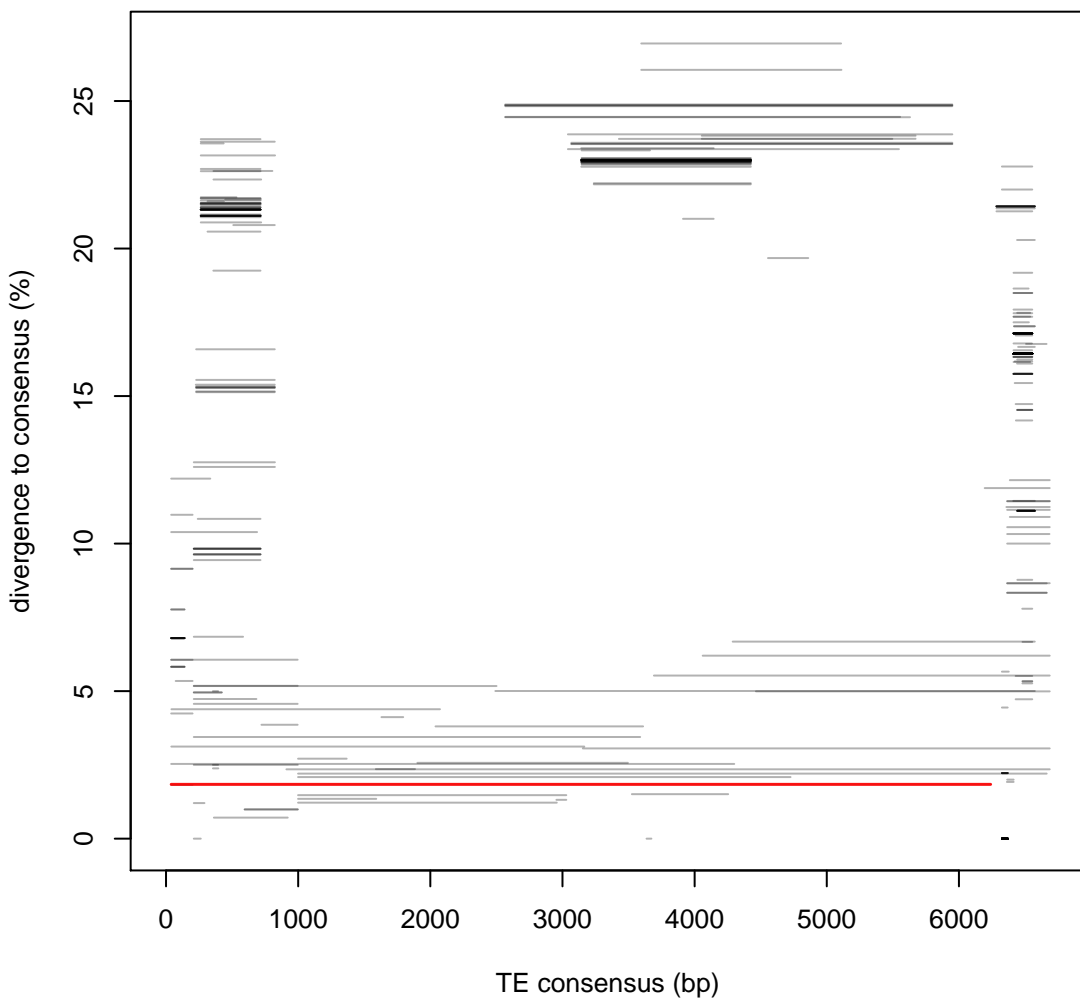
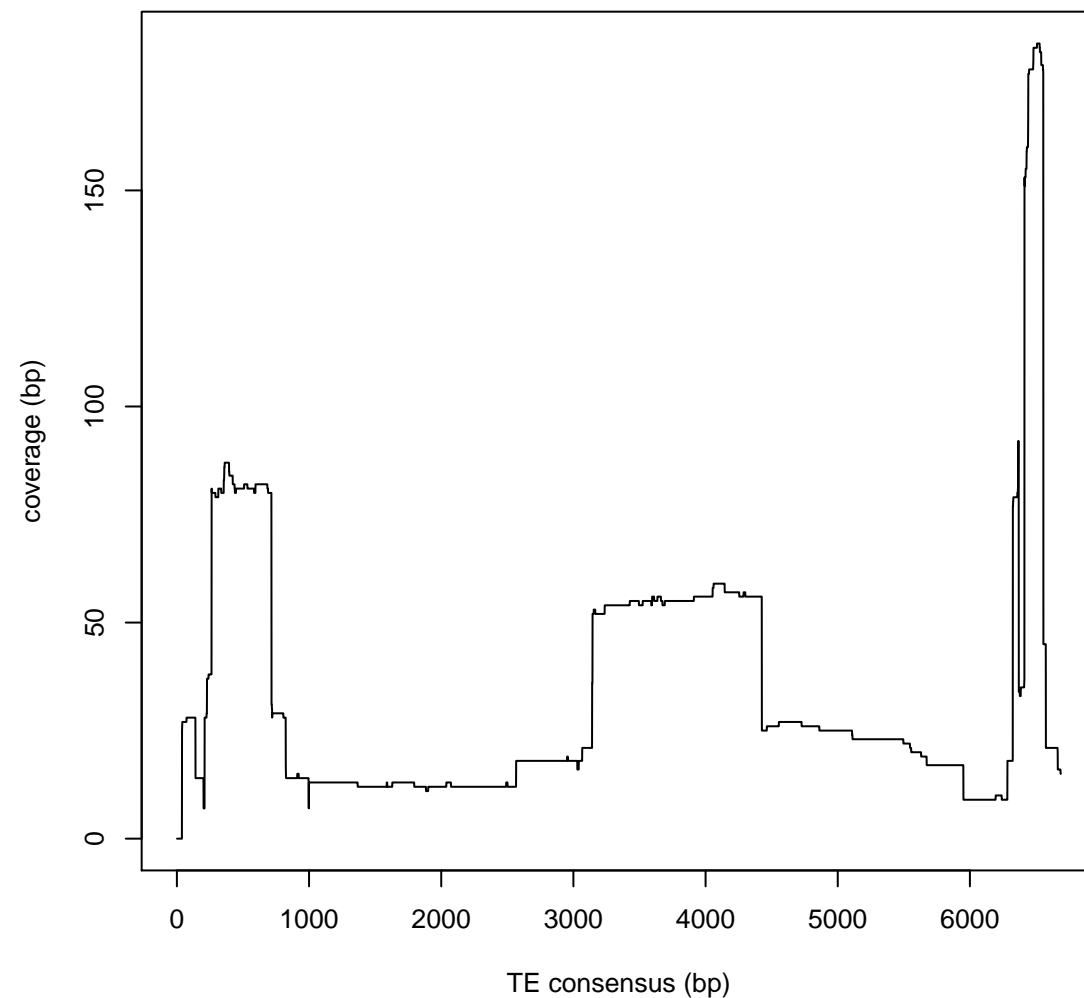


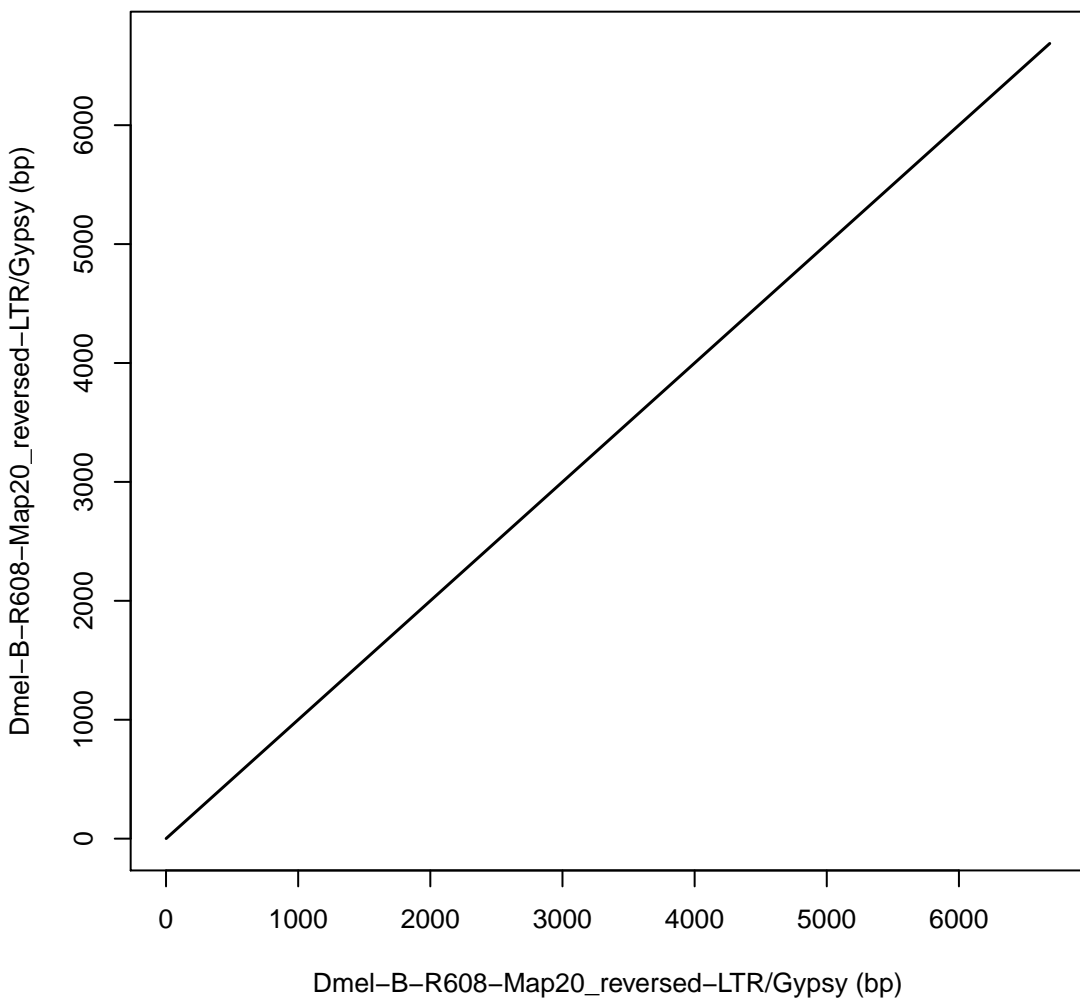
TE: Dmel-B-R608-Map20_reversed-LTR/Gypsy
consensus size: 6687bp; fragments: 424; full length: 1 (>=6018.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

