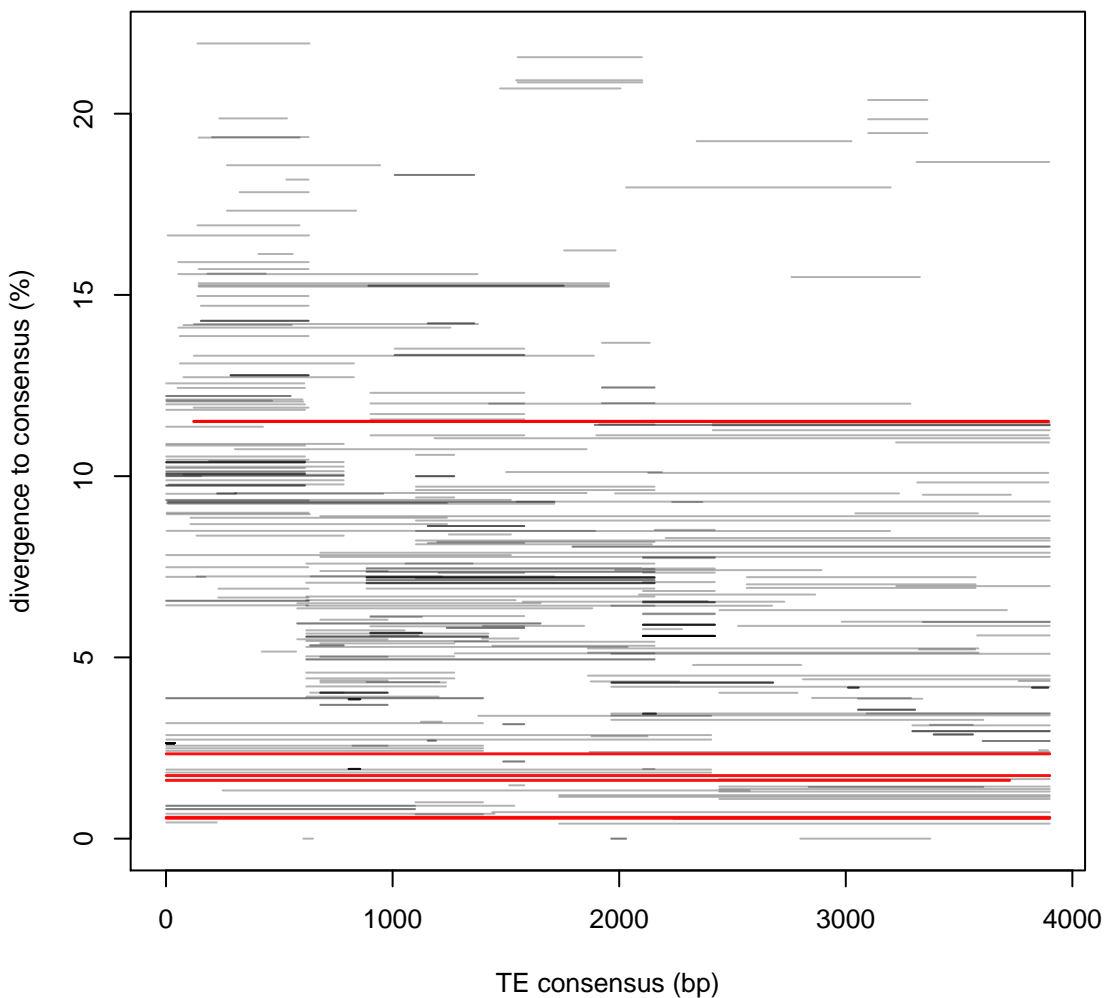
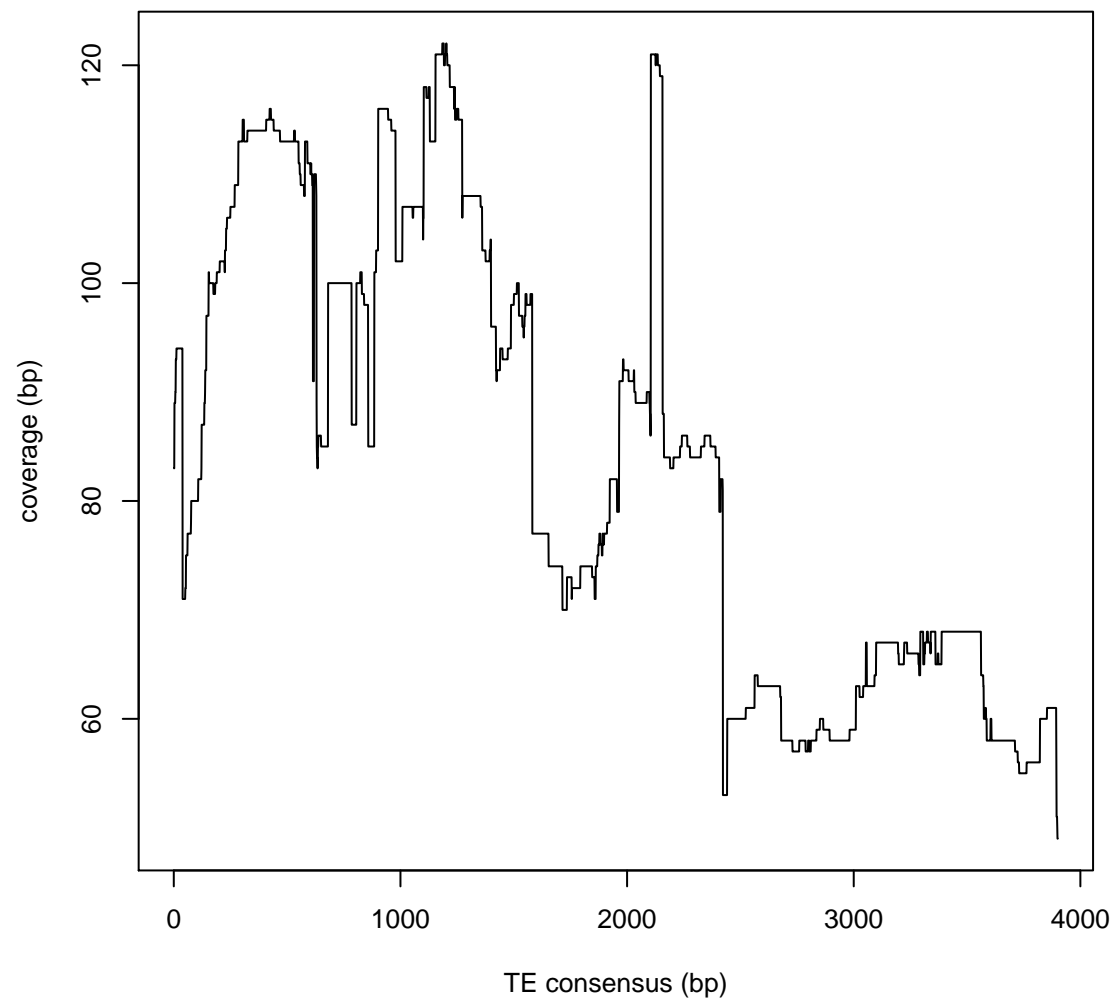


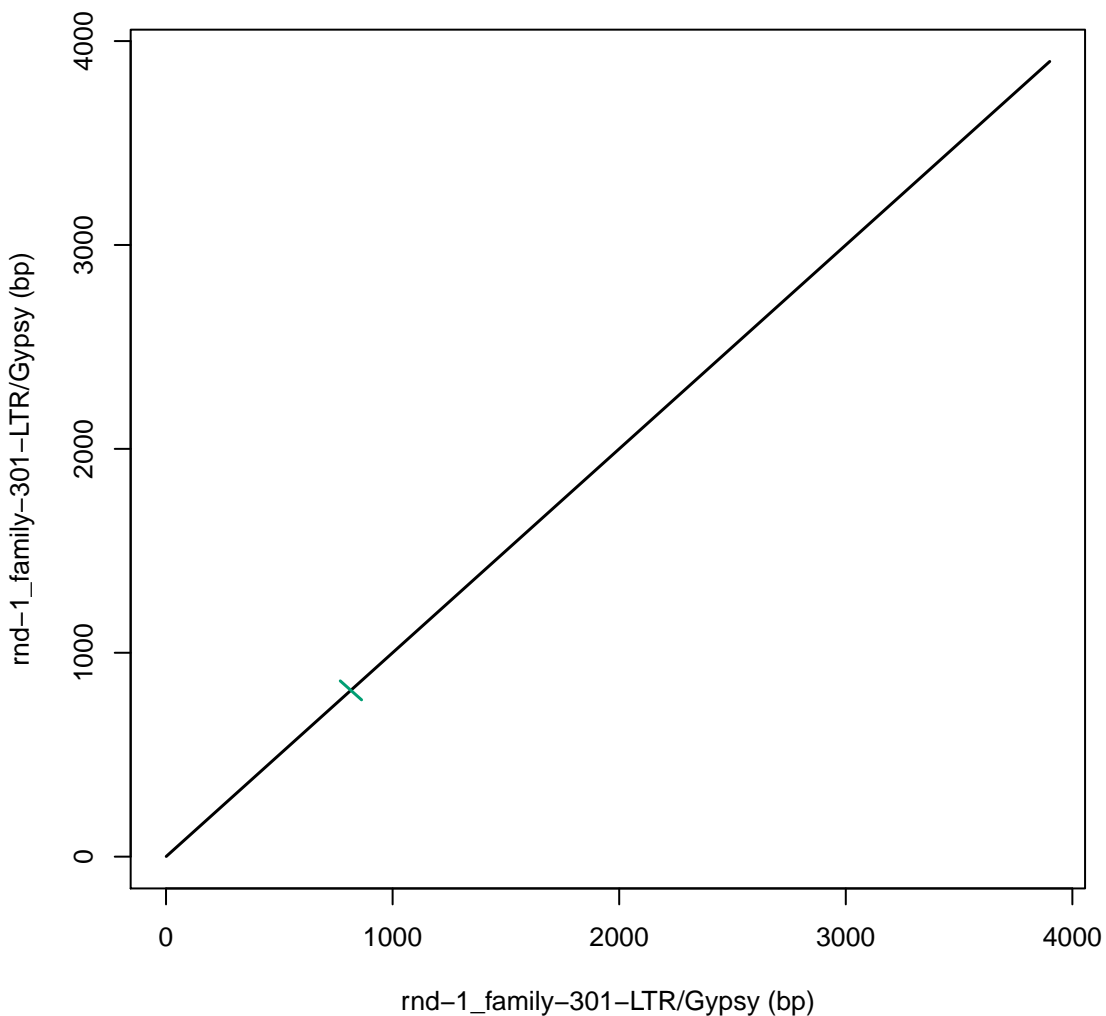
TE: rnd-1_family-301-LTR/Gypsy
consensus size: 3900bp; fragments: 424; full length: 7 (>=3510bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

