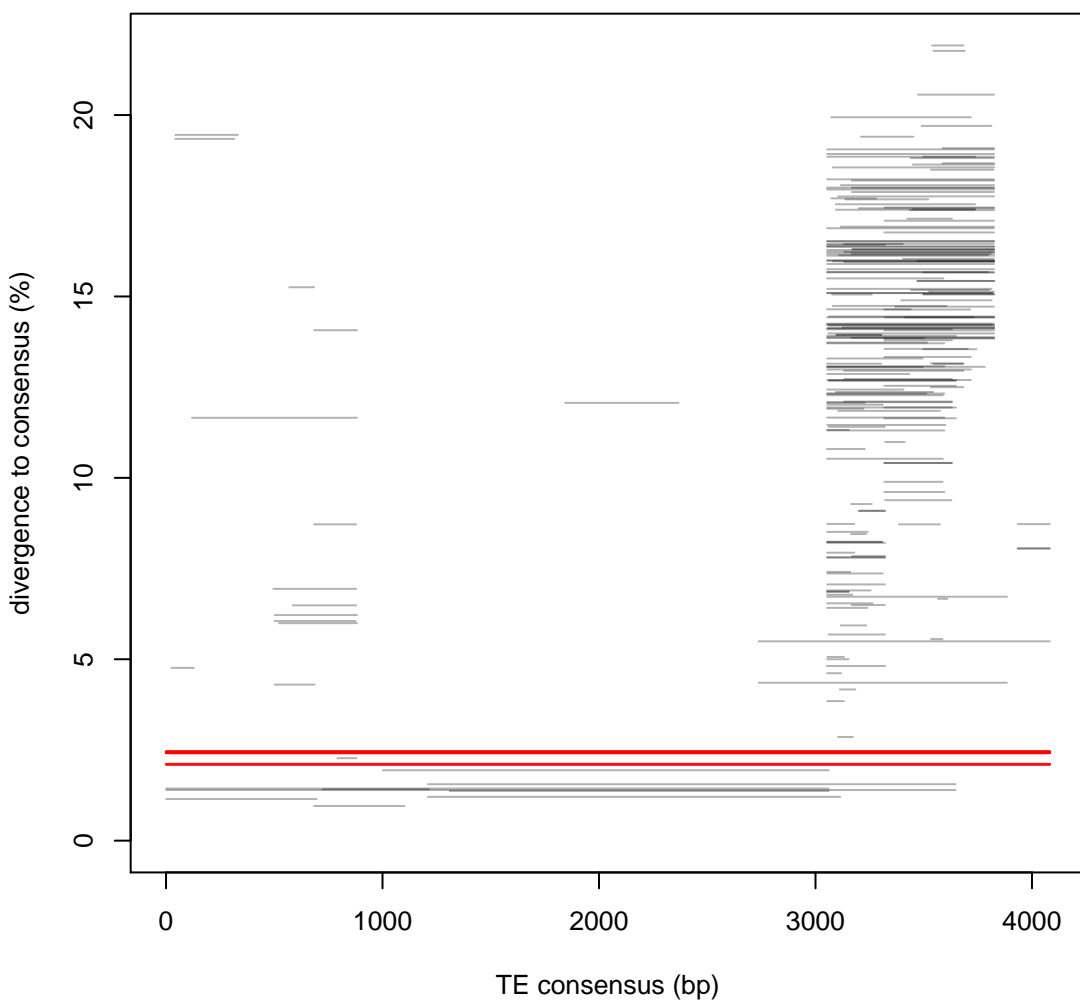
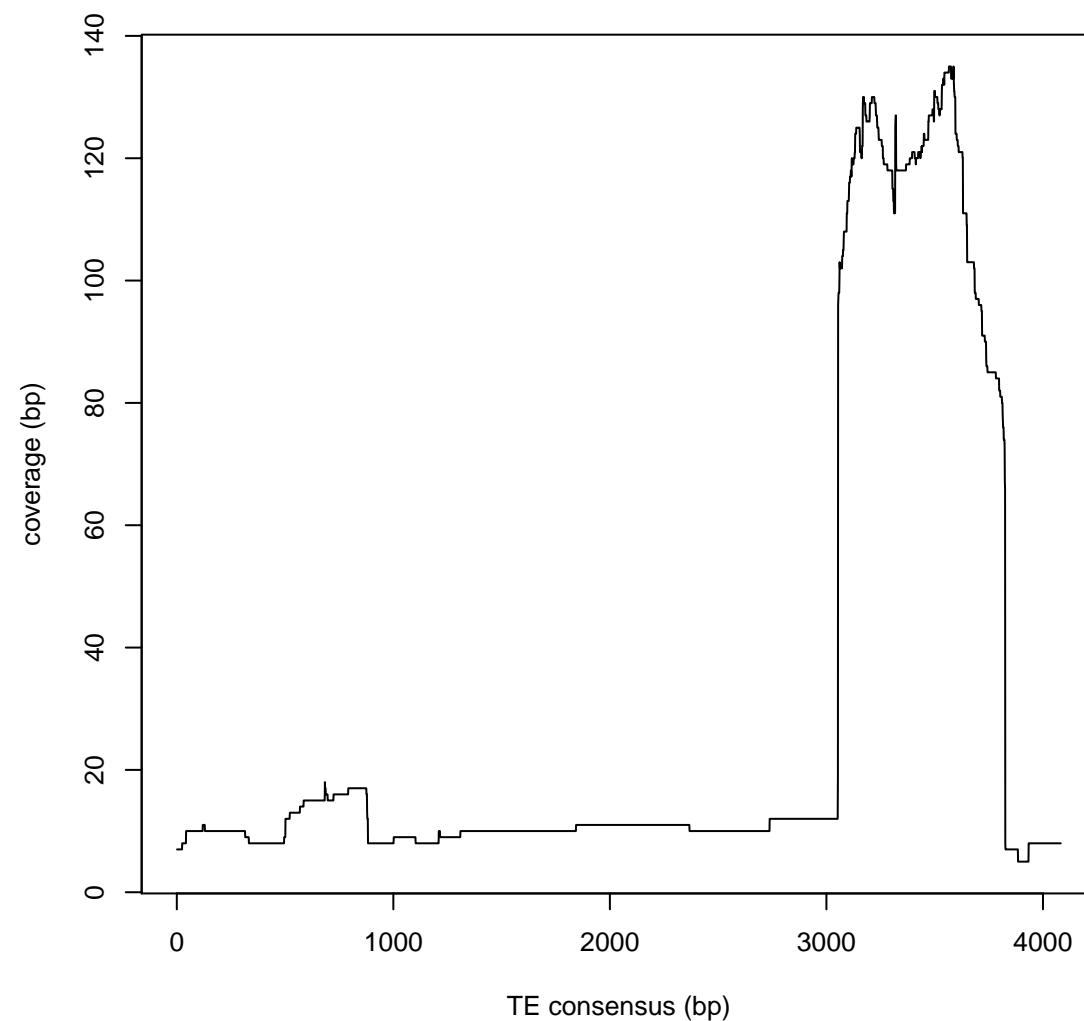


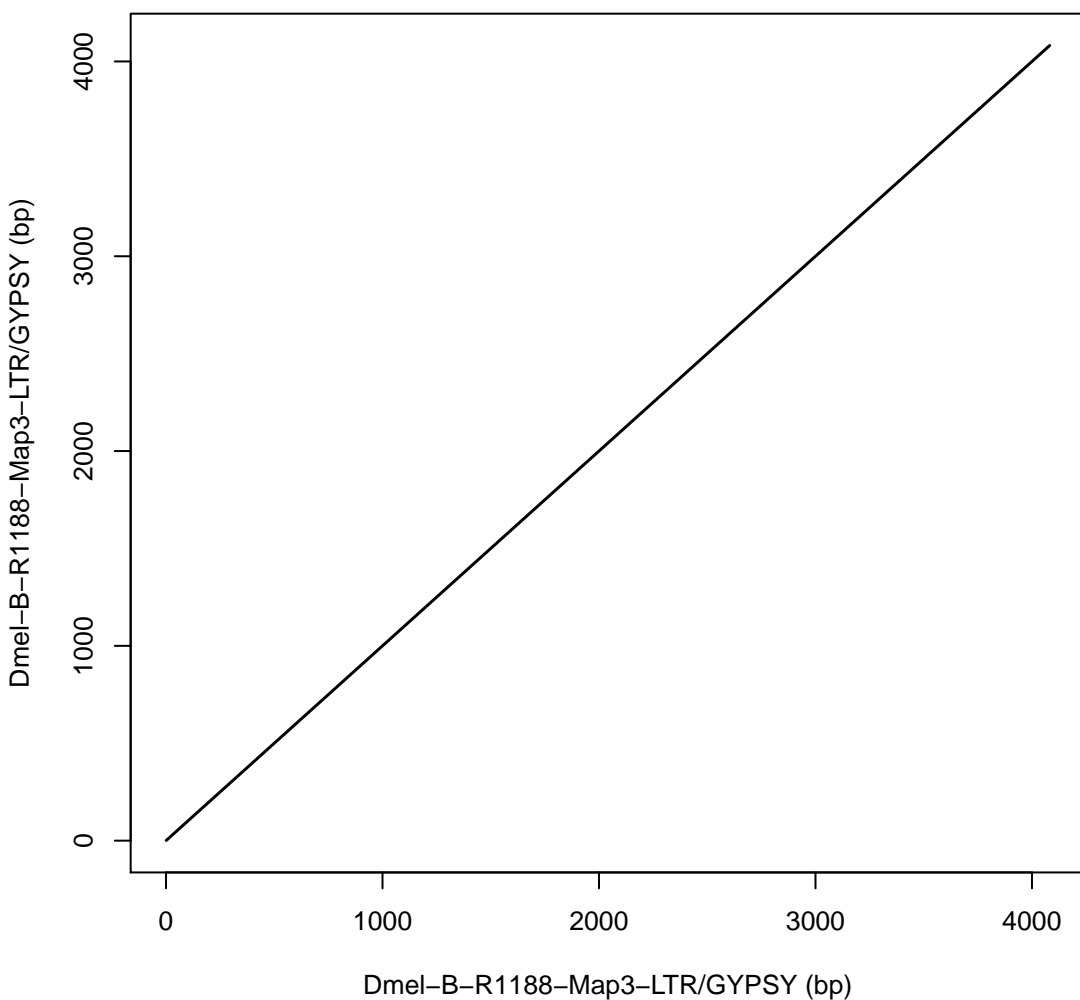
TE: Dmel-B-R1188-Map3-LTR/Gypsy
consensus size: 4082bp; fragments: 219; full length: 4 (>=3673.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

