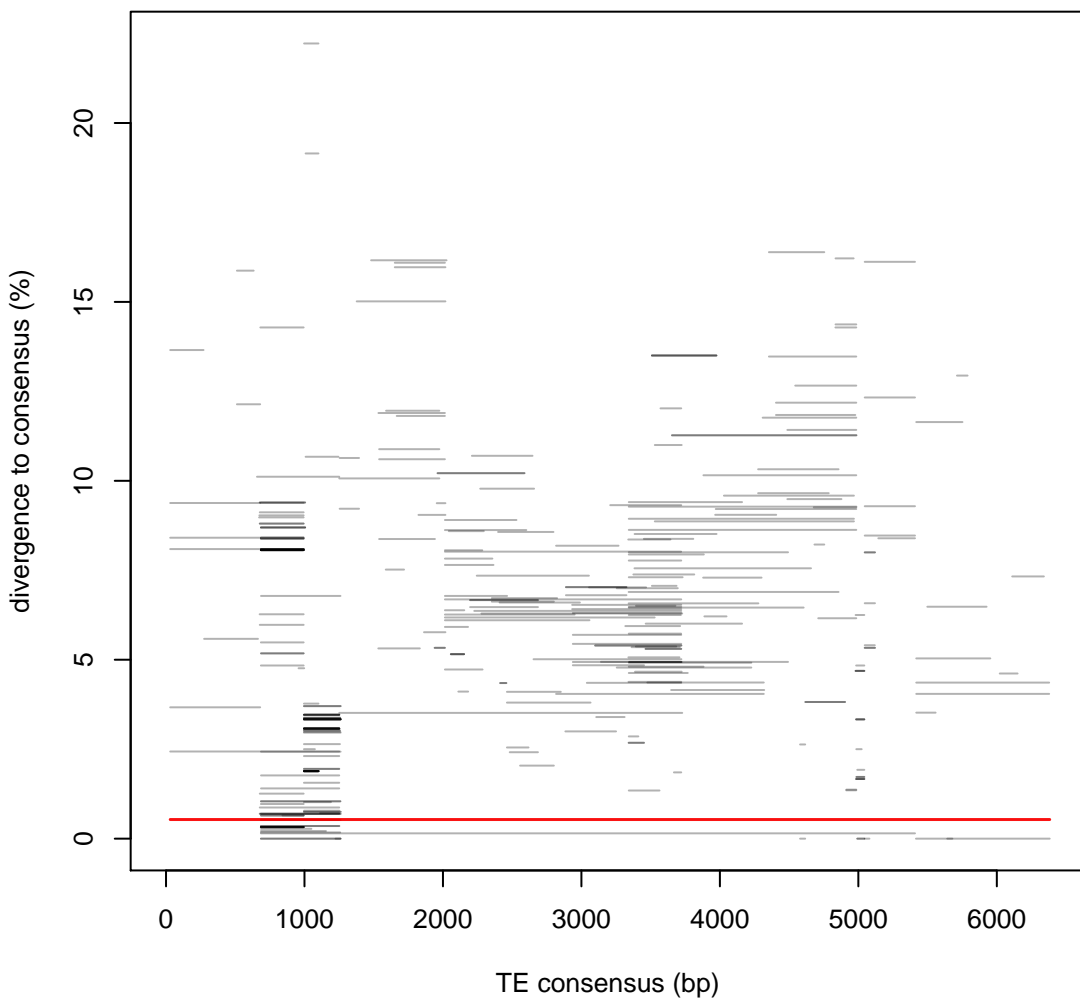
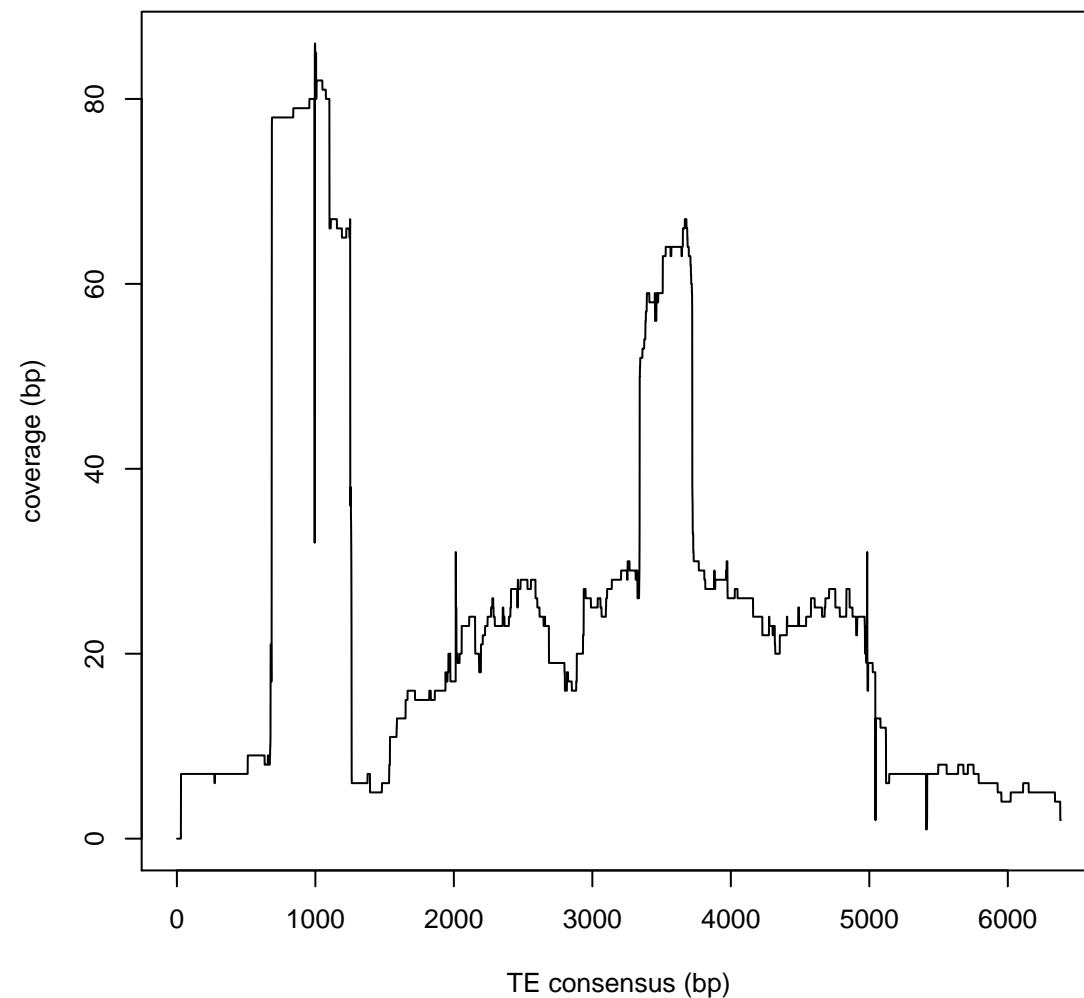


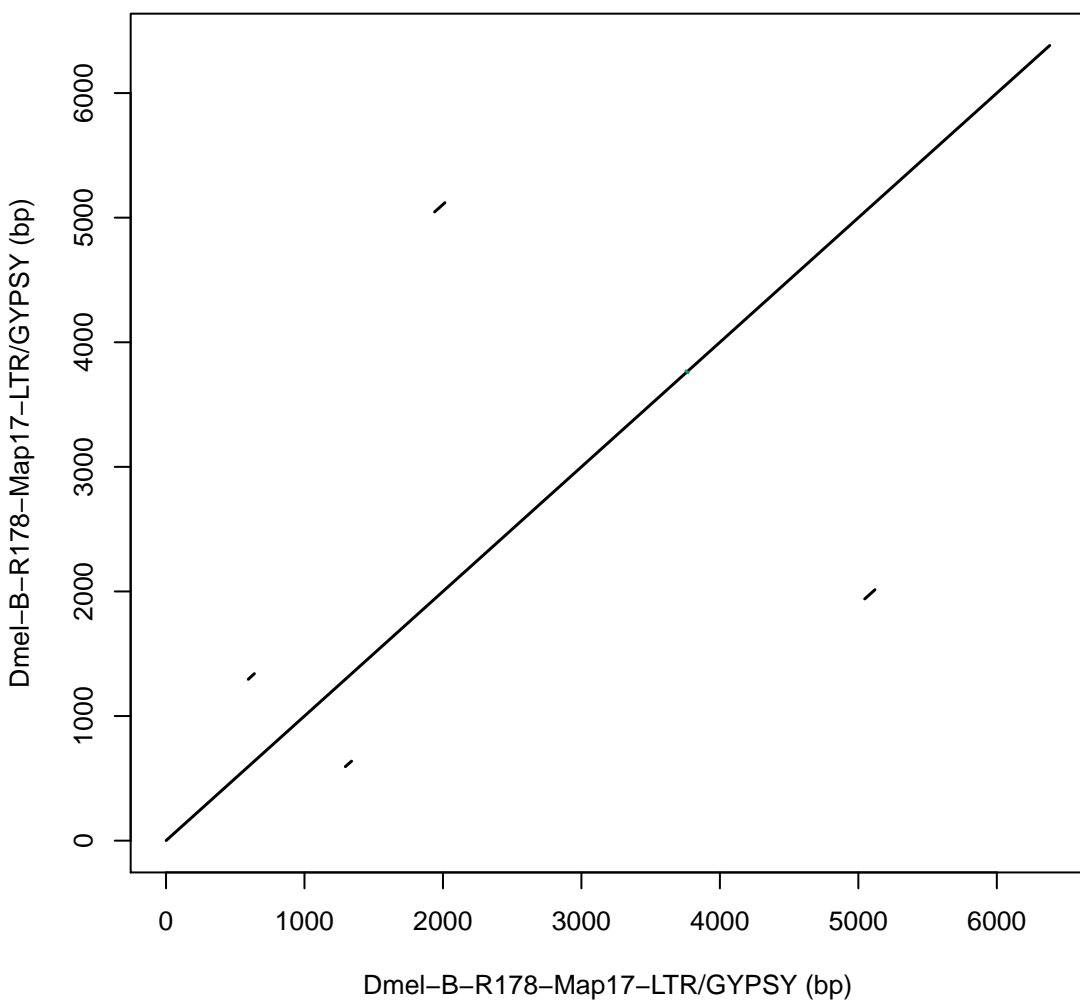
TE: Dmel-B-R178-Map17-LTR/GYPSY
consensus size: 6382bp; fragments: 348; full length: 1 (>=5743.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

