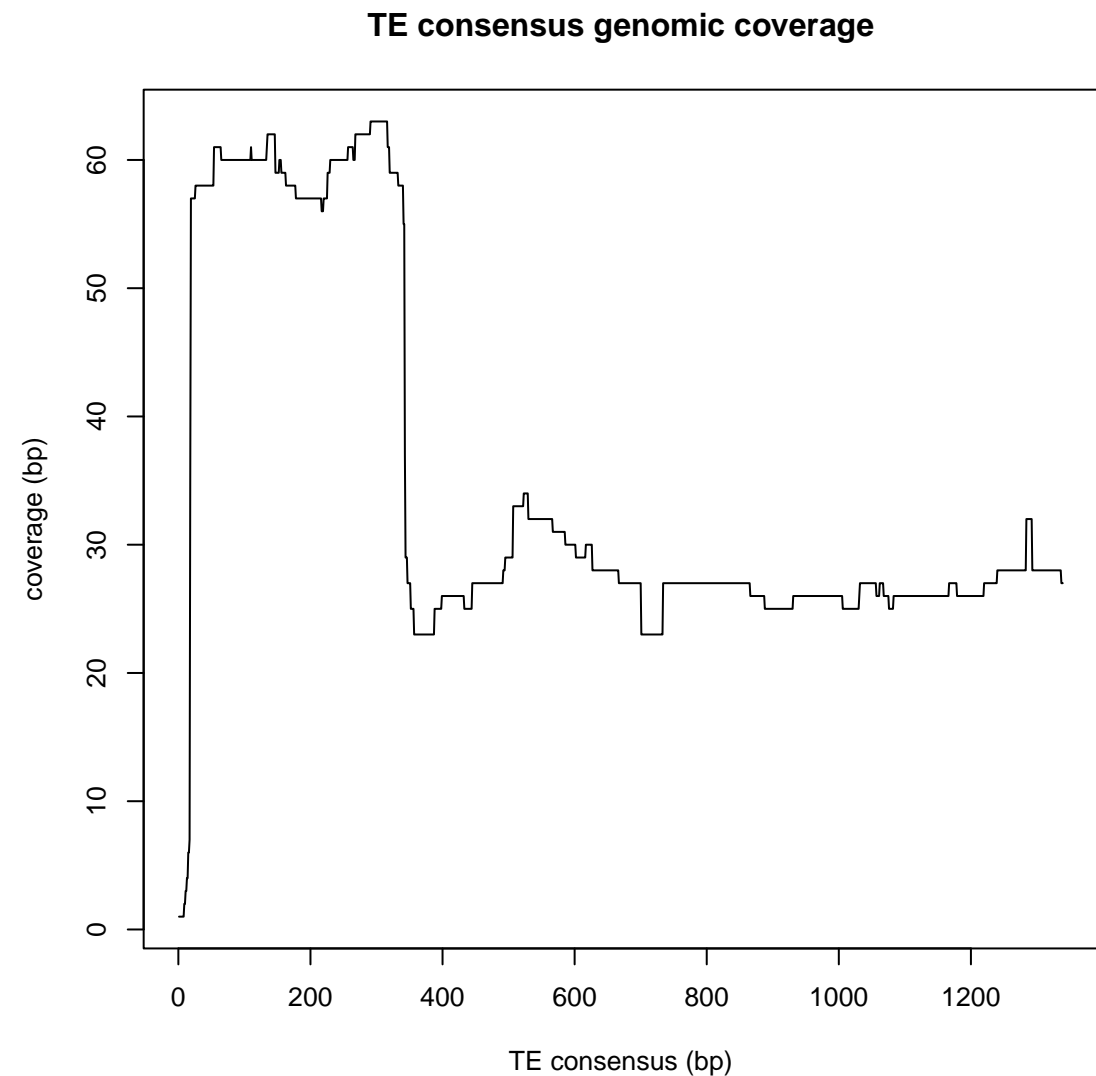
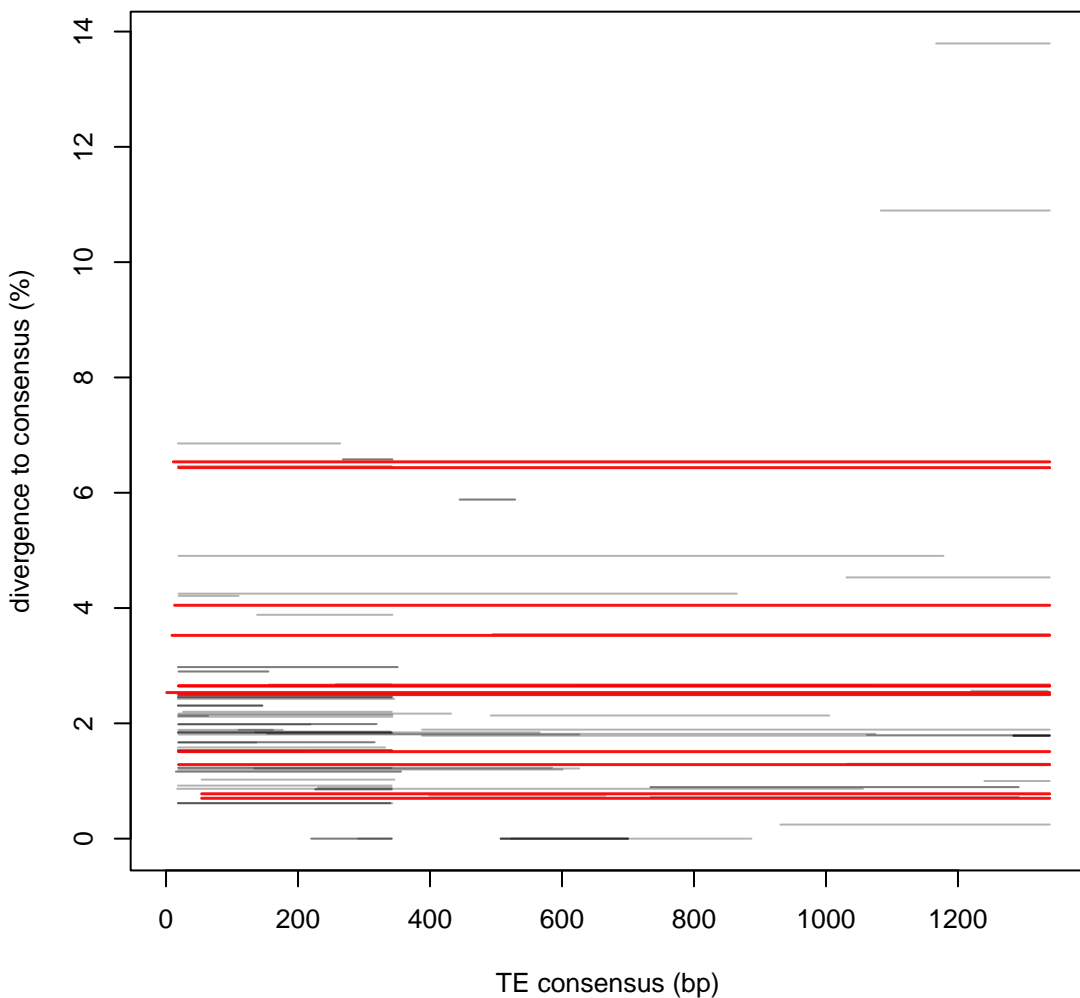
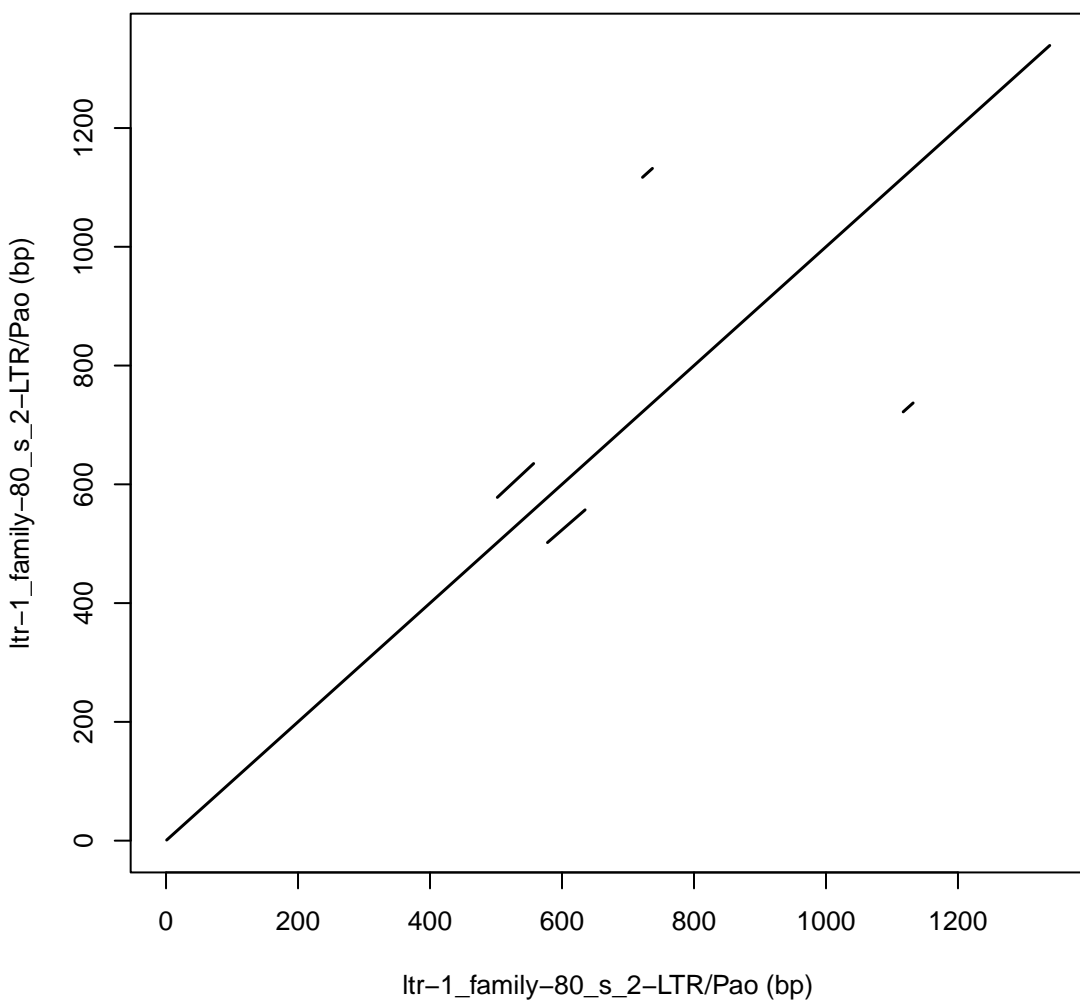


TE: ltr-1_family-80_s_2-LTR/Pao
consensus size: 1339bp; fragments: 105; full length: 12 (>=1205.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

