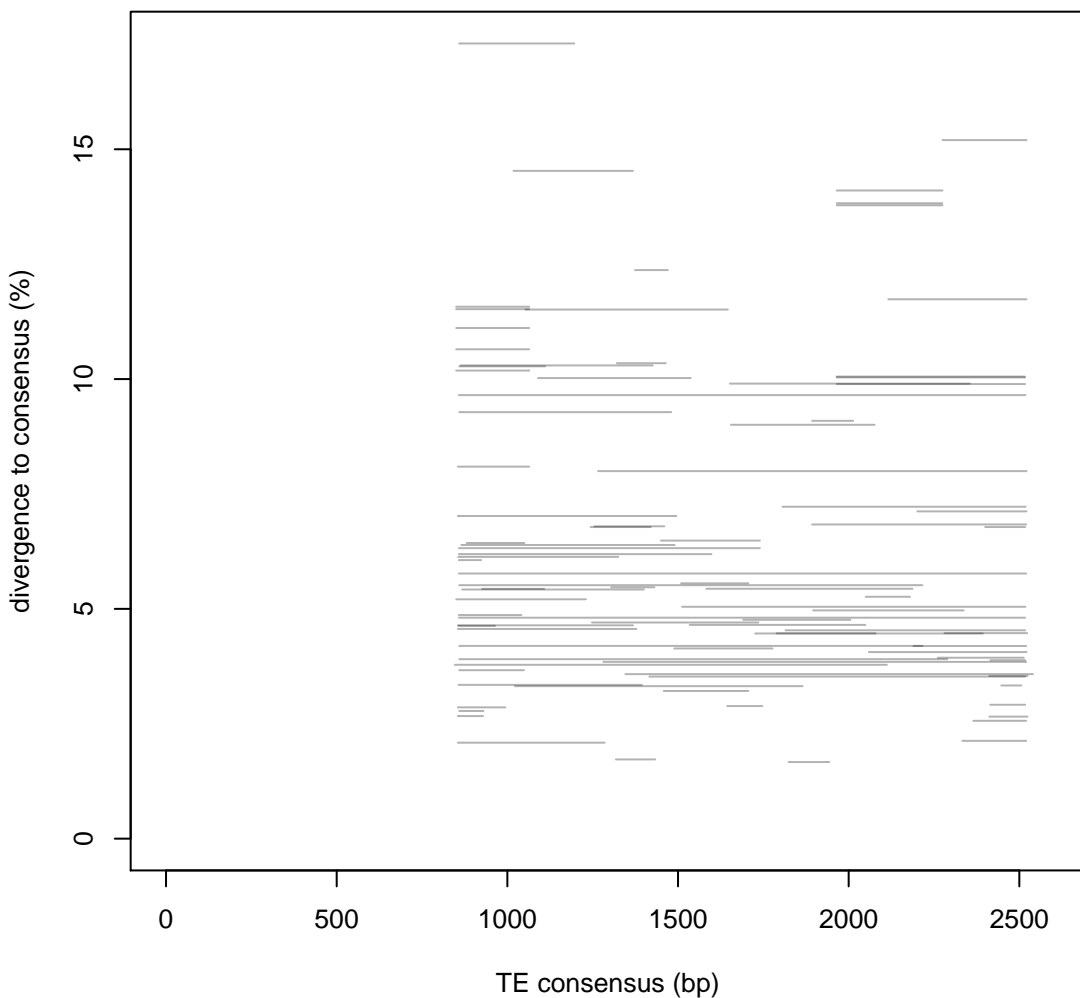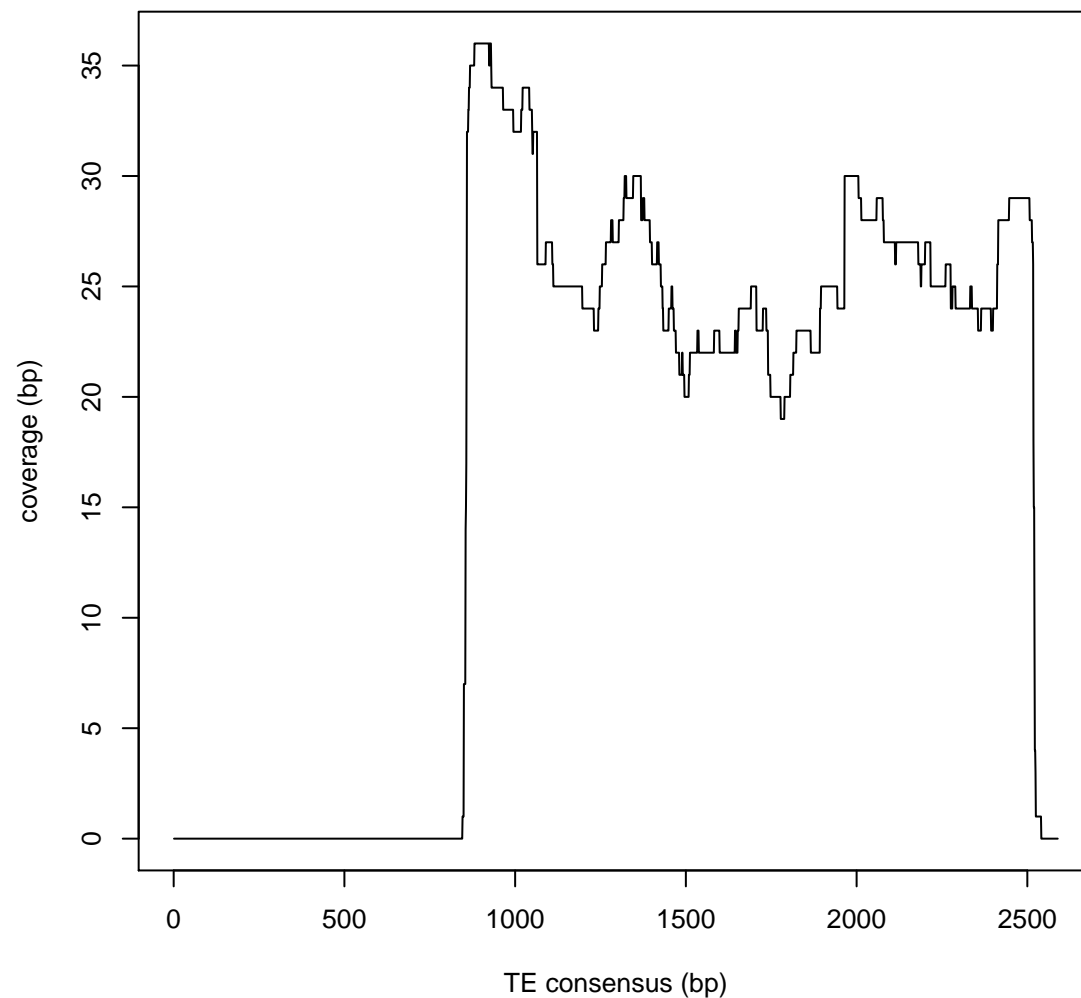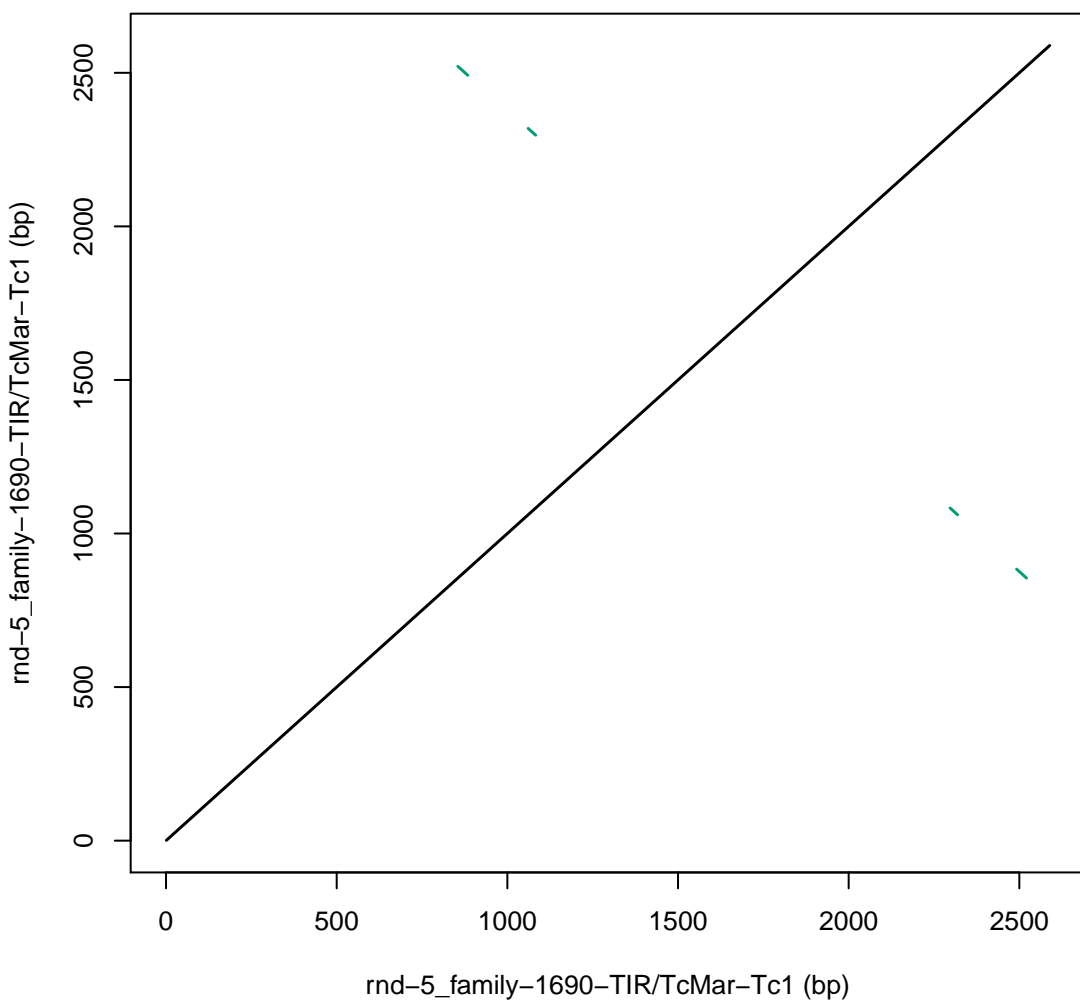**TE: rnd−5_family−1690−TIR/TcMar−Tc1**
consensus size: 2589bp; fragments: 93; full length: 0 (>=2330.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Tc1_DM_tp TIR/TcMar−Tc1

Tc1_DM_tp TIR/TcMar−Tc1