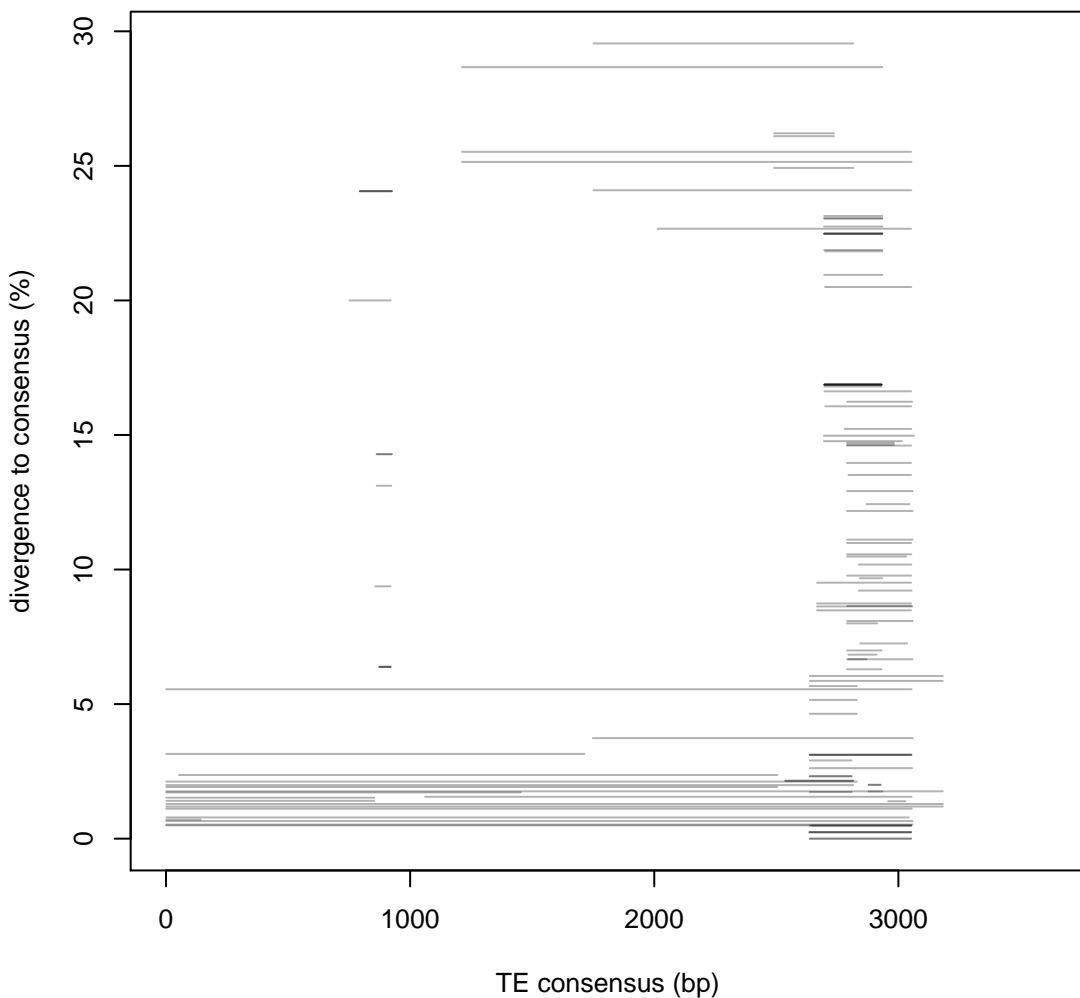
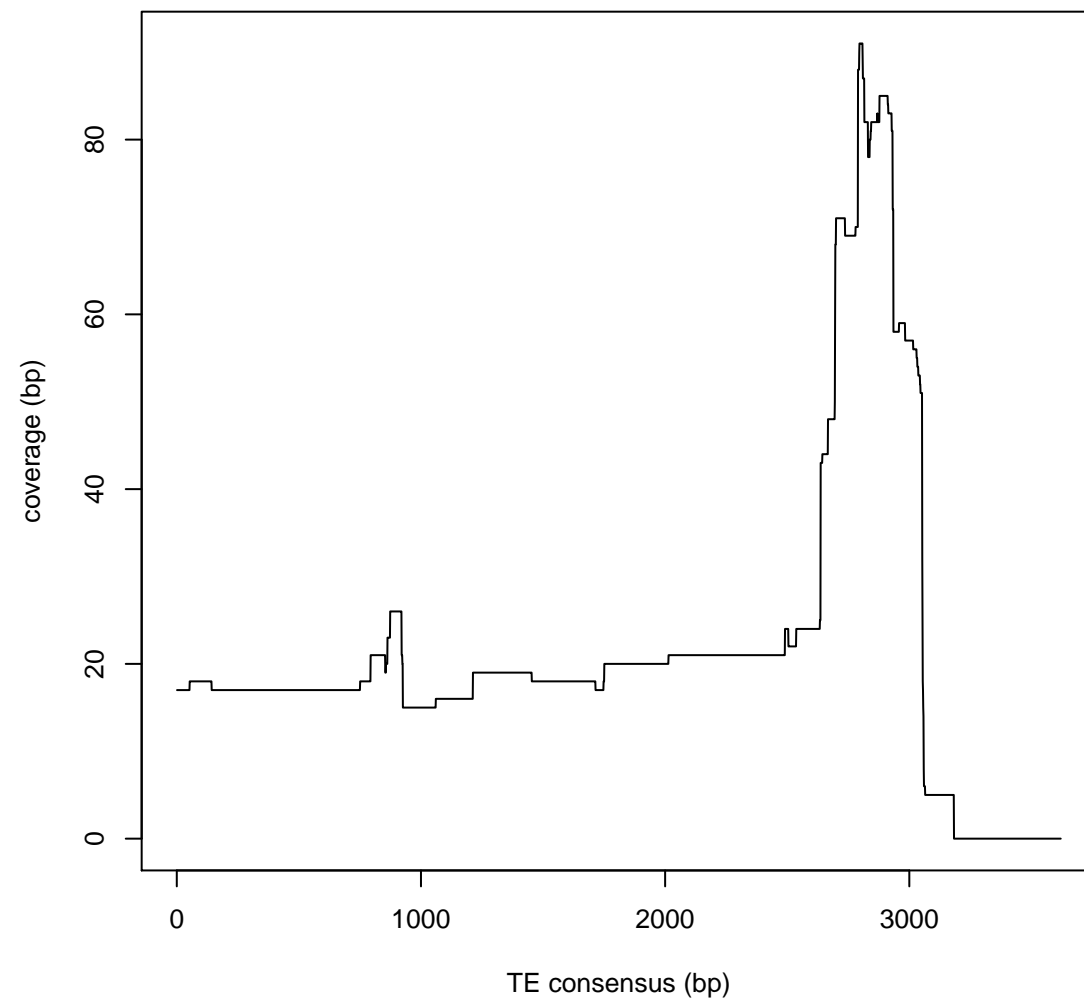


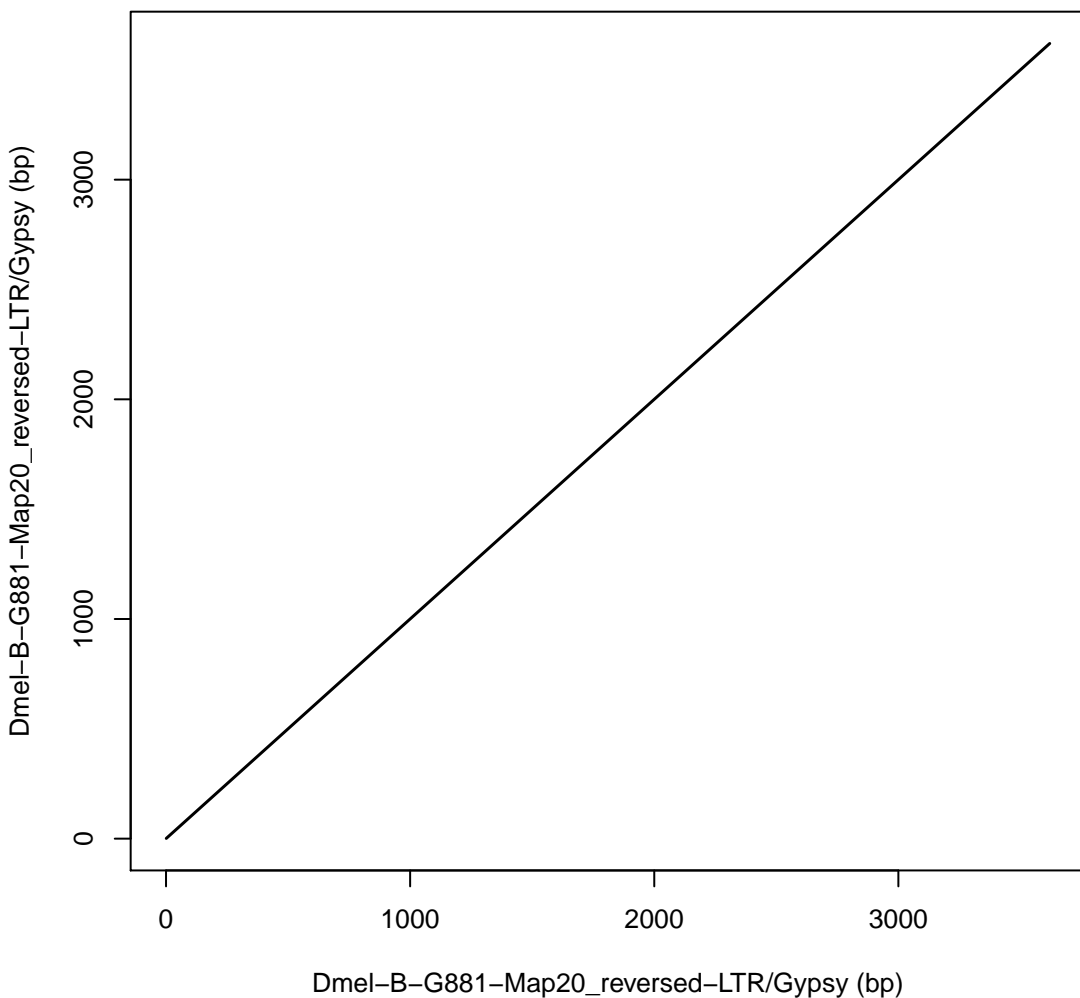
TE: Dmel-B-G881-Map20_reversed-LTR/Gypsy
consensus size: 3620bp; fragments: 120; full length: 0 (>=3258bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

