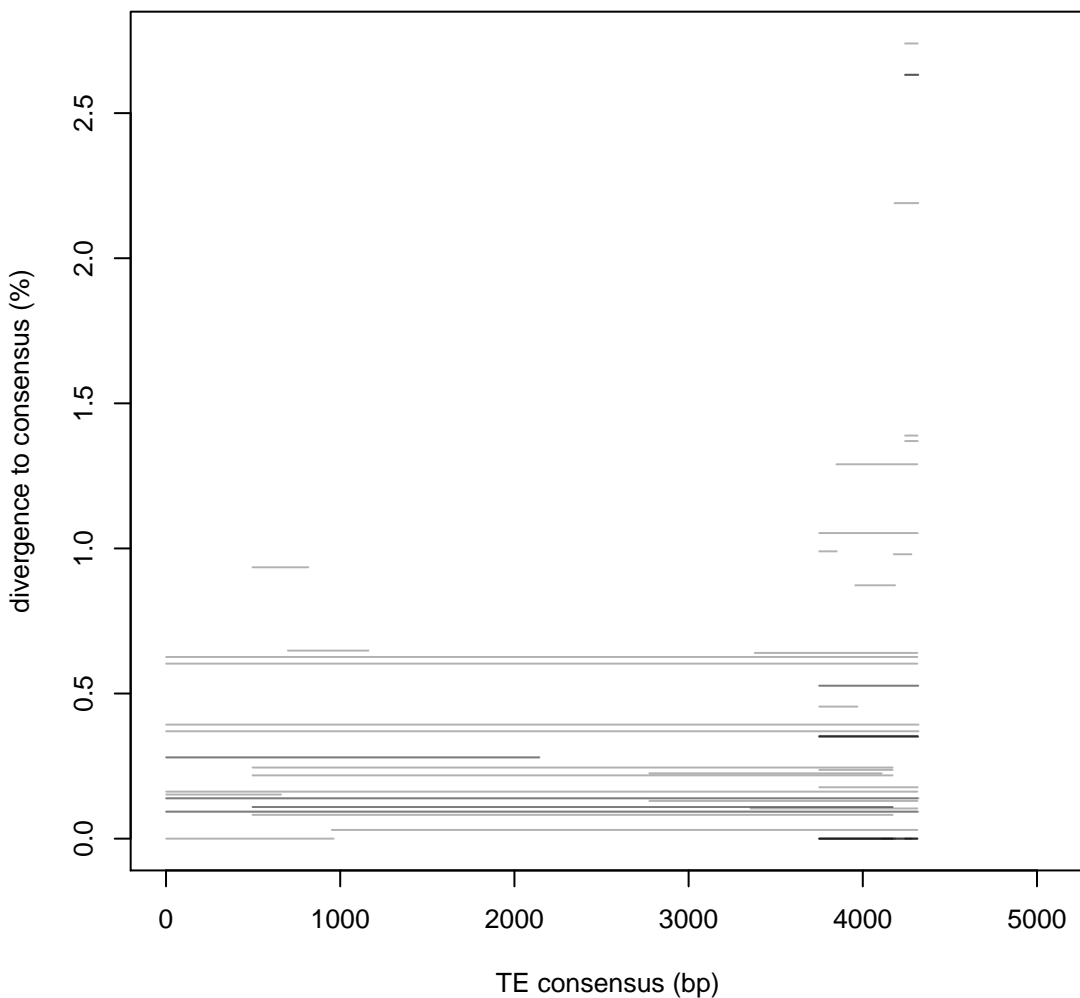
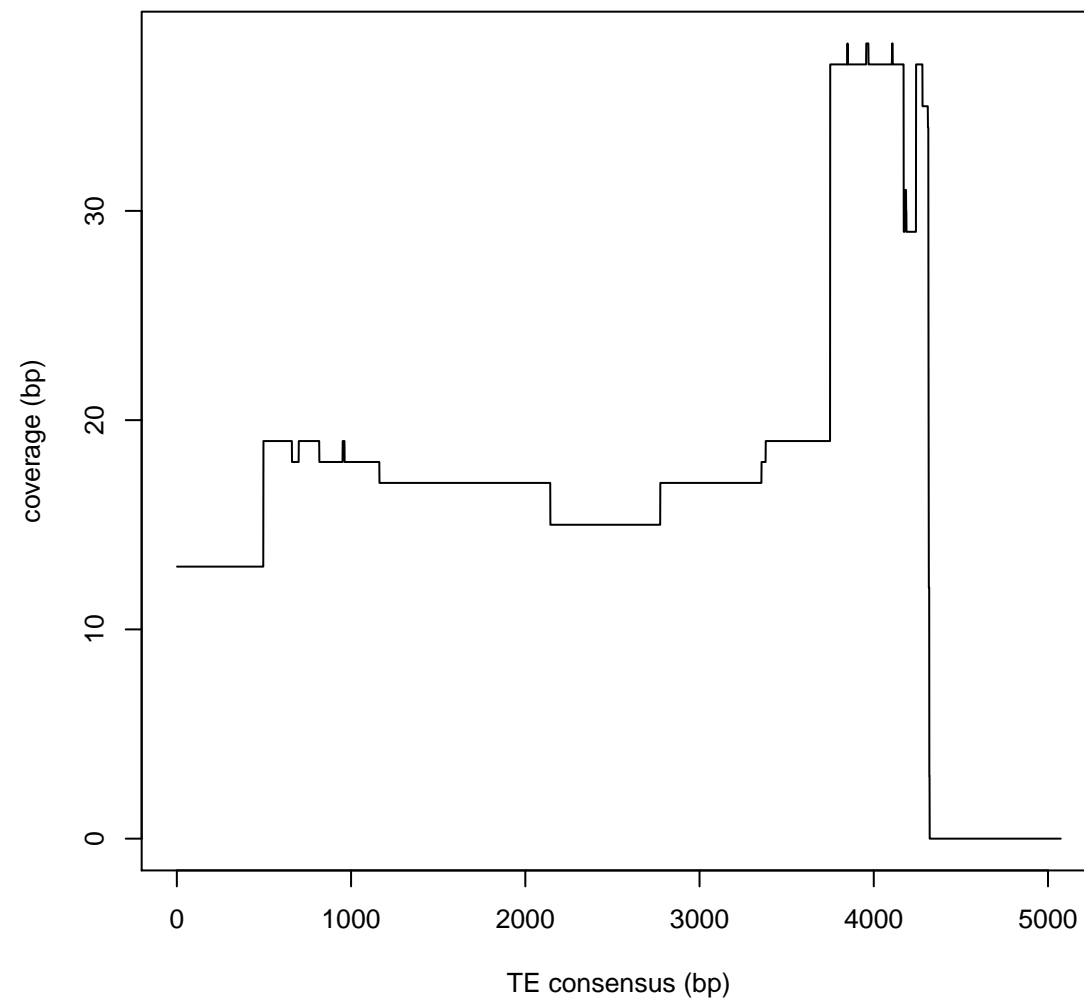


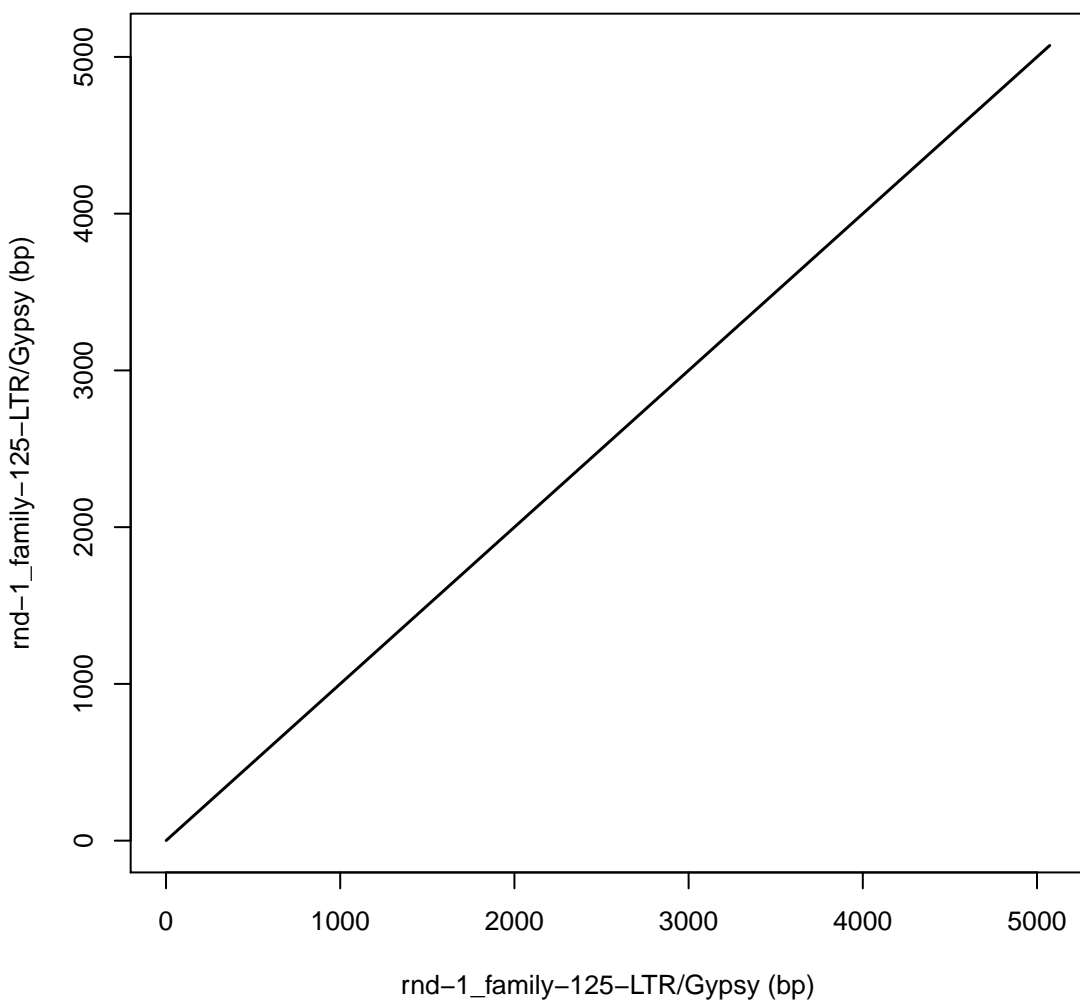
TE: rnd-1_family-125-LTR/Gypsy
consensus size: 5073bp; fragments: 56; full length: 0 (≥ 4565.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

