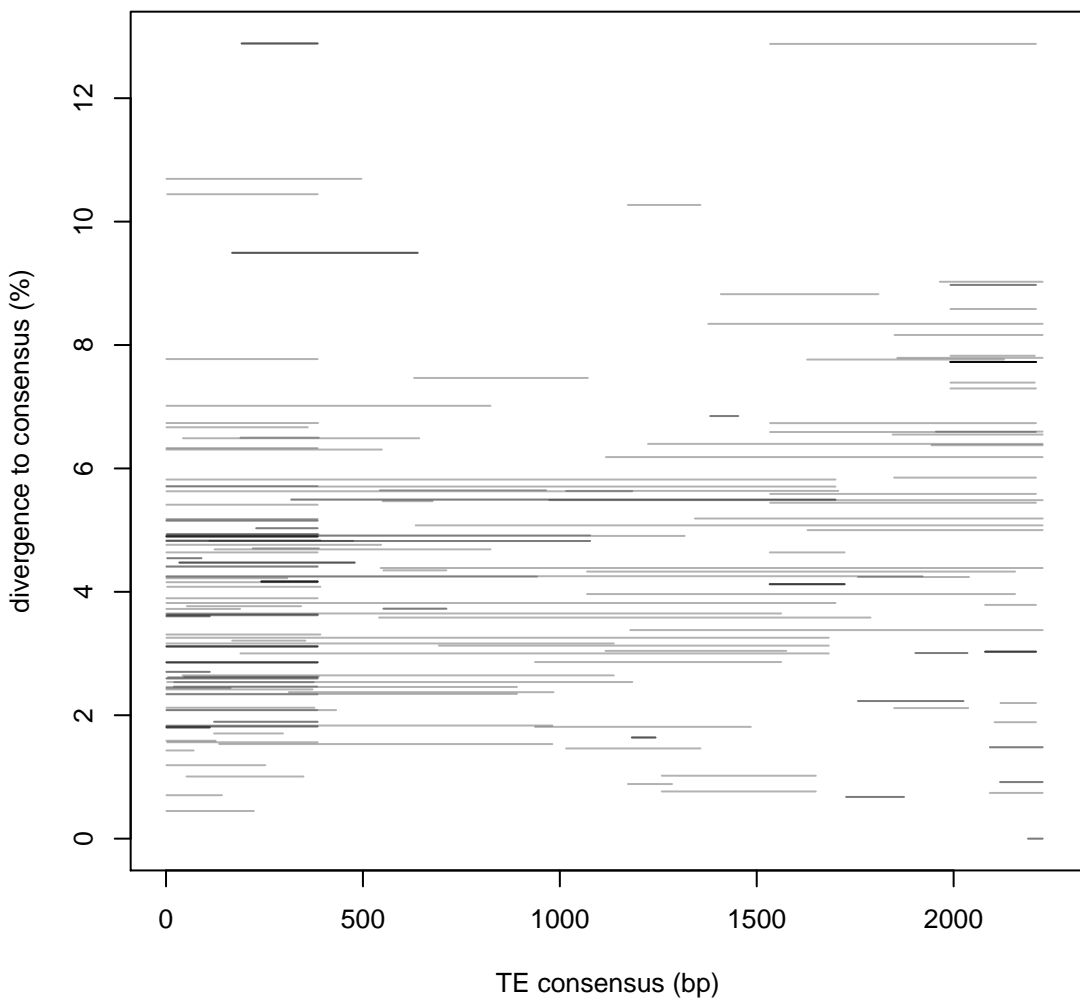
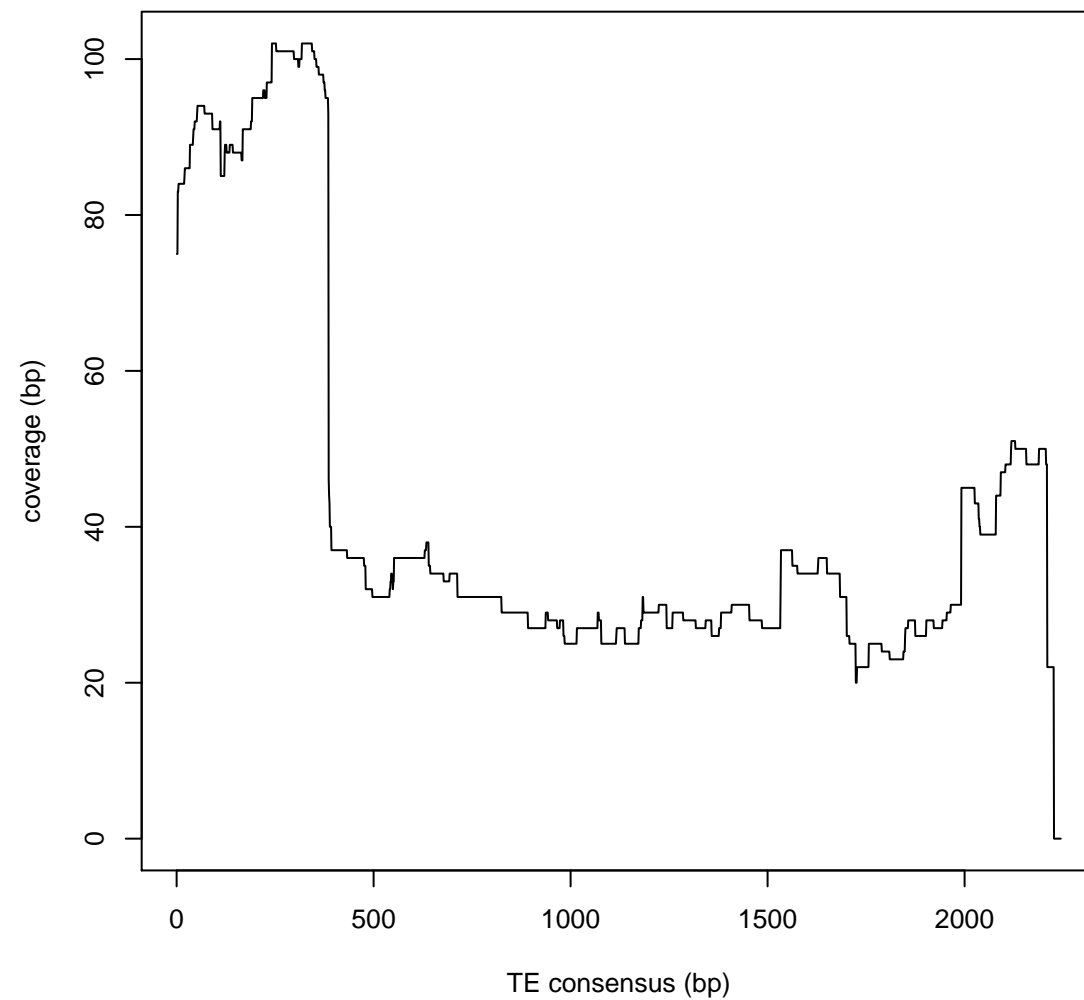


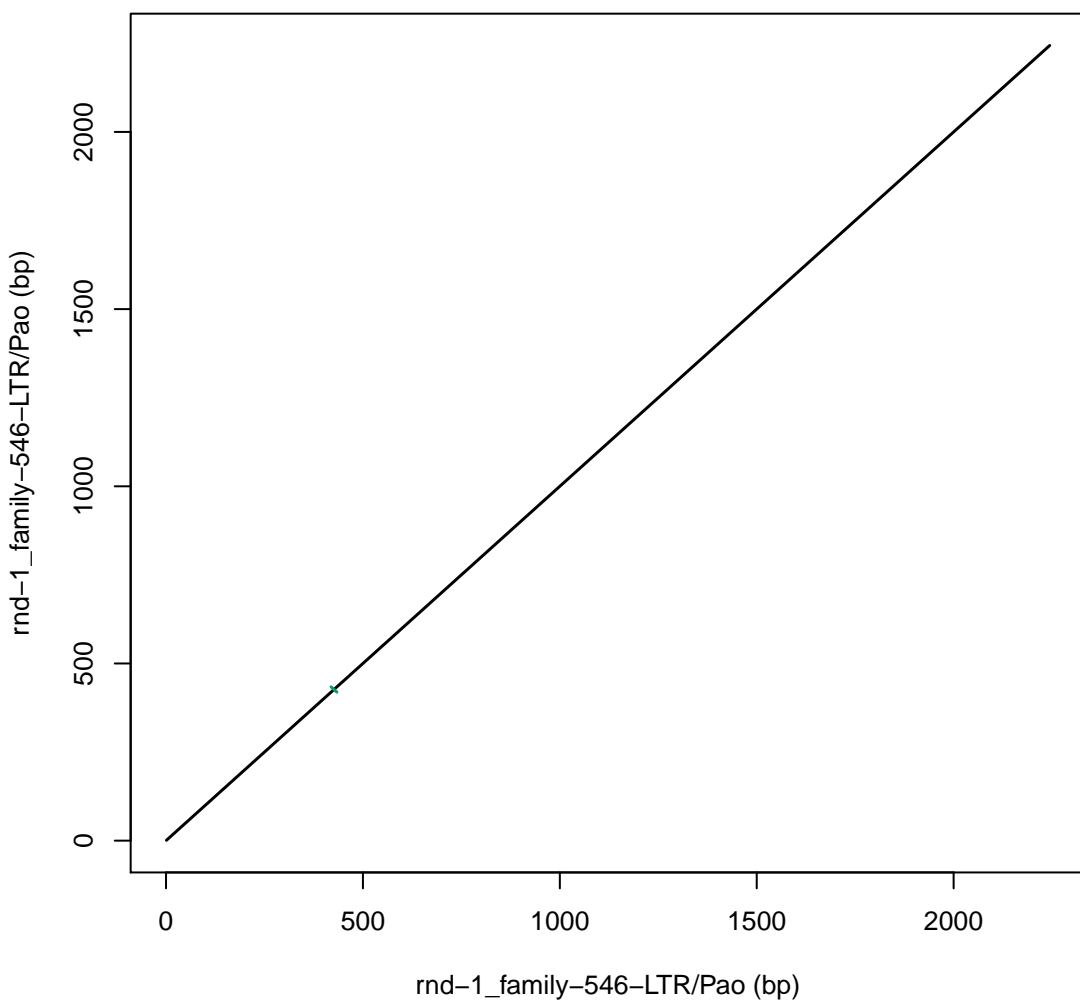
TE: rnd-1\_family-546-LTR/Pao  
consensus size: 2244bp; fragments: 209; full length: 0 (>=2019.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

