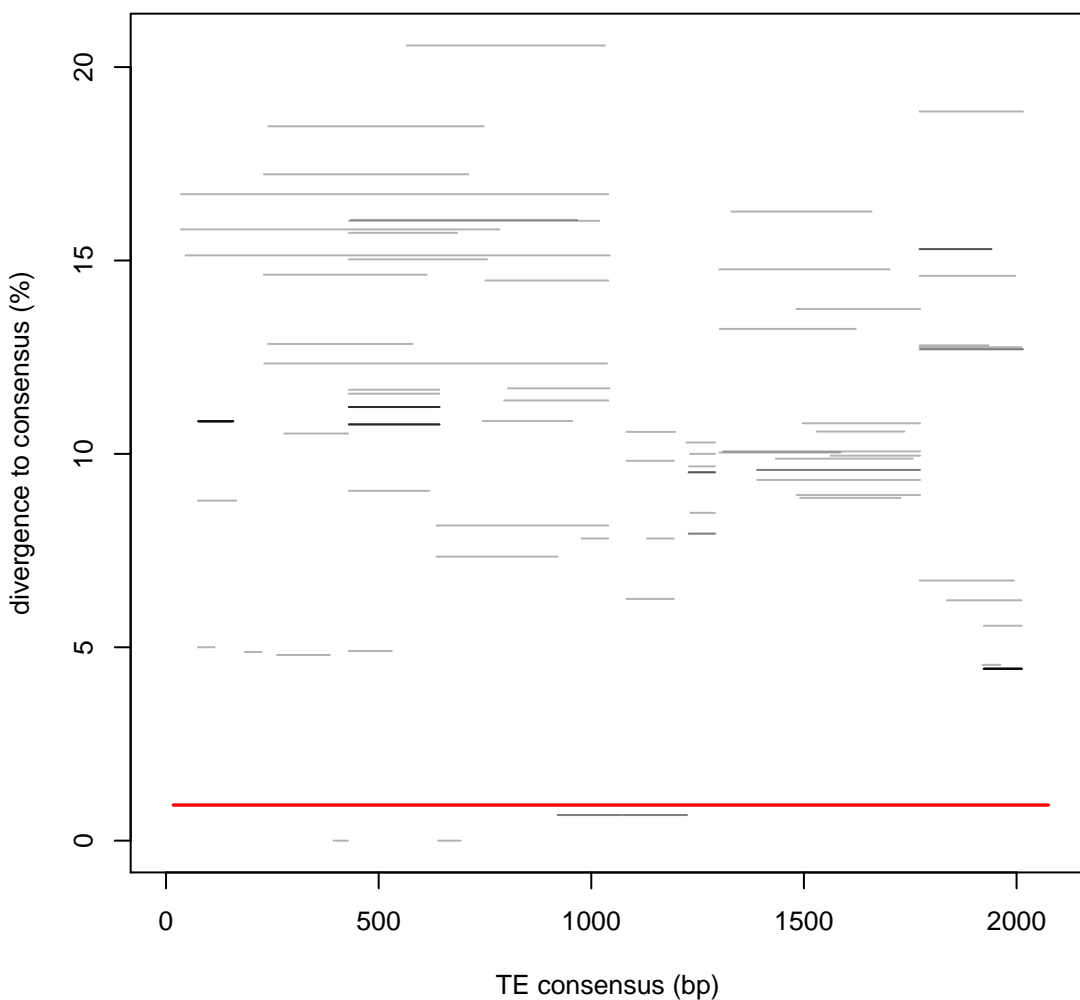
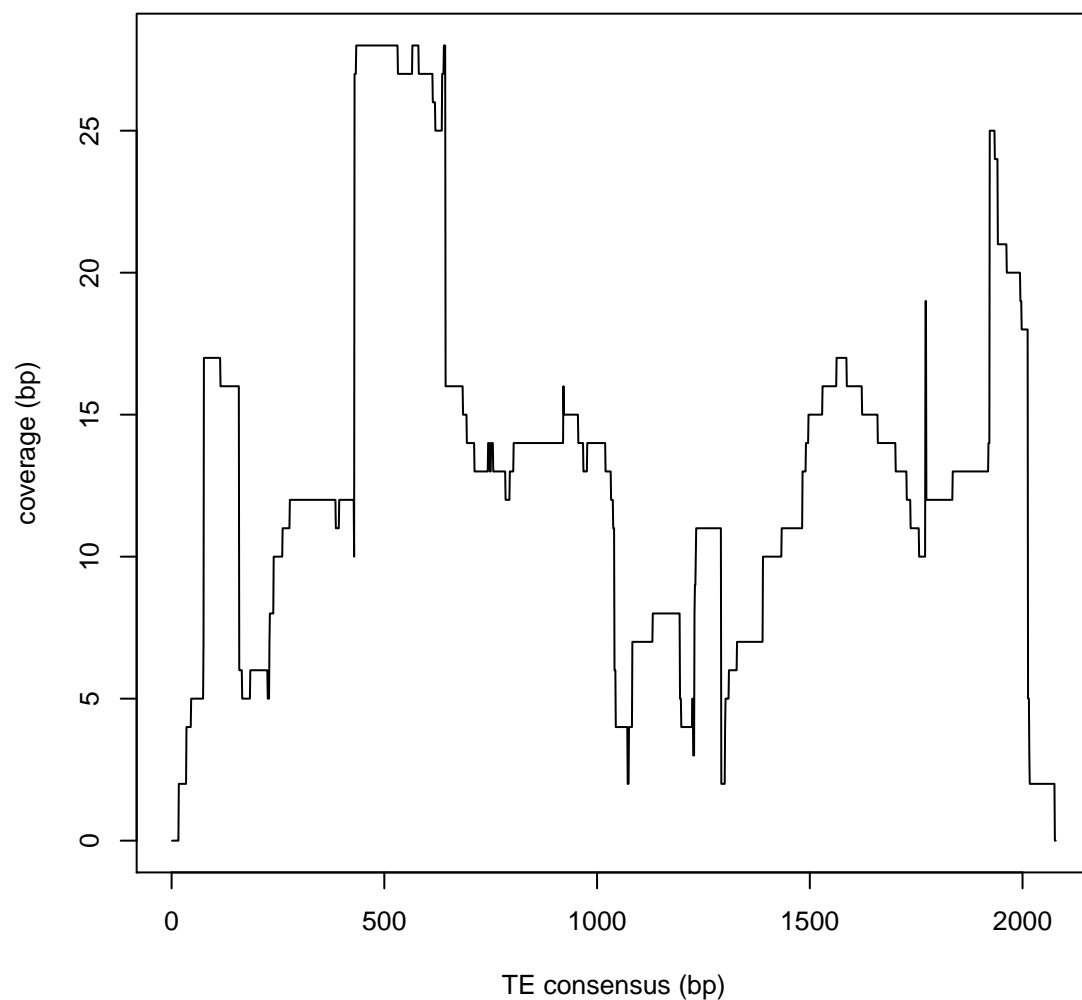


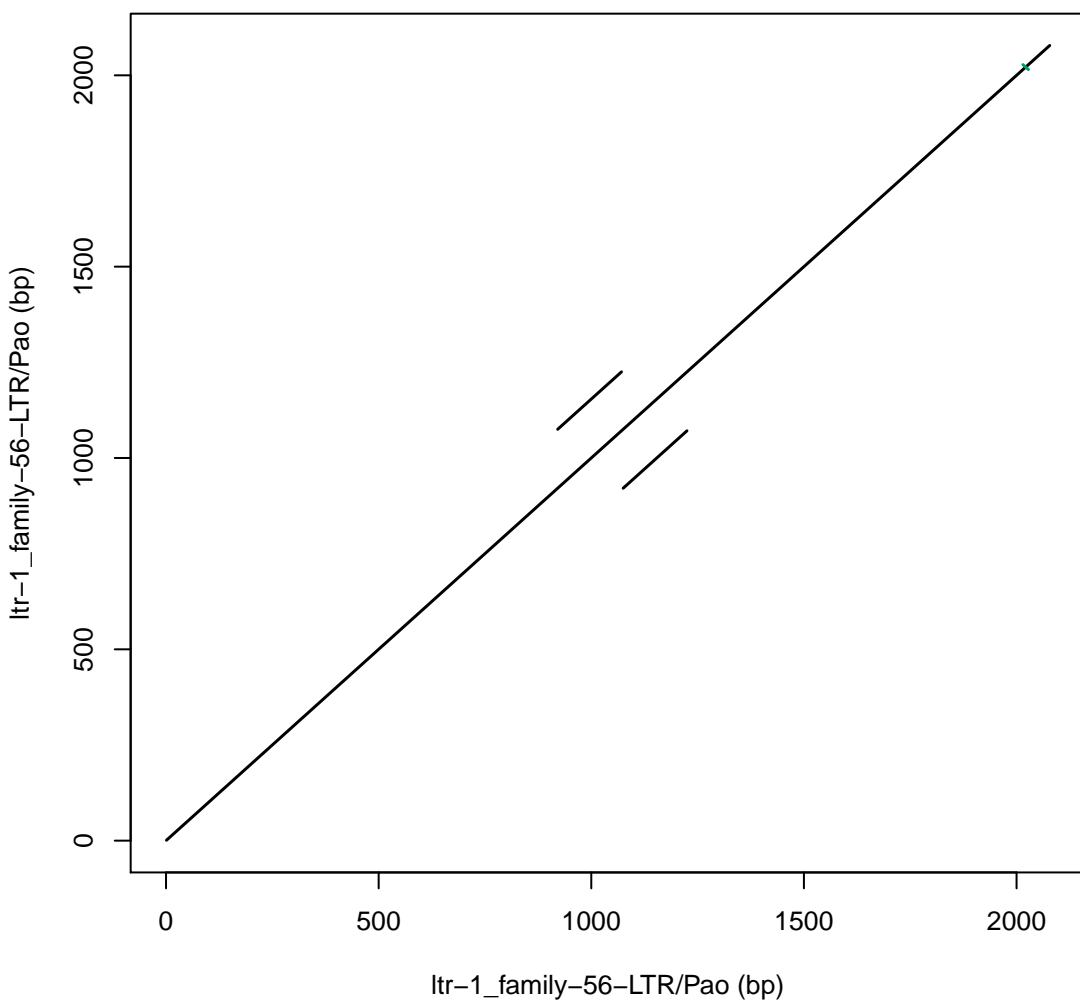
TE: ltr-1_family-56-LTR/Pao
consensus size: 2078bp; fragments: 108; full length: 2 (≥ 1870.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

