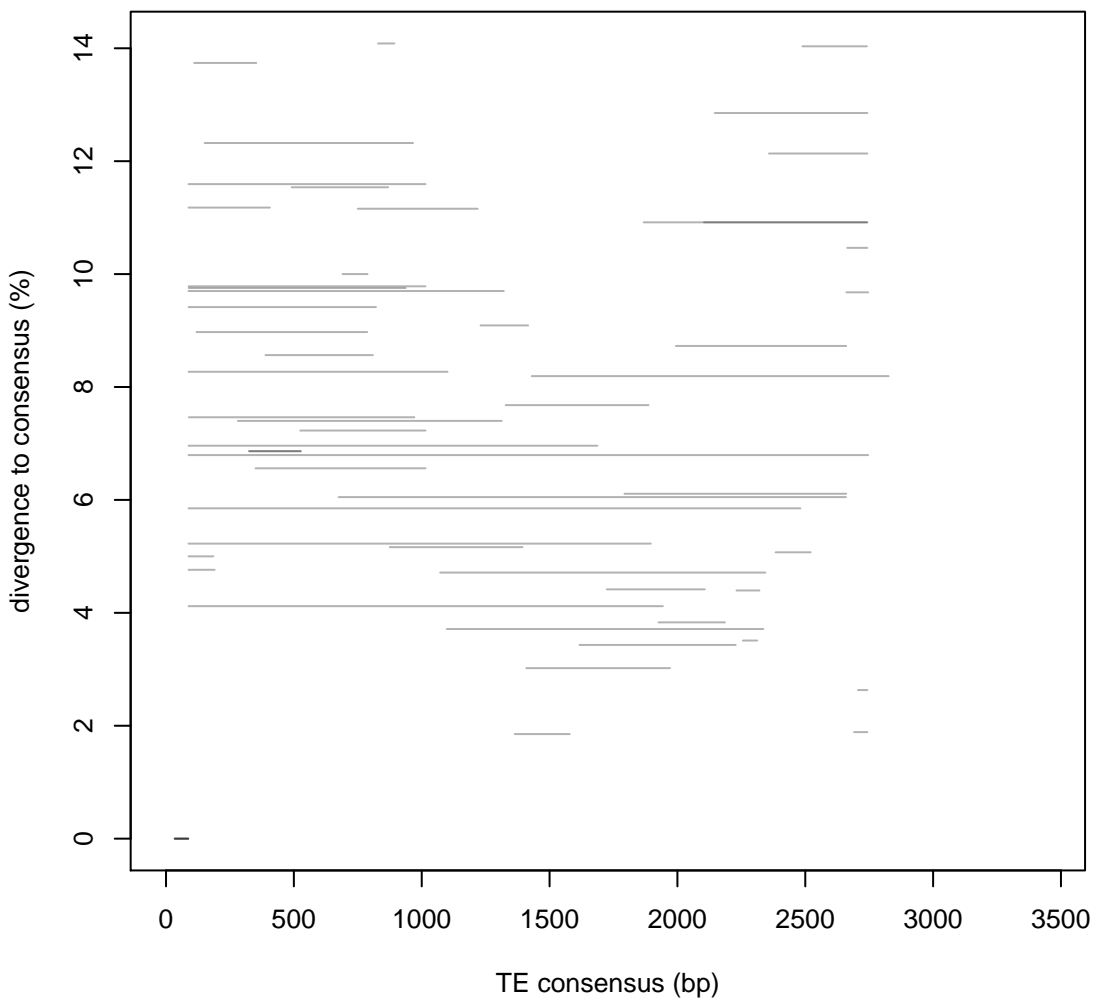
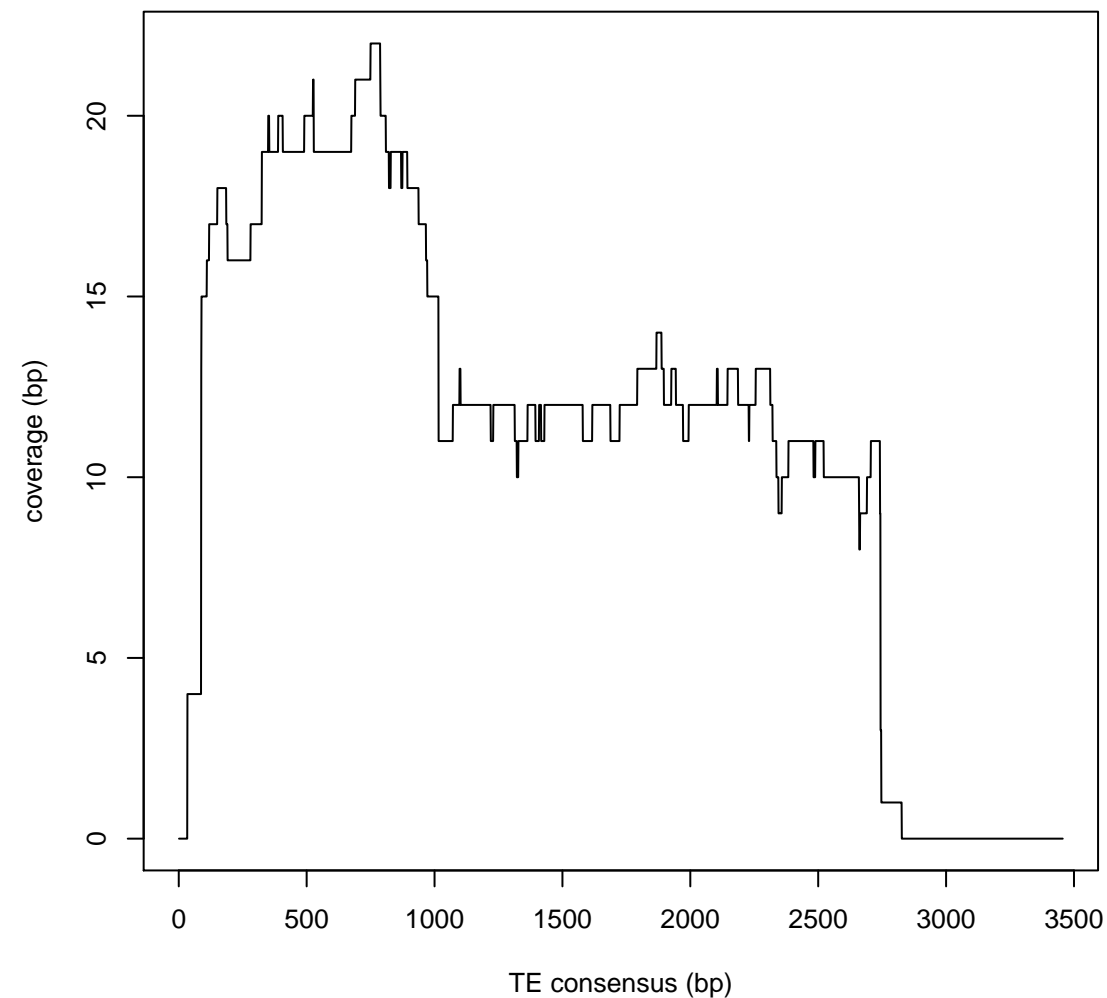


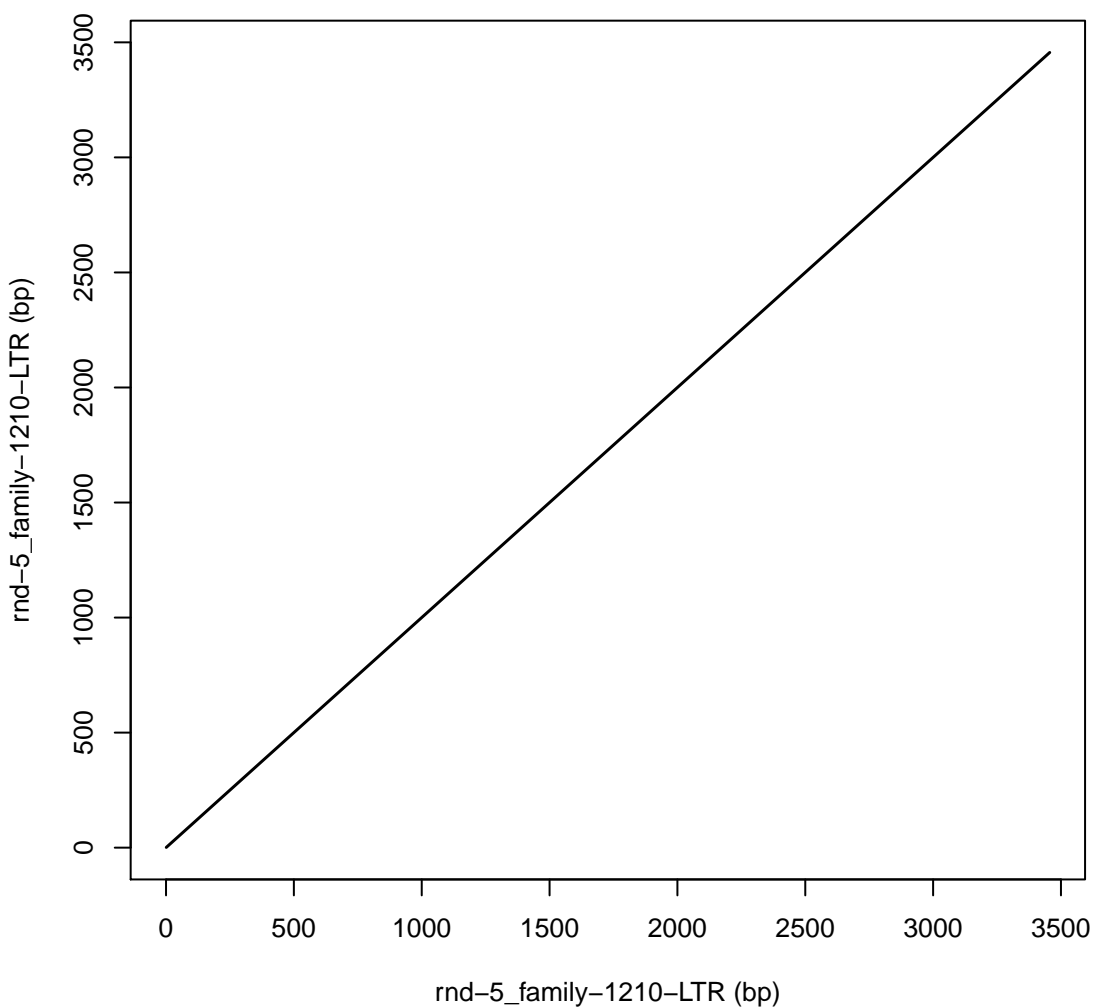
TE: rnd-5_family-1210-LTR
consensus size: 3456bp; fragments: 58; full length: 0 (>=3110.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

