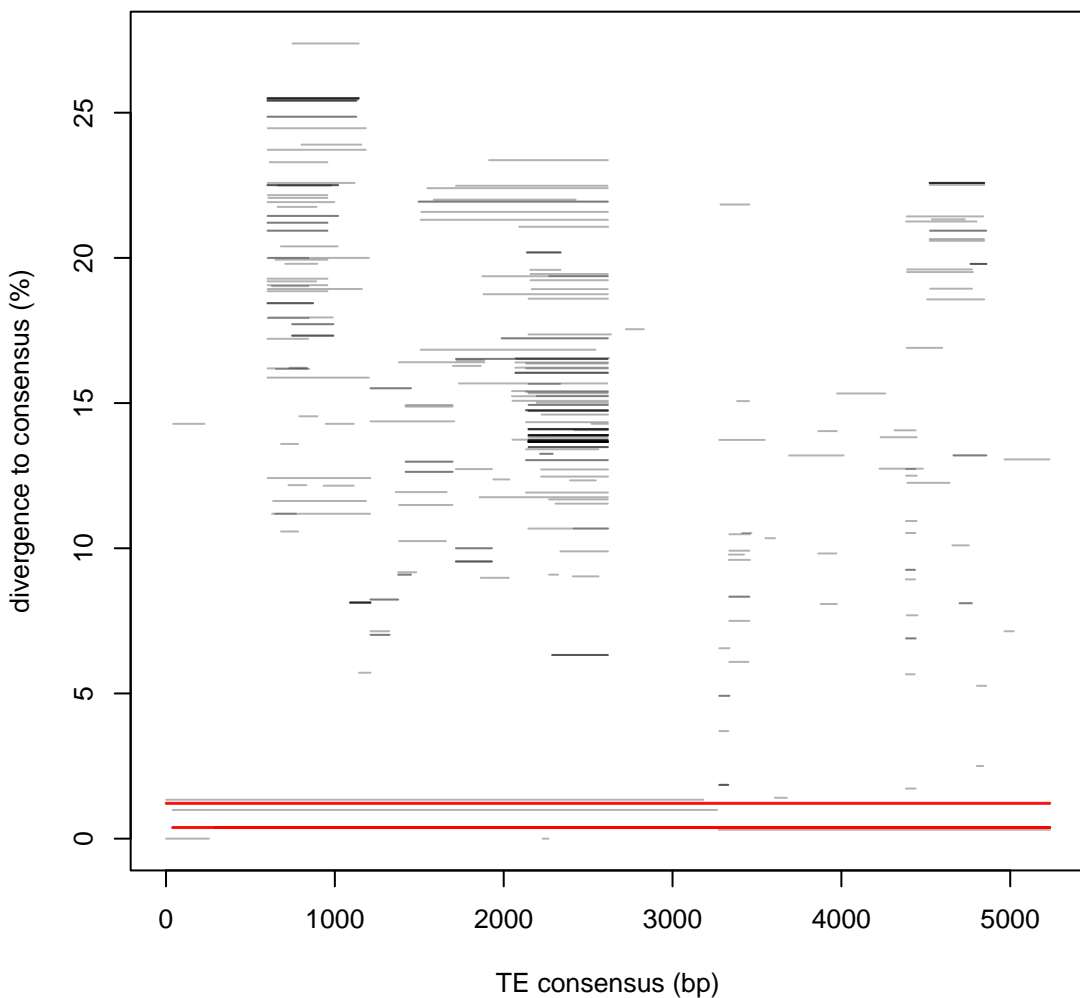
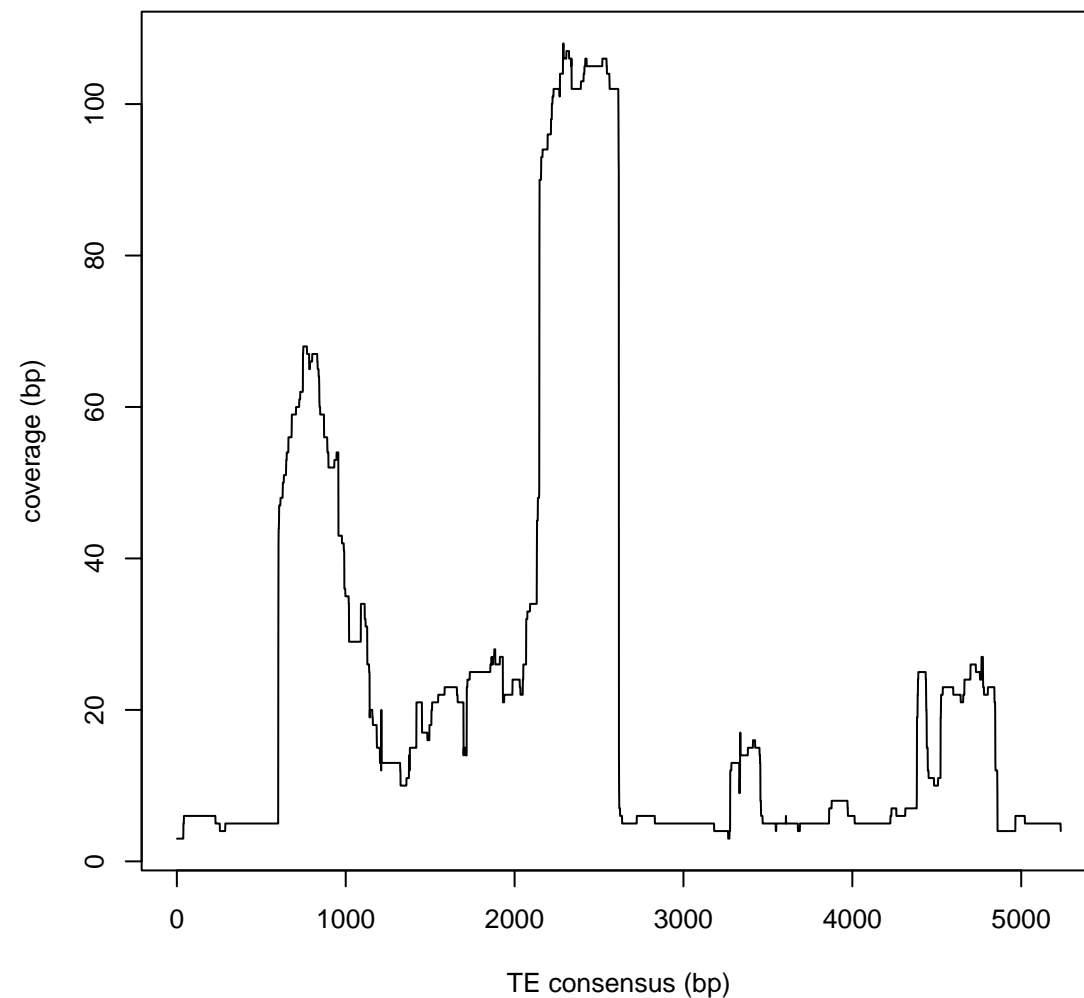


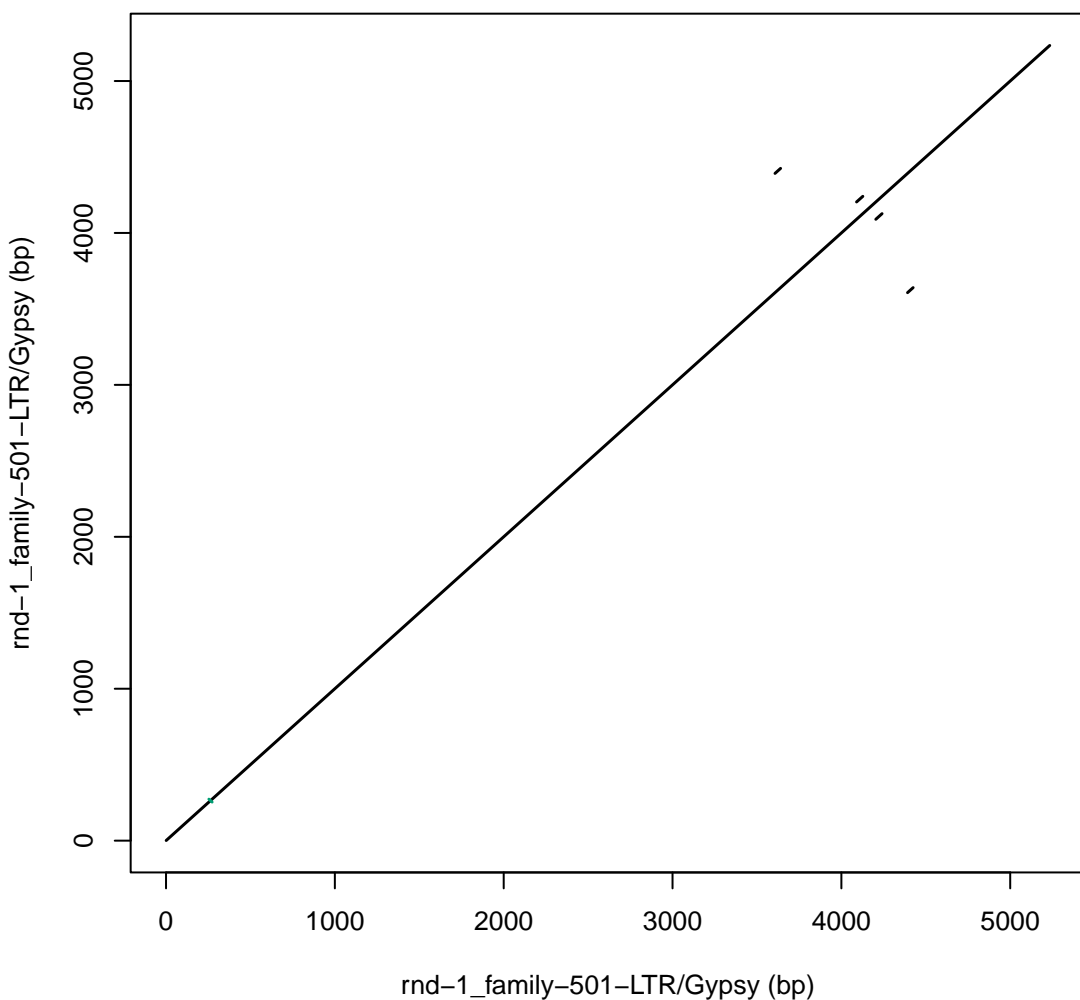
TE: rnd-1_family-501-LTR/Gypsy
consensus size: 5234bp; fragments: 296; full length: 3 (≥ 4710.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

