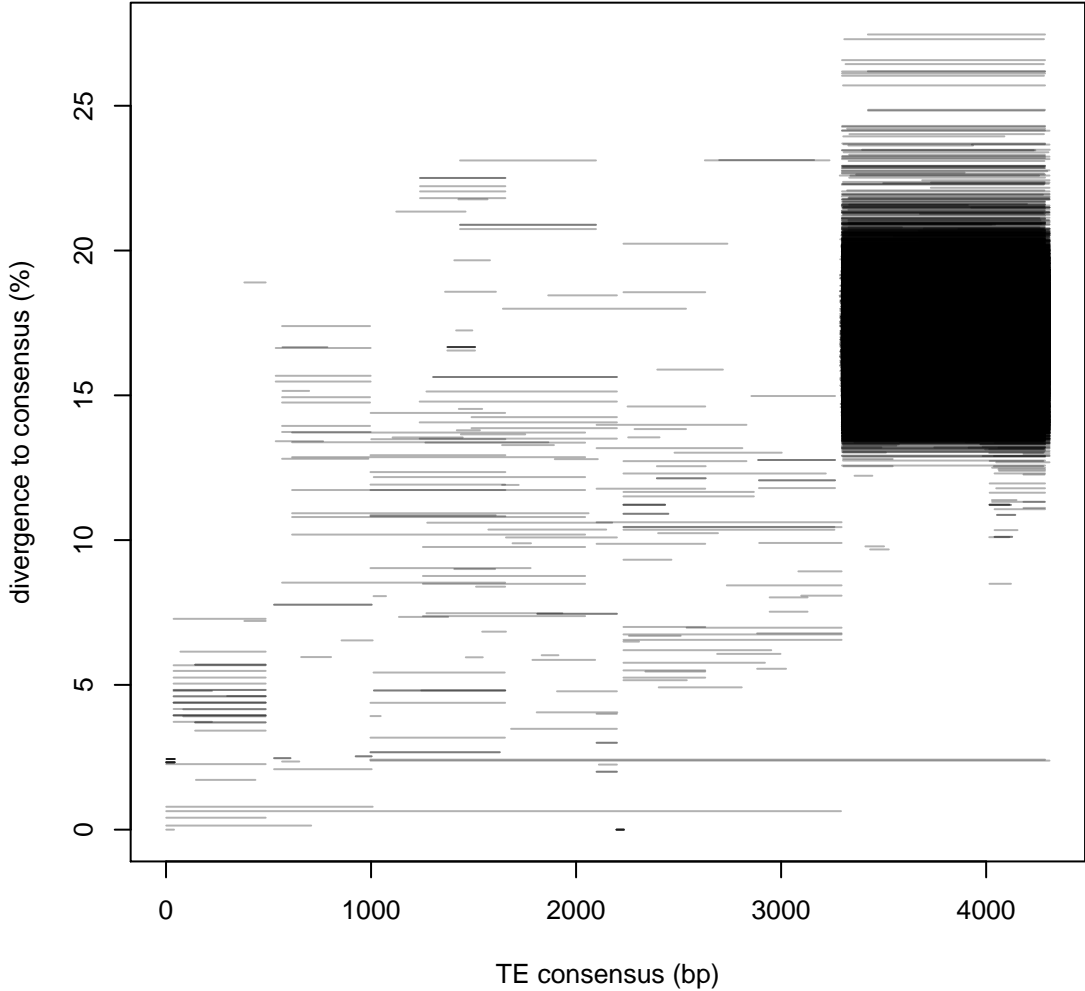
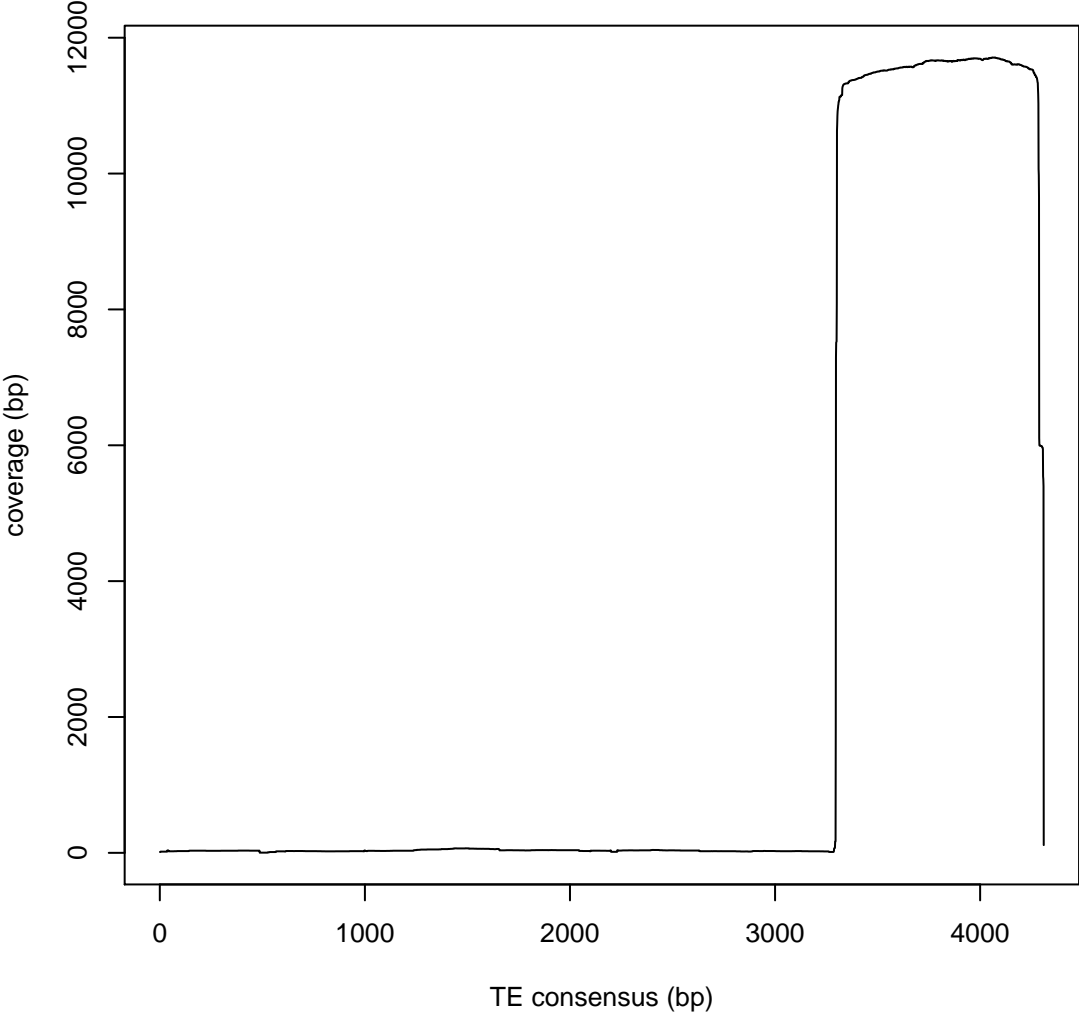


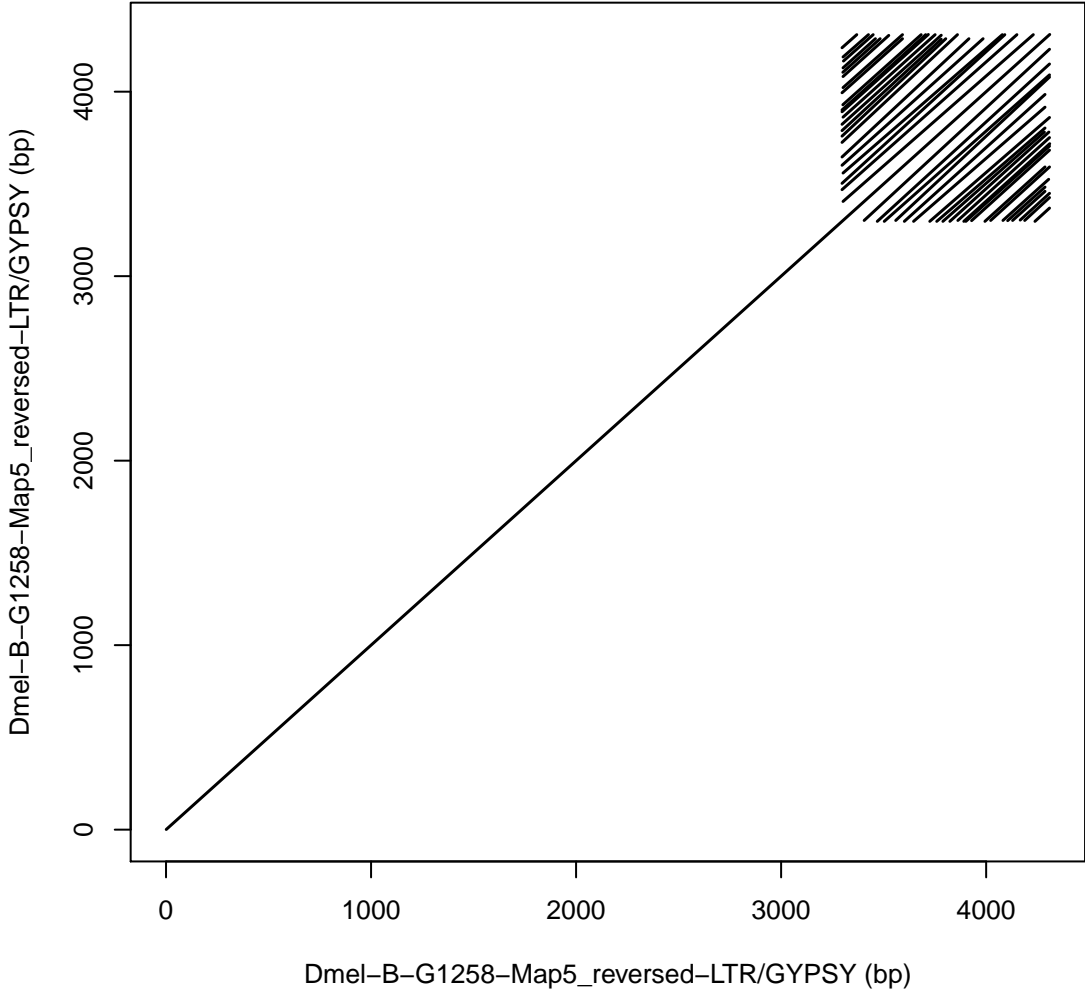
TE: Dmel-B-G1258-Map5_reversed-LTR/GYPSY
consensus size: 4310bp; fragments: 12241; full length: 0 (>=3879bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

