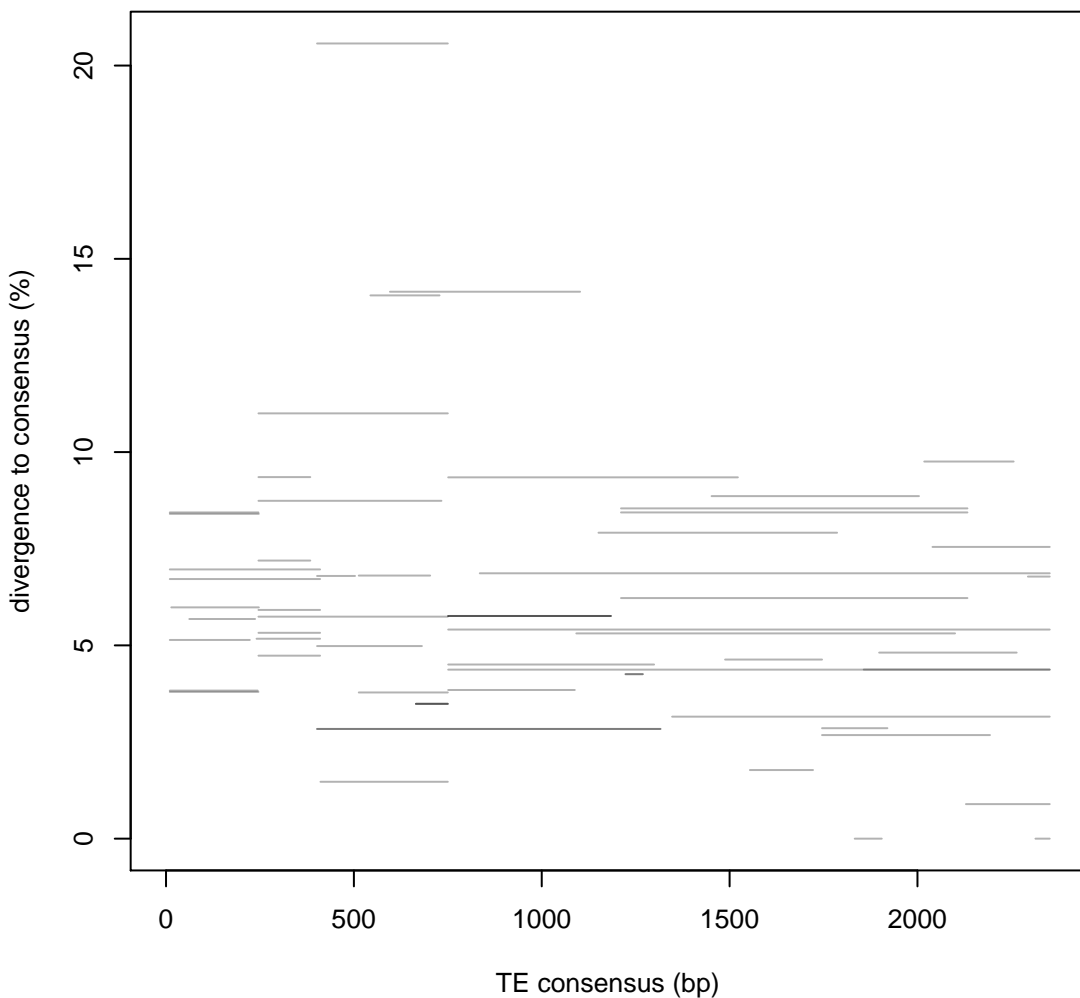
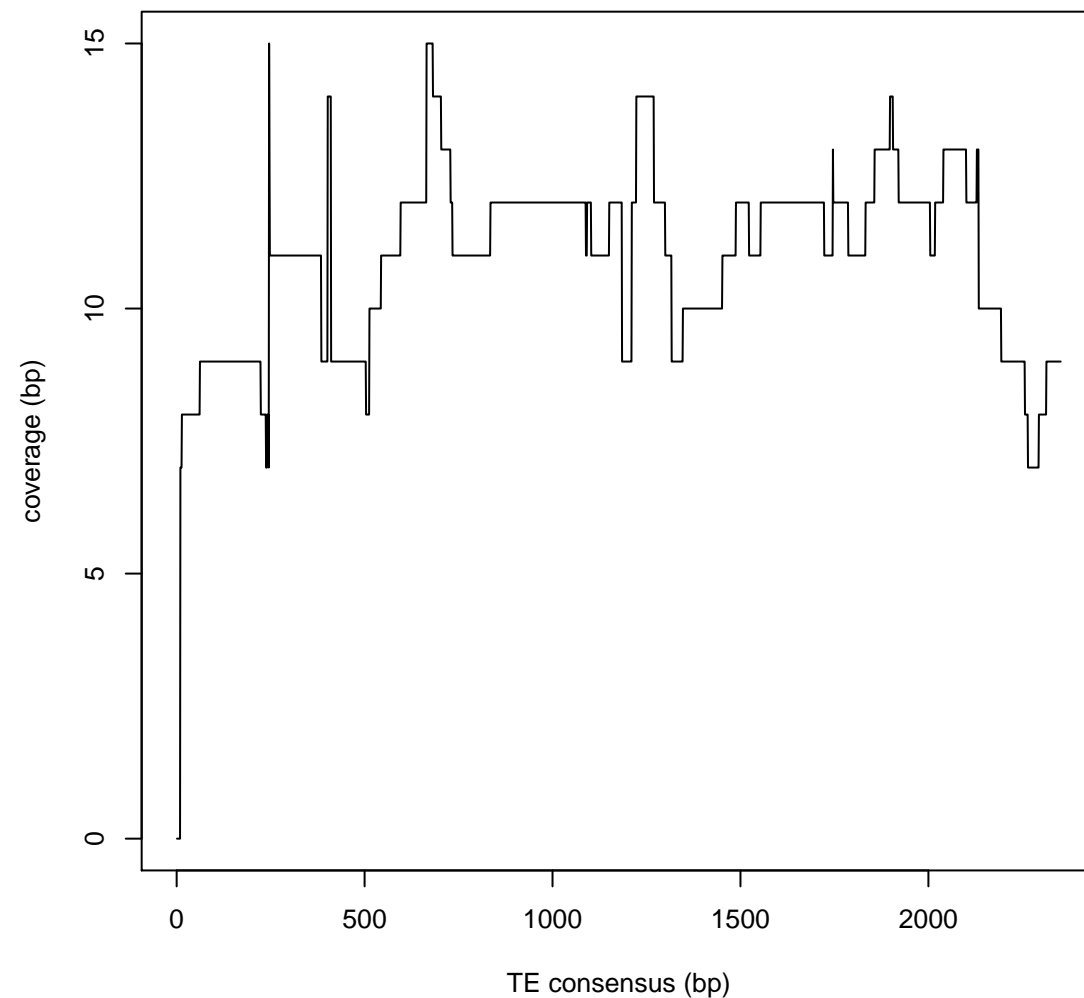


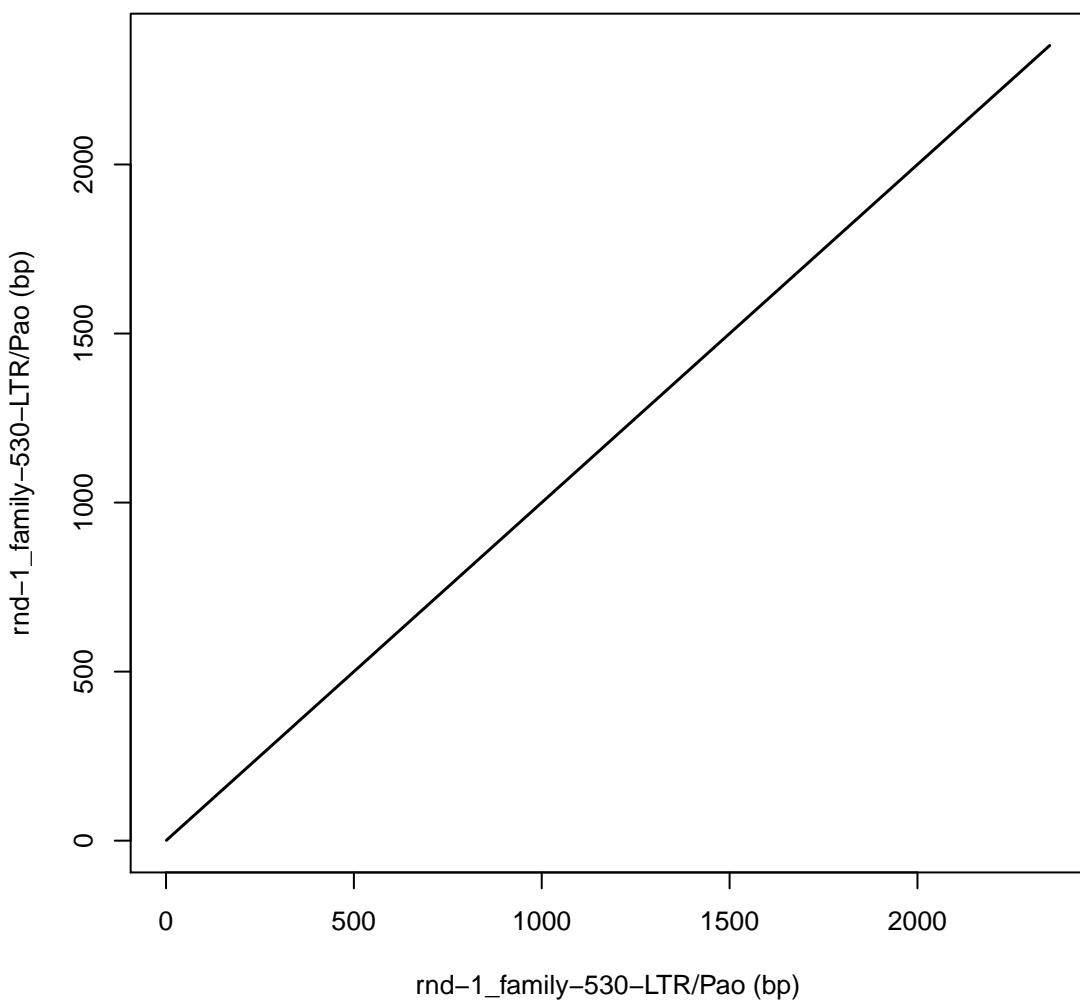
TE: rnd-1\_family-530-LTR/Pao  
consensus size: 2352bp; fragments: 61; full length: 0 (>=2116.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

