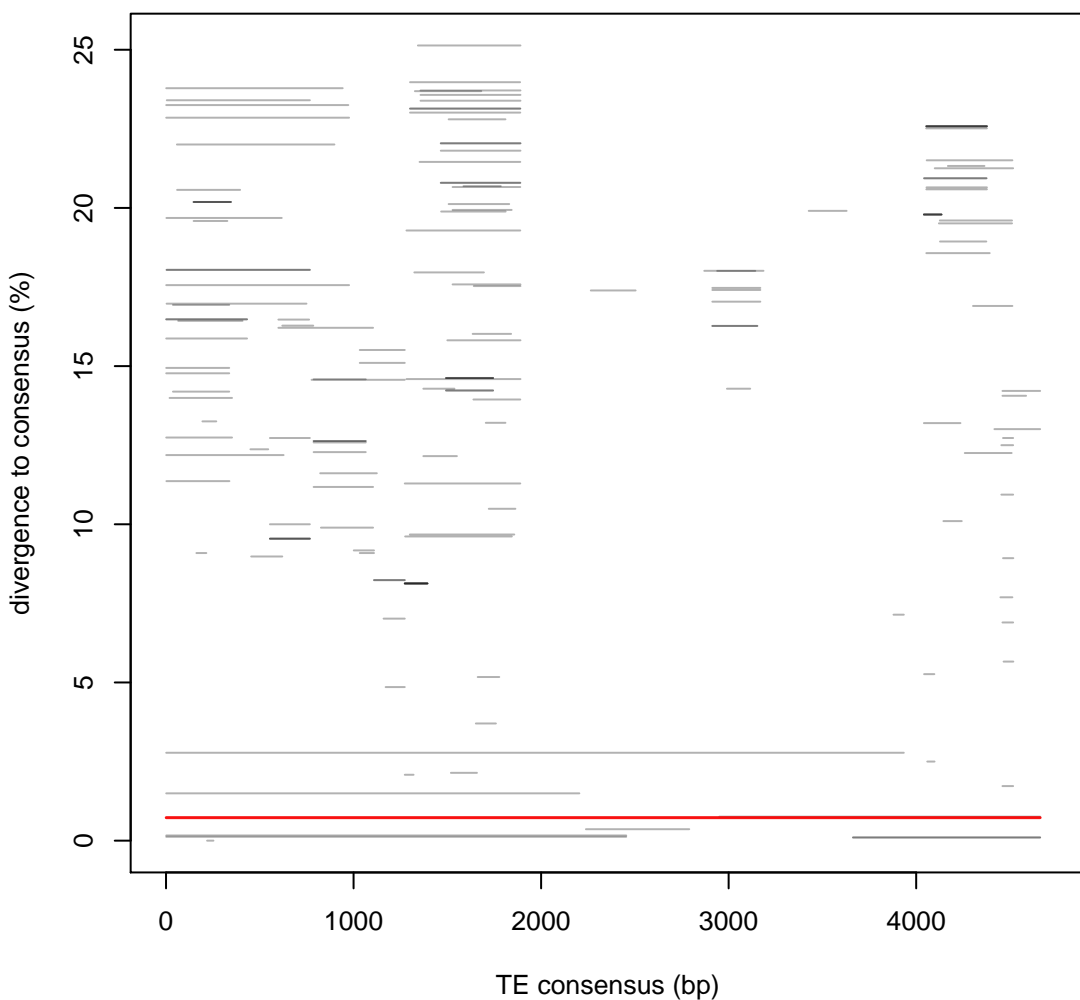
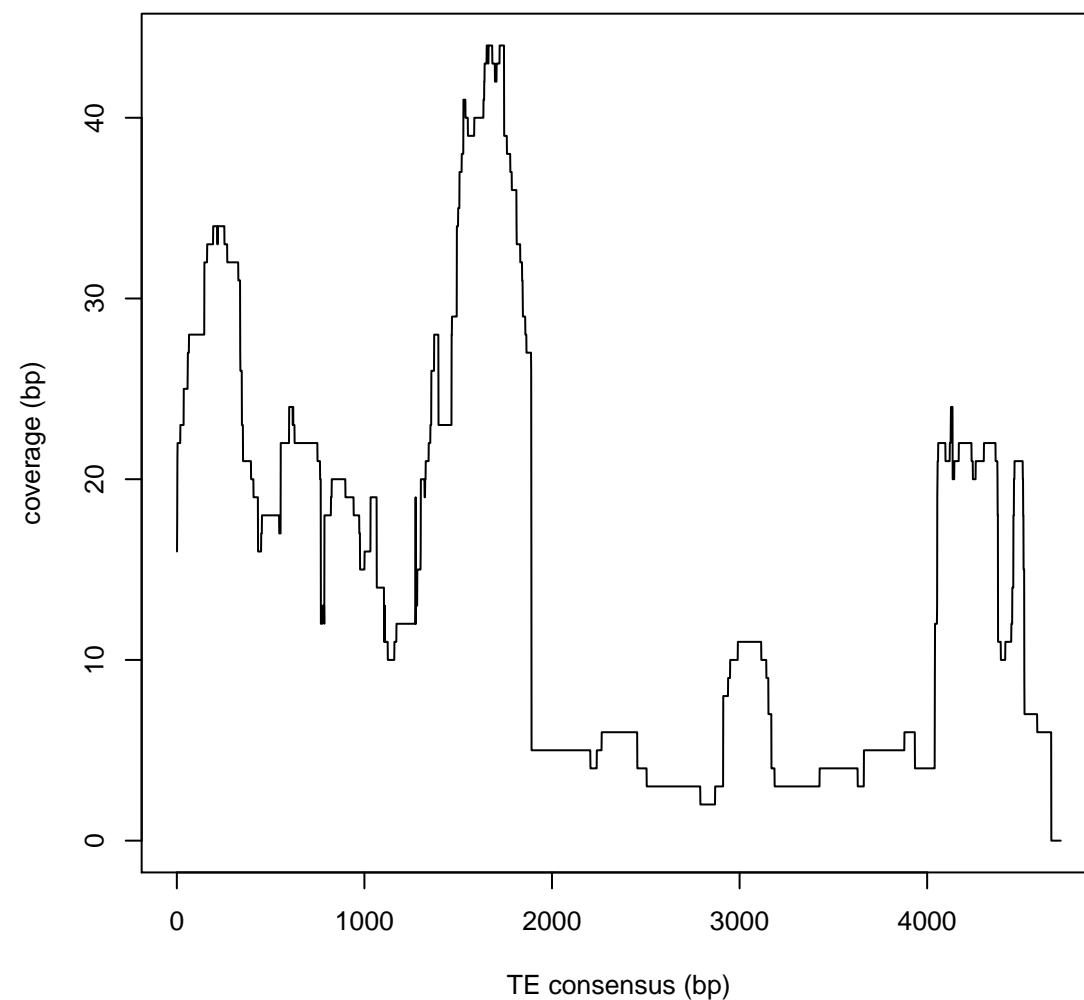


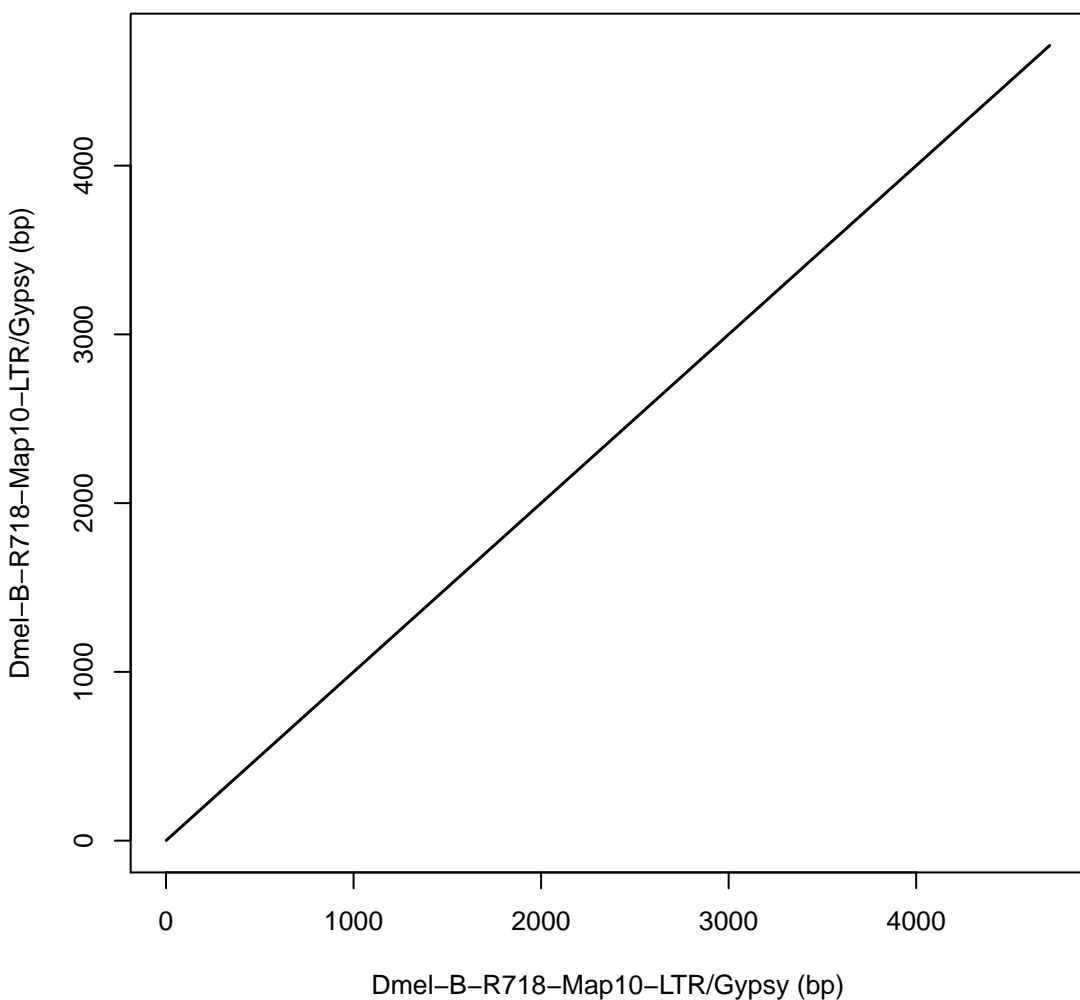
TE: Dmel-B-R718-Map10-LTR/Gypsy
consensus size: 4712bp; fragments: 164; full length: 1 (>=4240.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

