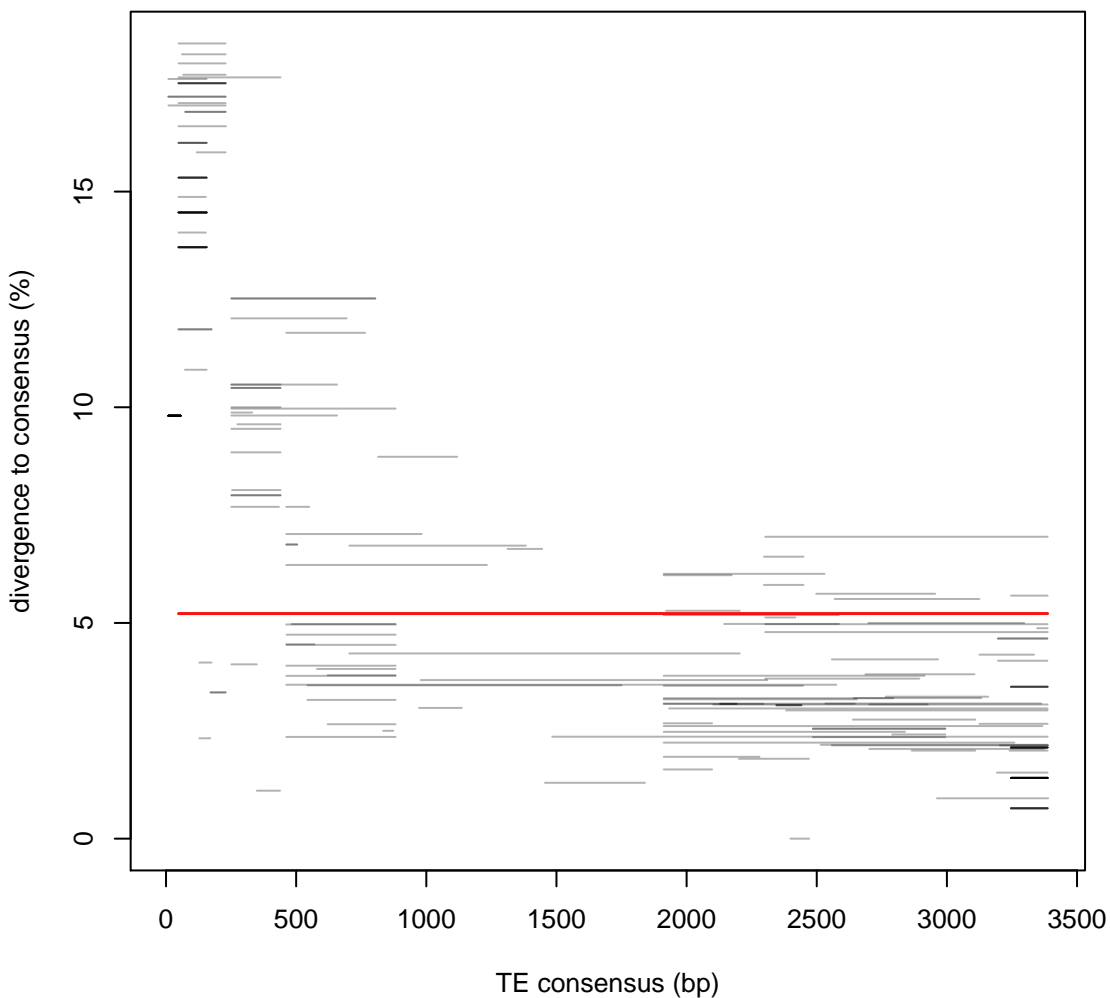
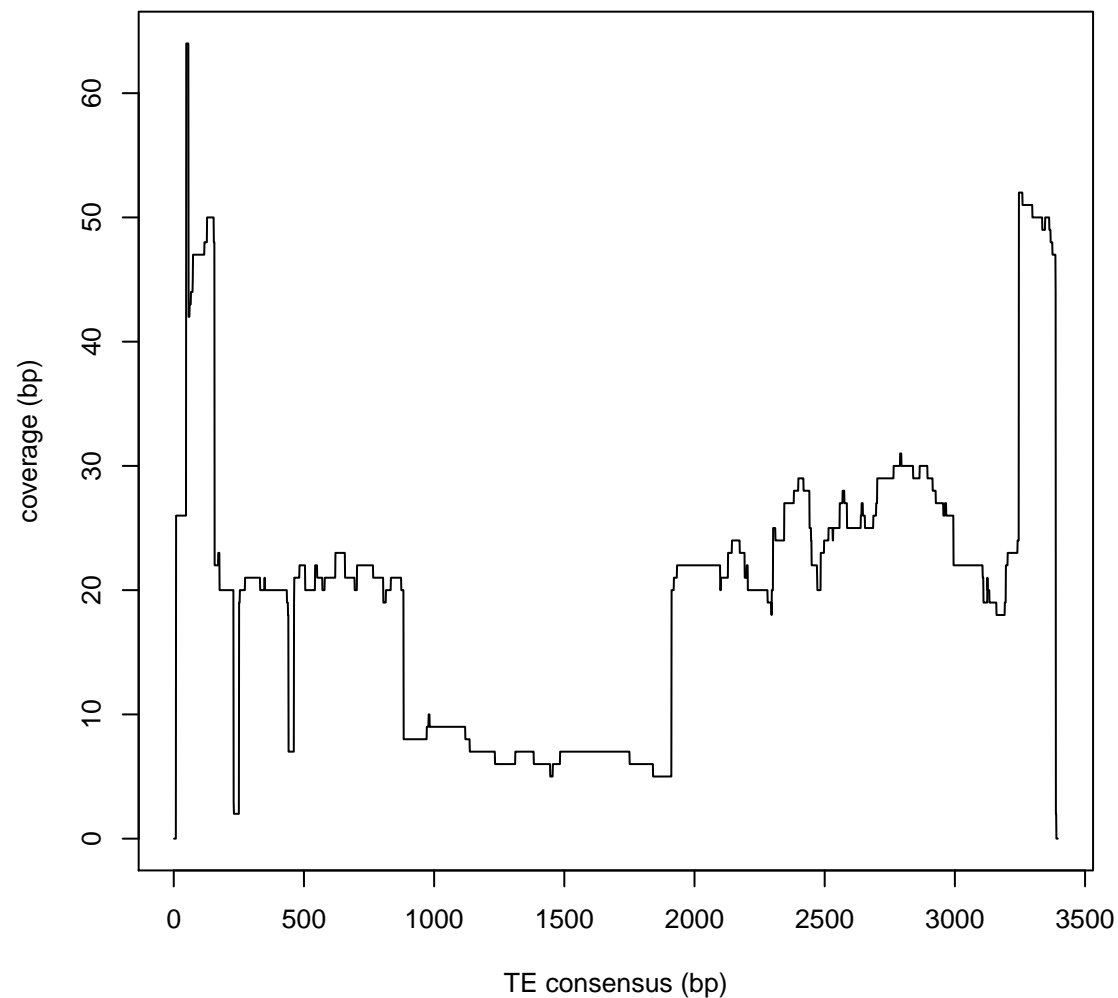


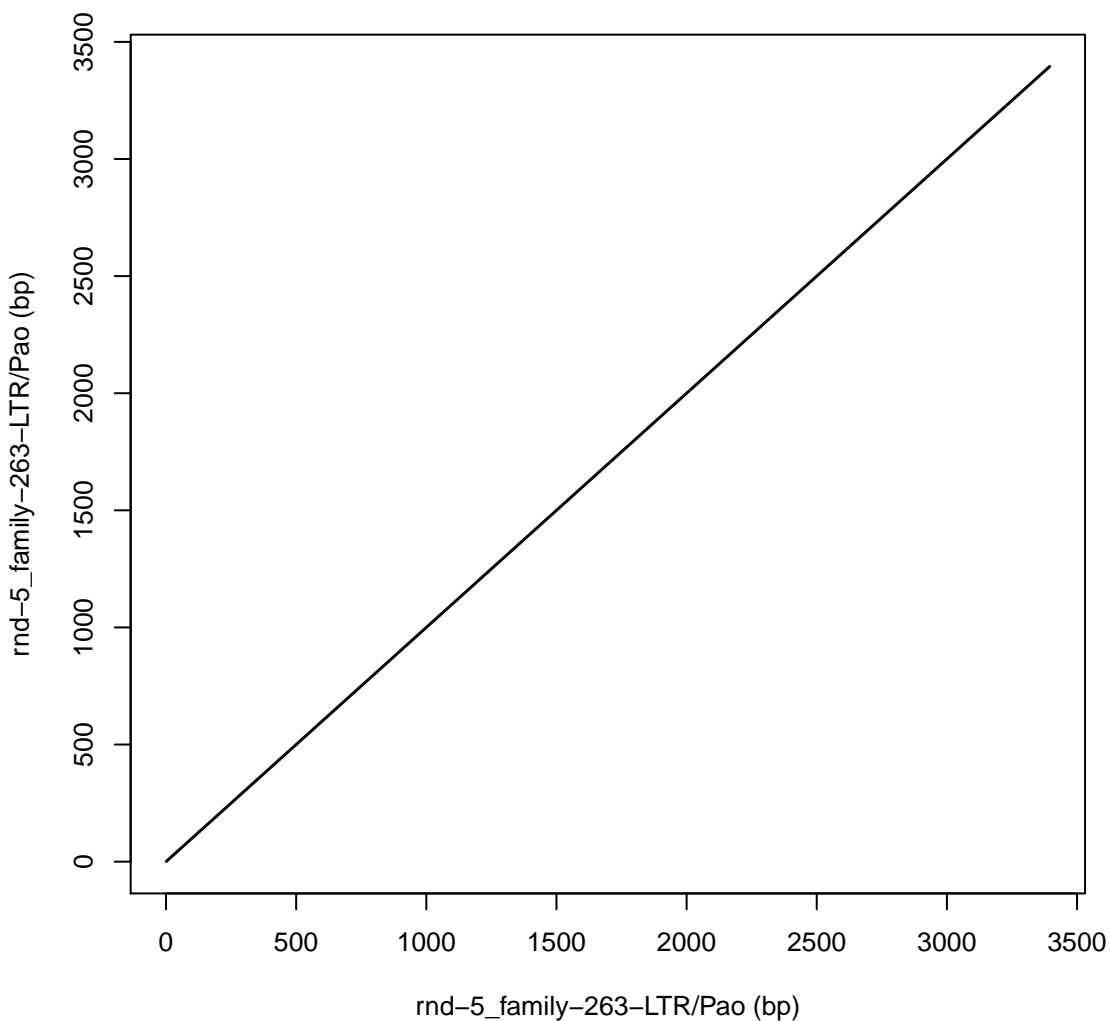
TE: rnd-5_family-263-LTR/Pao
consensus size: 3395bp; fragments: 213; full length: 1 (≥ 3055.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

