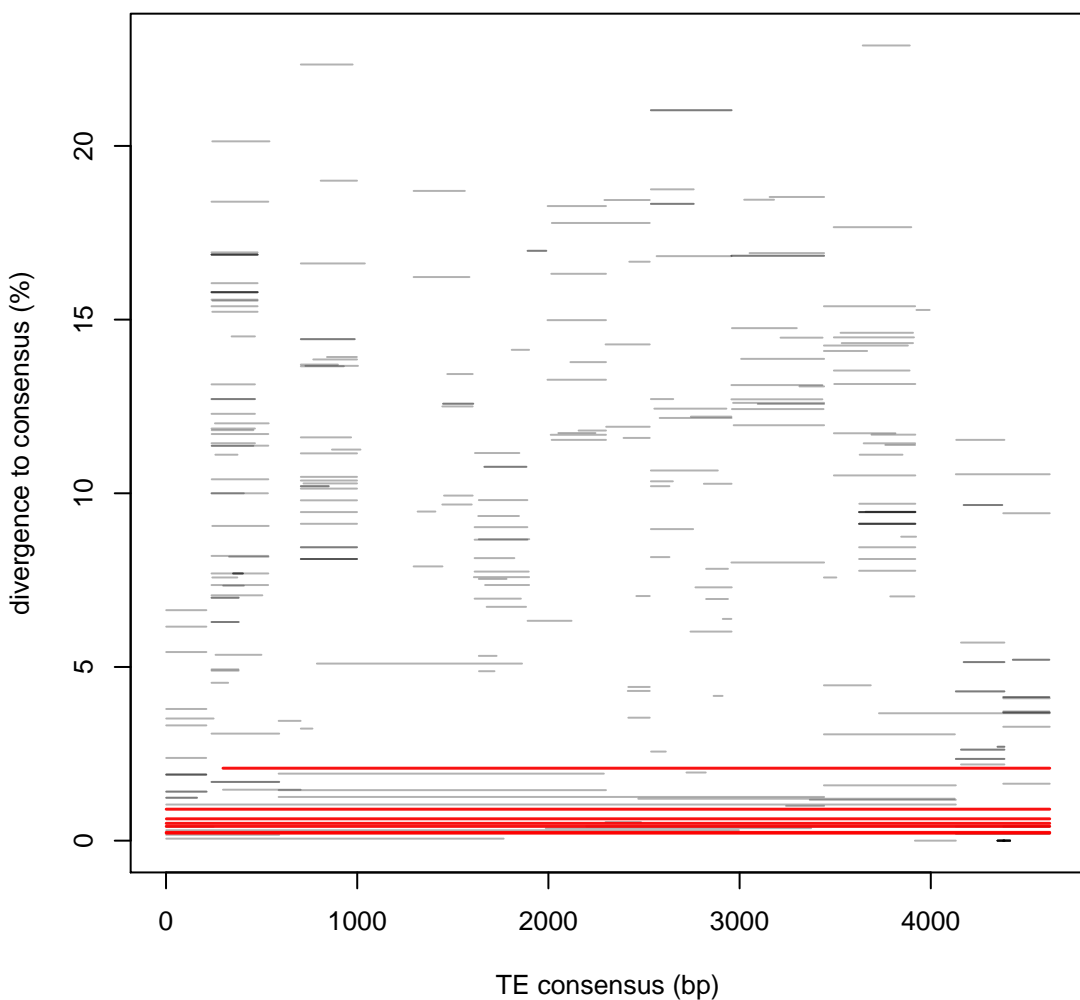
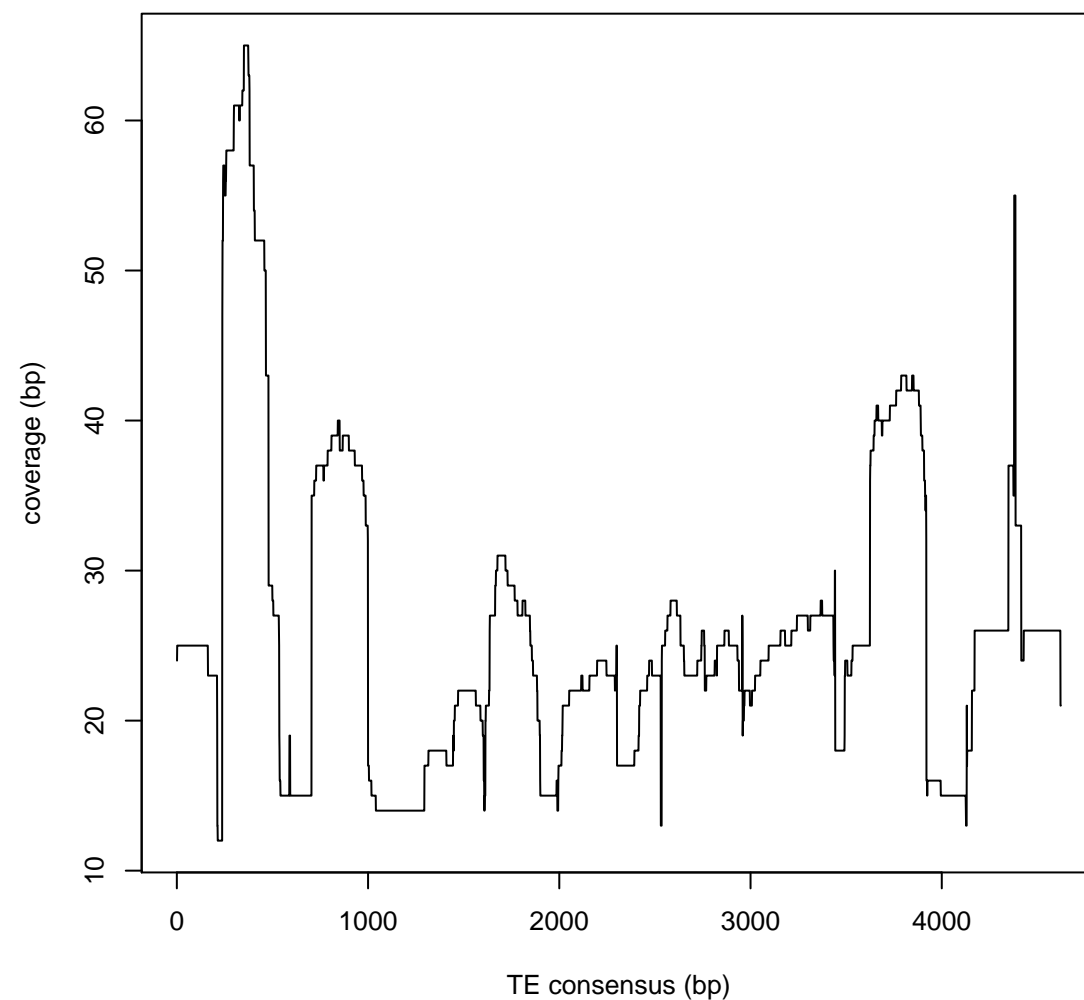


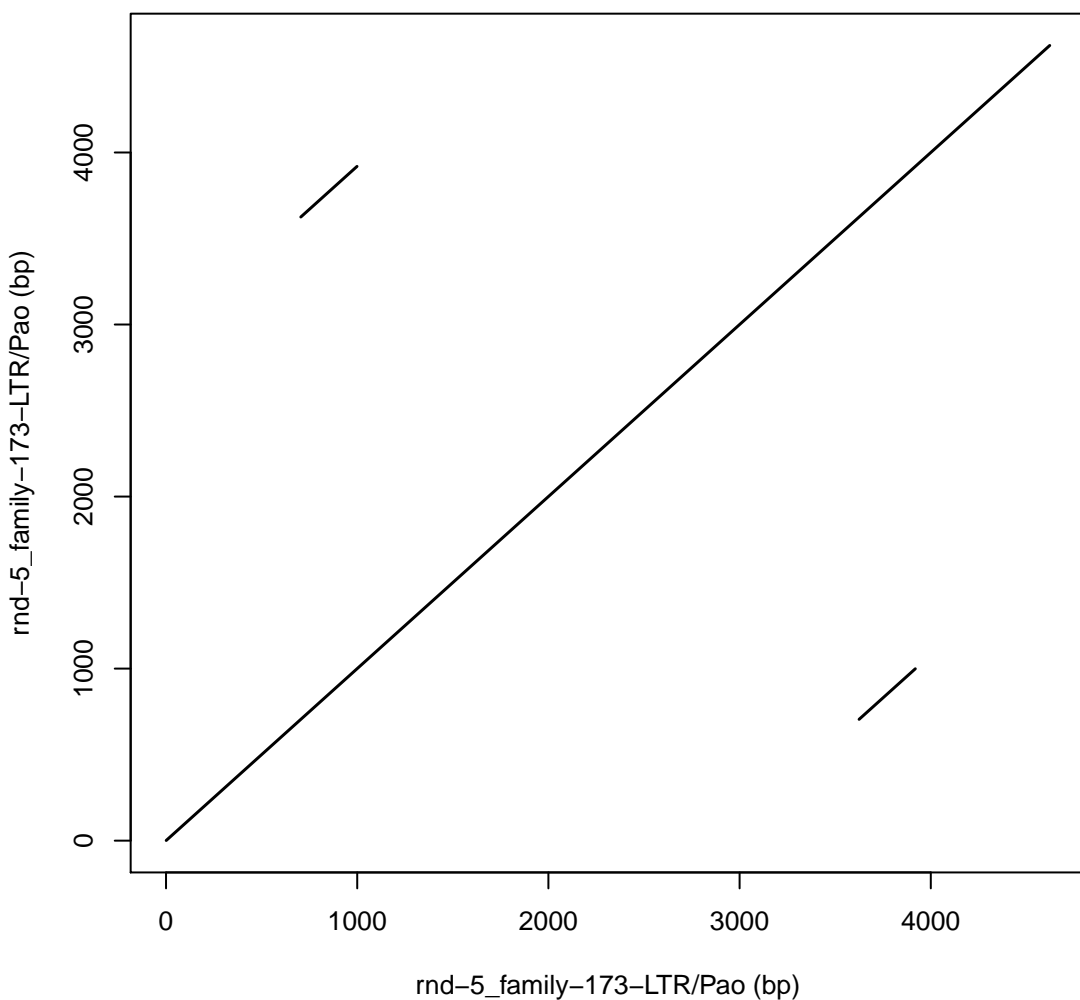
TE: rnd-5_family-173-LTR/Pao
consensus size: 4622bp; fragments: 298; full length: 7 (>=4159.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

