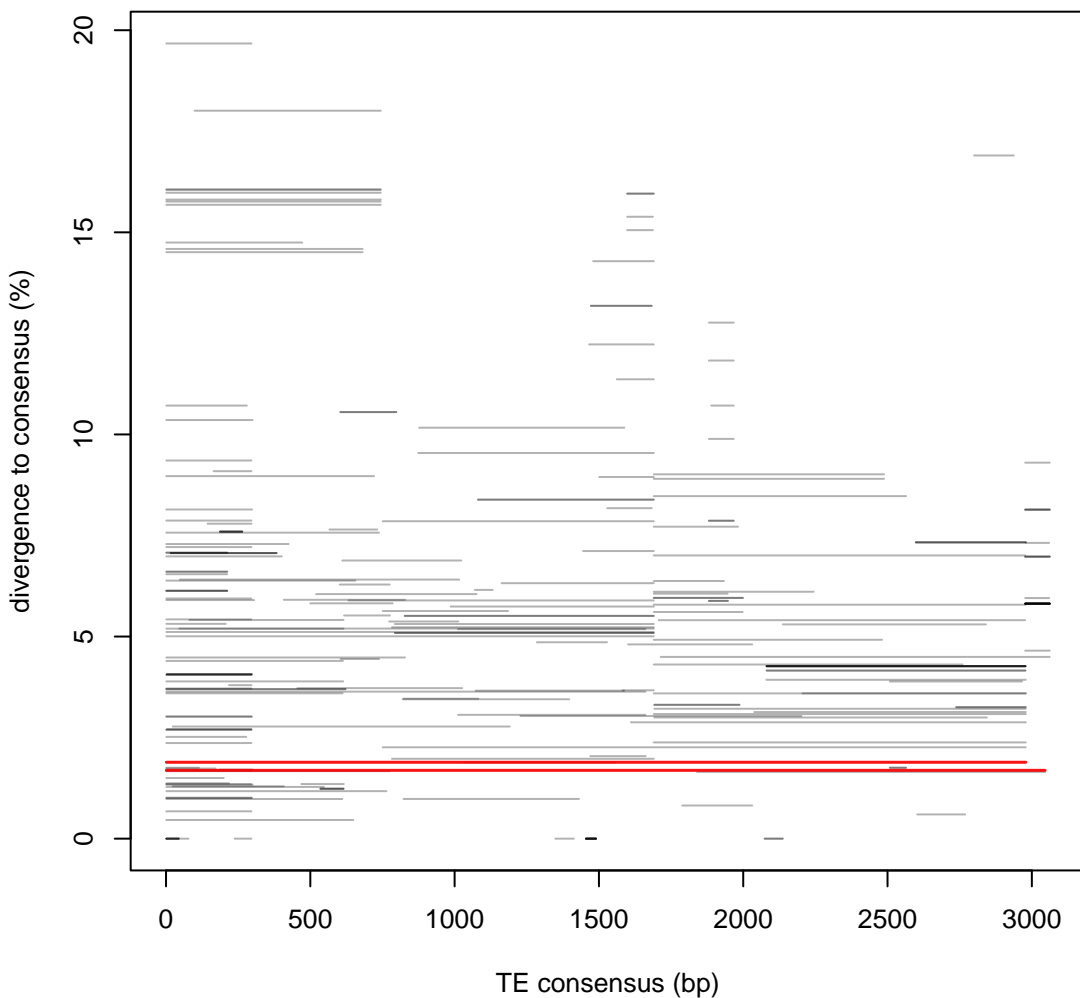
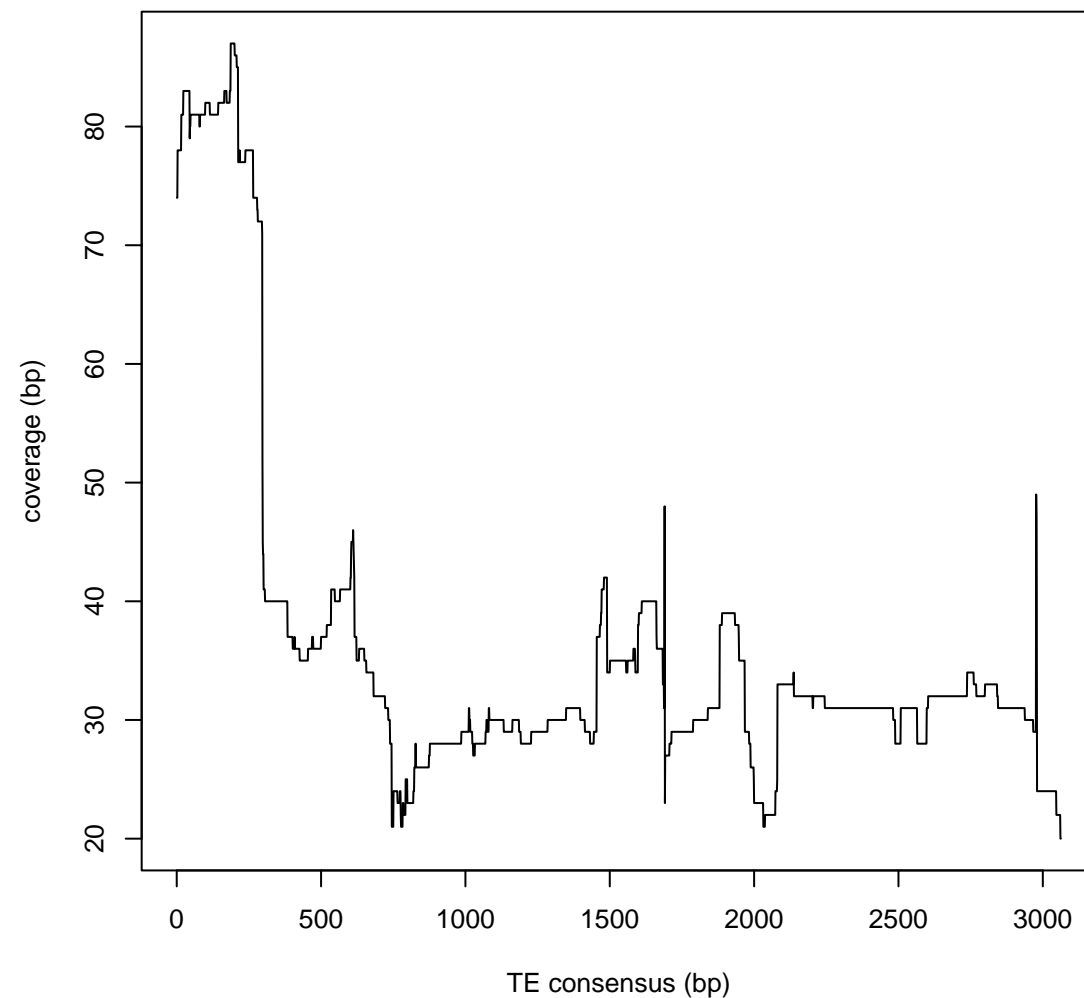


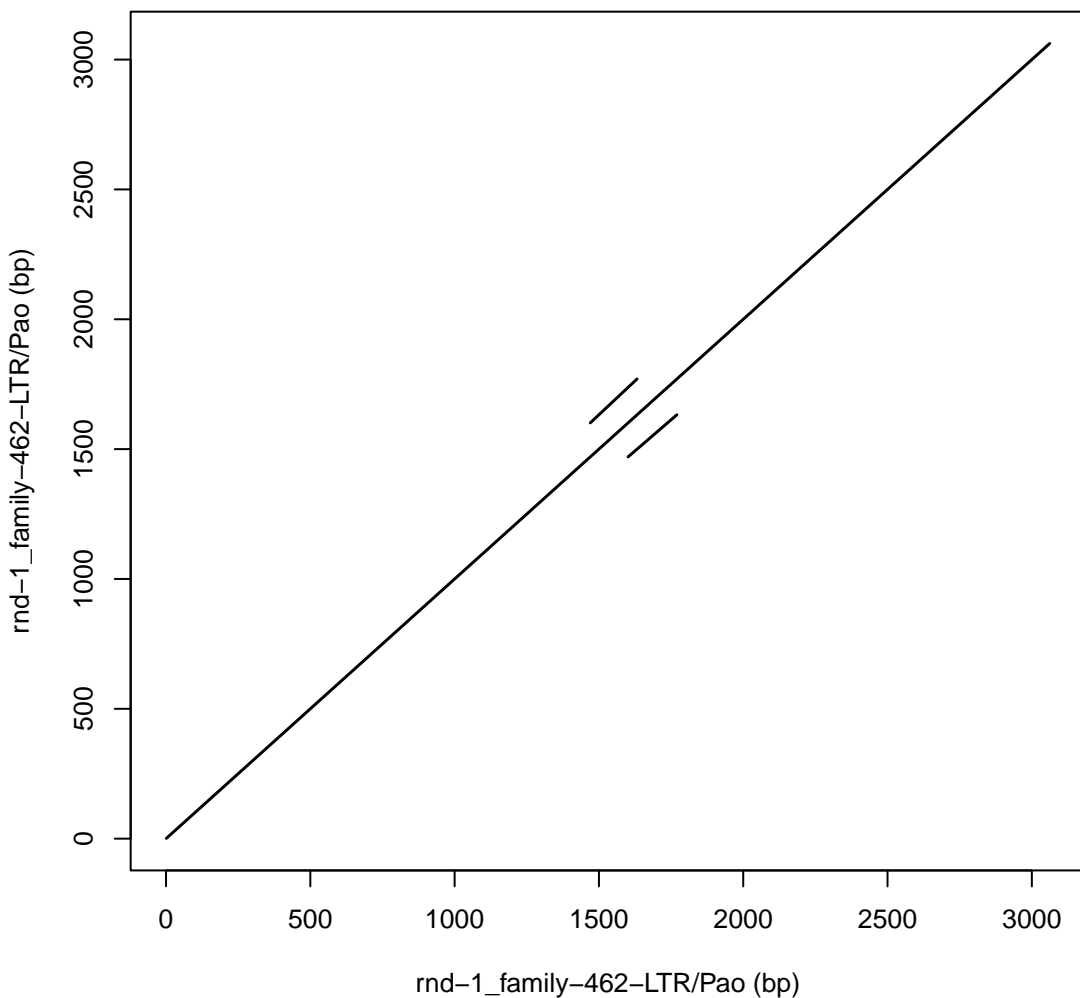
TE: rnd-1\_family-462-LTR/Pao  
consensus size: 3062bp; fragments: 242; full length: 2 (>=2755.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

