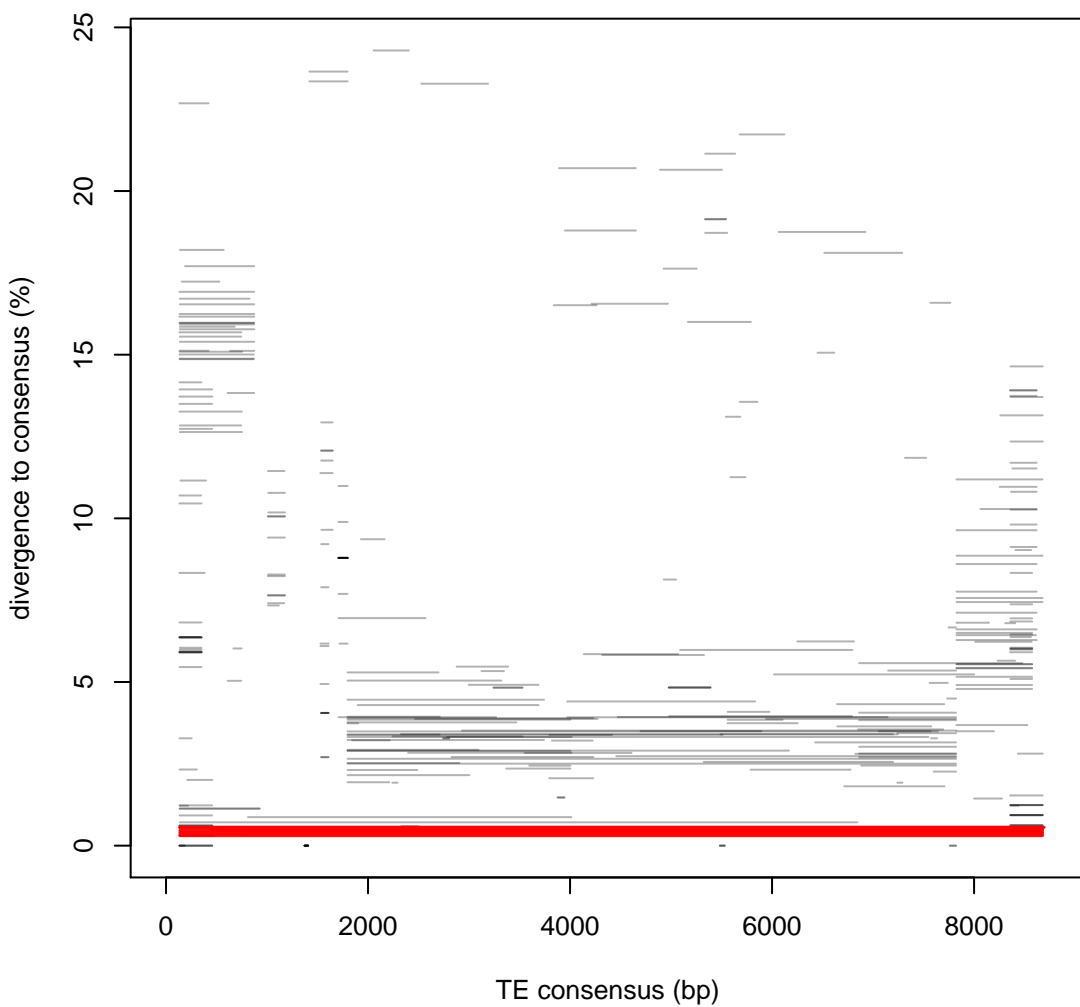
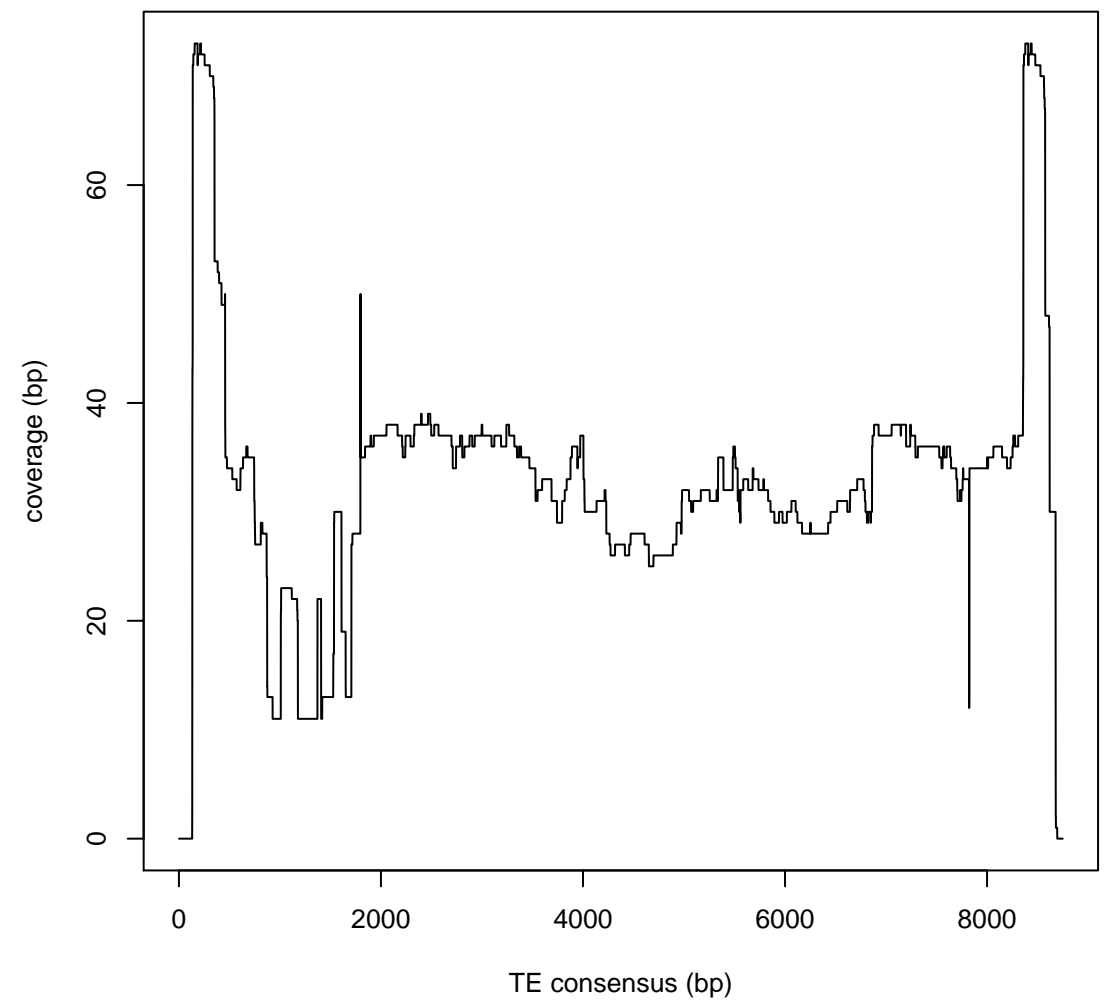


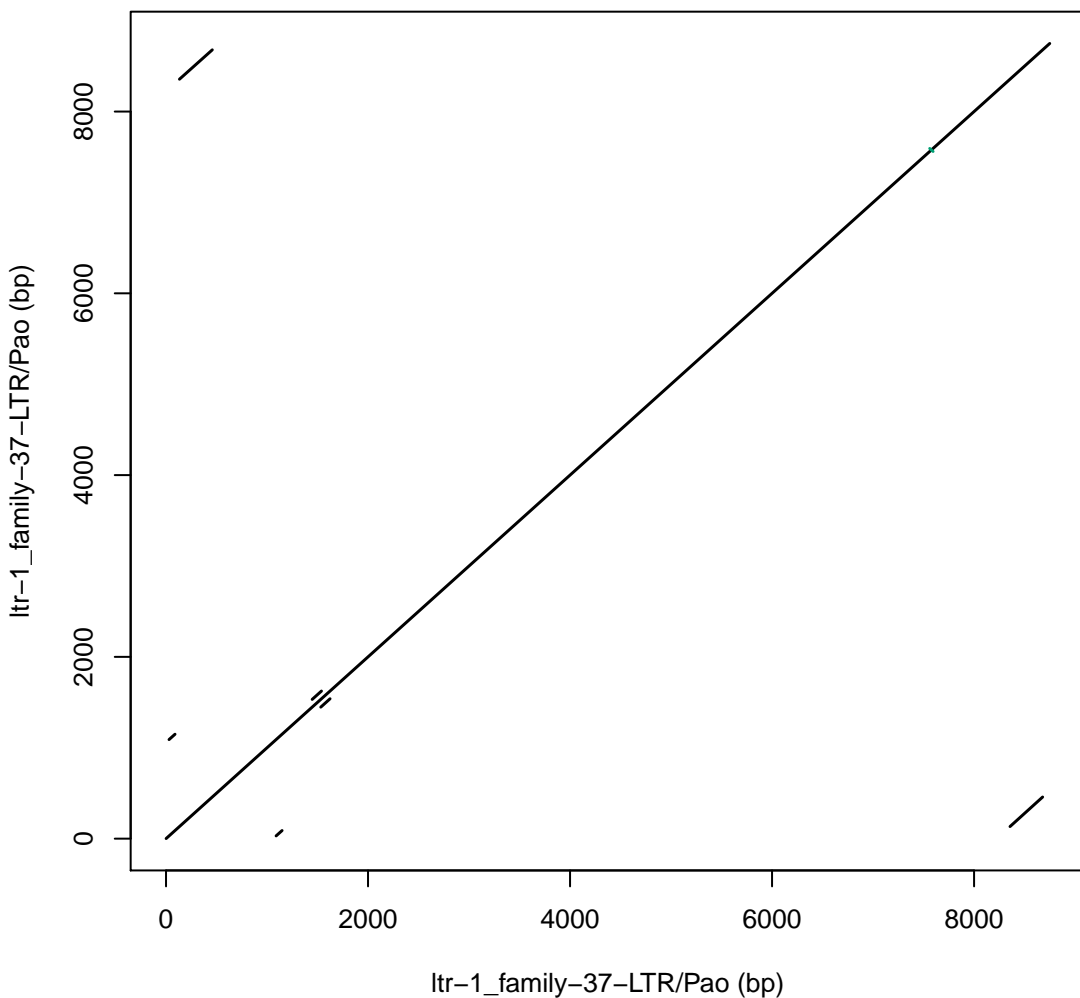
TE: ltr-1_family-37-LTR/Pao
consensus size: 8749bp; fragments: 327; full length: 8 (>=7874.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

