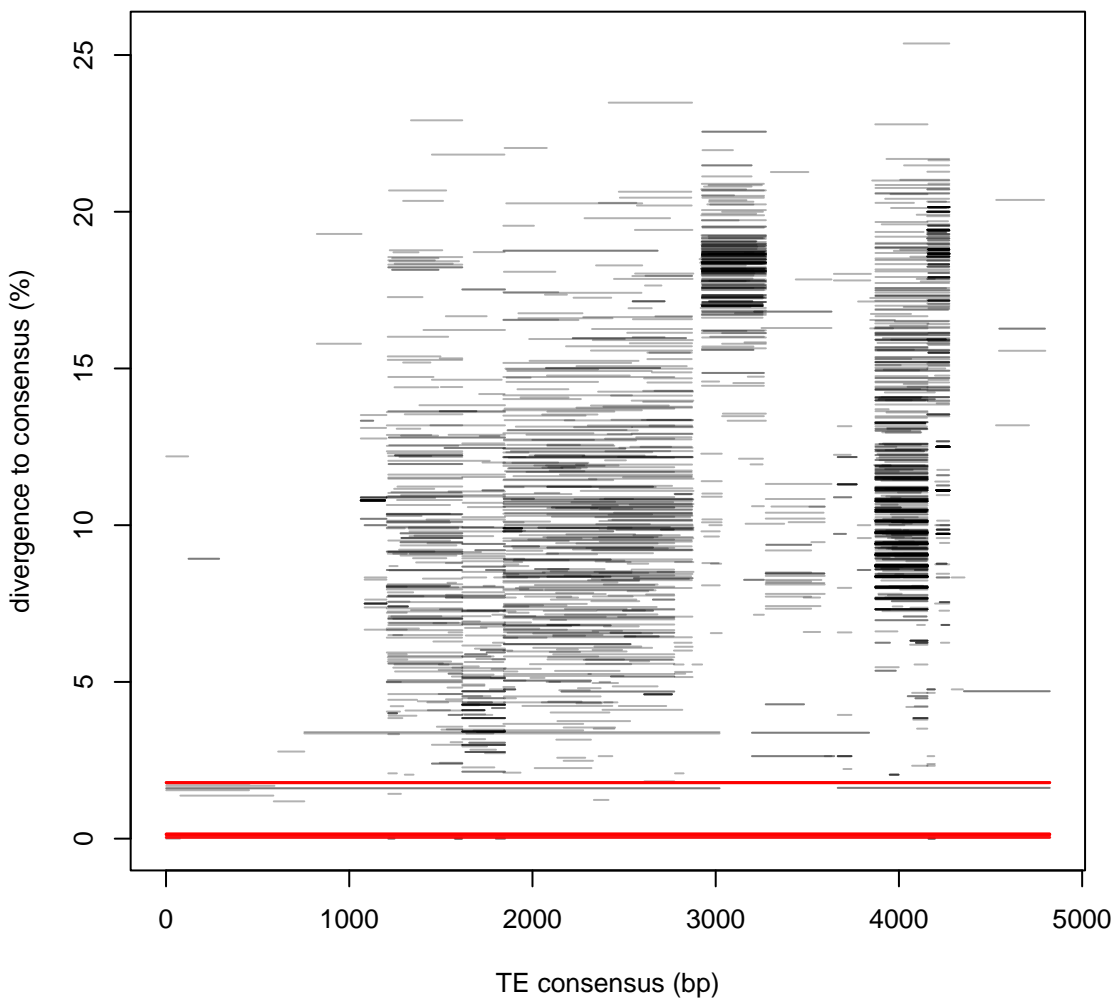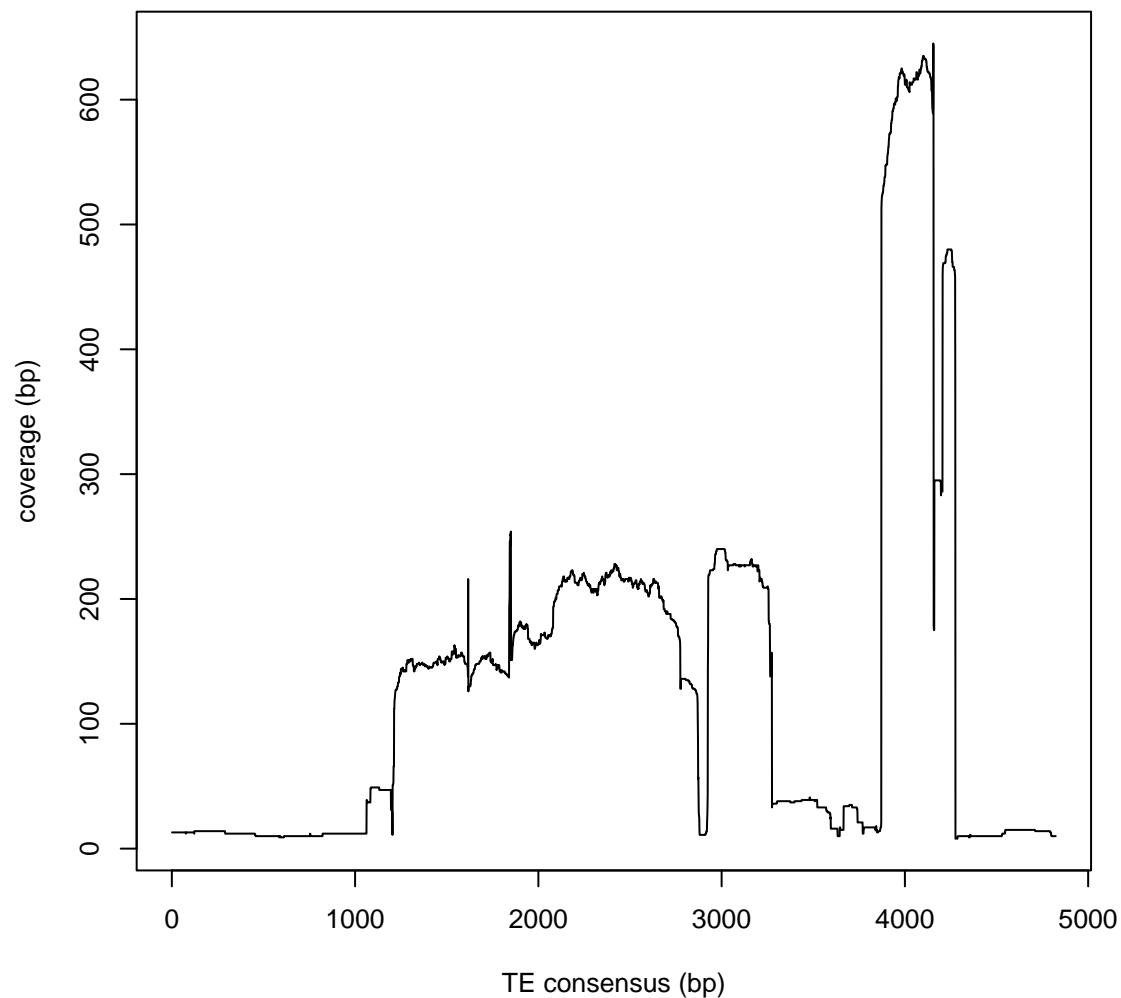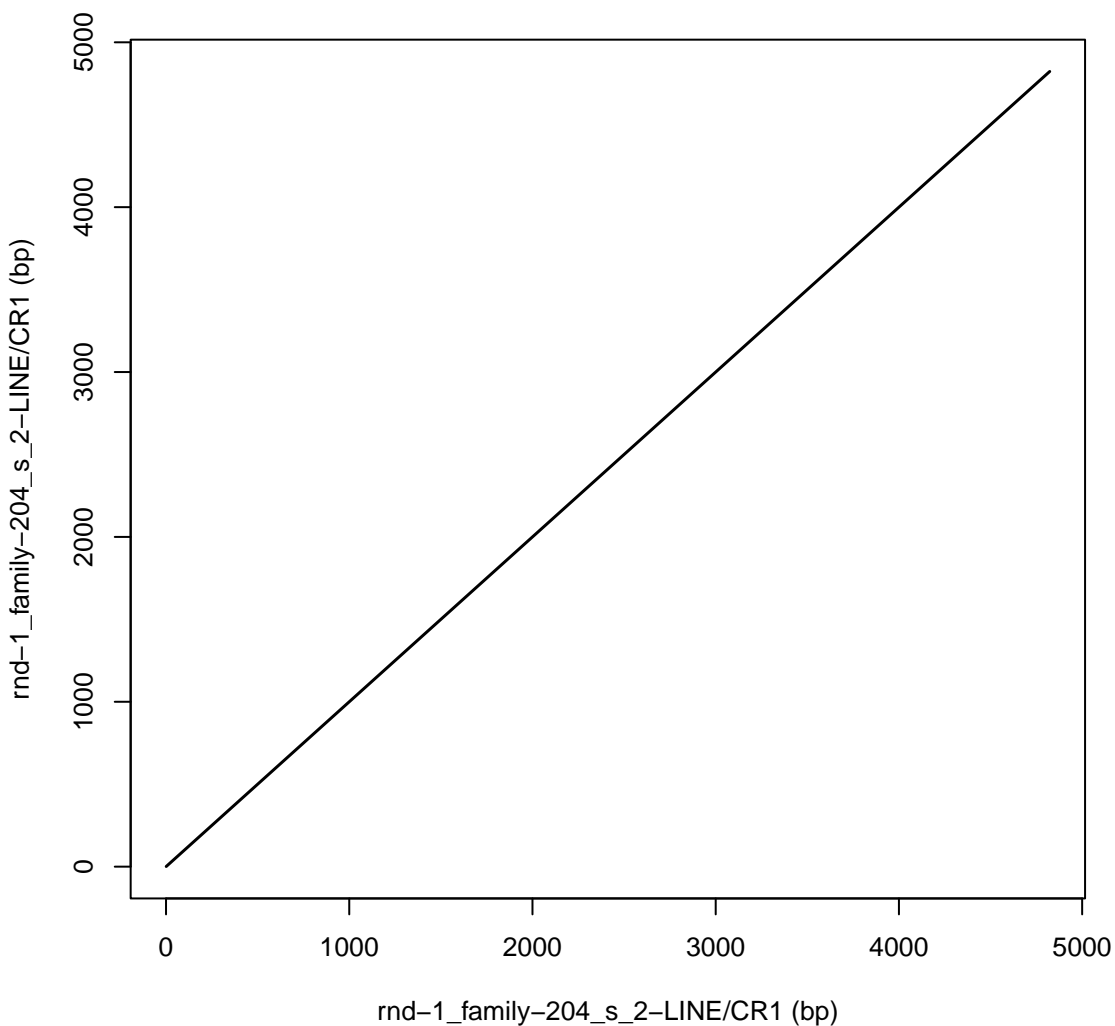**TE: rnd−1_family−204_s_2−LINE/CR1**
consensus size: 4823bp; fragments: 2272; full length: 6 (>=4340.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

CR1−8_AMi_pol LINE/CR1