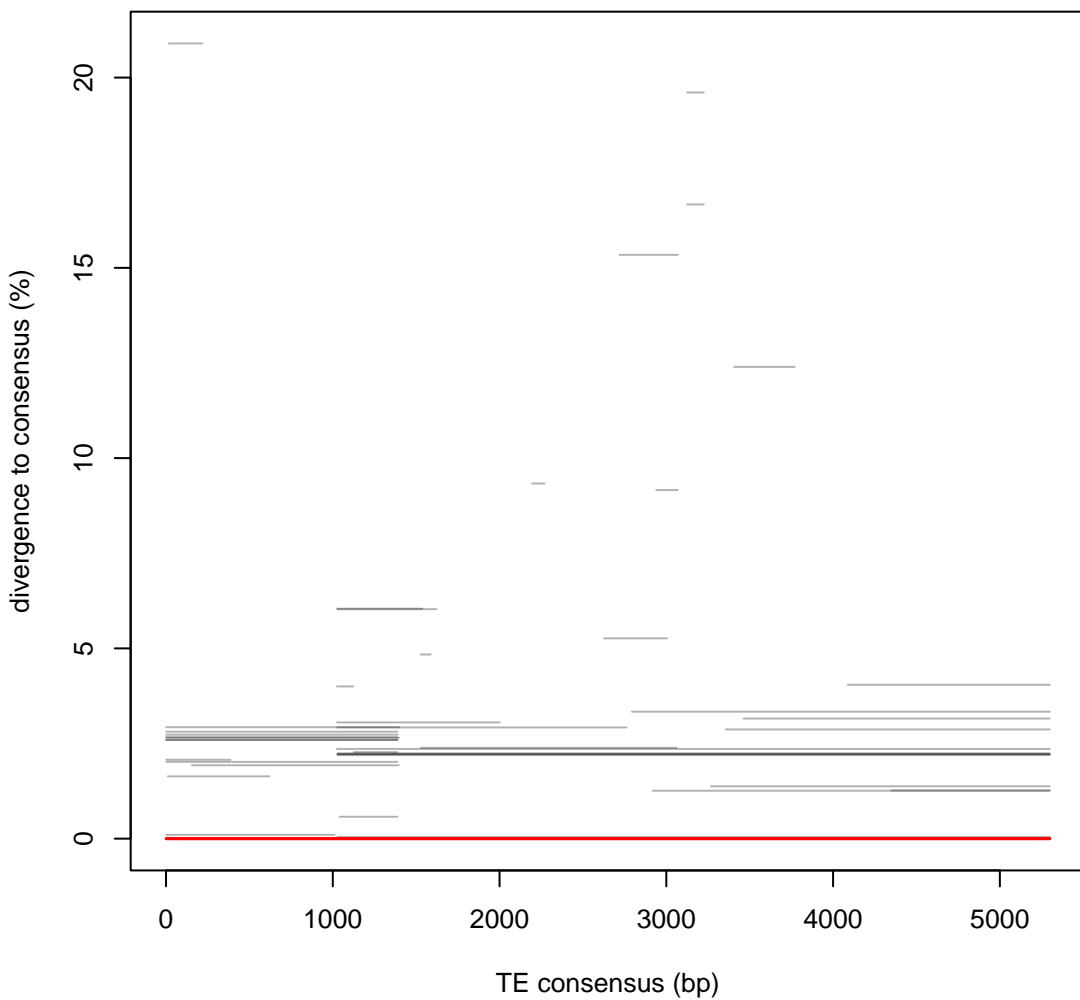
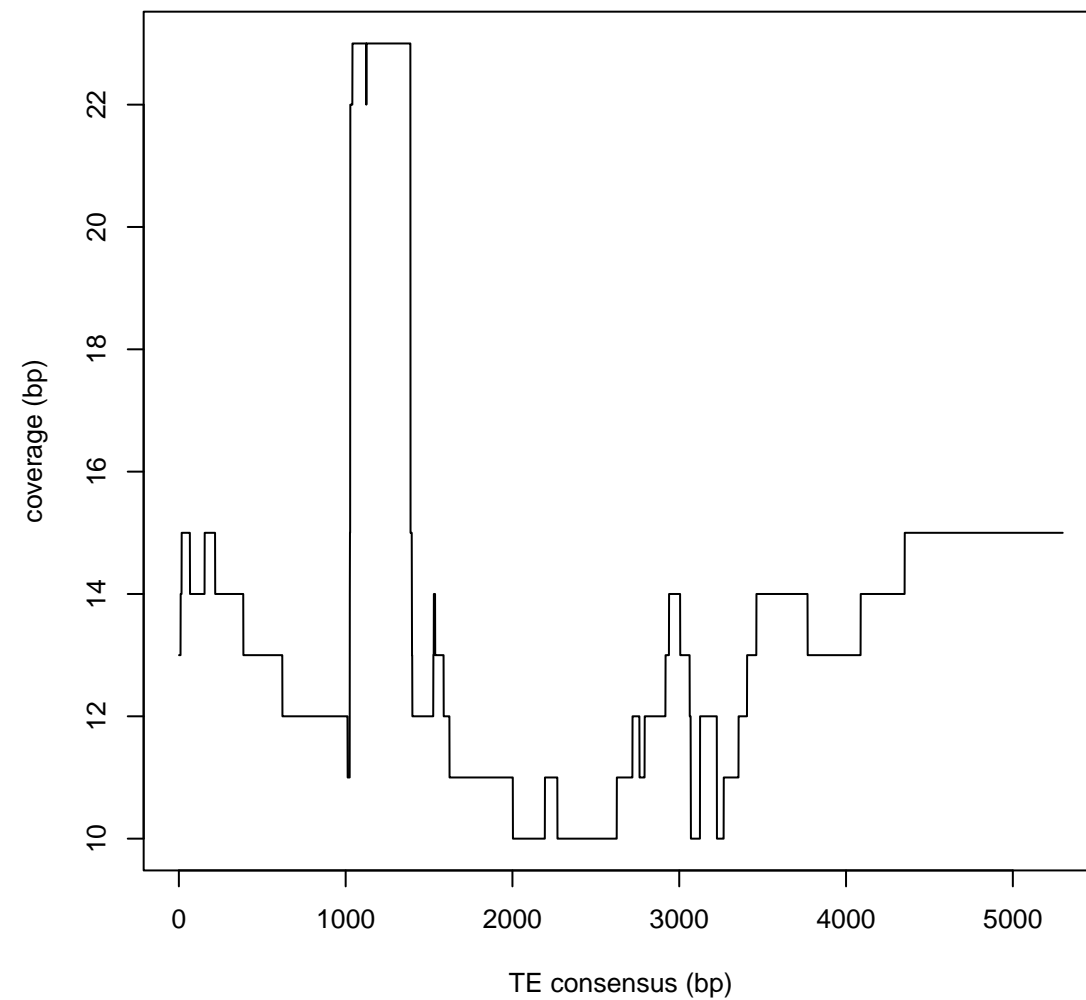


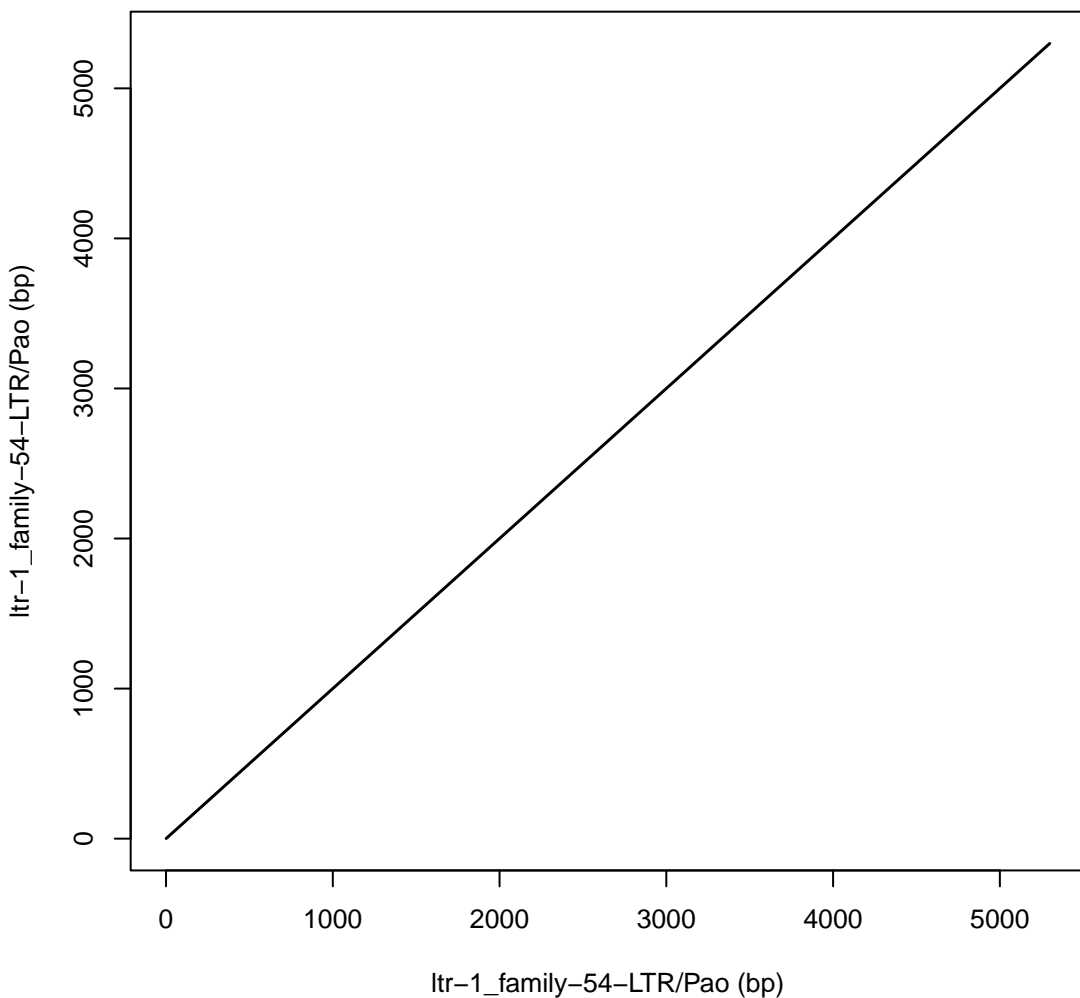
TE: ltr-1_family-54-LTR/Pao
consensus size: 5299bp; fragments: 45; full length: 2 (≥ 4769.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

