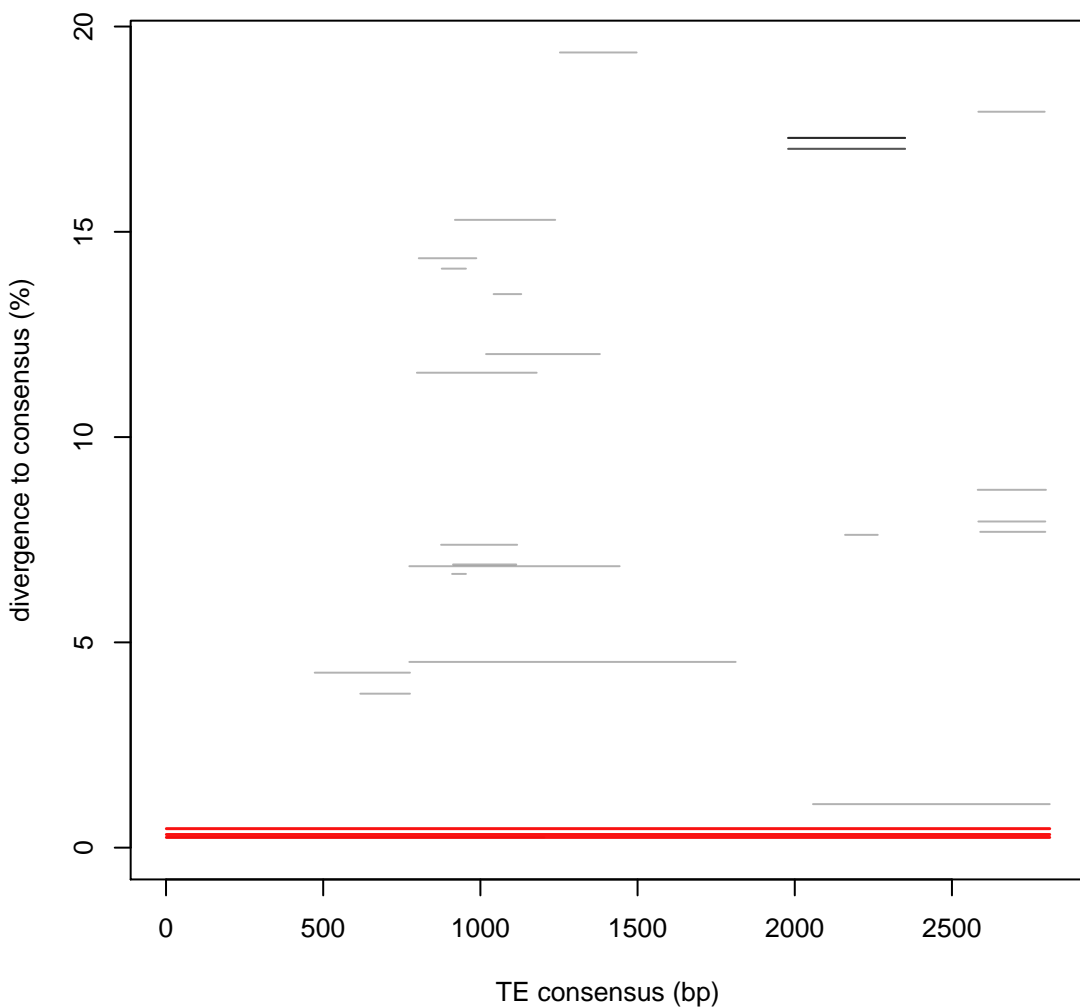
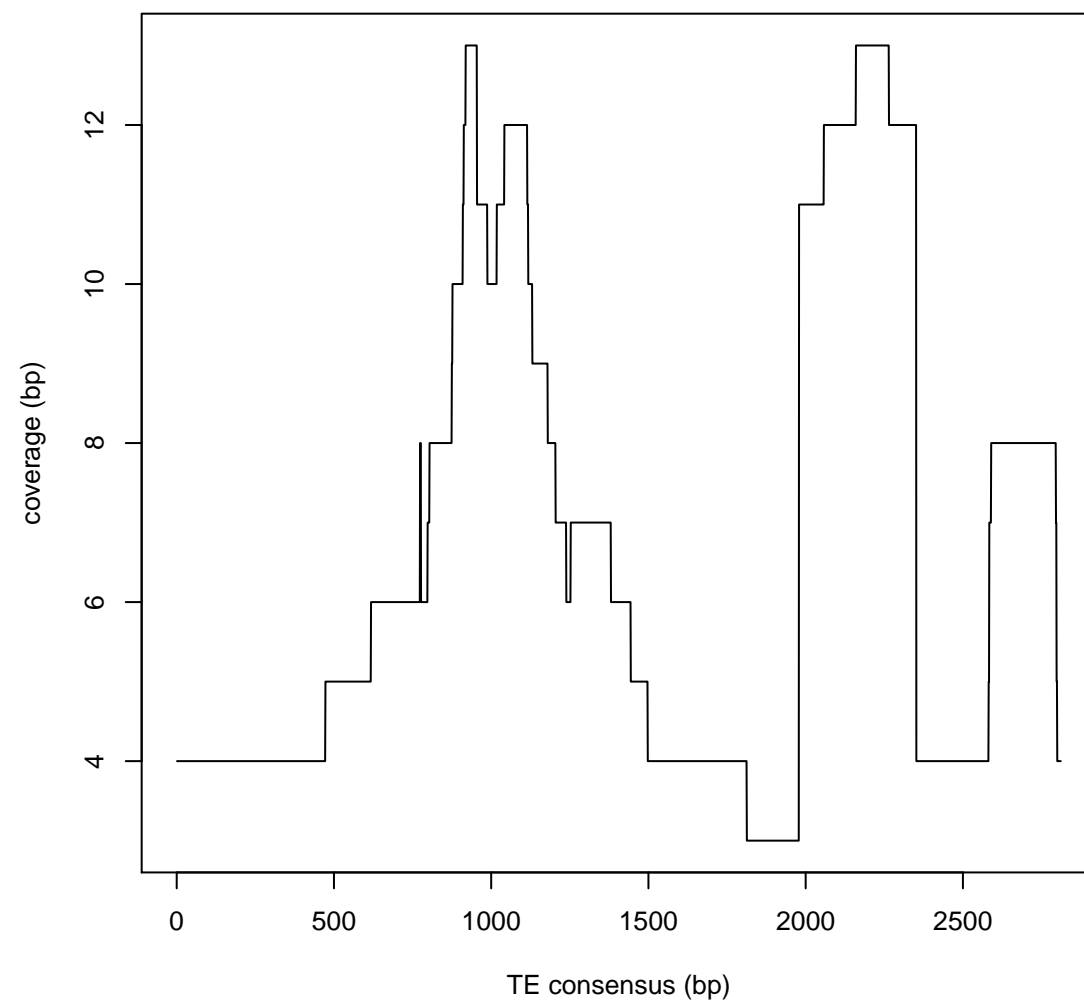


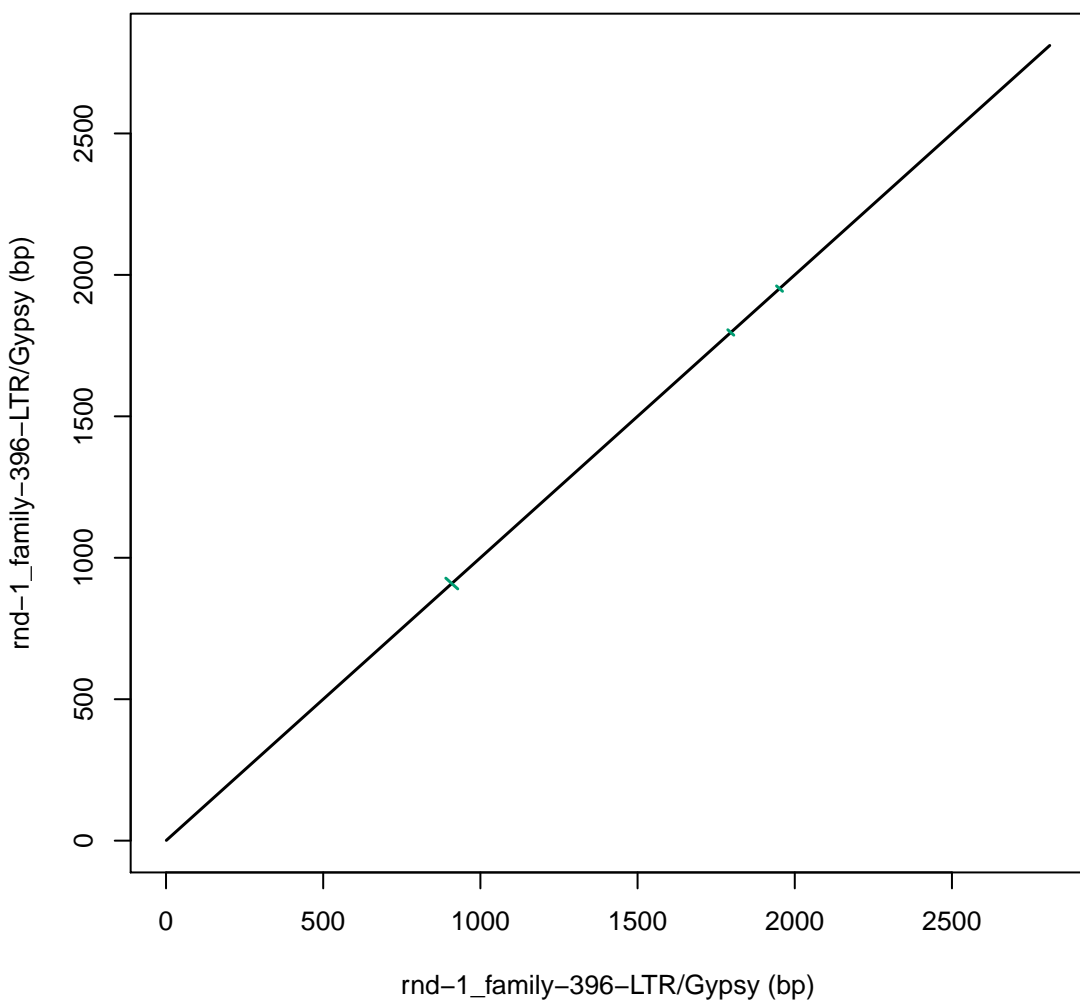
TE: rnd-1_family-396-LTR/Gypsy
consensus size: 2811bp; fragments: 32; full length: 3 (>=2529.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

