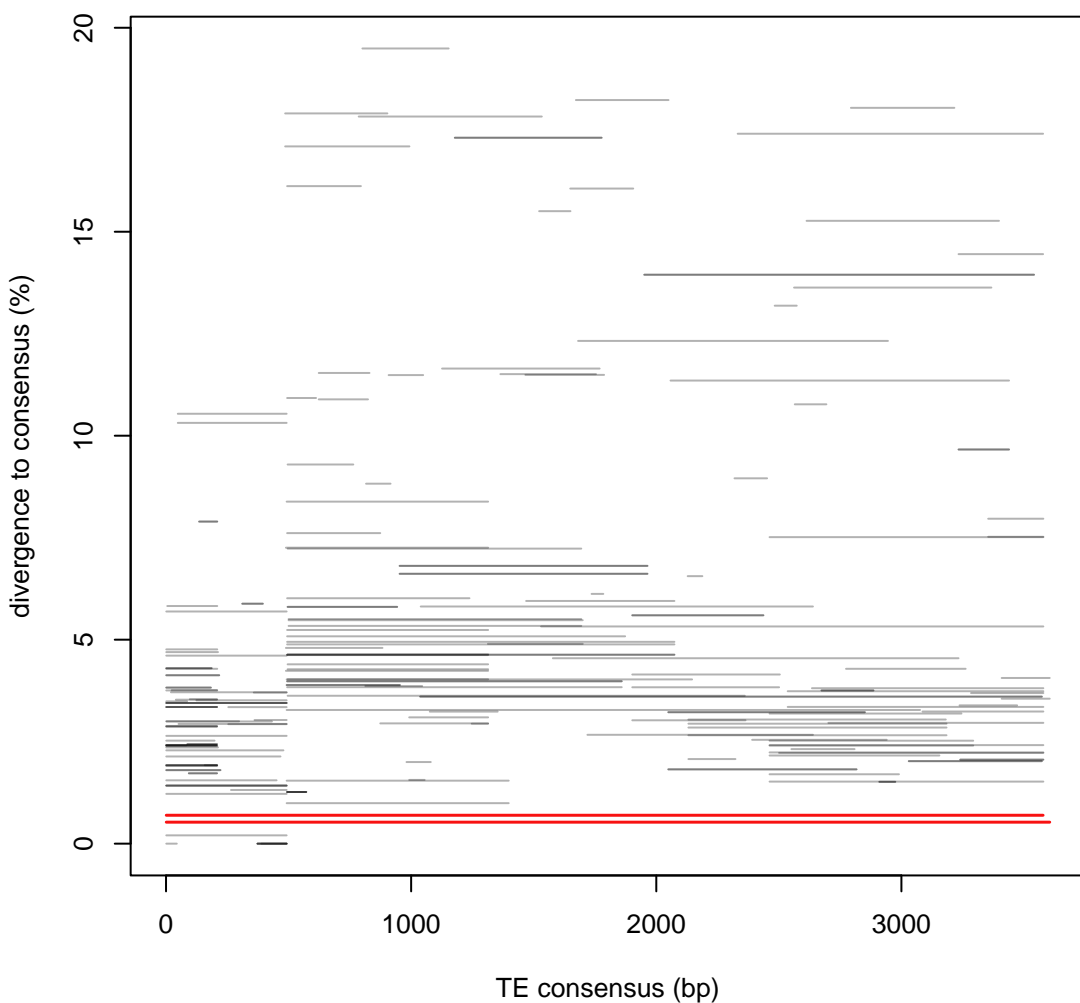
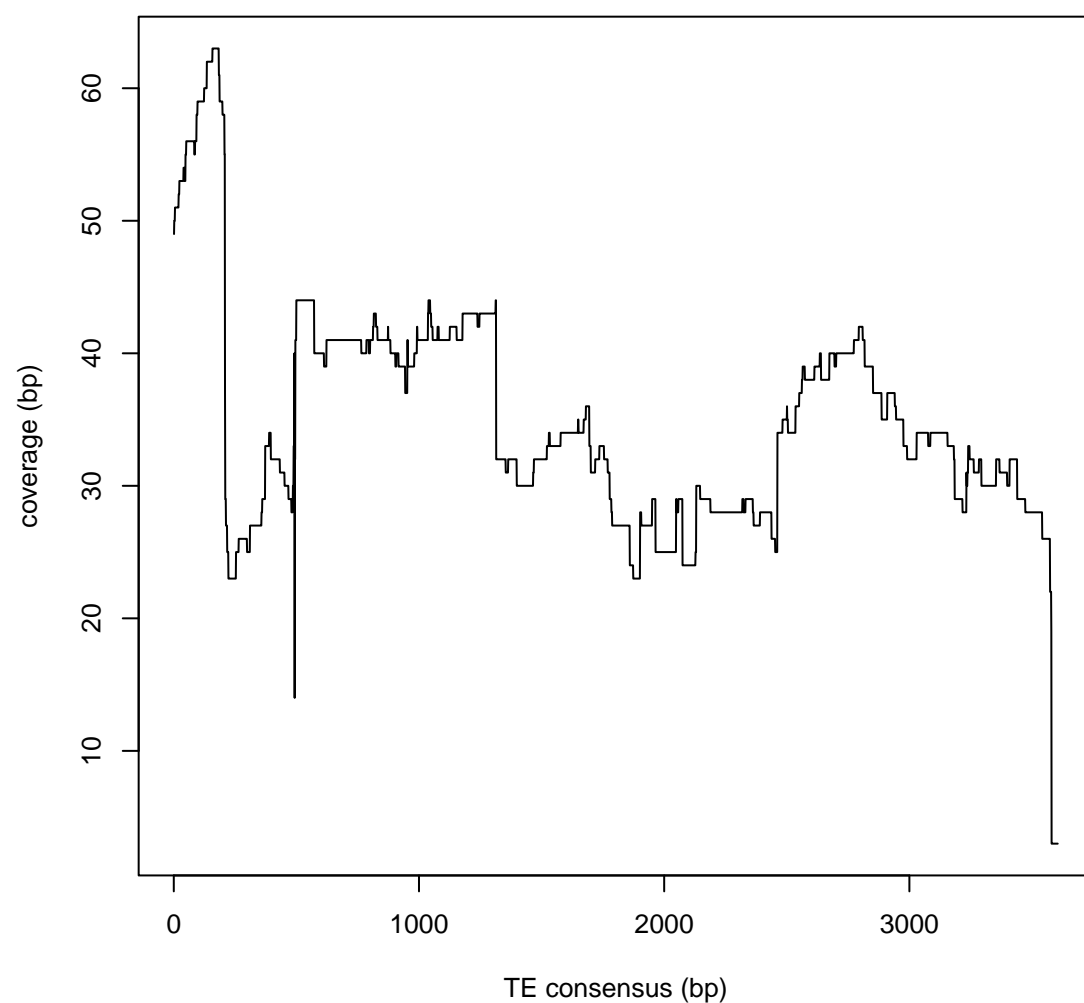


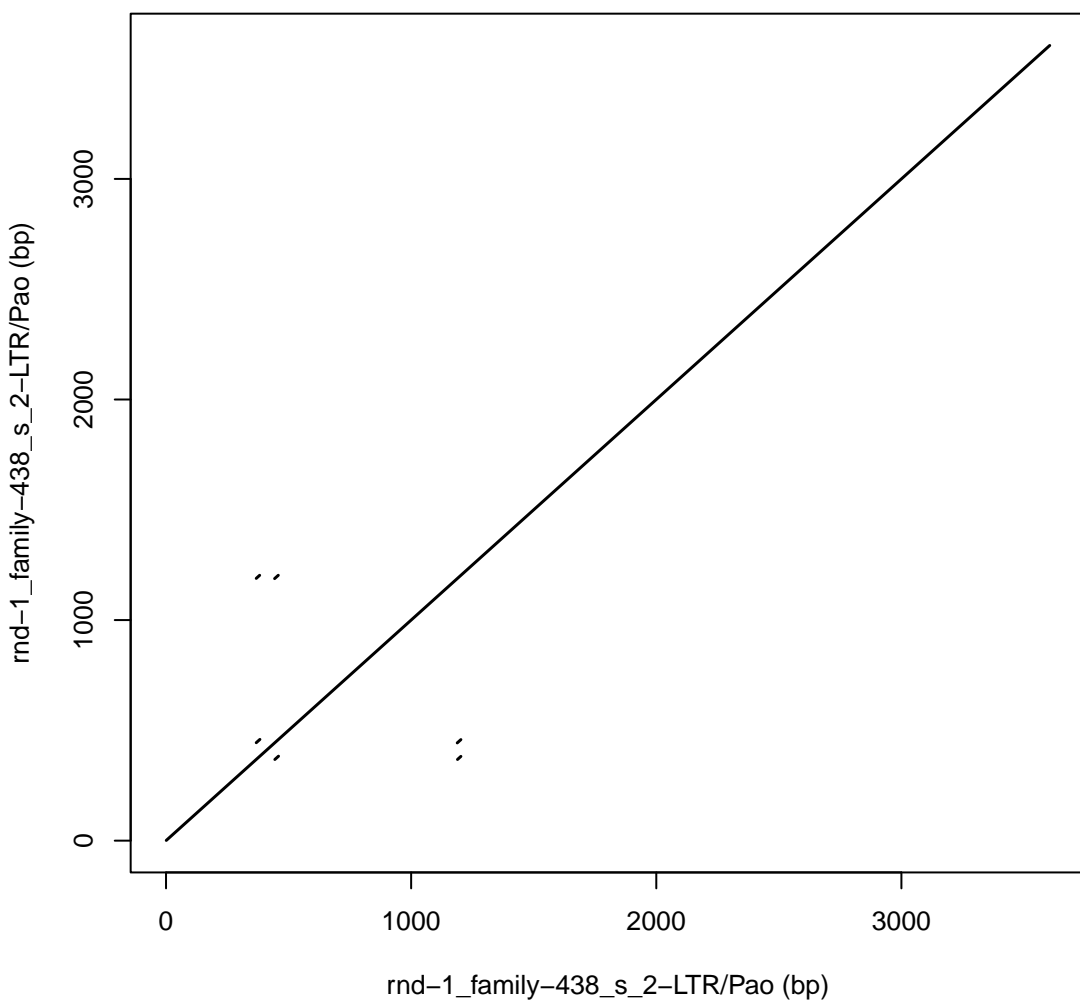
TE: rnd-1_family-438_s_2-LTR/Pao
consensus size: 3605bp; fragments: 215; full length: 2 (>=3244.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

