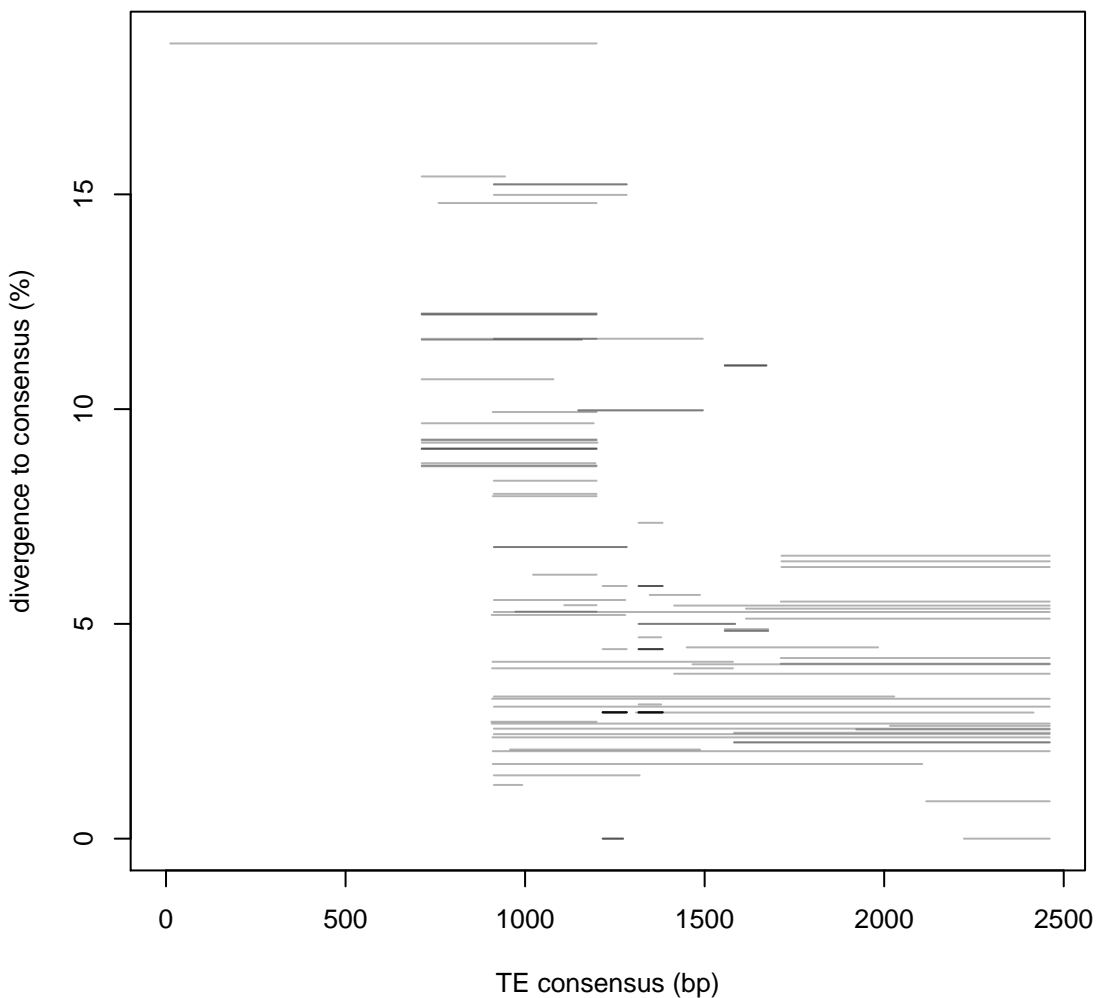
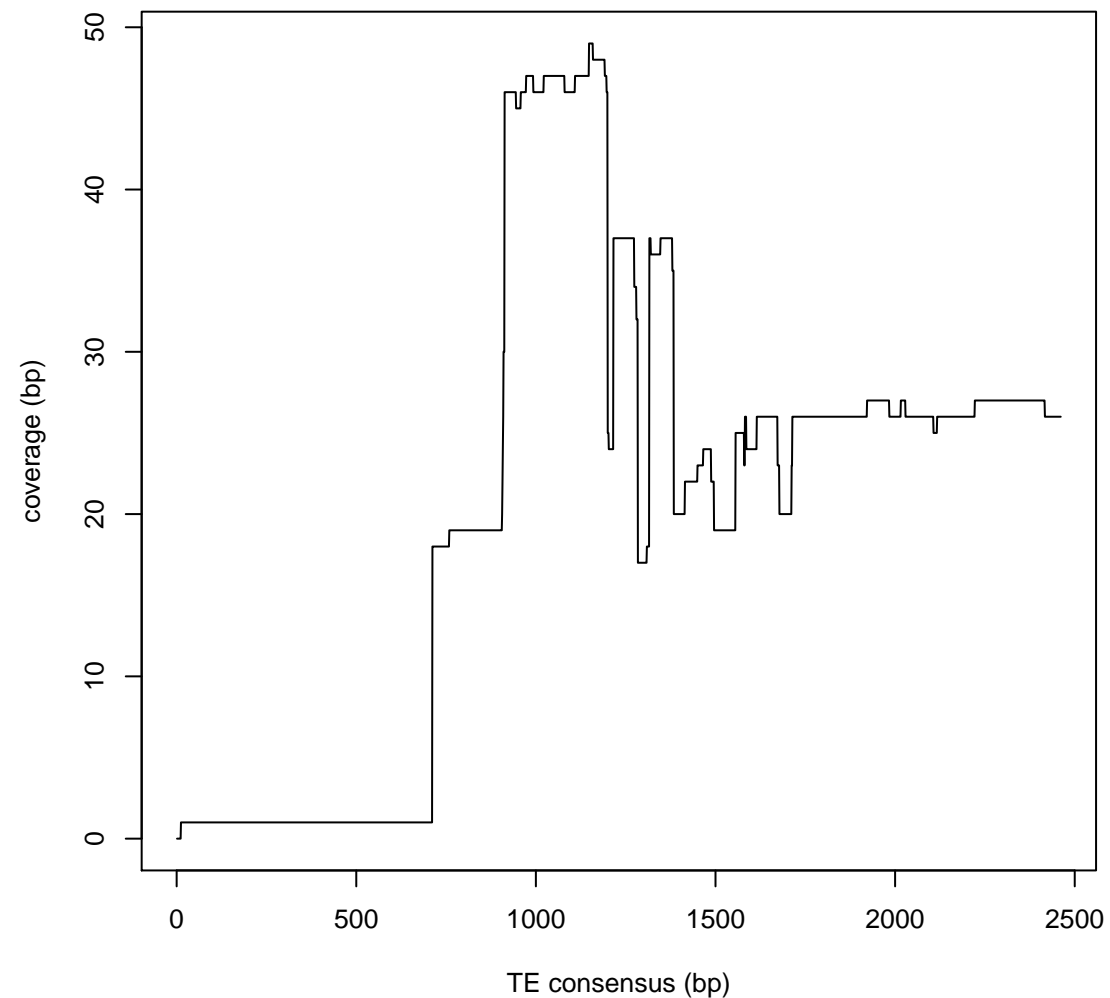


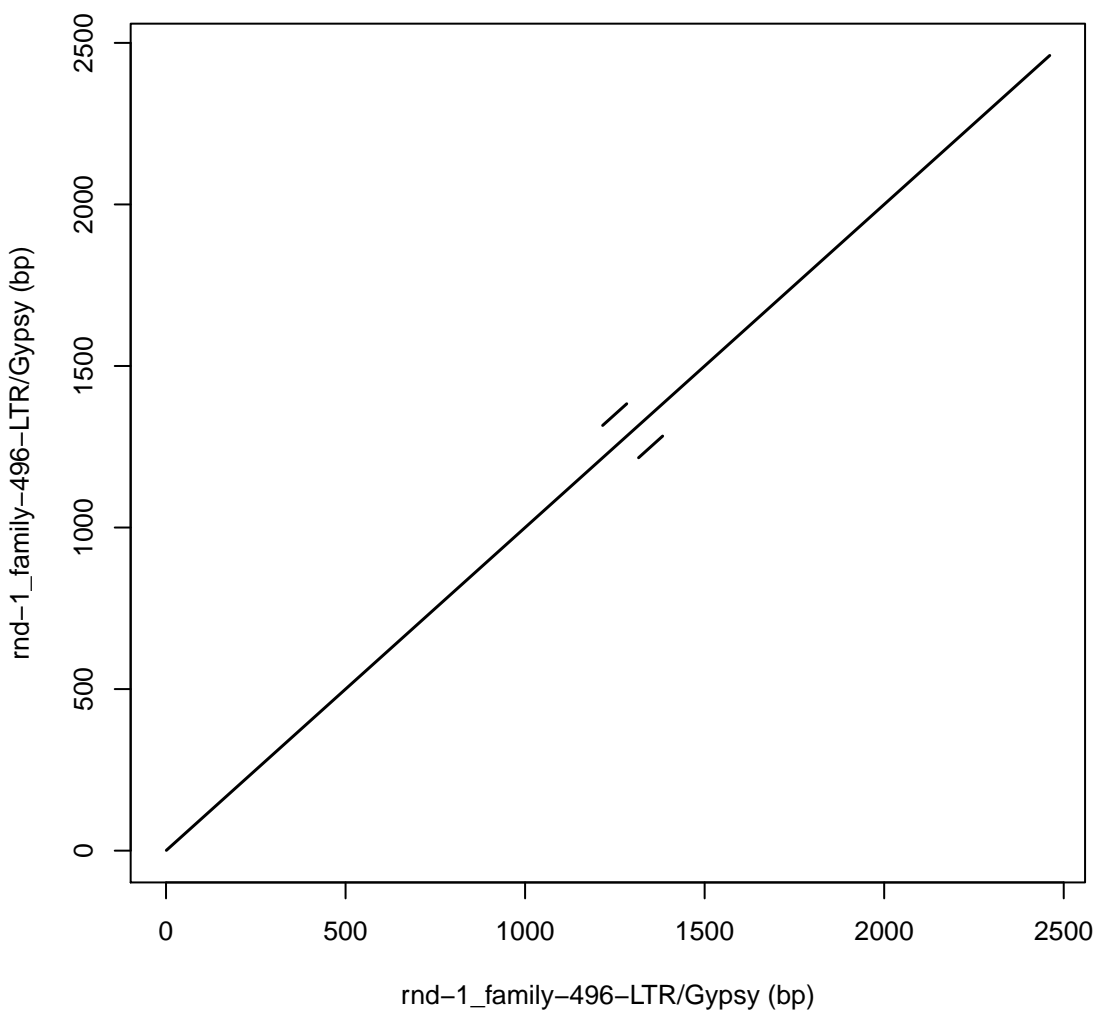
TE: rnd-1_family-496-LTR/Gypsy
consensus size: 2461bp; fragments: 111; full length: 0 (>=2214.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

