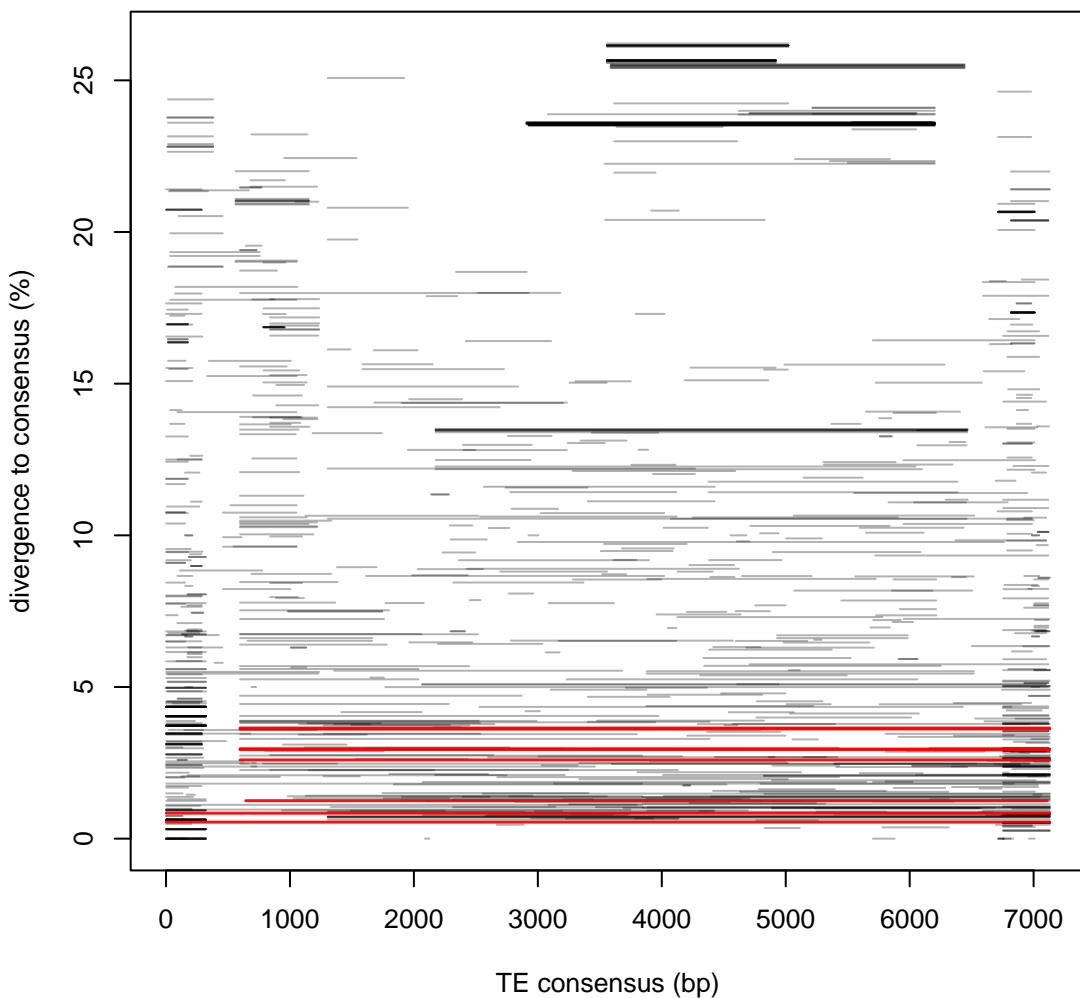
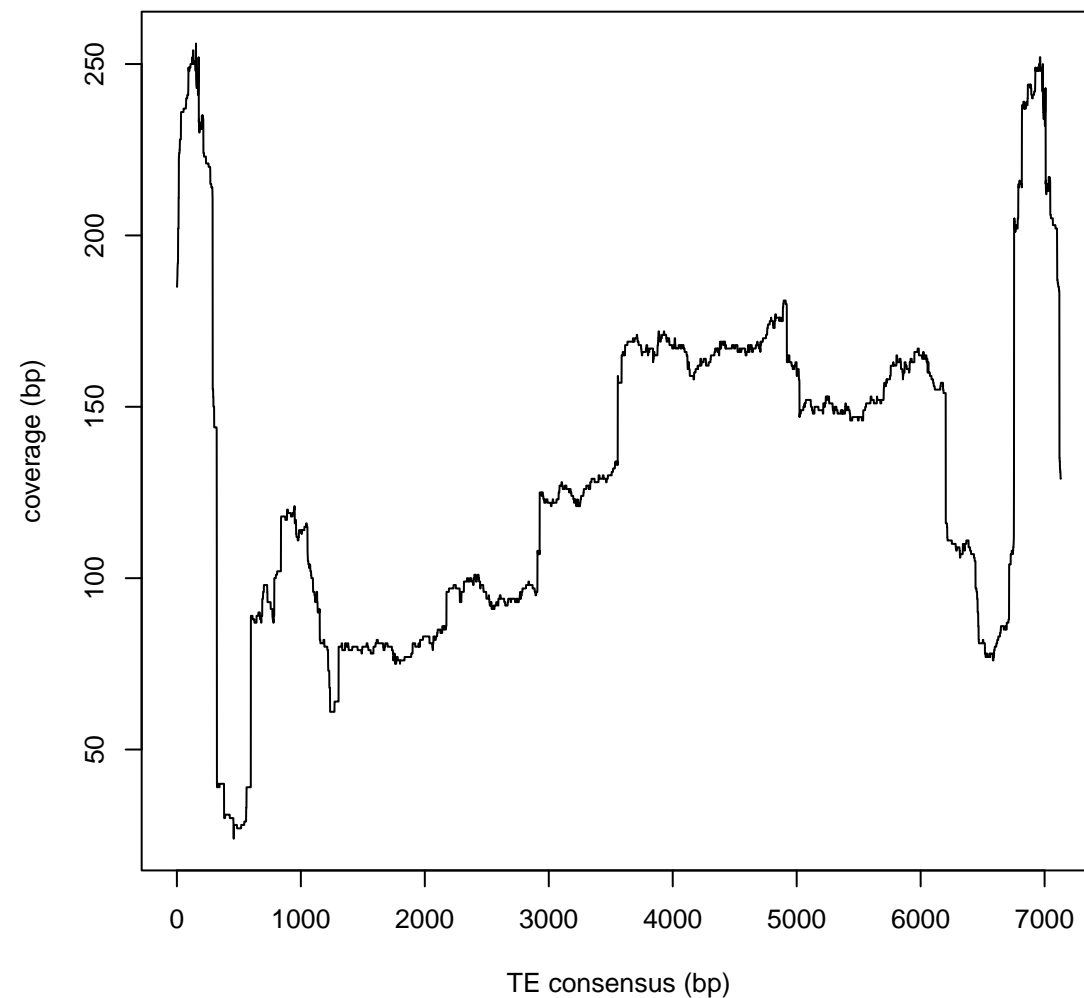


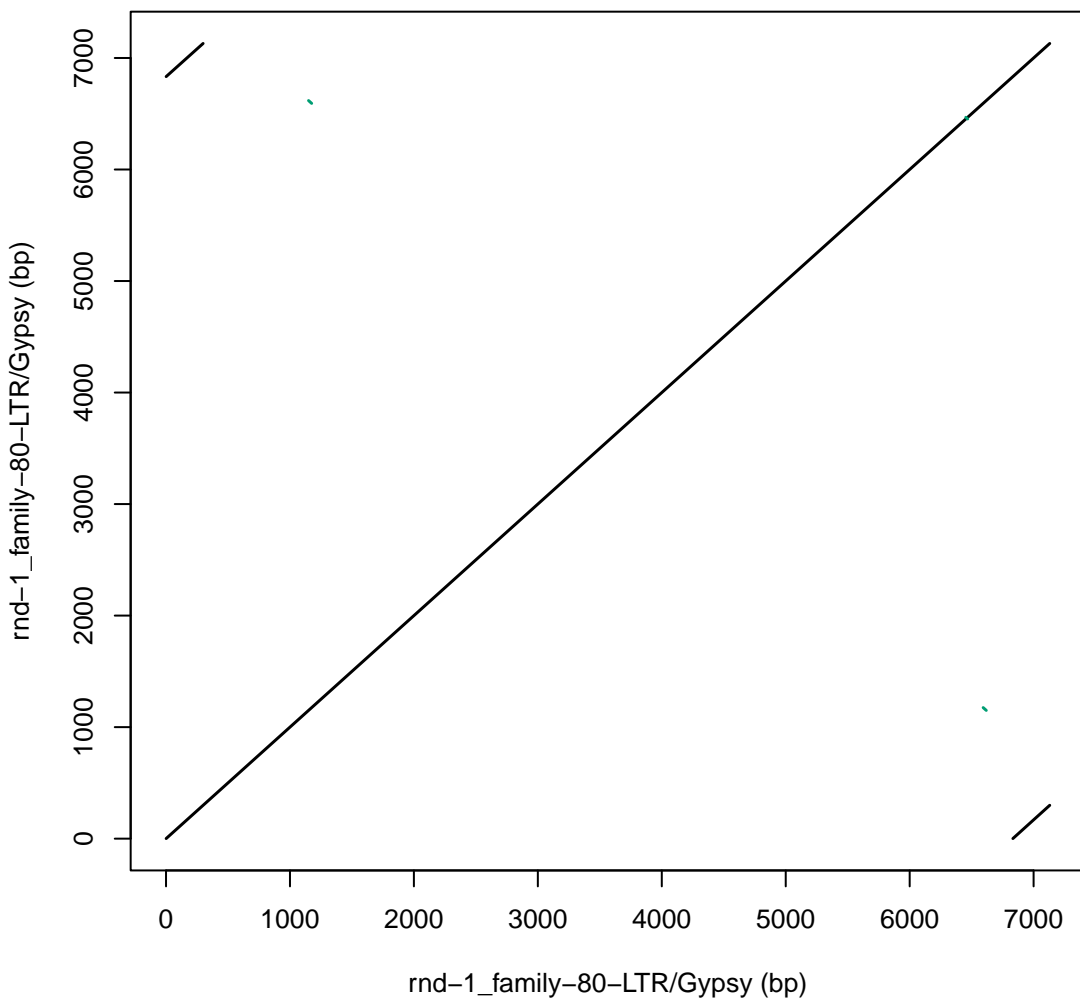
TE: rnd-1_family-80-LTR/Gypsy
consensus size: 7130bp; fragments: 1063; full length: 8 (>=6417bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

