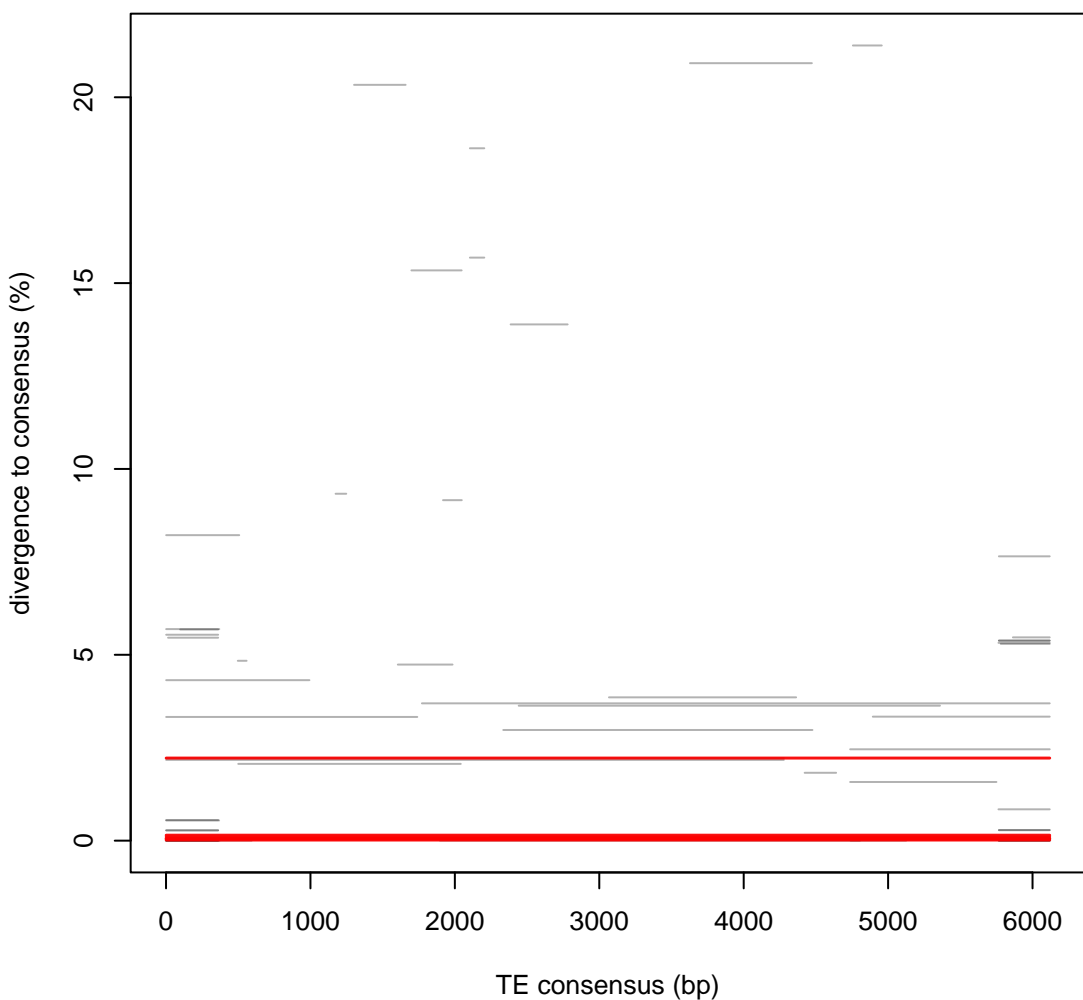
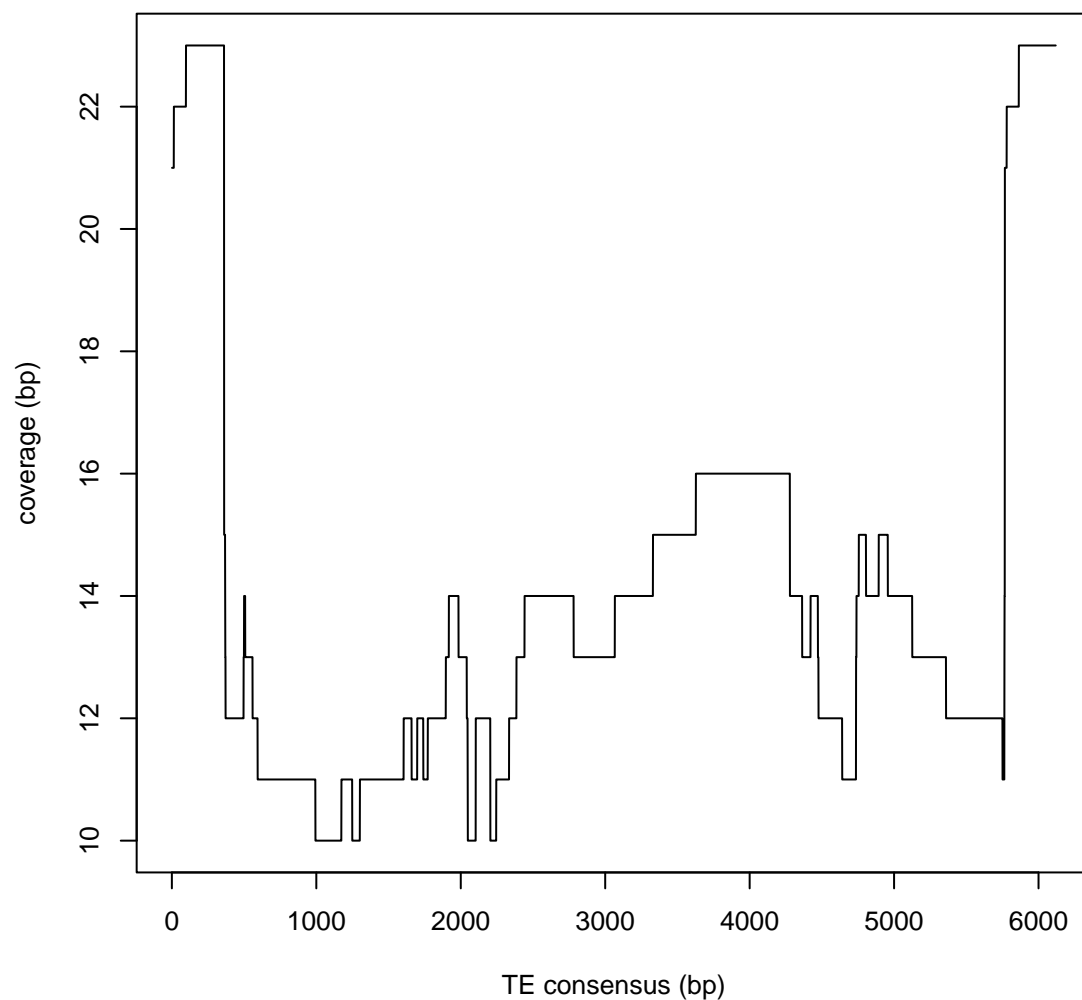


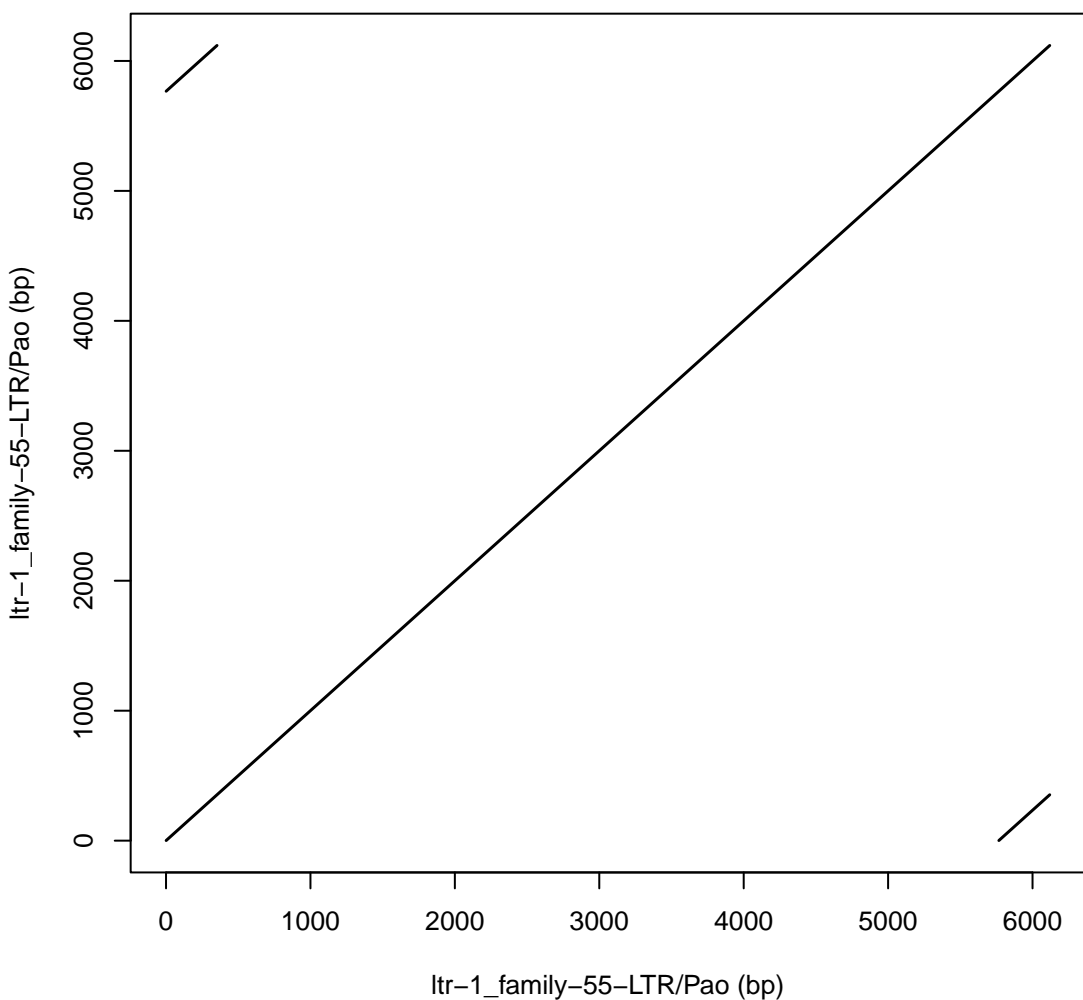
TE: ltr-1_family-55-LTR/Pao
consensus size: 6119bp; fragments: 59; full length: 6 (≥ 5507.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

