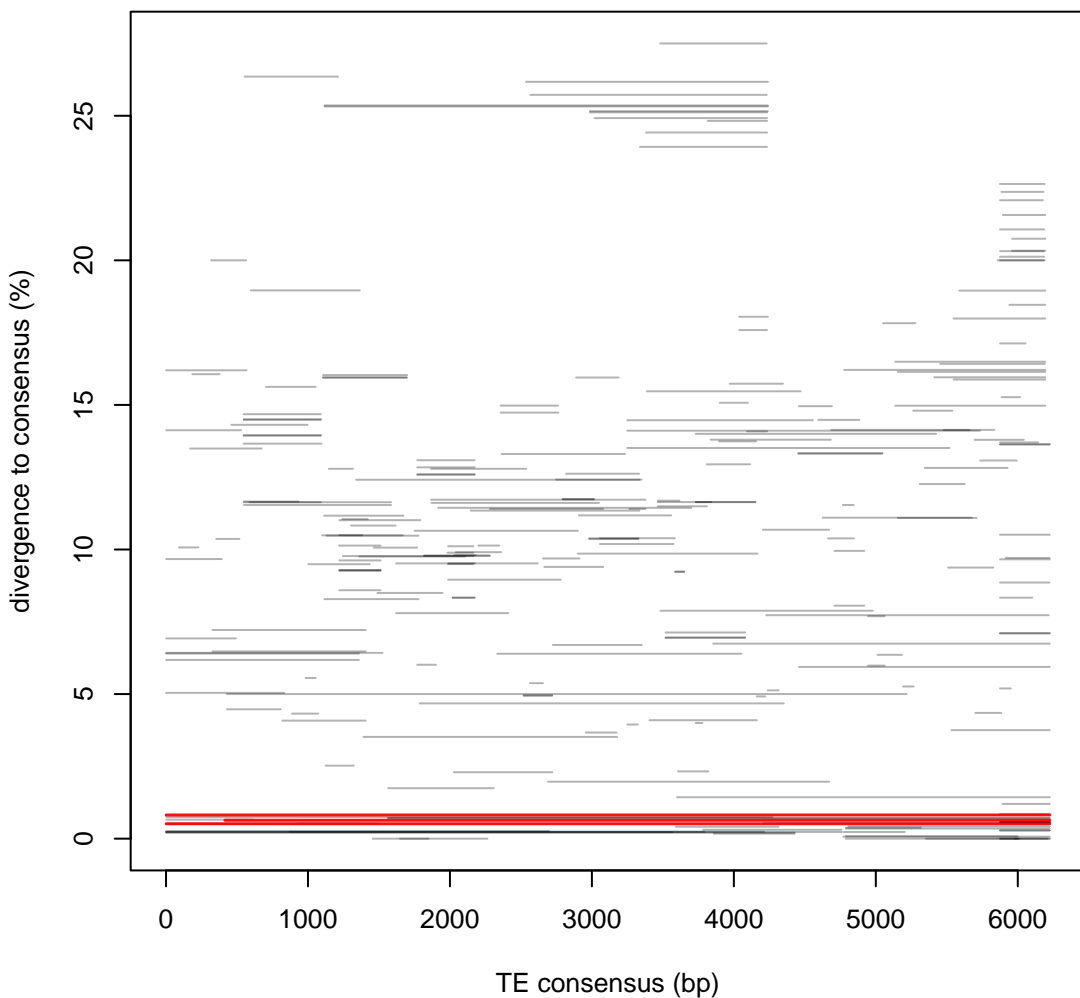
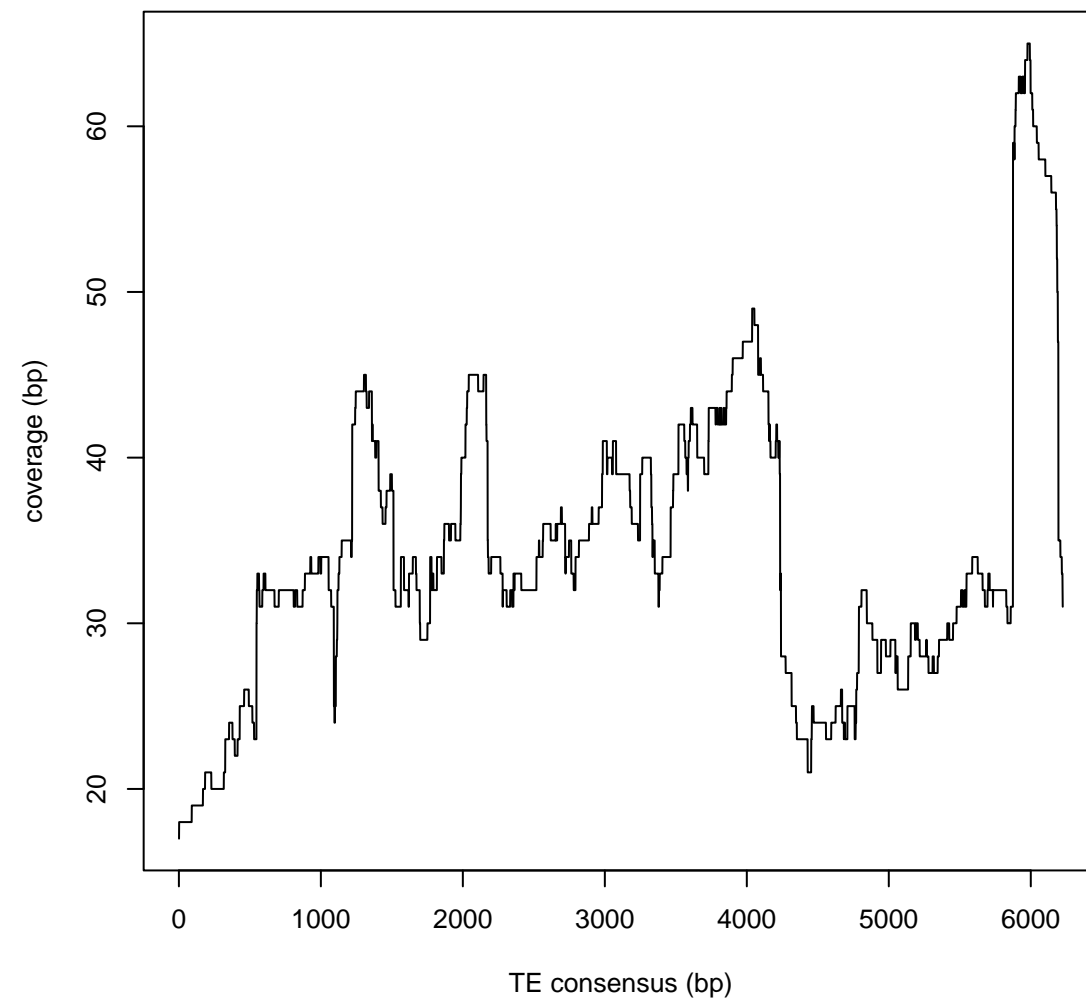


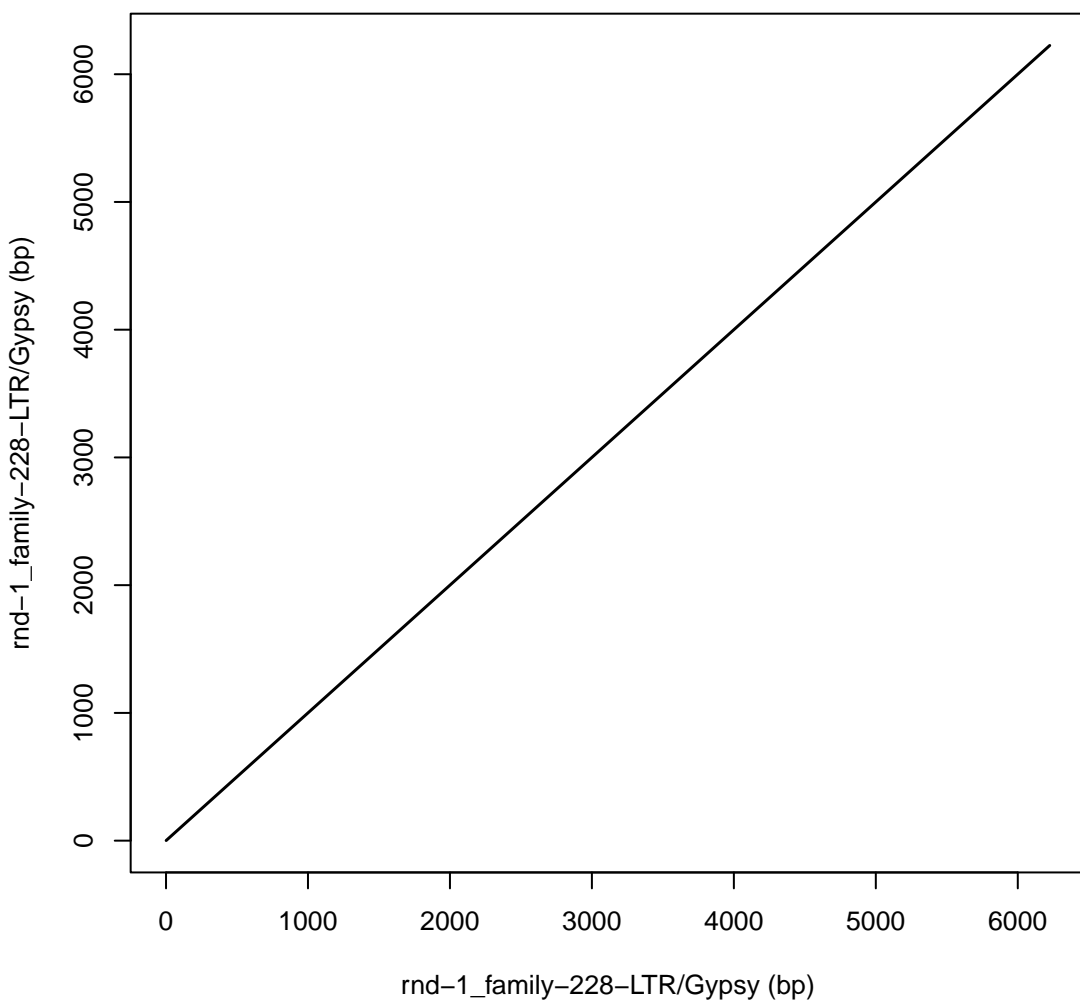
TE: rnd-1_family-228-LTR/Gypsy
consensus size: 6225bp; fragments: 260; full length: 3 (>=5602.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

