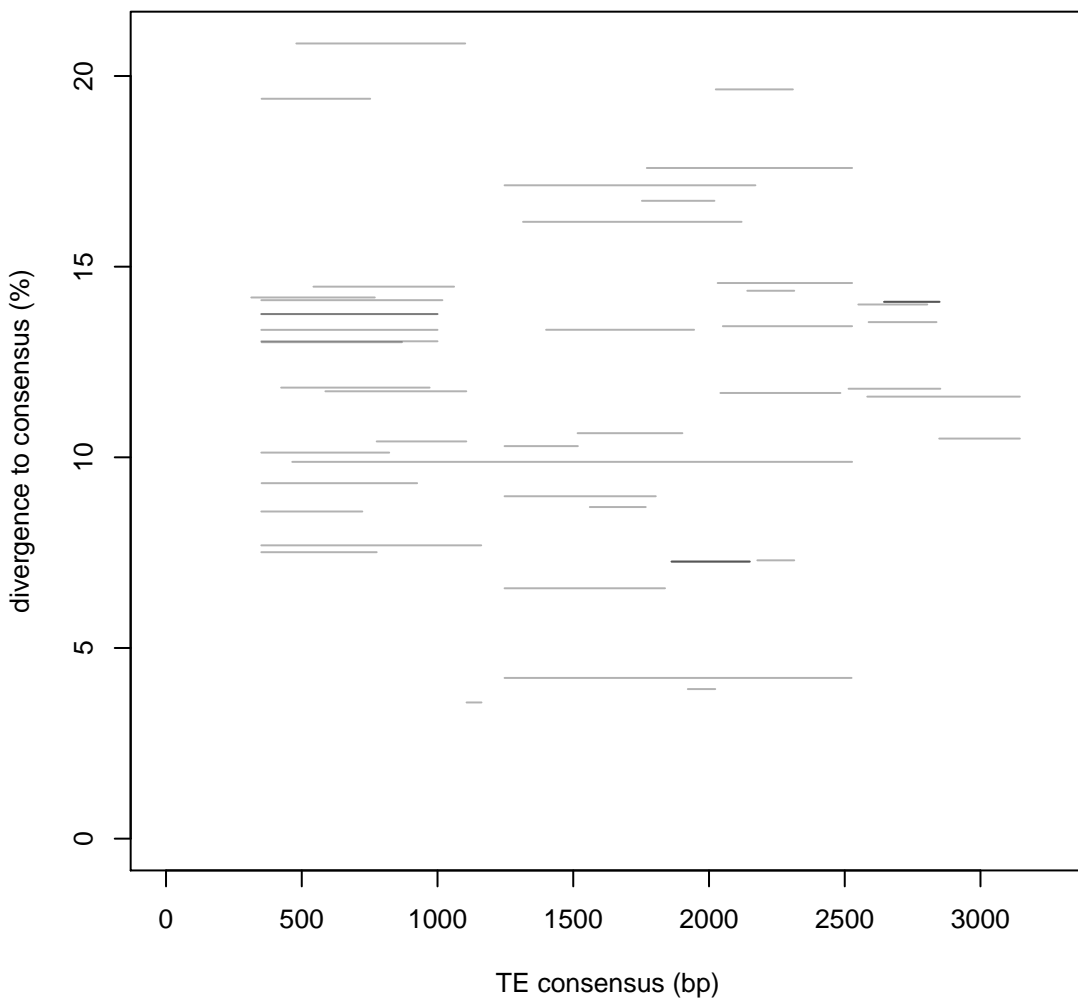
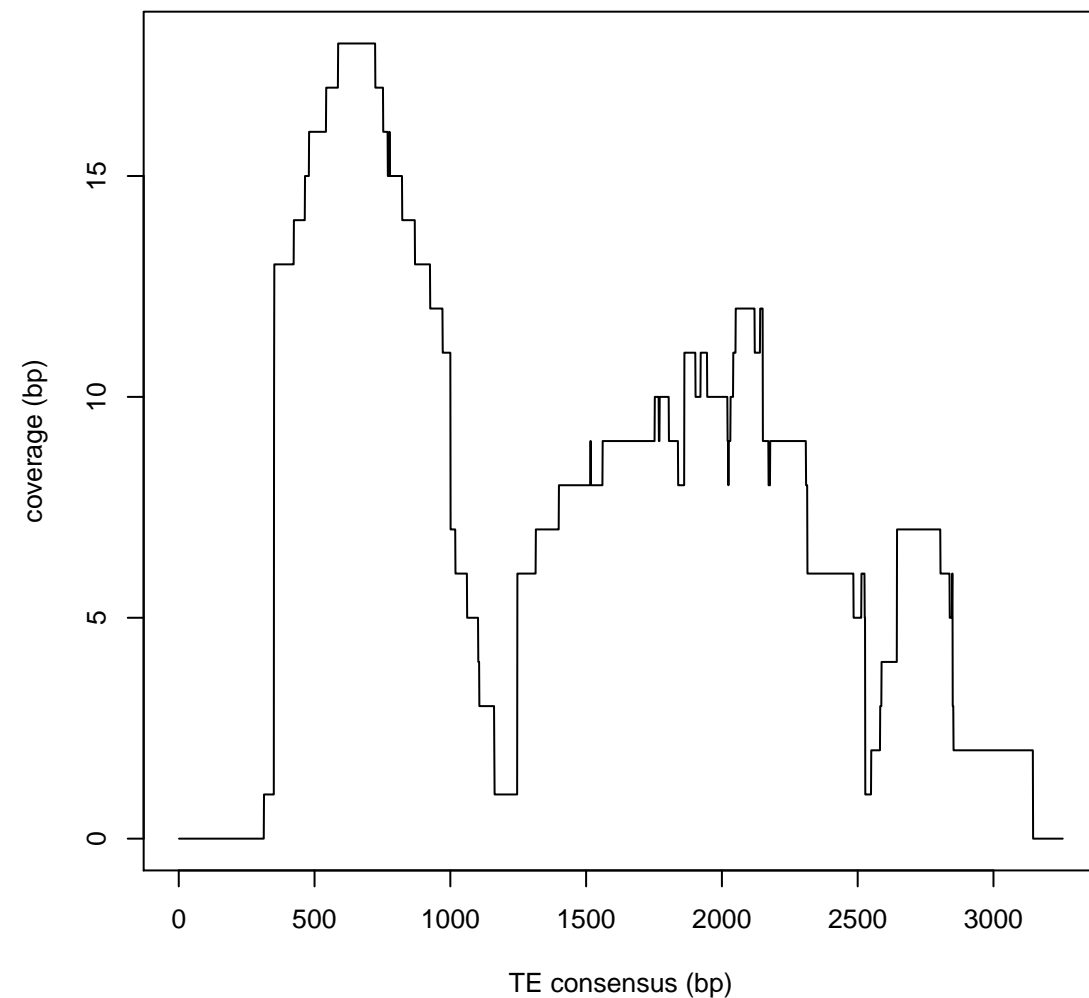


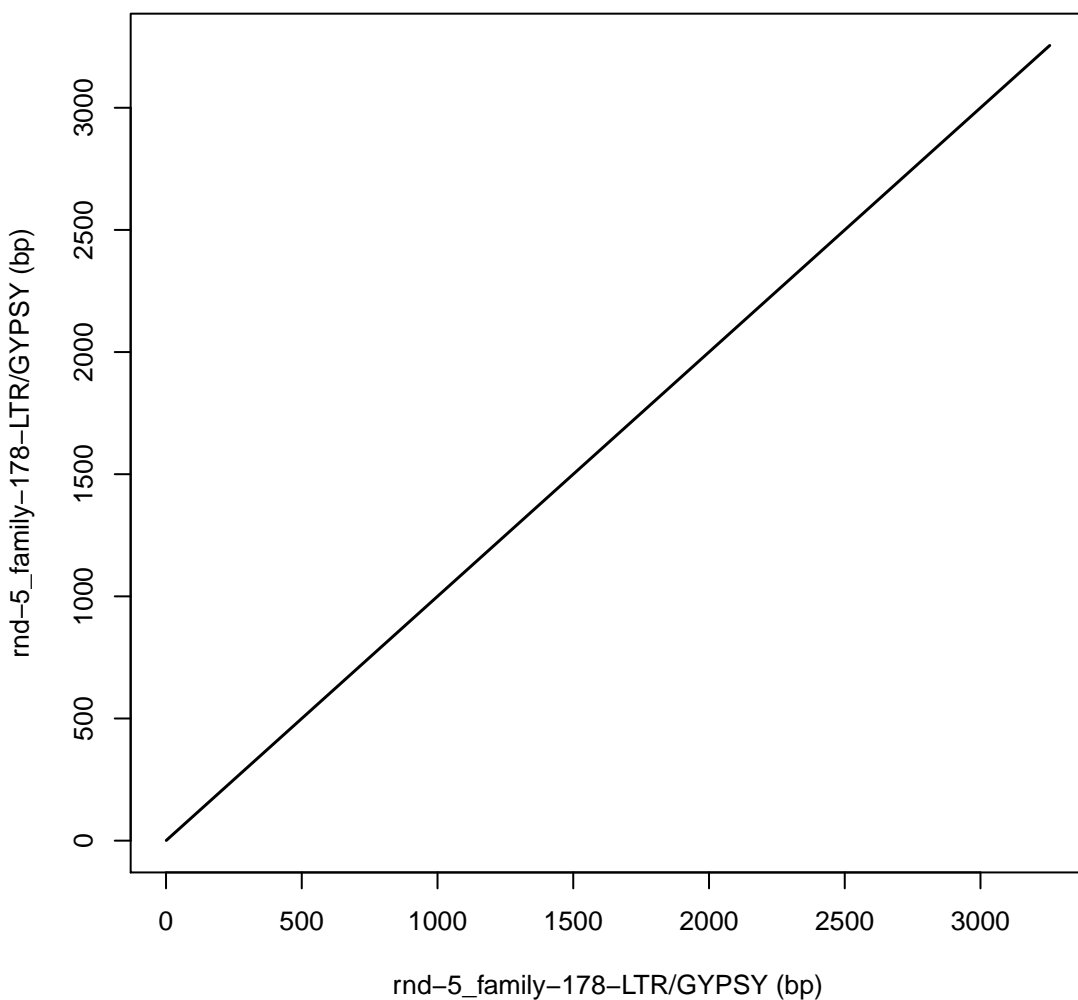
TE: rnd-5\_family-178-LTR/GYPY  
consensus size: 3255bp; fragments: 49; full length: 0 (>=2929.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

