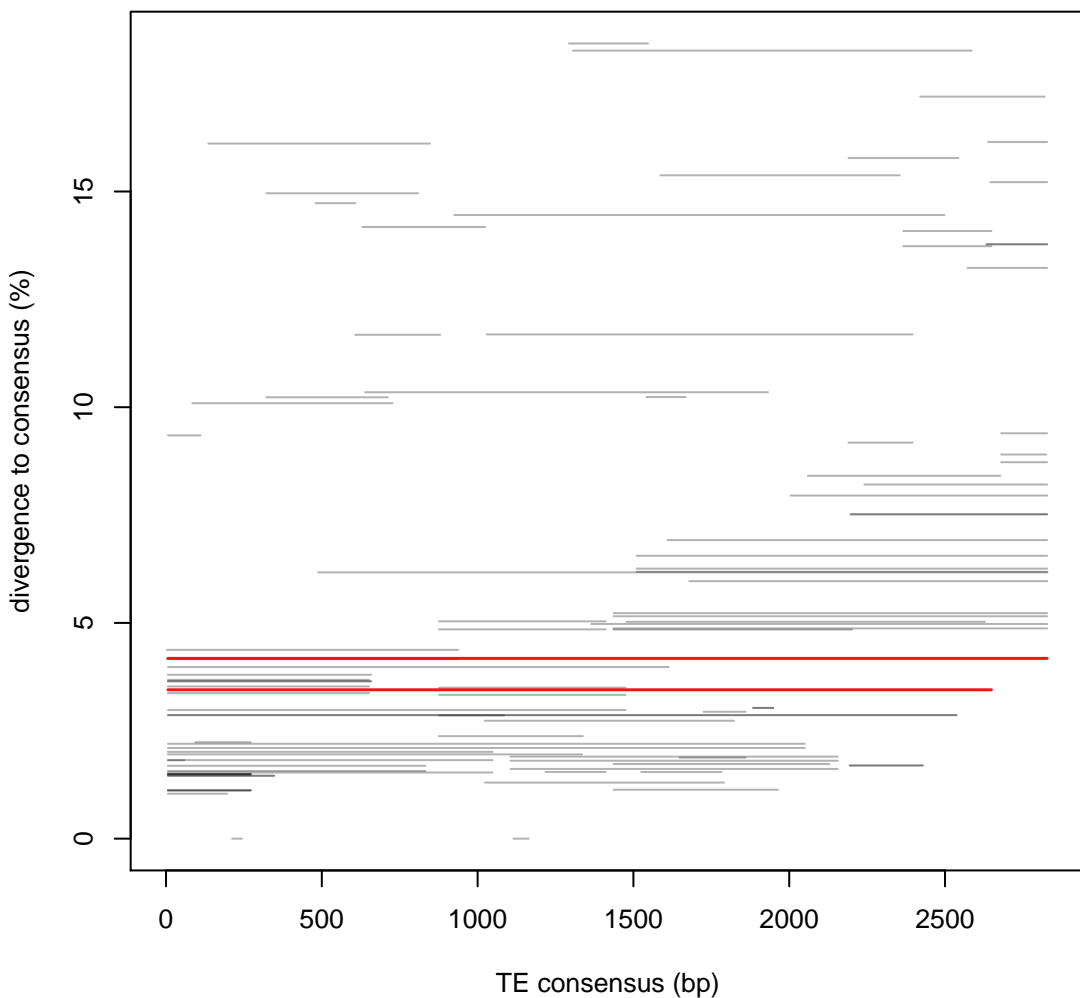
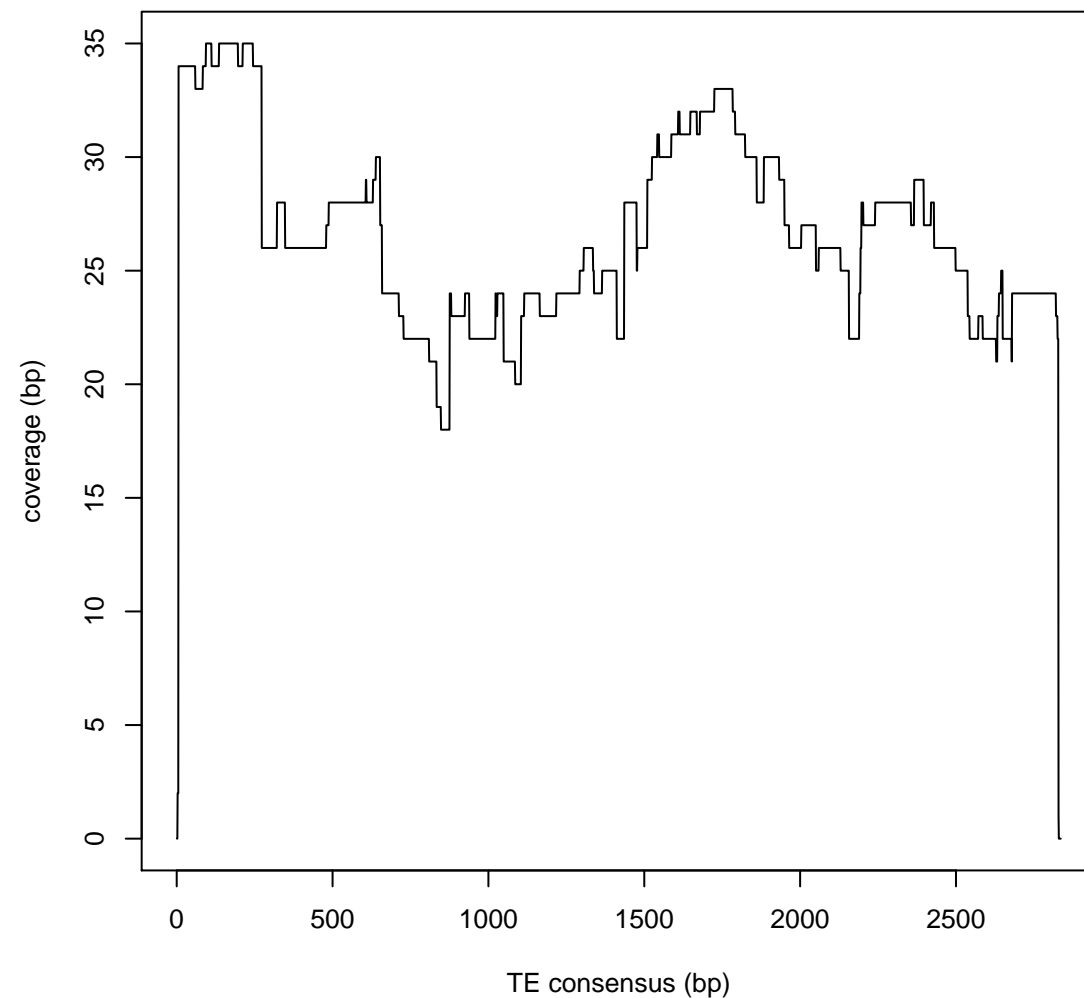


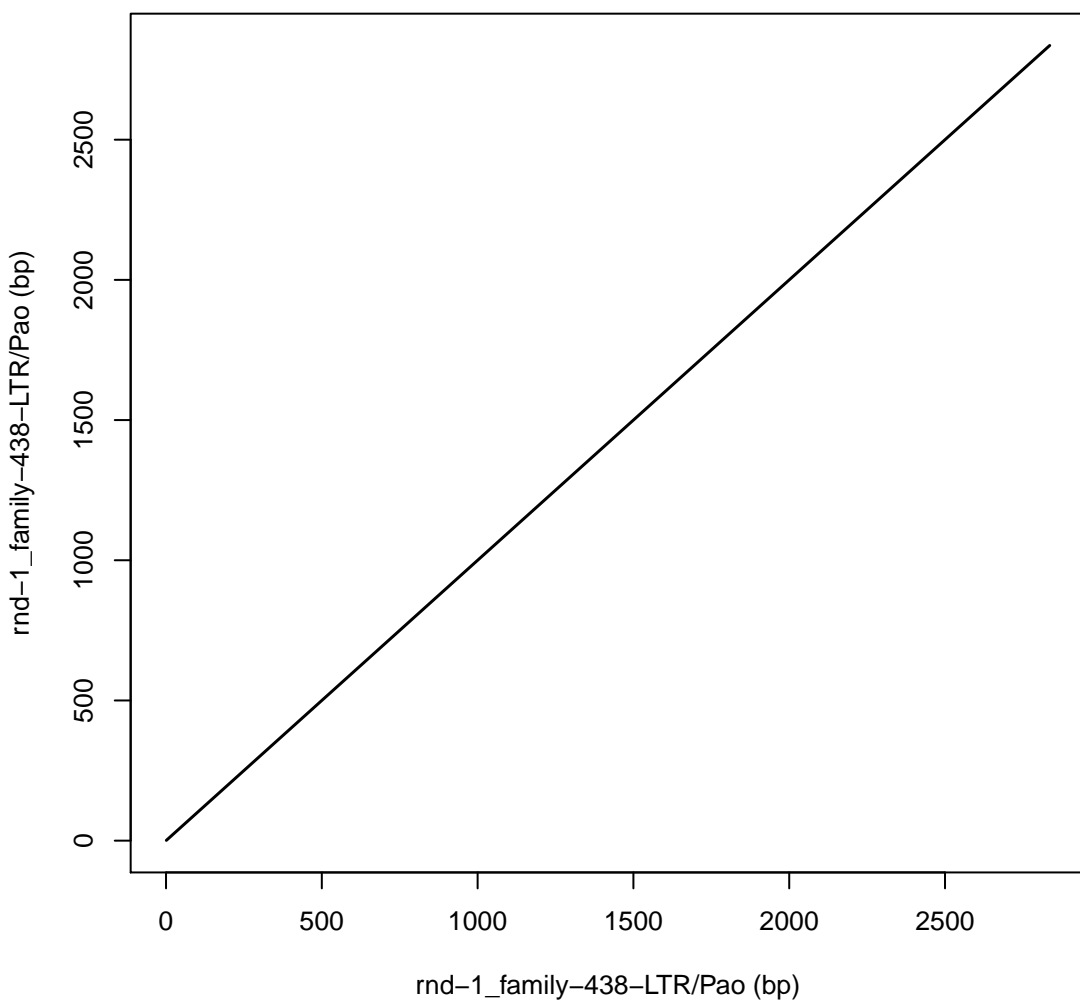
TE: rnd-1_family-438-LTR/Pao
consensus size: 2836bp; fragments: 102; full length: 2 (≥ 2552.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

