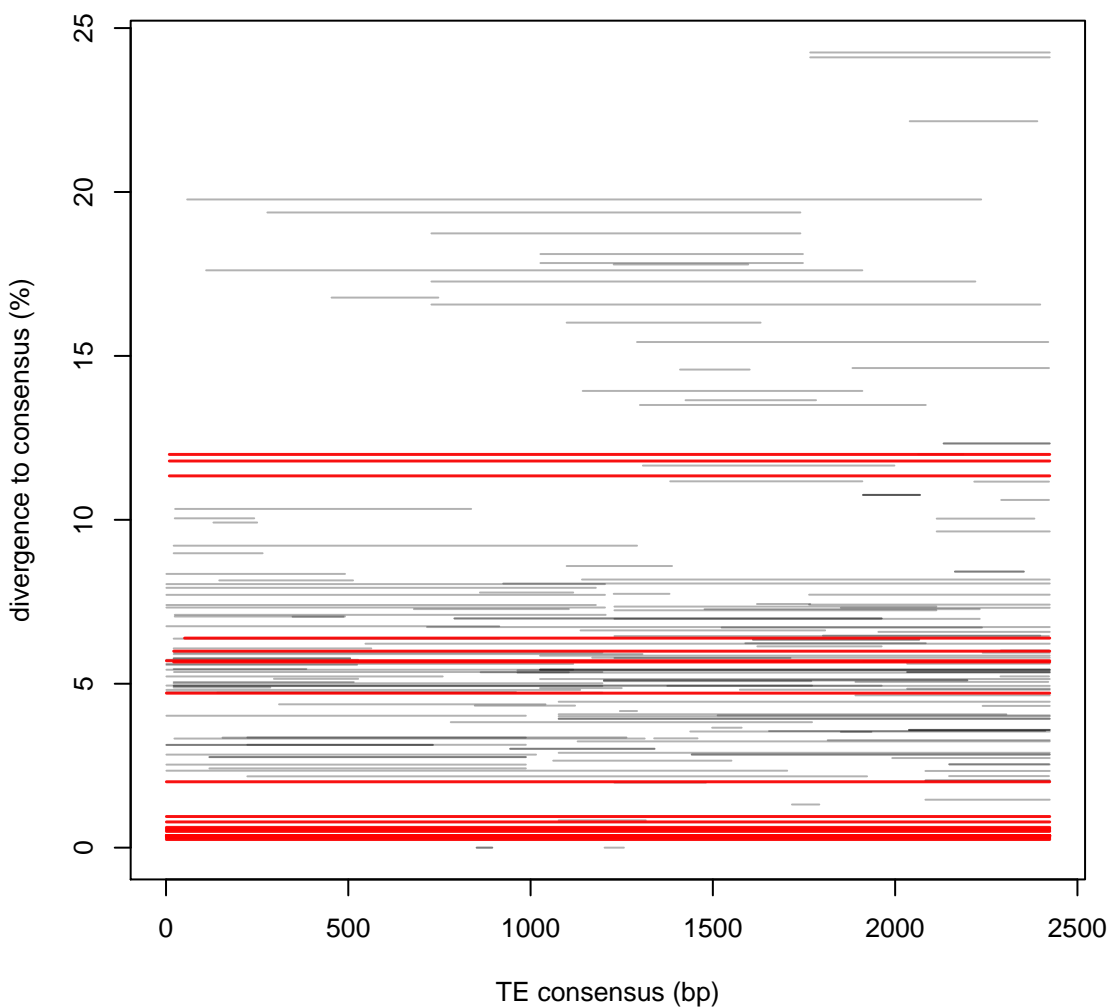
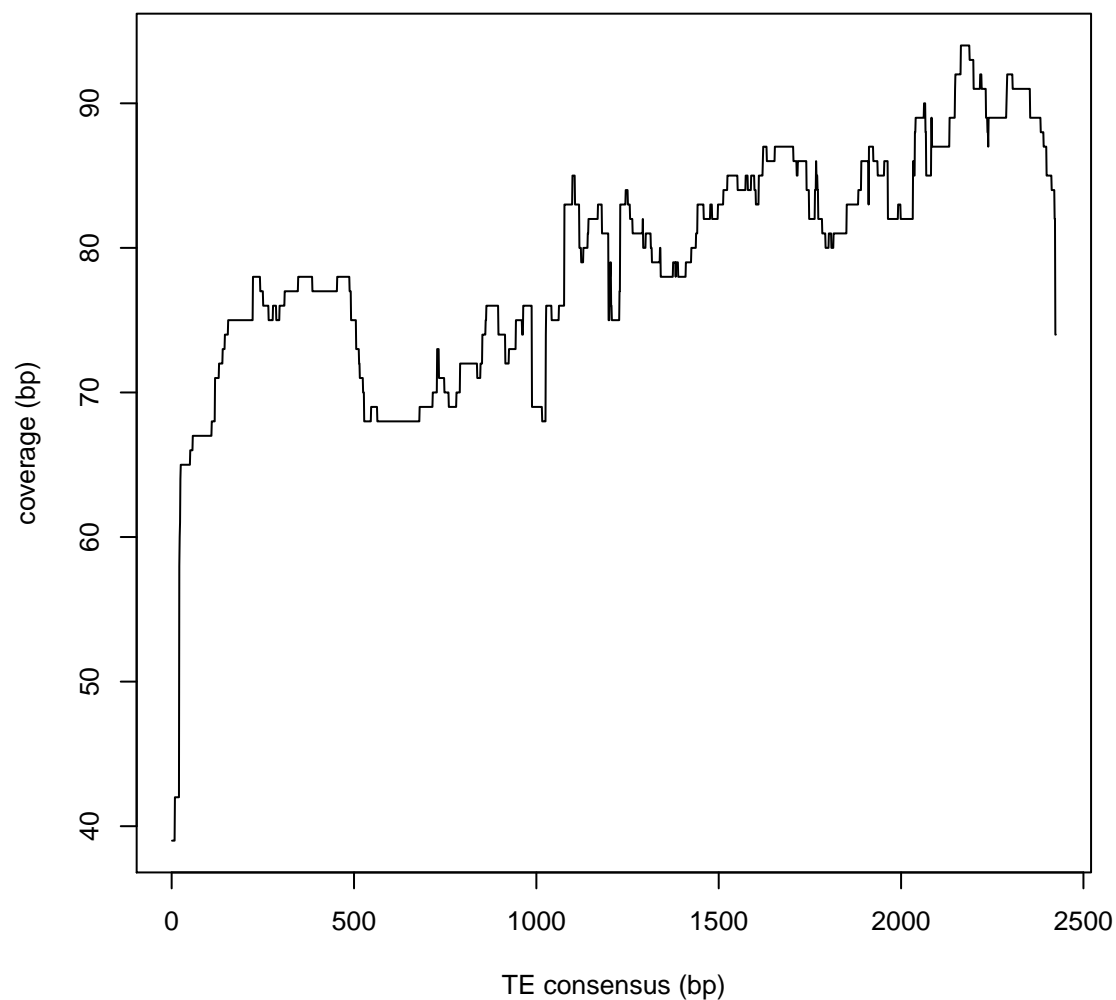


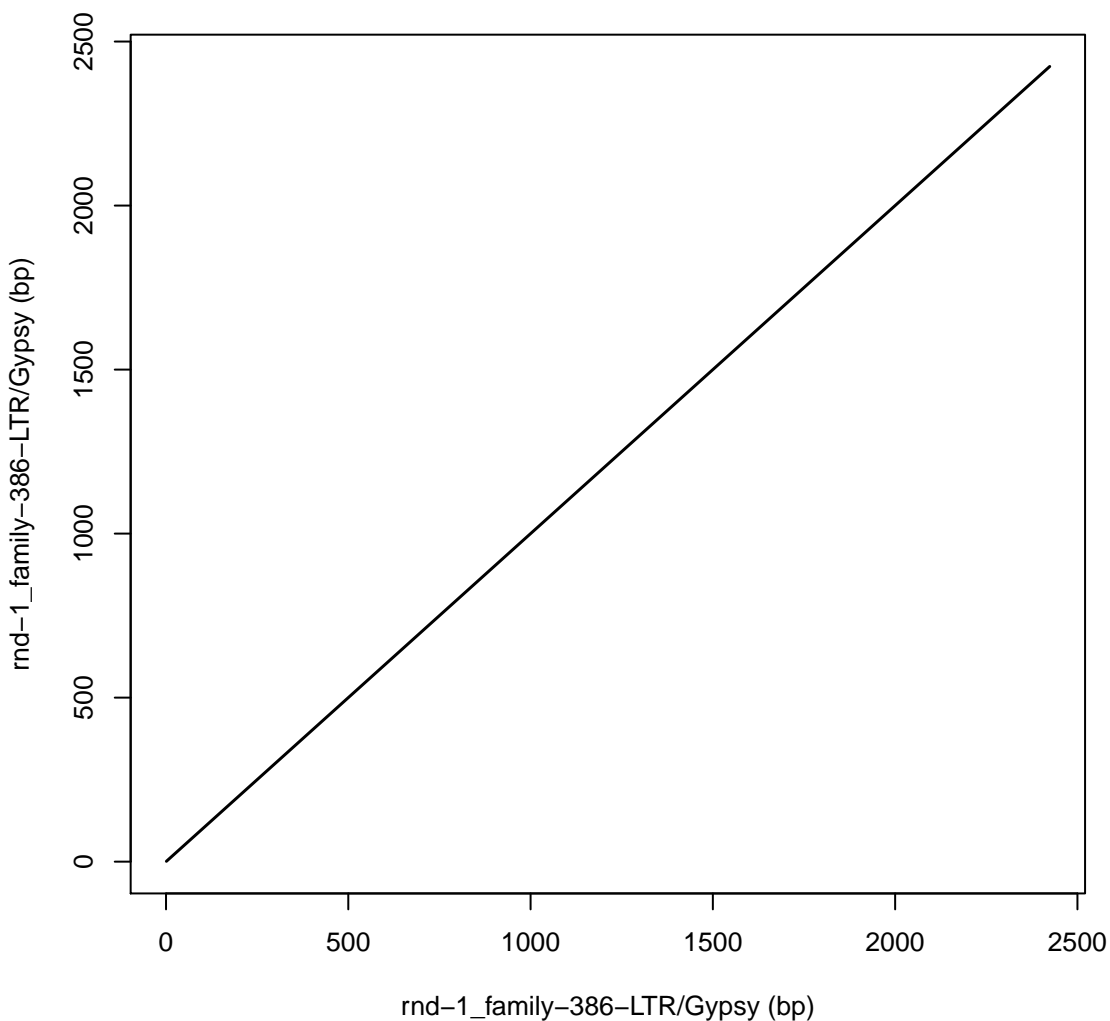
TE: rnd-1_family-386-LTR/Gypsy
consensus size: 2424bp; fragments: 206; full length: 24 (>=2181.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

