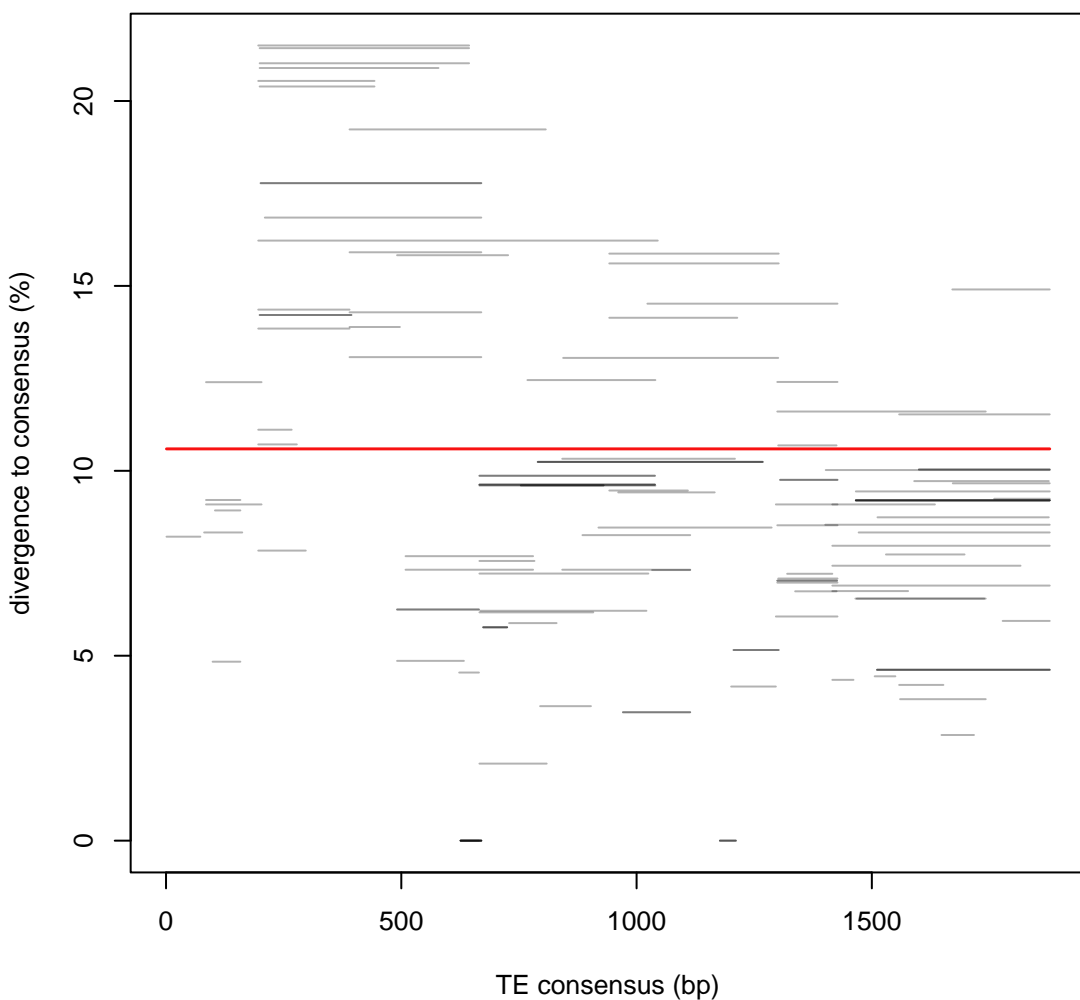
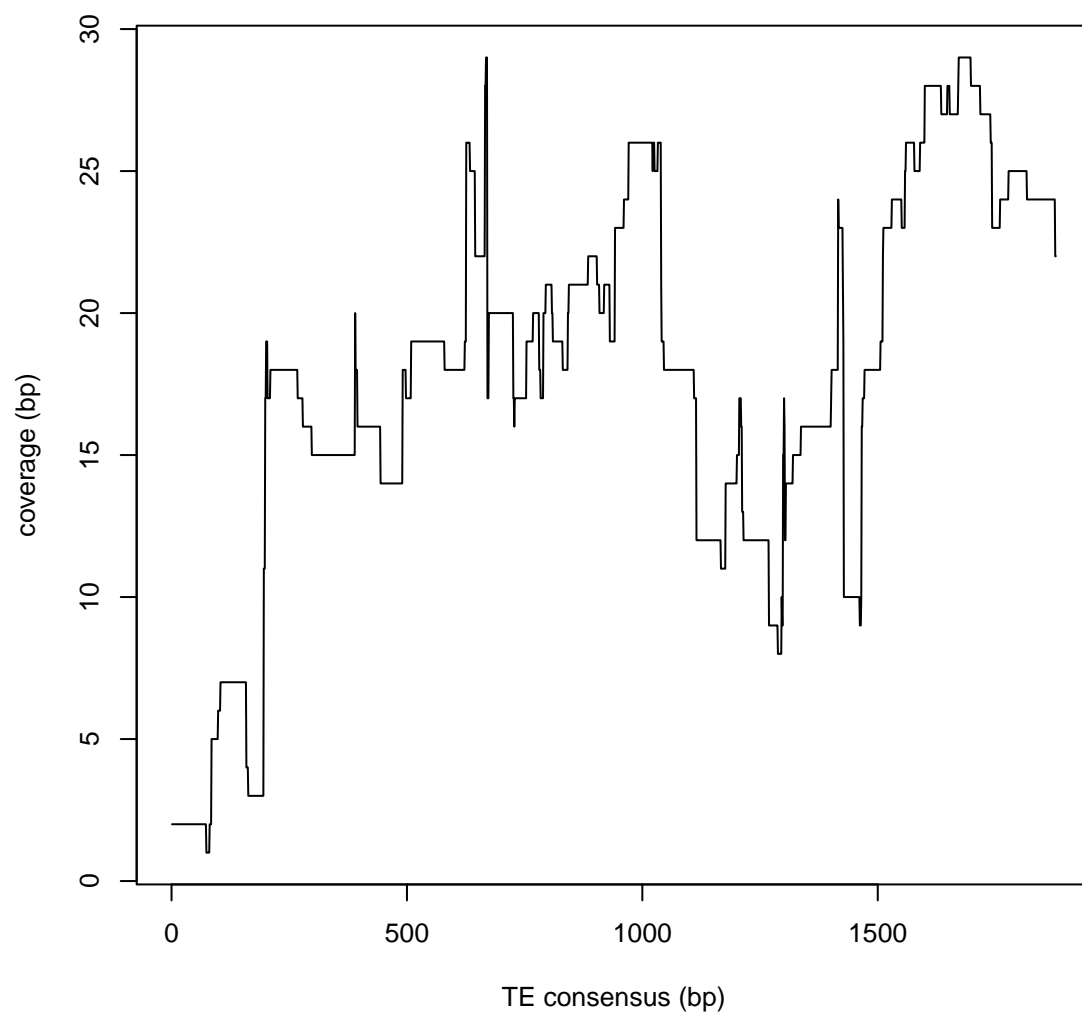


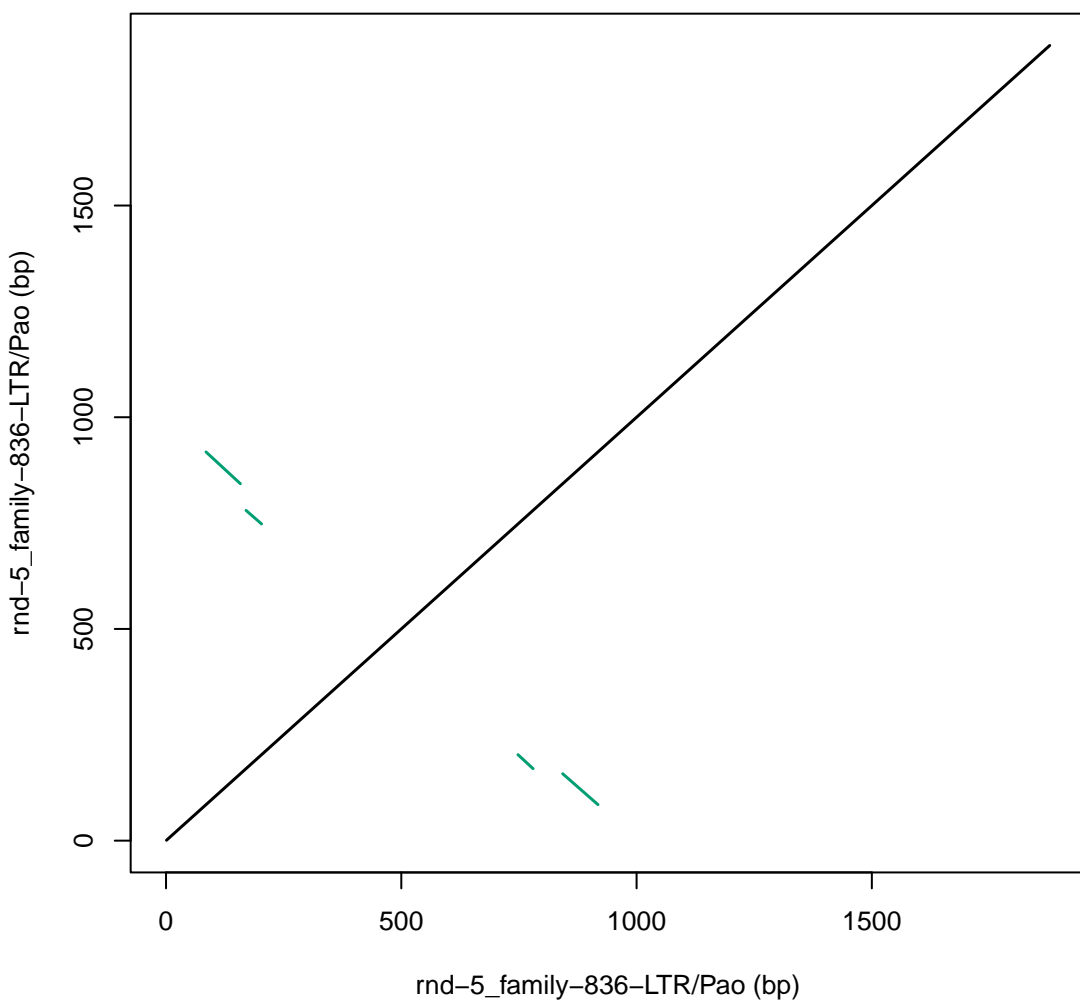
TE: rnd-5_family-836-LTR/Pao
consensus size: 1878bp; fragments: 134; full length: 1 (>=1690.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

