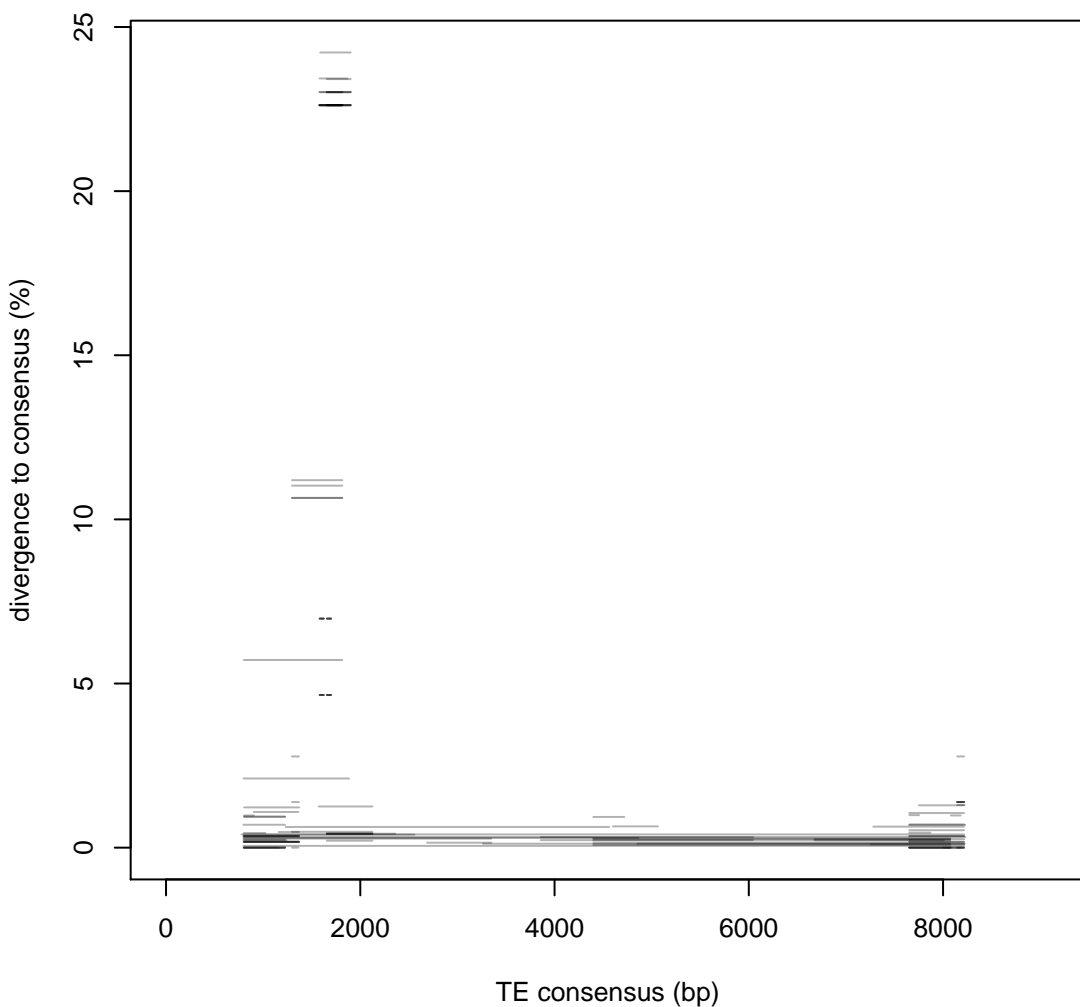
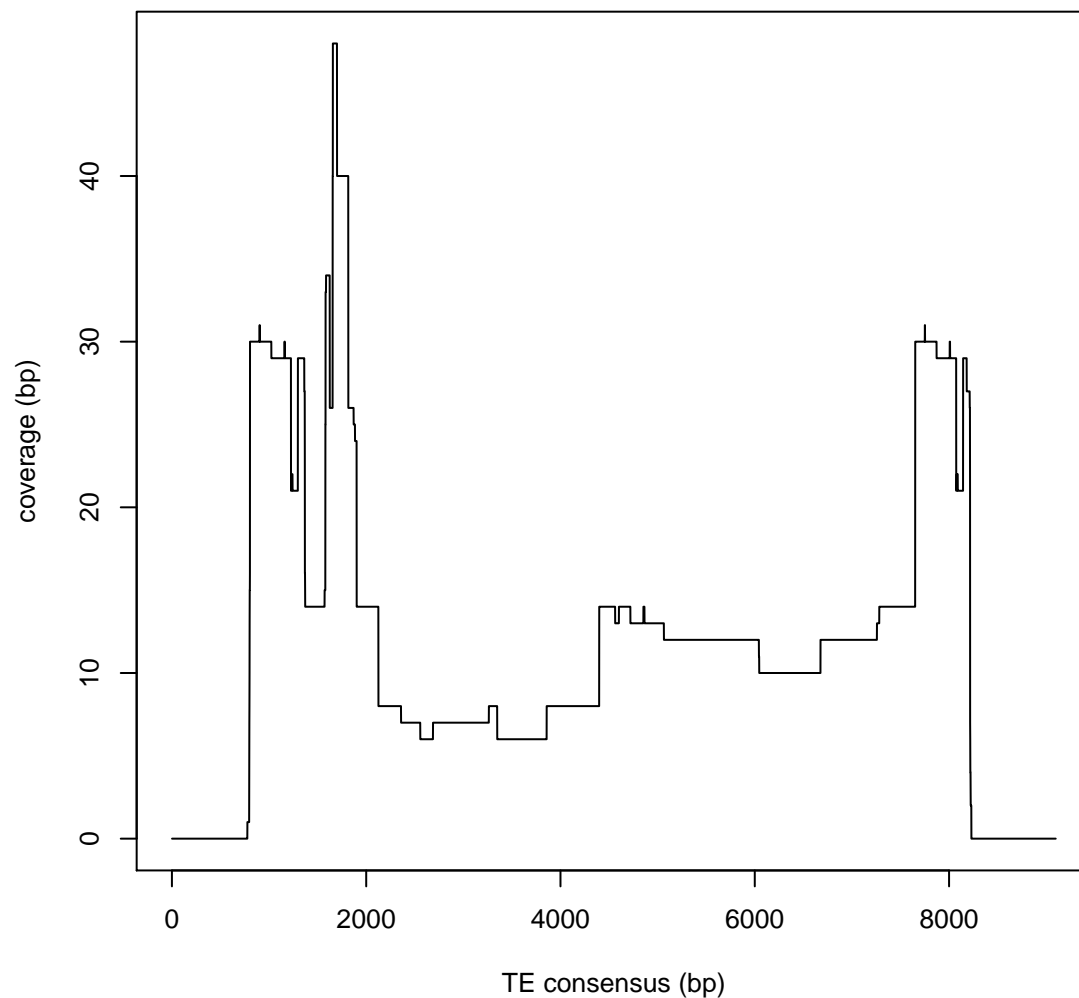


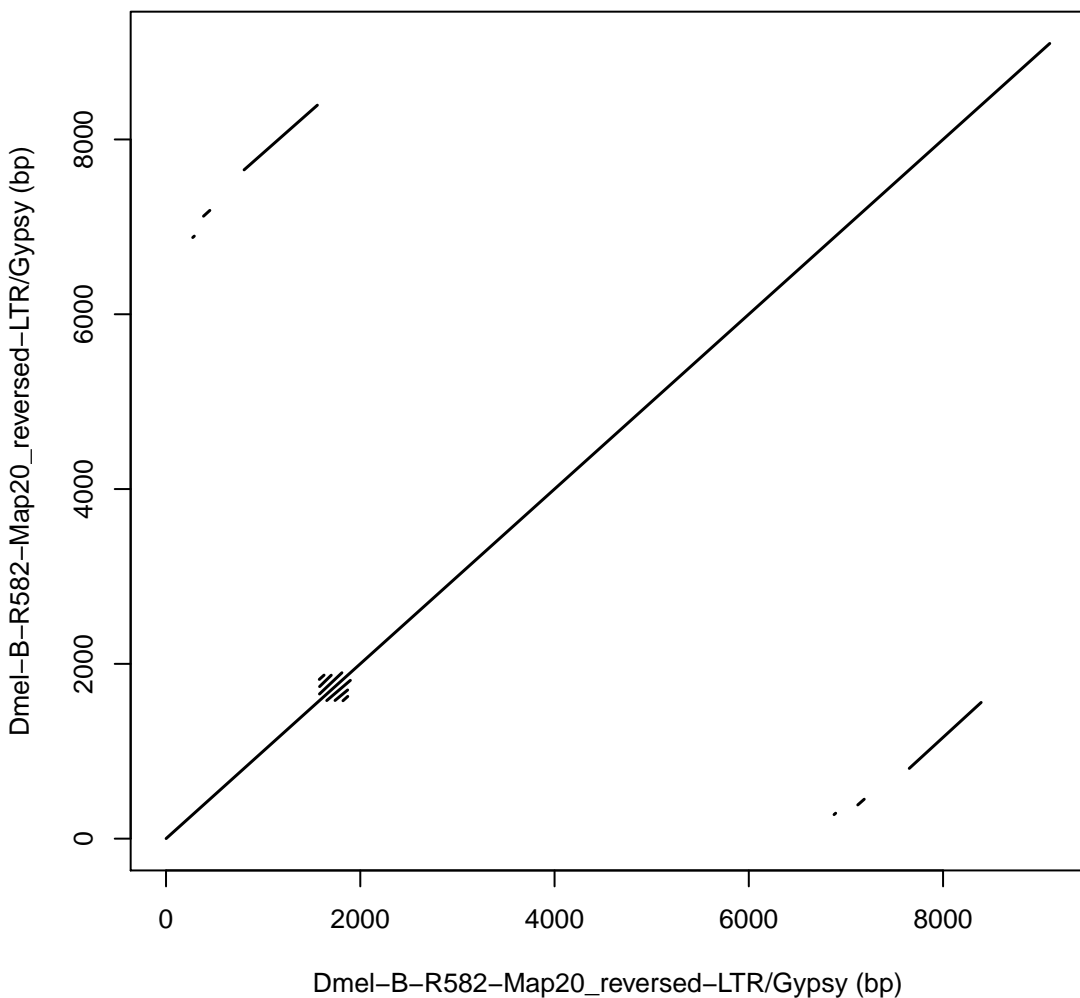
TE: Dmel-B-R582-Map20\_reversed-LTR/Gypsy  
consensus size: 9098bp; fragments: 126; full length: 0 (>=8188.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

