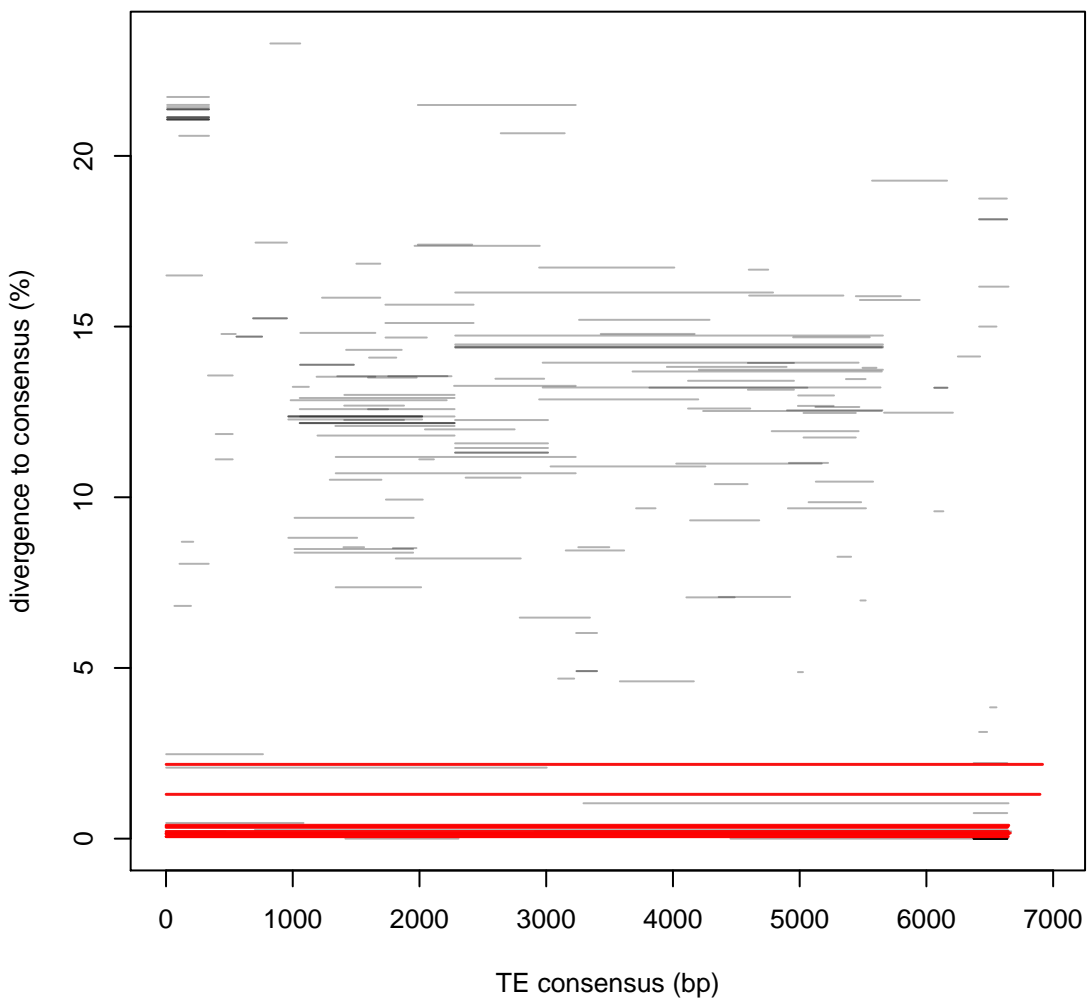
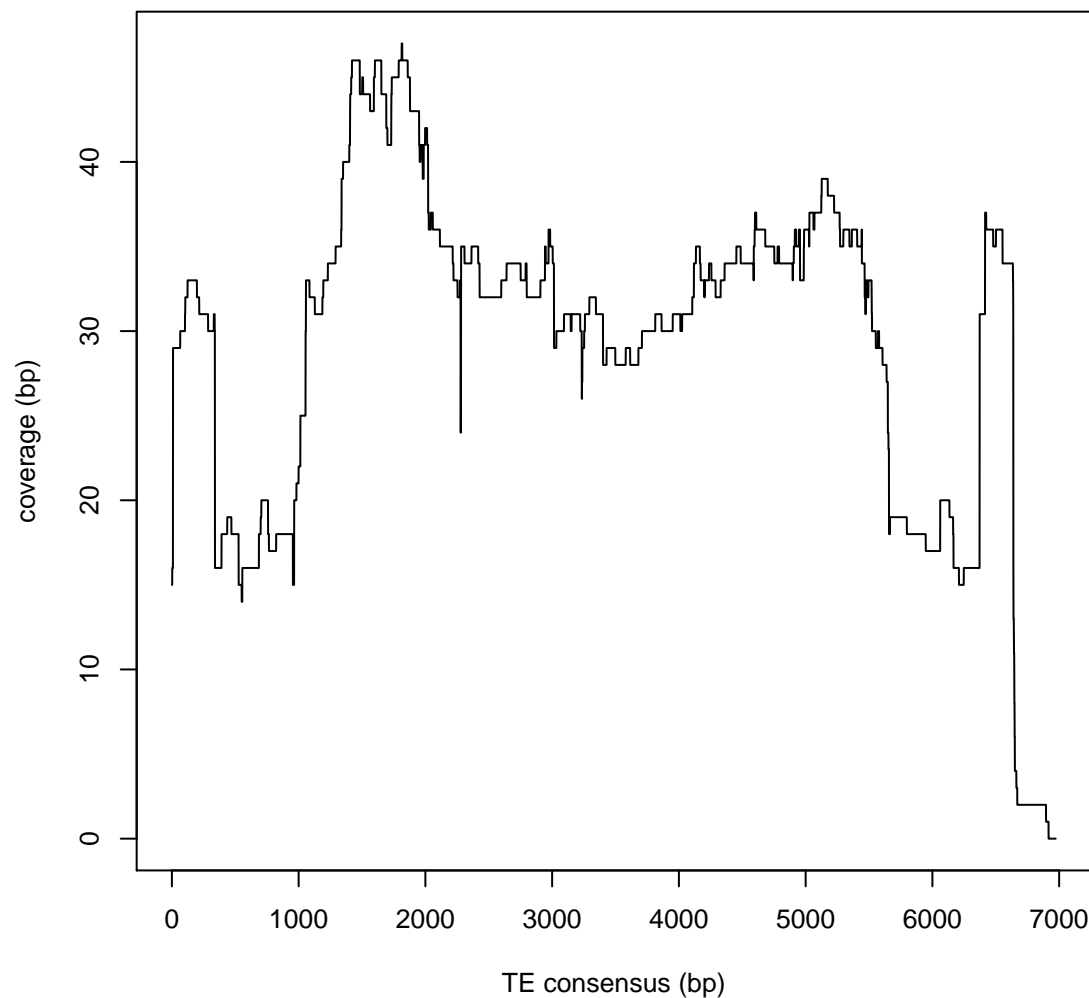


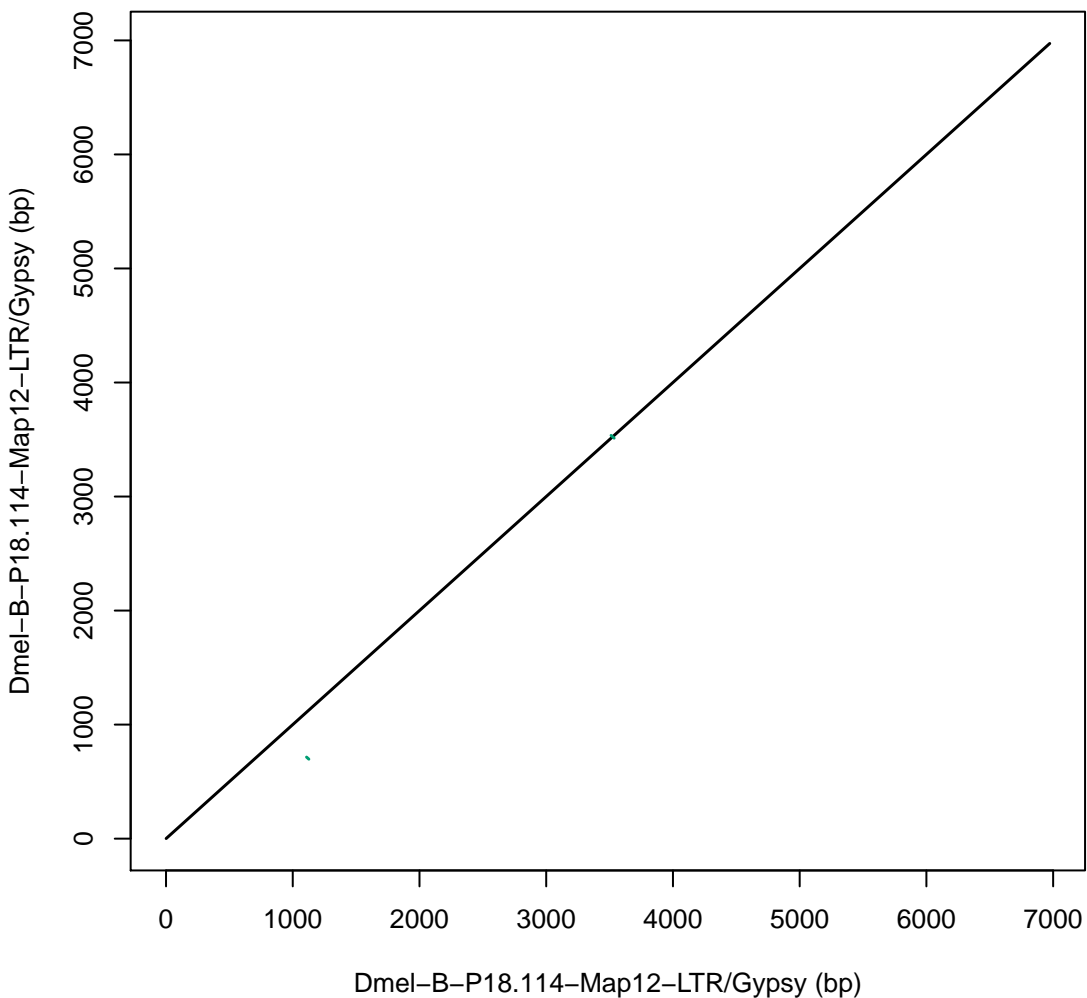
TE: Dmel-B-P18.114-Map12-LTR/Gypsy  
consensus size: 6973bp; fragments: 191; full length: 11 (>=6275.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

