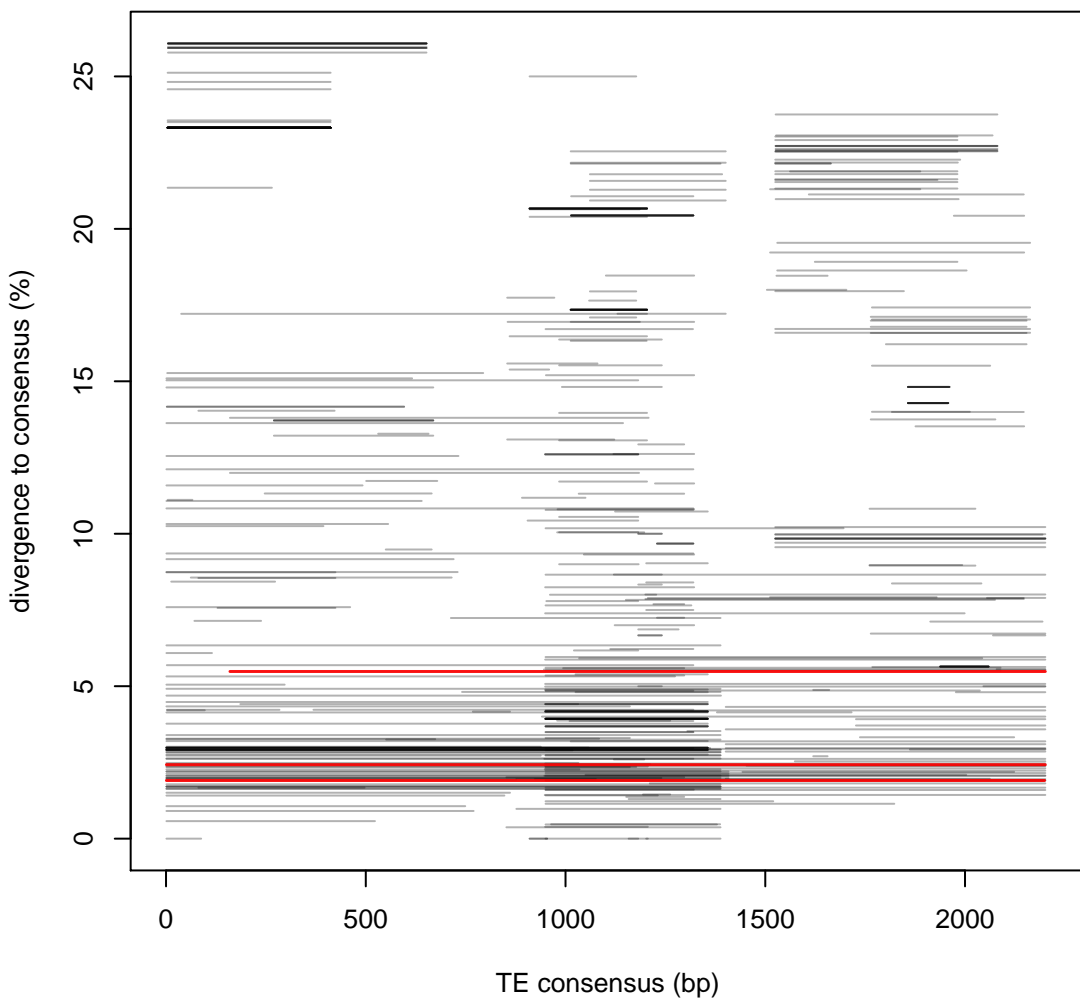
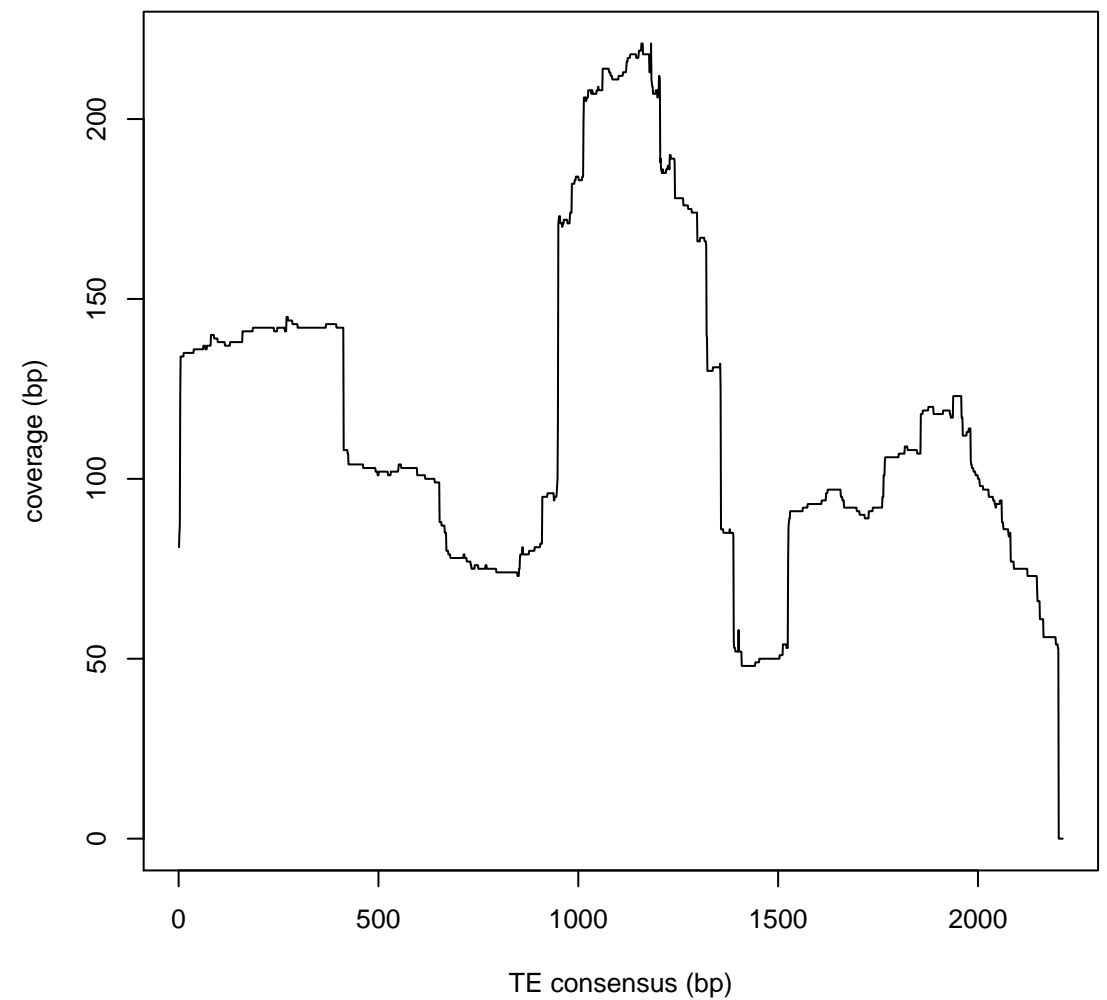


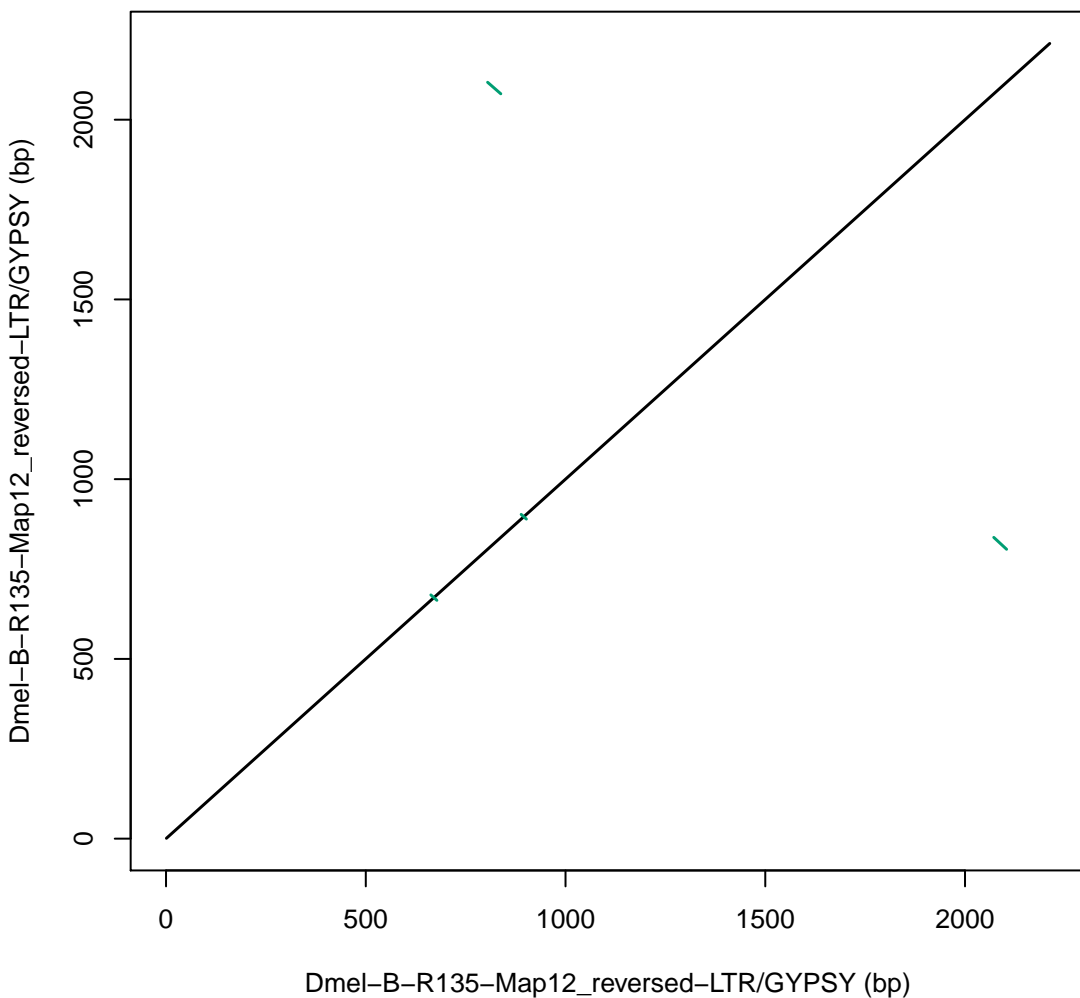
TE: Dmel-B-R135-Map12_reversed-LTR/GYPsy
consensus size: 2212bp; fragments: 457; full length: 3 (>=1990.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

