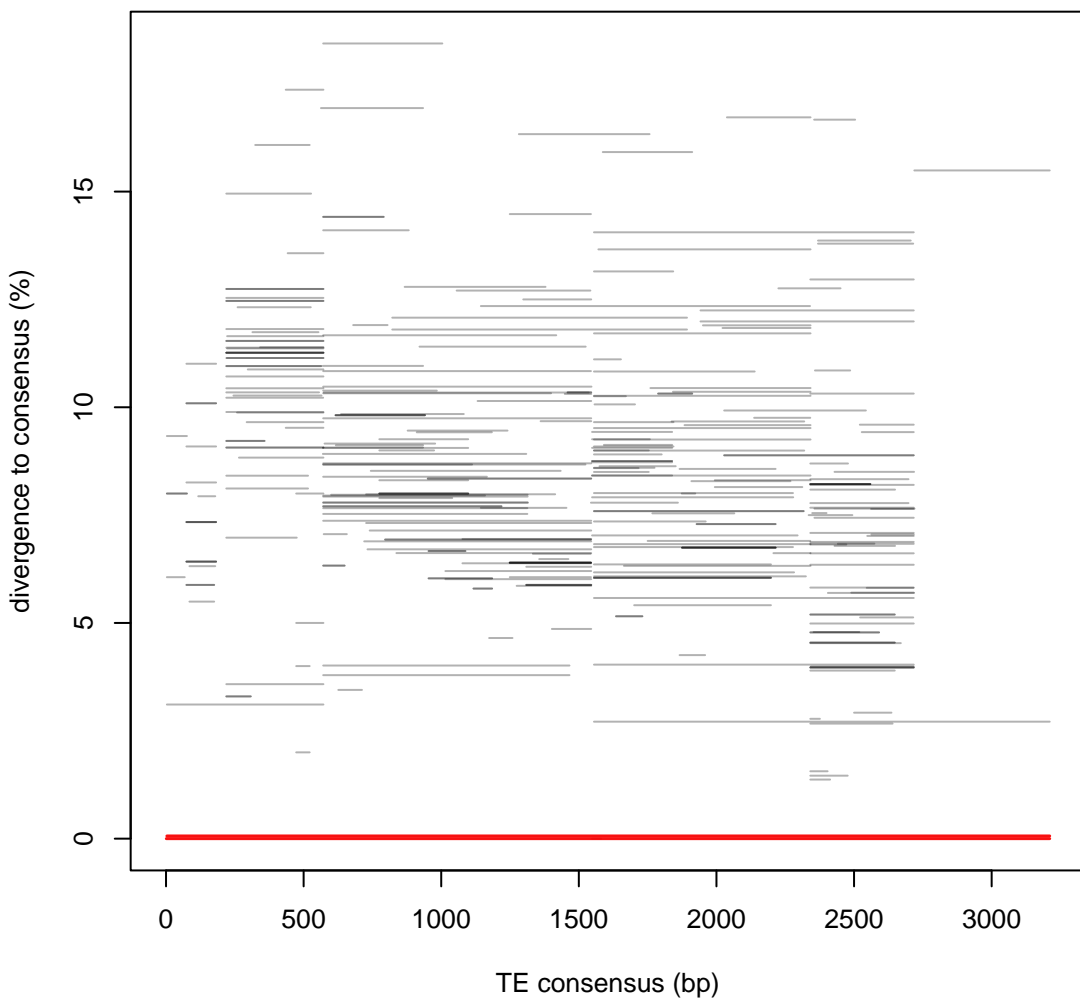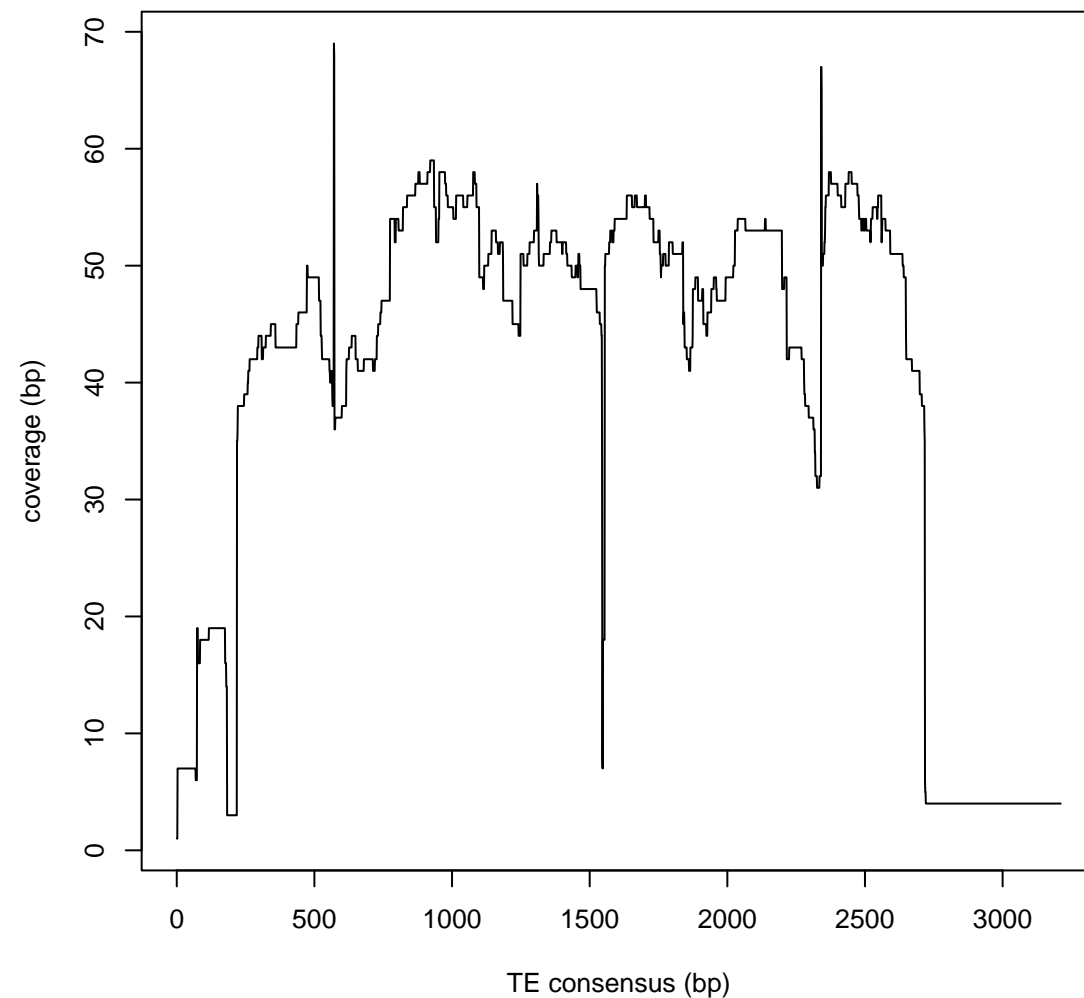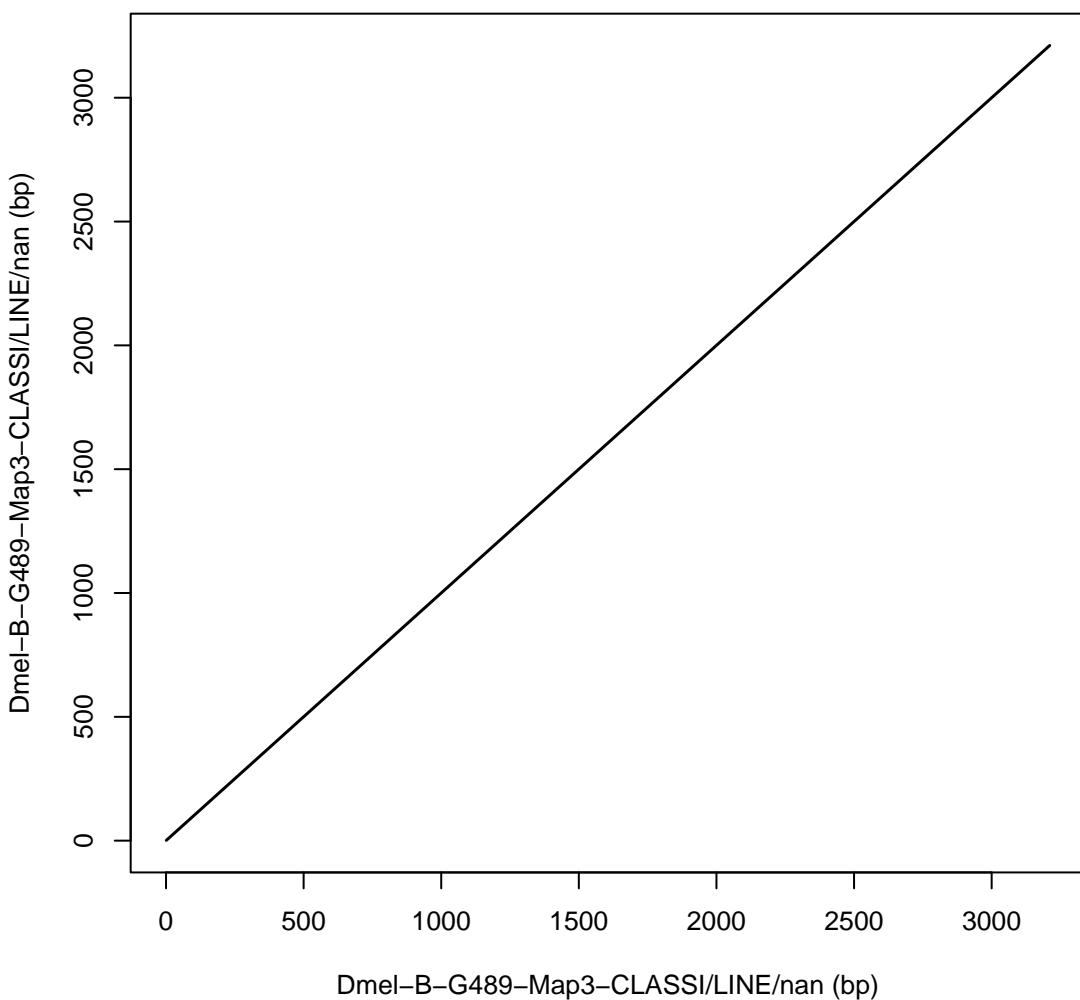**TE: Dmel−B−G489−Map3−CLASSI/LINE/nan**
consensus size: 3211bp; fragments: 321; full length: 2 (>=2889.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Copia−23_BG LTR/Copia

Jockey−4_DEu_pol LINE/I−Jockey

DOC5_DM_pol LINE/I−Jockey

Jockey−11B_DBp_pol LINE/I−Jockey