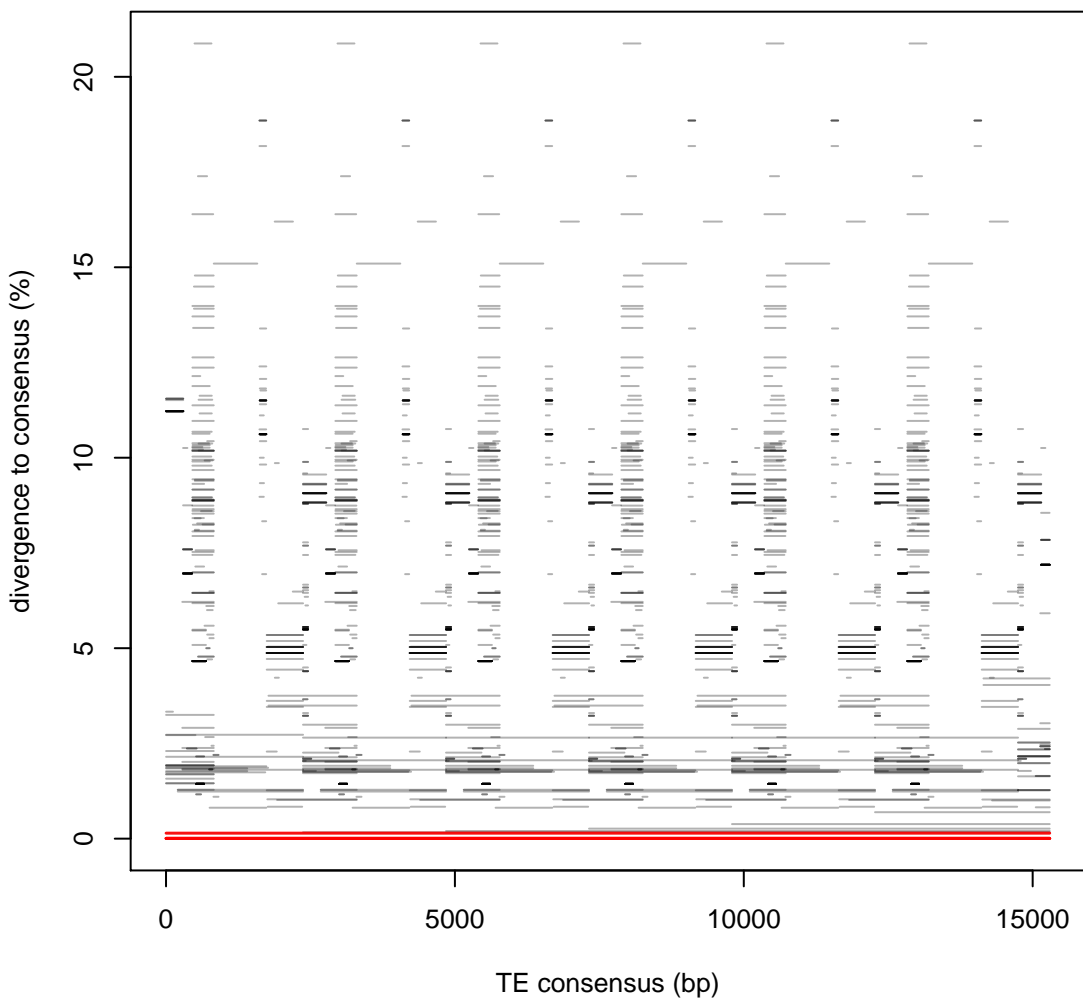
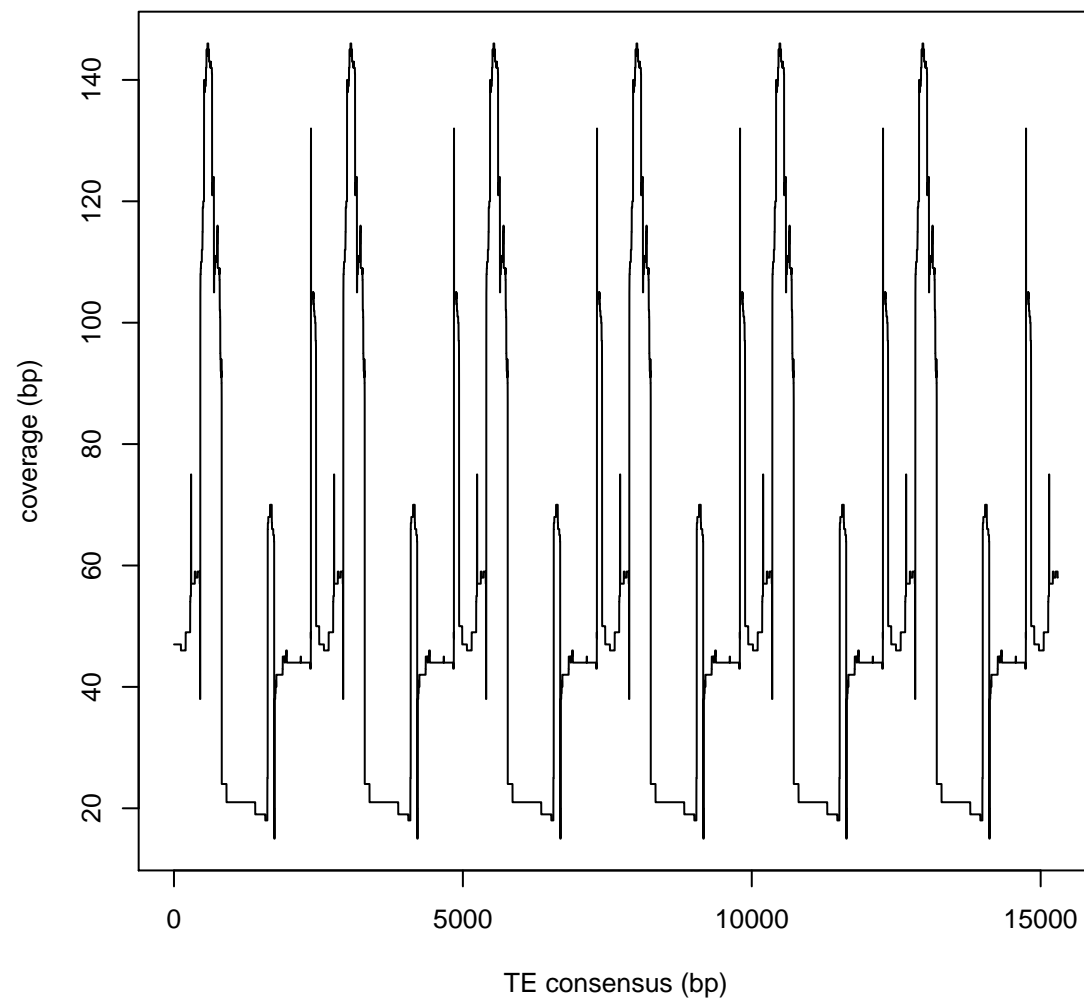


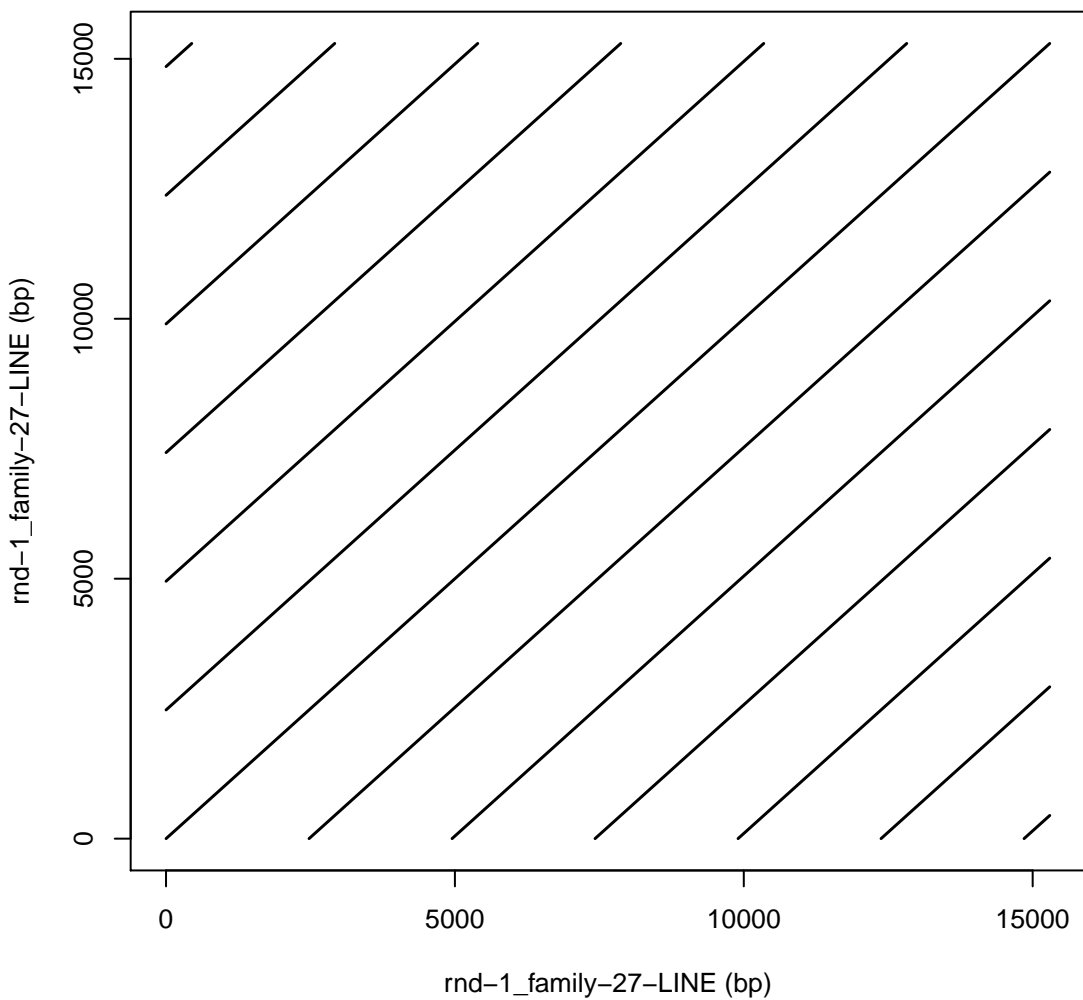
TE: rnd-1_family-27-LINE
consensus size: 15295bp; fragments: 2155; full length: 4 (≥ 13765.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

