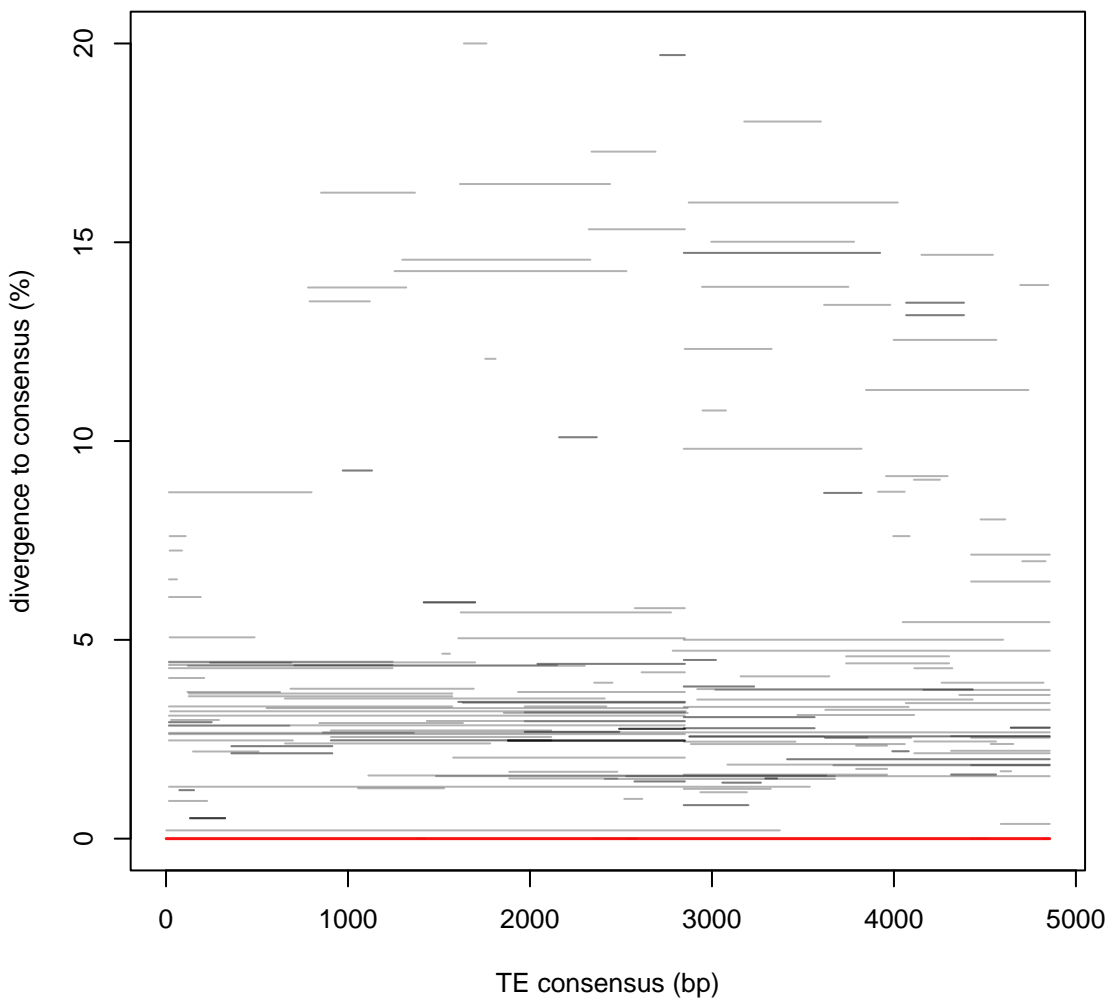
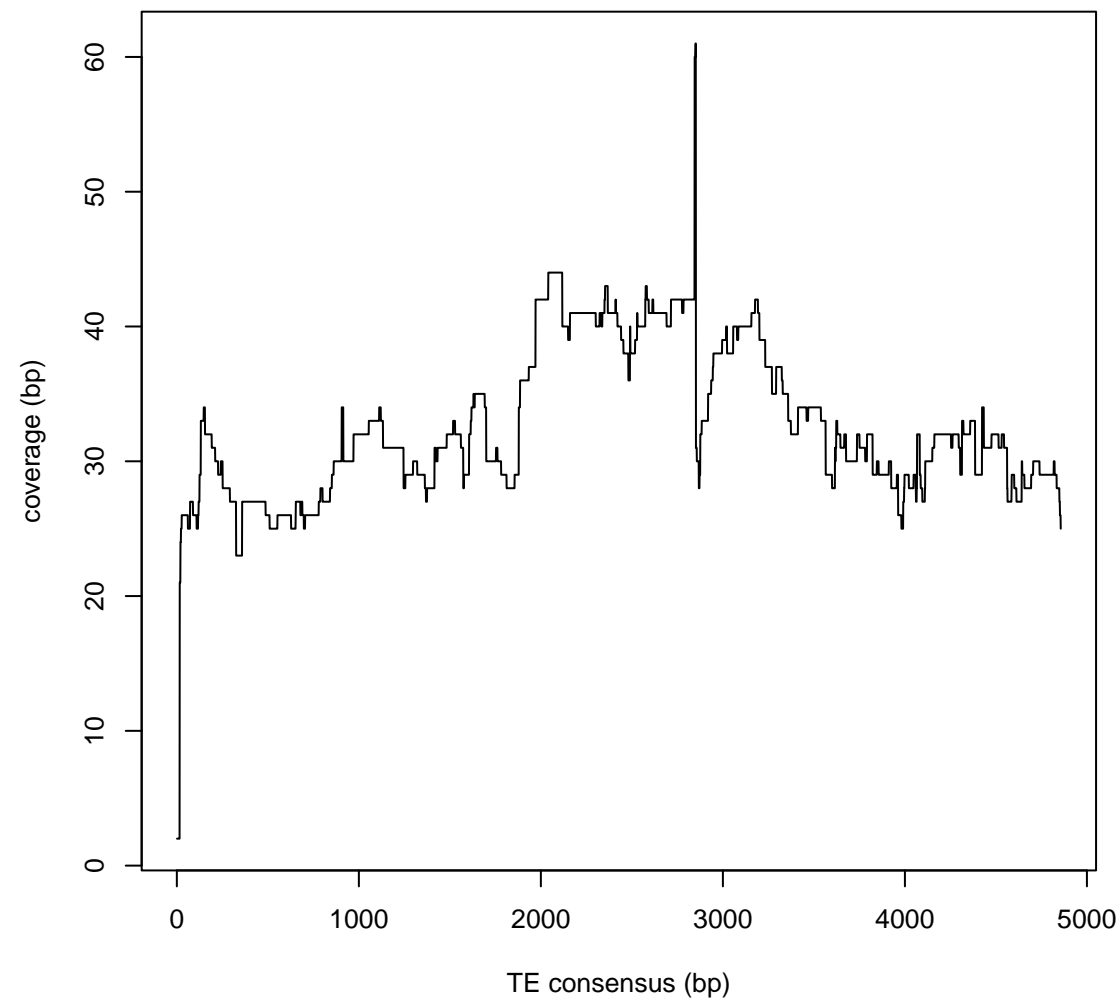


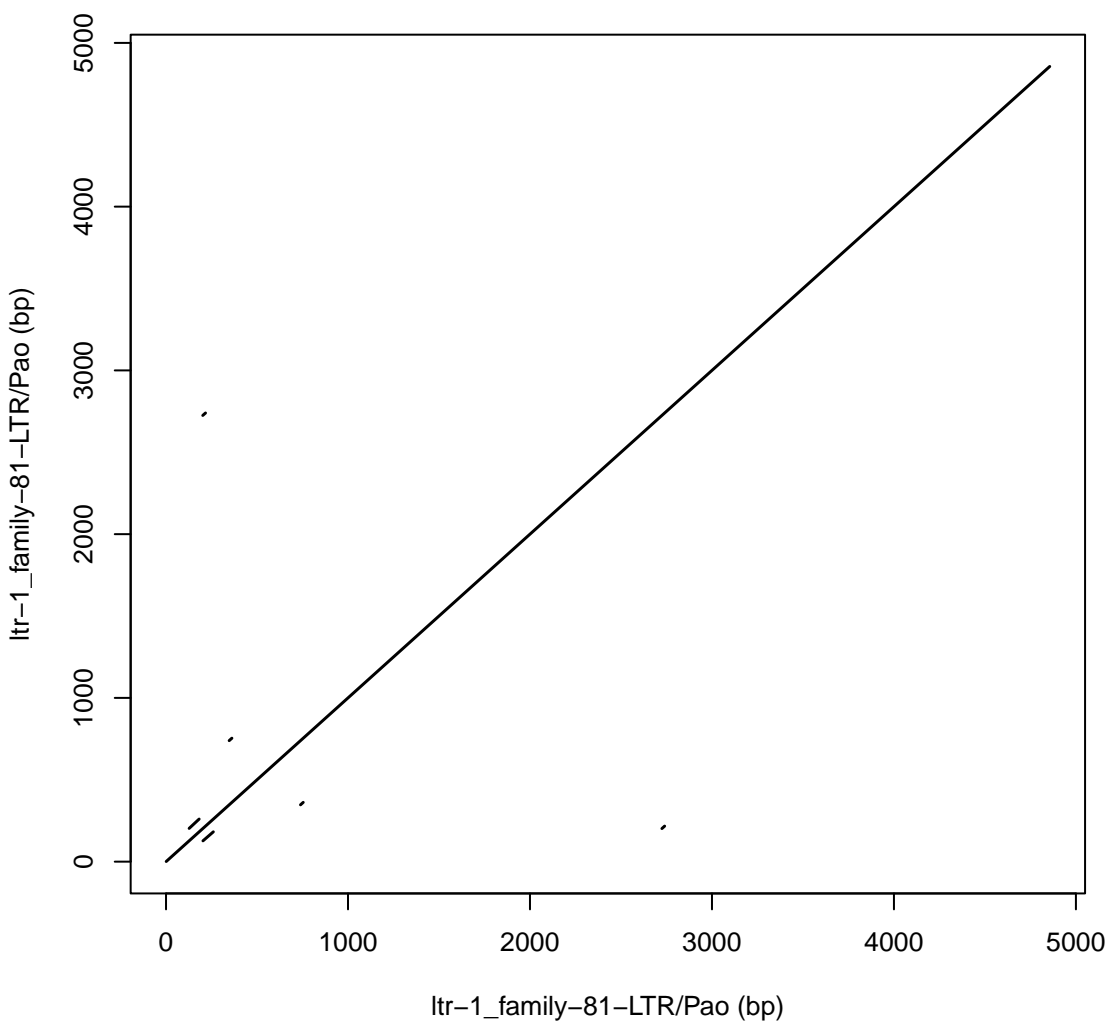
TE: ltr-1_family-81-LTR/Pao
consensus size: 4856bp; fragments: 207; full length: 1 (≥ 4370.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

