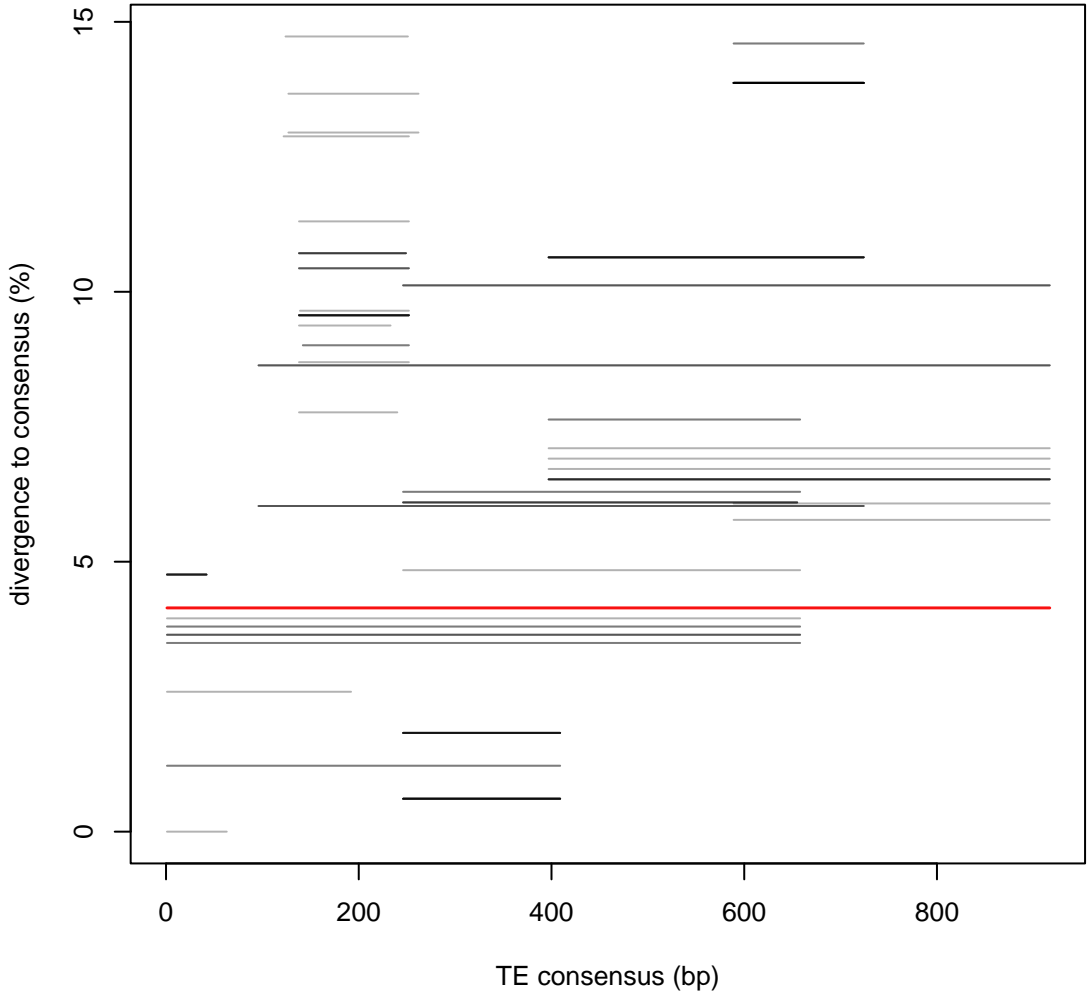
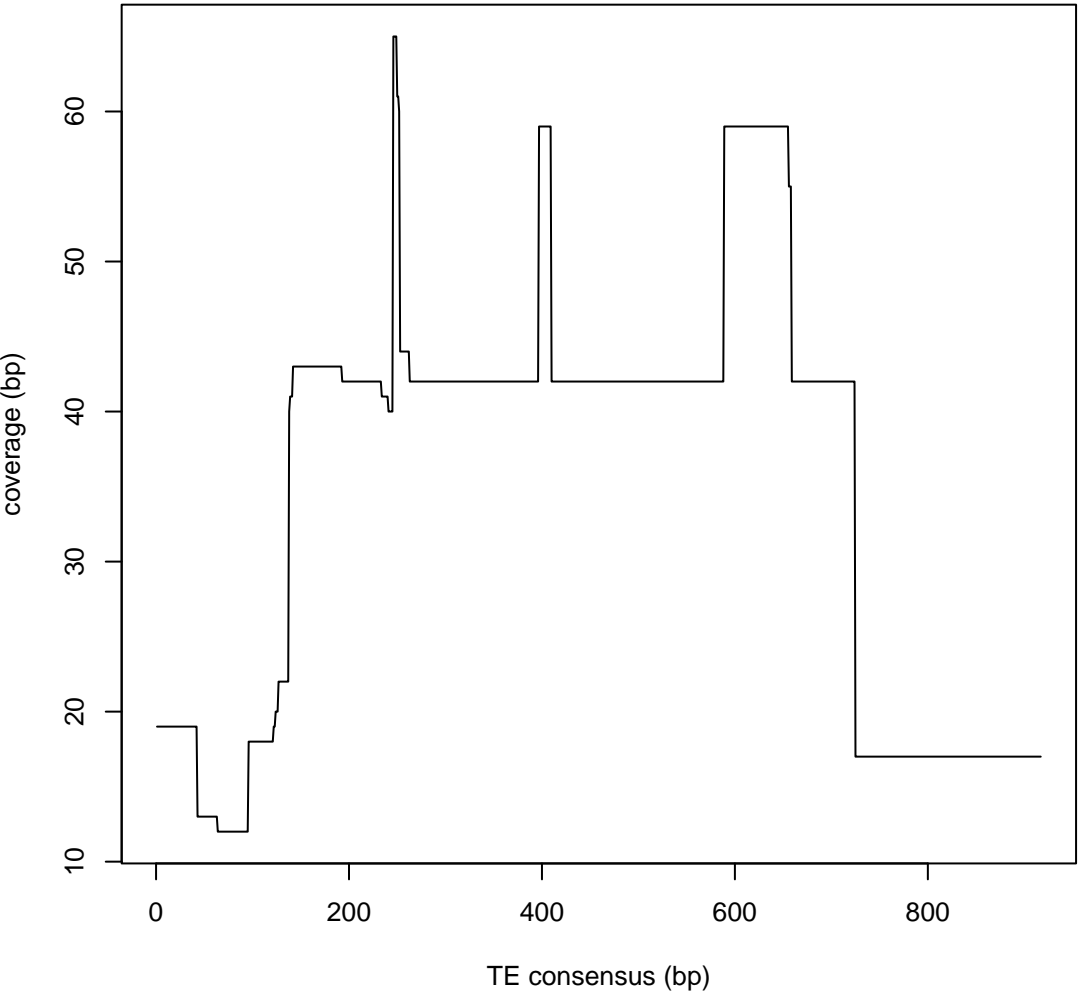


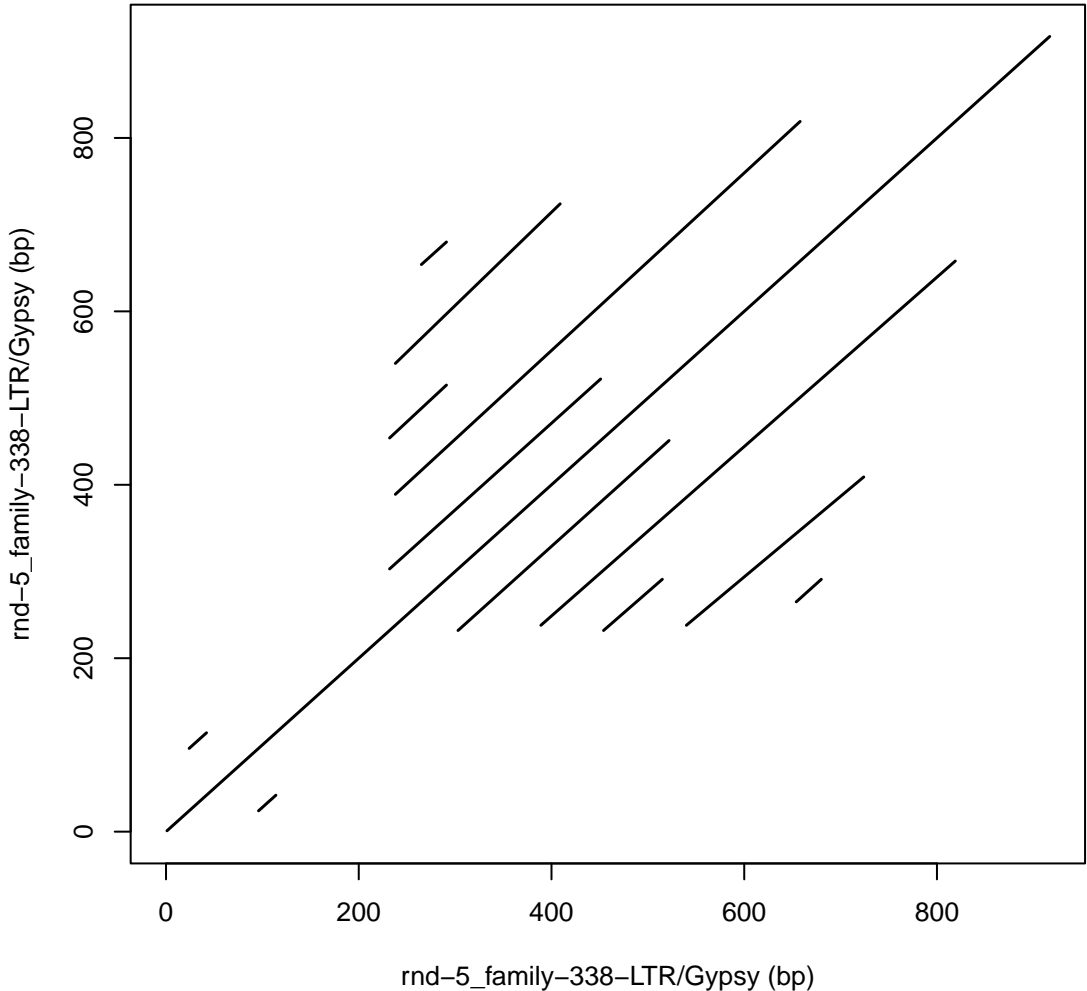
TE: rnd-5_family-338-LTR/Gypsy
consensus size: 917bp; fragments: 109; full length: 1 (>=825.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

