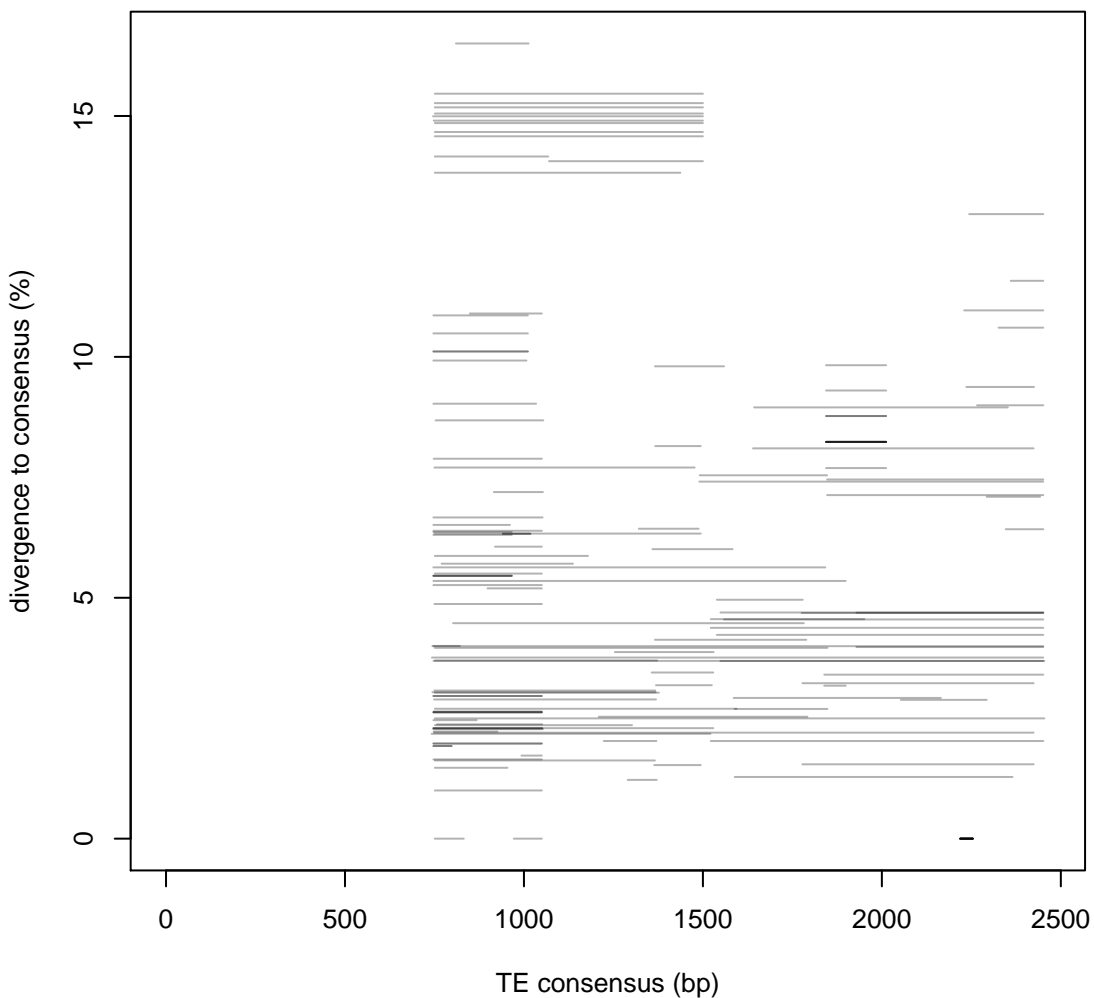
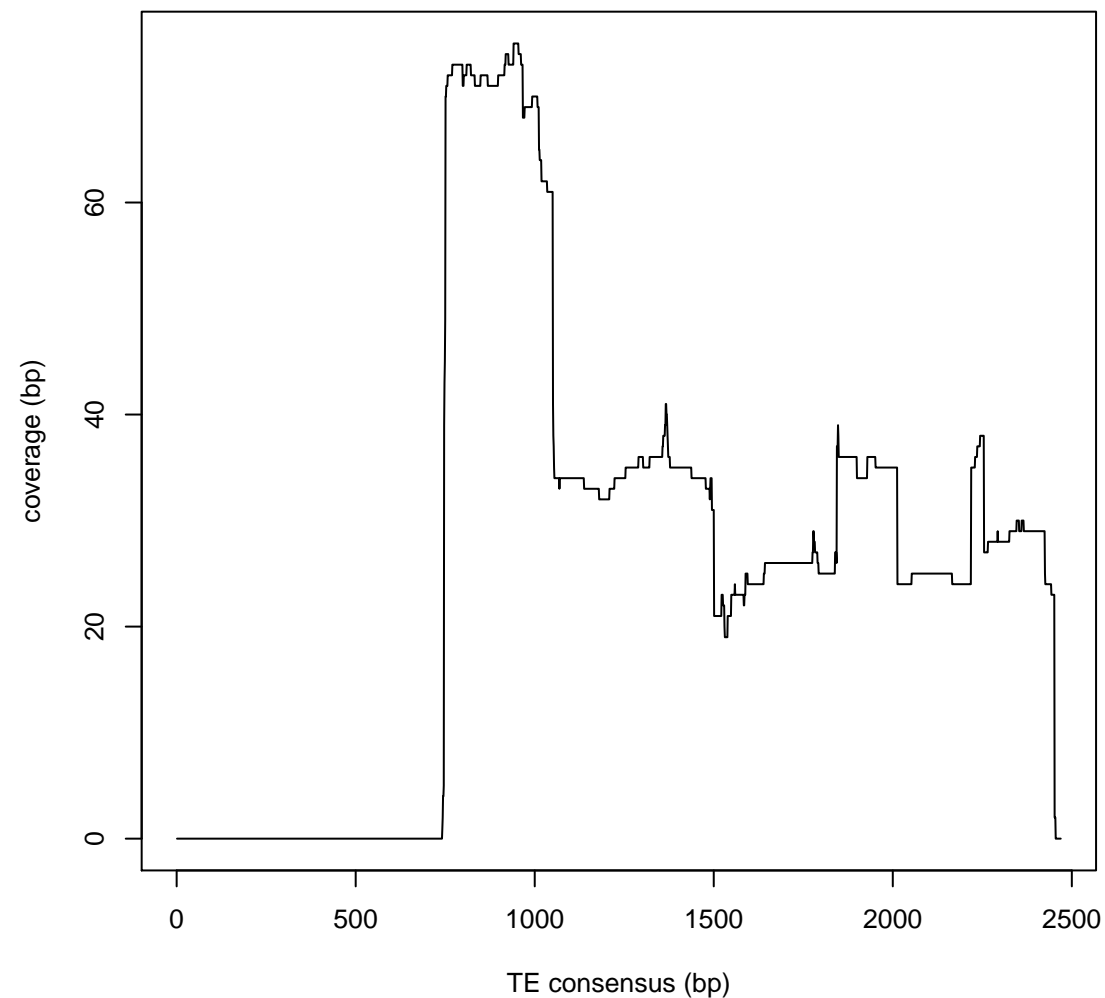


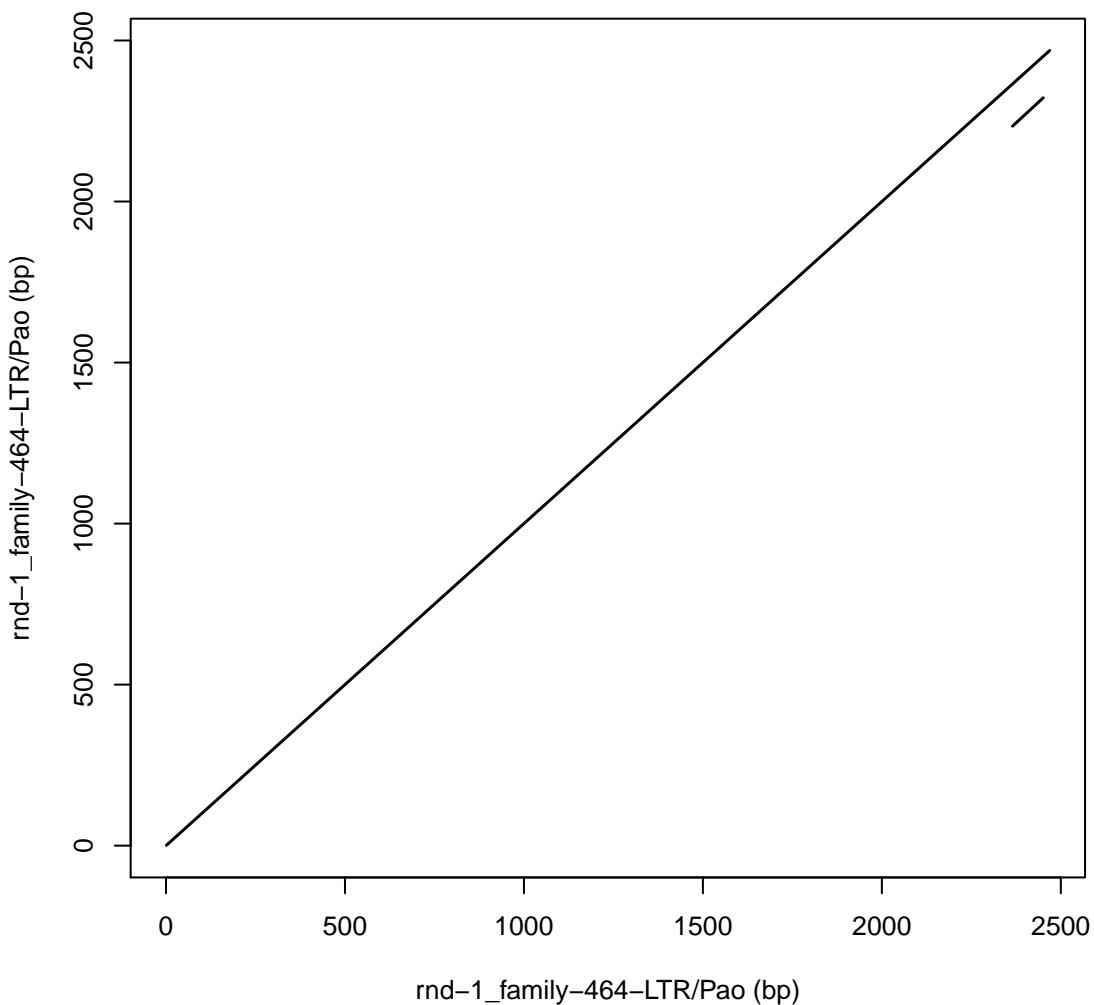
TE: rnd-1_family-464-LTR/Pao
consensus size: 2469bp; fragments: 151; full length: 0 (>=2222.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

