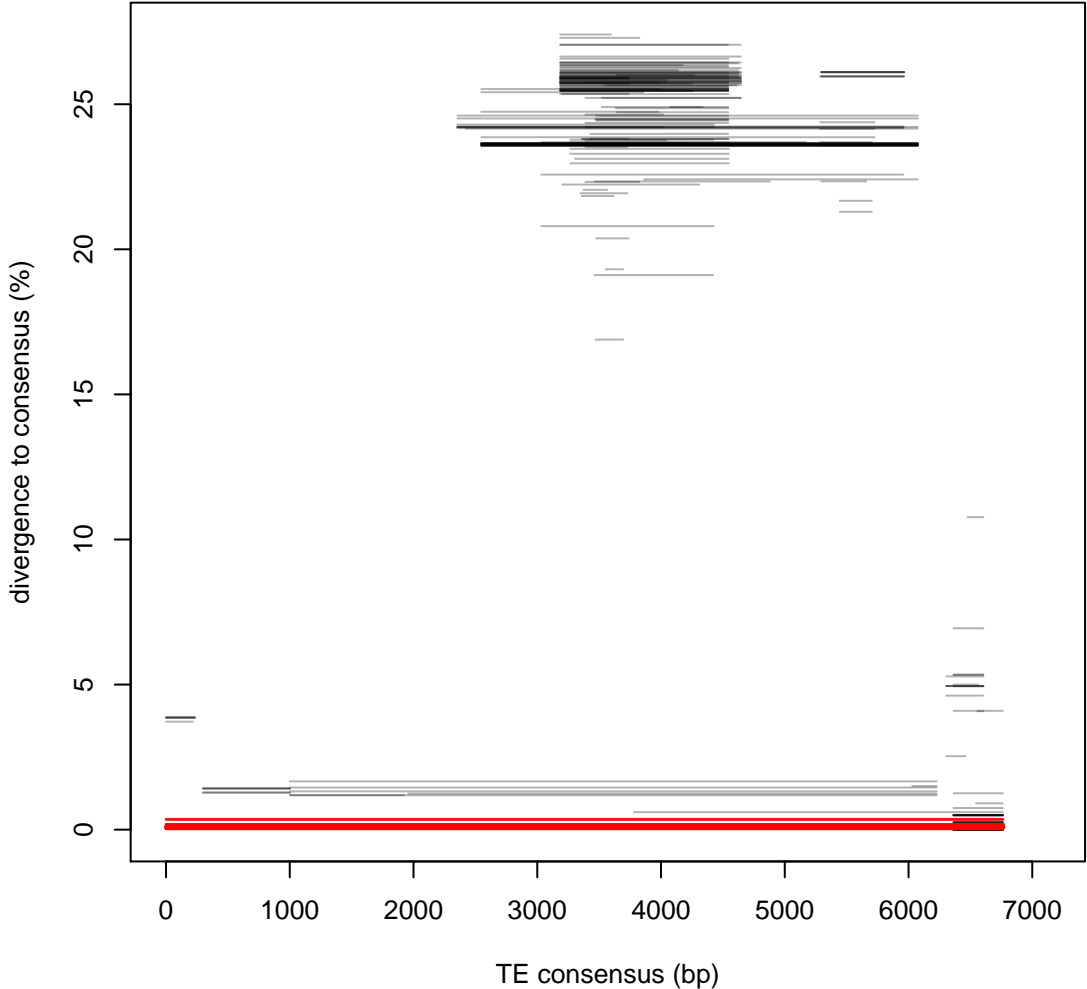
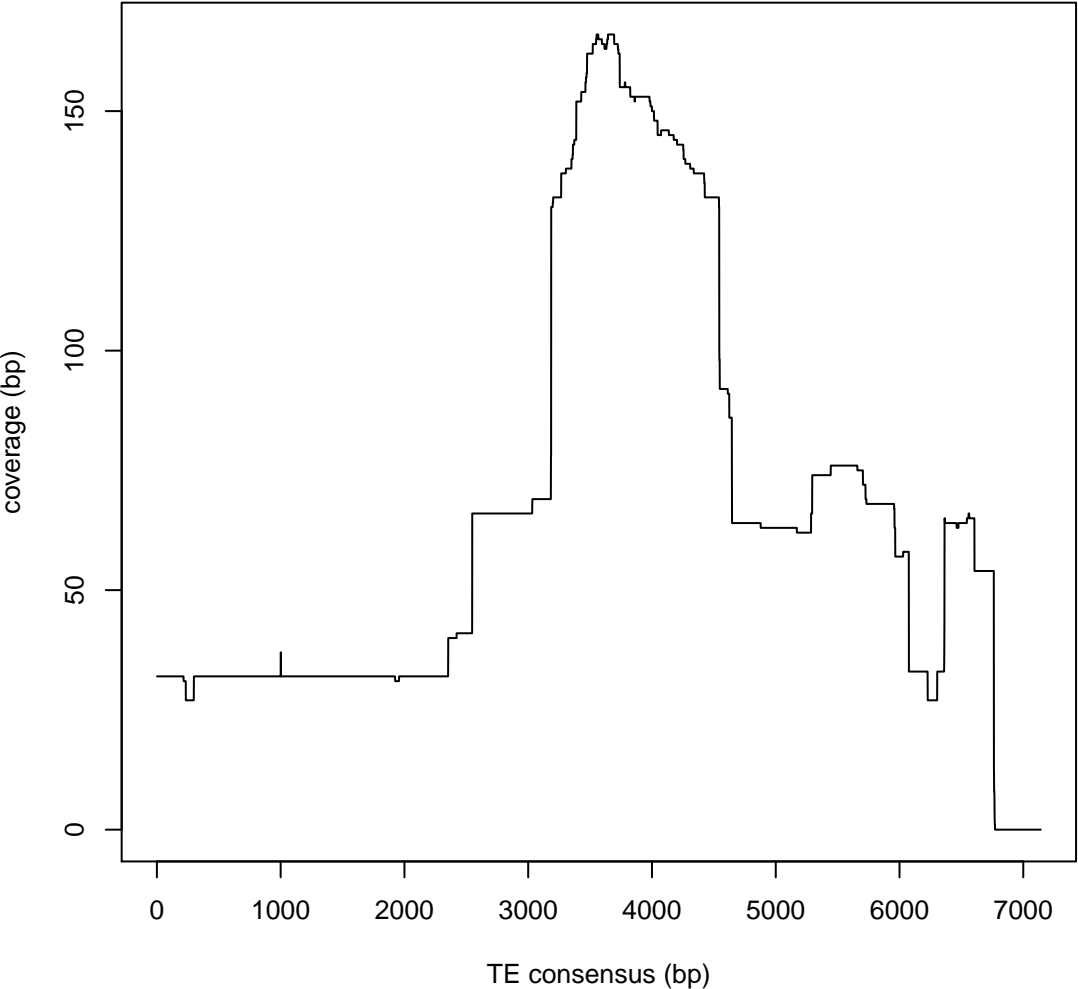


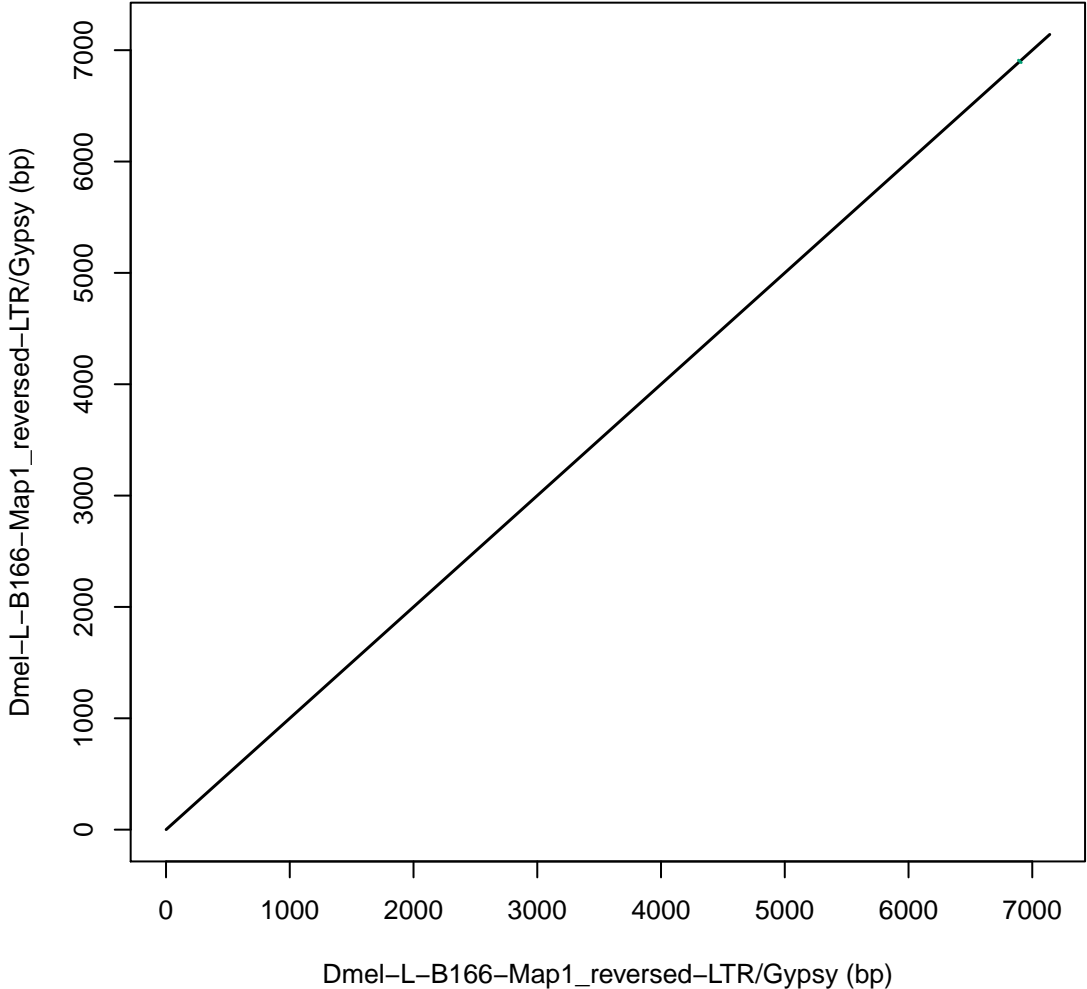
TE: Dmel-L-B166-Map1_reversed-LTR/Gypsy
consensus size: 7141bp; fragments: 239; full length: 25 (>=6426.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

