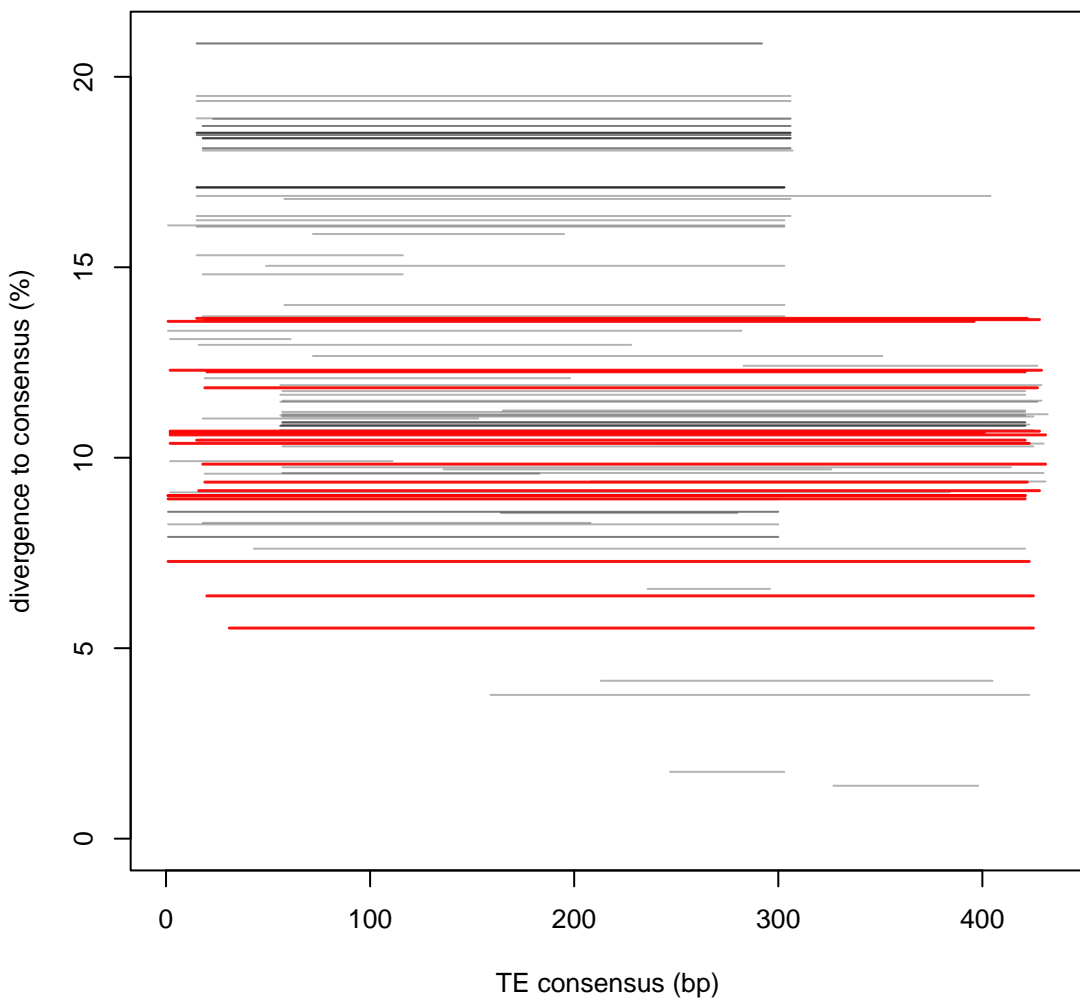
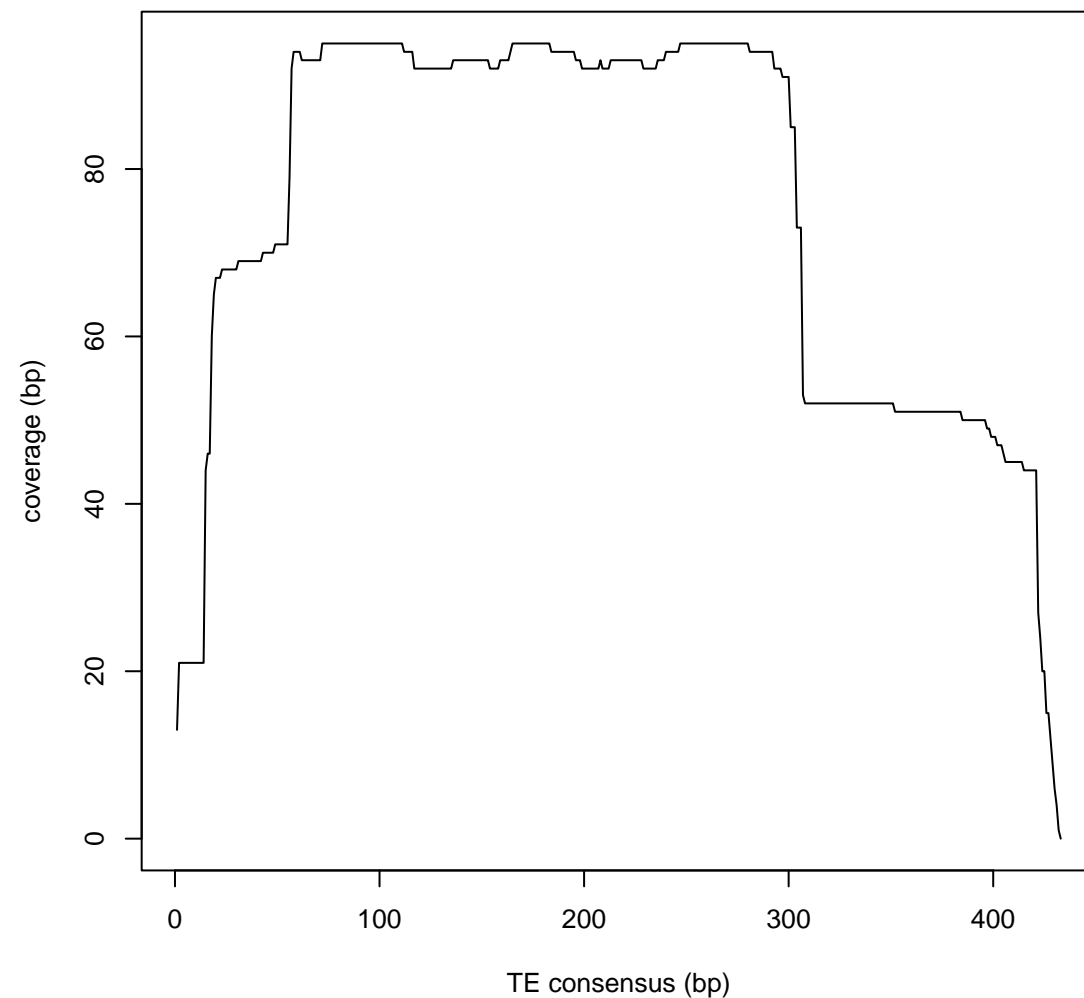


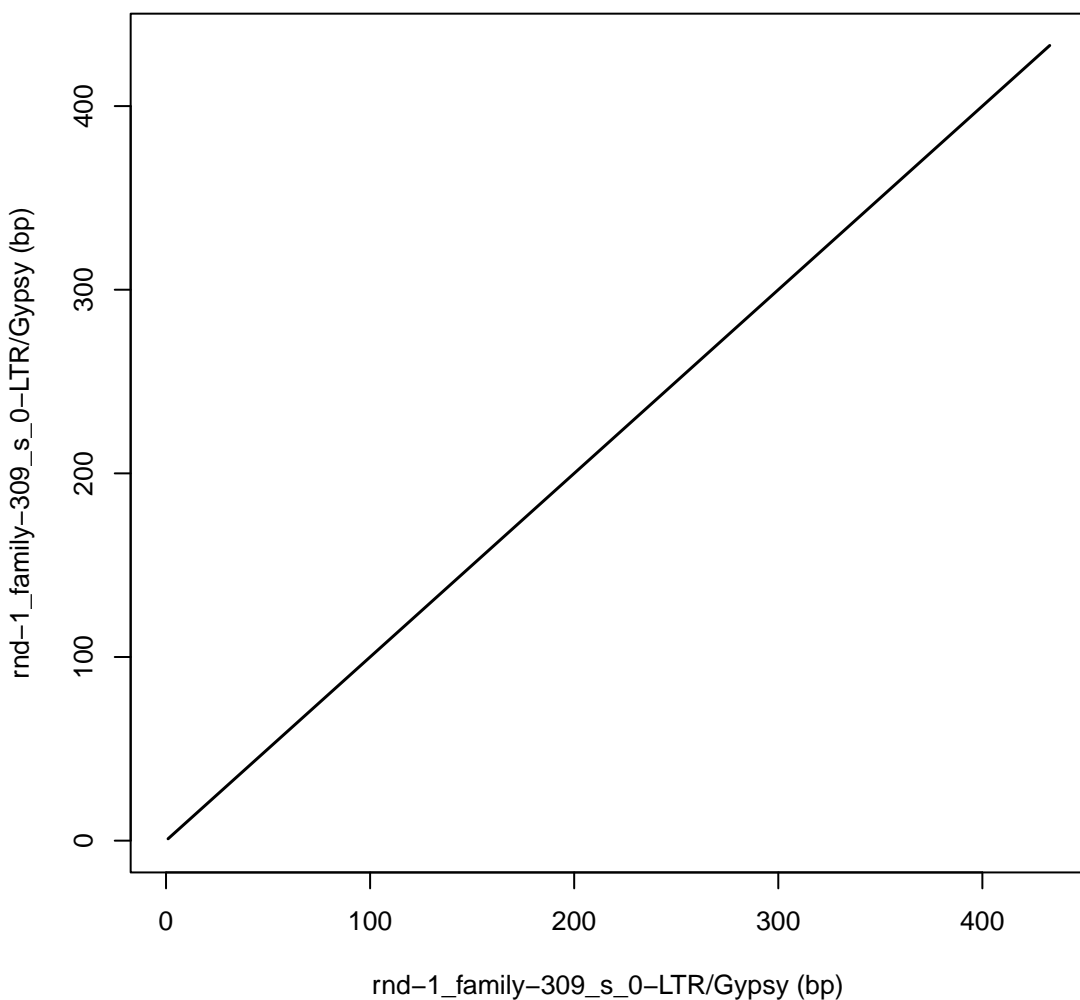
TE: rnd-1_family-309_s_0-LTR/Gypsy
consensus size: 433bp; fragments: 107; full length: 20 (≥ 389.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

