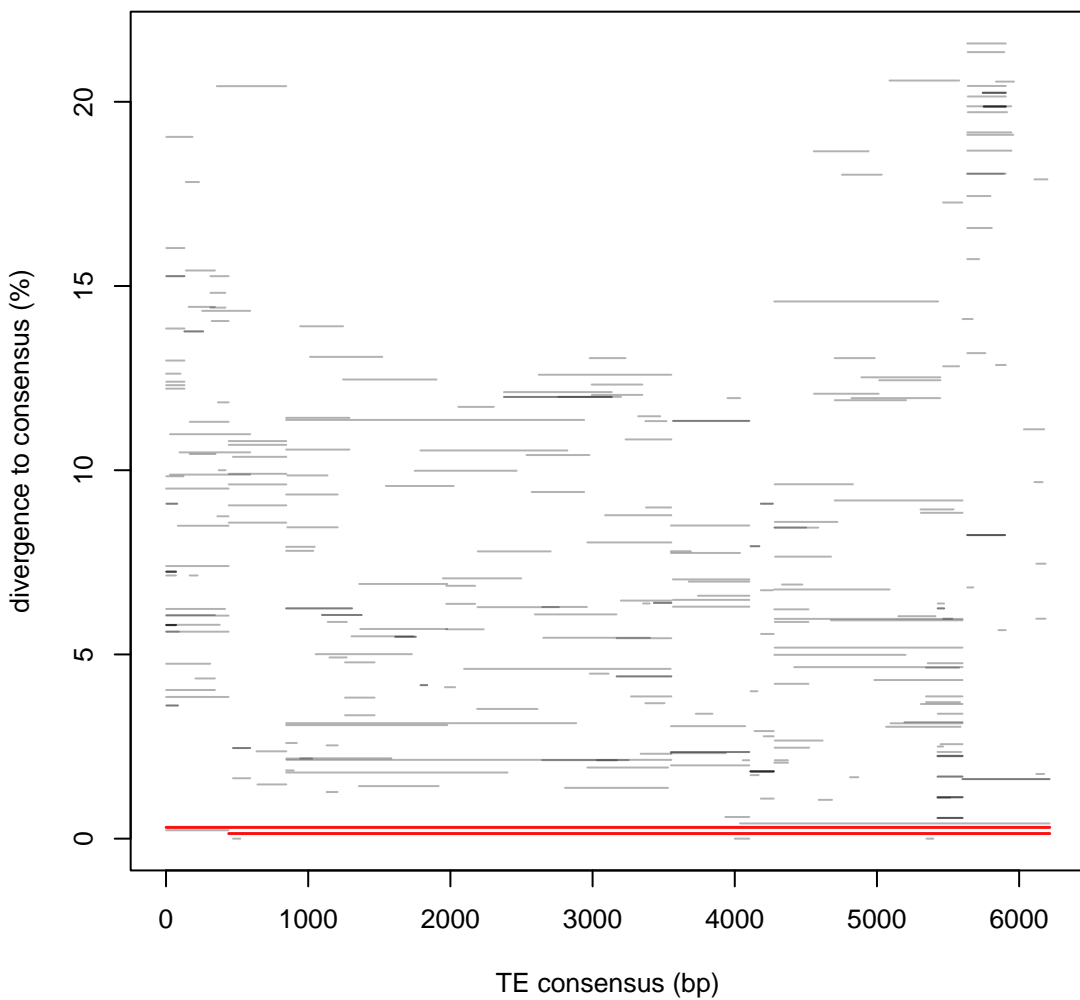
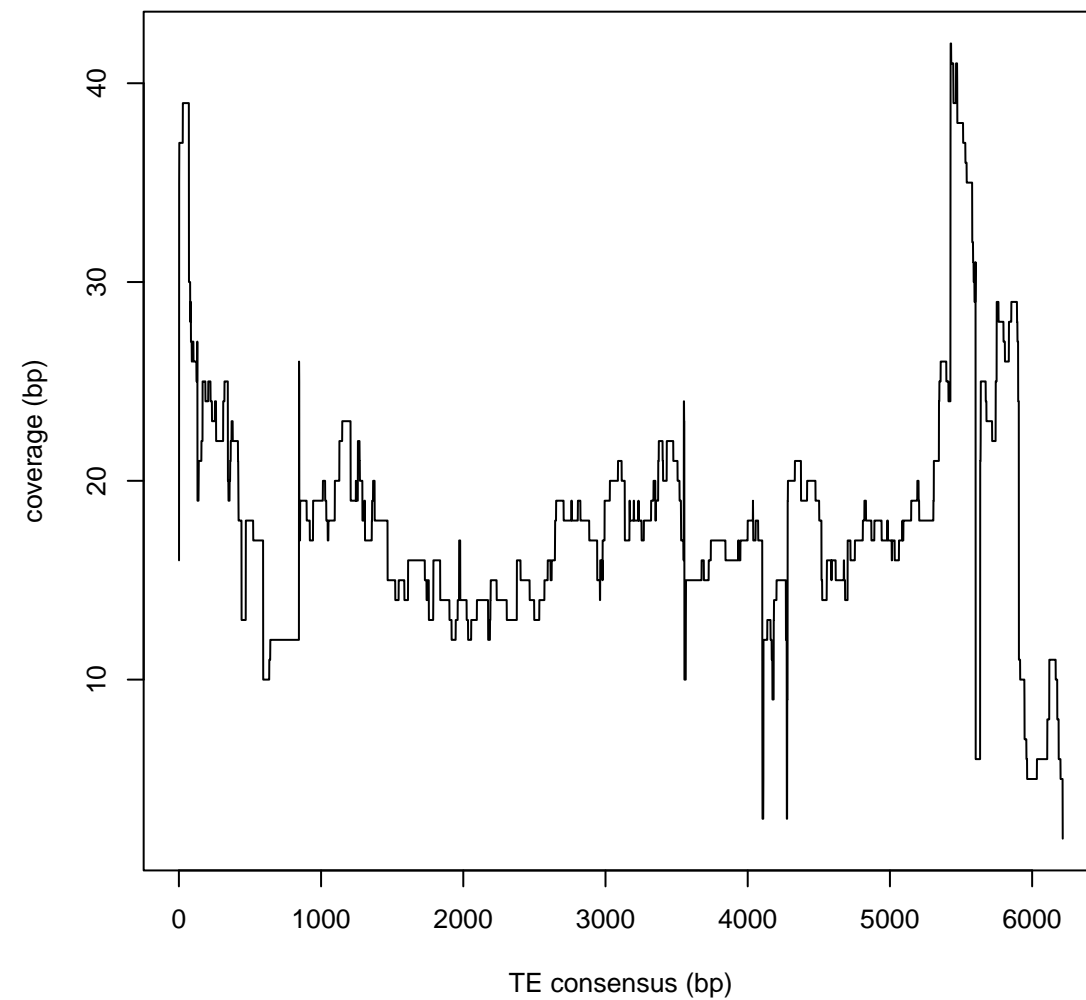


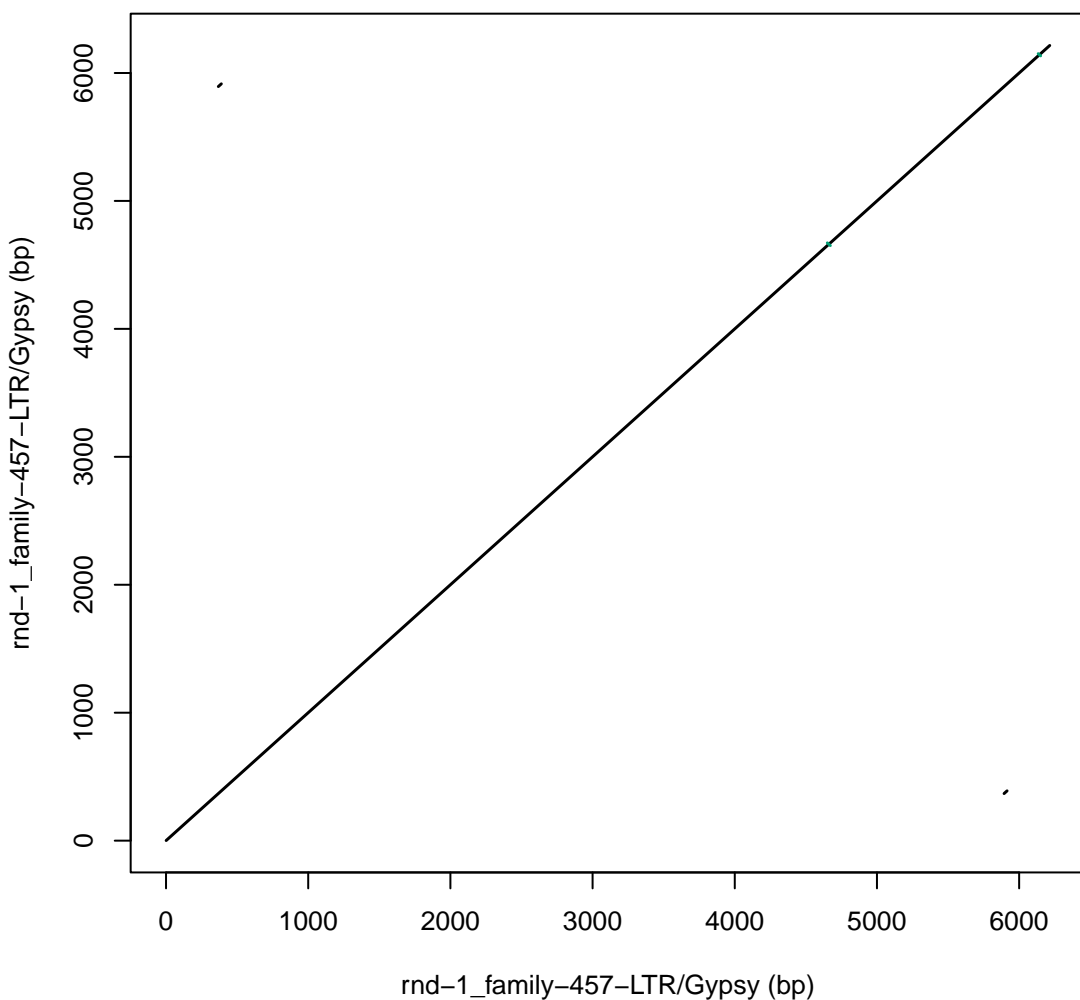
TE: rnd-1_family-457-LTR/Gypsy
consensus size: 6215bp; fragments: 301; full length: 2 (>=5593.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

