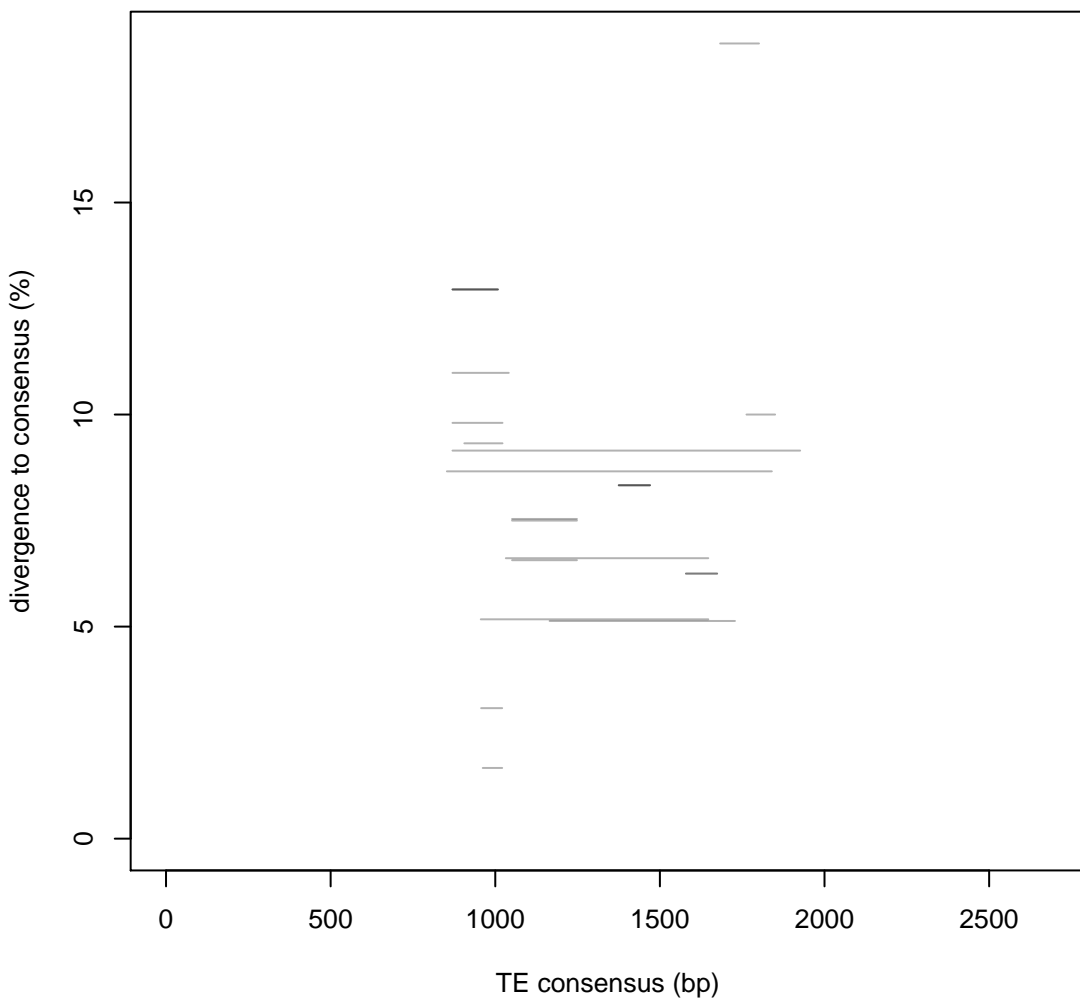
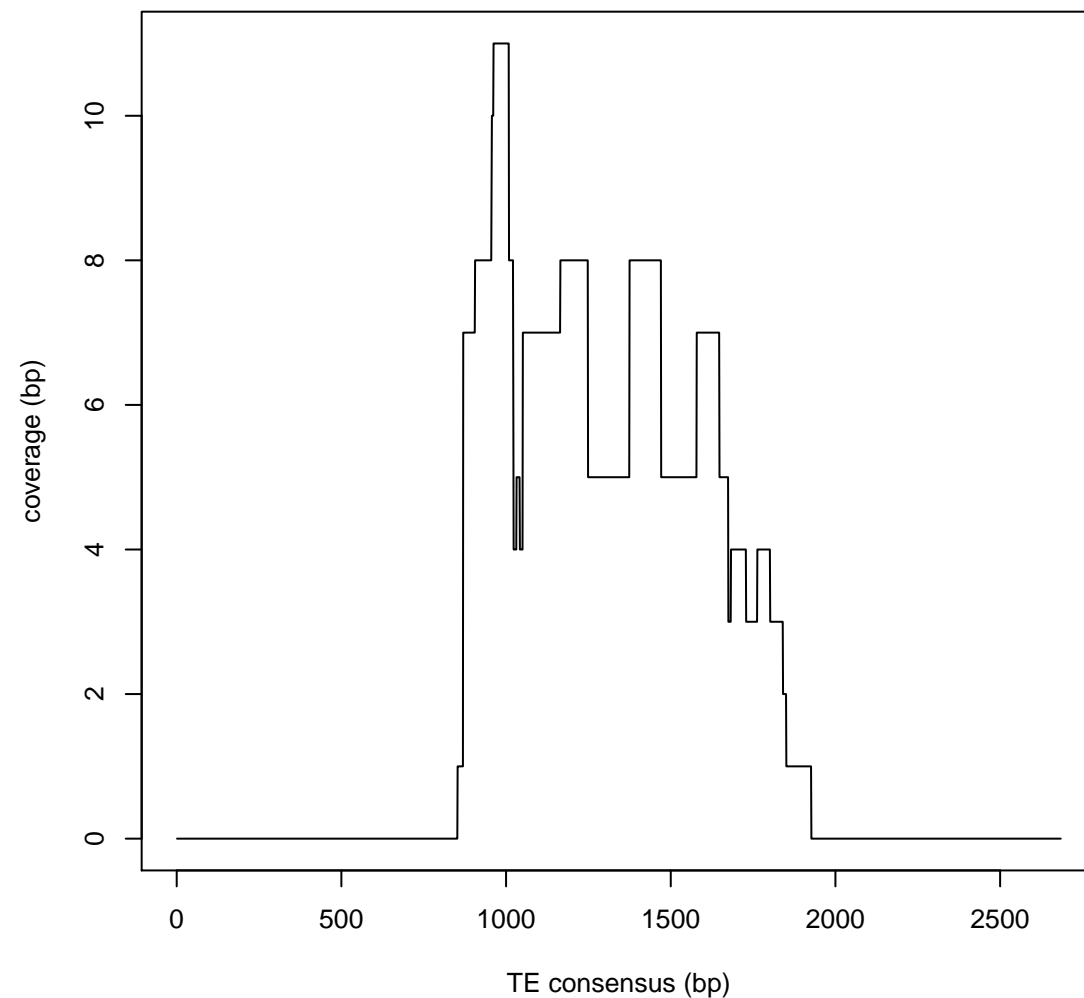


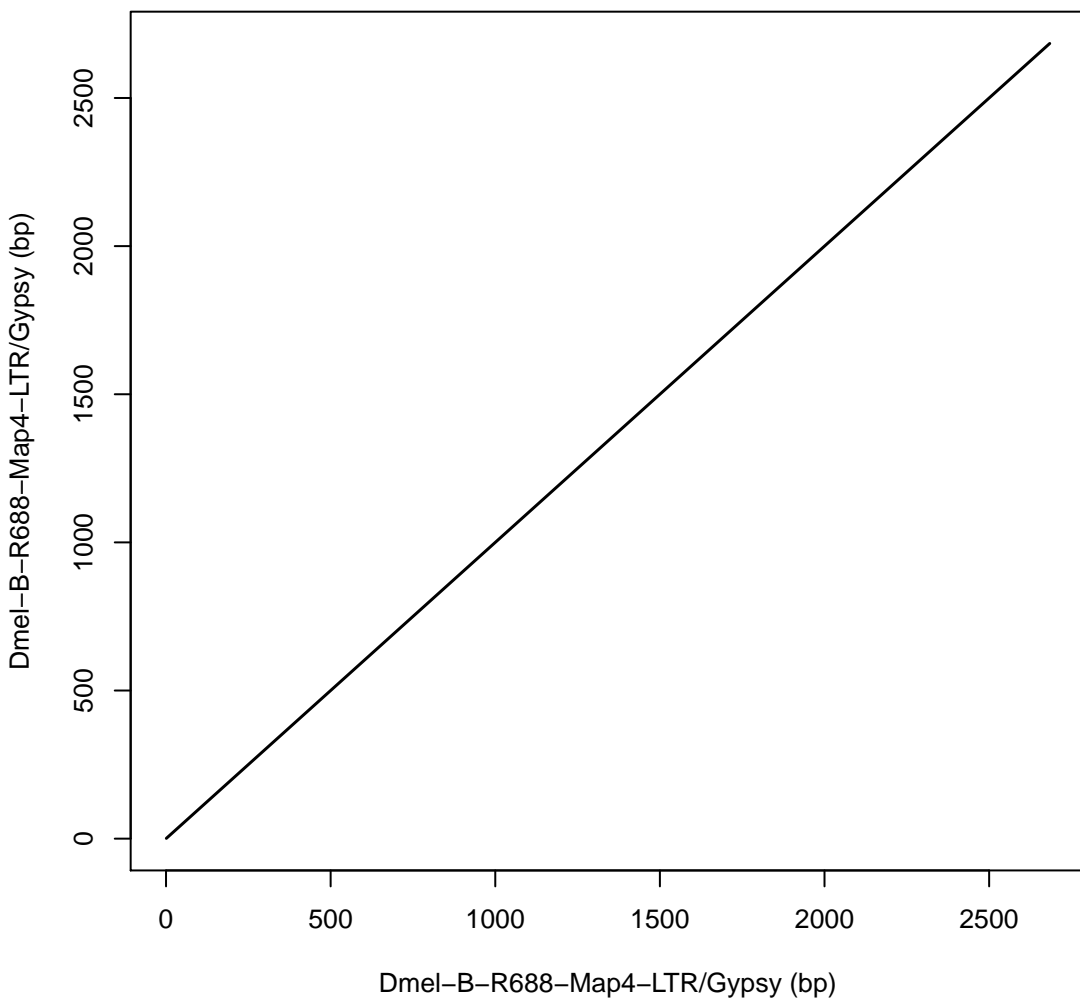
TE: Dmel-B-R688-Map4-LTR/Gypsy
consensus size: 2684bp; fragments: 23; full length: 0 (>=2415.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

