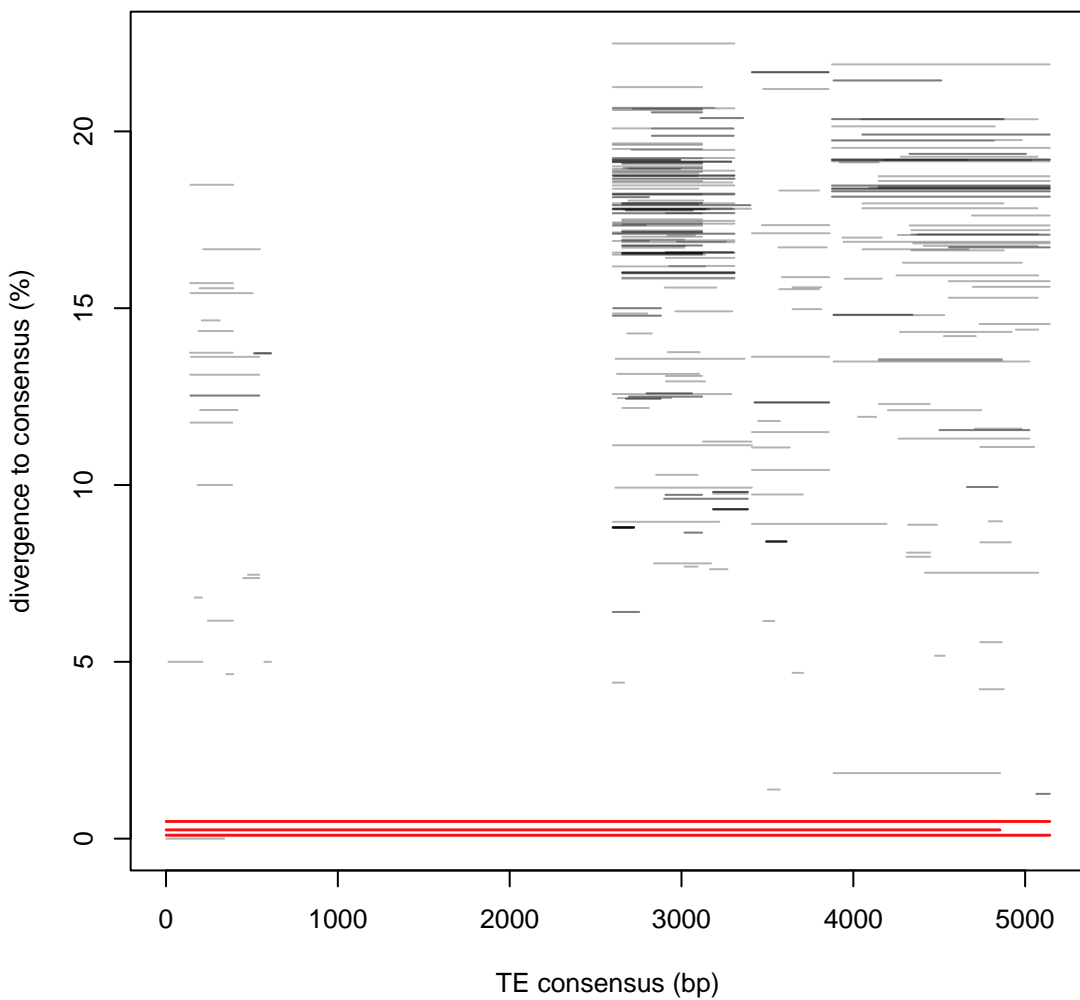
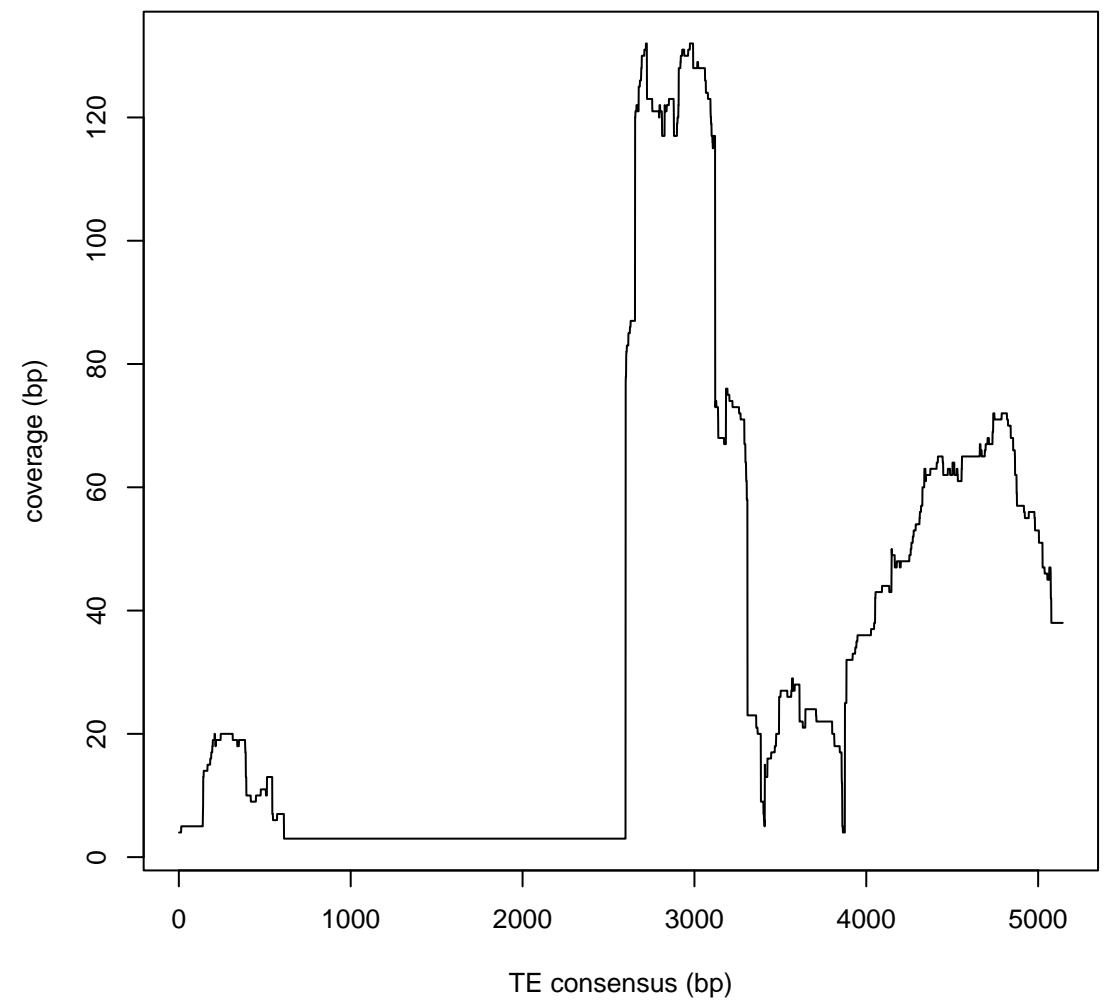


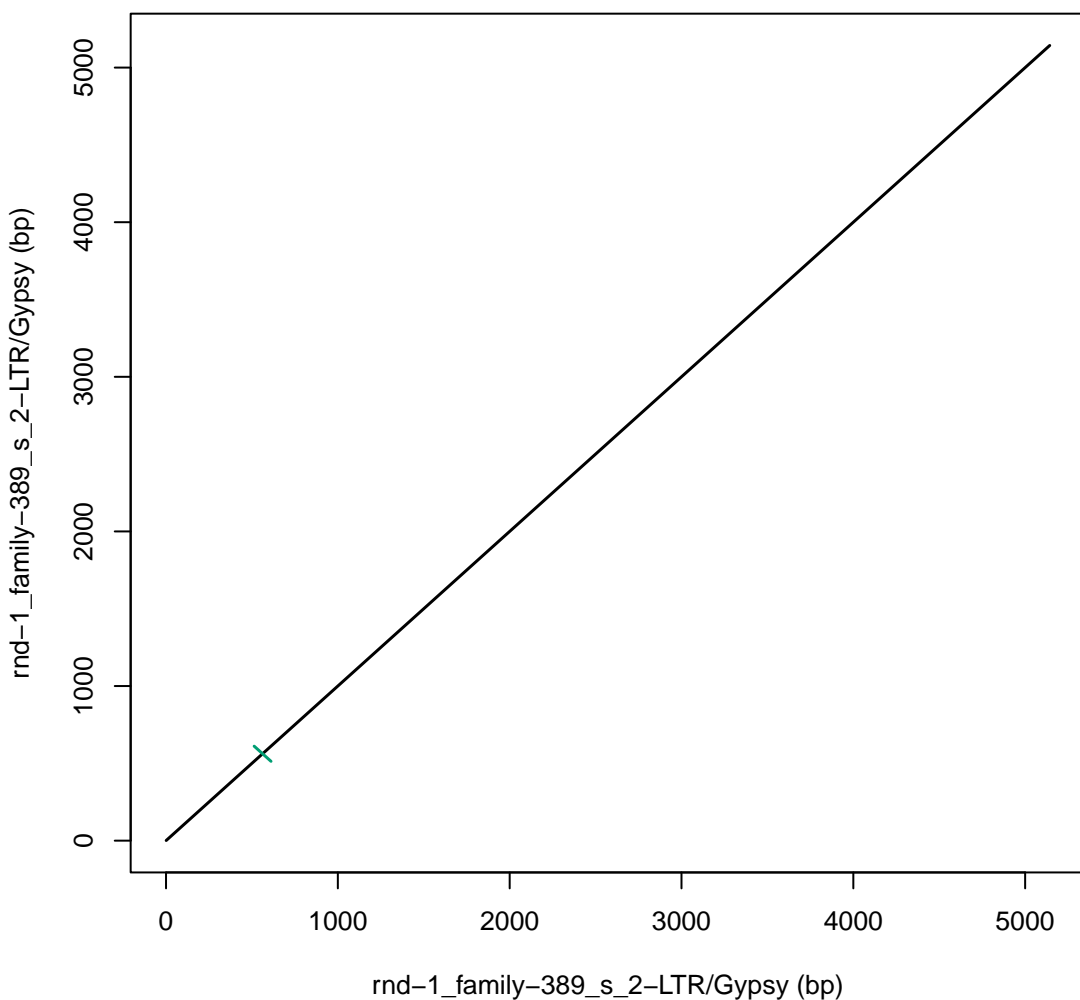
TE: rnd-1_family-389_s_2-LTR/Gypsy
consensus size: 5143bp; fragments: 321; full length: 3 (>=4628.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

