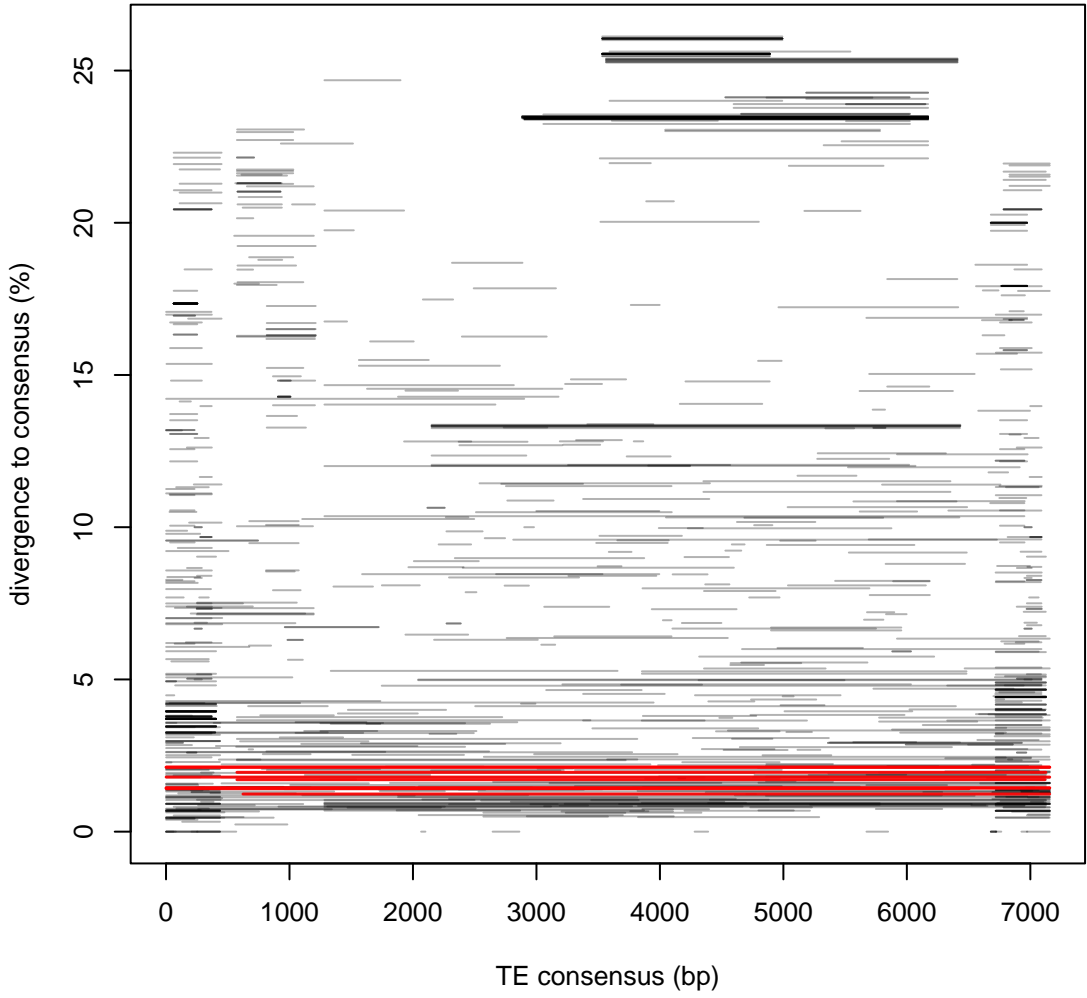
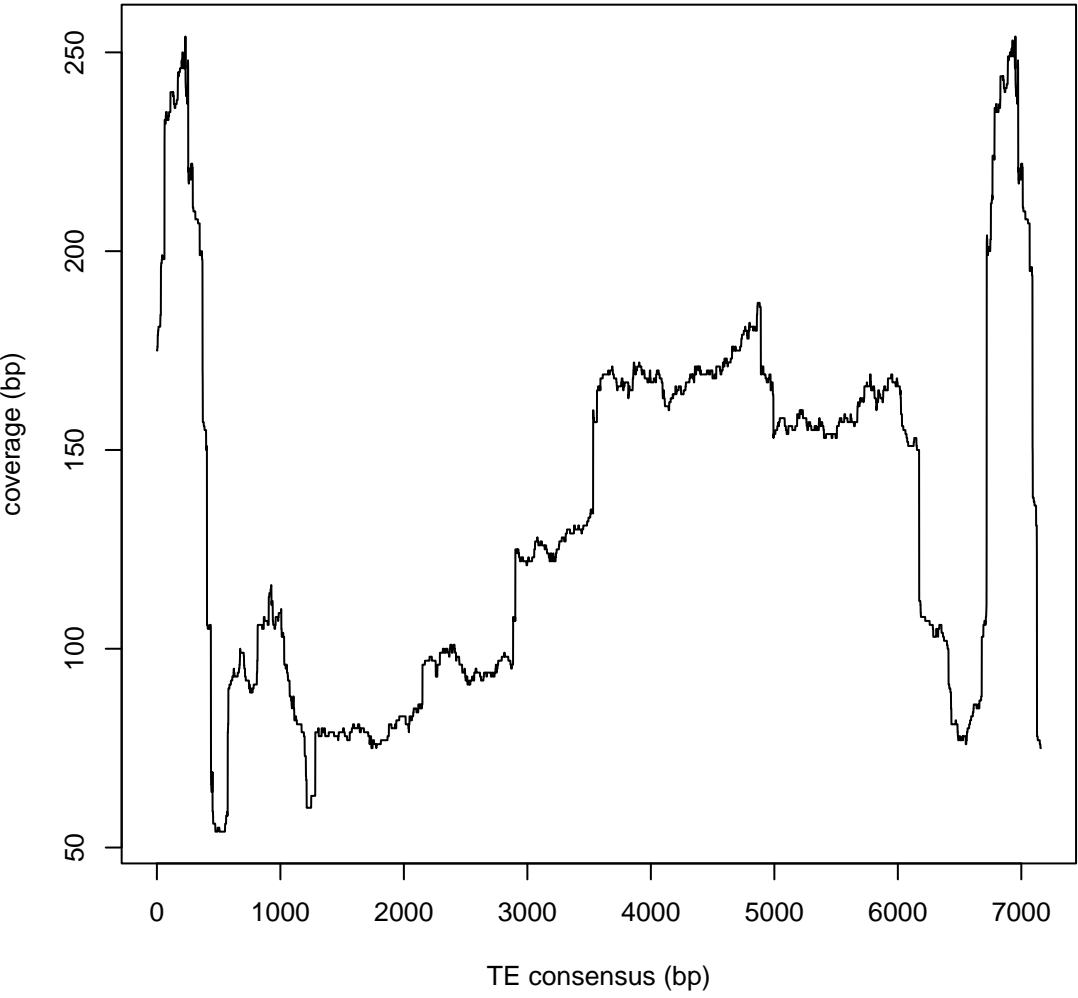


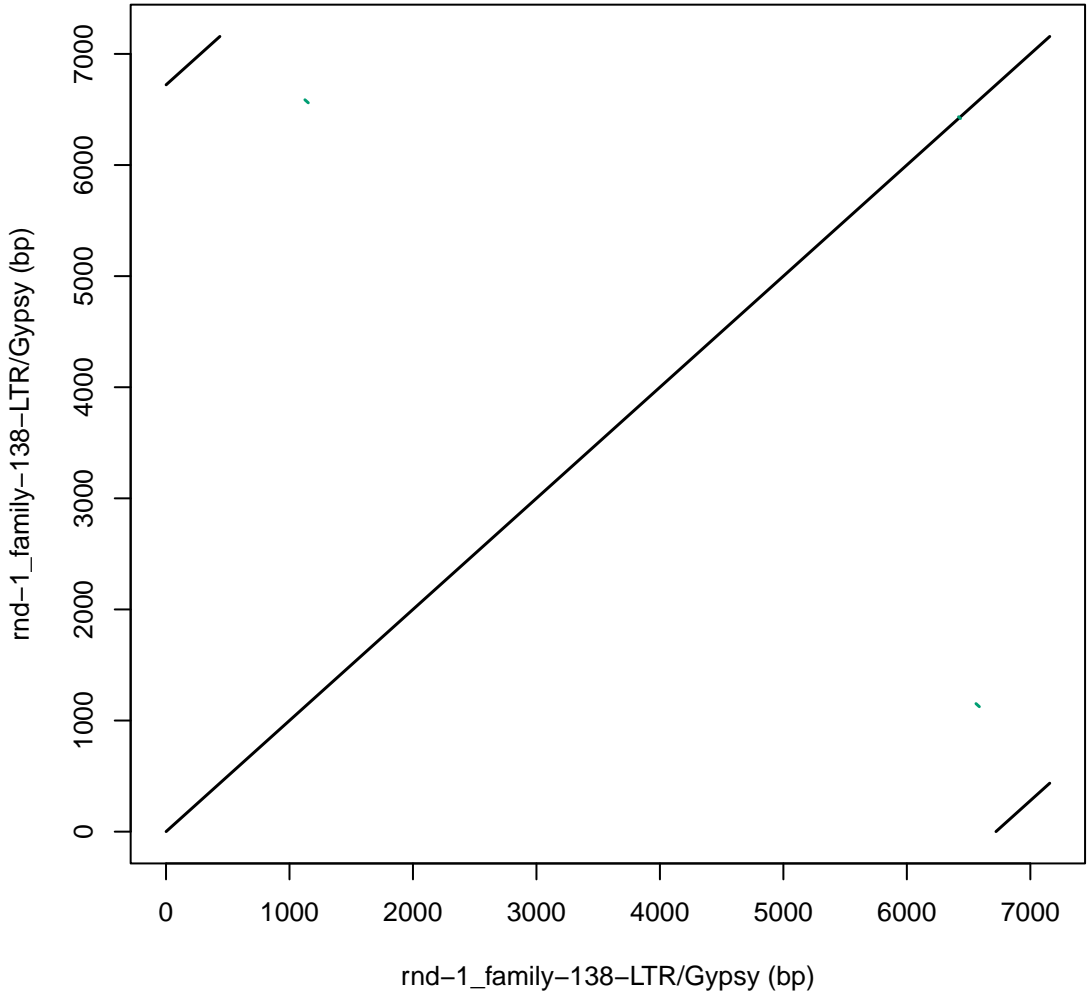
TE: rnd-1_family-138-LTR/Gypsy
consensus size: 7157bp; fragments: 1044; full length: 9 (>=6441.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

