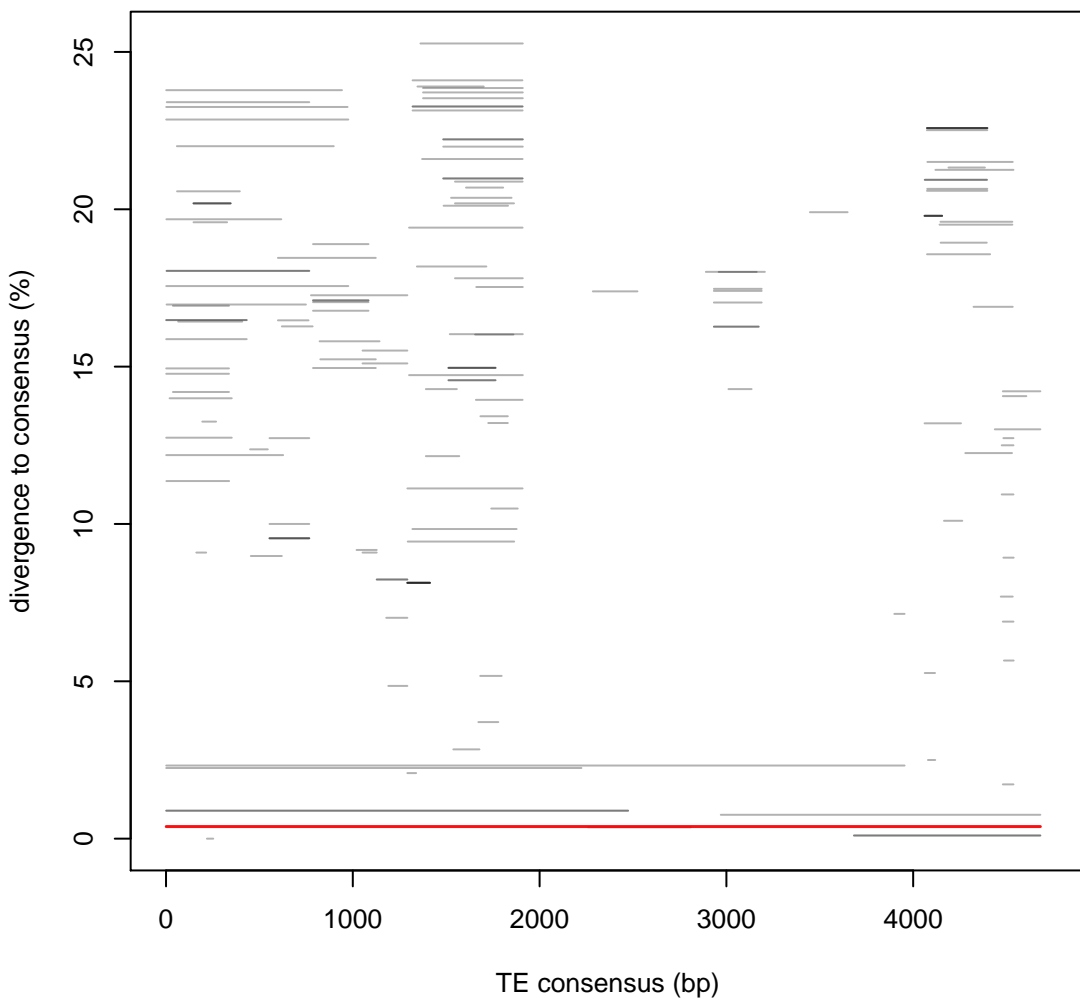
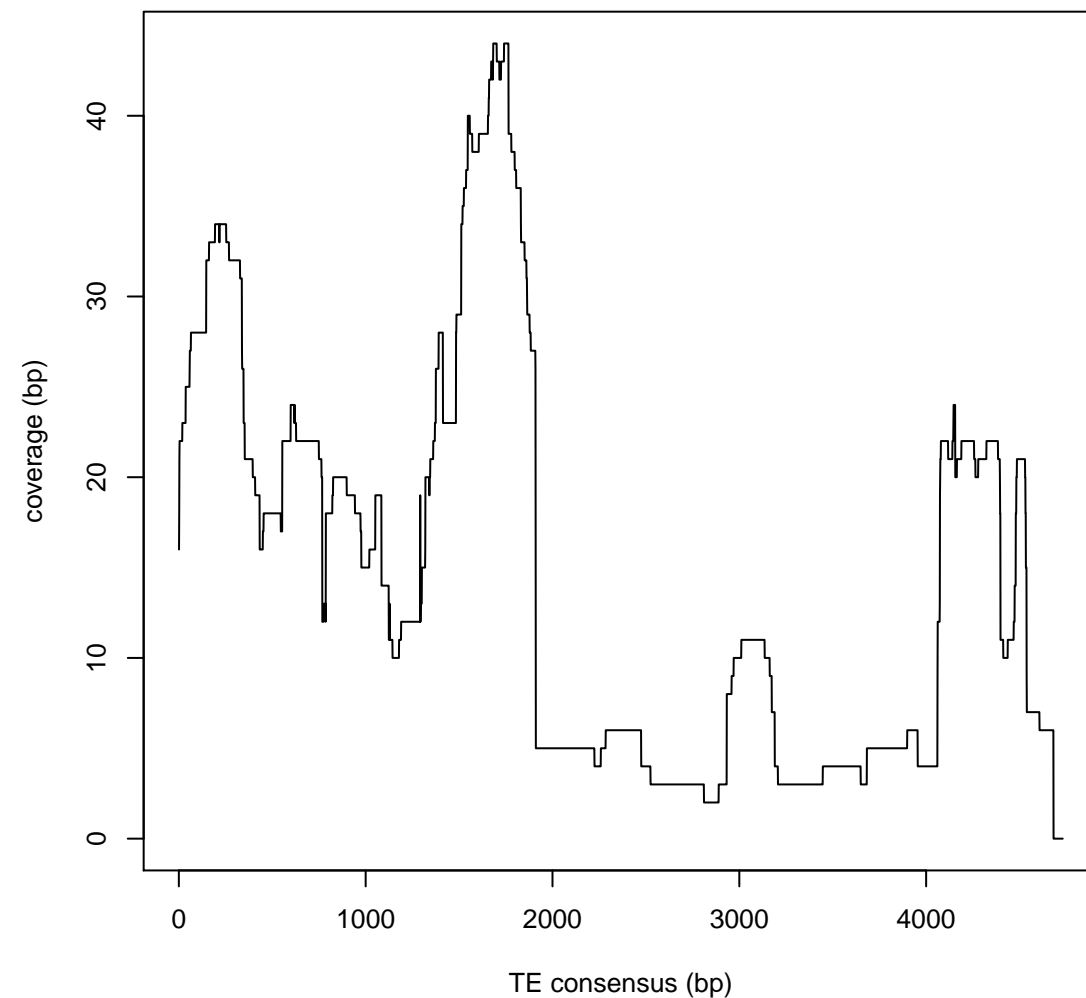


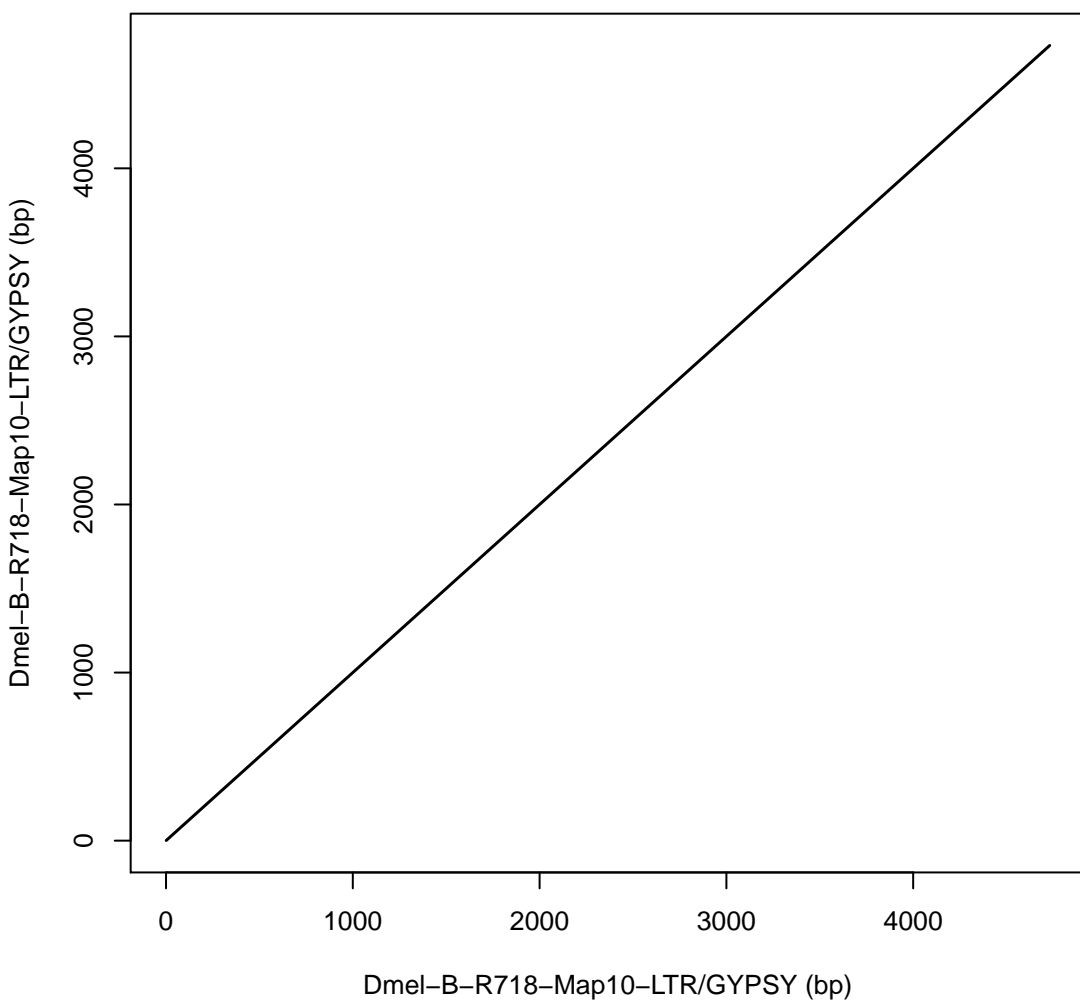
TE: Dmel-B-R718-Map10-LTR/GYPSY
consensus size: 4731bp; fragments: 164; full length: 1 (>=4257.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

