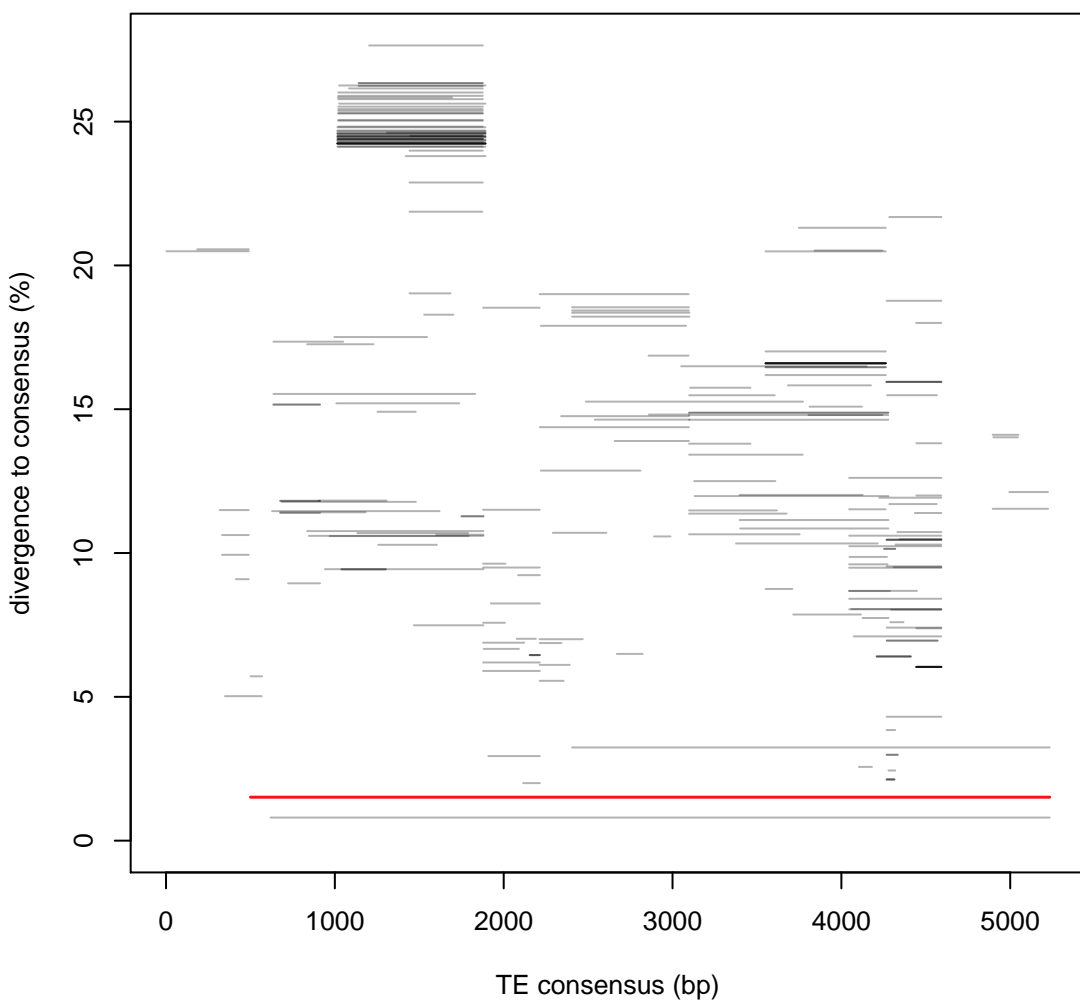
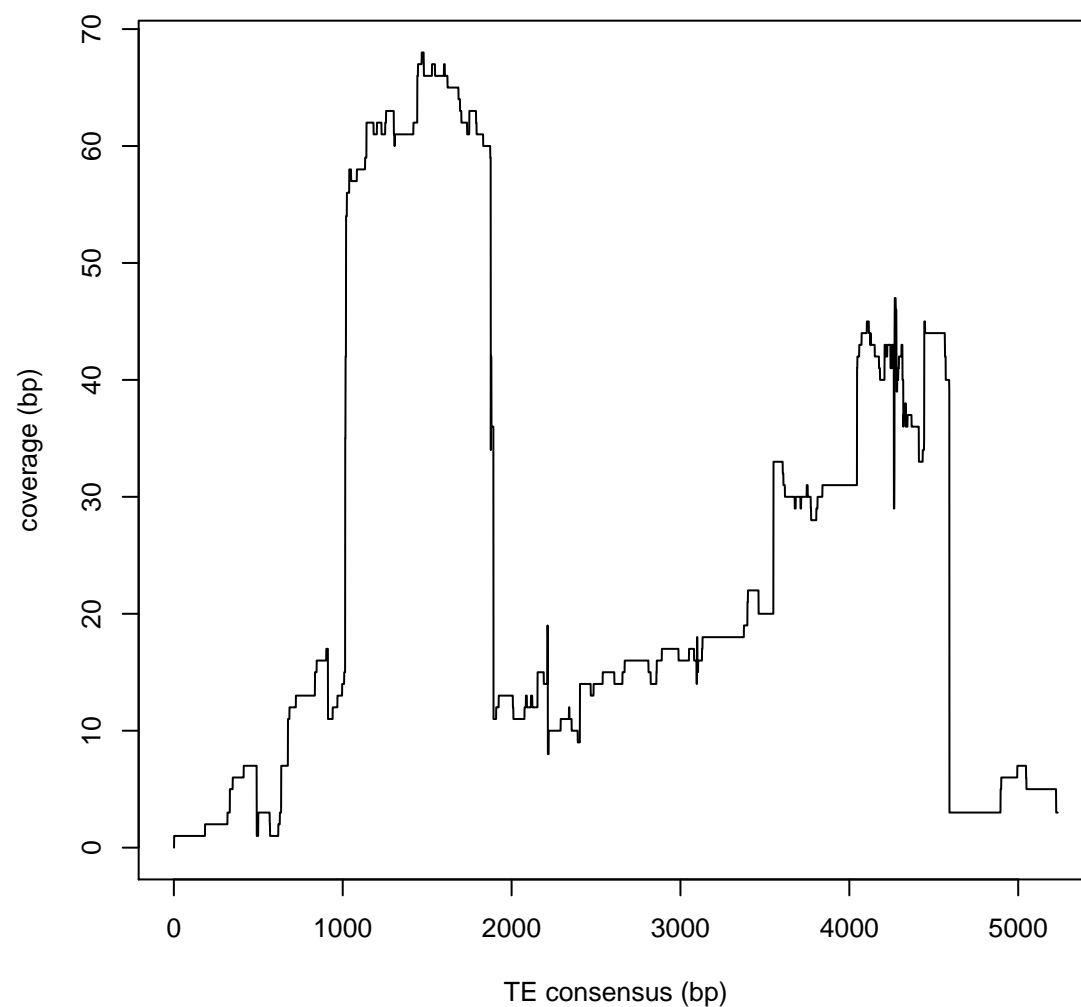


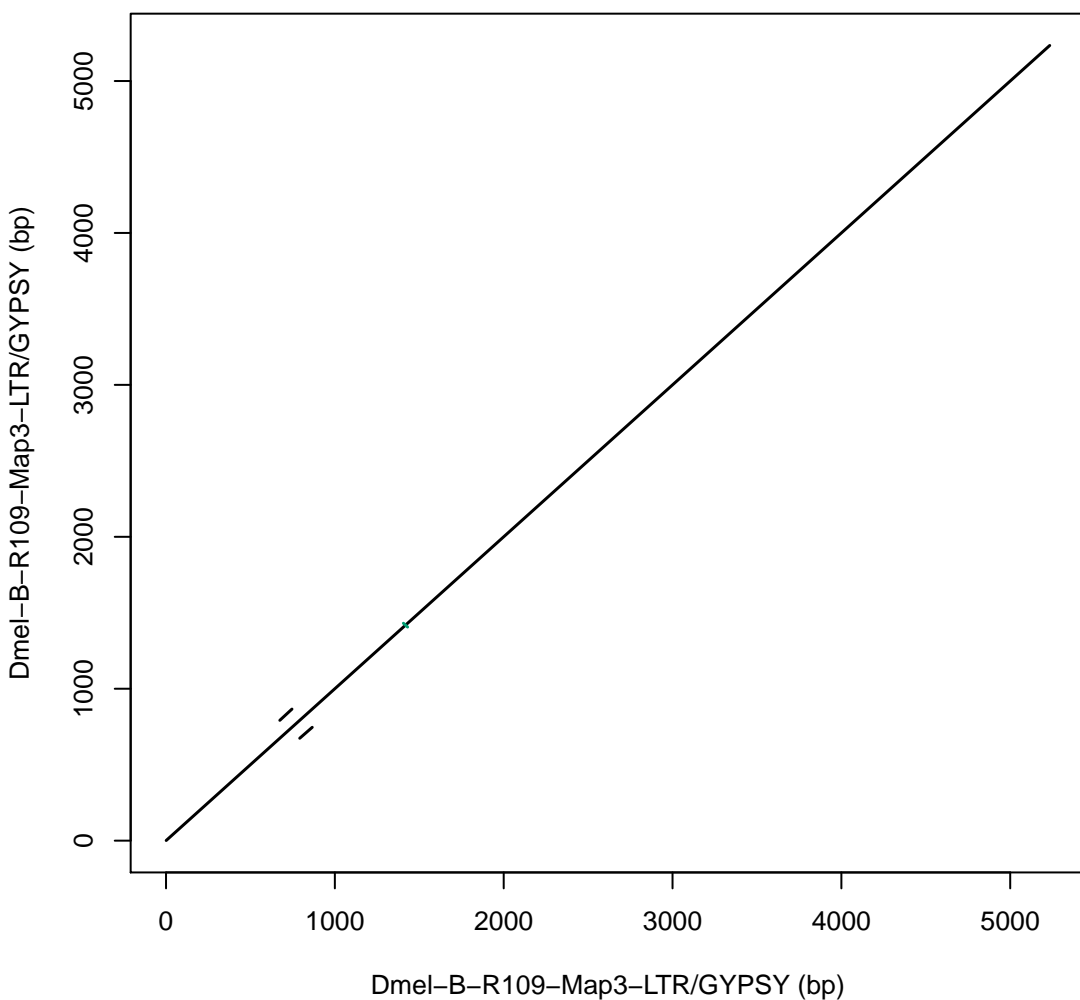
TE: Dmel-B-R109-Map3-LTR/GYPsy
consensus size: 5234bp; fragments: 233; full length: 1 (>=4710.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

