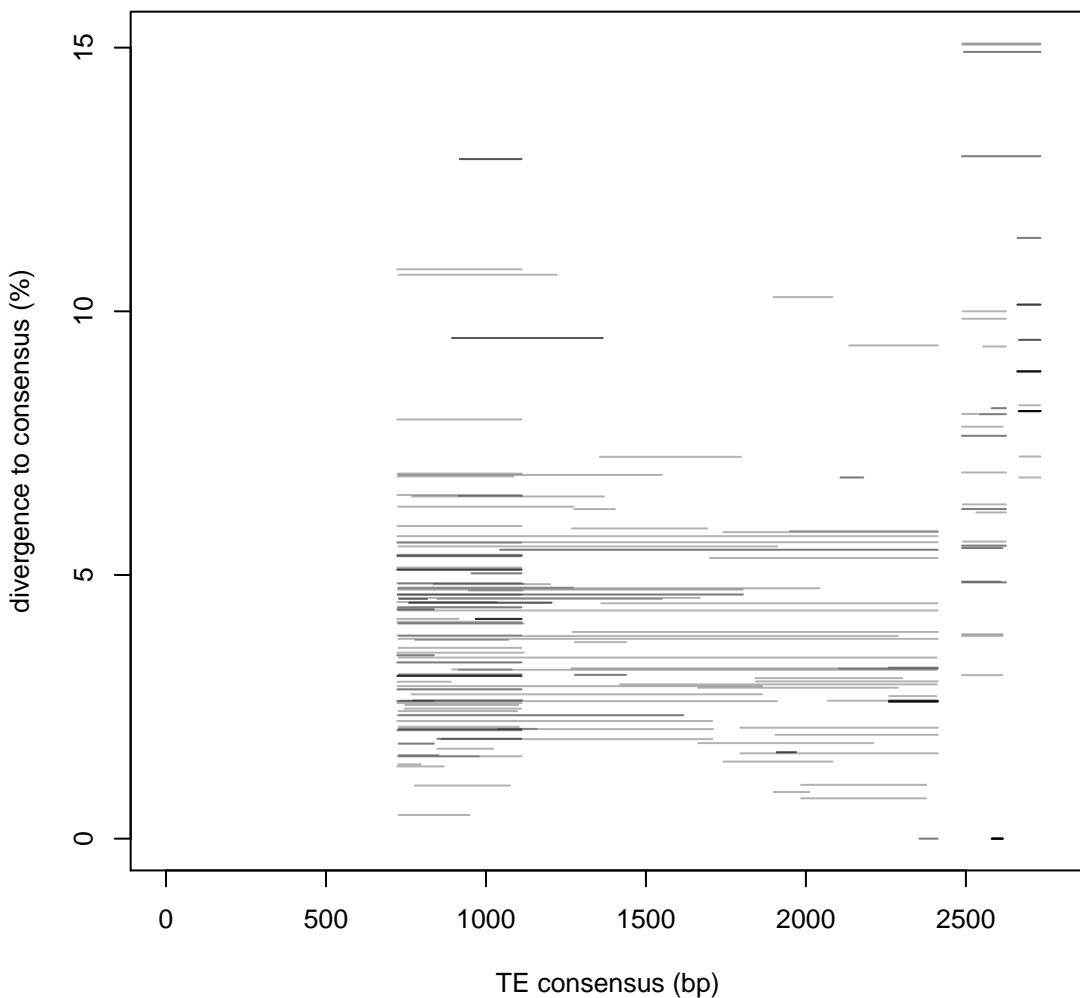
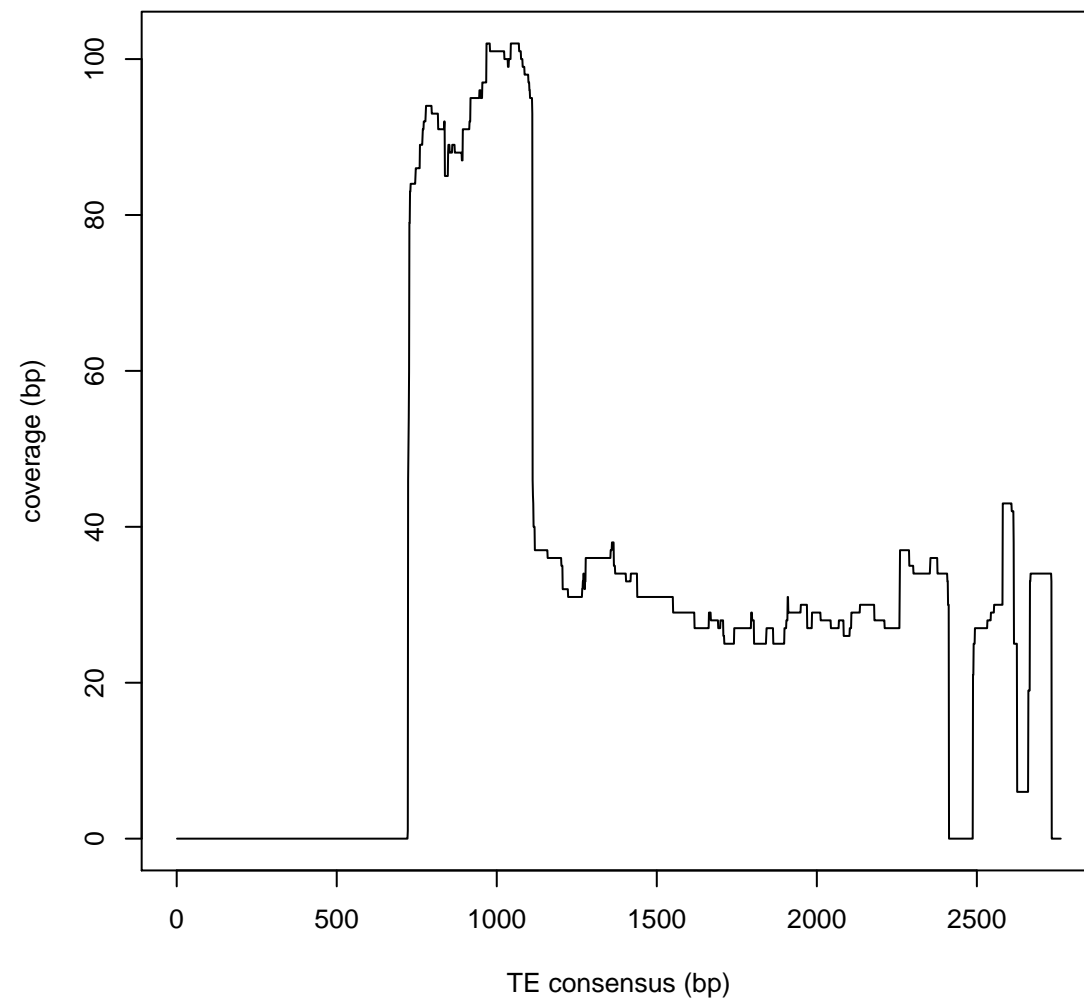


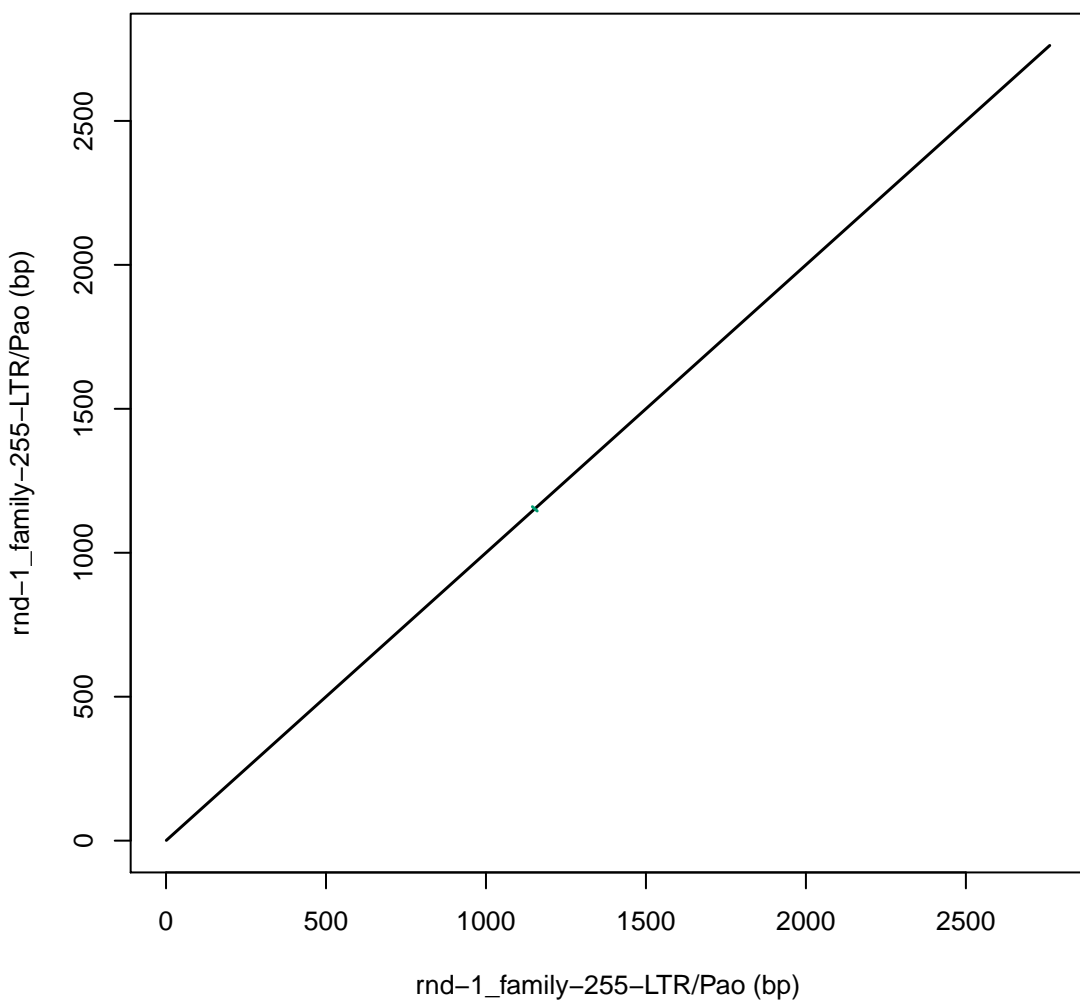
TE: rnd-1_family-255-LTR/Pao
consensus size: 2762bp; fragments: 236; full length: 0 (≥ 2485.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

