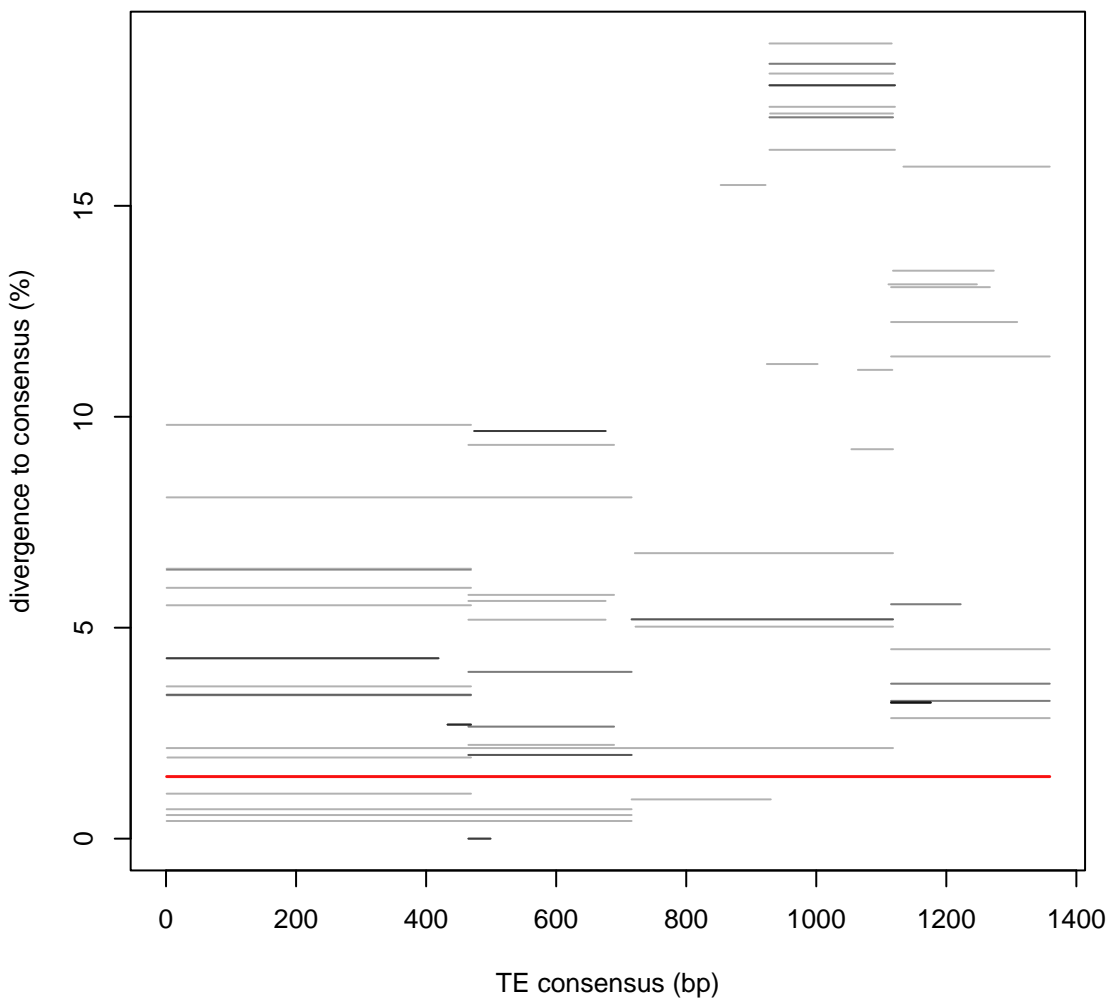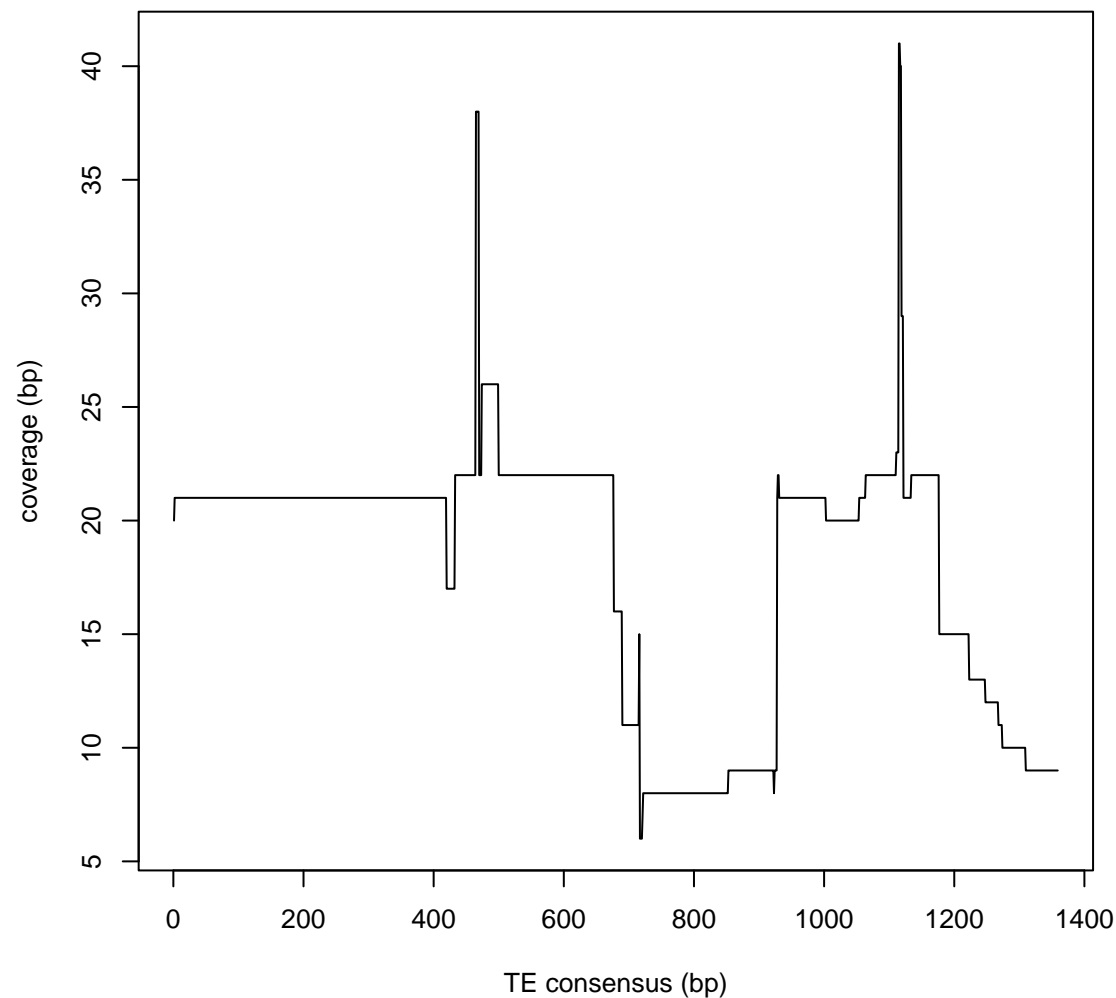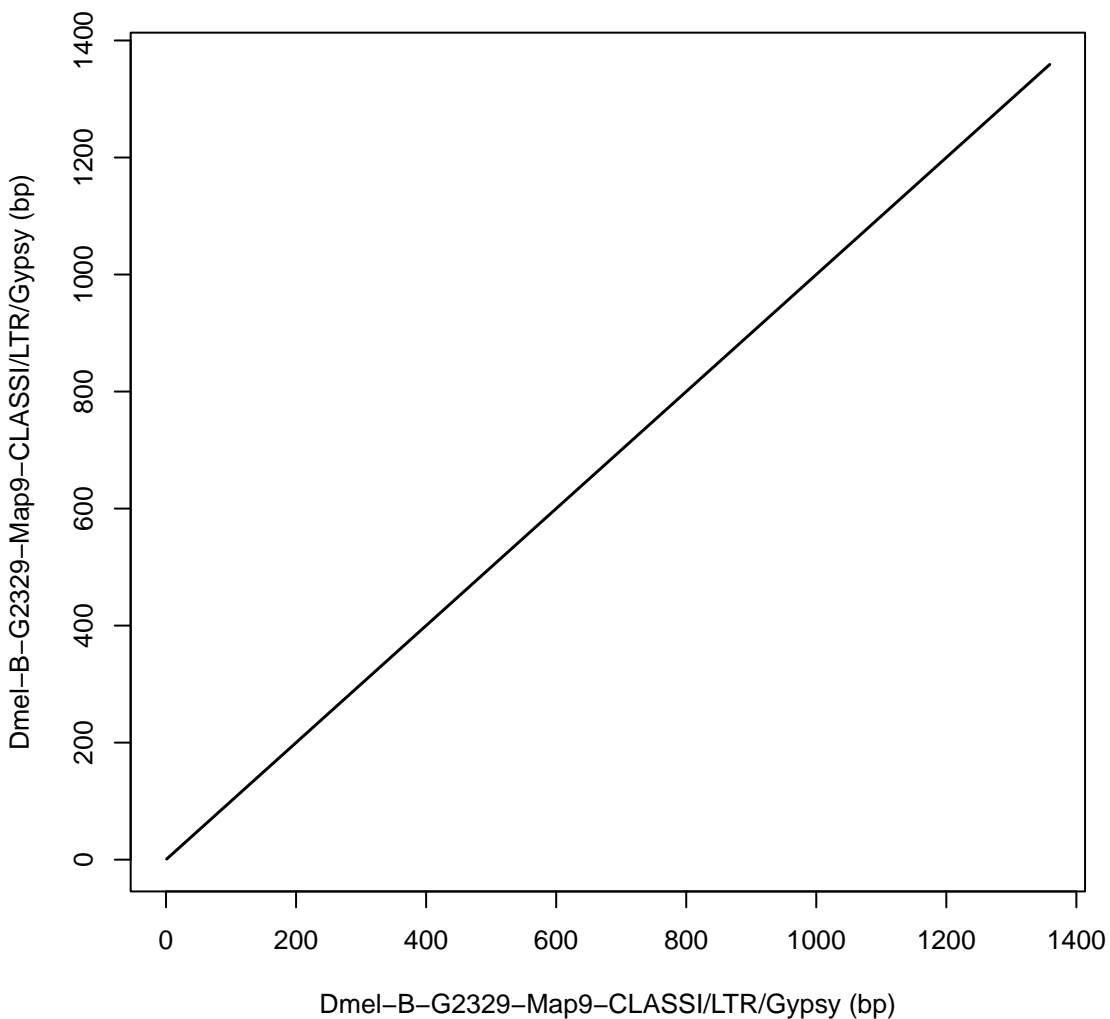**TE: Dmel−B−G2329−Map9−CLASSI/LTR/Gypsy**
consensus size: 1359bp; fragments: 90; full length: 1 (>=1223.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits