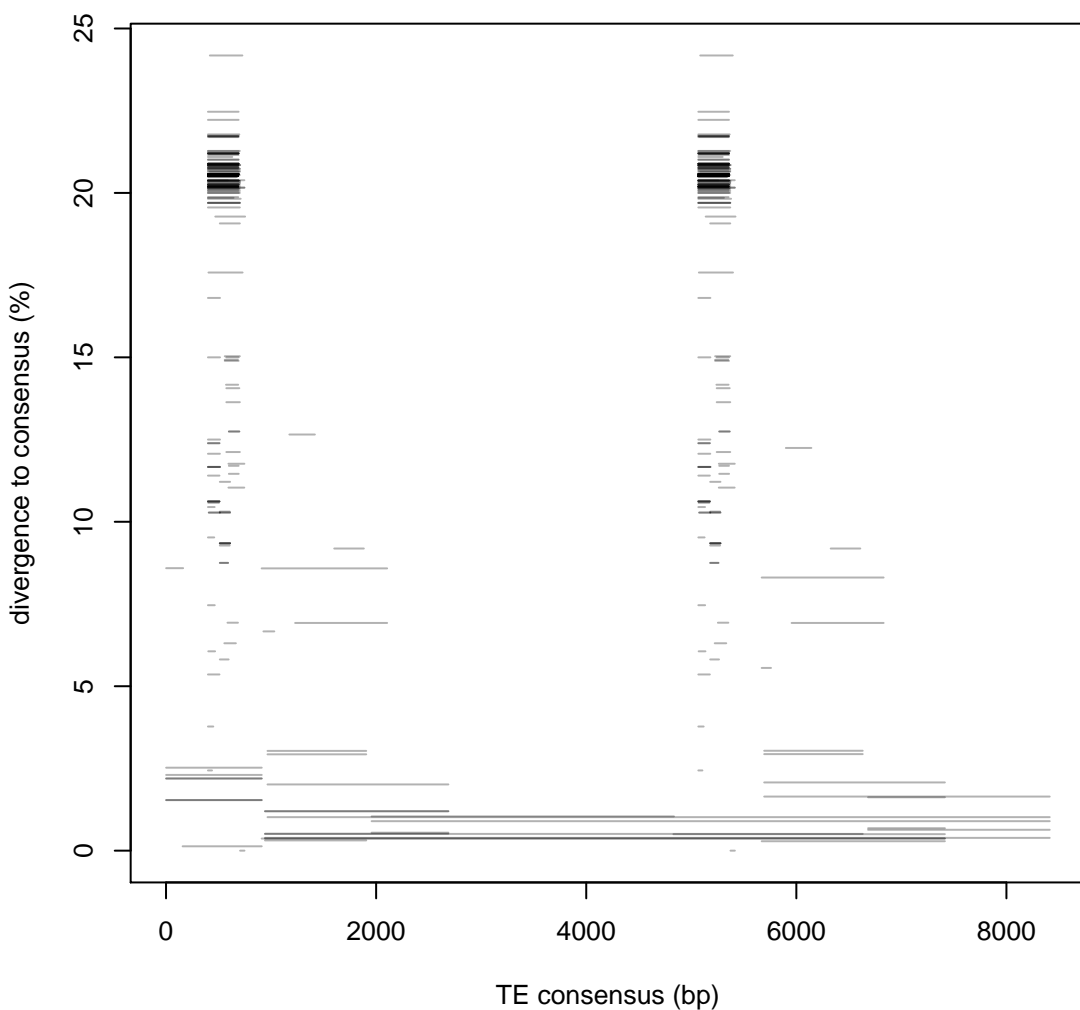
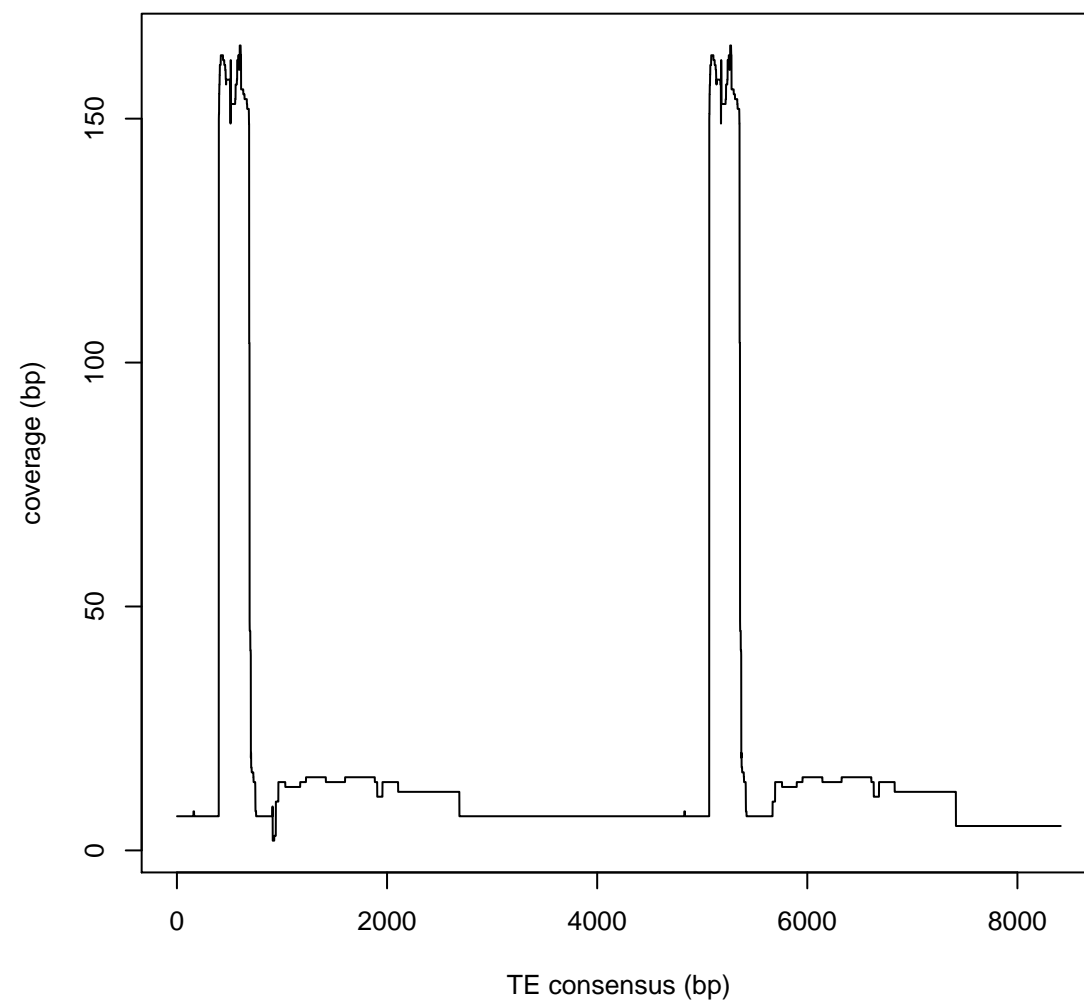


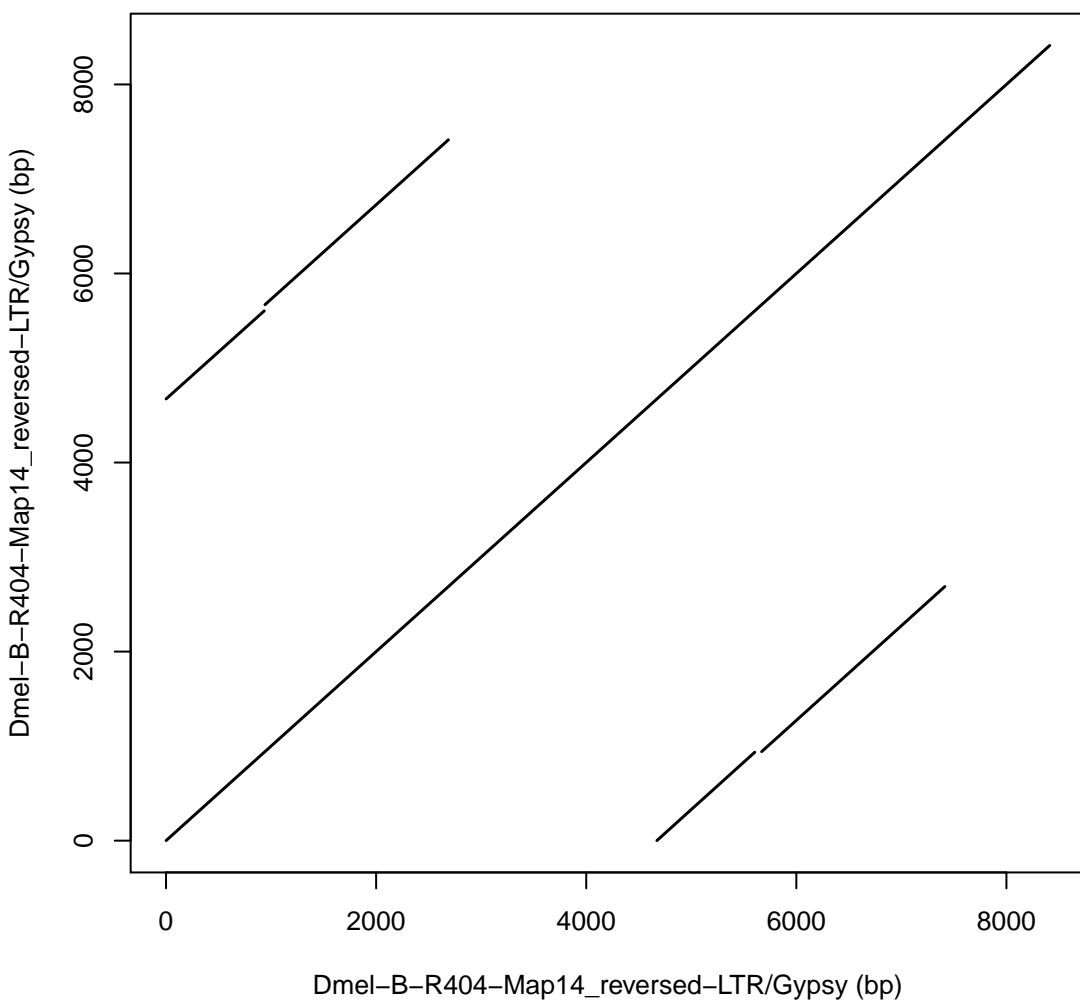
TE: Dmel-B-R404-Map14_reversed-LTR/Gypsy
consensus size: 8412bp; fragments: 416; full length: 0 (>=7570.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

