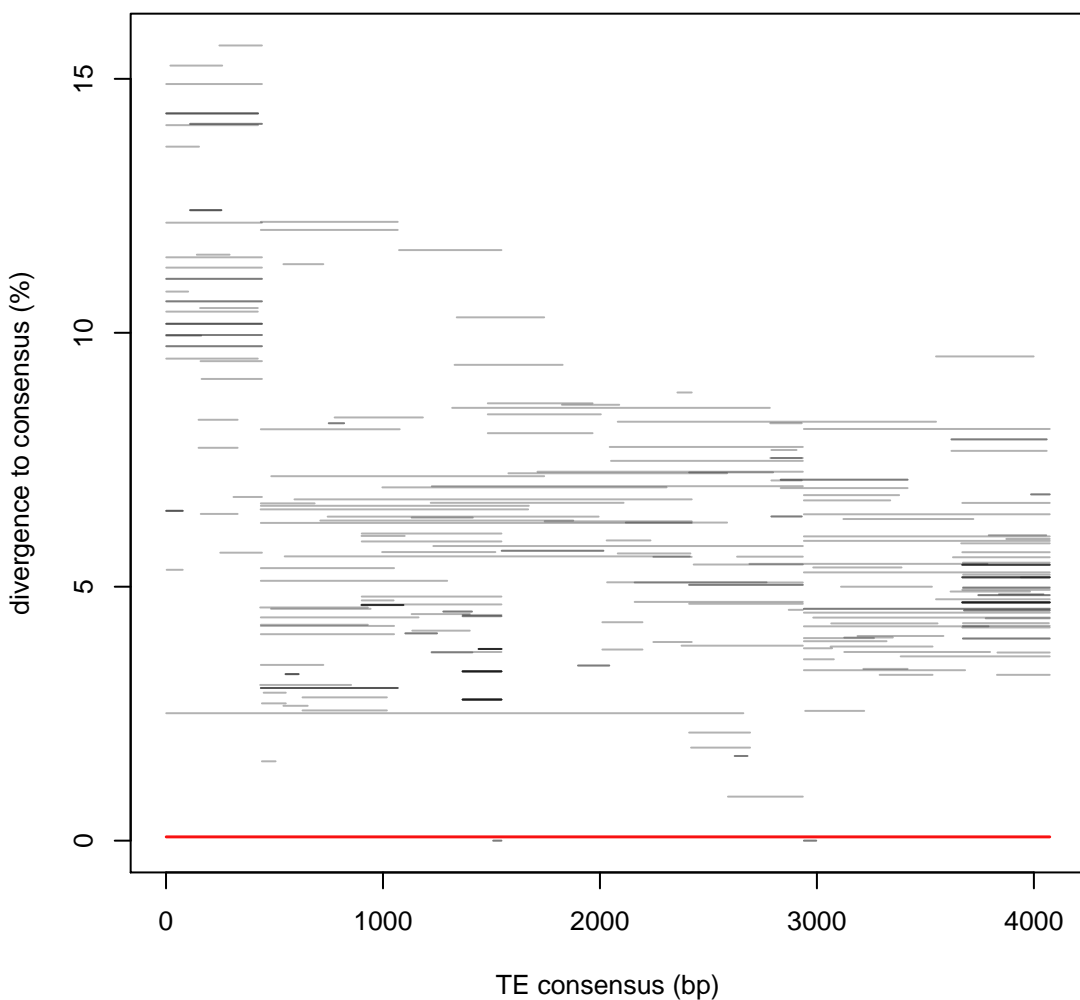
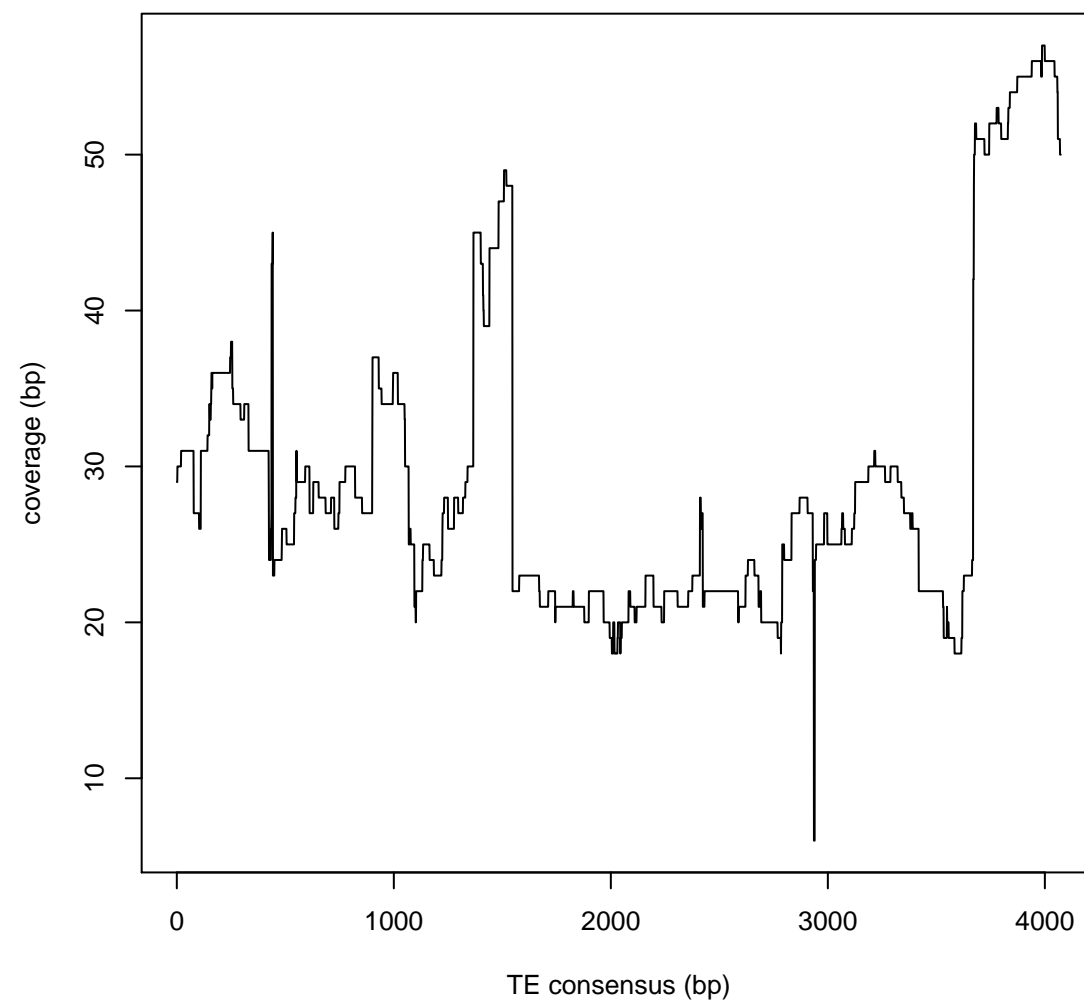


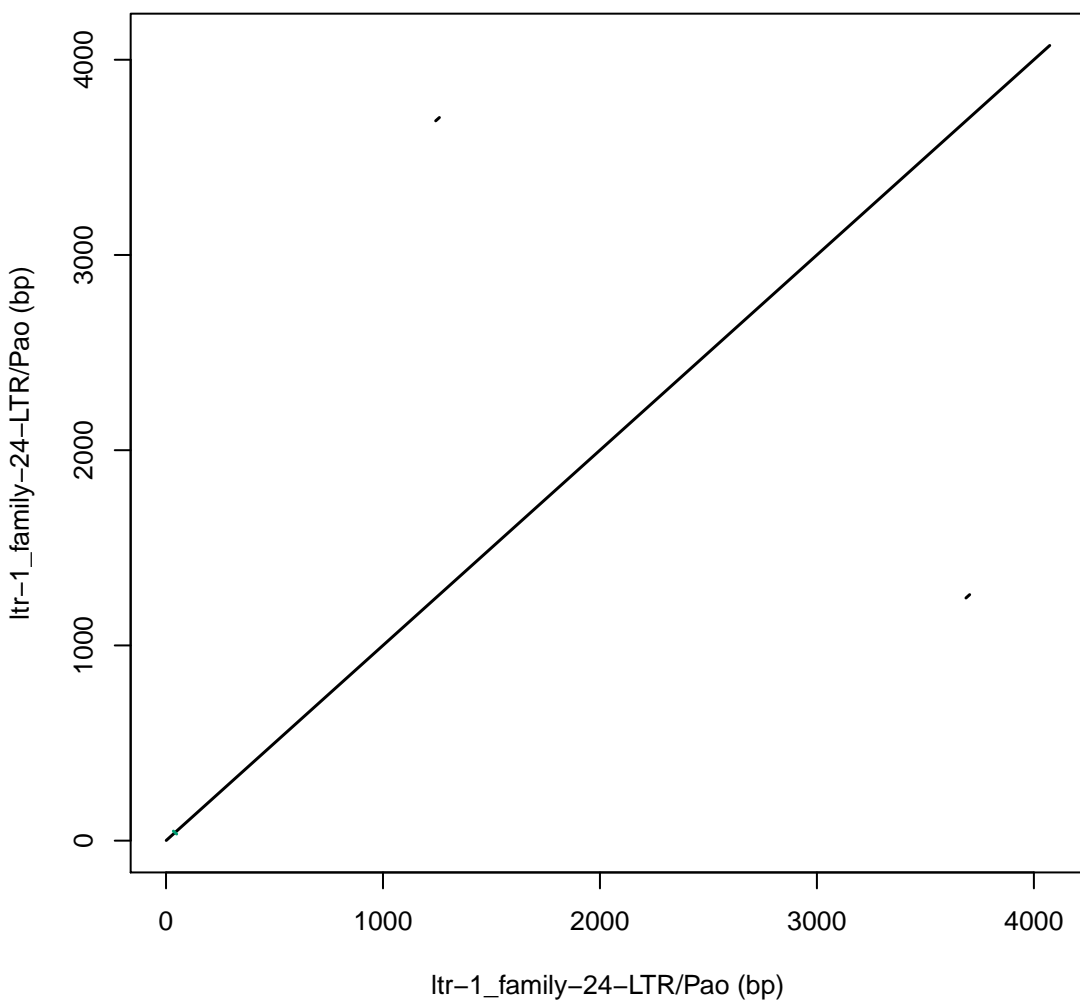
TE: ltr-1_family-24-LTR/Pao
consensus size: 4073bp; fragments: 261; full length: 1 (>=3665.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

