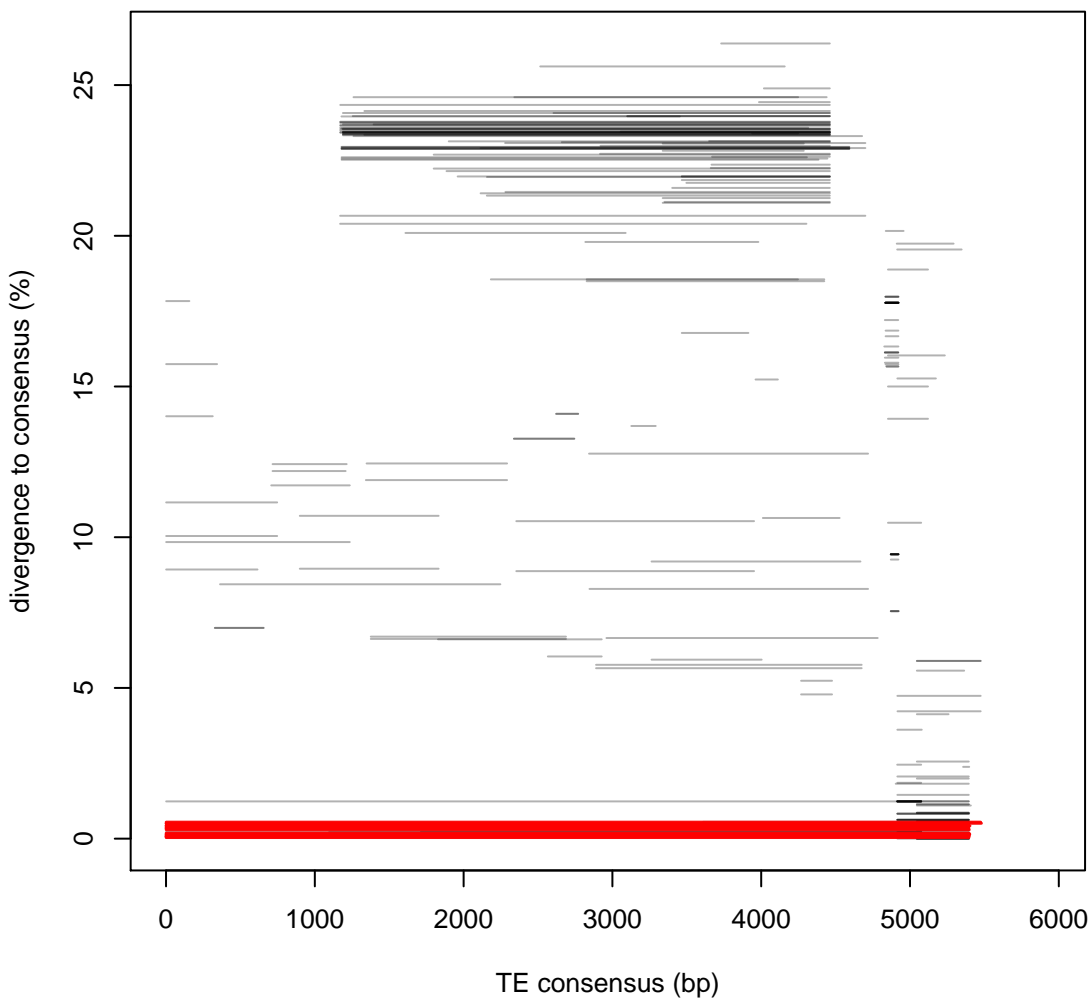
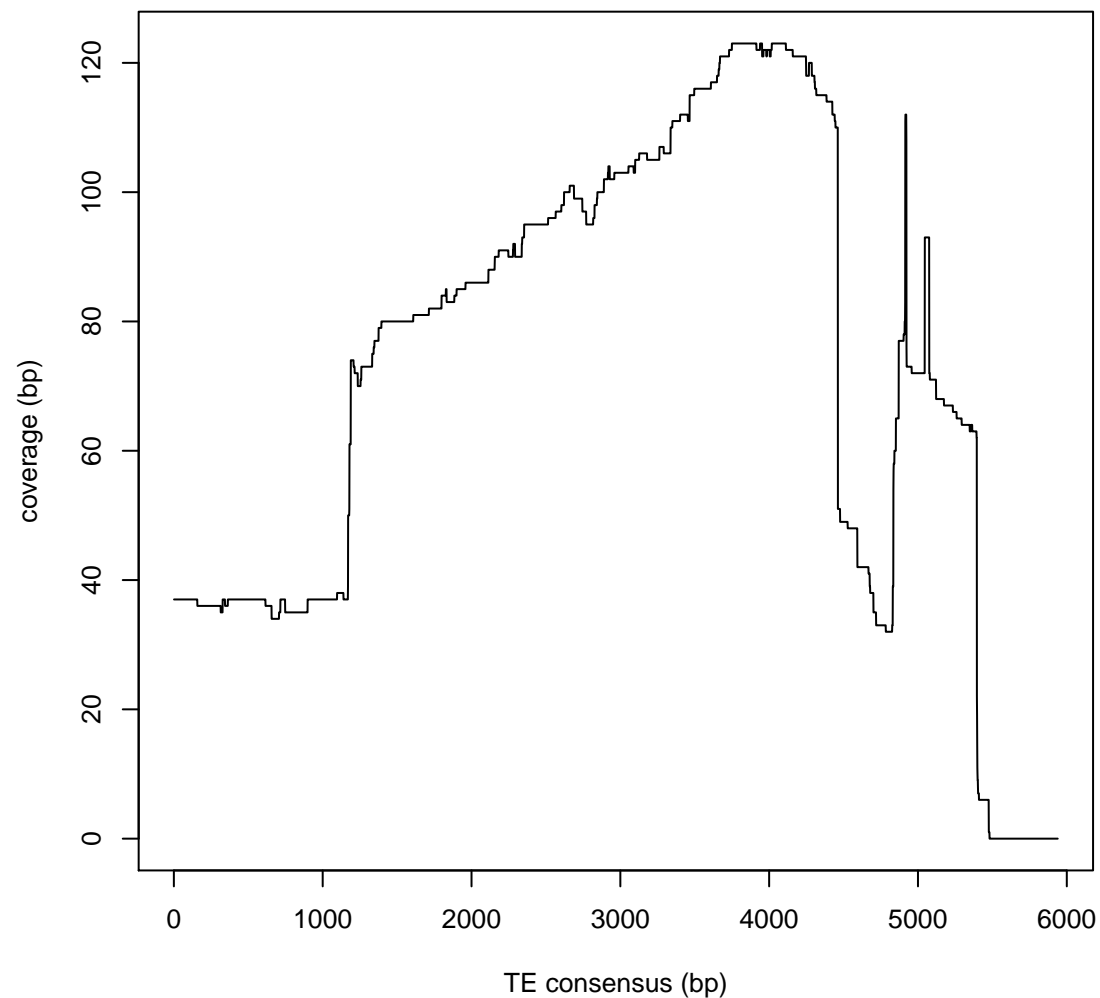


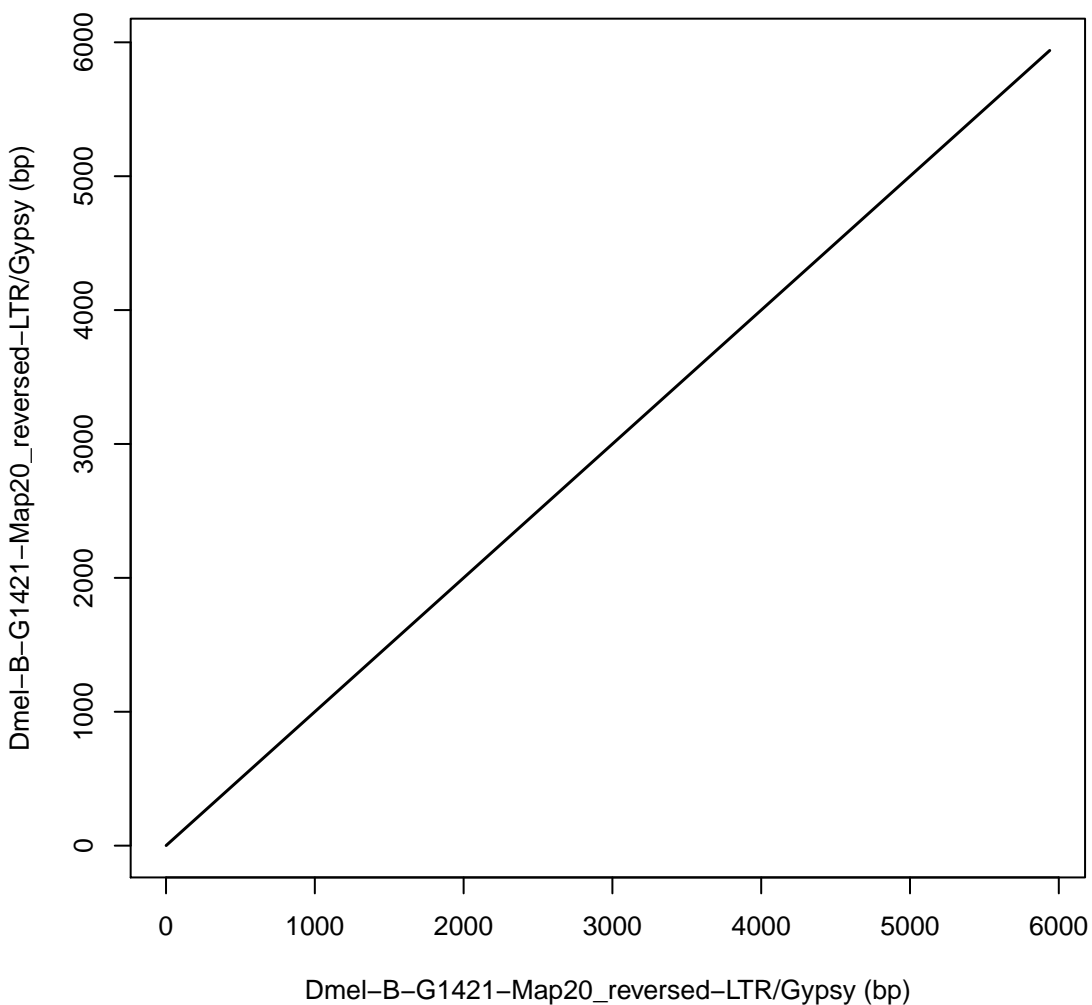
TE: Dmel-B-G1421-Map20\_reversed-LTR/Gypsy  
consensus size: 5939bp; fragments: 262; full length: 19 (>=5345.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

