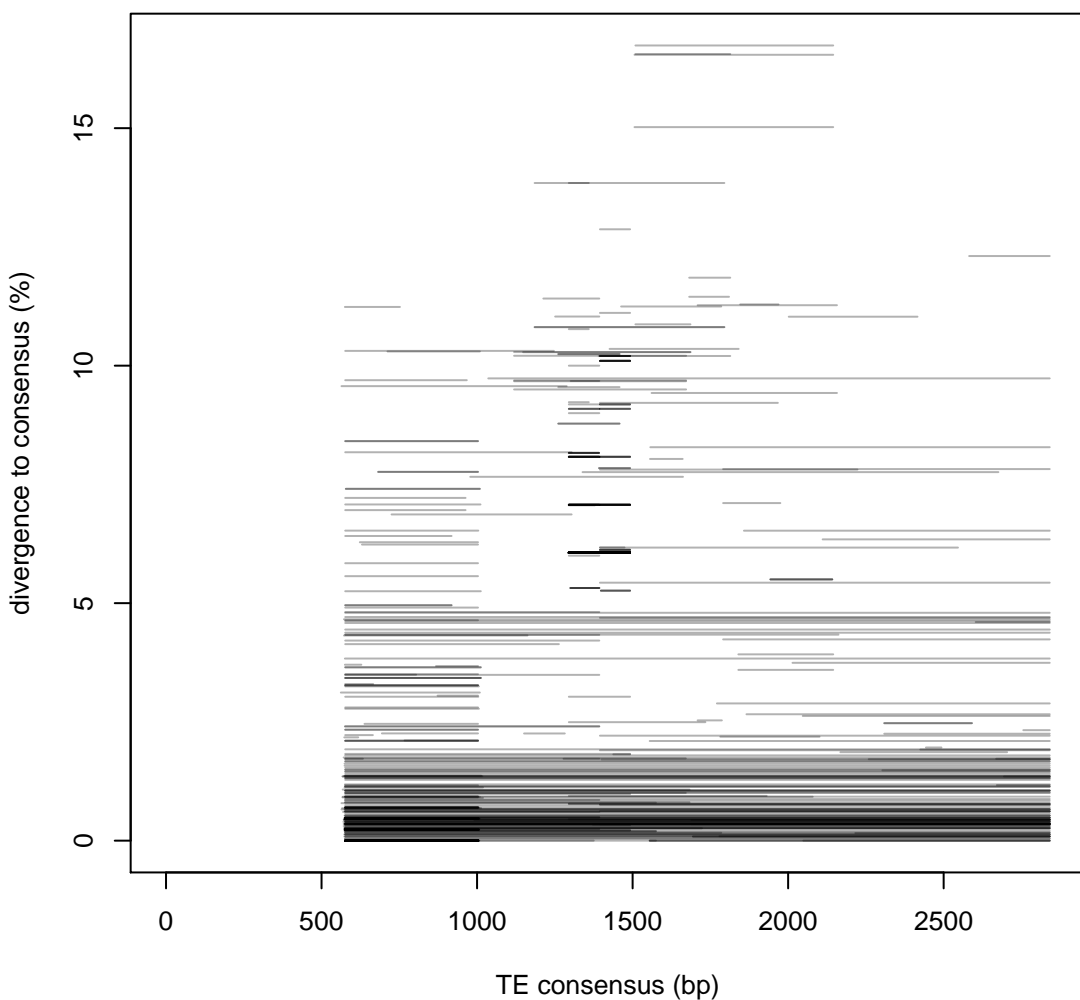
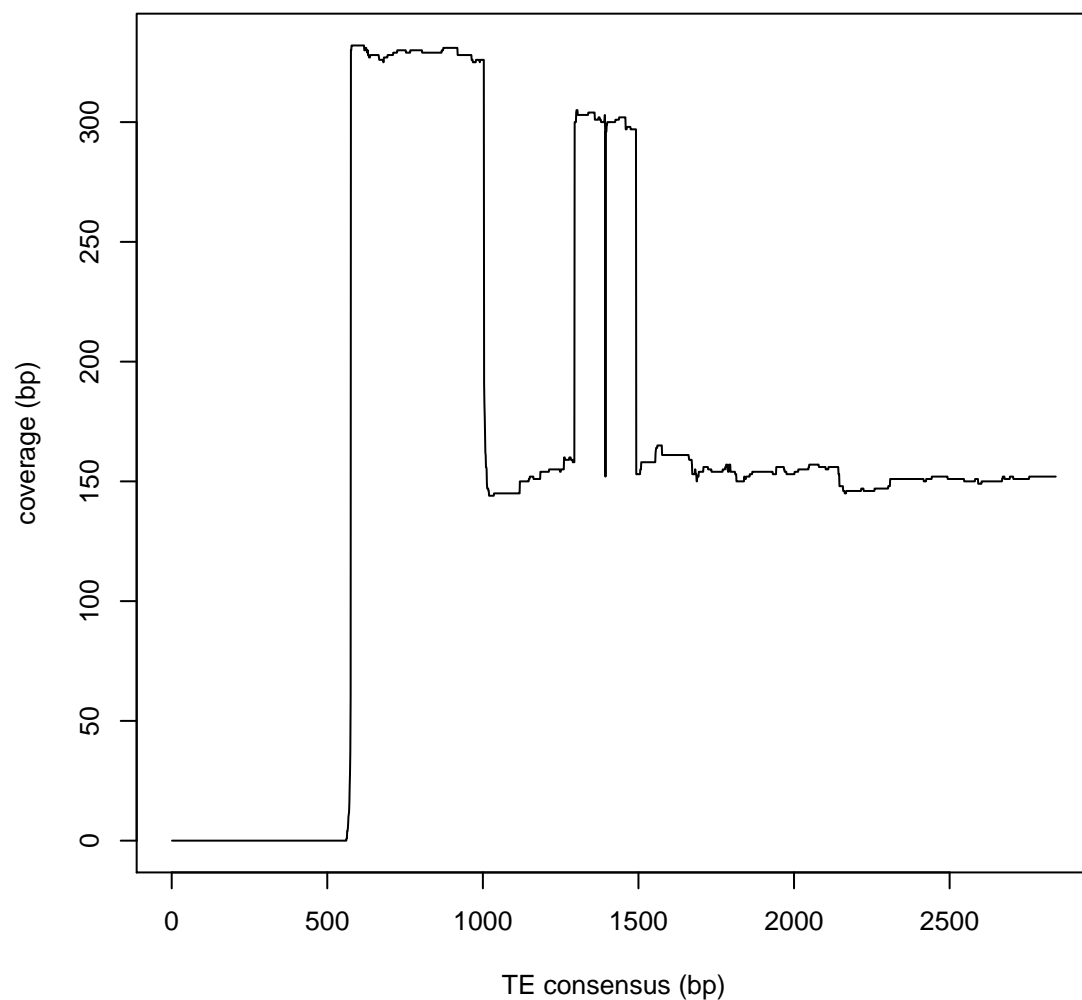


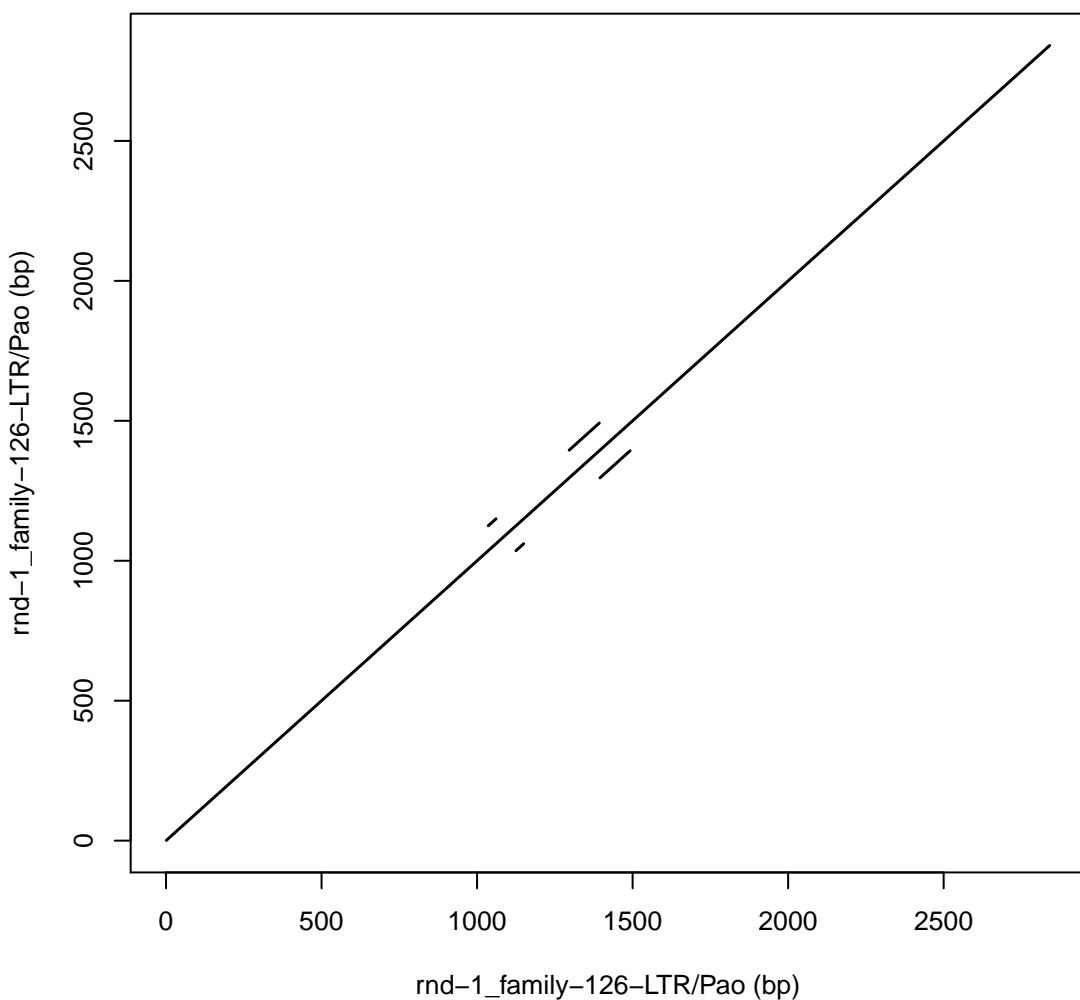
TE: rnd-1_family-126-LTR/Pao
consensus size: 2841bp; fragments: 721; full length: 0 (>=2556.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

