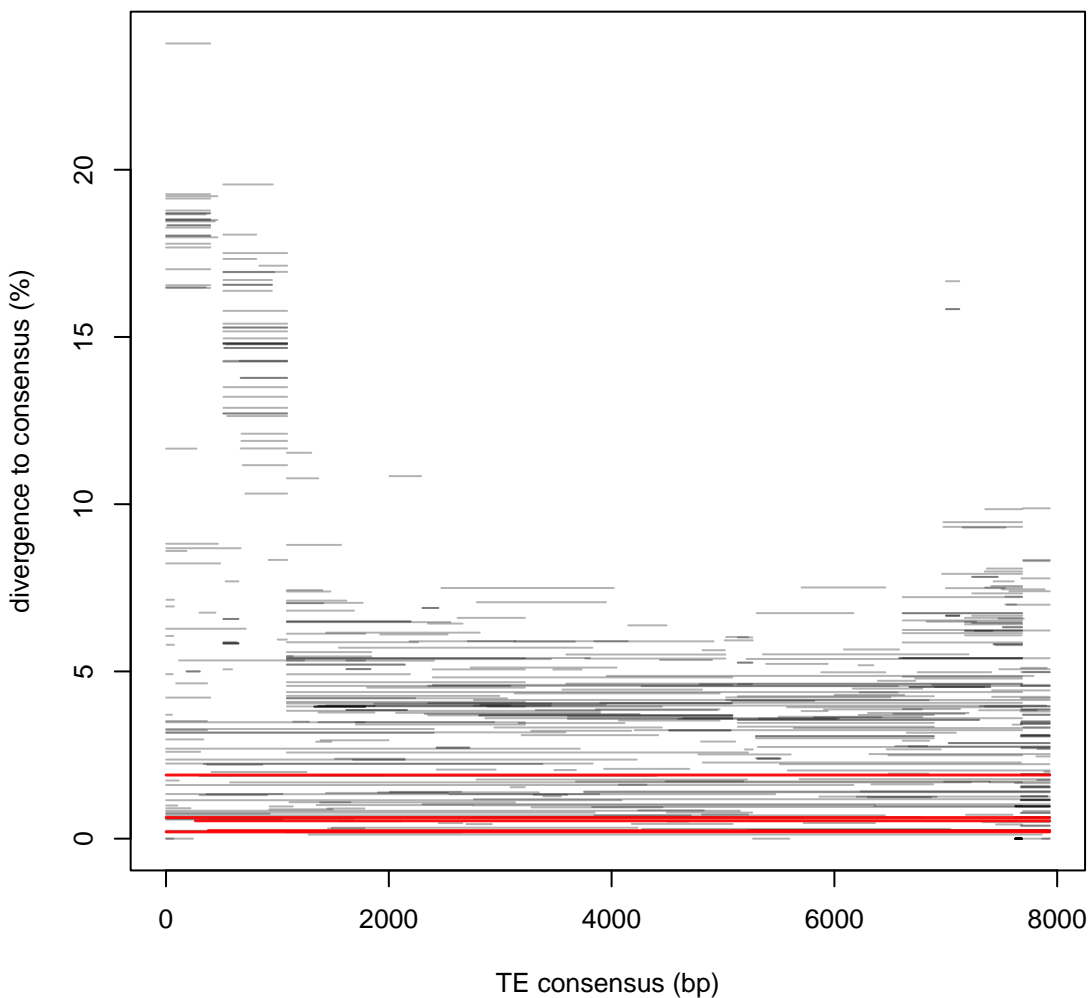
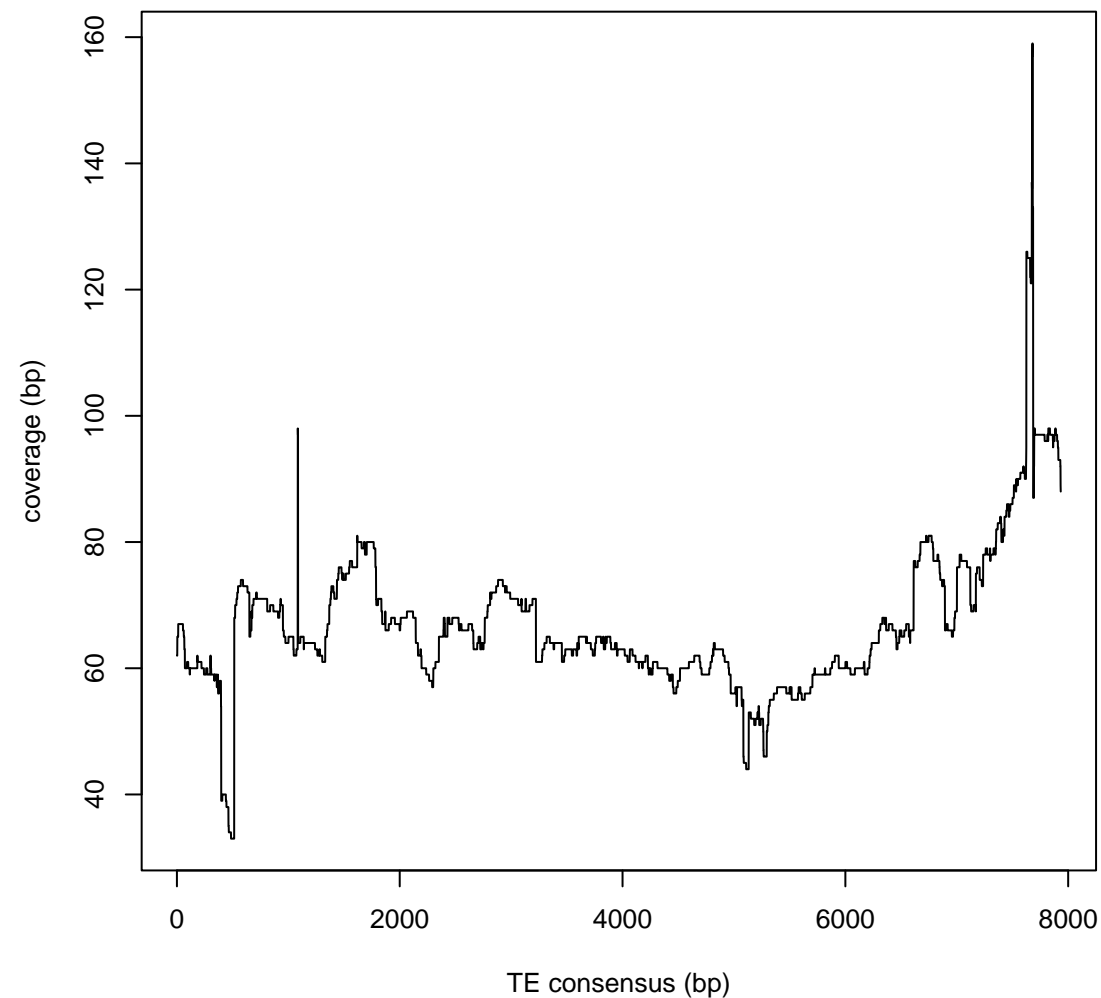


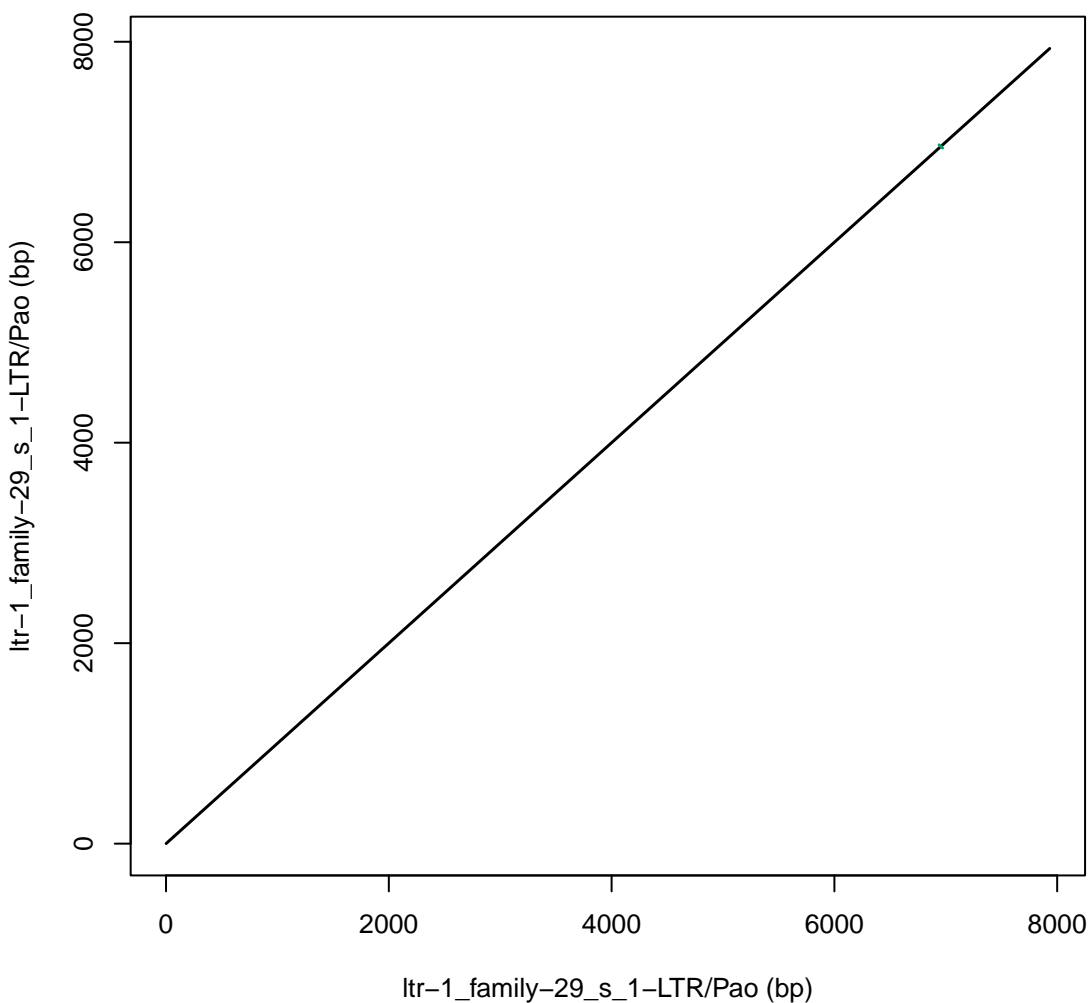
TE: ltr-1_family-29_s_1-LTR/Pao
consensus size: 7933bp; fragments: 529; full length: 5 (>=7139.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

