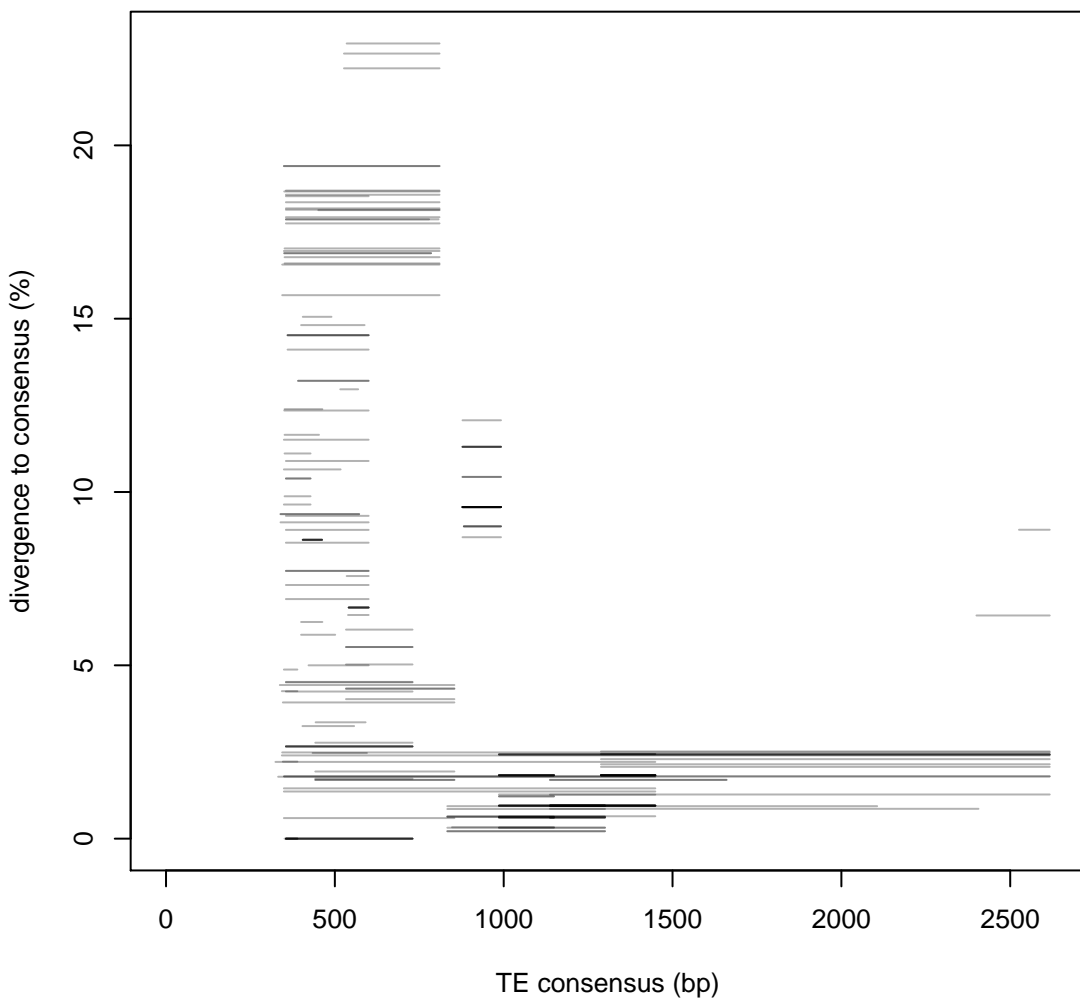
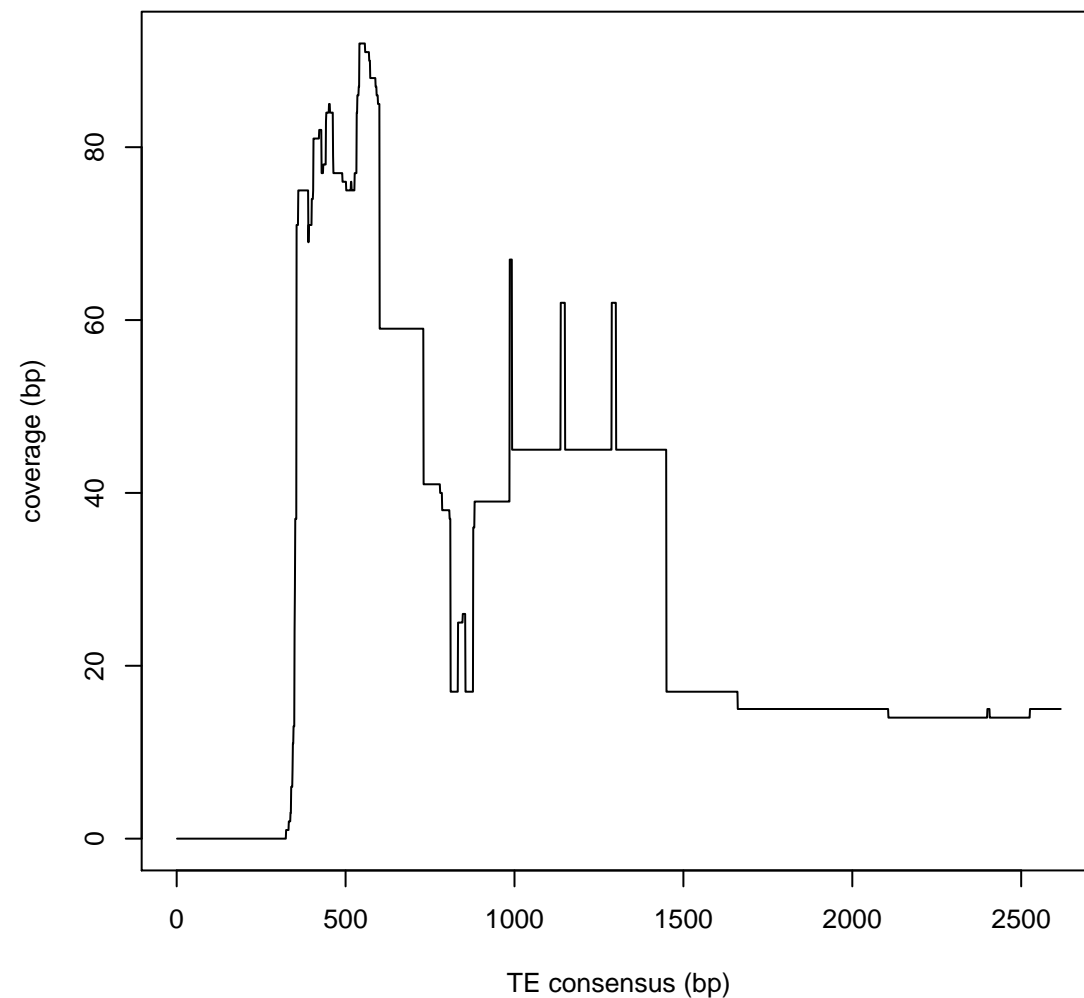


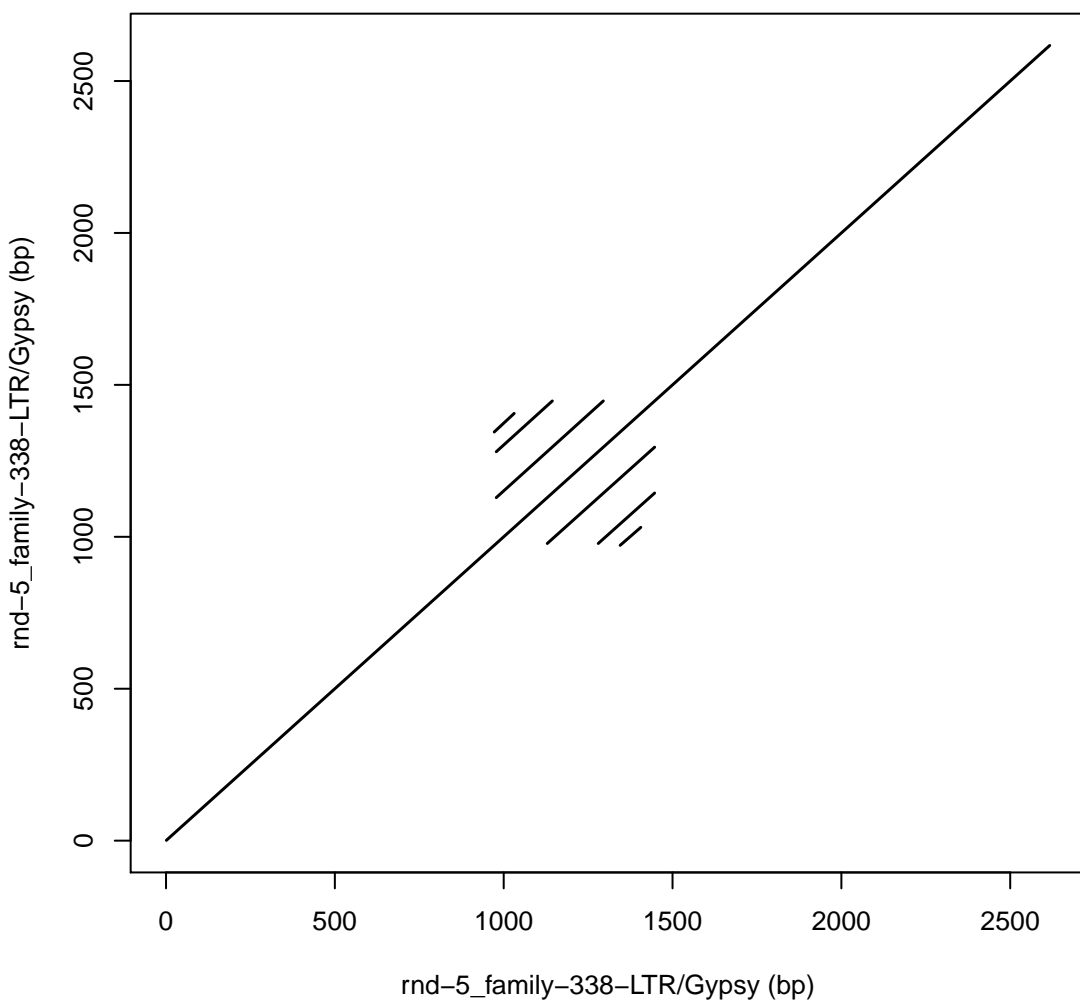
TE: rnd-5_family-338-LTR/Gypsy
consensus size: 2617bp; fragments: 209; full length: 0 (>=2355.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

