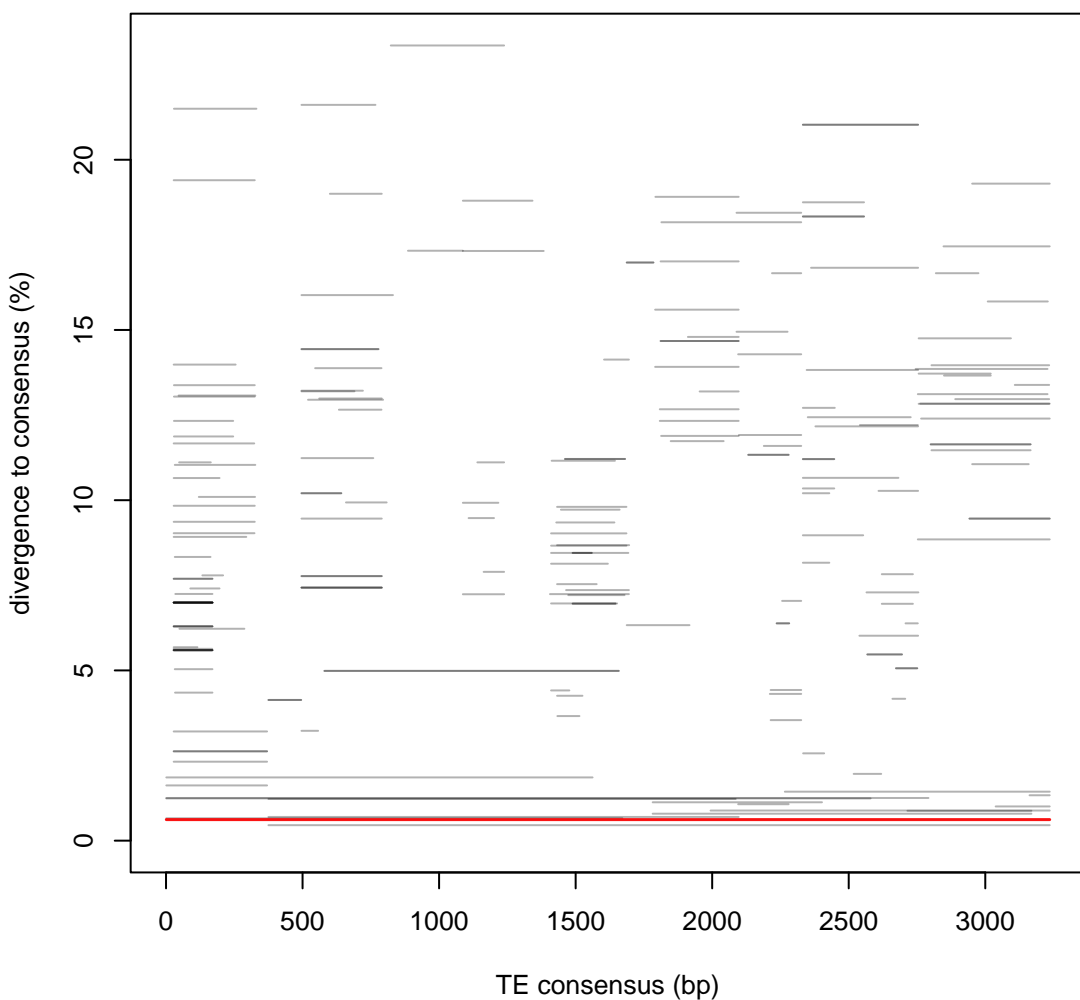
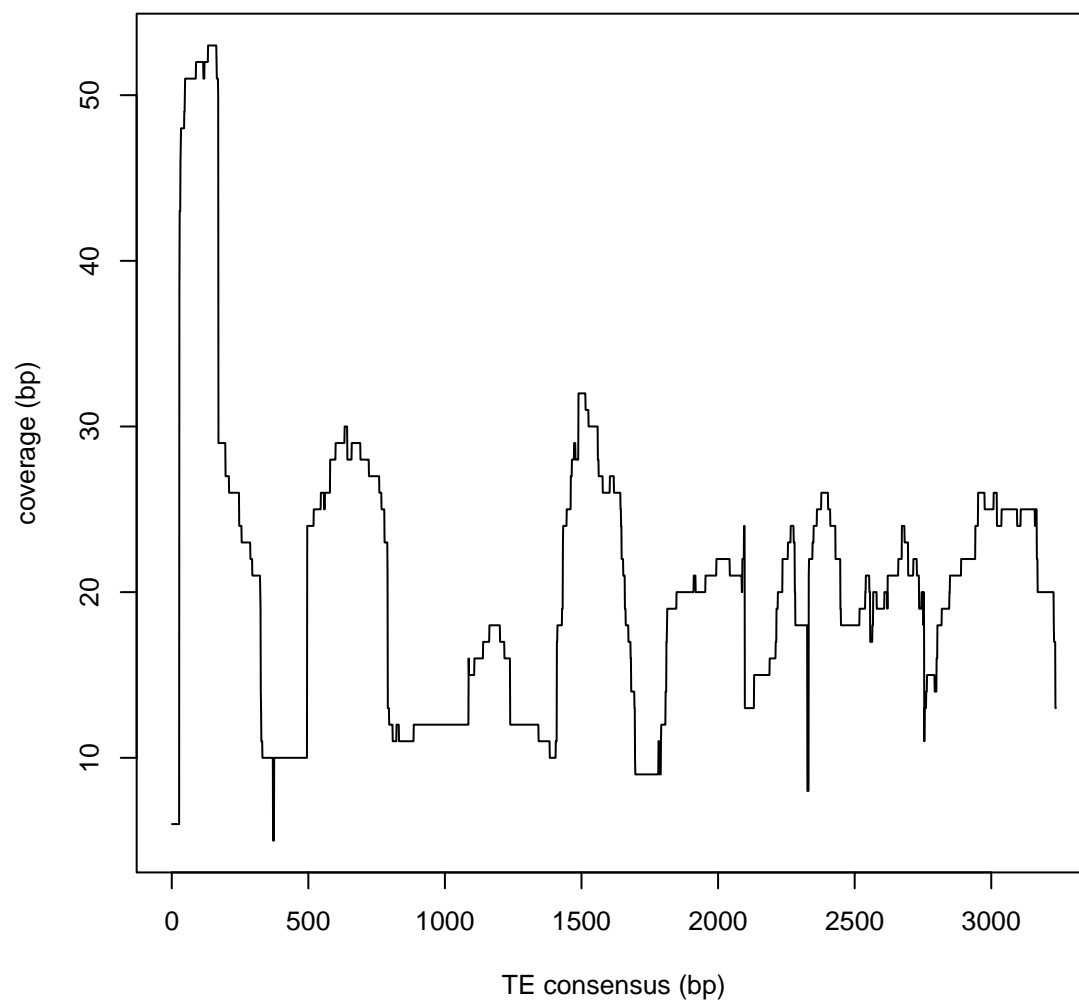


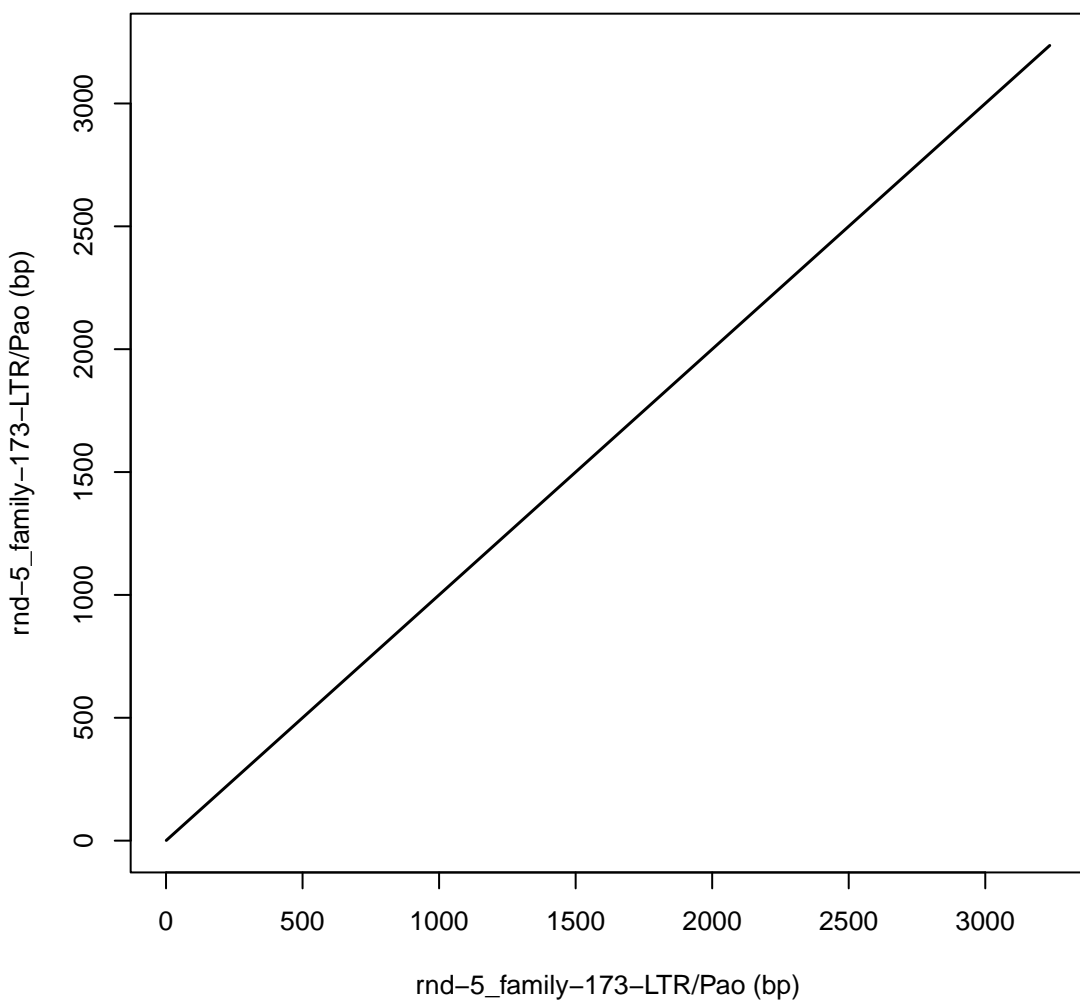
TE: rnd-5_family-173-LTR/Pao
consensus size: 3236bp; fragments: 207; full length: 1 (>=2912.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

