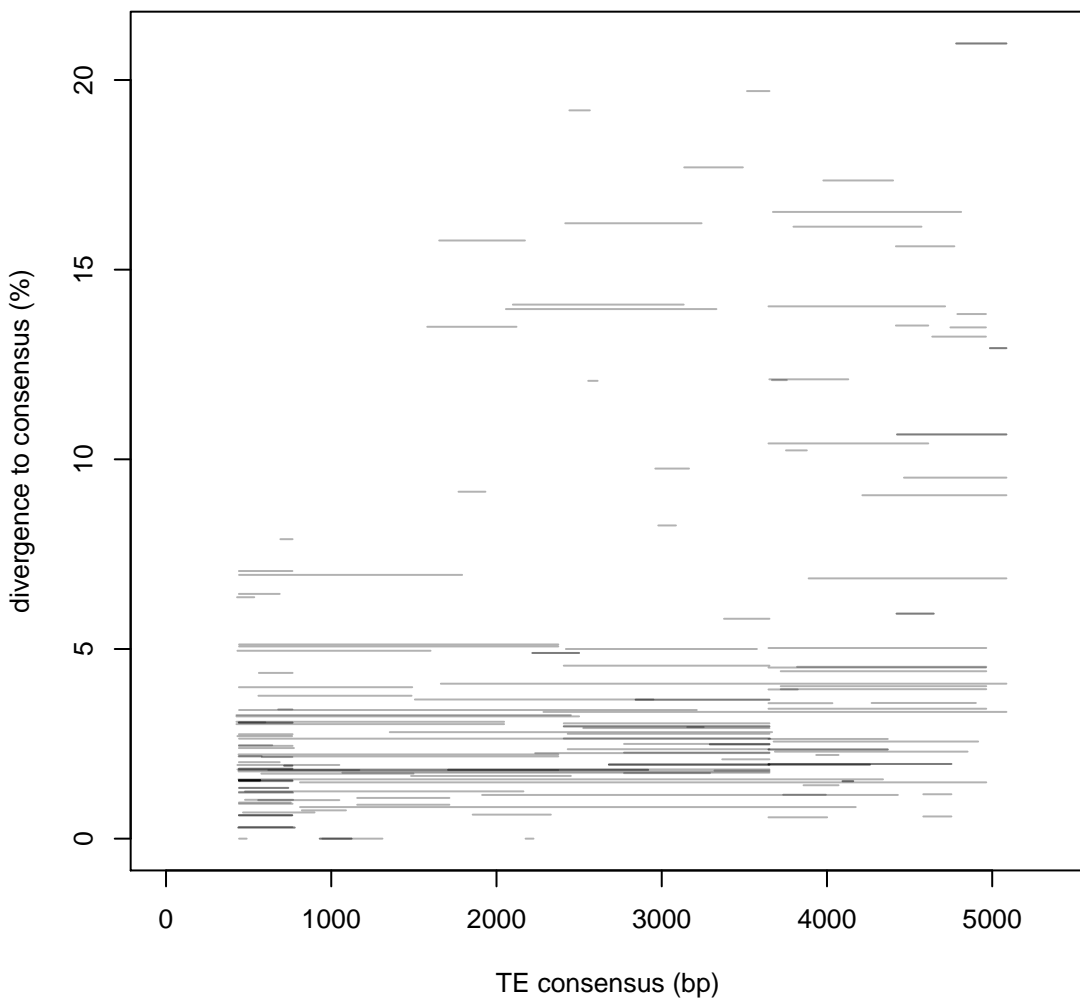
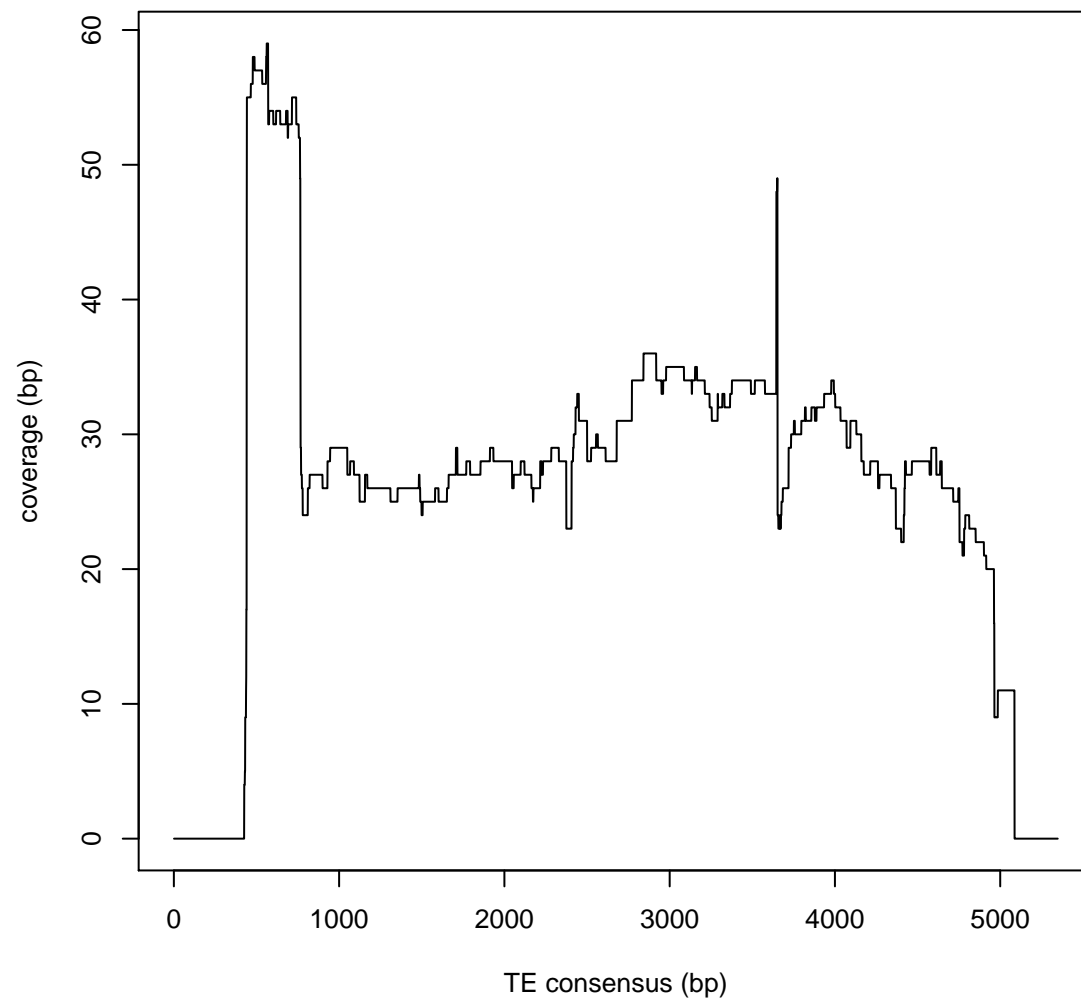


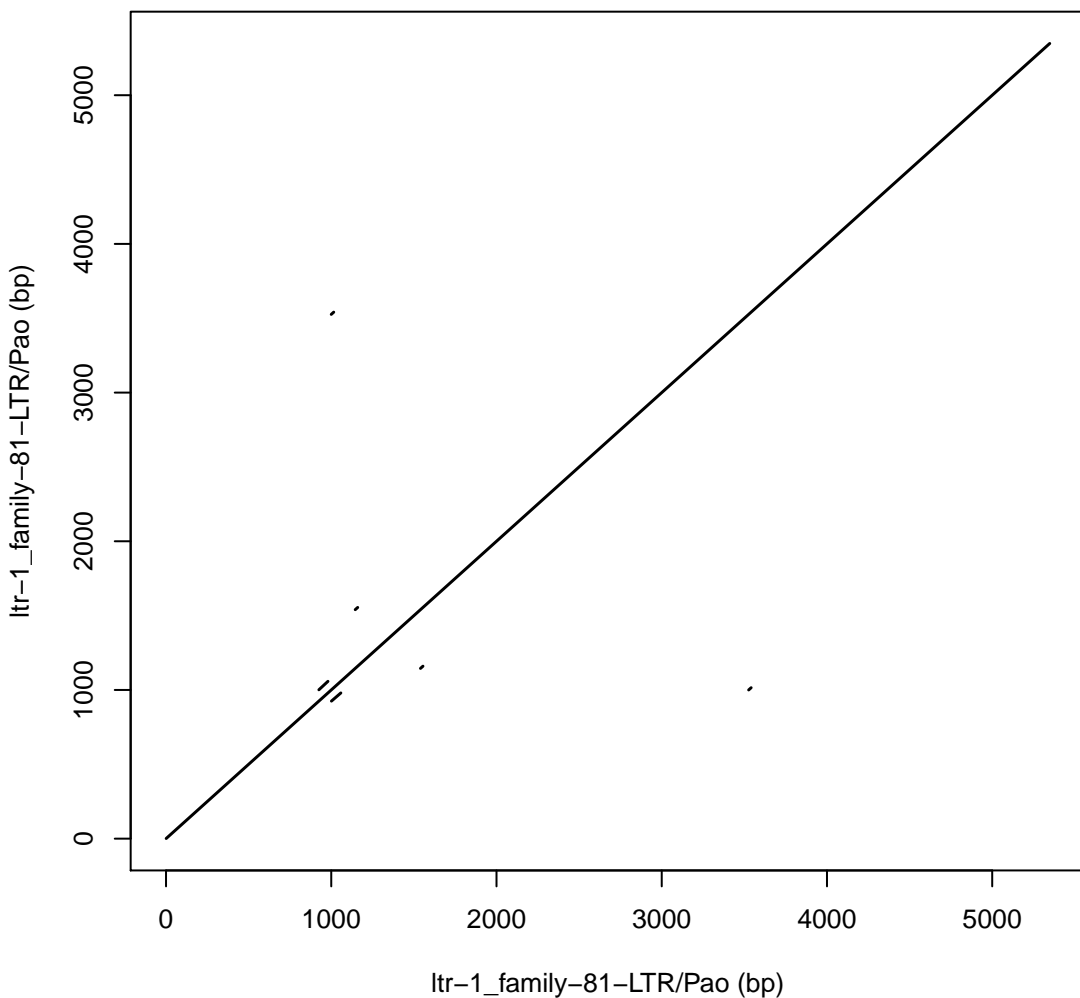
TE: ltr-1_family-81-LTR/Pao
consensus size: 5348bp; fragments: 176; full length: 0 (>=4813.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

