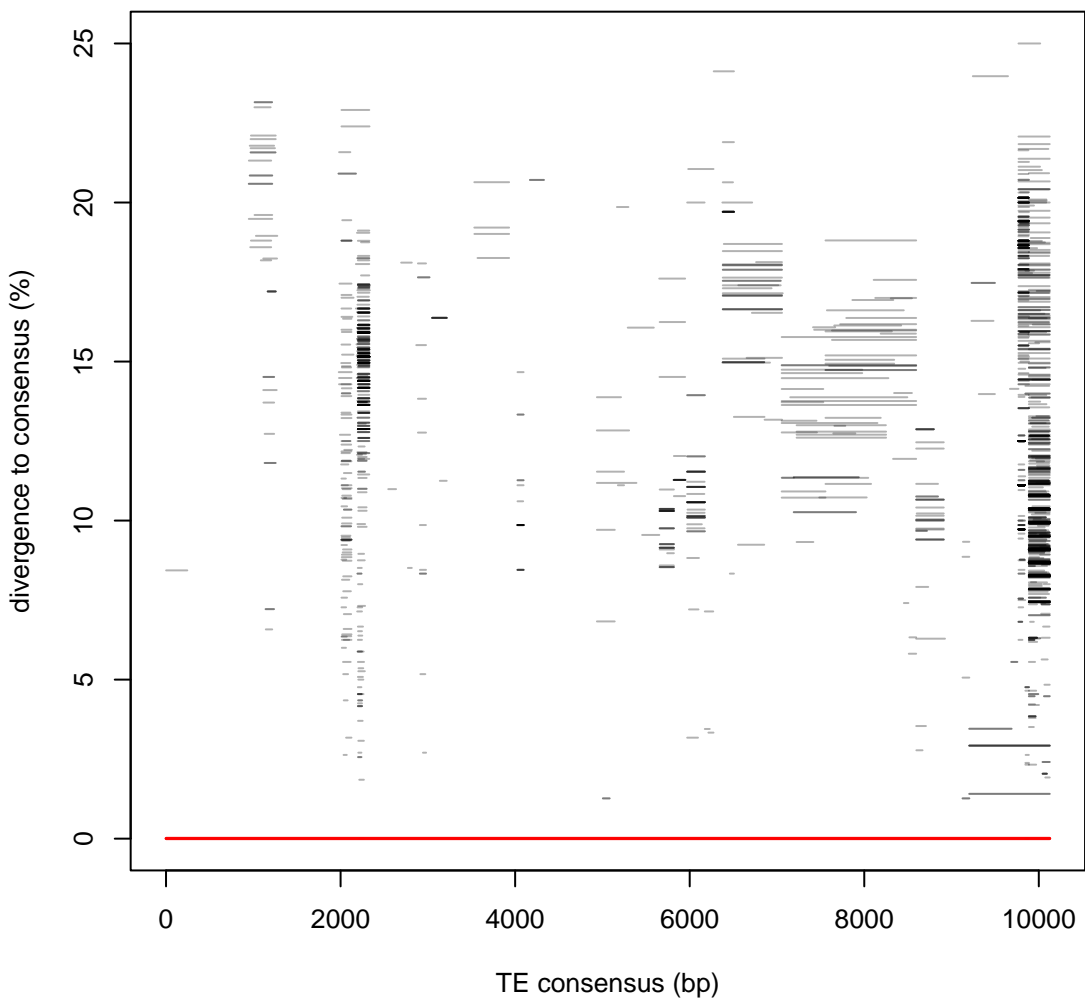
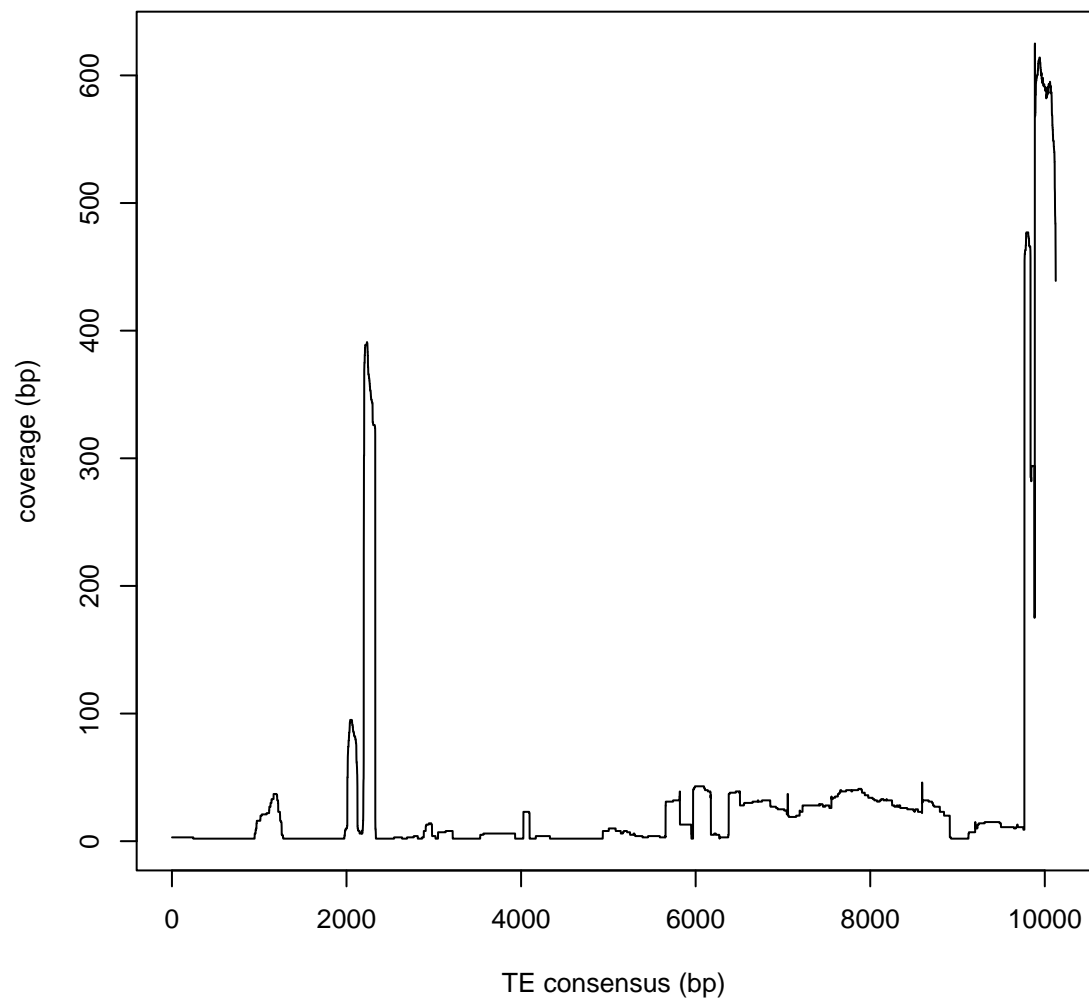


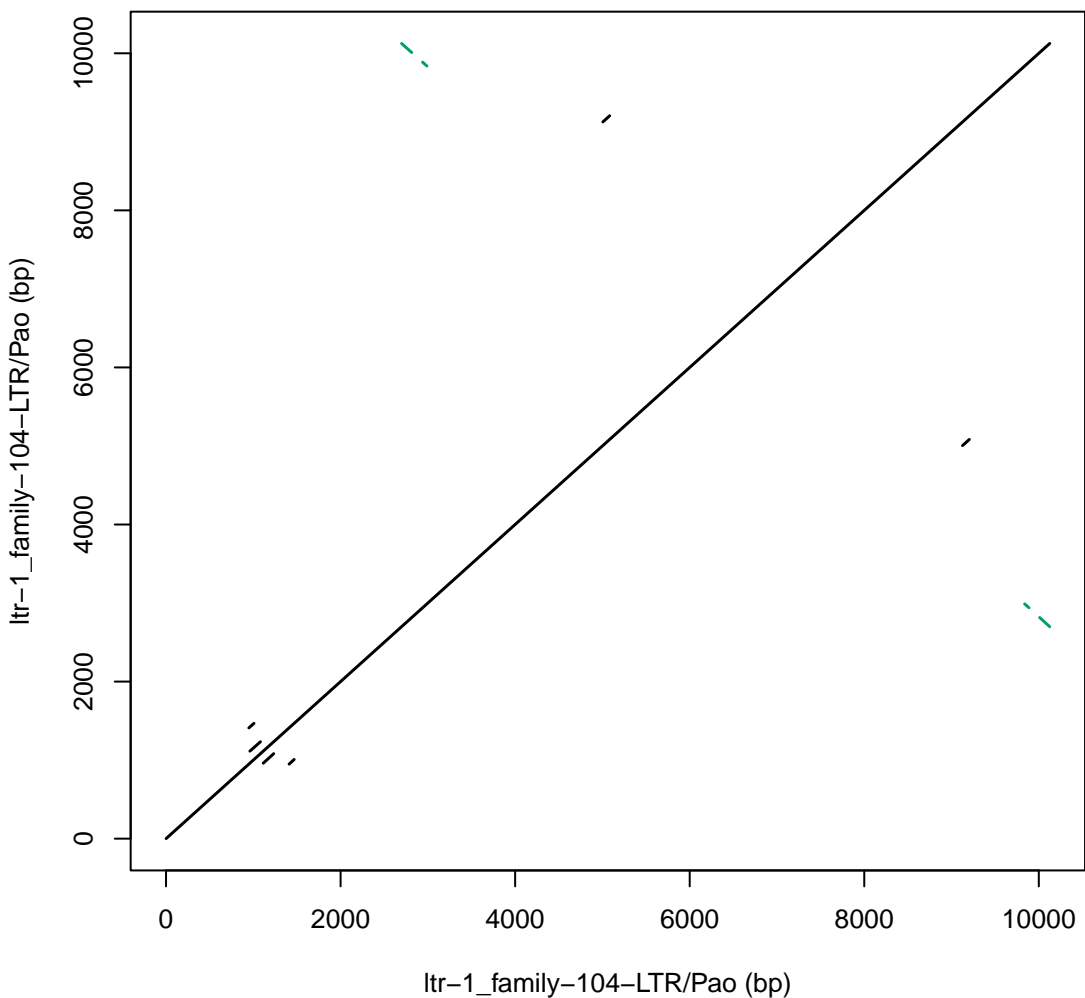
TE: ltr-1_family-104-LTR/Pao
consensus size: 10125bp; fragments: 1852; full length: 2 (>=9112.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

