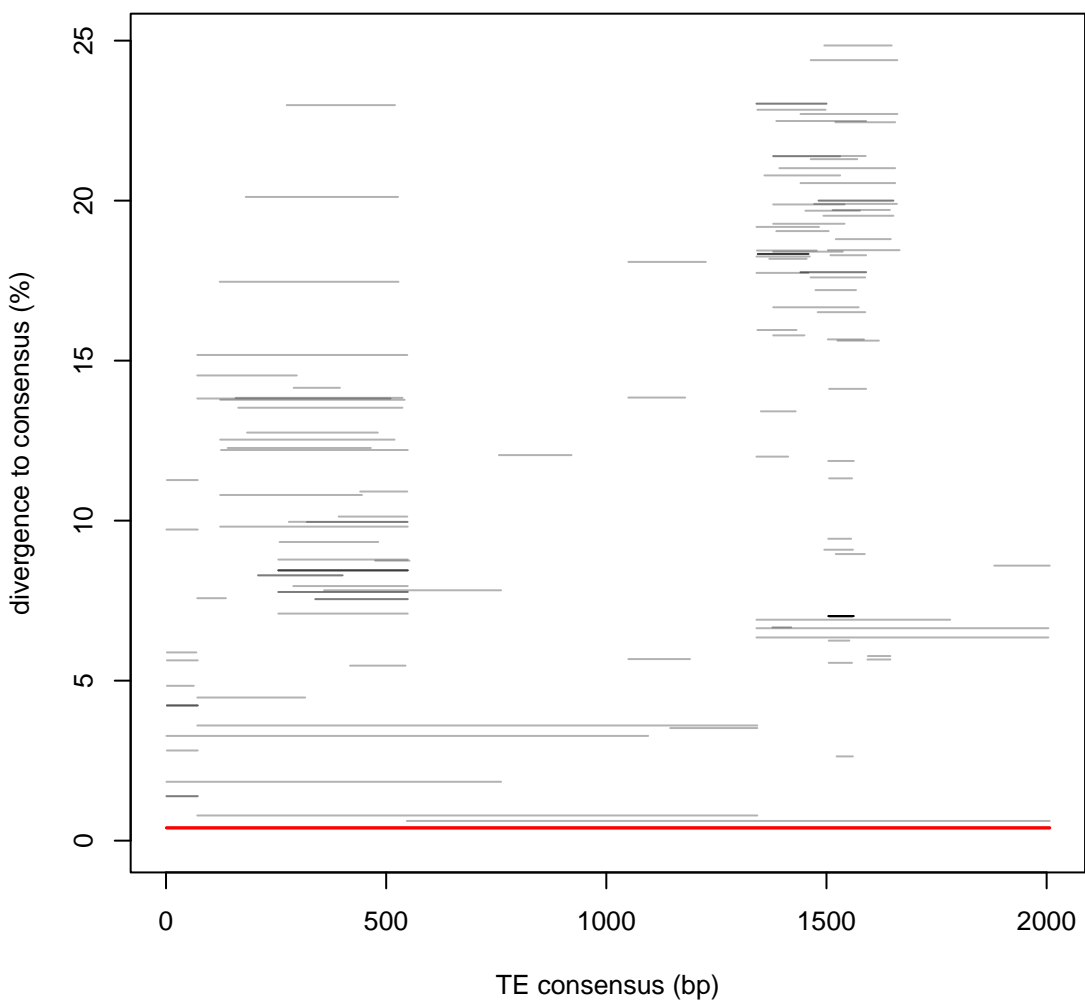
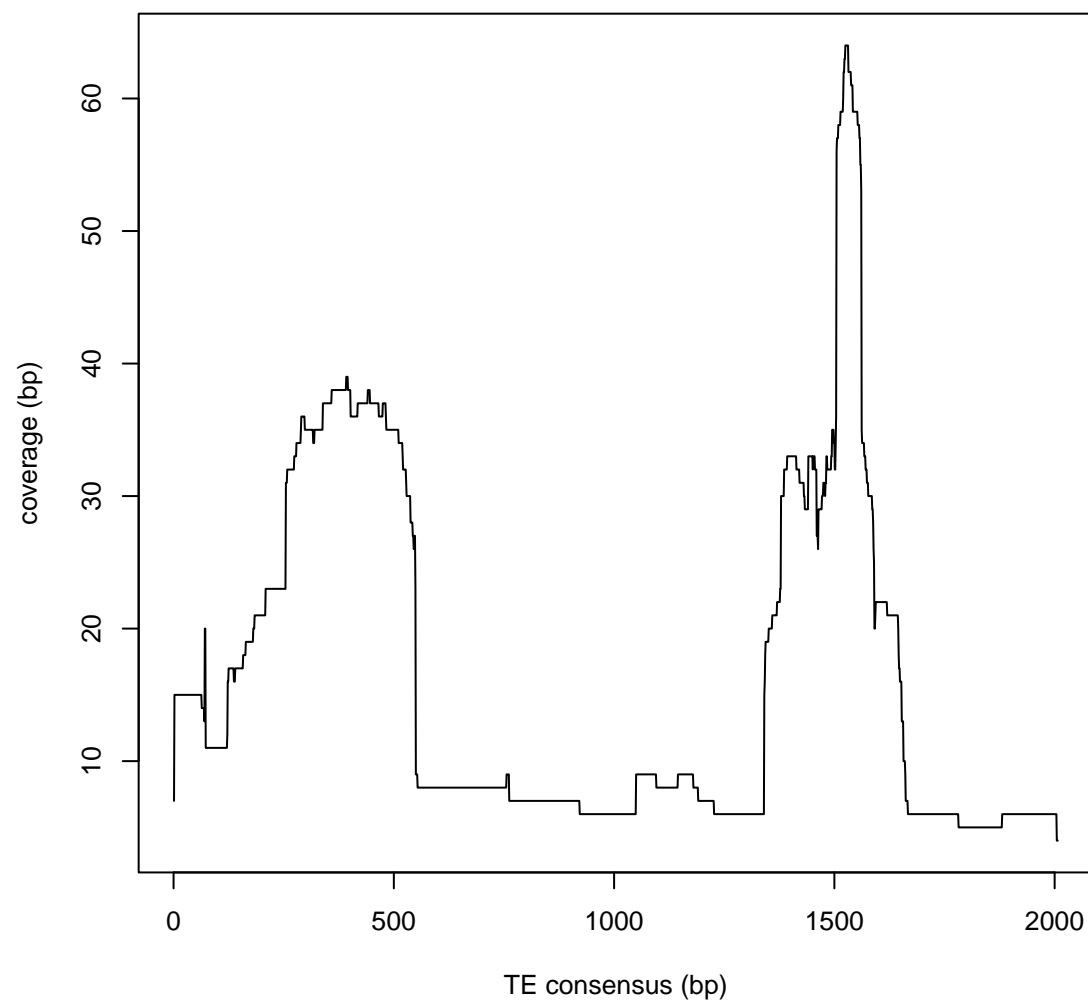


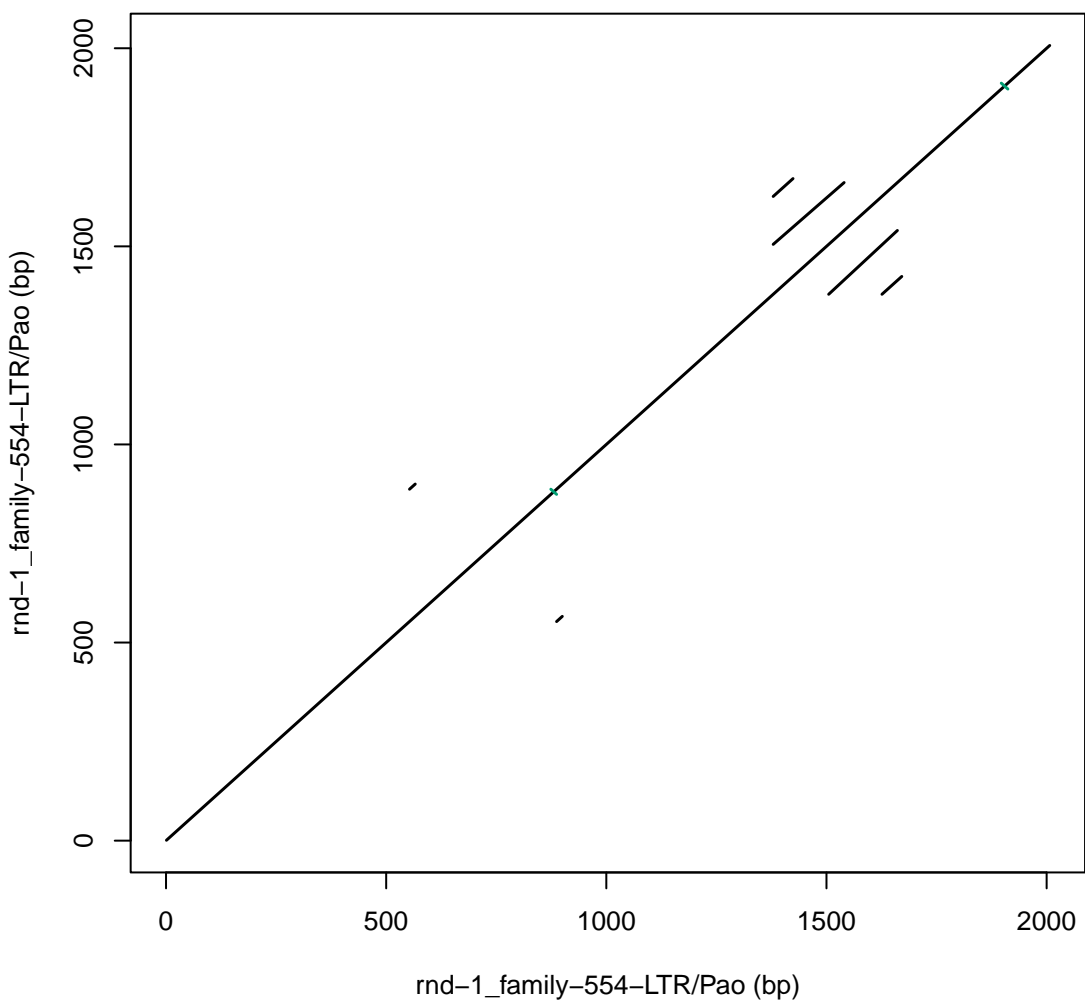
TE: rnd-1_family-554-LTR/Pao
consensus size: 2007bp; fragments: 144; full length: 2 (≥ 1806.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

