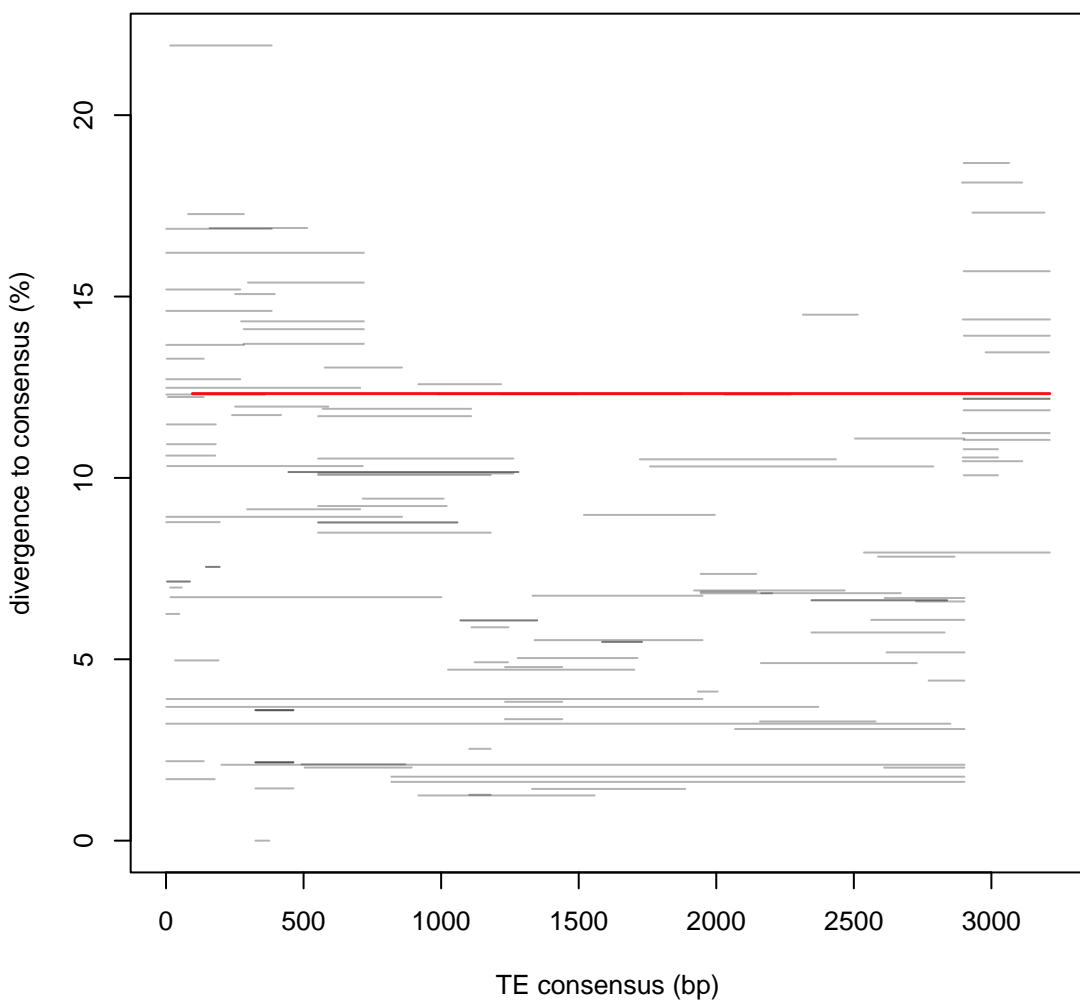
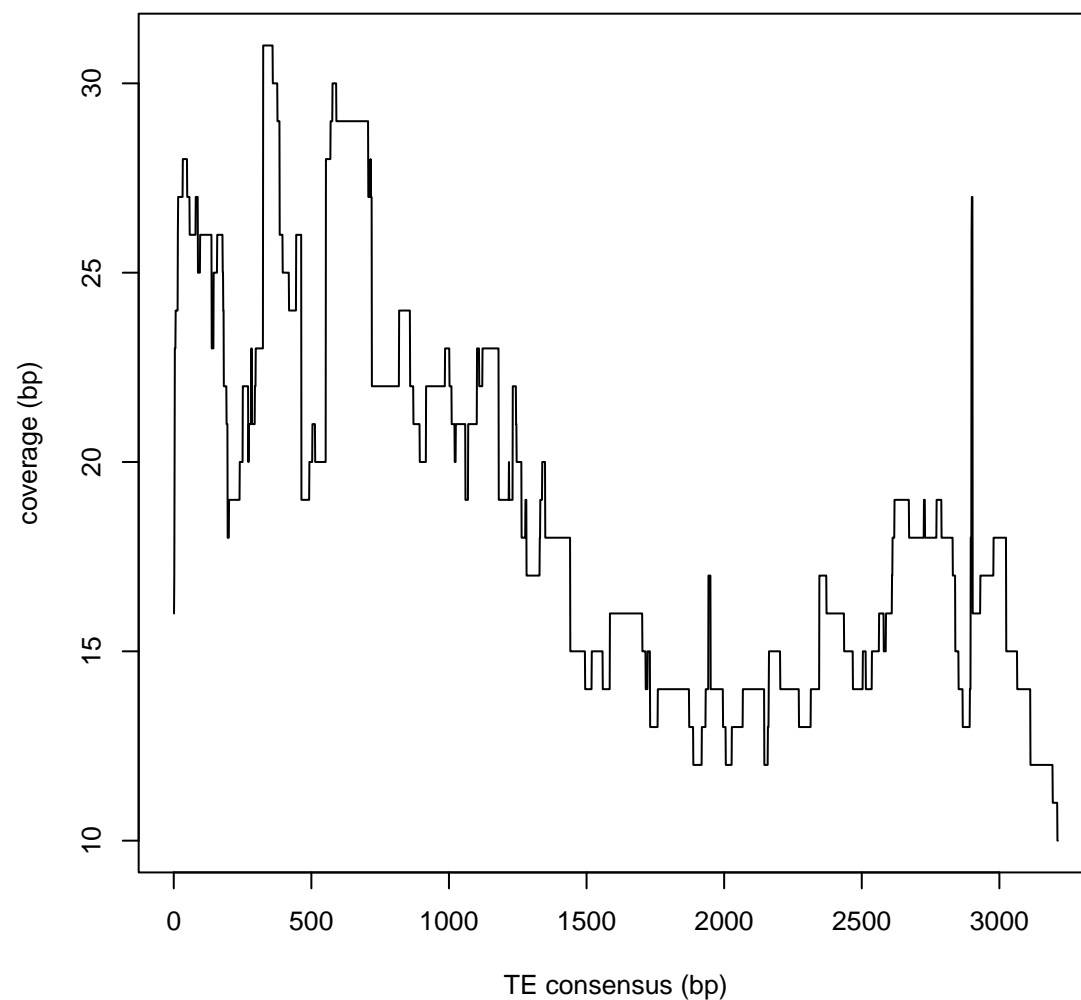


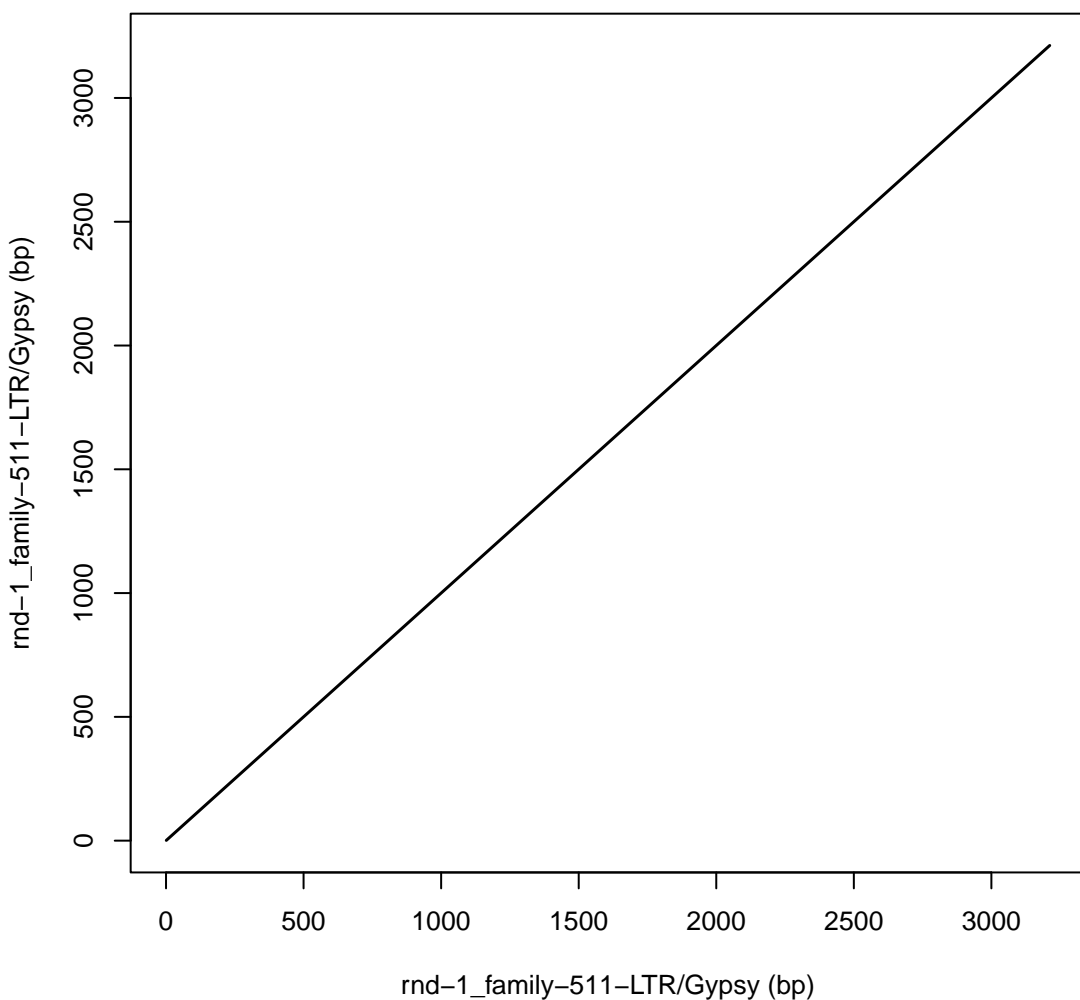
TE: rnd-1_family-511-LTR/Gypsy
consensus size: 3212bp; fragments: 129; full length: 1 (>=2890.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

