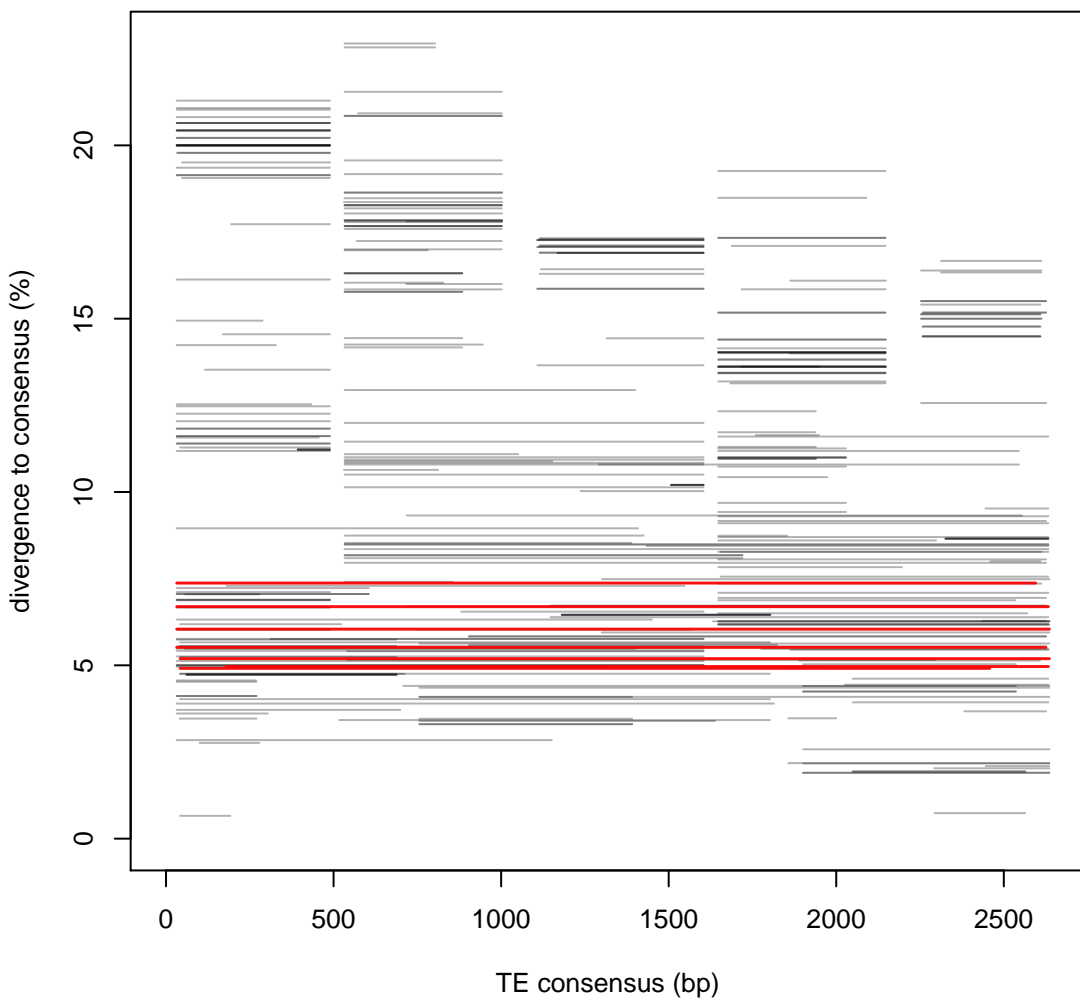
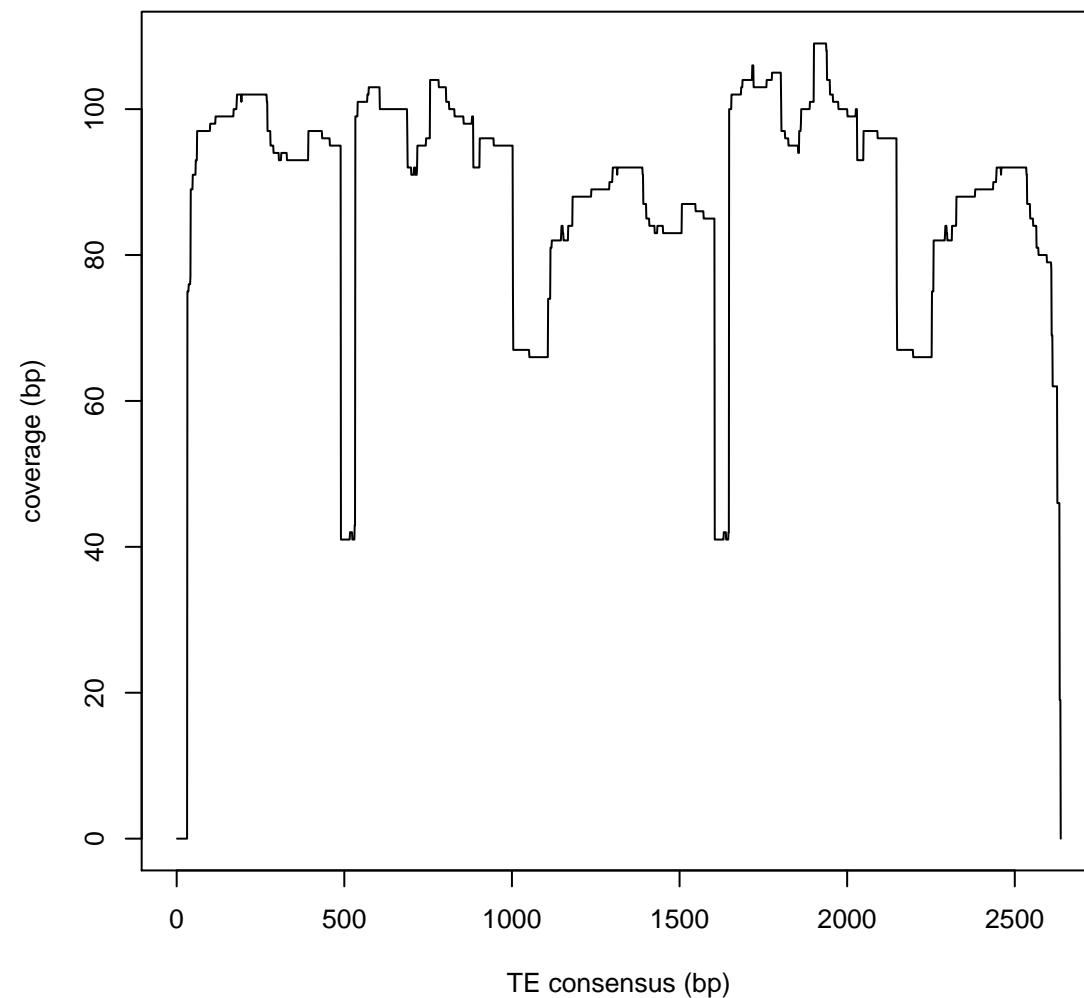


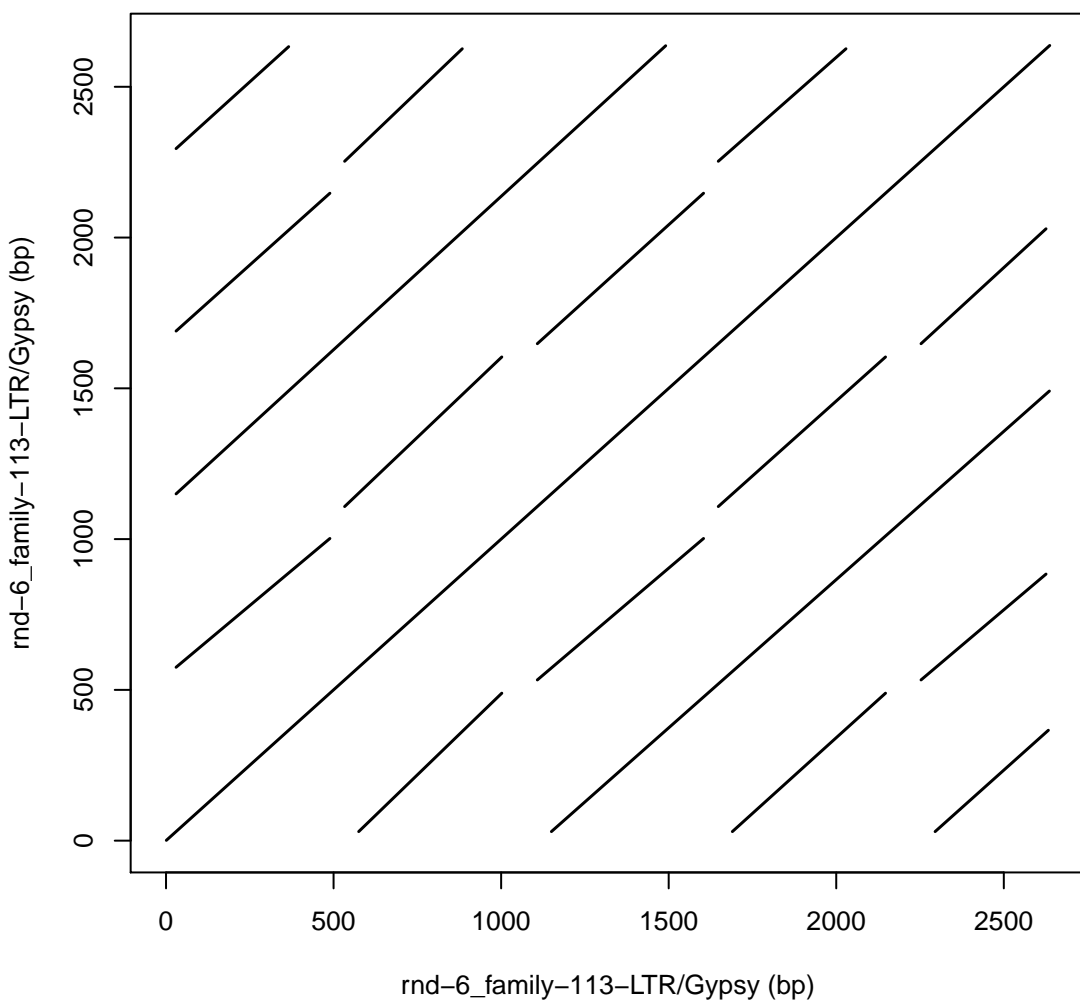
TE: rnd-6_family-113-LTR/Gypsy
consensus size: 2637bp; fragments: 341; full length: 7 (≥ 2373.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

