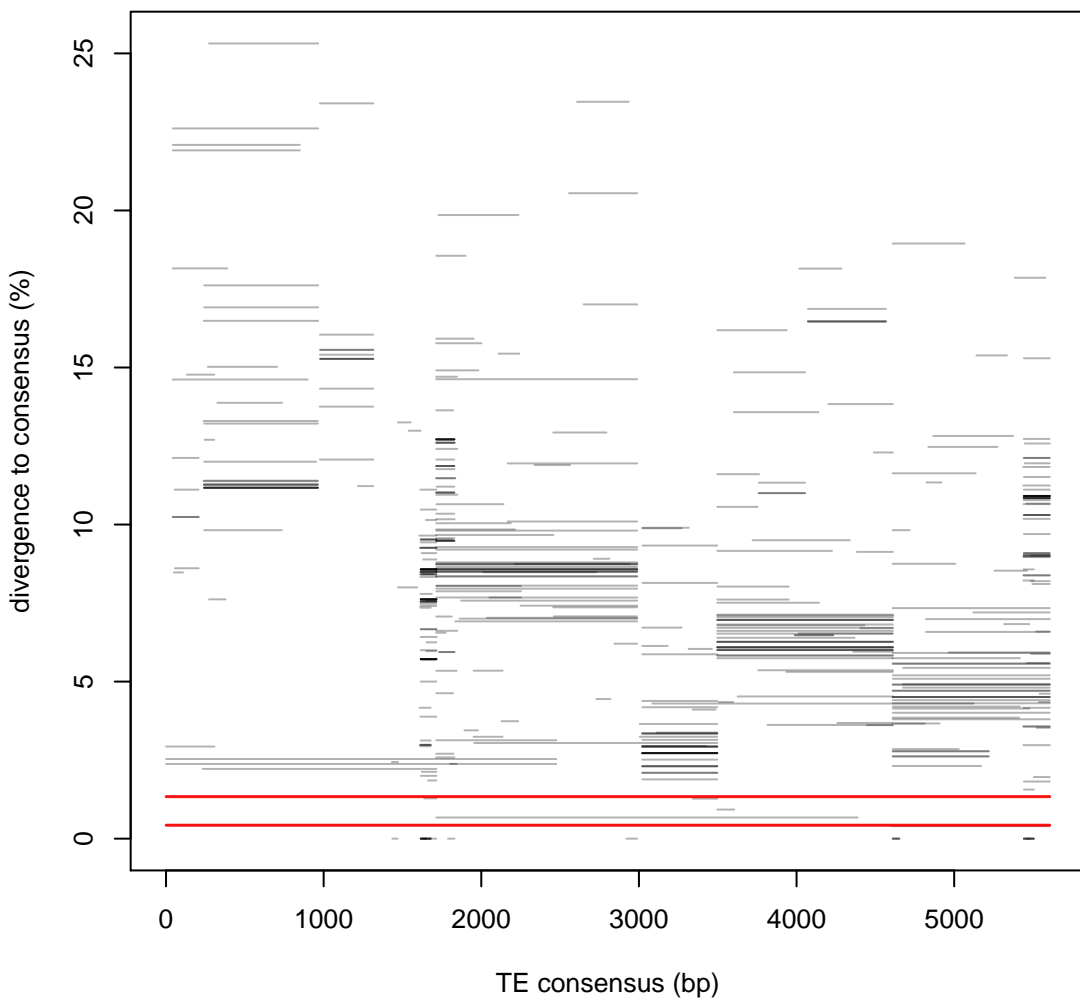
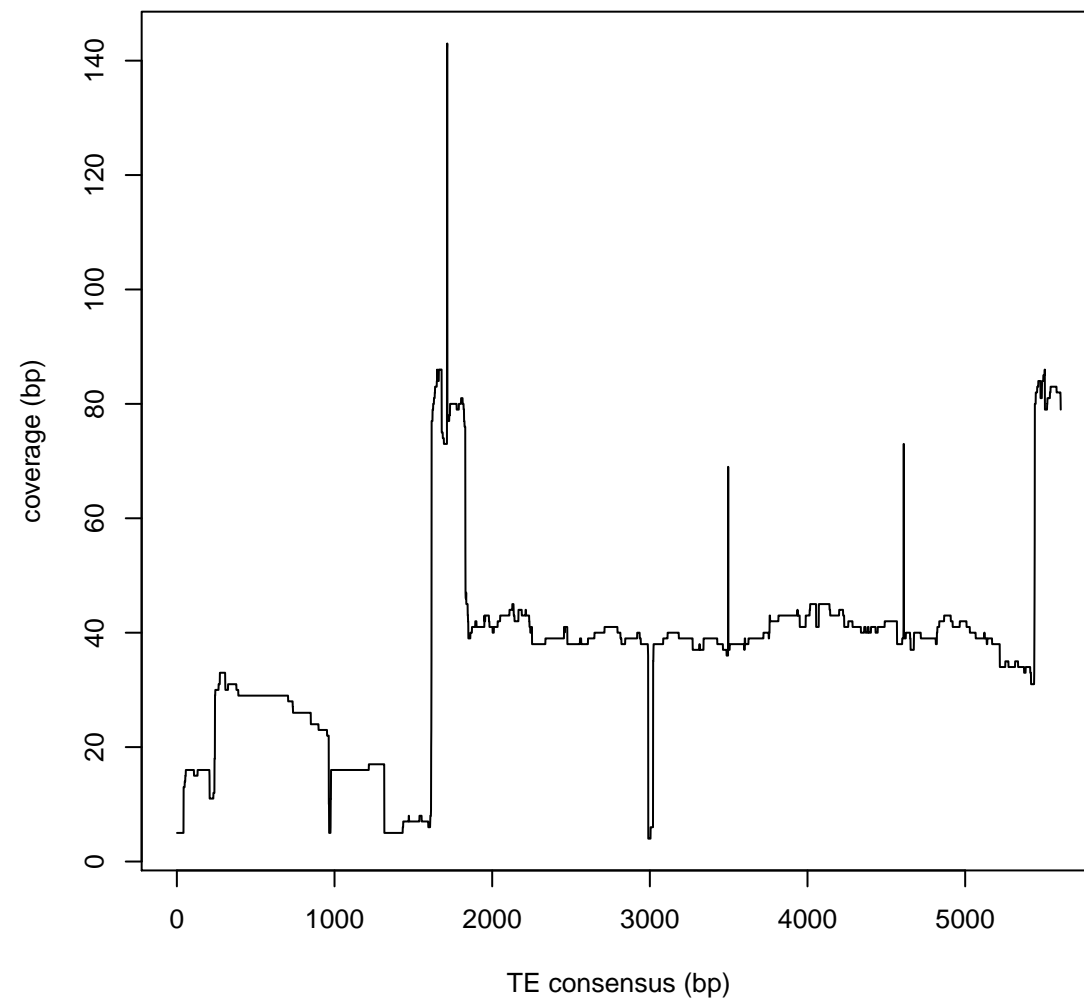


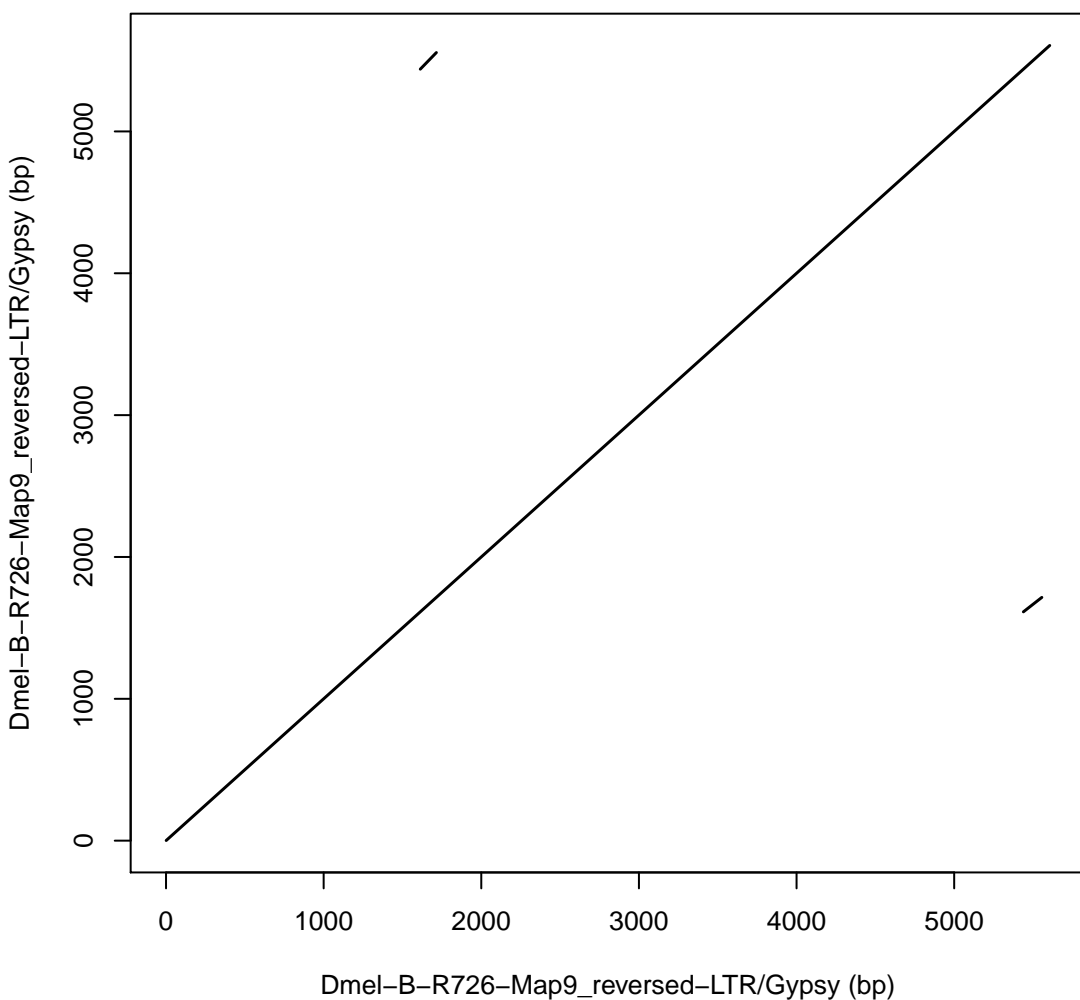
TE: Dmel-B-R726-Map9_reversed-LTR/Gypsy
consensus size: 5606bp; fragments: 458; full length: 2 (>=5045.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

