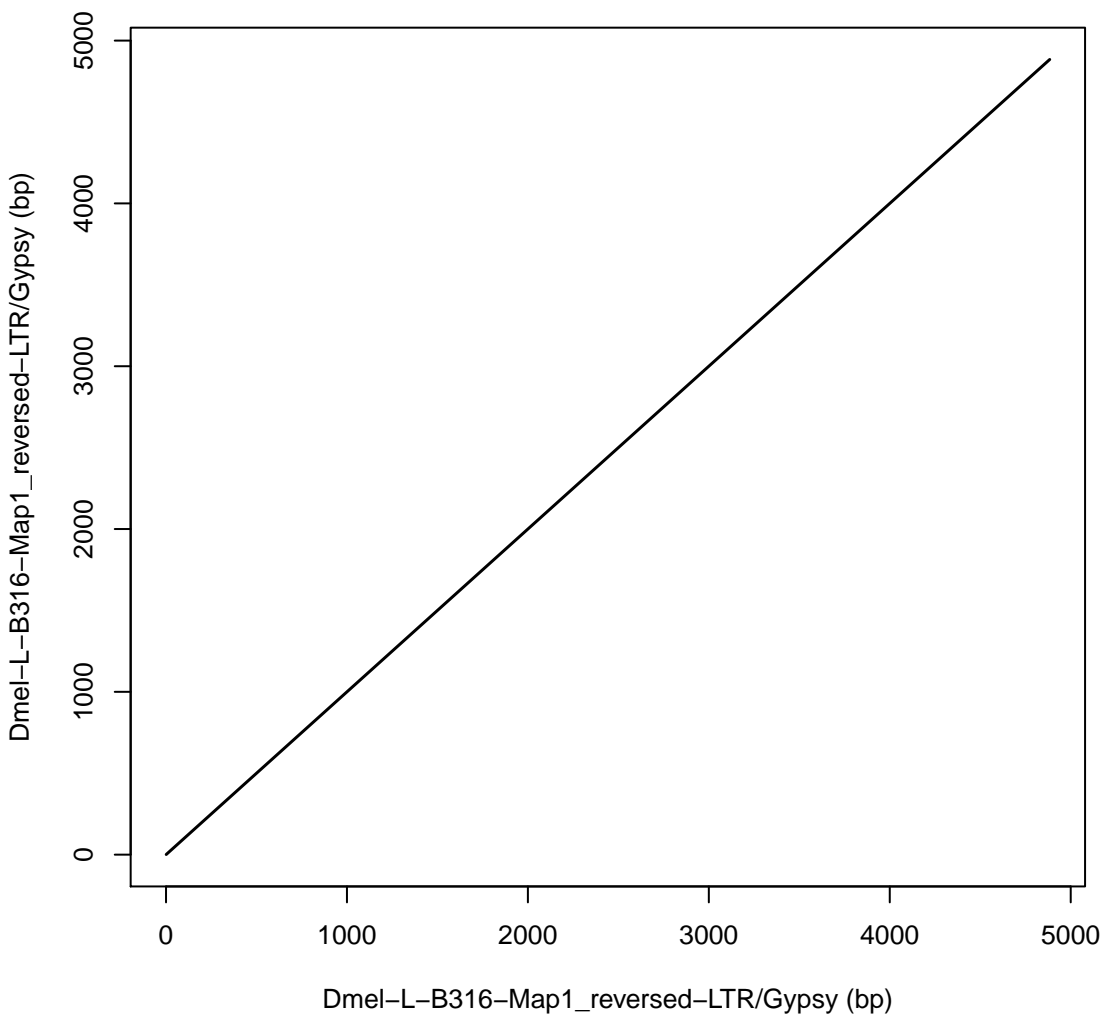
TE: Dmel-L-B316-Map1_reversed-LTR/Gypsy
consensus size: 4884bp; fragments: 191; full length: 8 (≥ 4395.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

