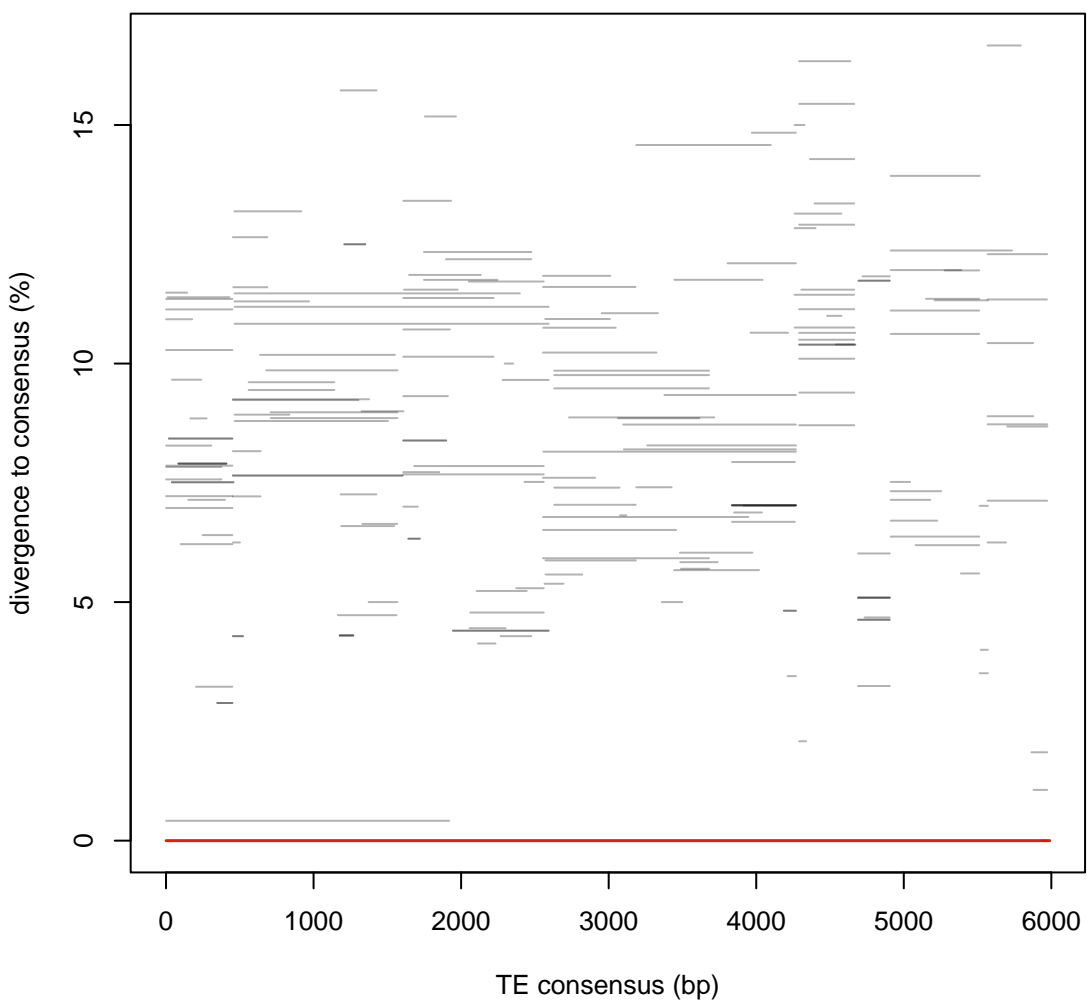
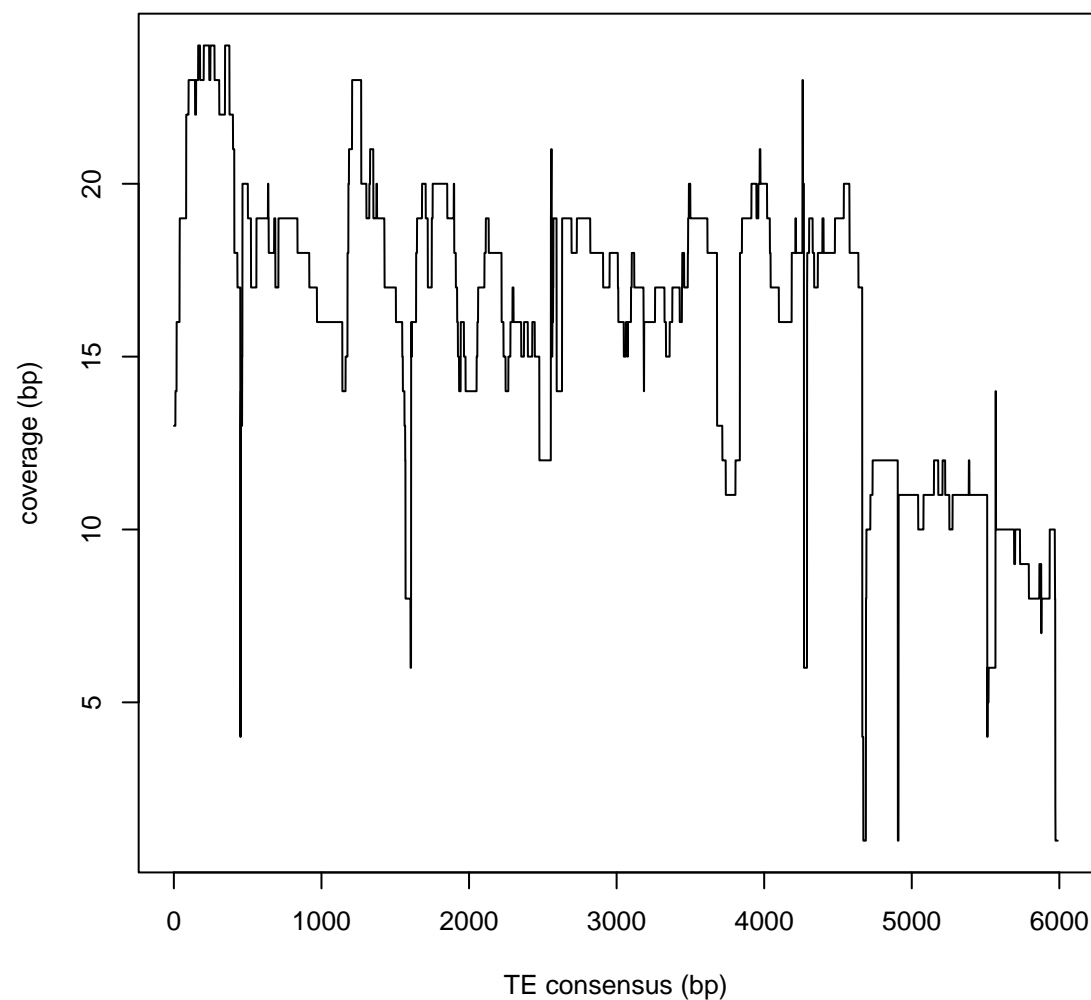


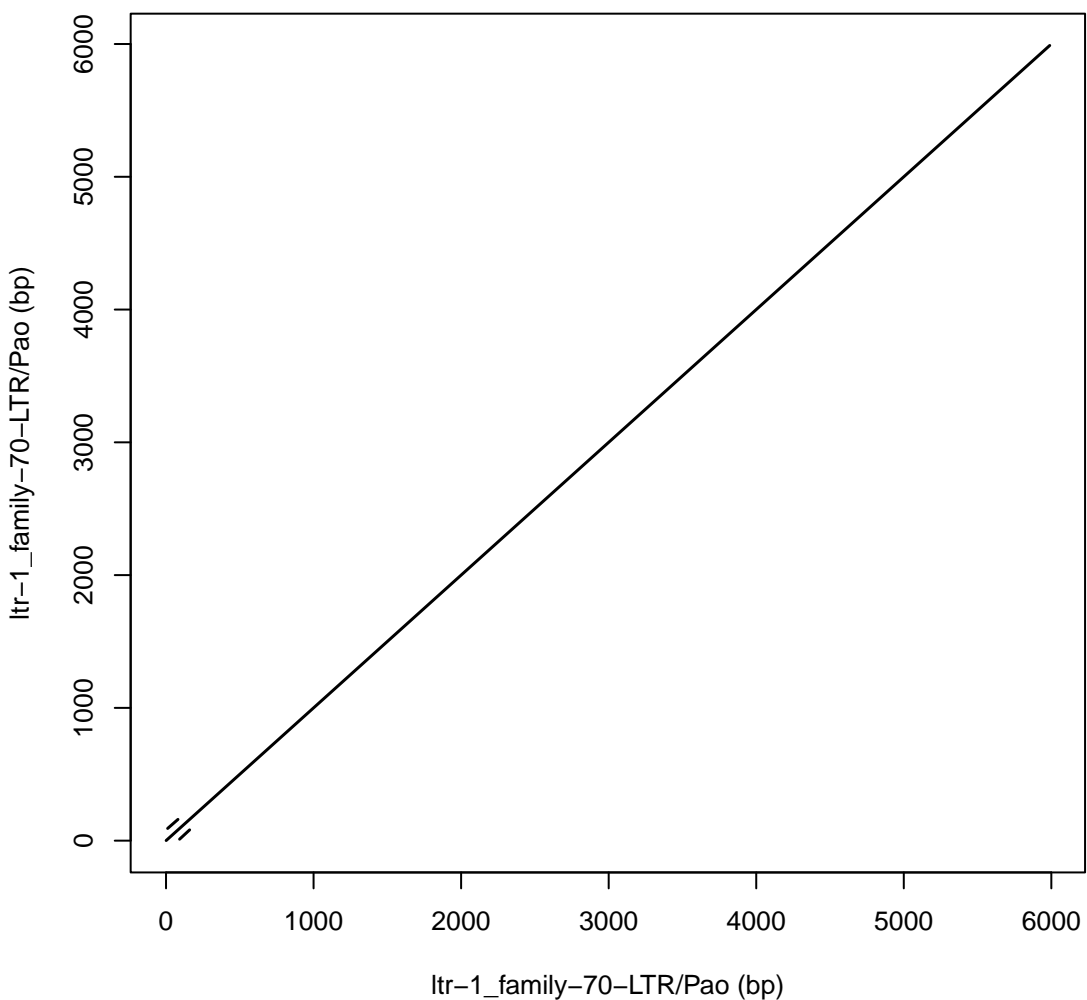
TE: ltr-1_family-70-LTR/Pao
consensus size: 5989bp; fragments: 208; full length: 1 (>=5390.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

