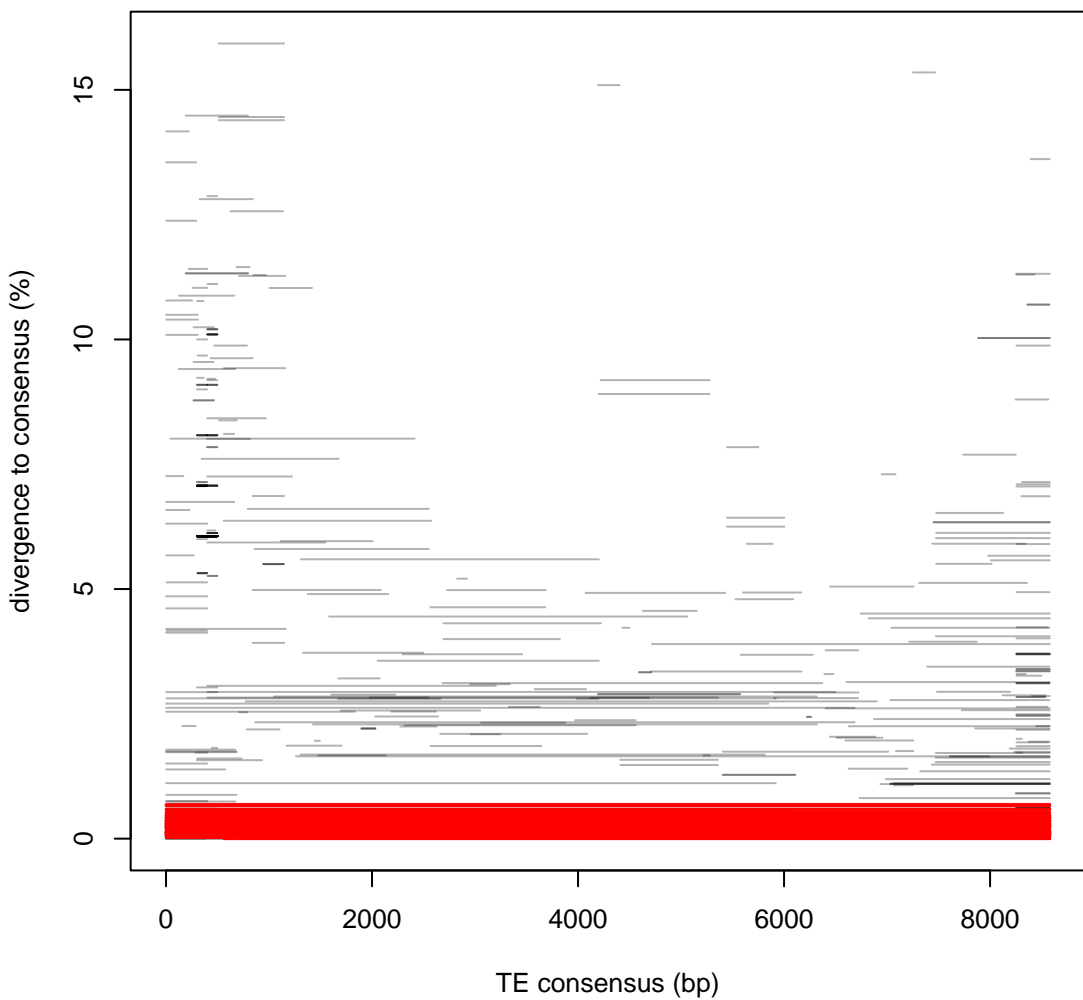
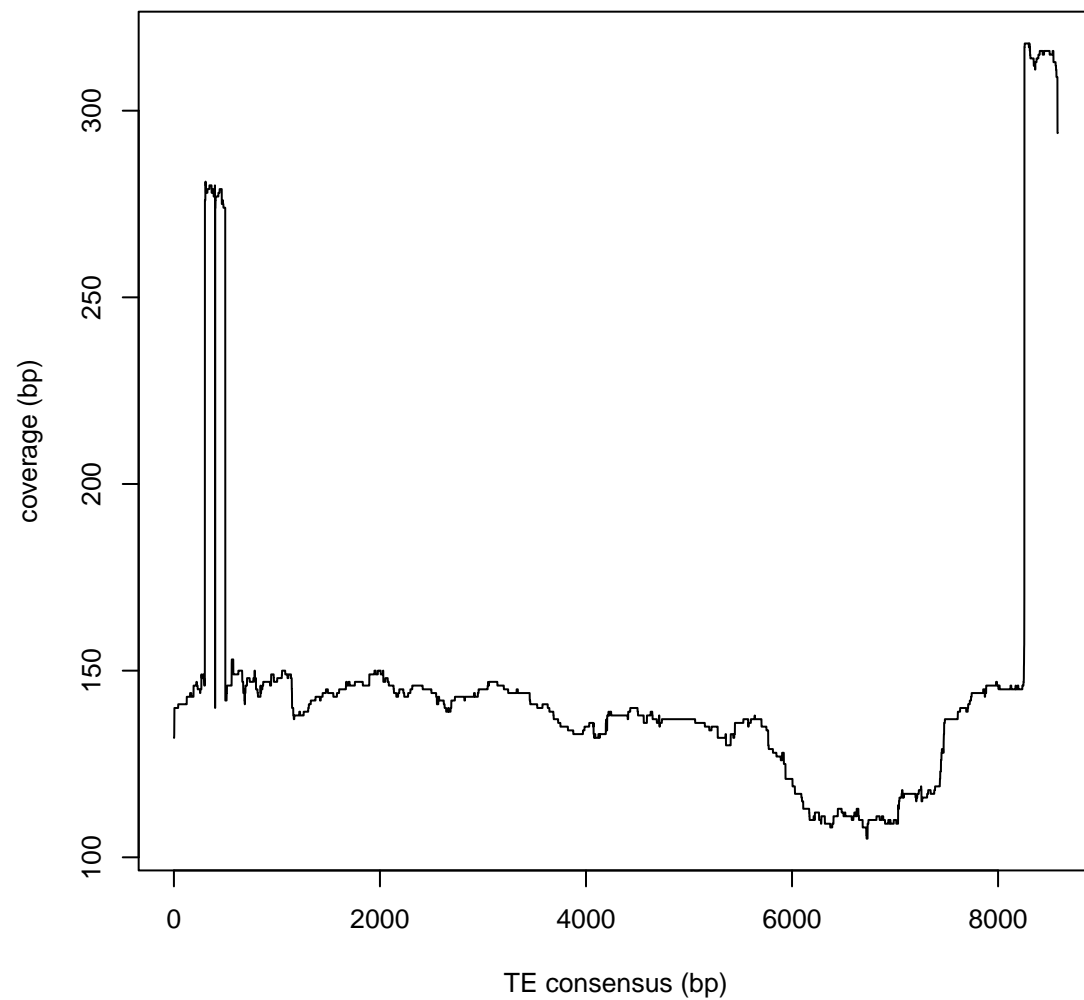


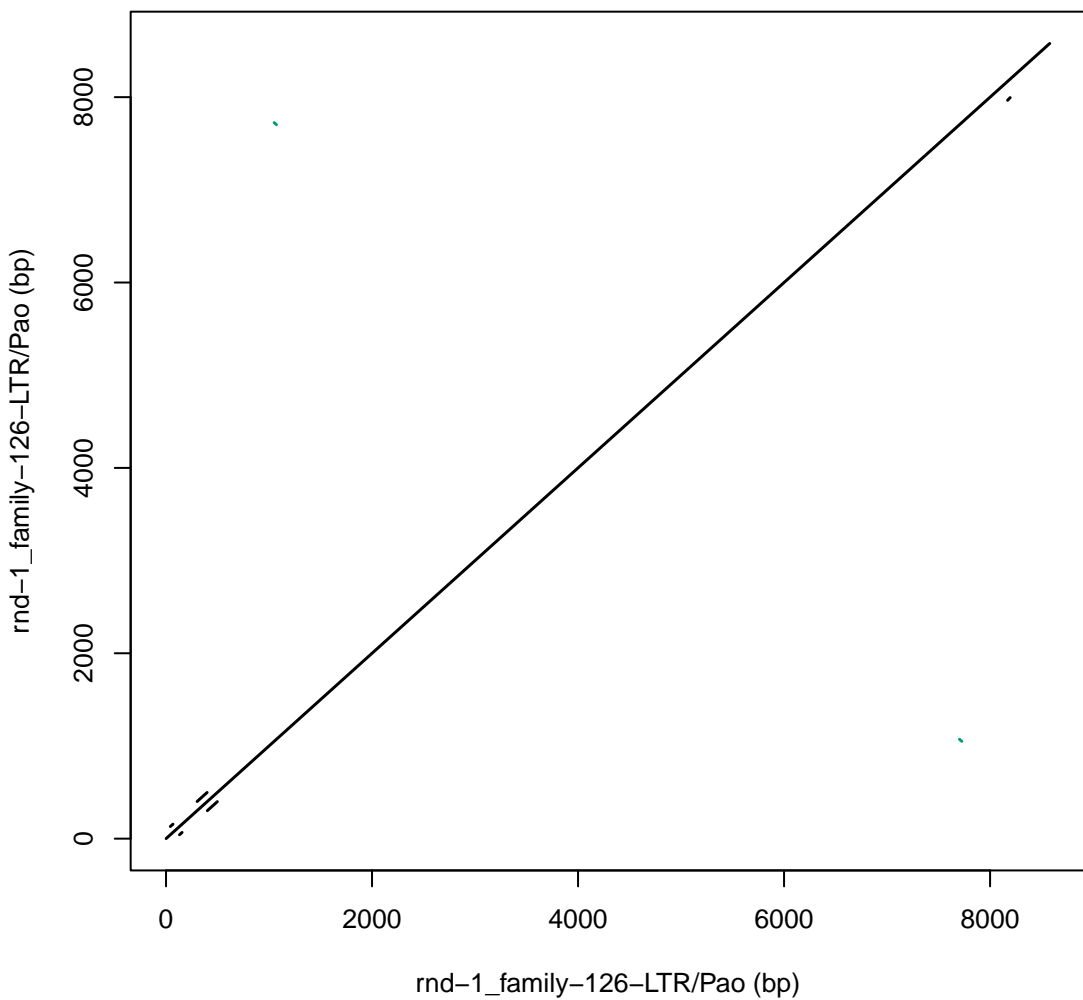
TE: rnd-1_family-126-LTR/Pao
consensus size: 8579bp; fragments: 823; full length: 72 (>=7721.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

