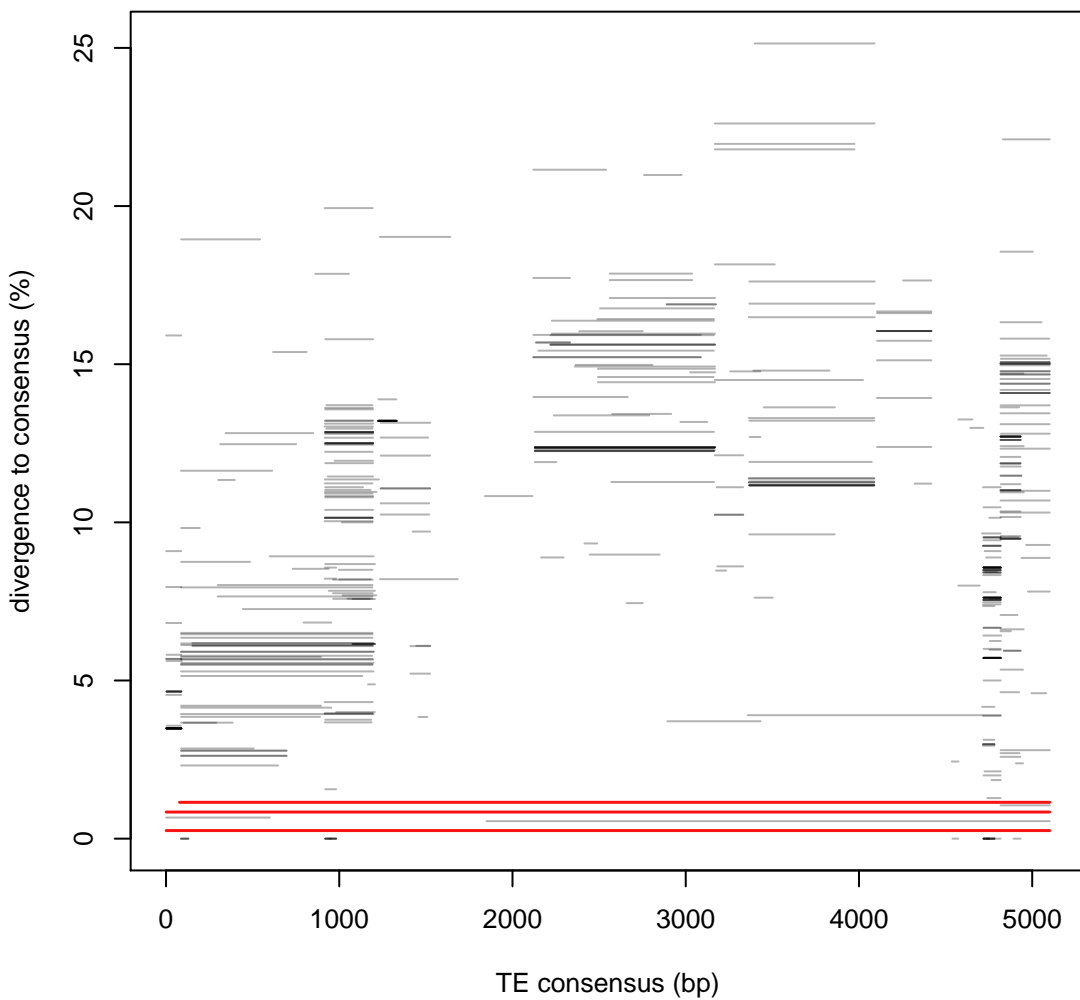
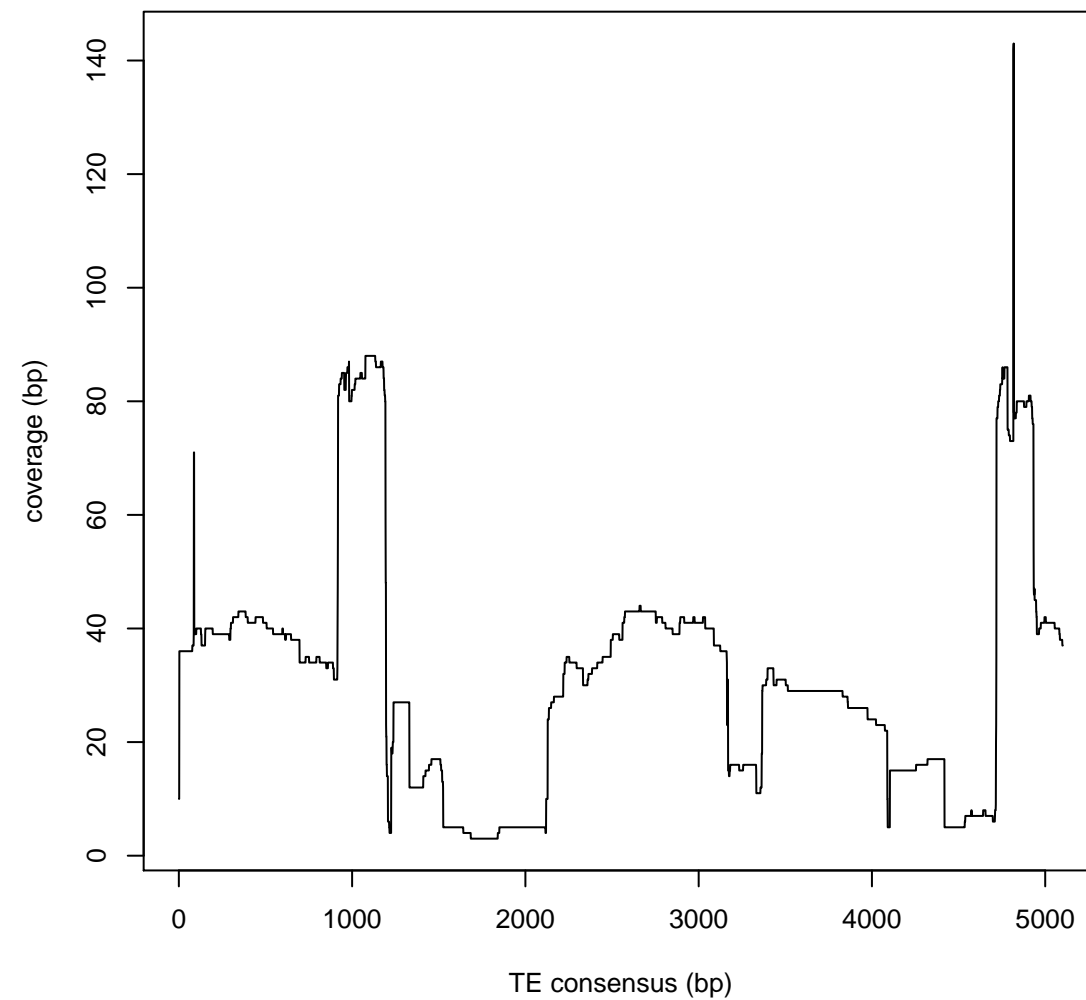


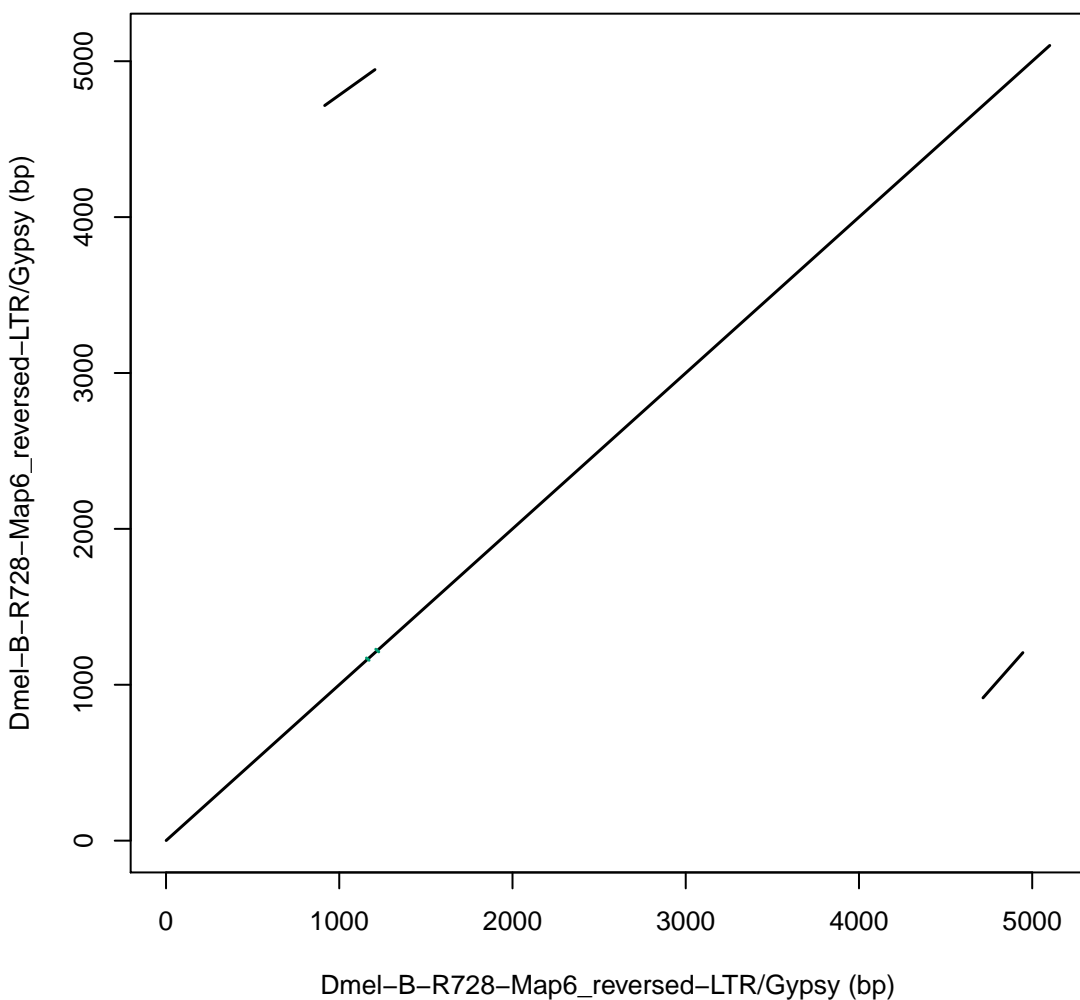
TE: Dmel-B-R728-Map6_reversed-LTR/Gypsy
consensus size: 5101bp; fragments: 459; full length: 3 (>=4590.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

