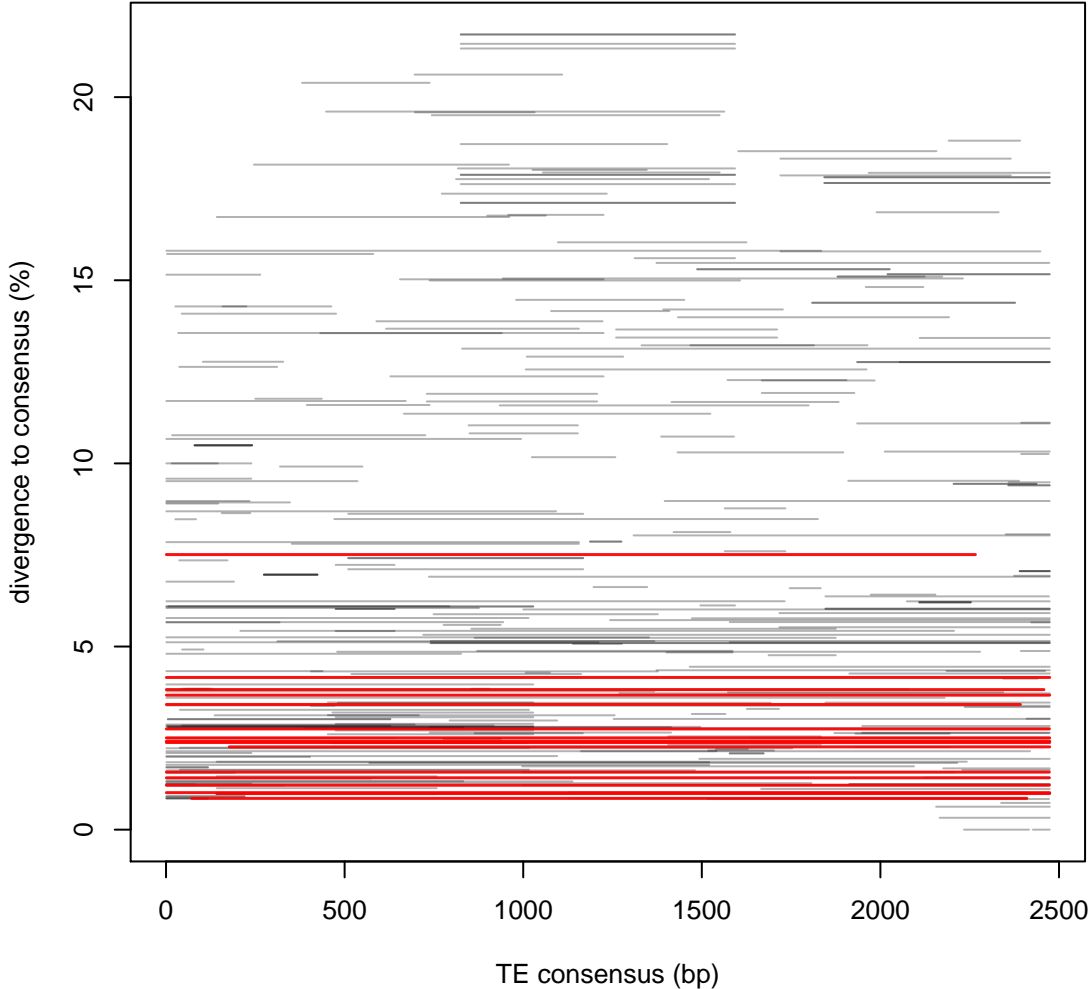
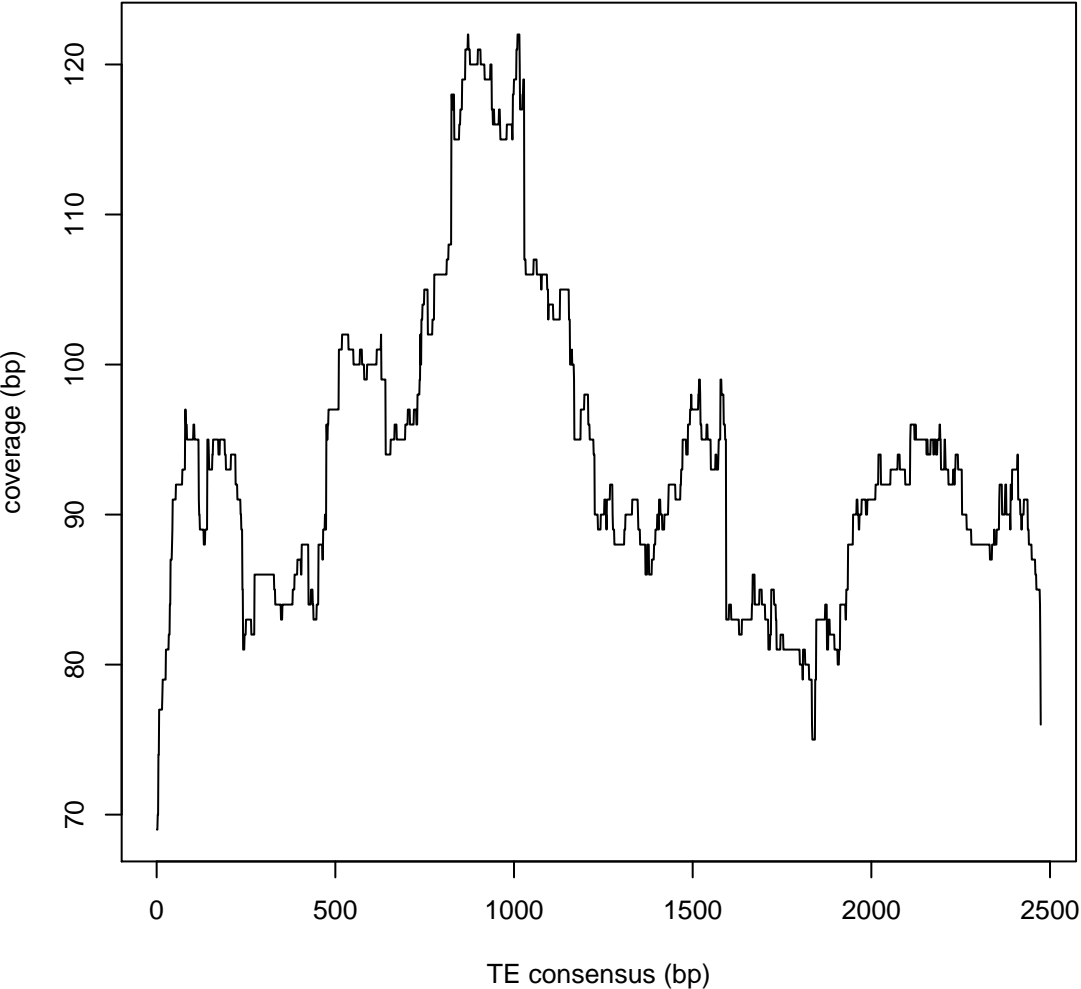


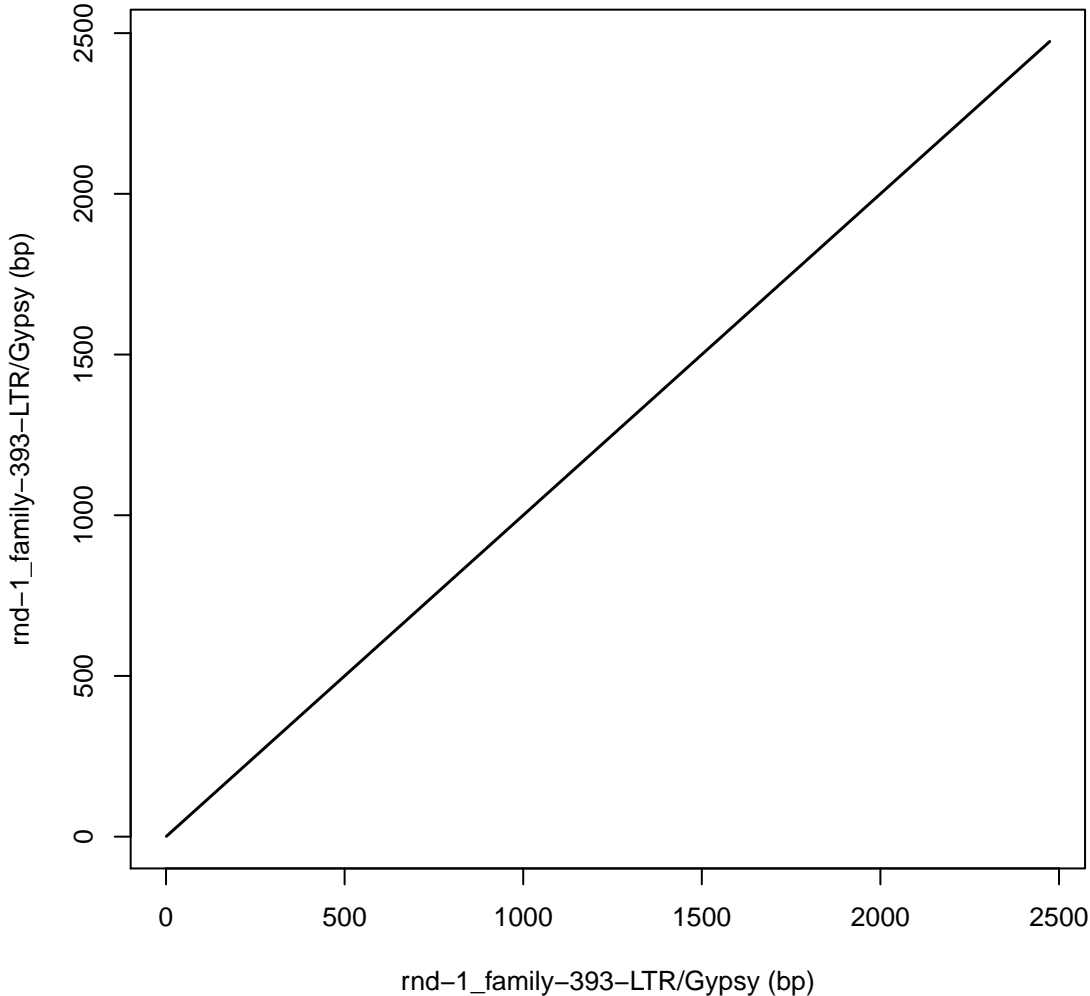
TE: rnd-1_family-393-LTR/Gypsy
consensus size: 2474bp; fragments: 333; full length: 16 (>=2226.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

