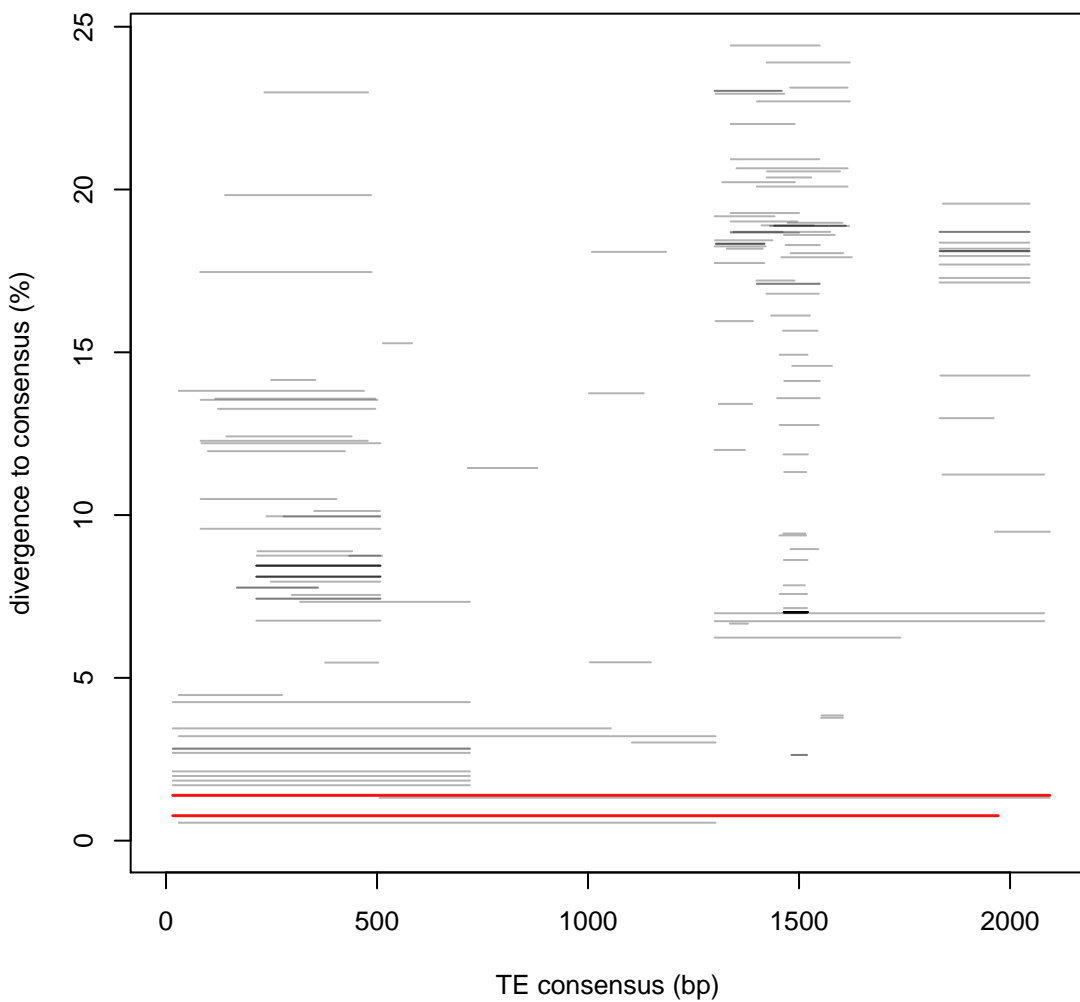
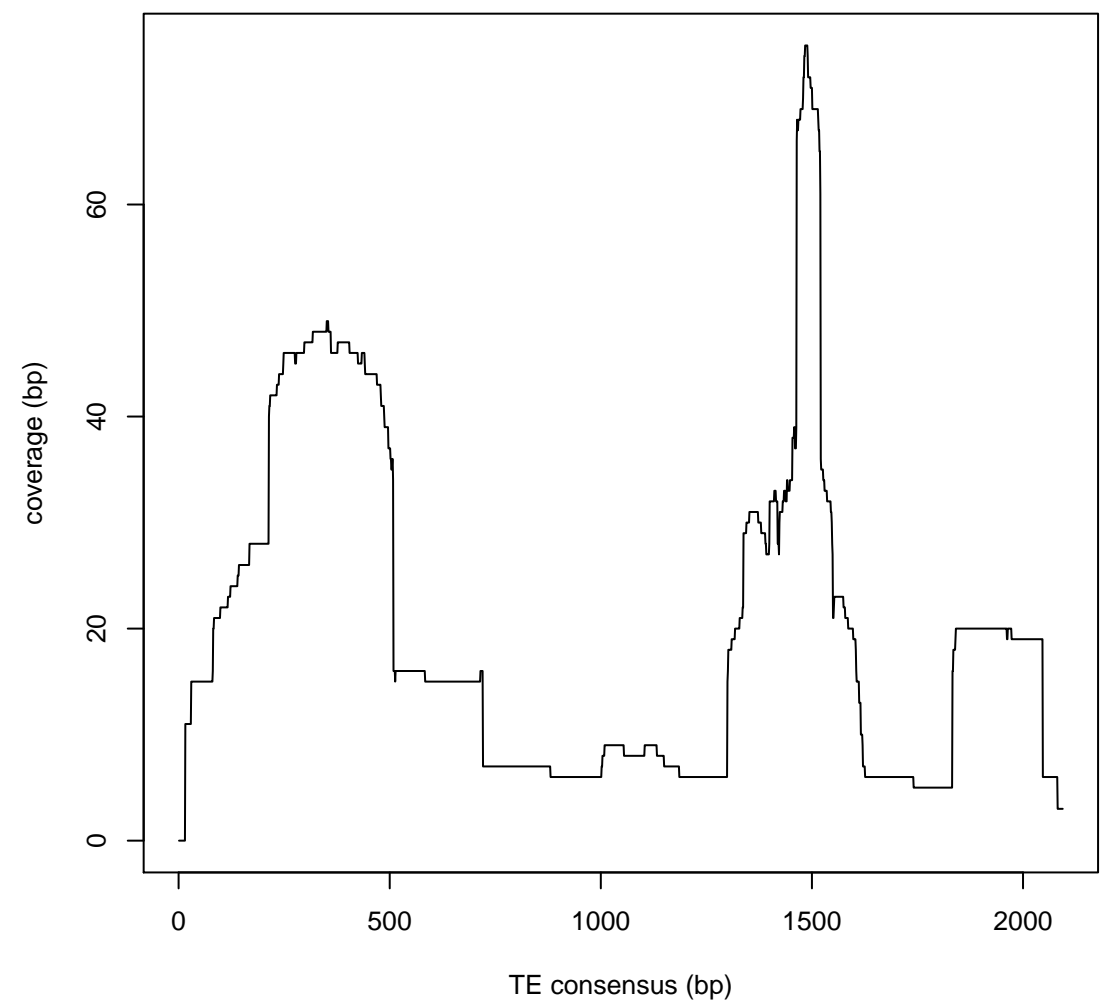


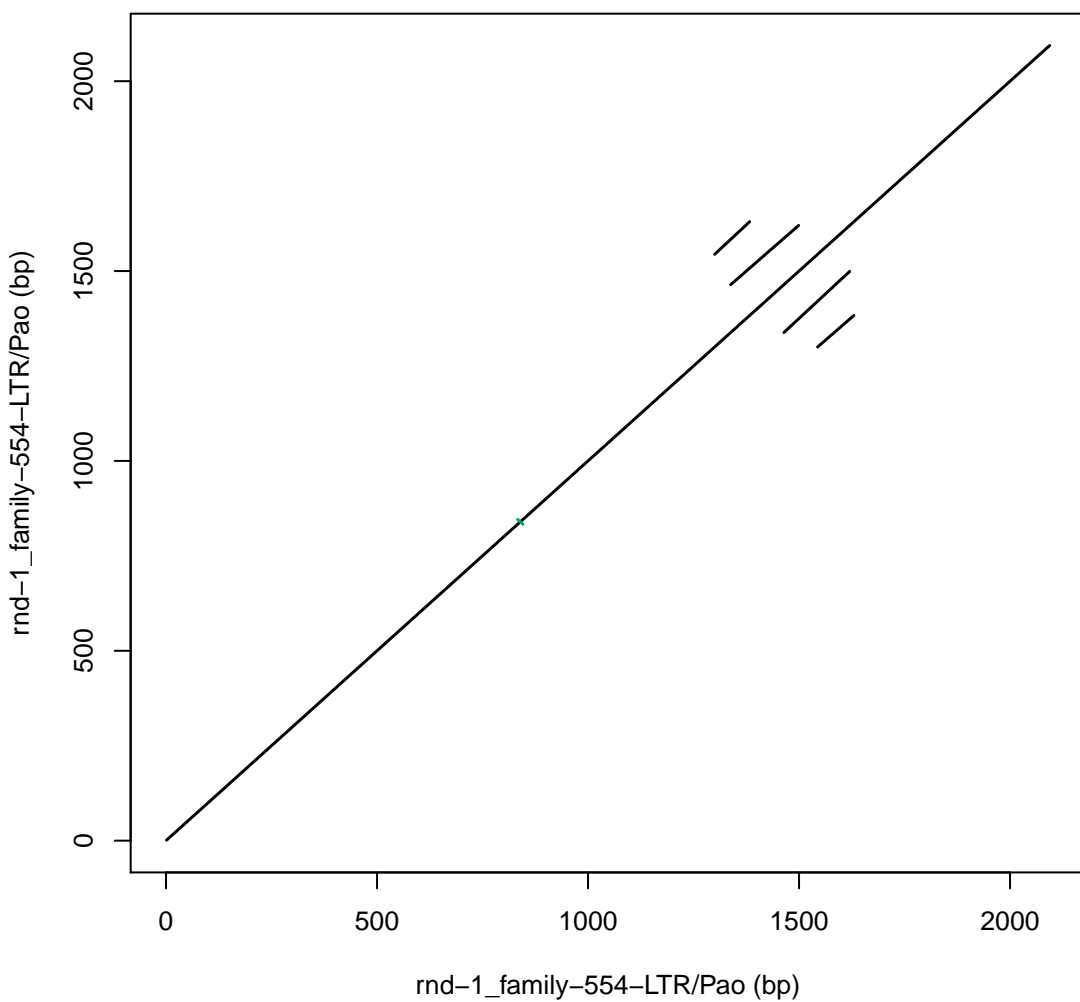
TE: rnd-1_family-554-LTR/Pao
consensus size: 2094bp; fragments: 165; full length: 2 (>=1884.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

