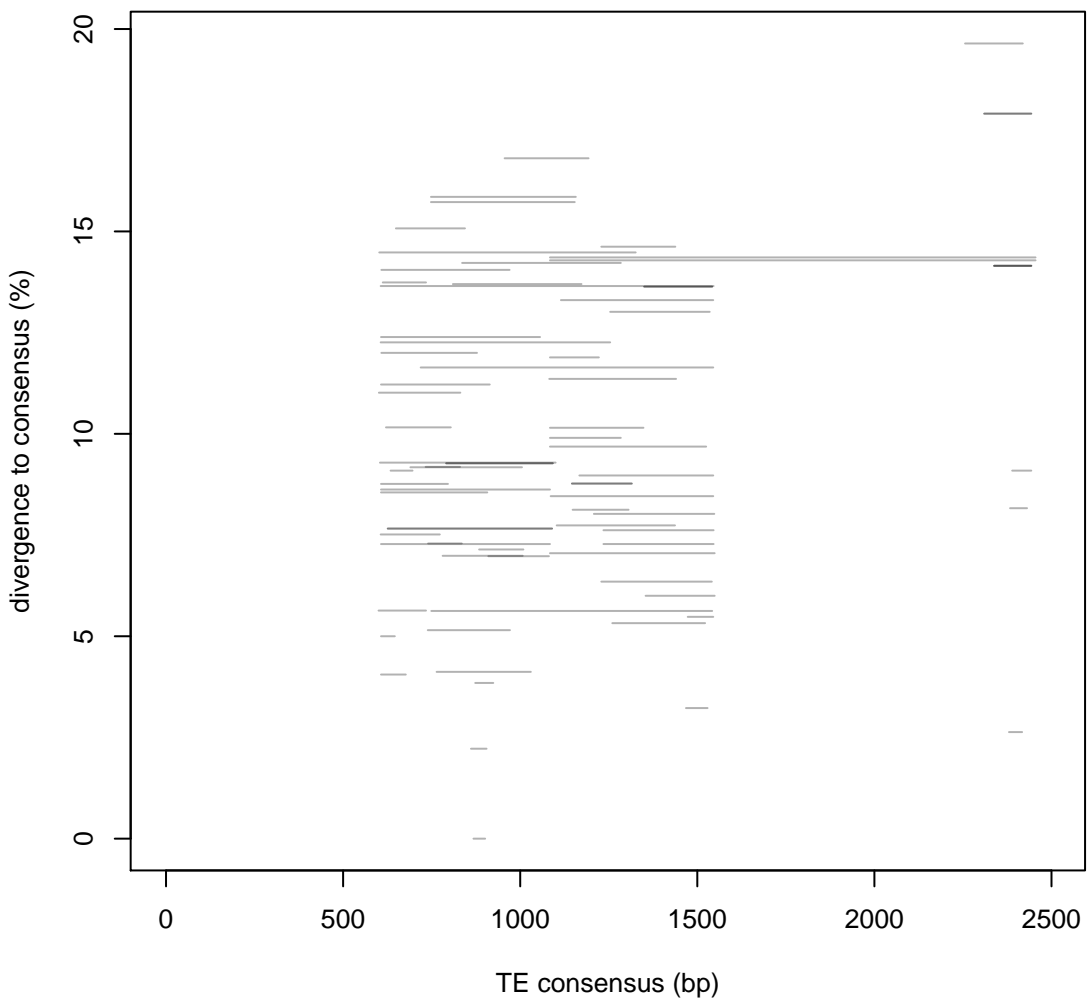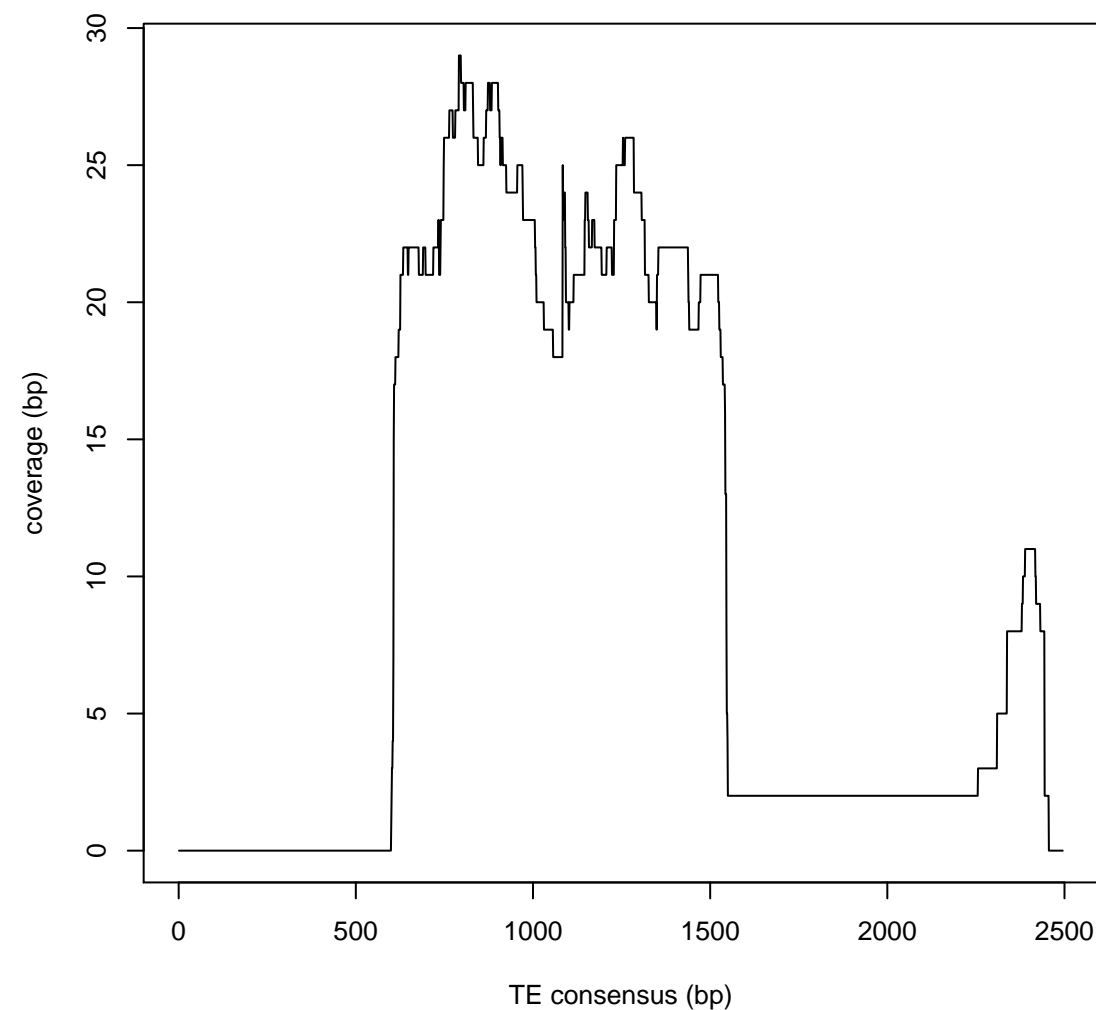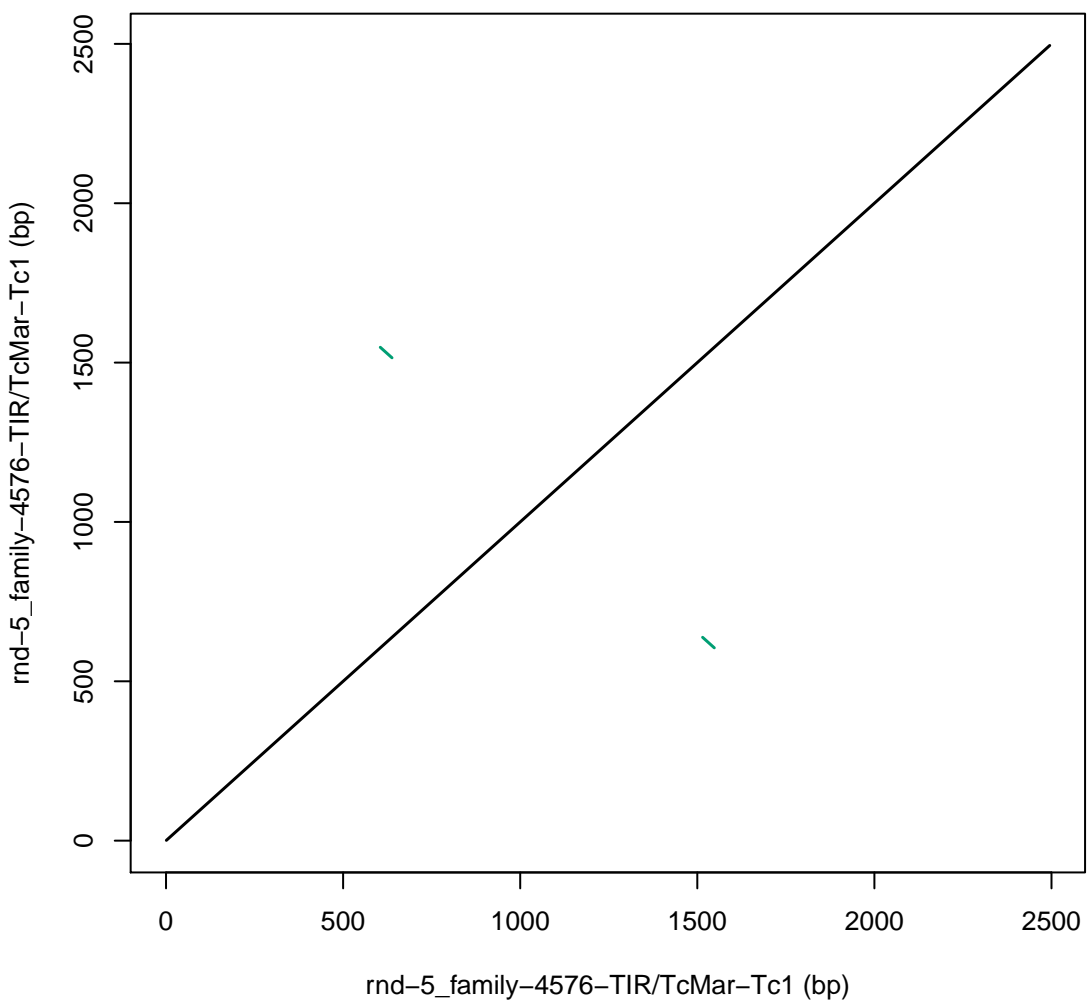**TE: rnd−5_family−4576−TIR/TcMar−Tc1**
consensus size: 2495bp; fragments: 79; full length: 0 (>=2245.5bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

Mariner2_DM_tp TIR/TcMar−Tc1