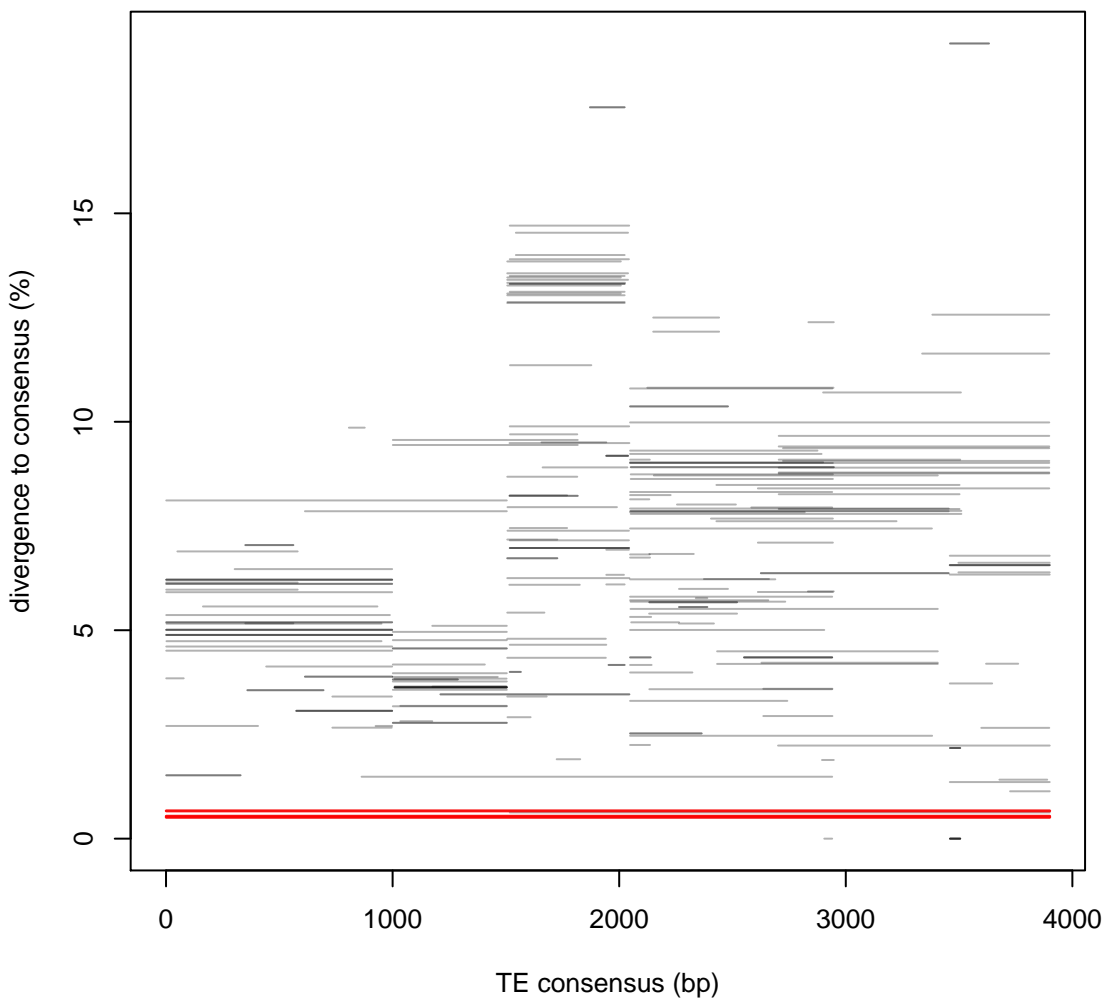
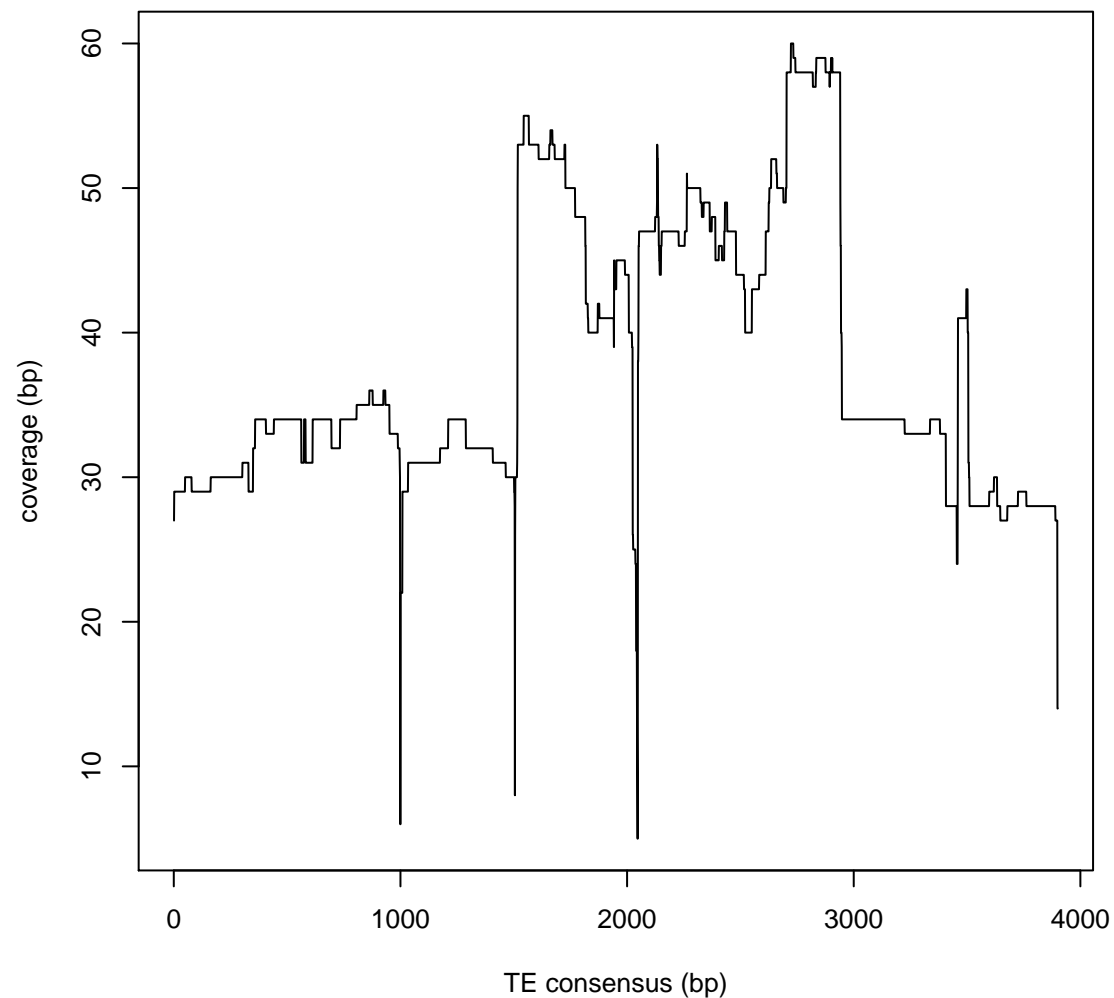


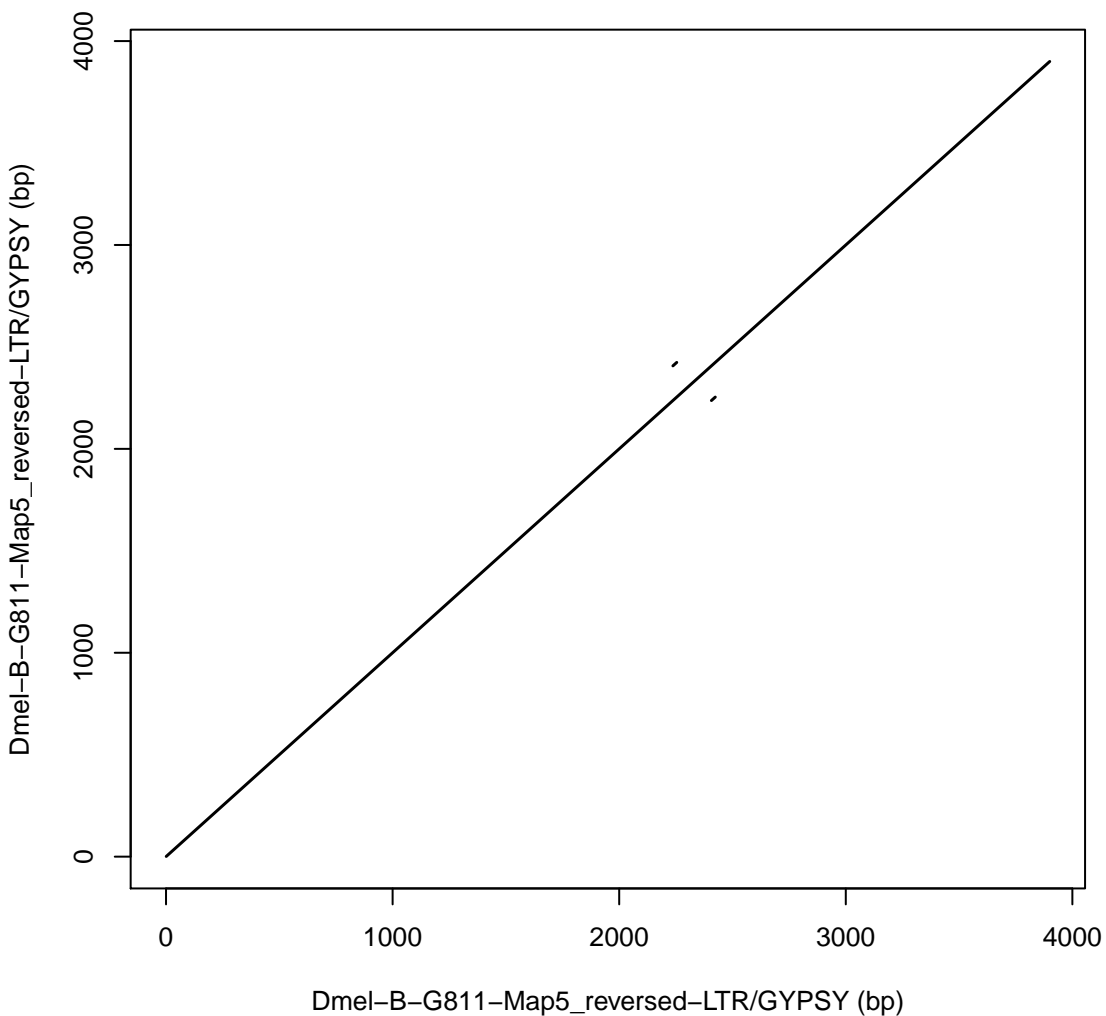
TE: Dmel-B-G811-Map5_reversed-LTR/Gypsy
consensus size: 3900bp; fragments: 254; full length: 3 (>=3510bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

