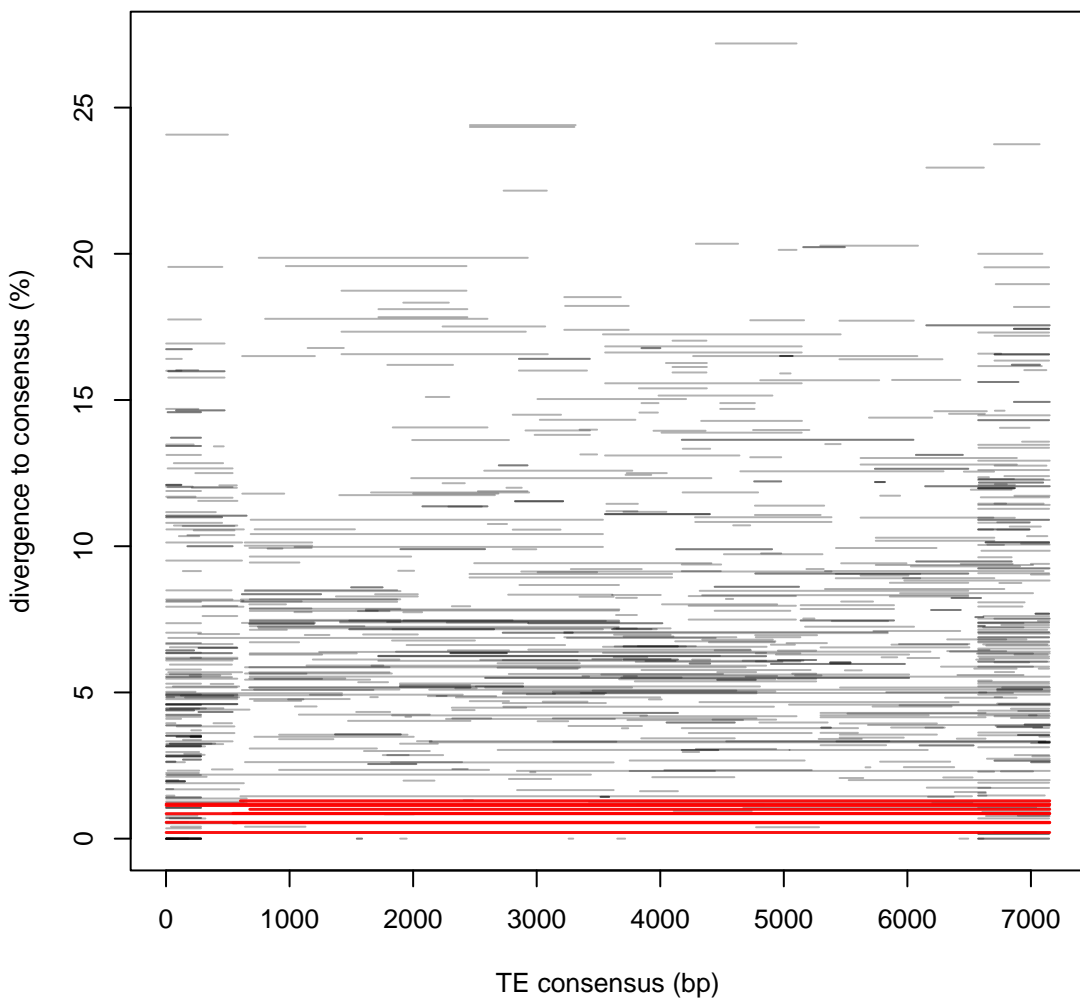
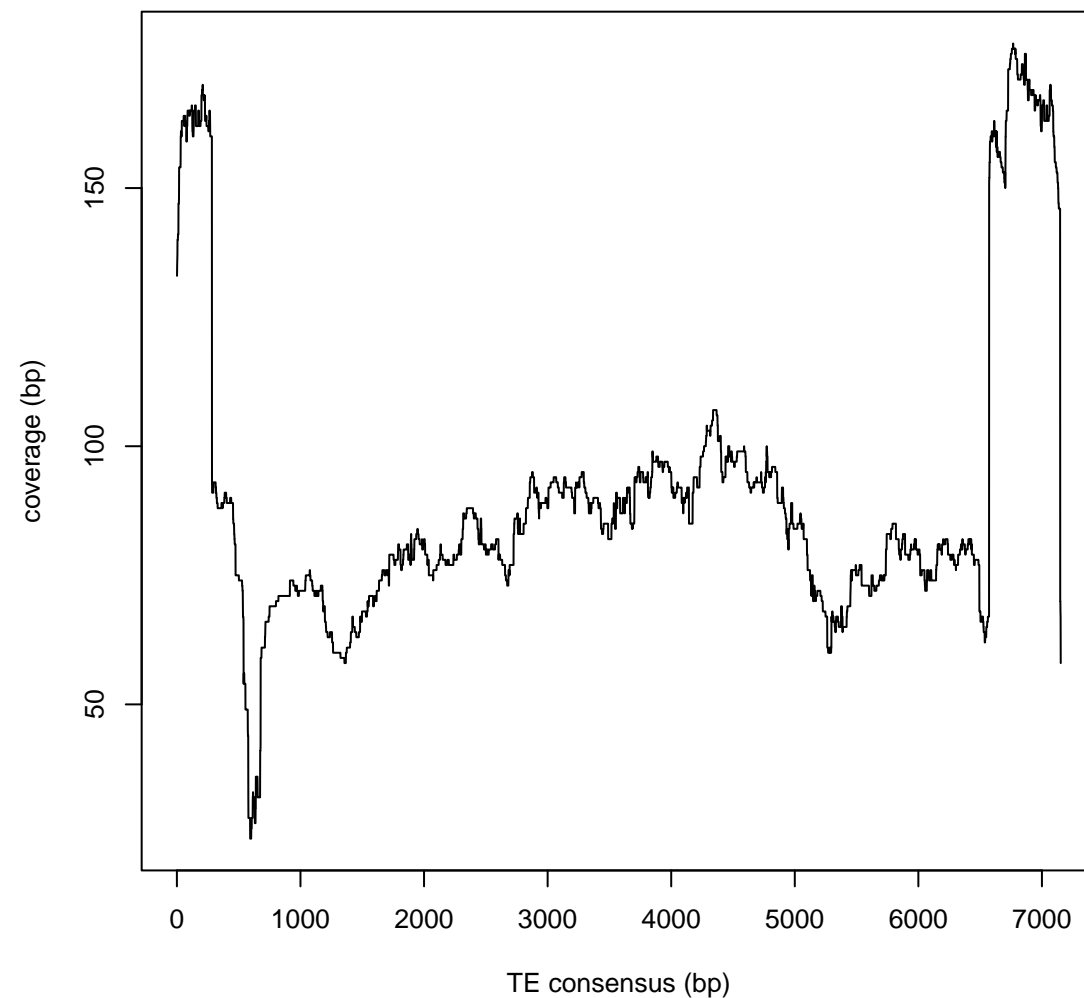


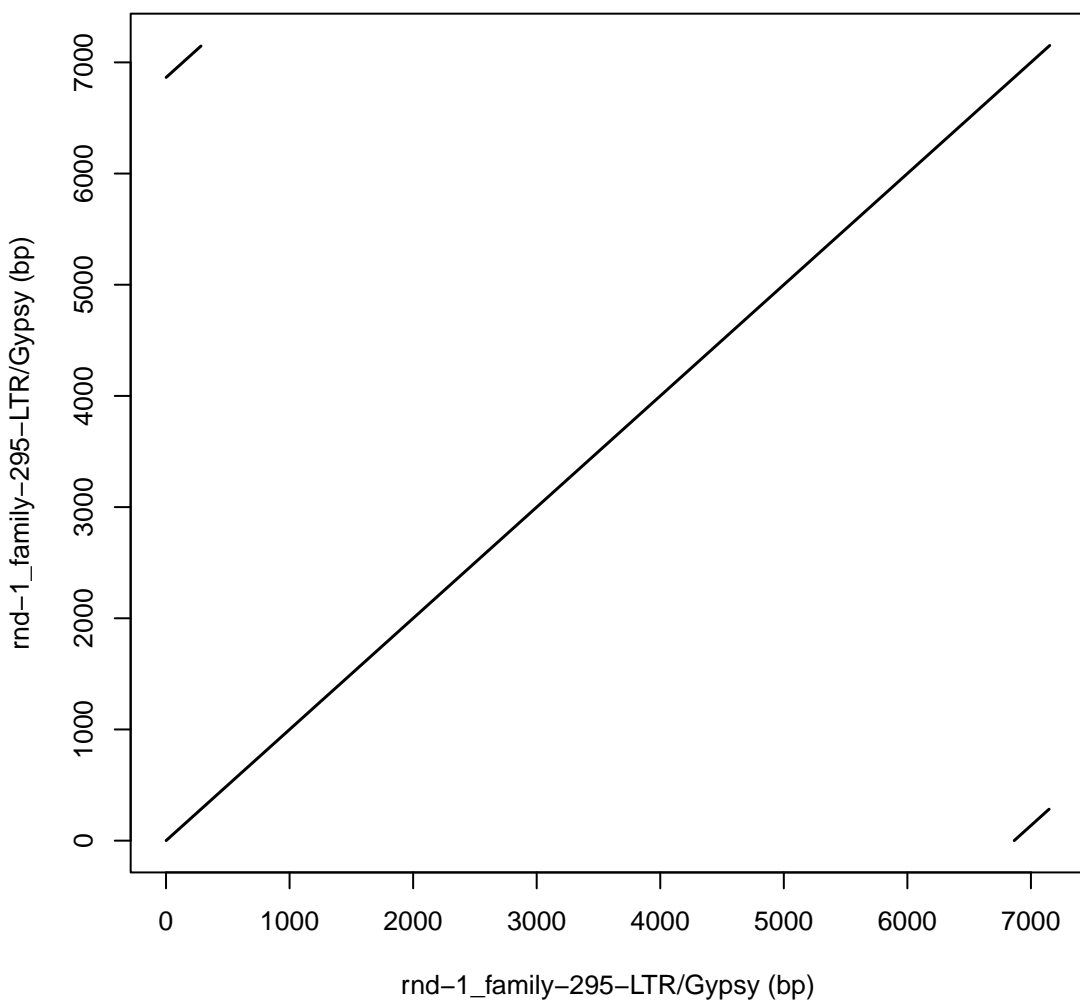
TE: rnd-1_family-295-LTR/Gypsy
consensus size: 7152bp; fragments: 913; full length: 9 (≥ 6436.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

