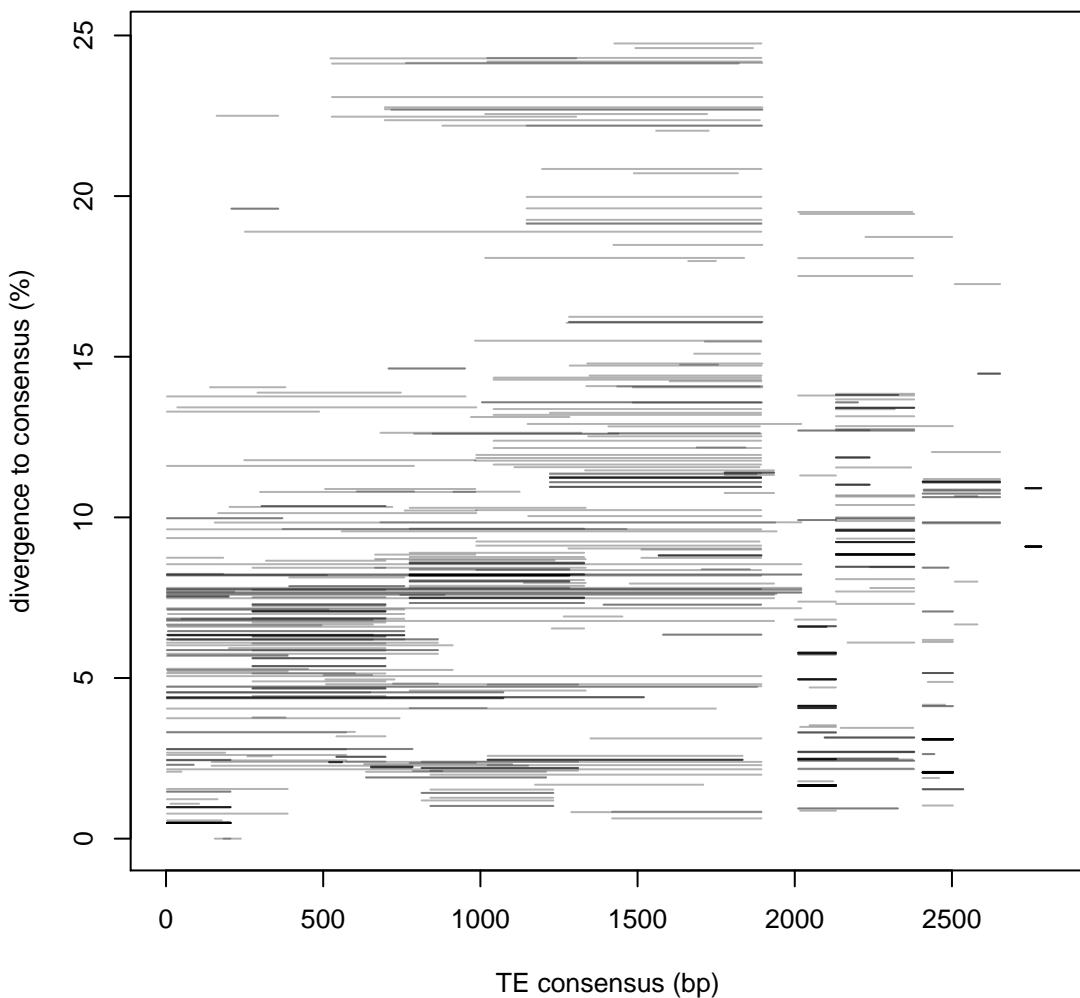
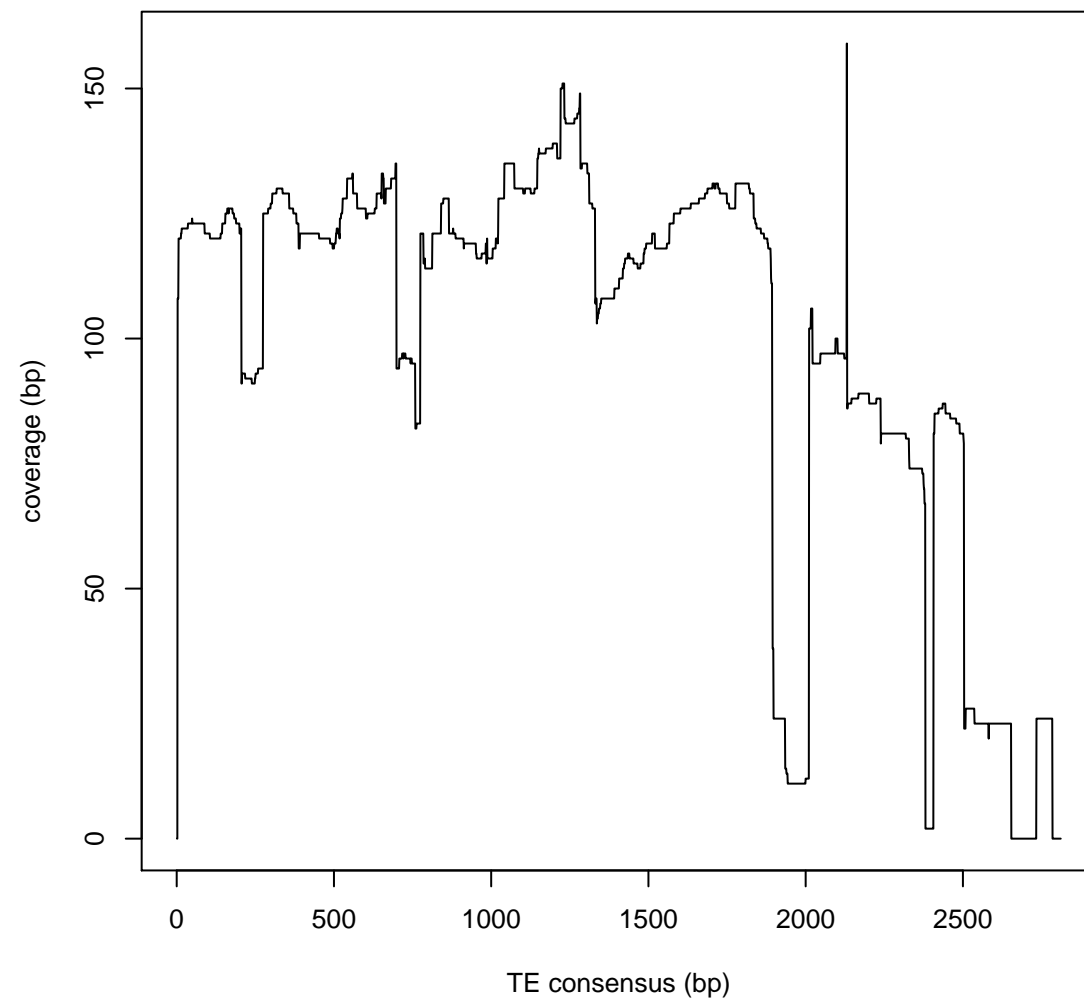


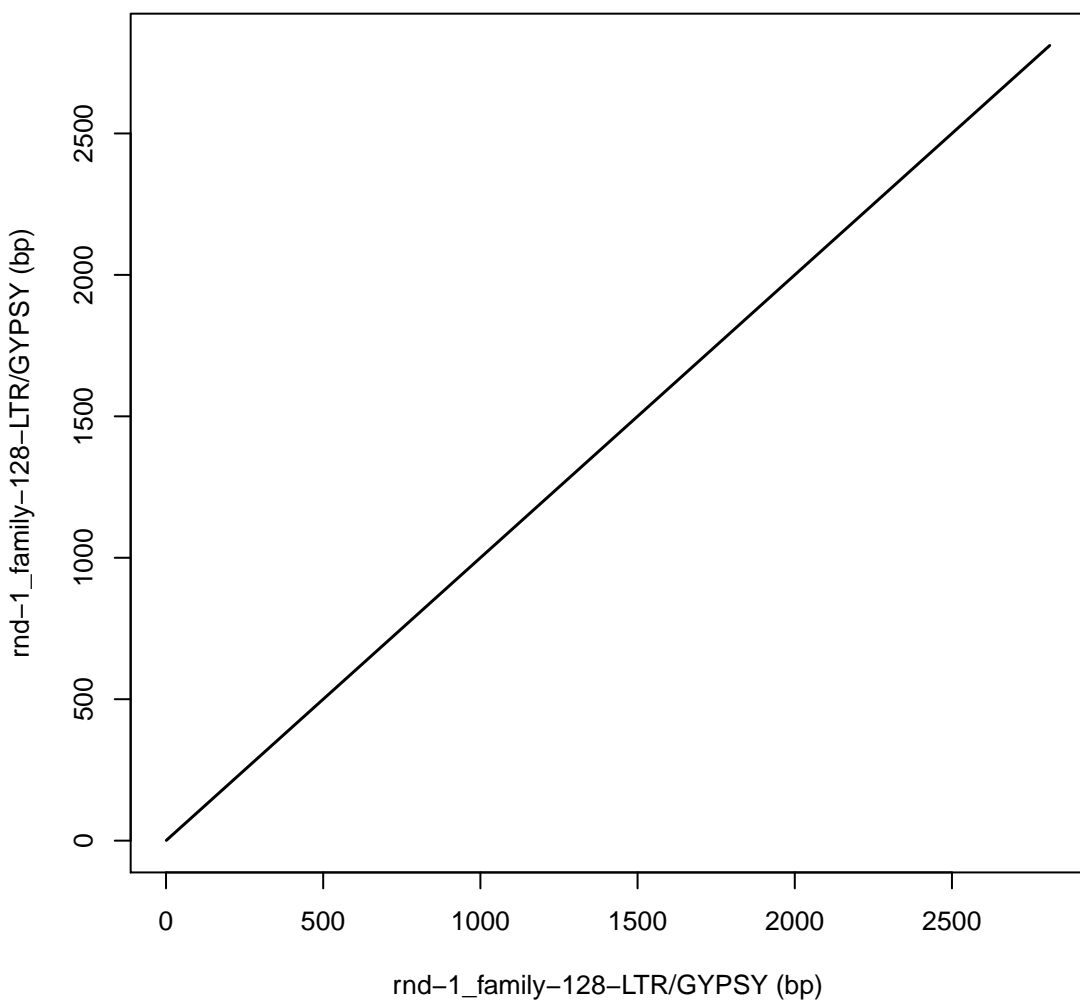
TE: rnd-1_family-128-LTR/GYPSY
consensus size: 2811bp; fragments: 668; full length: 0 (>=2529.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

