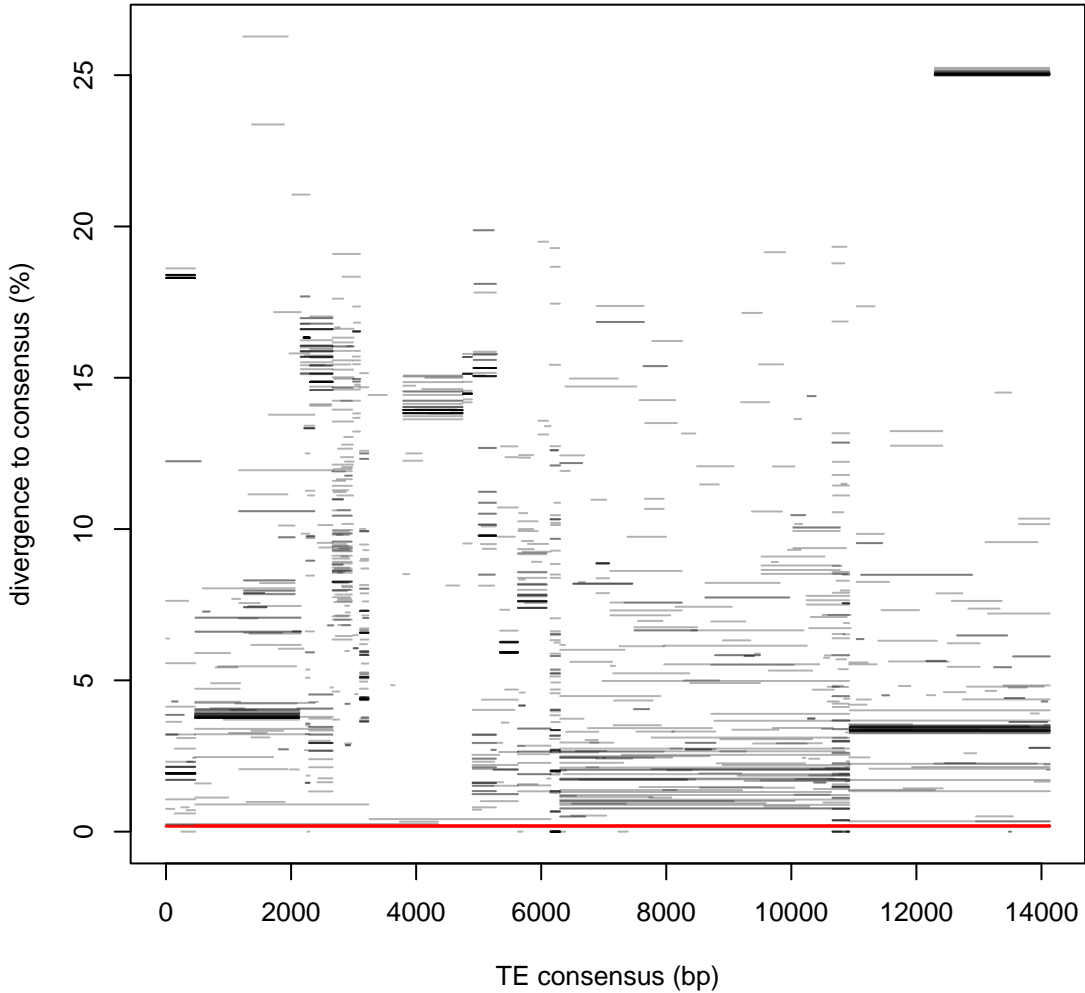
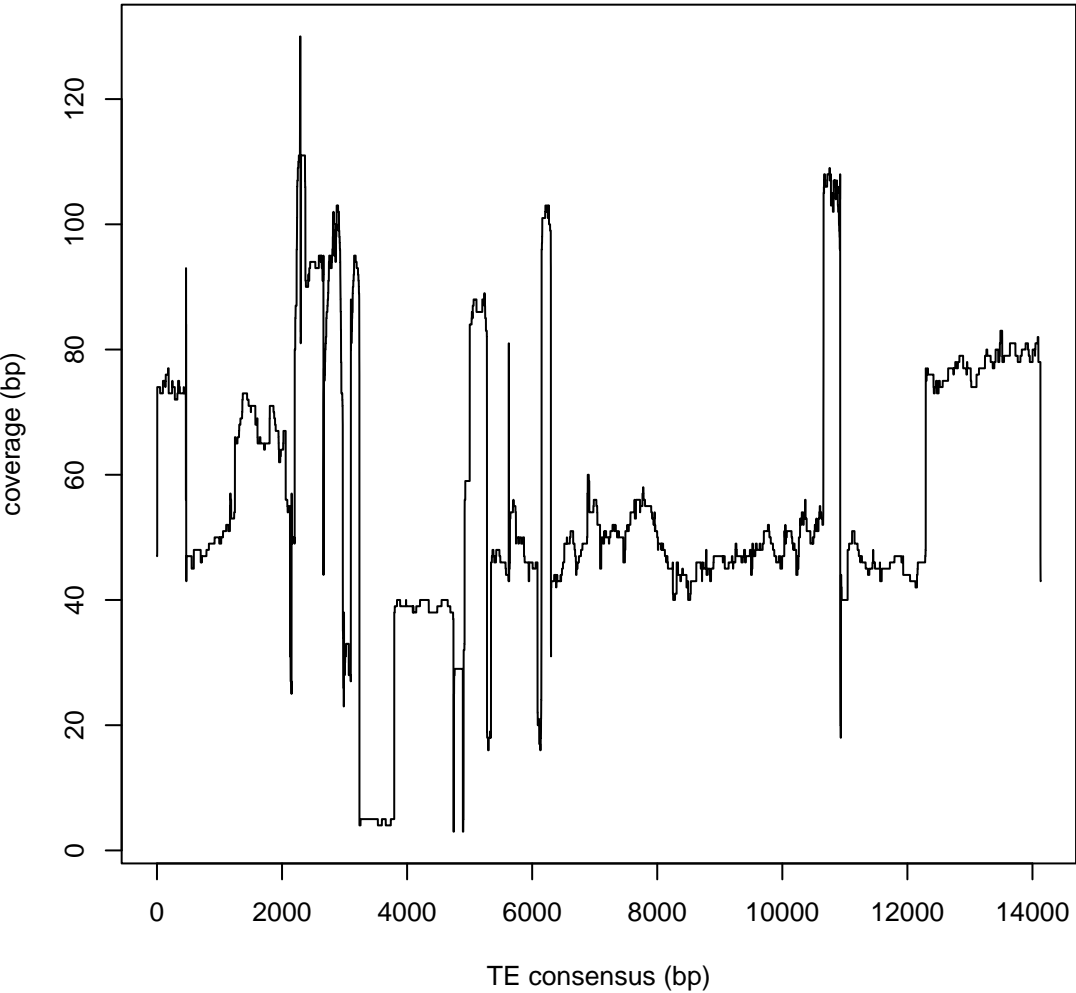


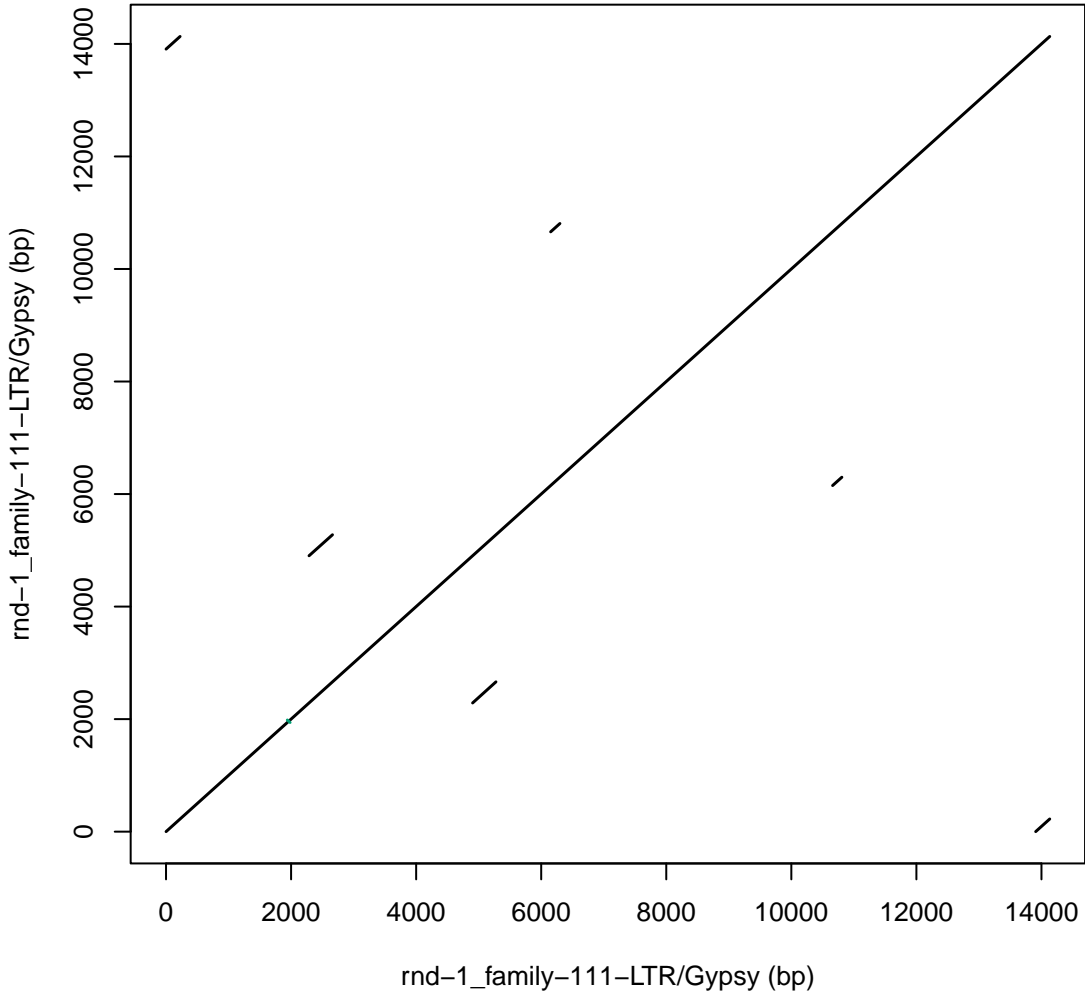
TE: rnd-1_family-111-LTR/Gypsy
consensus size: 14132bp; fragments: 1343; full length: 2 (>=12718.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

