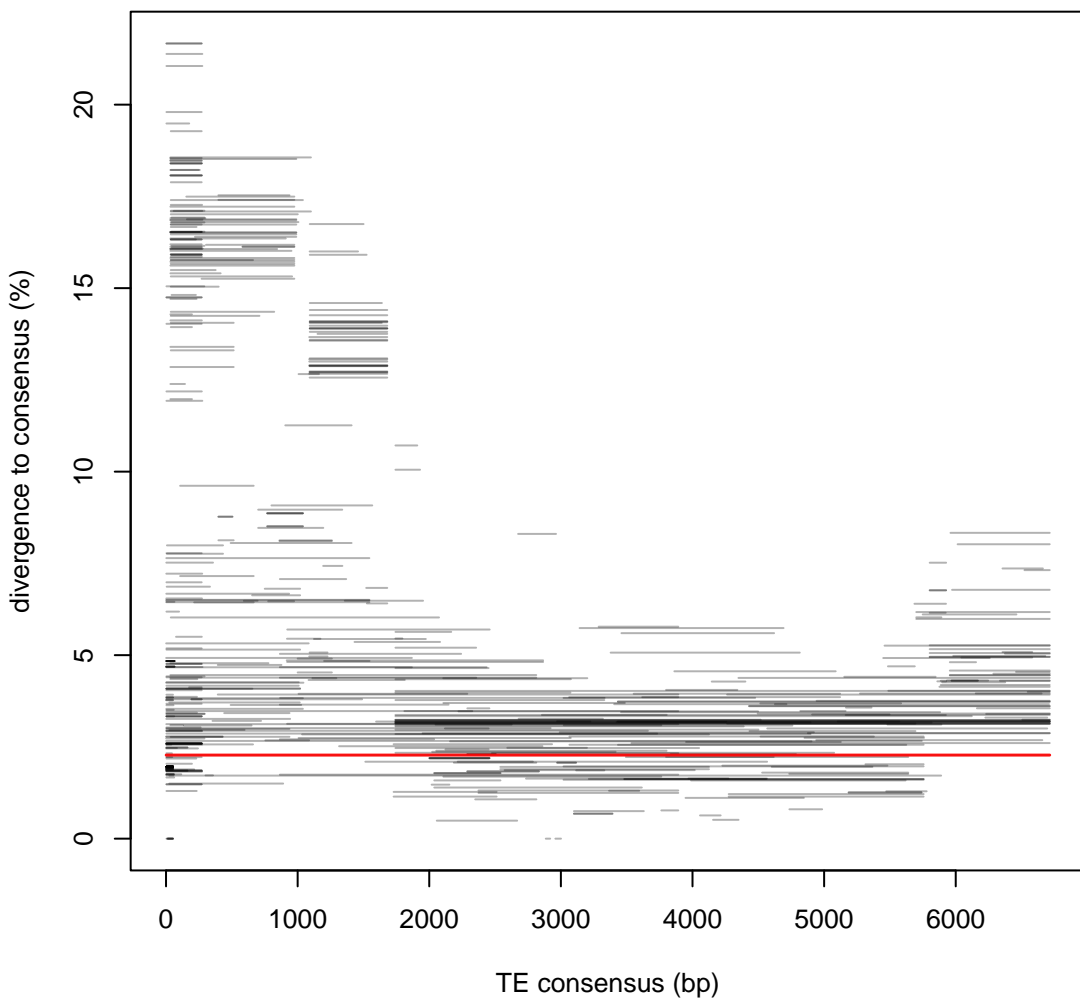
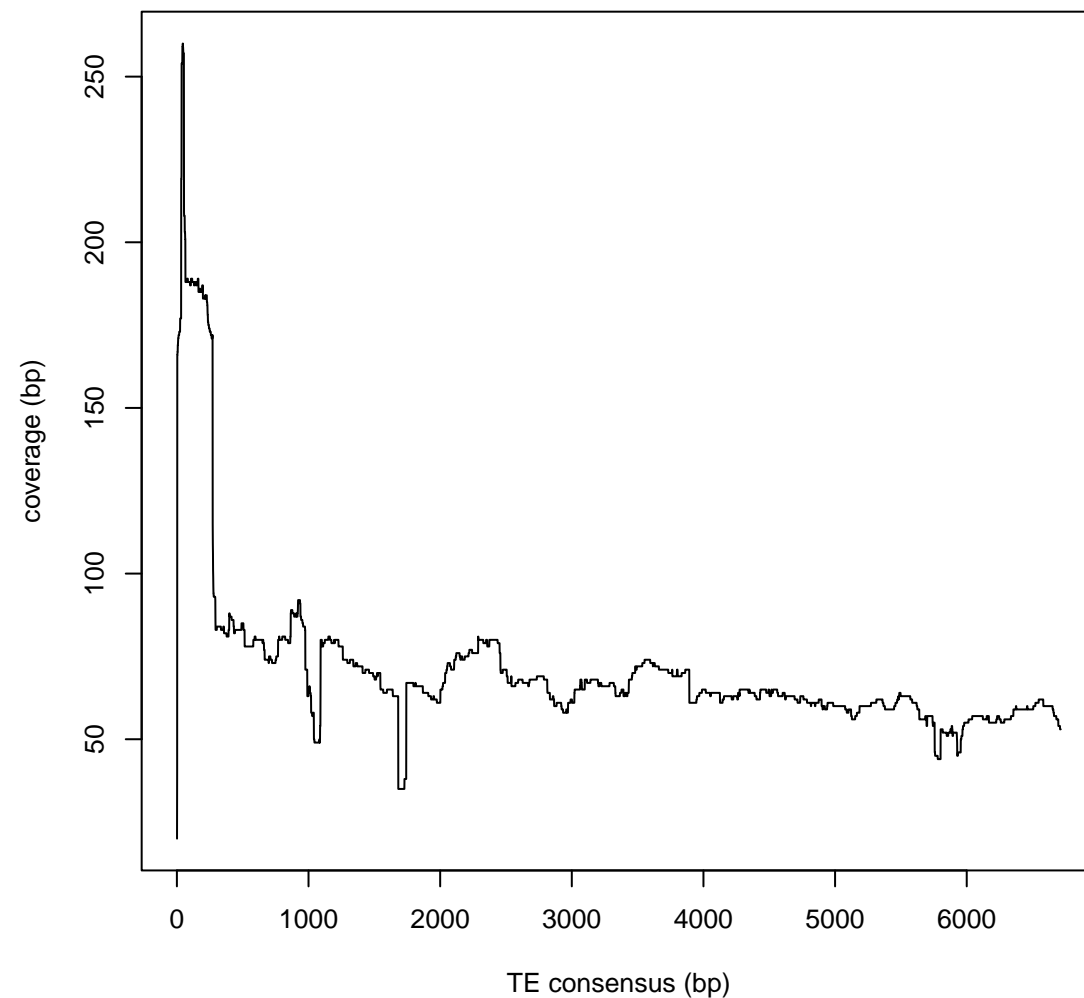


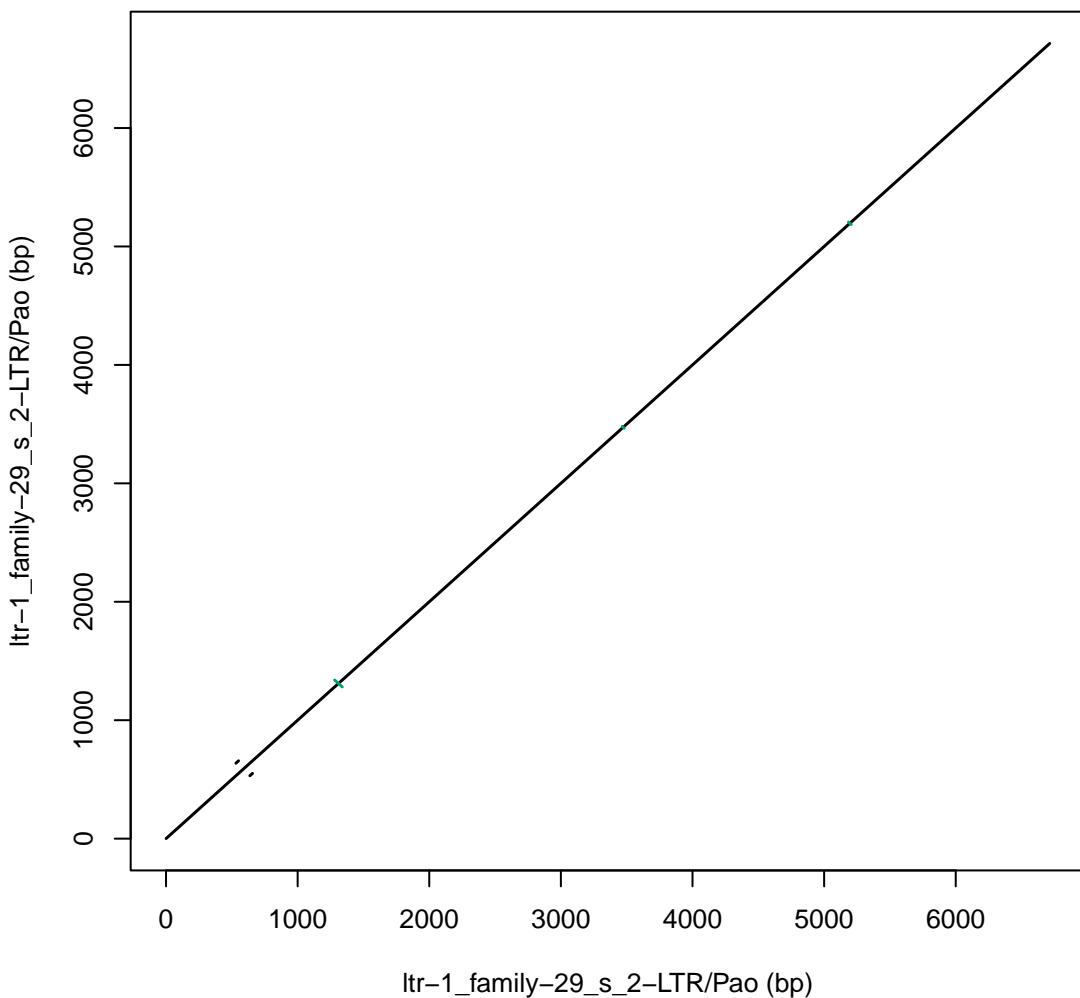
TE: ltr-1_family-29_s_2-LTR/Pao
consensus size: 6714bp; fragments: 572; full length: 1 (>=6042.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

