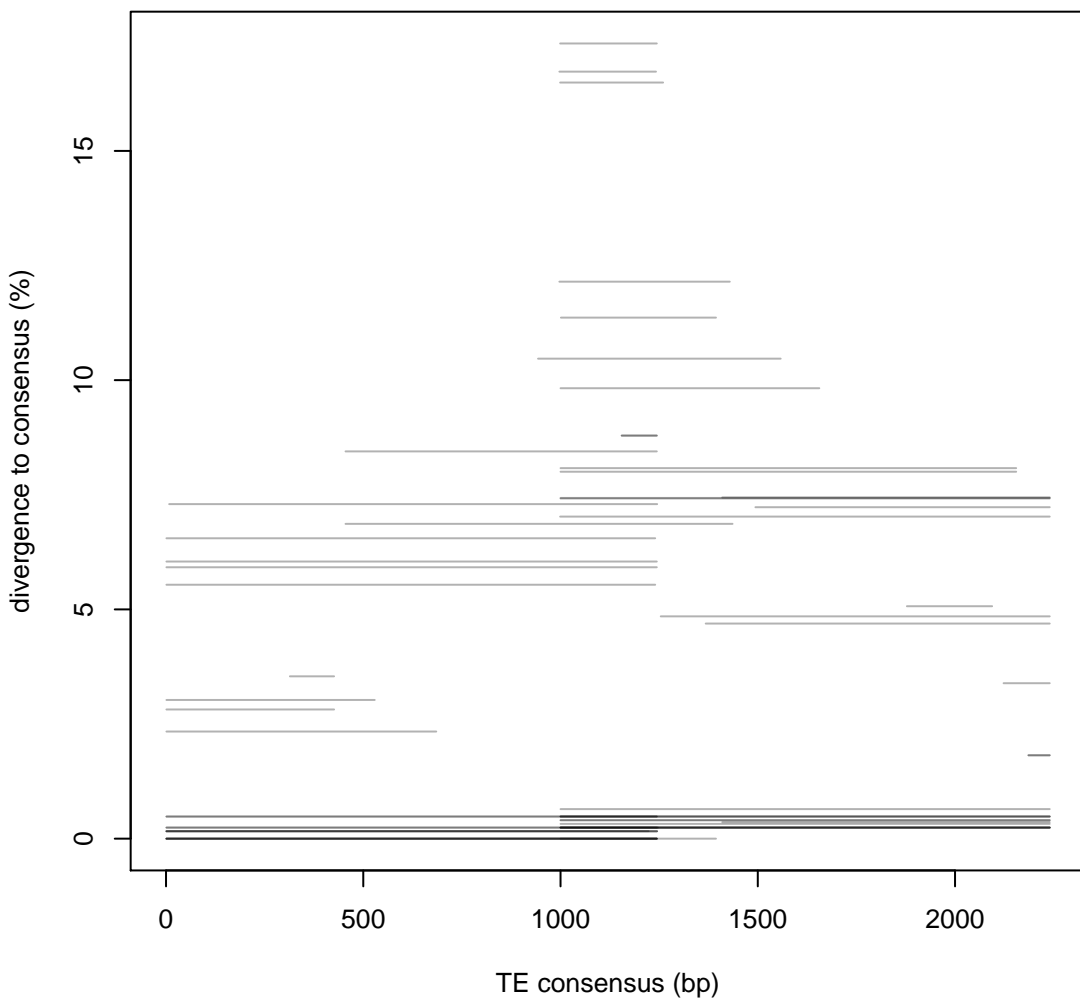
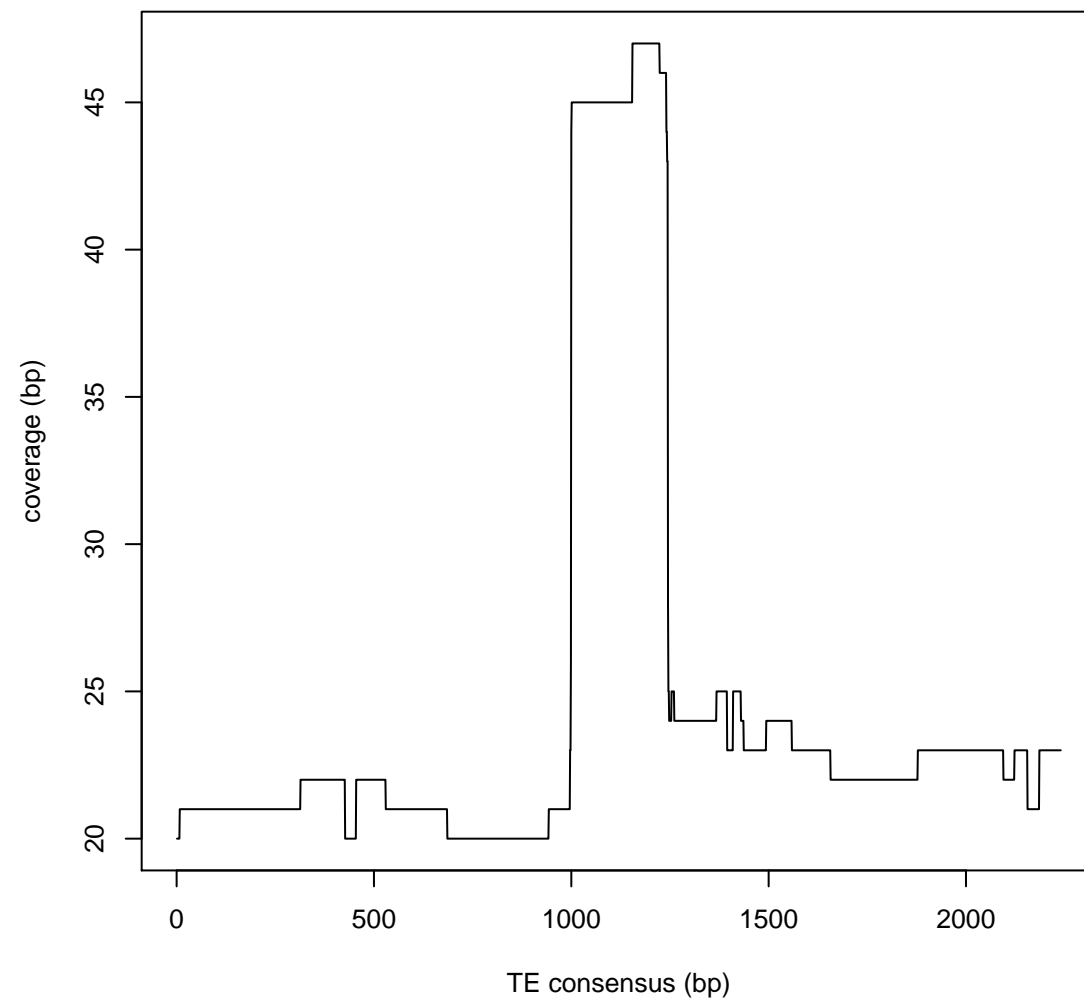


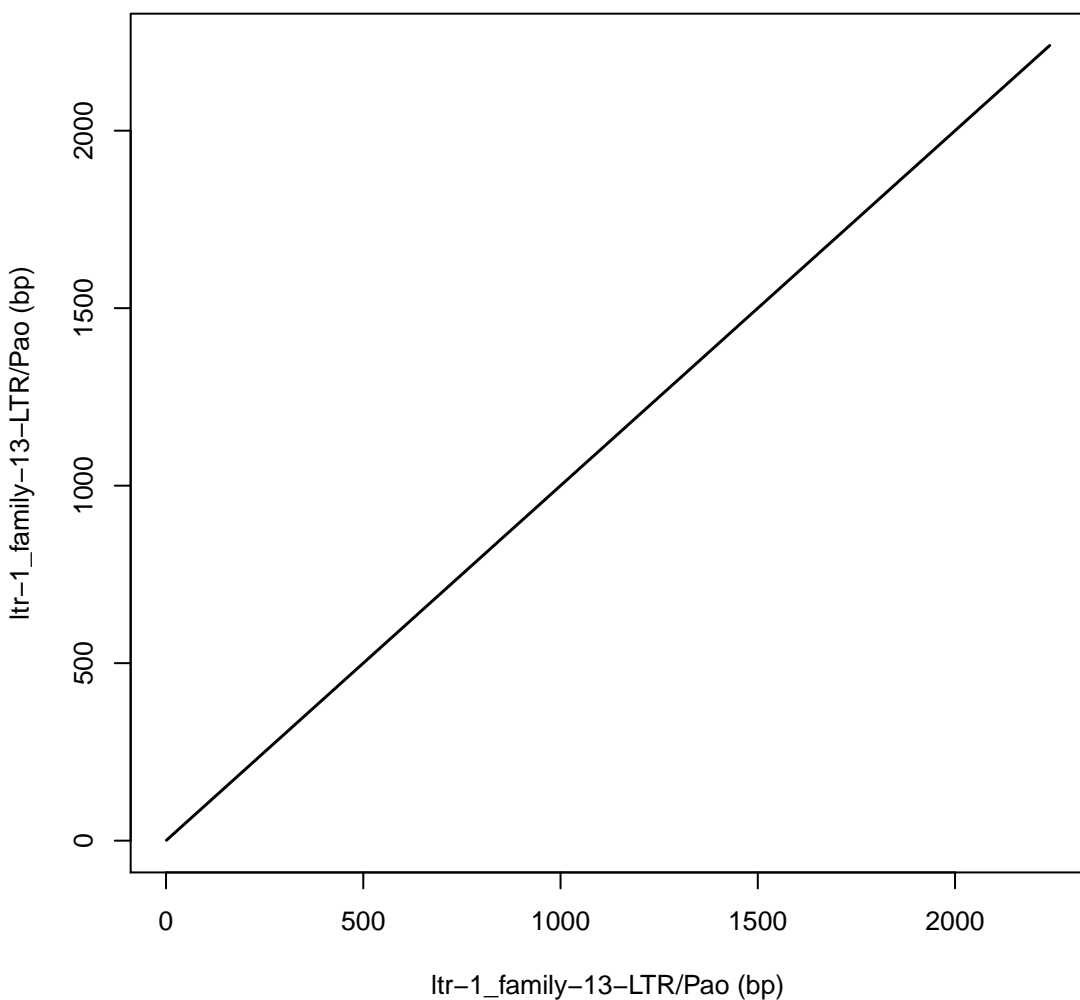
TE: ltr-1_family-13-LTR/Pao
consensus size: 2240bp; fragments: 60; full length: 0 (>=2016bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

