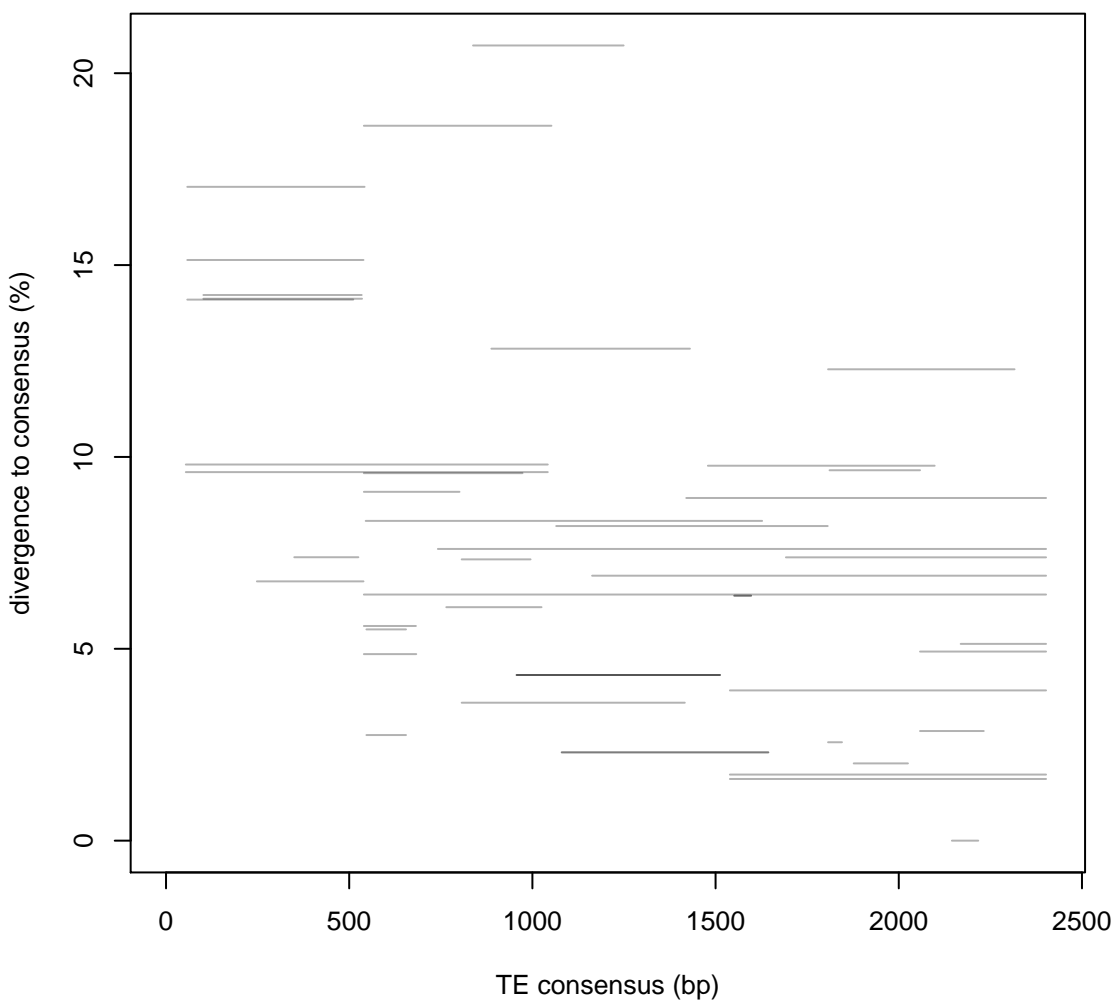
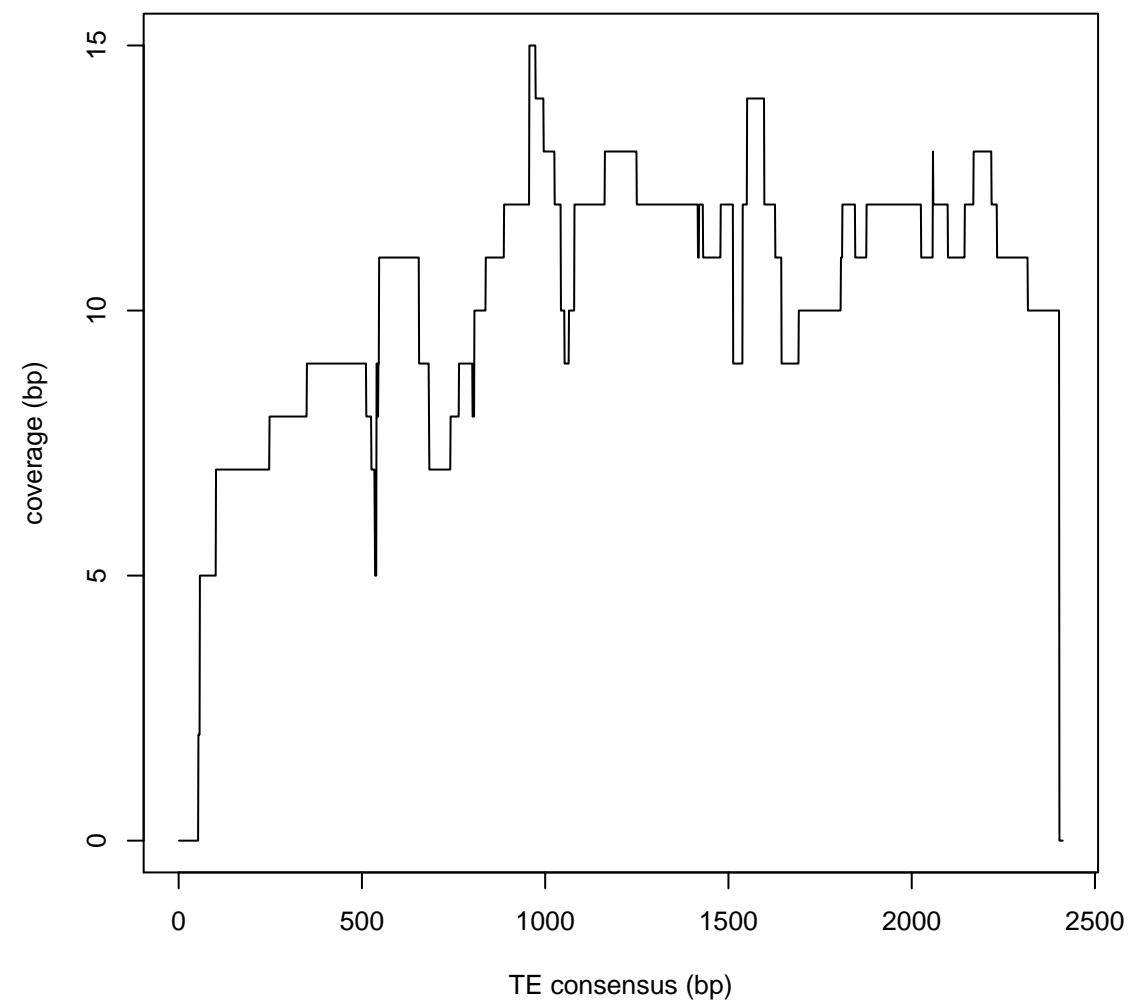


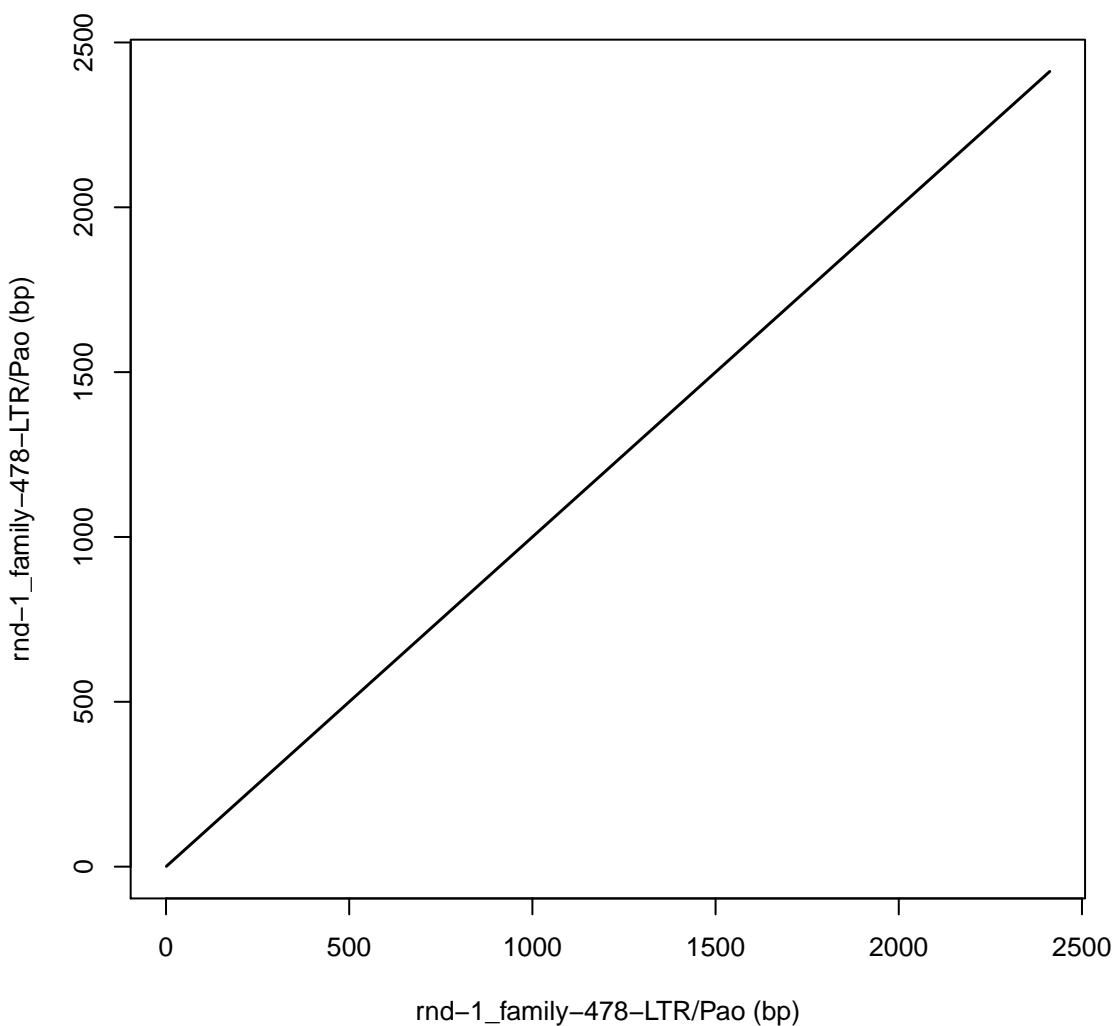
TE: rnd-1\_family-478-LTR/Pao  
consensus size: 2412bp; fragments: 47; full length: 0 (>=2170.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

