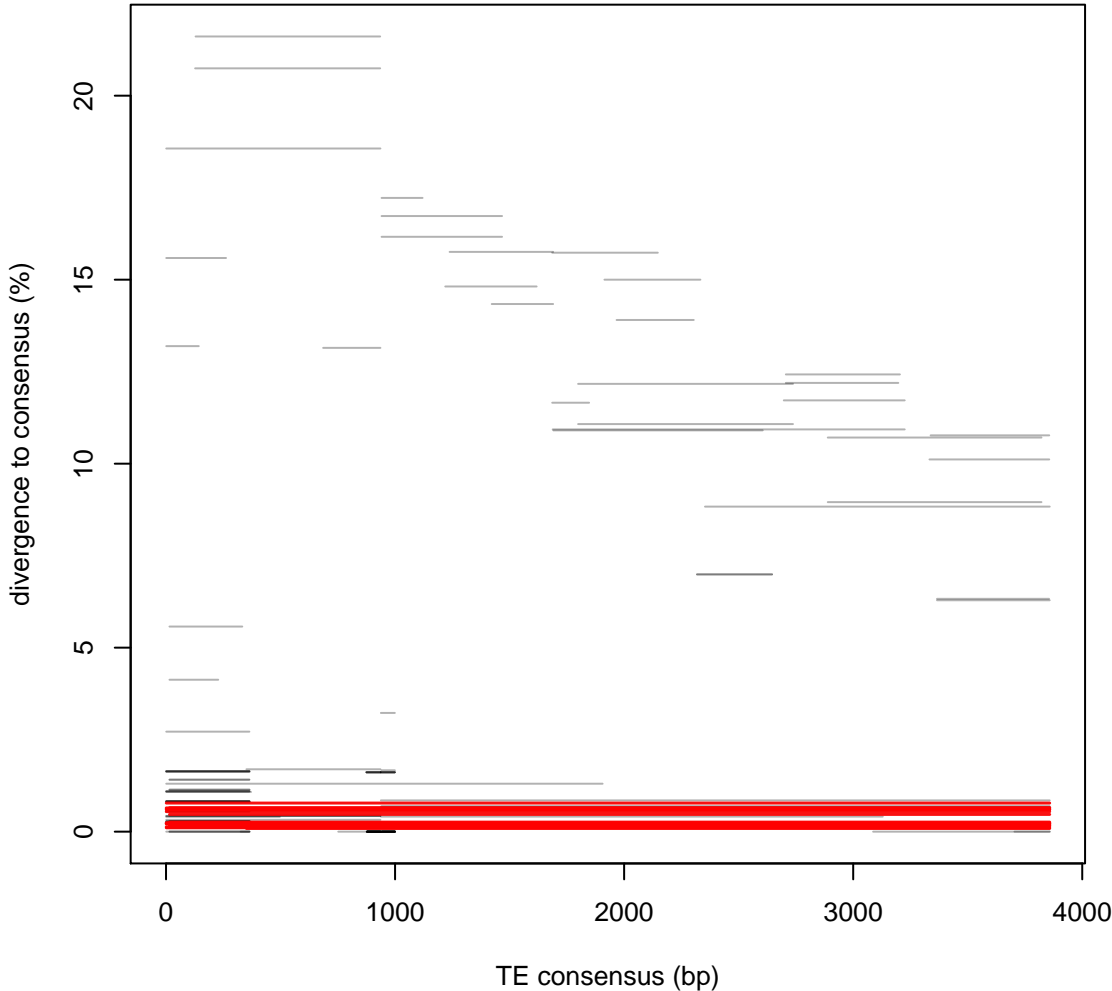
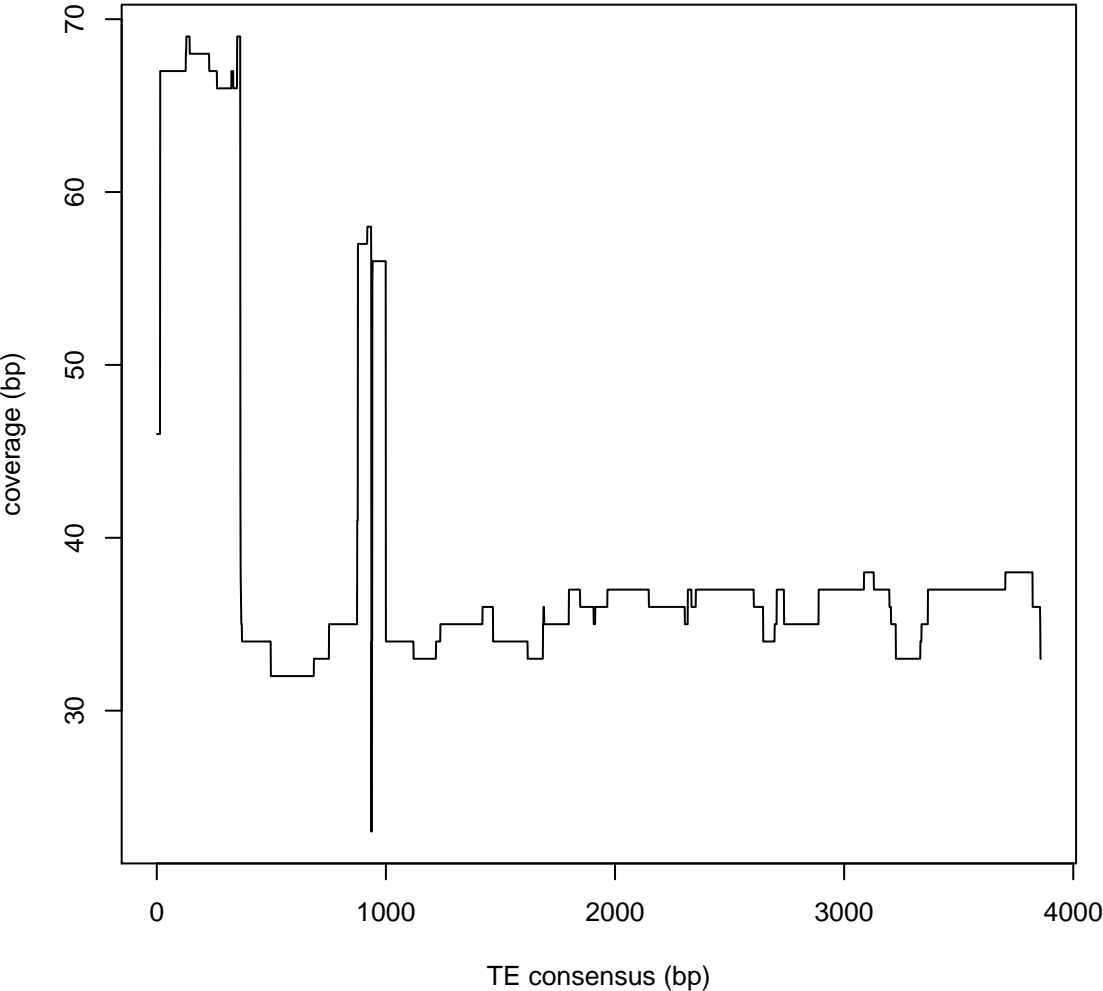


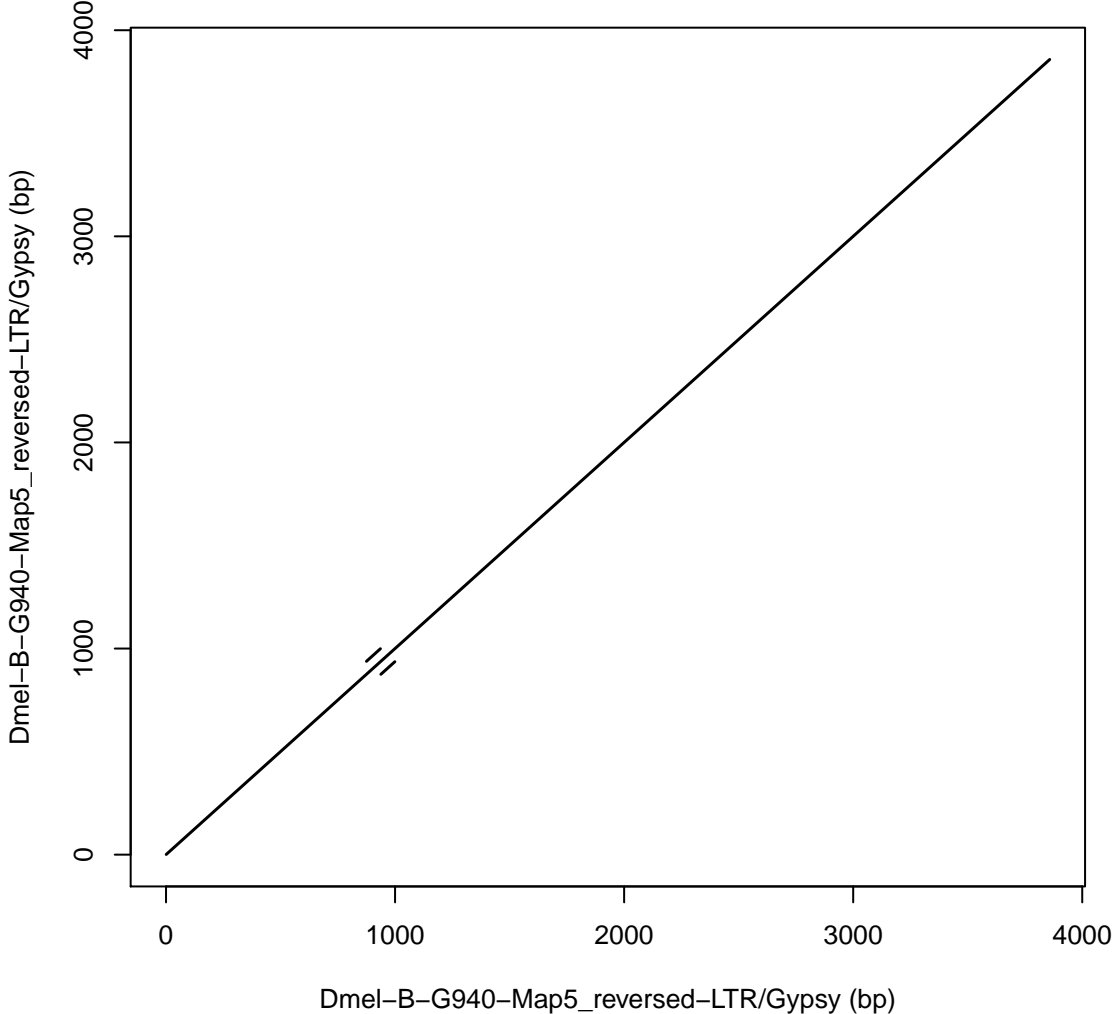
TE: Dmel-B-G940-Map5_reversed-LTR/Gypsy
consensus size: 3858bp; fragments: 157; full length: 20 (>=3472.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

