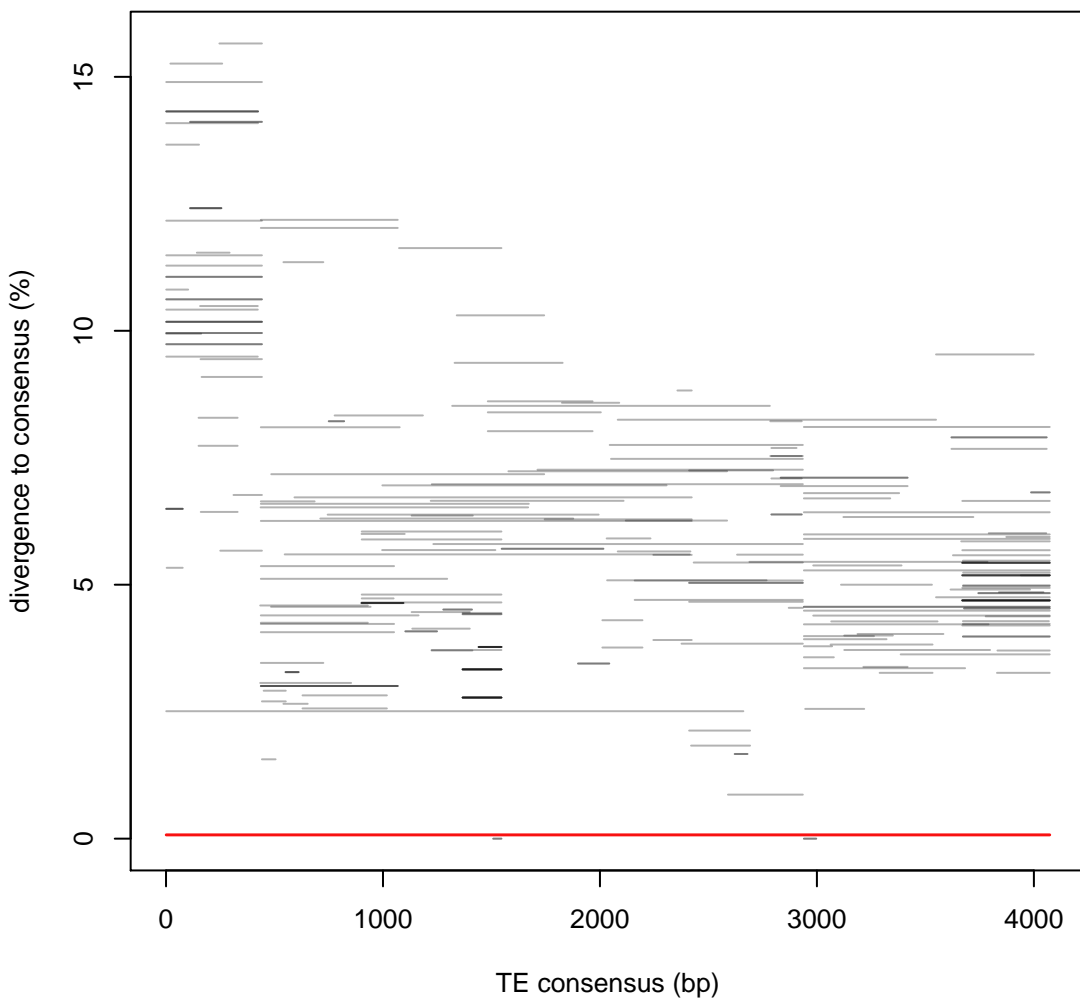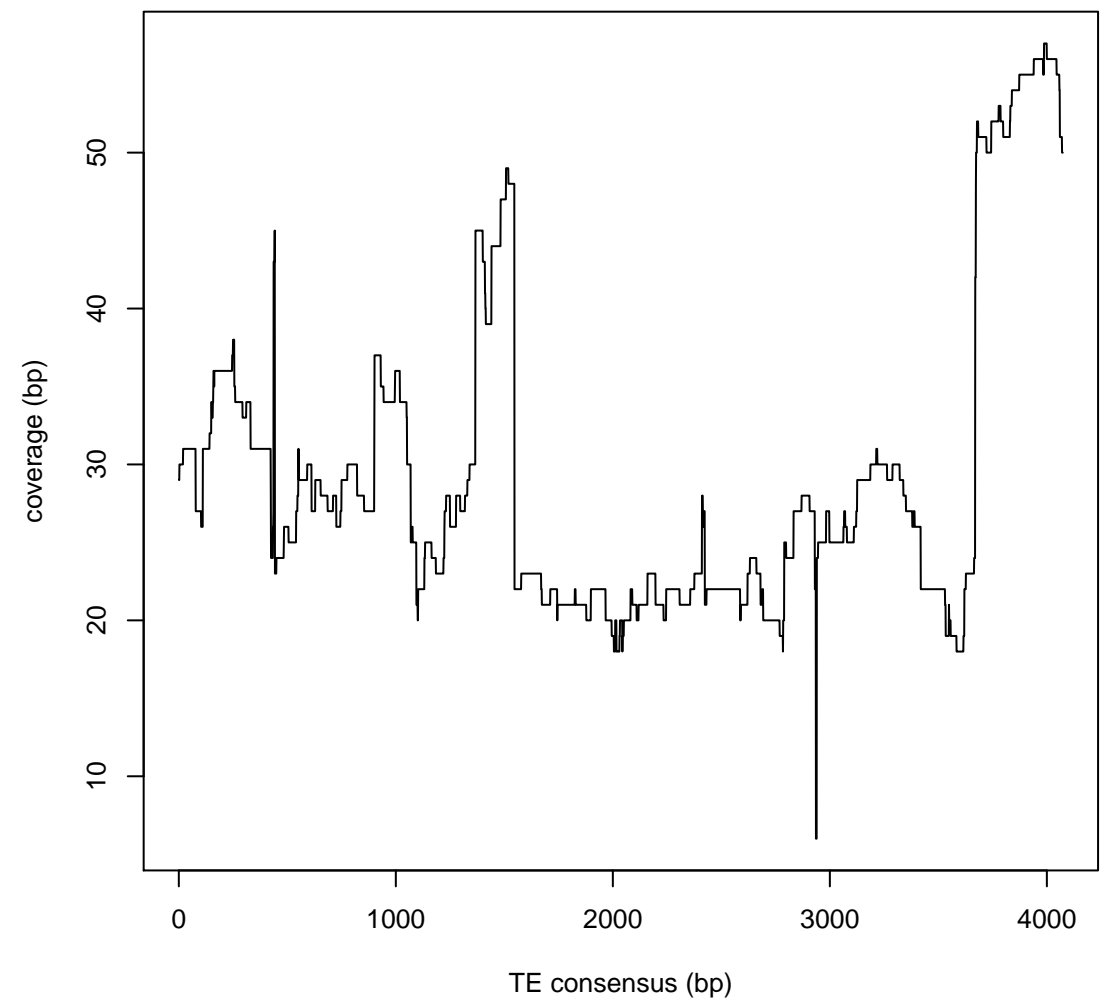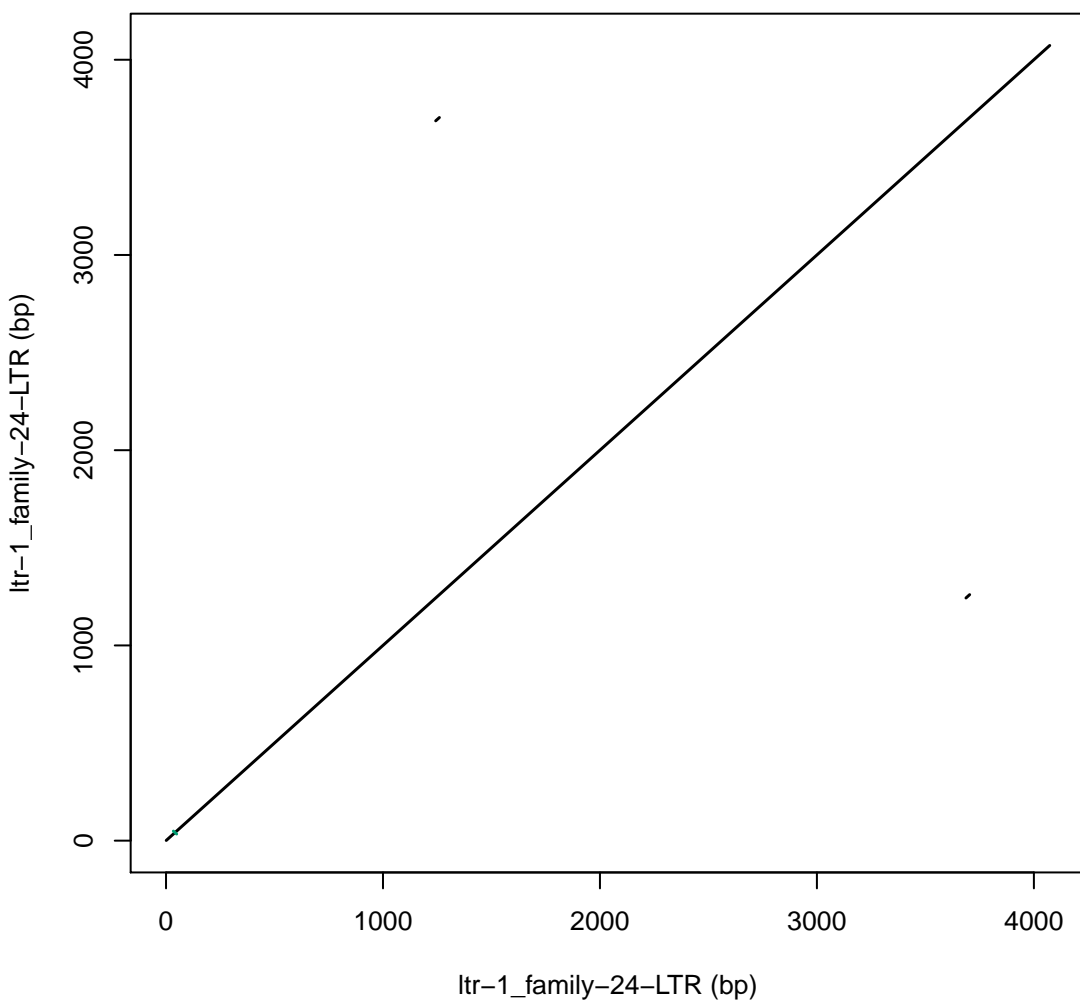# TE: ltr−1_family−24−LTR
consensus size: 4073bp; fragments: 261; full length: 1 (>=3665.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

63_CCri_tp TIR/PIF−Harbinger
BEL−4_AA_pol LTR/Pao
BEL−2_DFi_pol LTR/Pao
BEL−616_AA_pol LTR/Pao
BEL−5_DVir_pol LTR/Pao
BEL−17_DTa_pol LTR/Pao