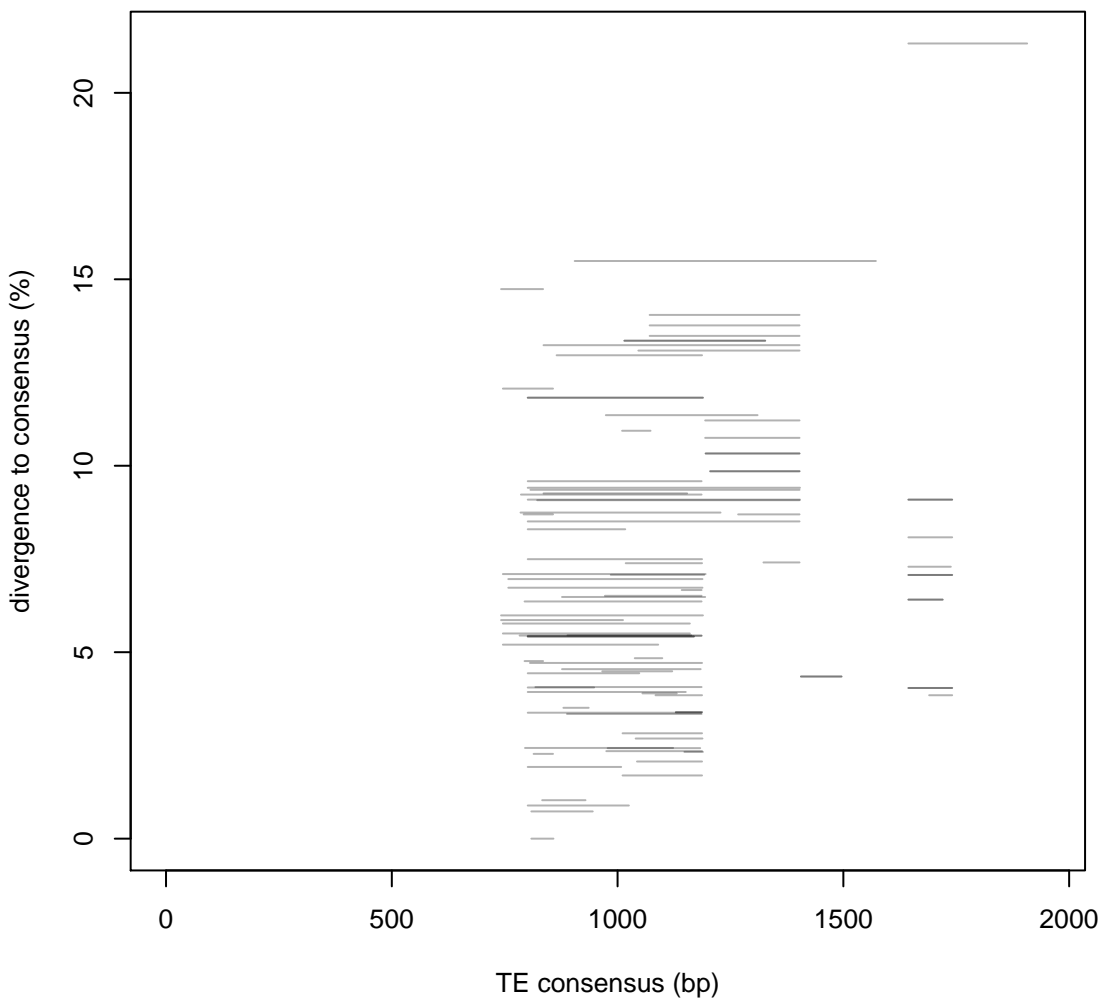
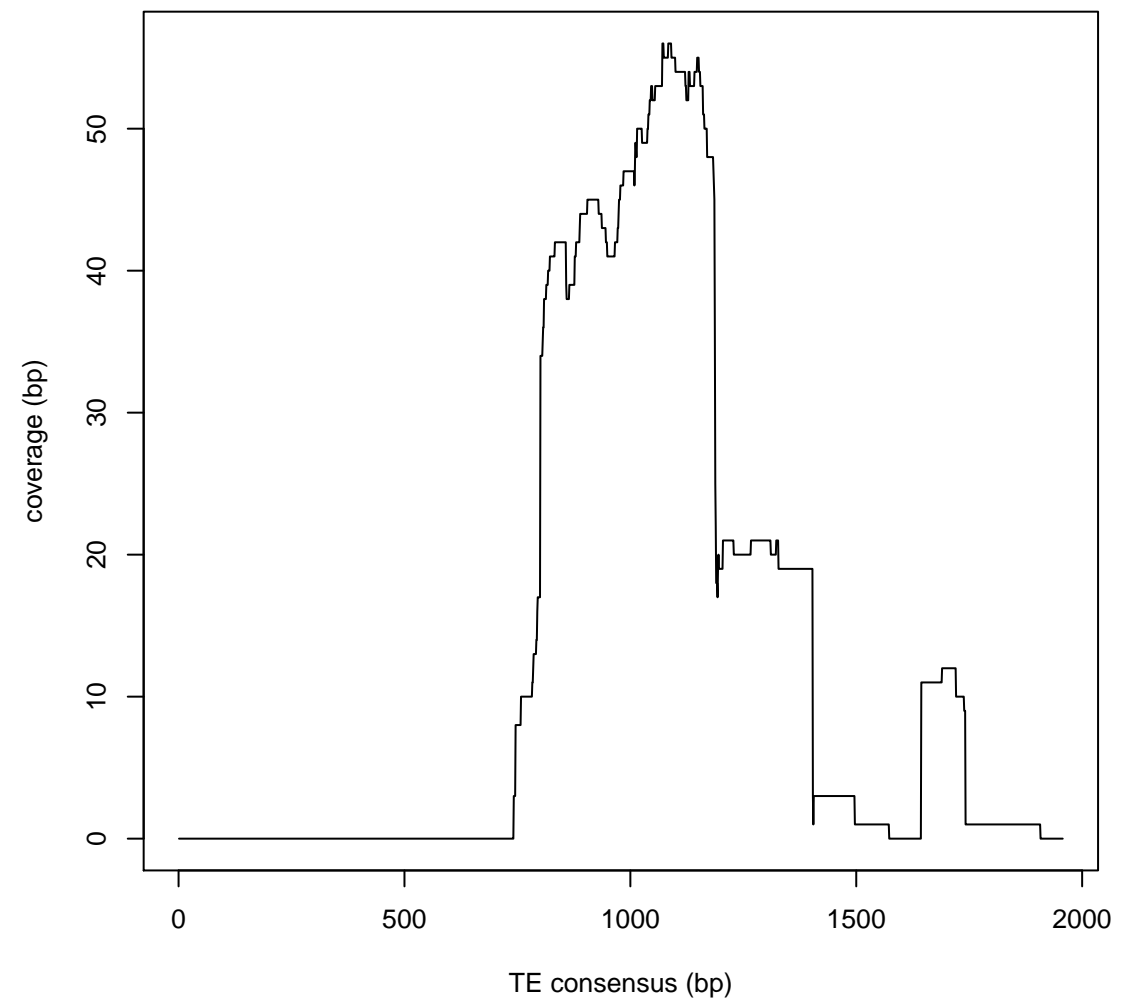


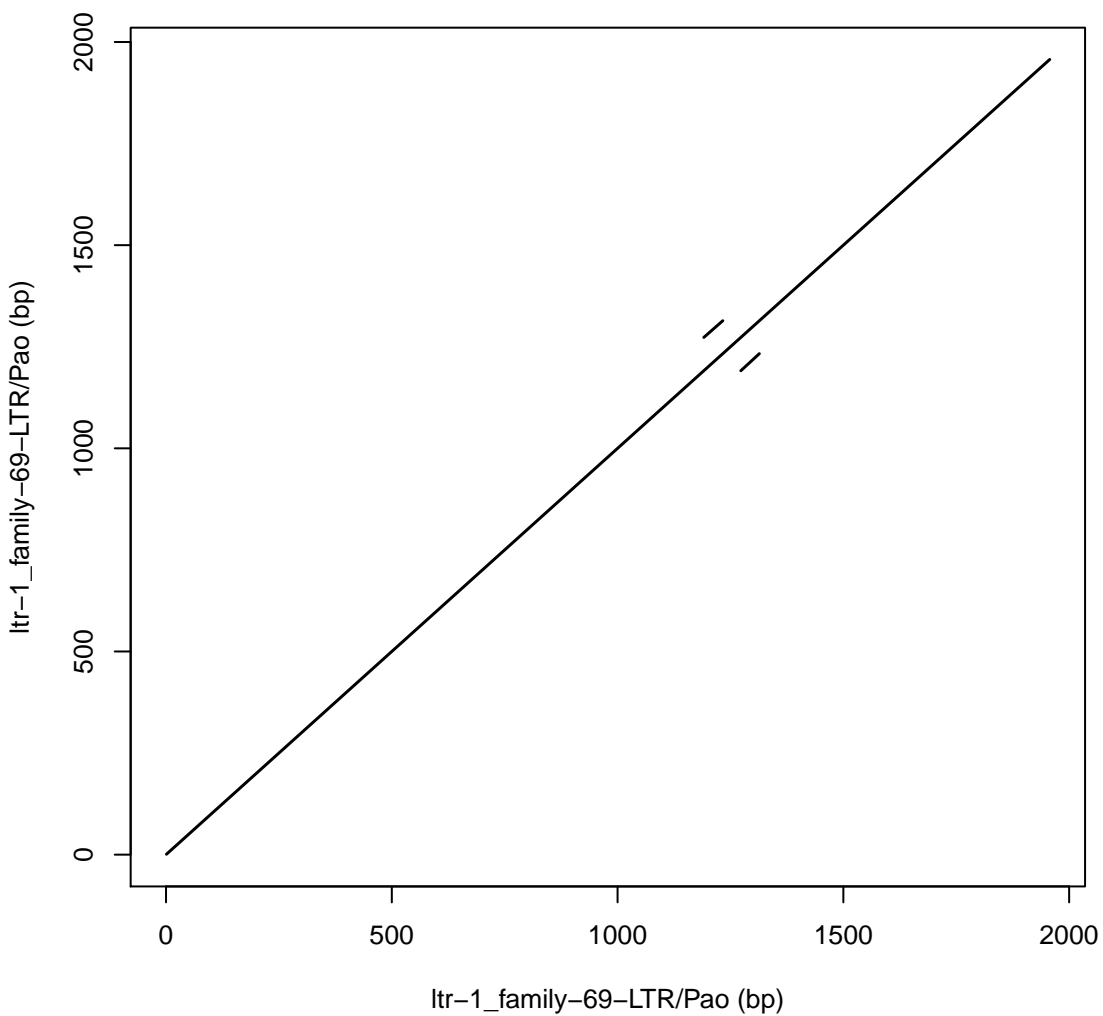
TE: ltr-1_family-69-LTR/Pao
consensus size: 1957bp; fragments: 98; full length: 0 (>=1761.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

