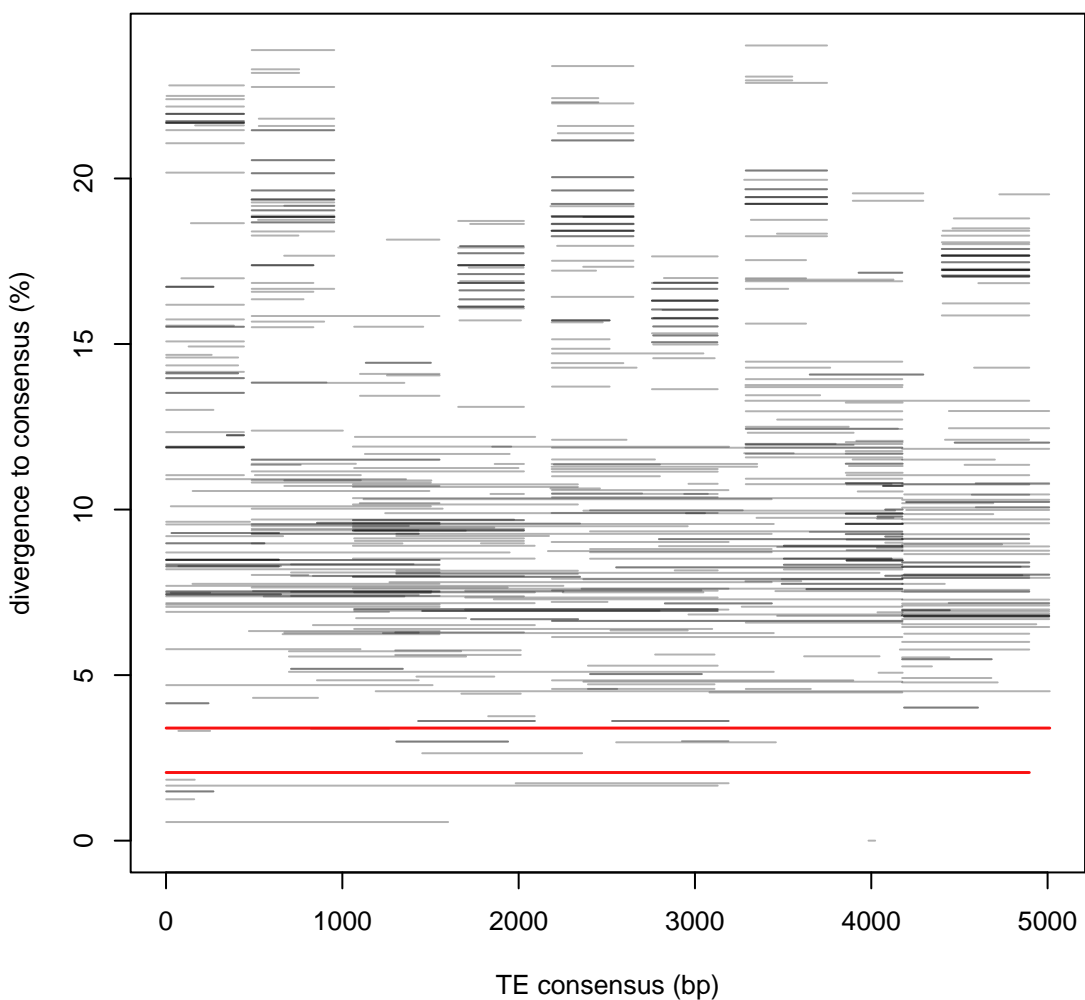
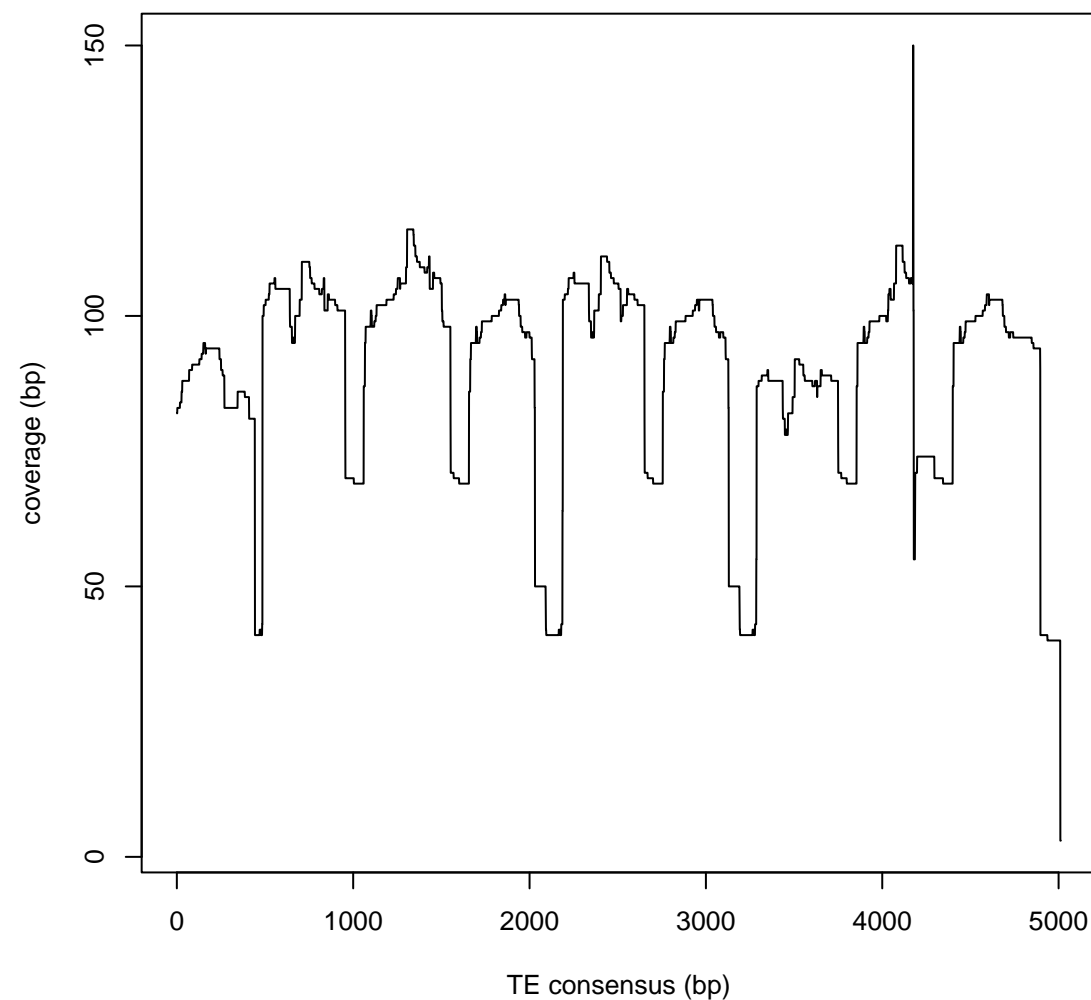


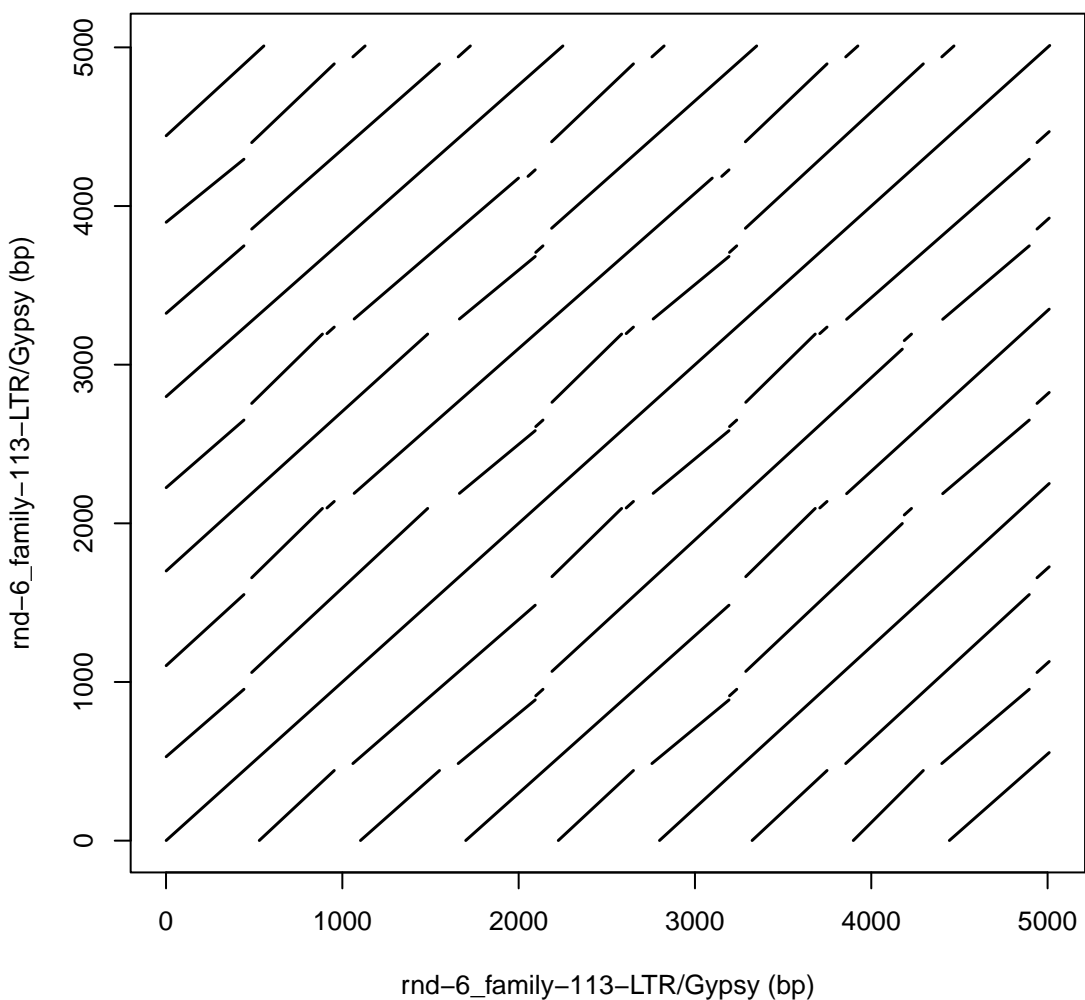
TE: rnd-6_family-113-LTR/Gypsy
consensus size: 5012bp; fragments: 646; full length: 2 (≥ 4510.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

