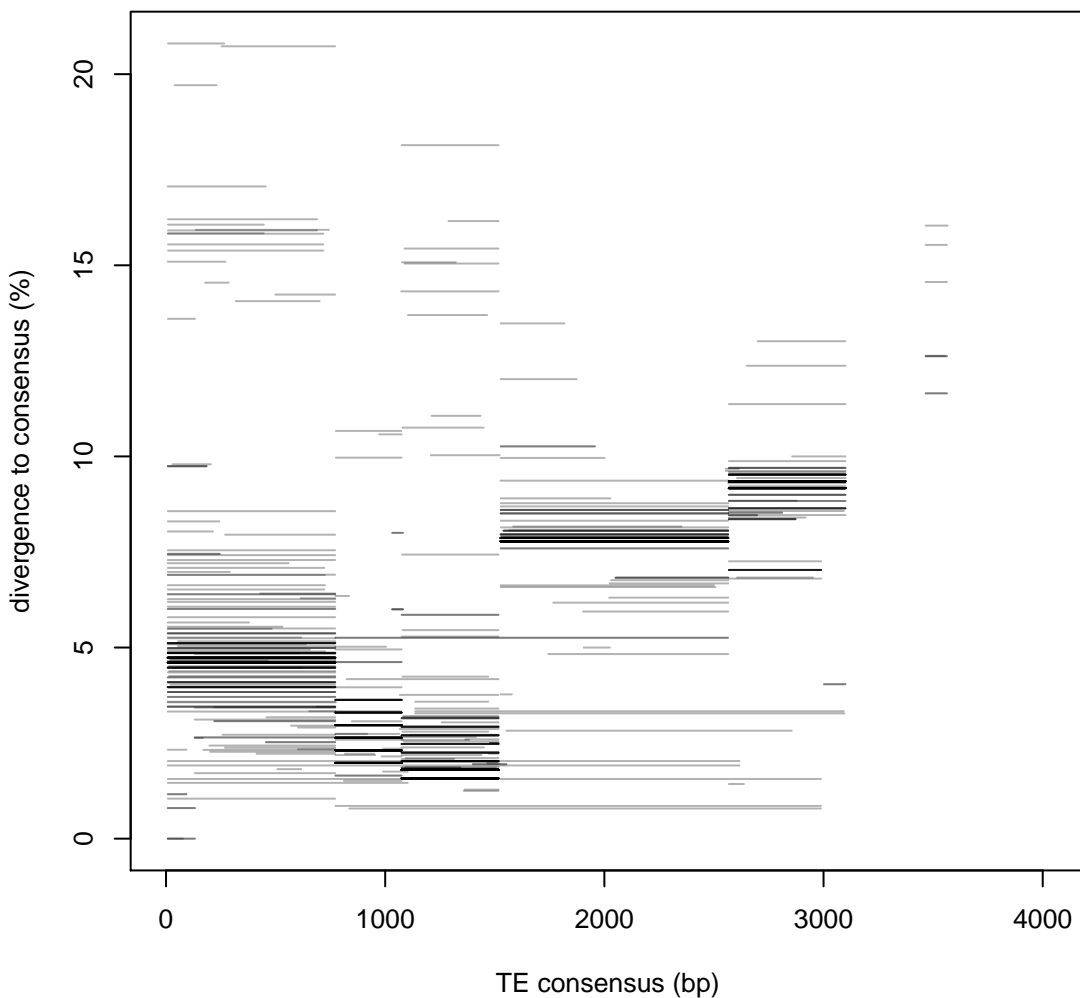
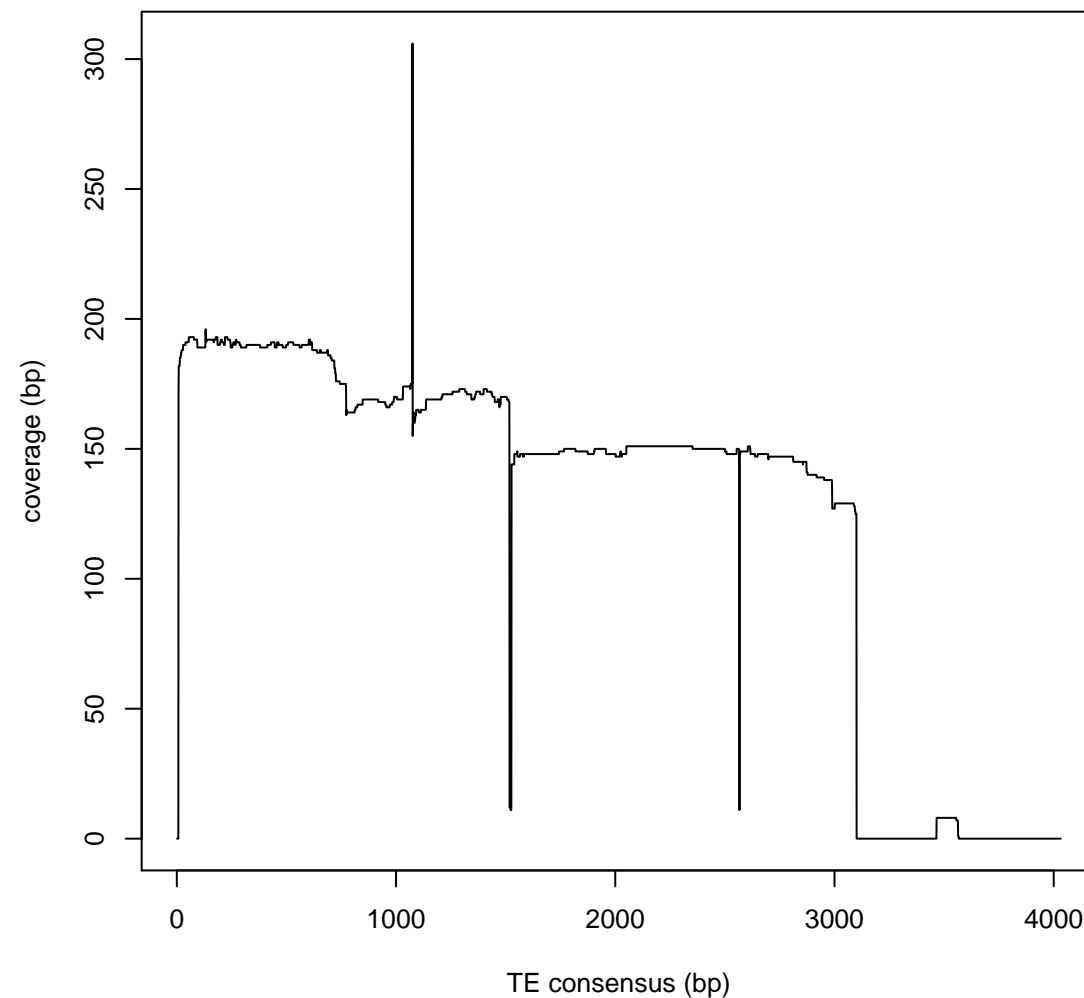


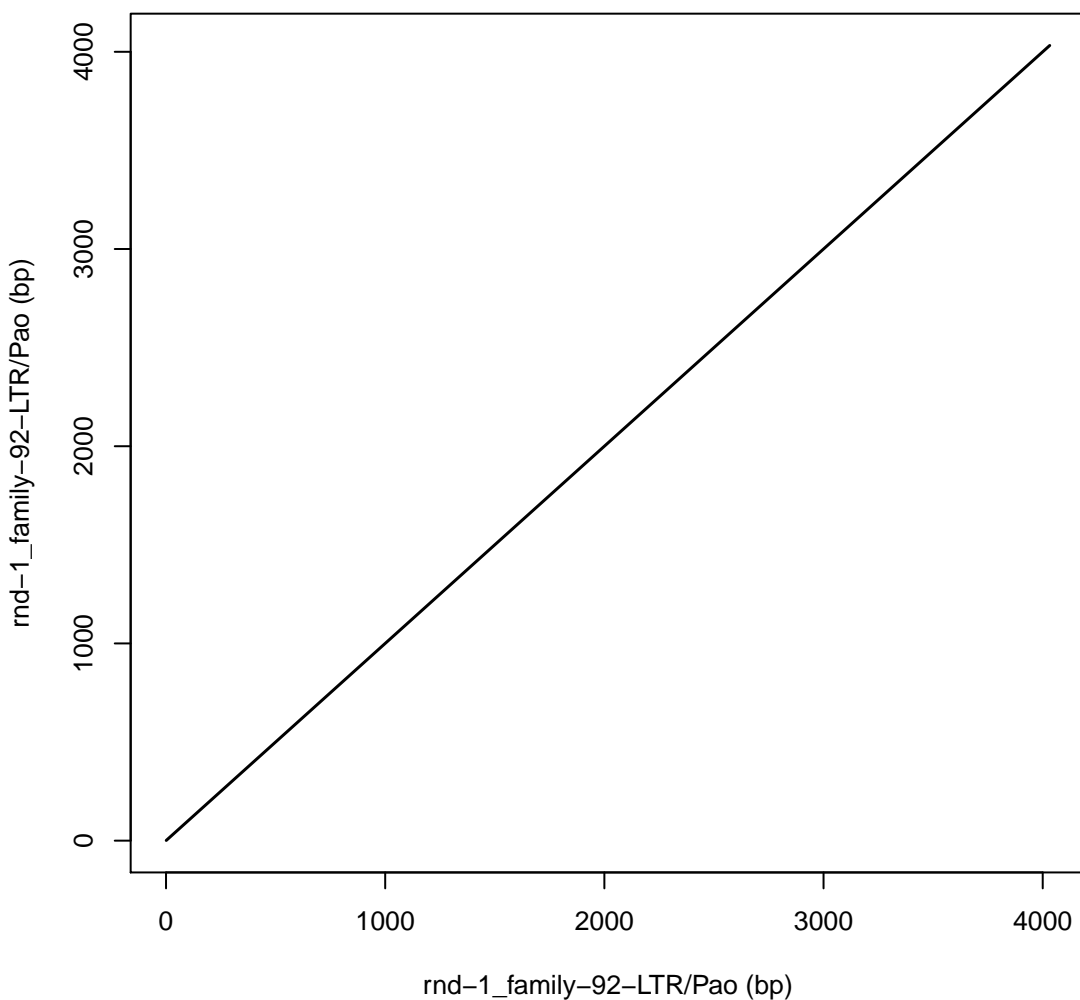
TE: rnd-1_family-92-LTR/Pao
consensus size: 4032bp; fragments: 883; full length: 0 (>=3628.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

