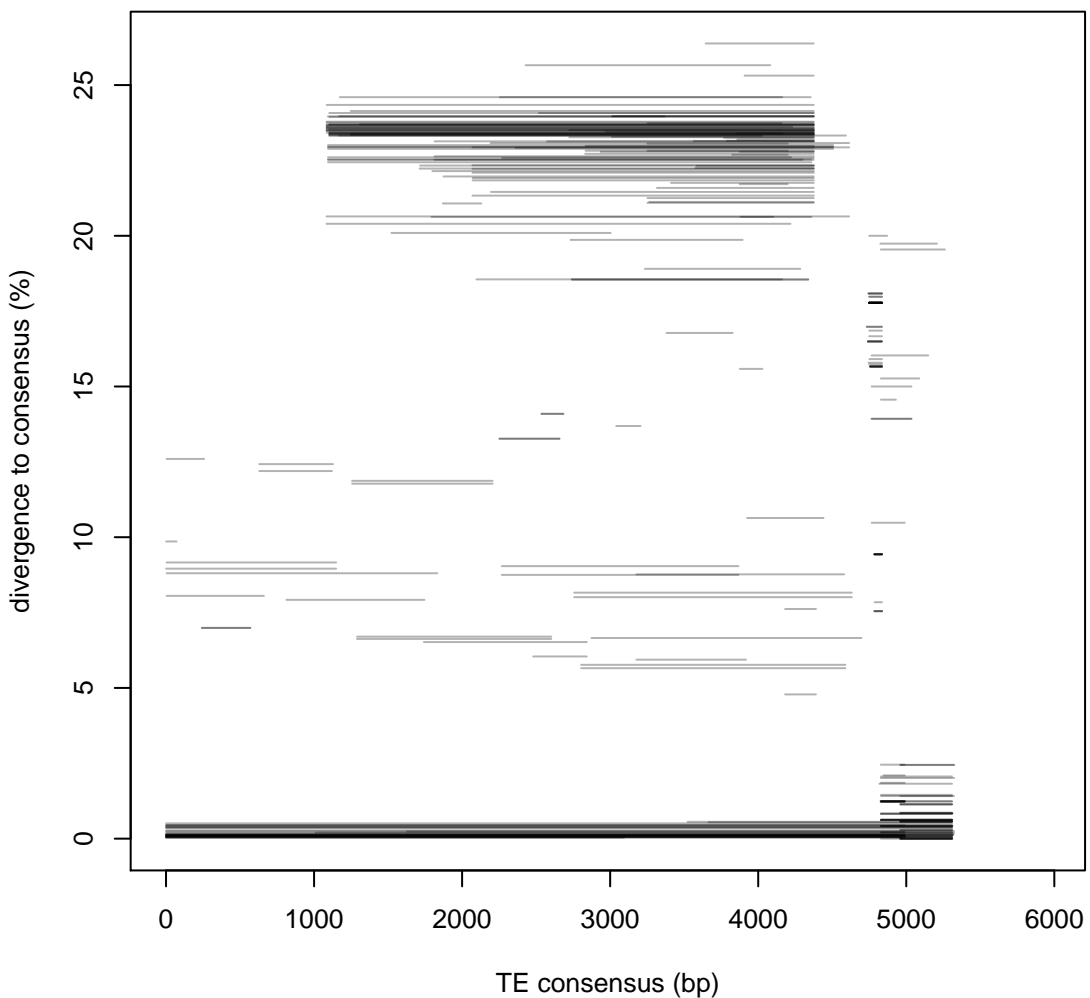
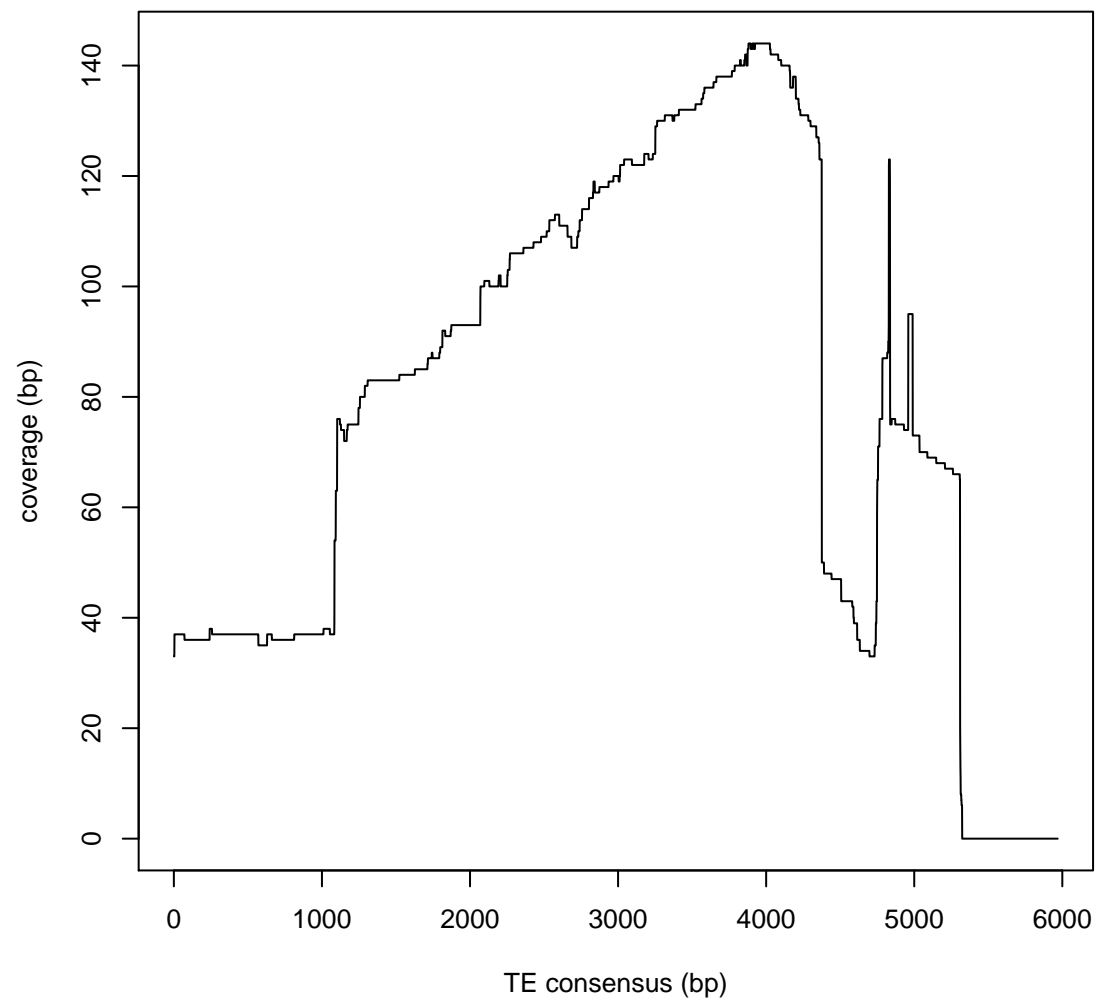


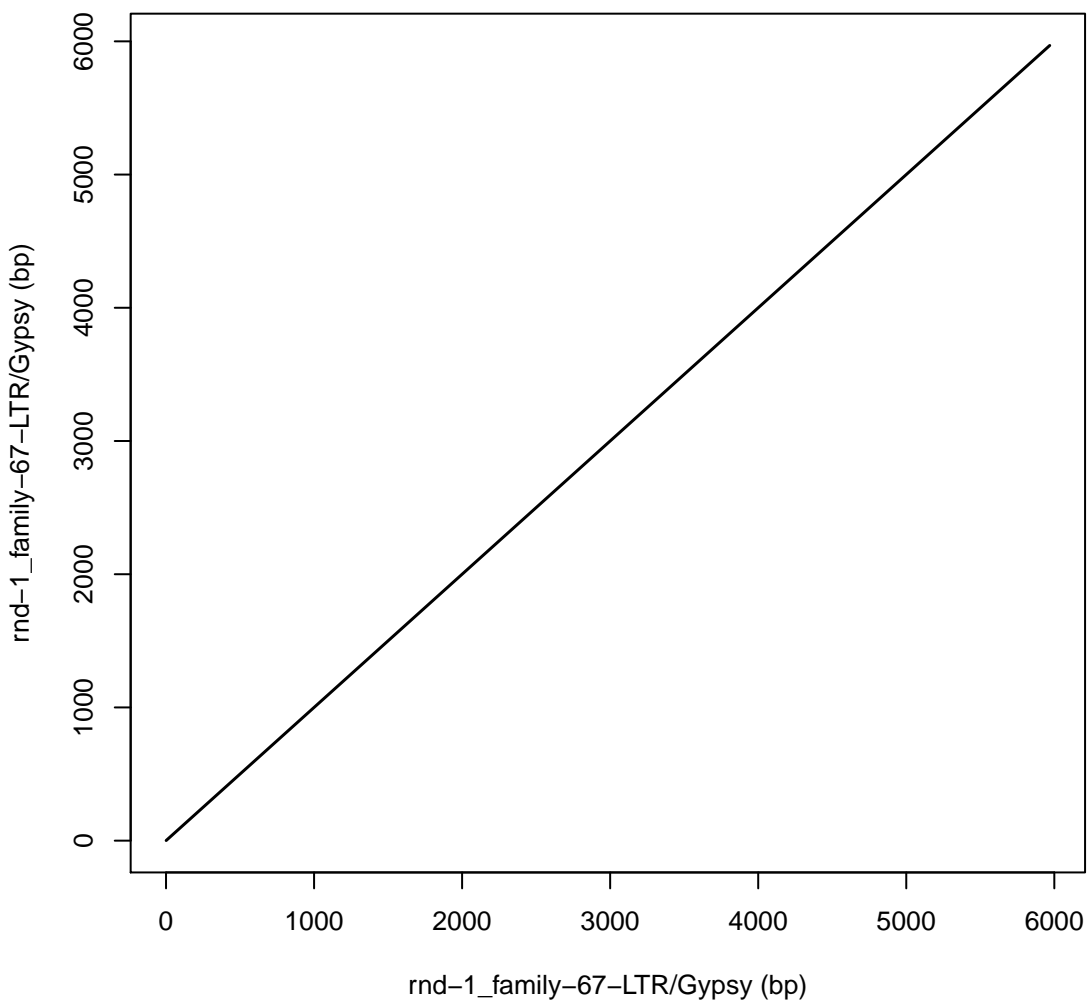
TE: rnd-1_family-67-LTR/Gypsy
consensus size: 5969bp; fragments: 290; full length: 0 (>=5372.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

