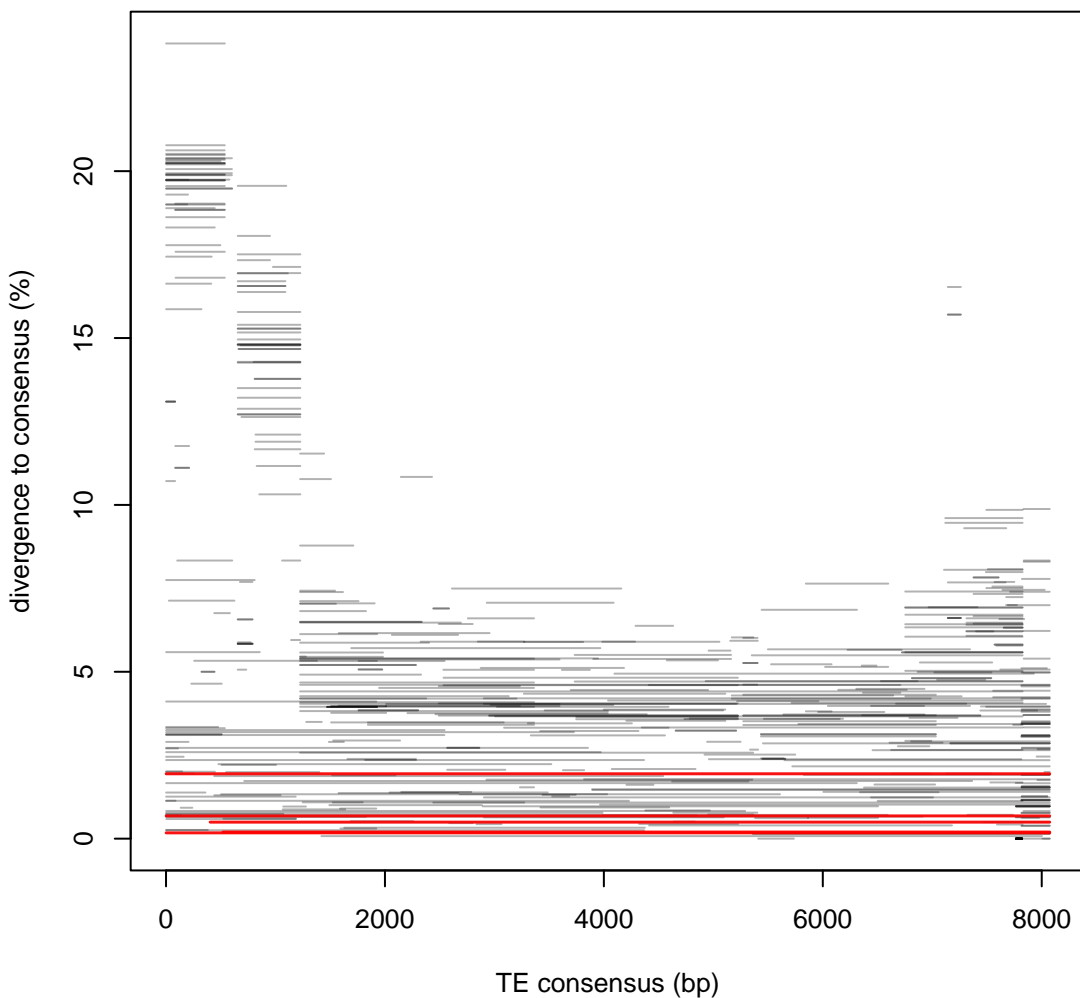
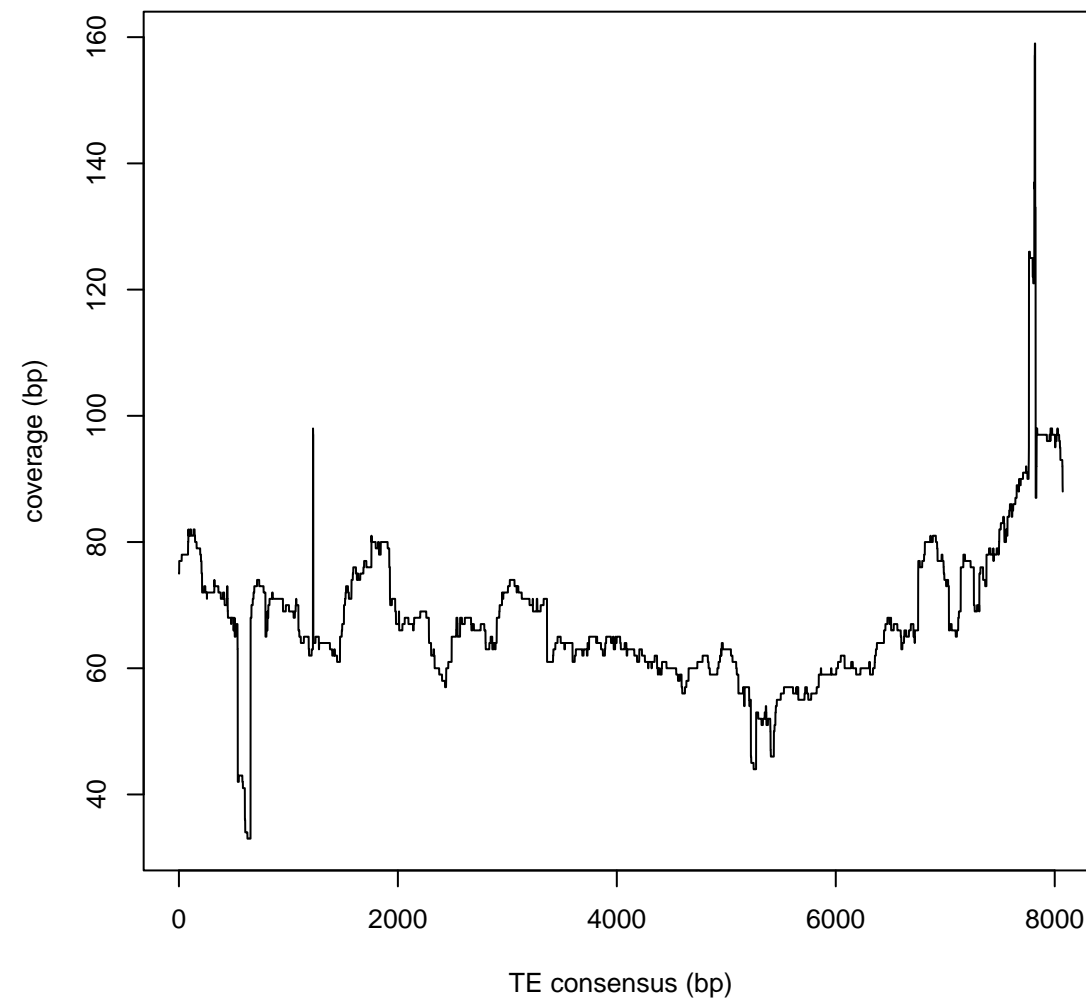


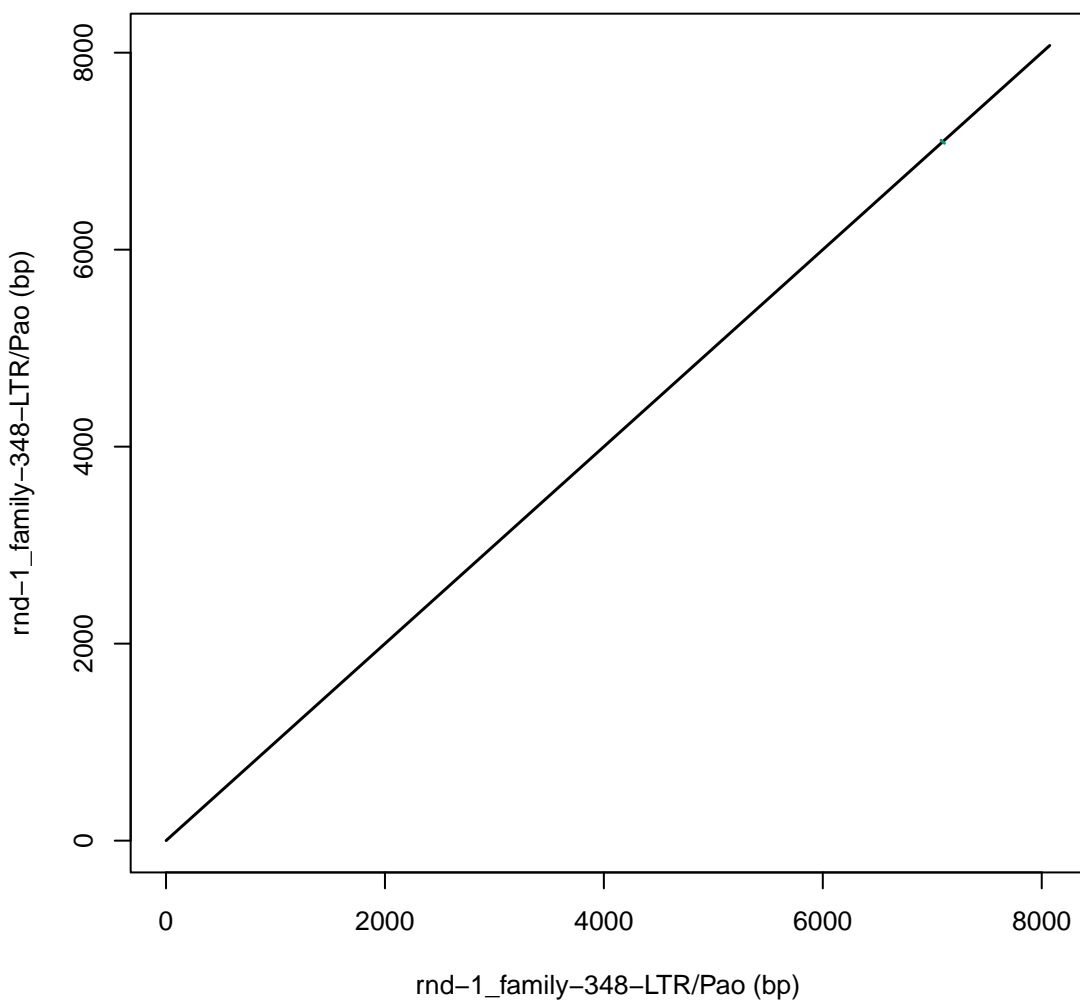
TE: rnd-1\_family-348-LTR/Pao  
consensus size: 8073bp; fragments: 550; full length: 5 ( $\geq 7265.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

