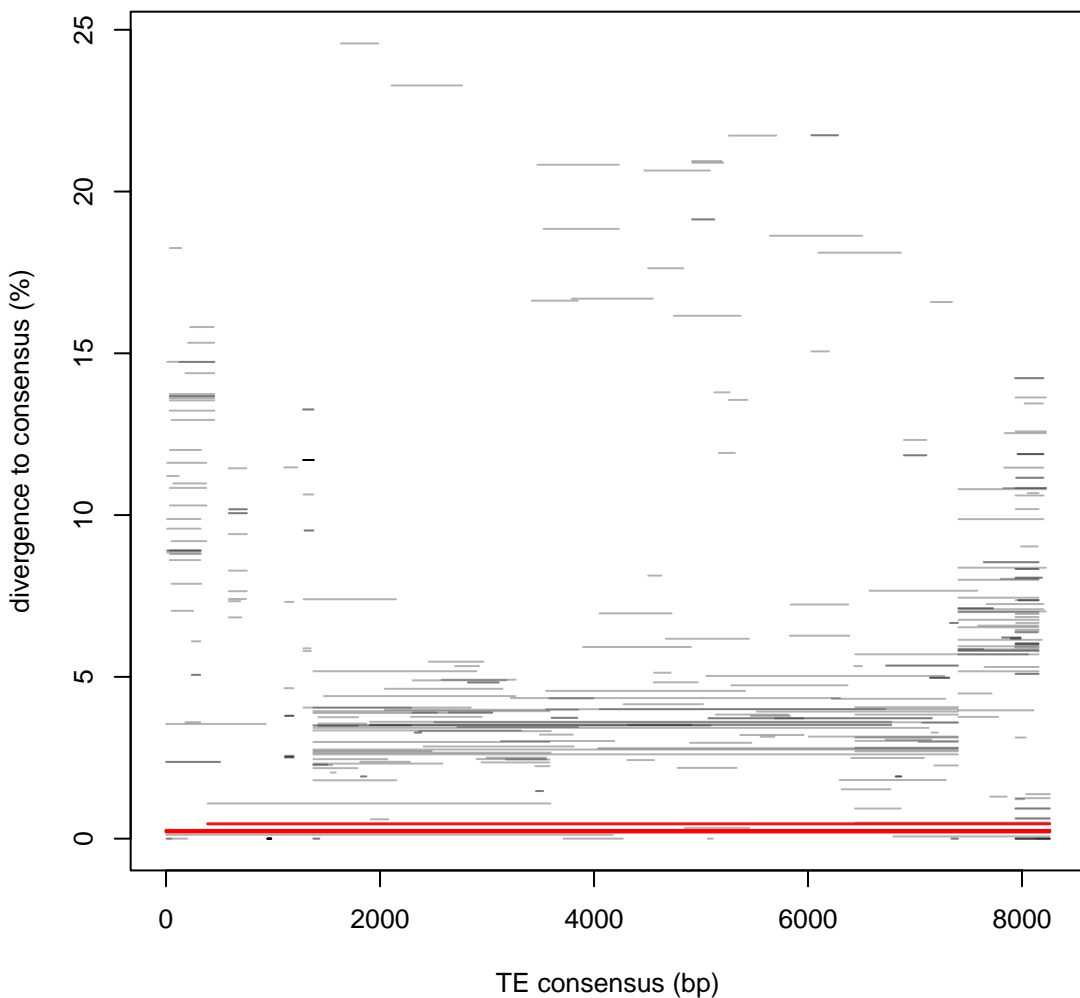
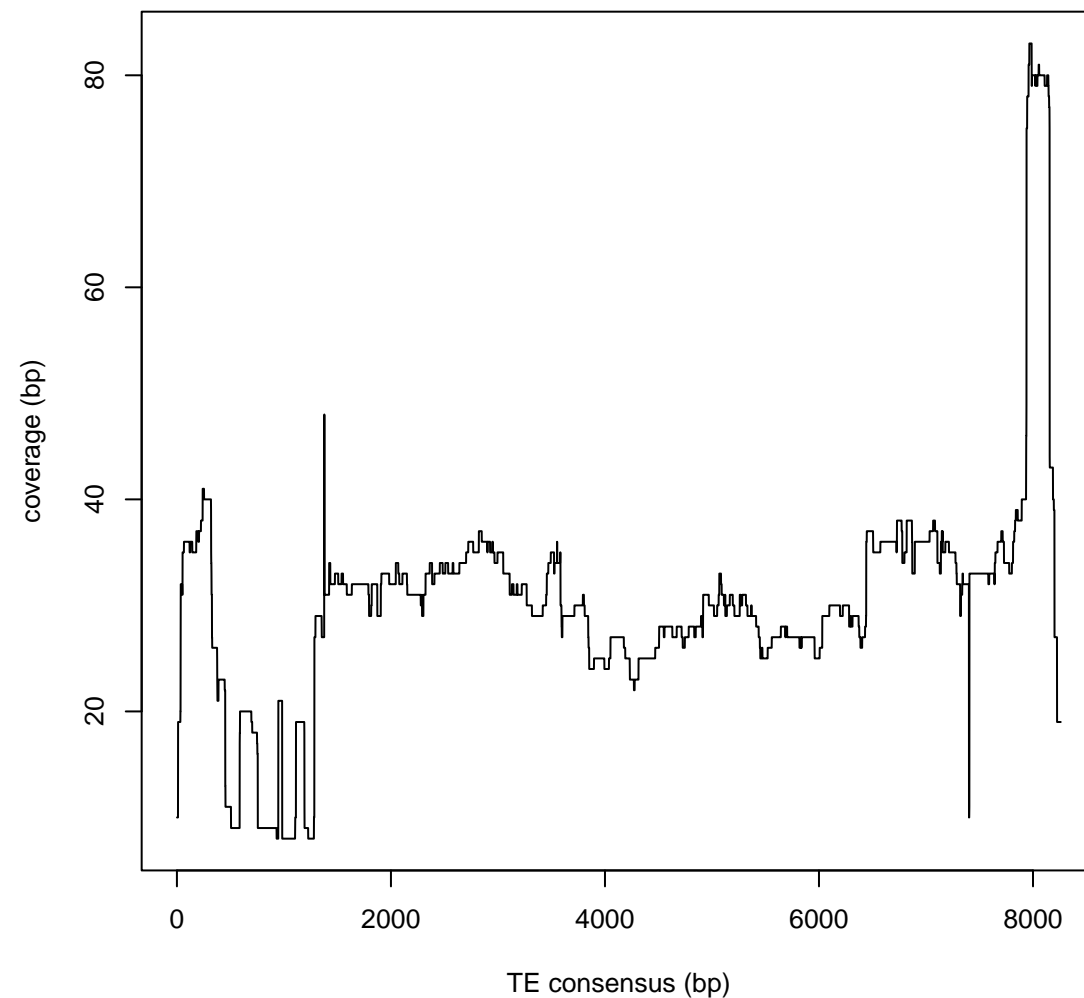


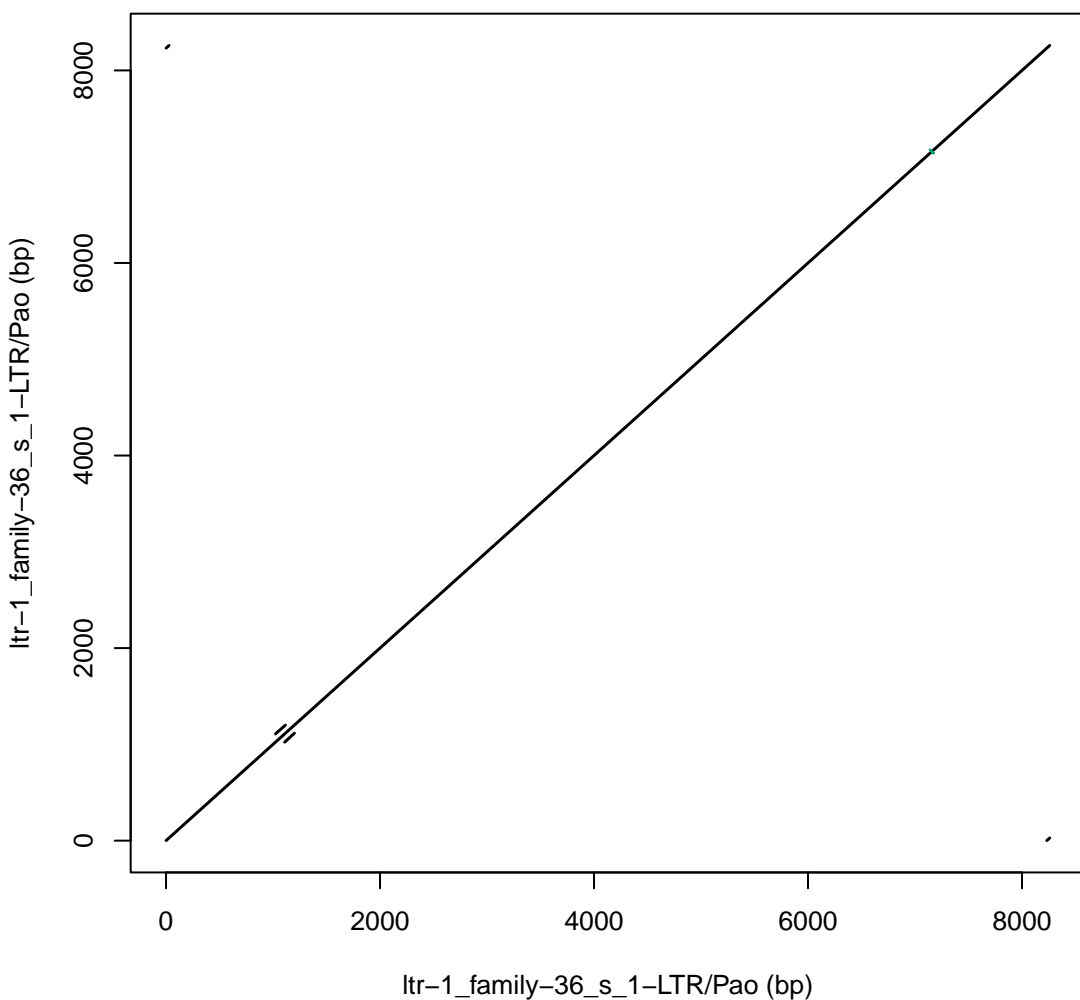
TE: ltr-1_family-36_s_1-LTR/Pao
consensus size: 8259bp; fragments: 341; full length: 6 (>=7433.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

