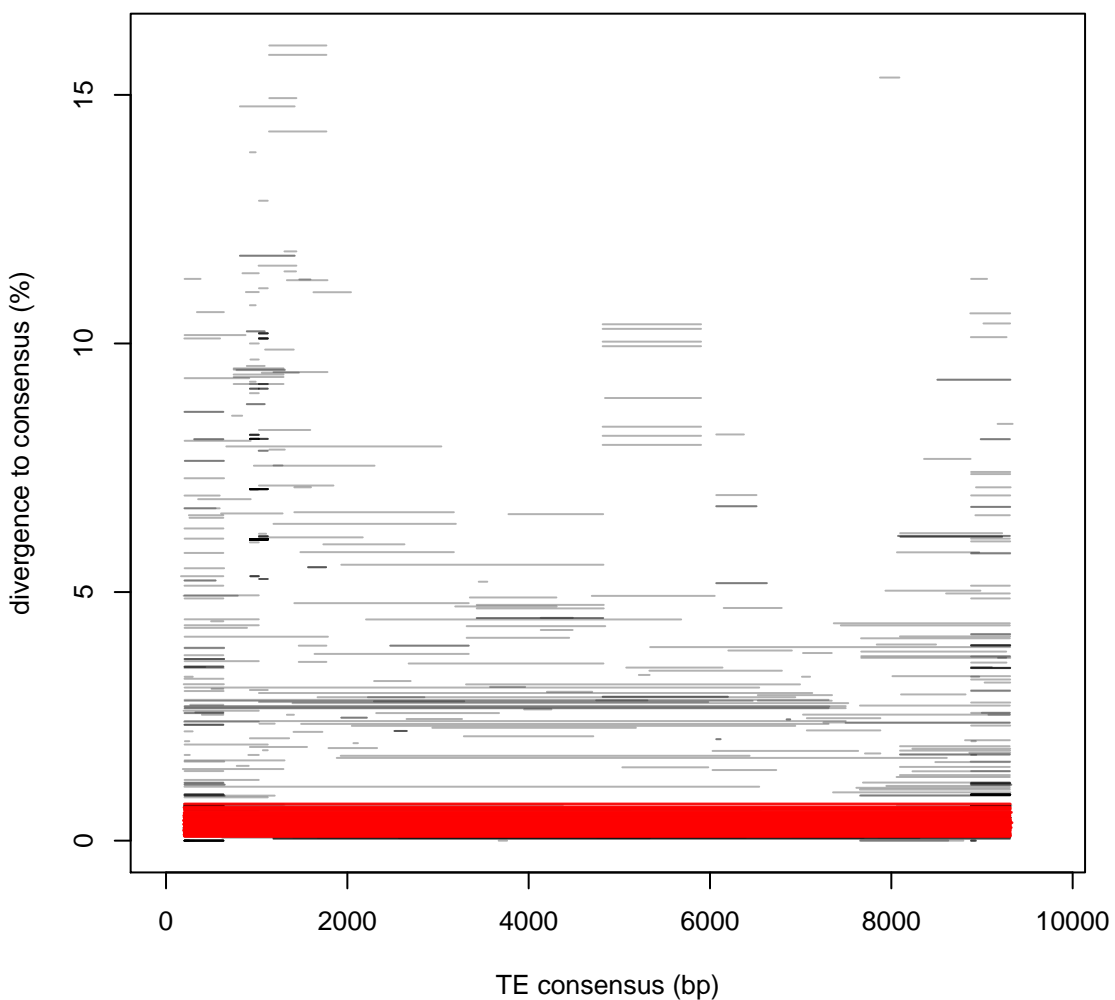
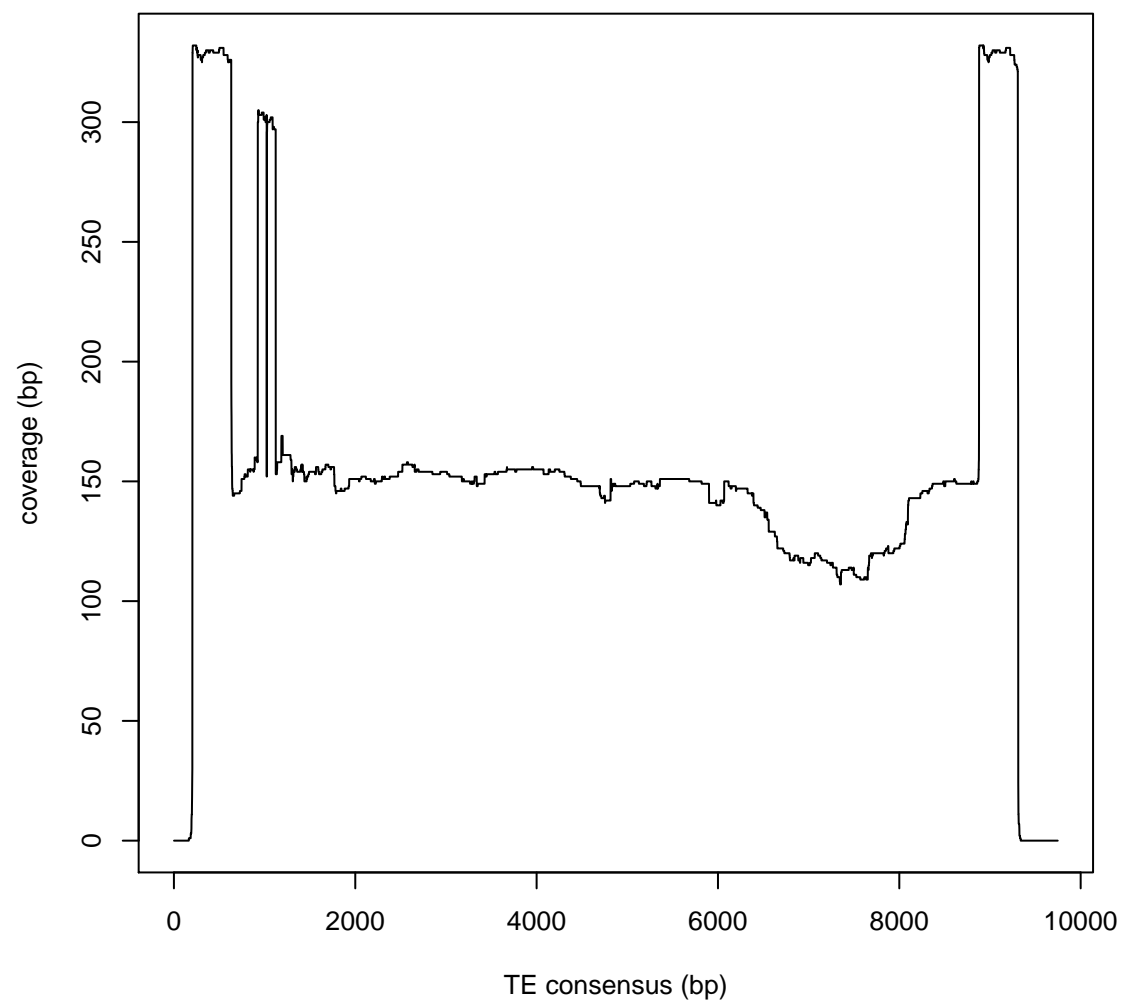


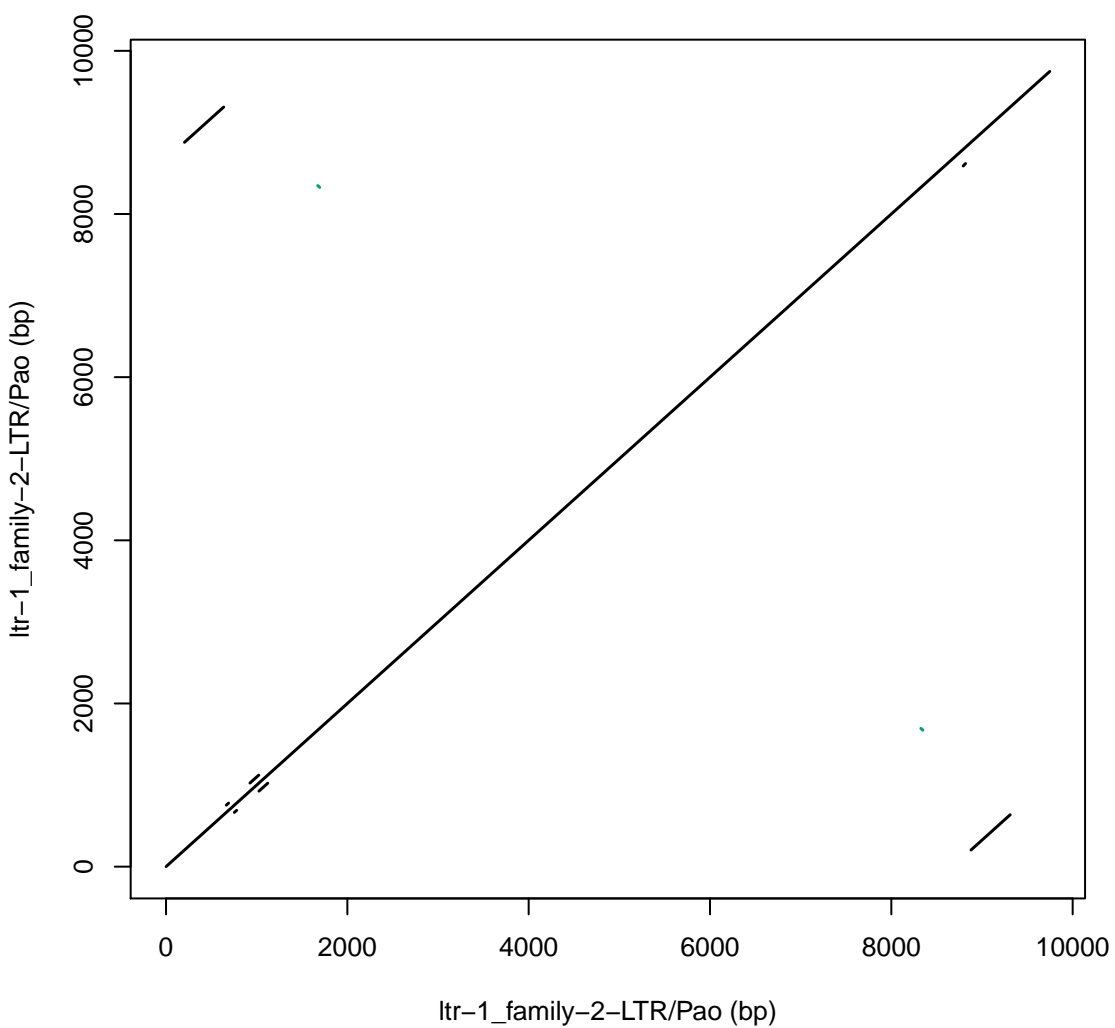
TE: ltr-1_family-2-LTR/Pao
consensus size: 9747bp; fragments: 1069; full length: 60 (>=8772.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

