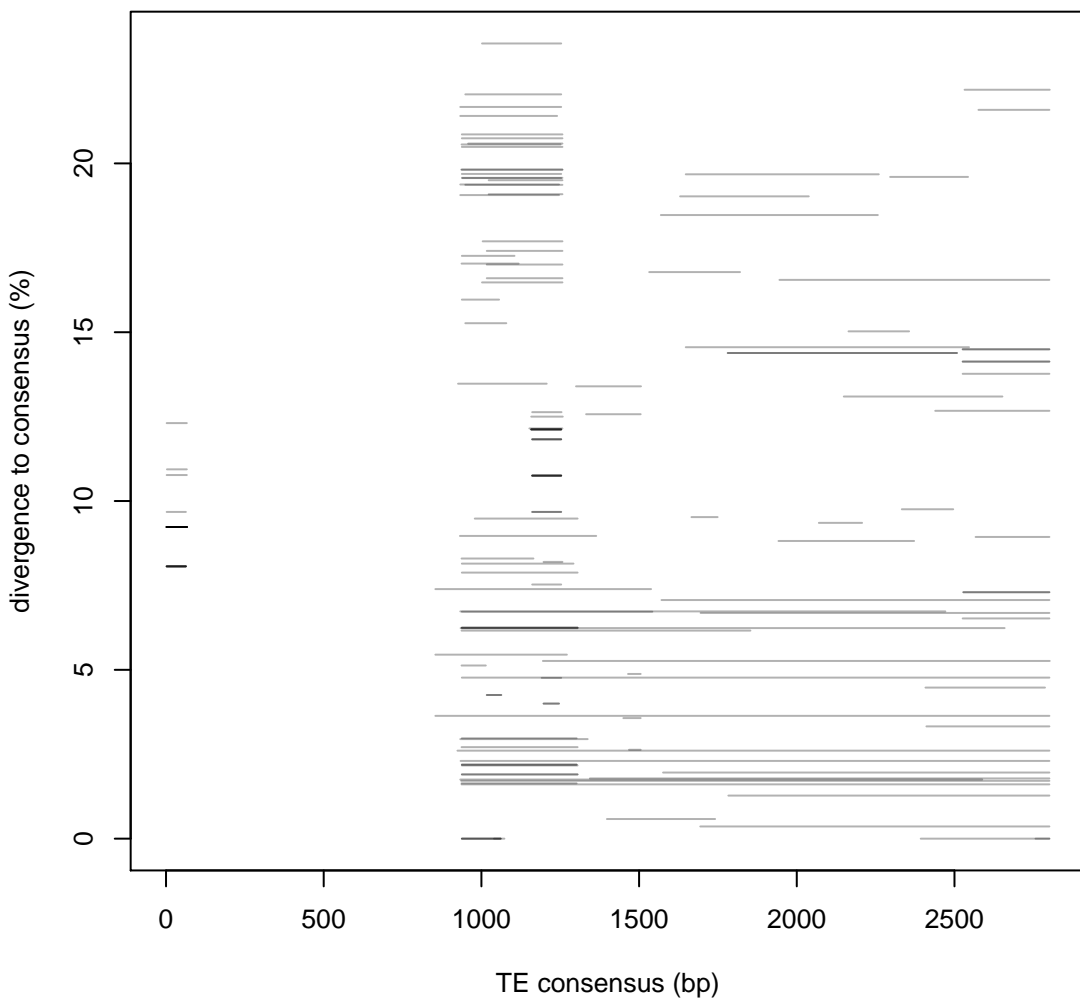
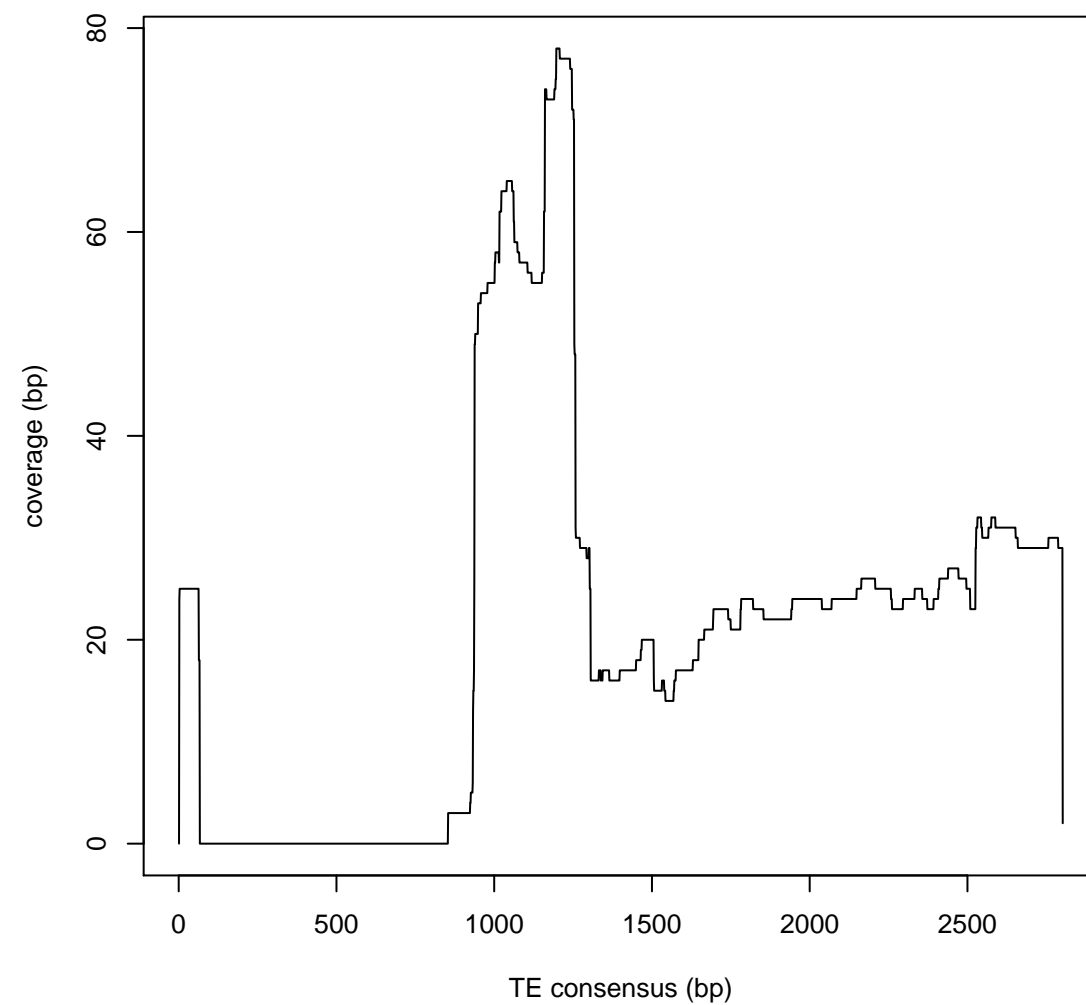


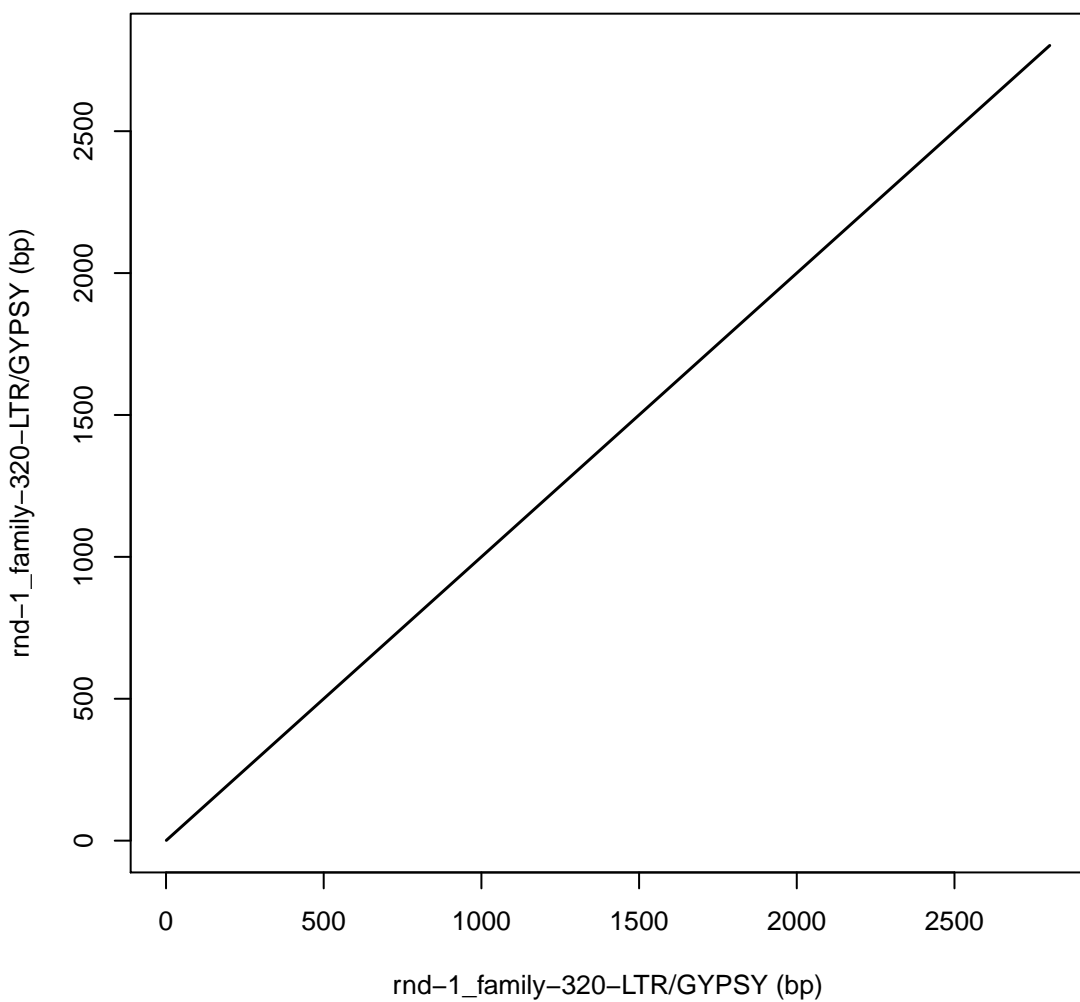
TE: rnd-1\_family-320-LTR/GYPSY  
consensus size: 2802bp; fragments: 158; full length: 0 (>=2521.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

