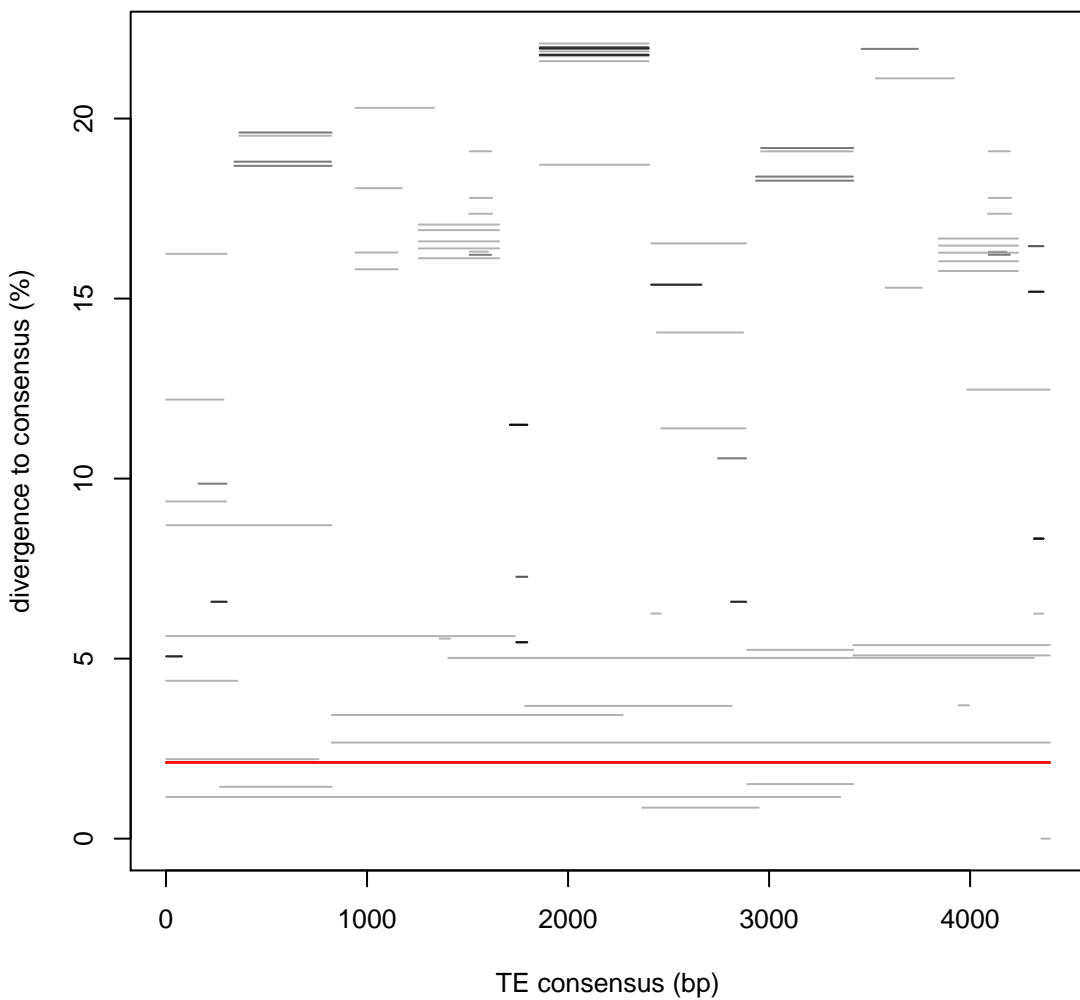
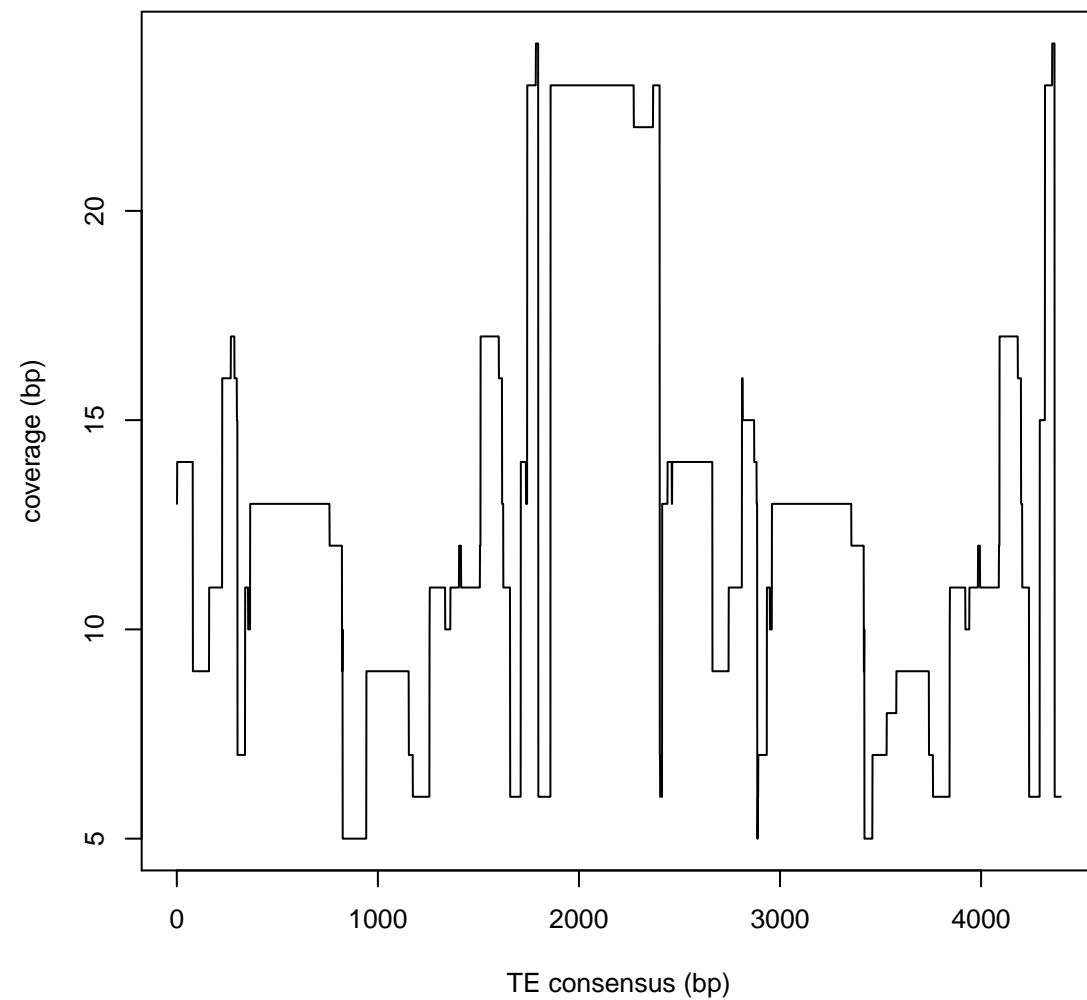


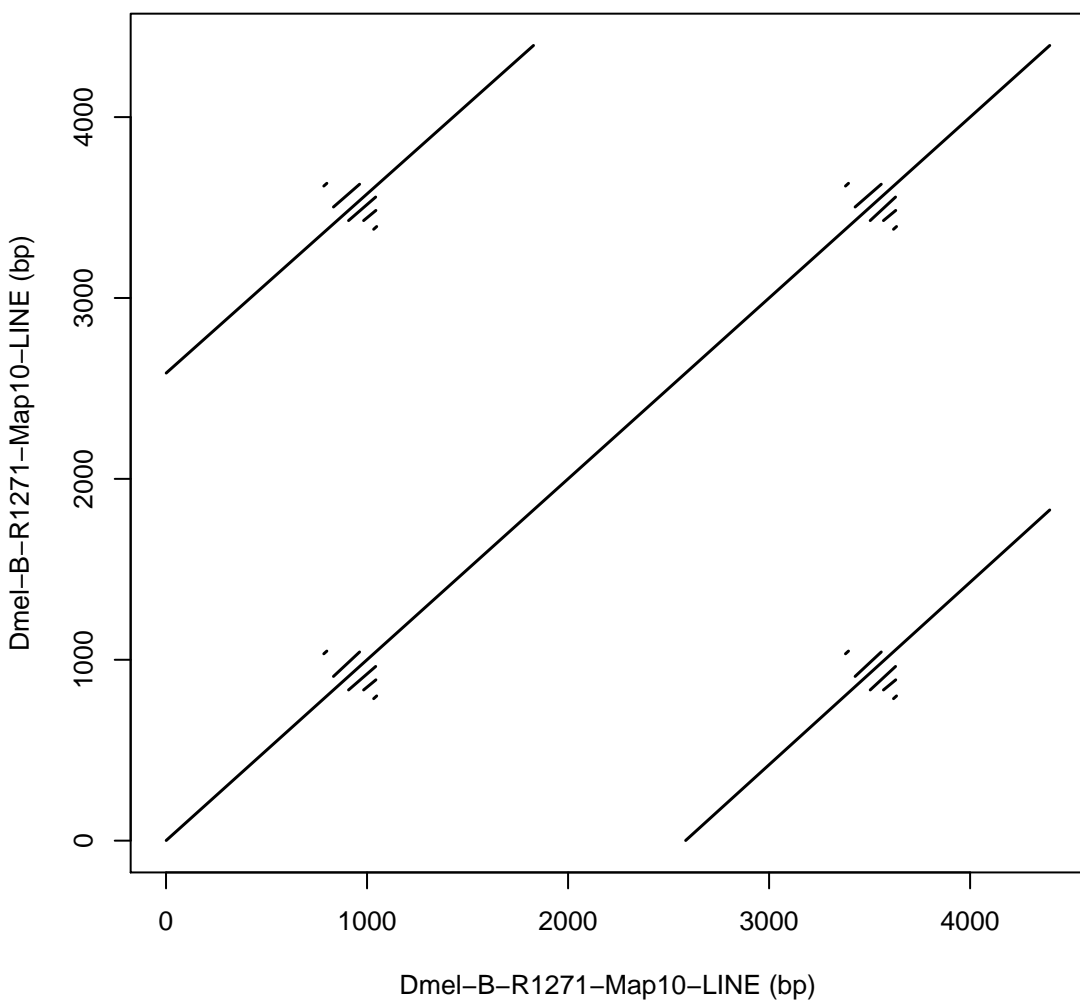
TE: Dmel-B-R1271-Map10-LINE
consensus size: 4396bp; fragments: 148; full length: 1 (>=3956.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

