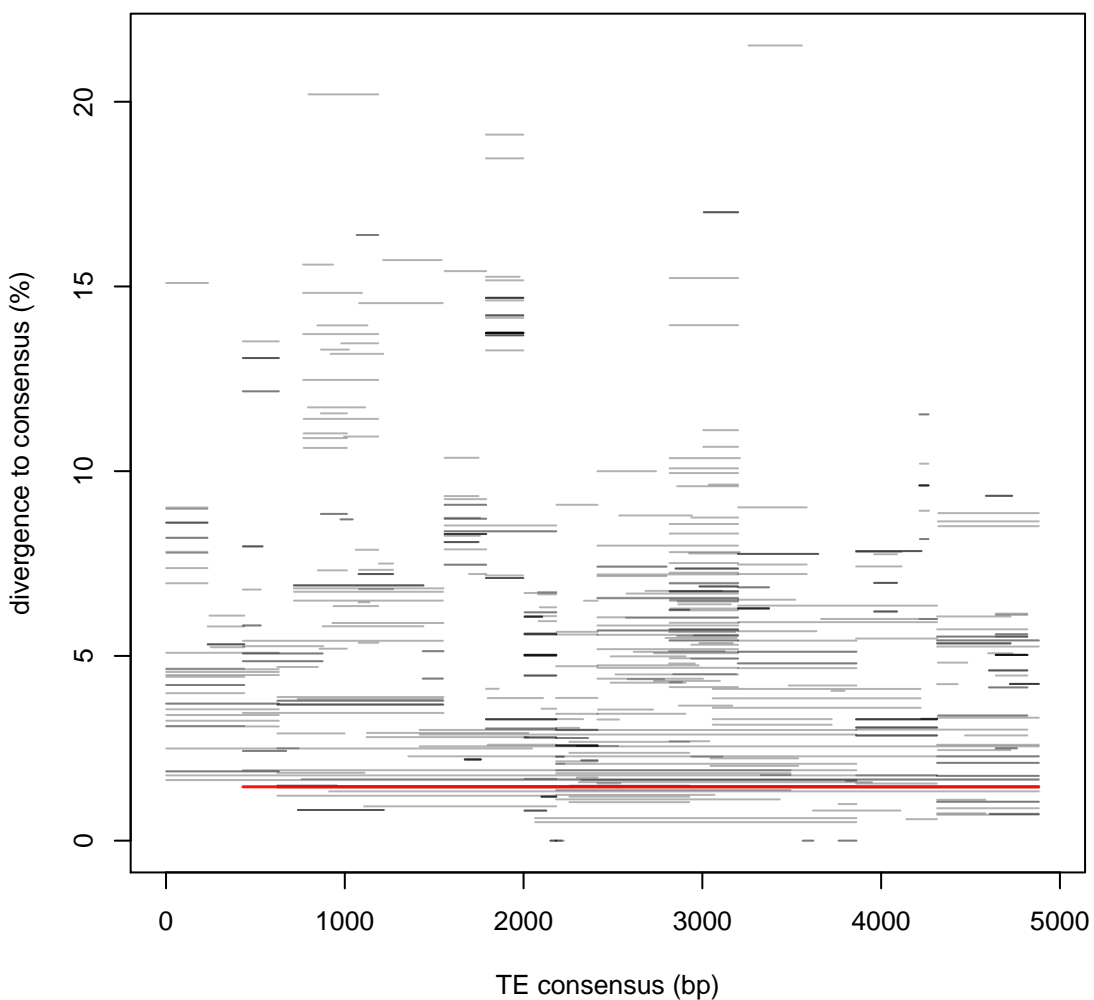
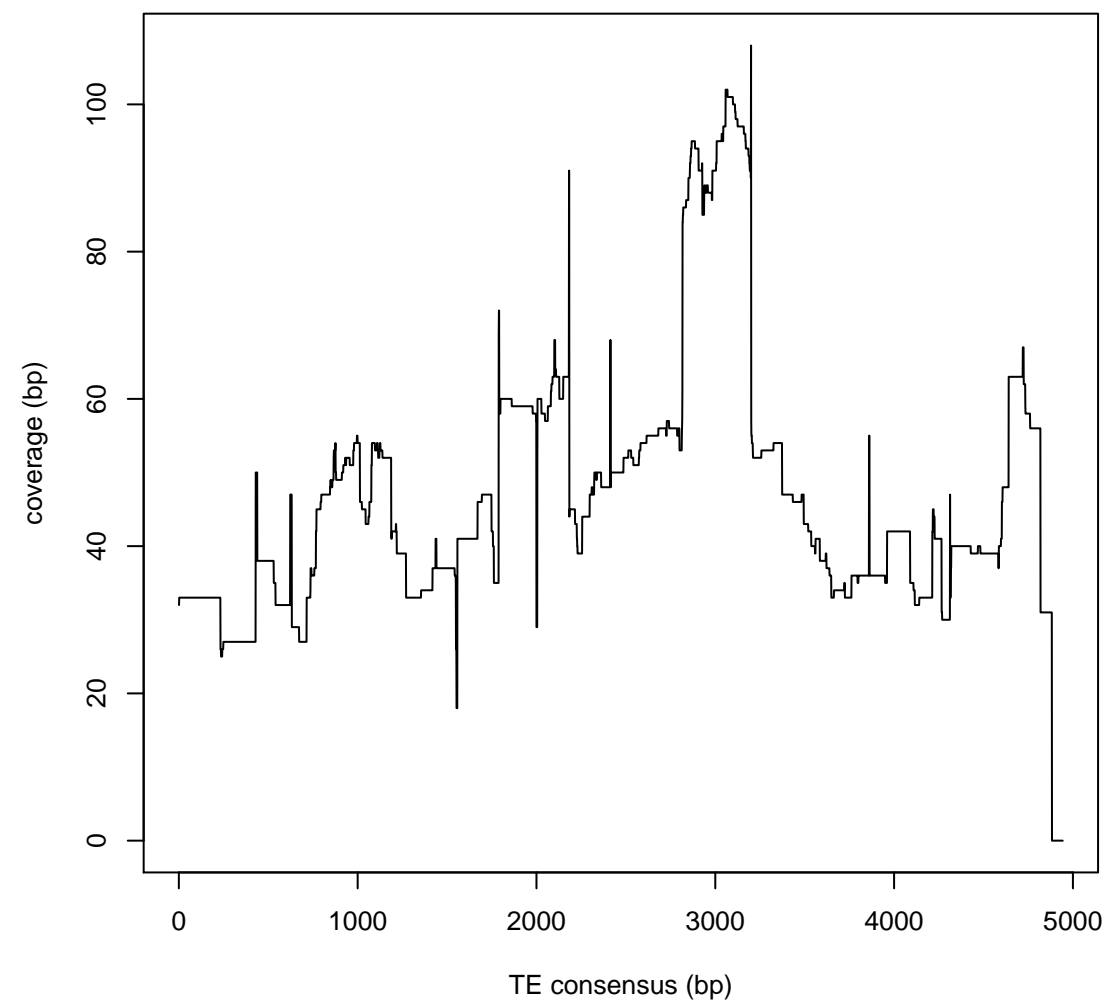


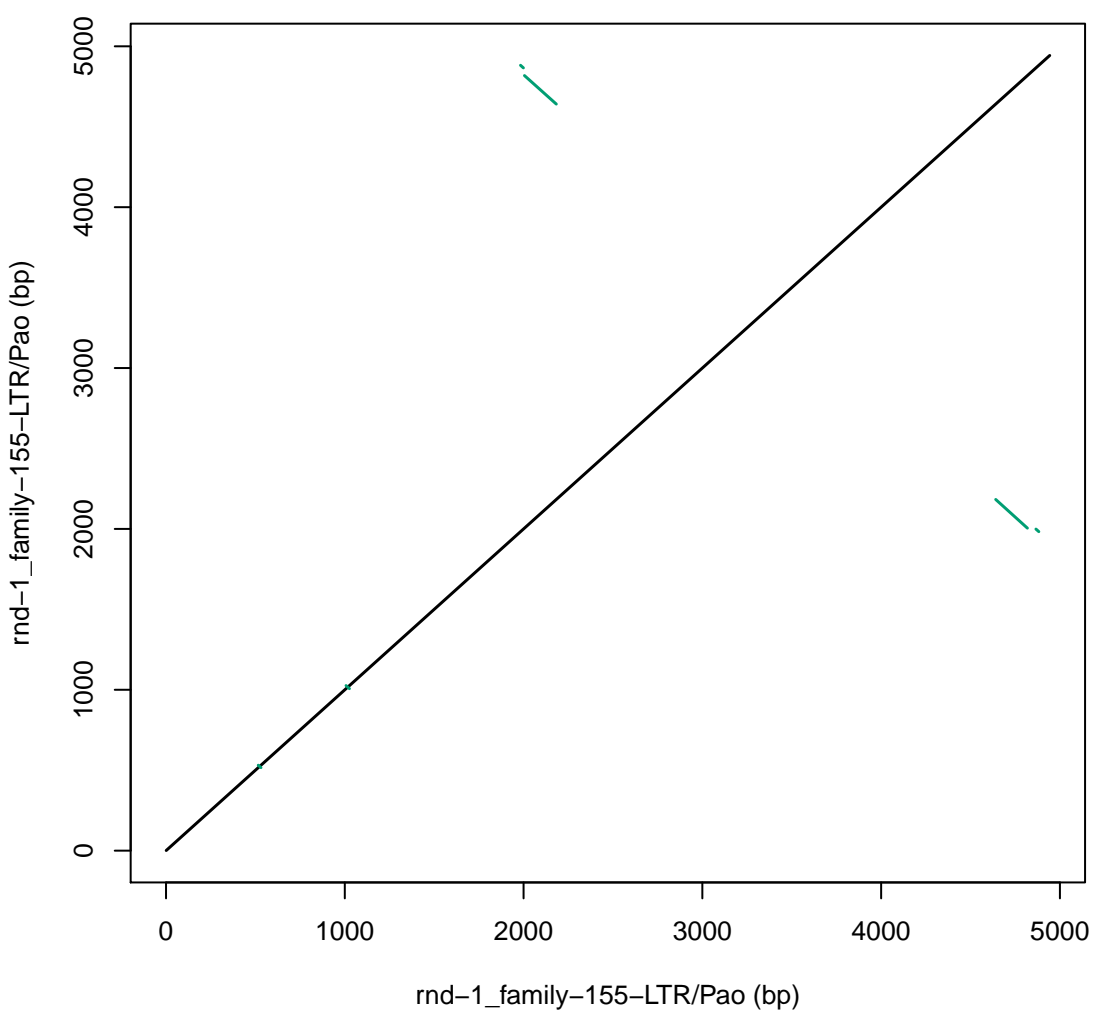
TE: rnd-1_family-155-LTR/Pao
consensus size: 4943bp; fragments: 531; full length: 1 (>=4448.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

