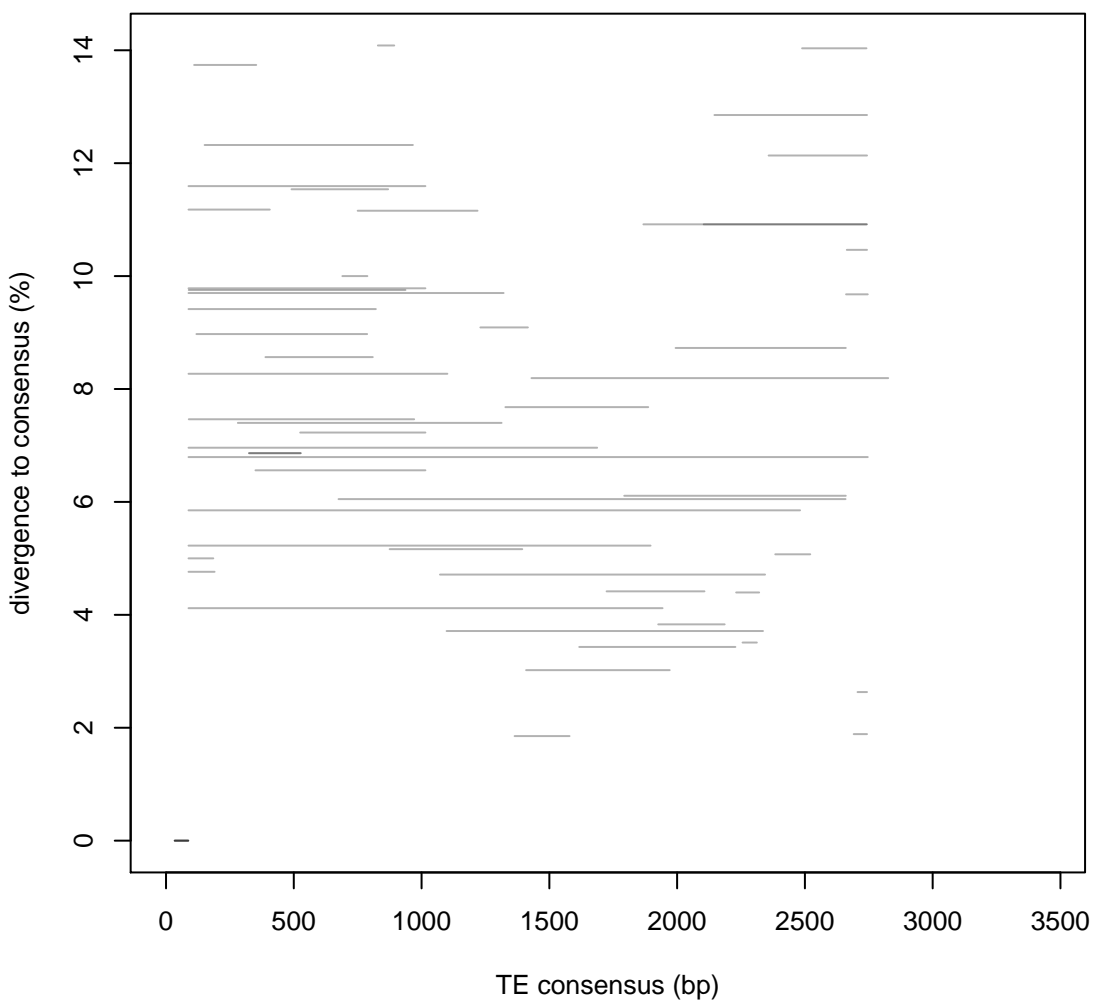
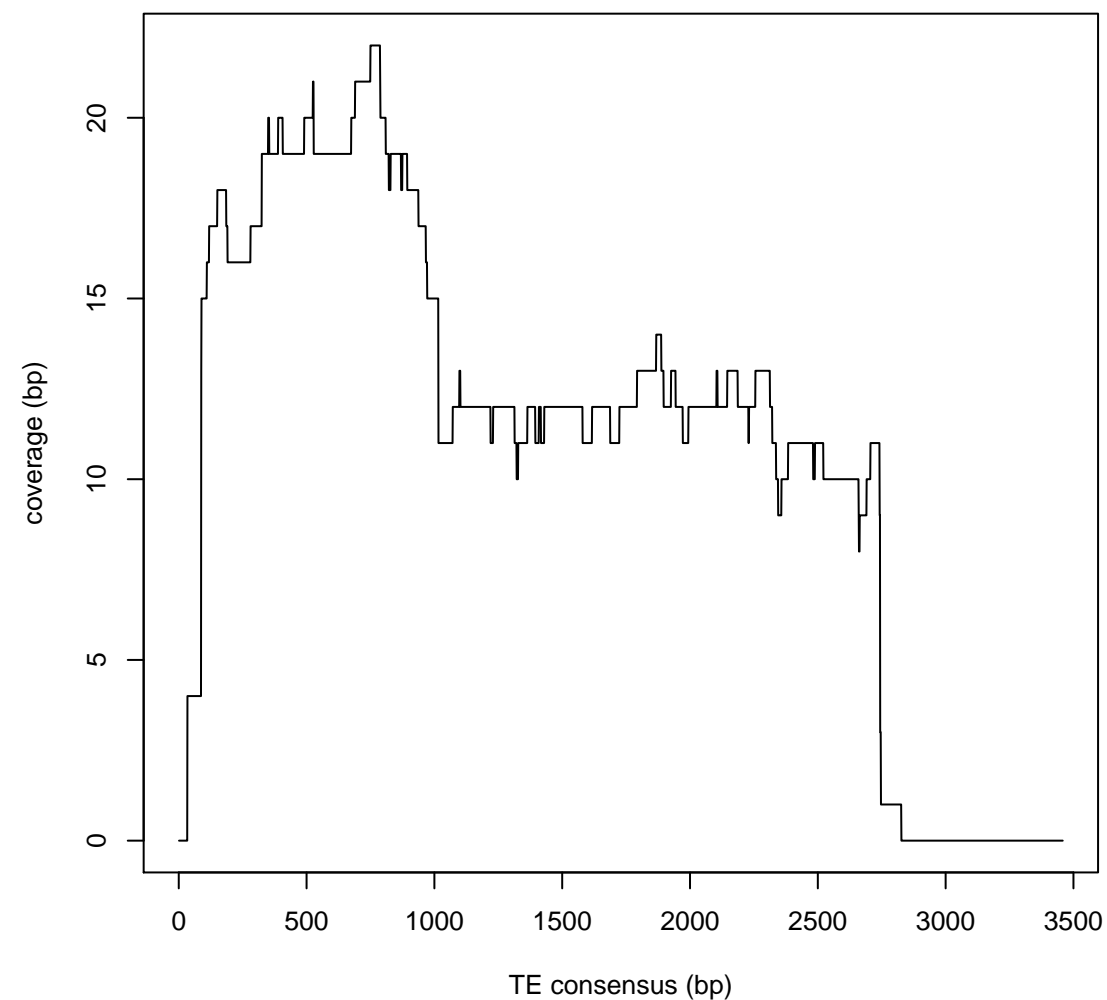


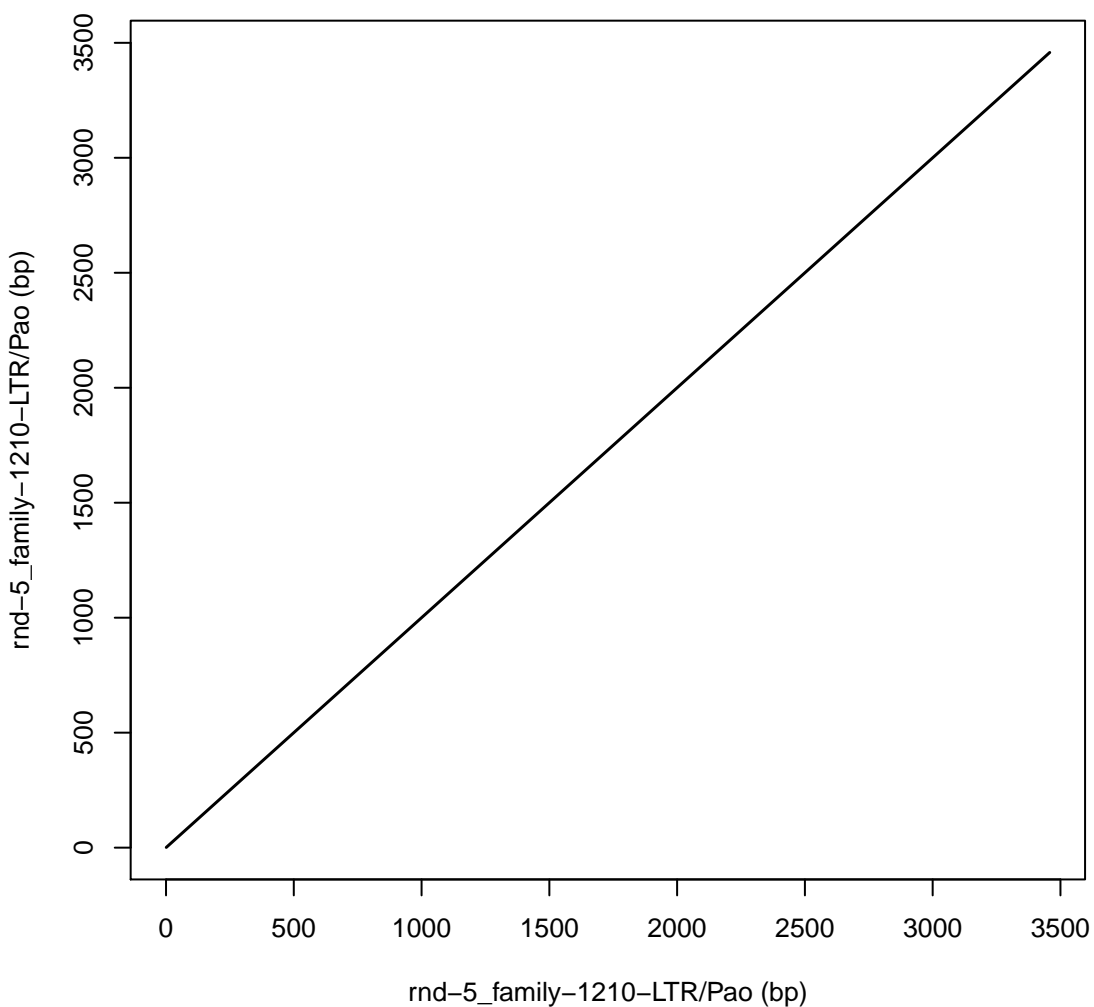
TE: rnd-5\_family-1210-LTR/Pao  
consensus size: 3458bp; fragments: 58; full length: 0 (>=3112.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

