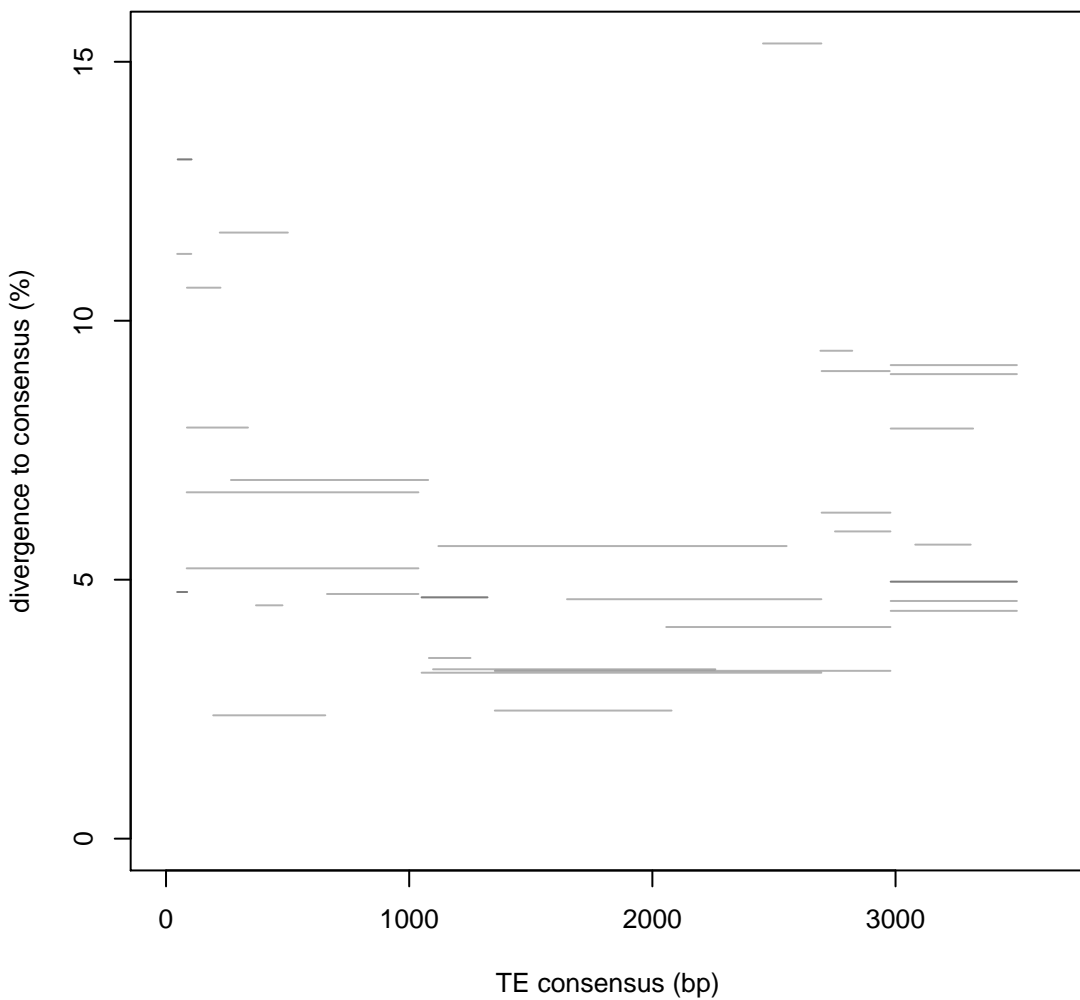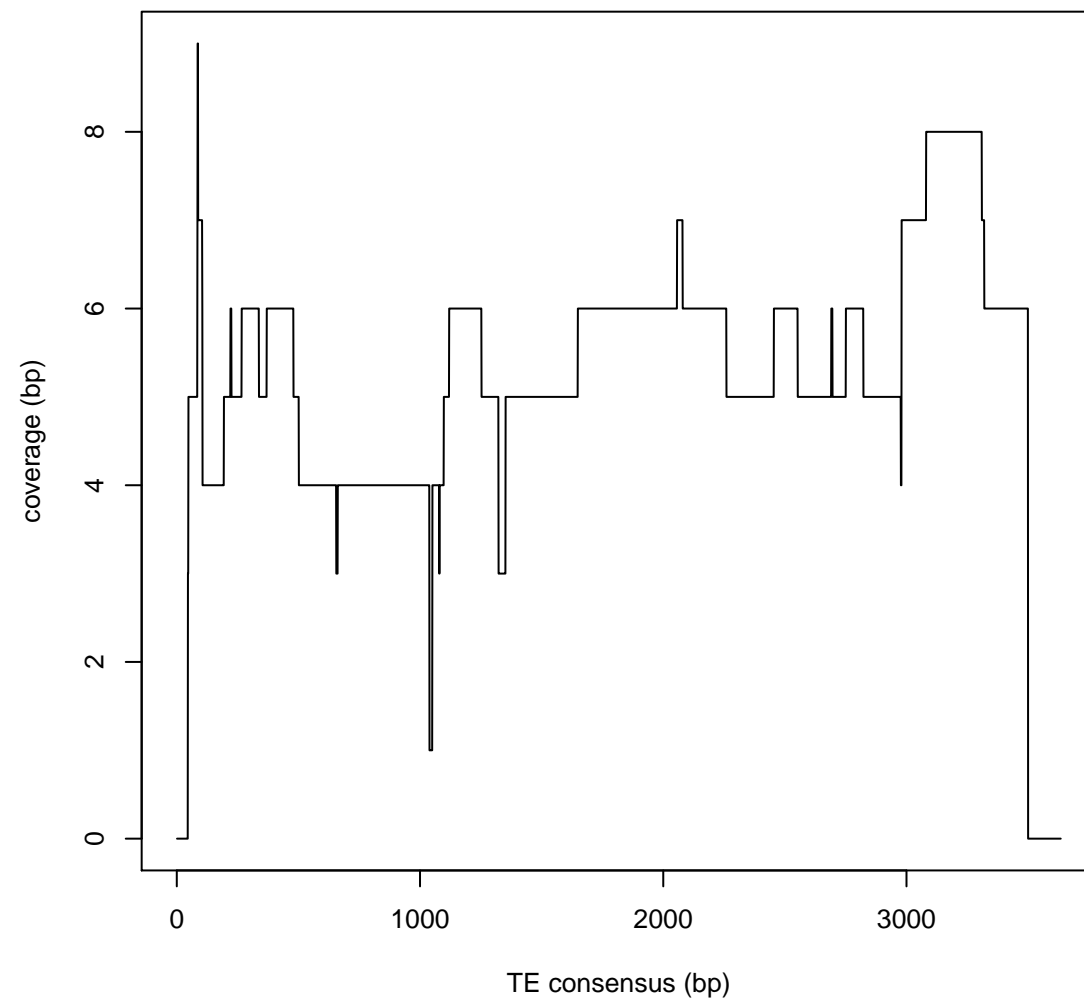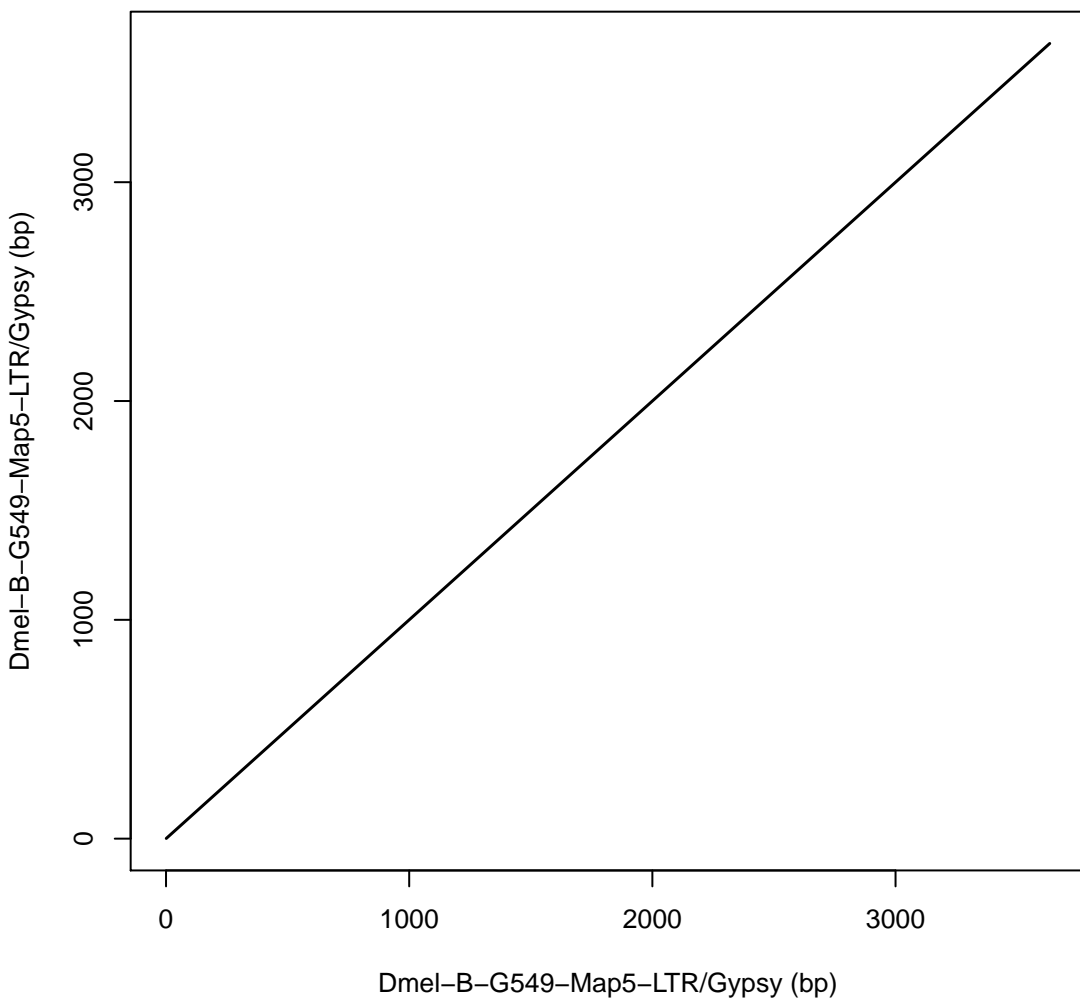**TE: Dmel−B−G549−Map5−LTR/Gypsy**
consensus size: 3634bp; fragments: 37; full length: 0 (>=3270.6bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits