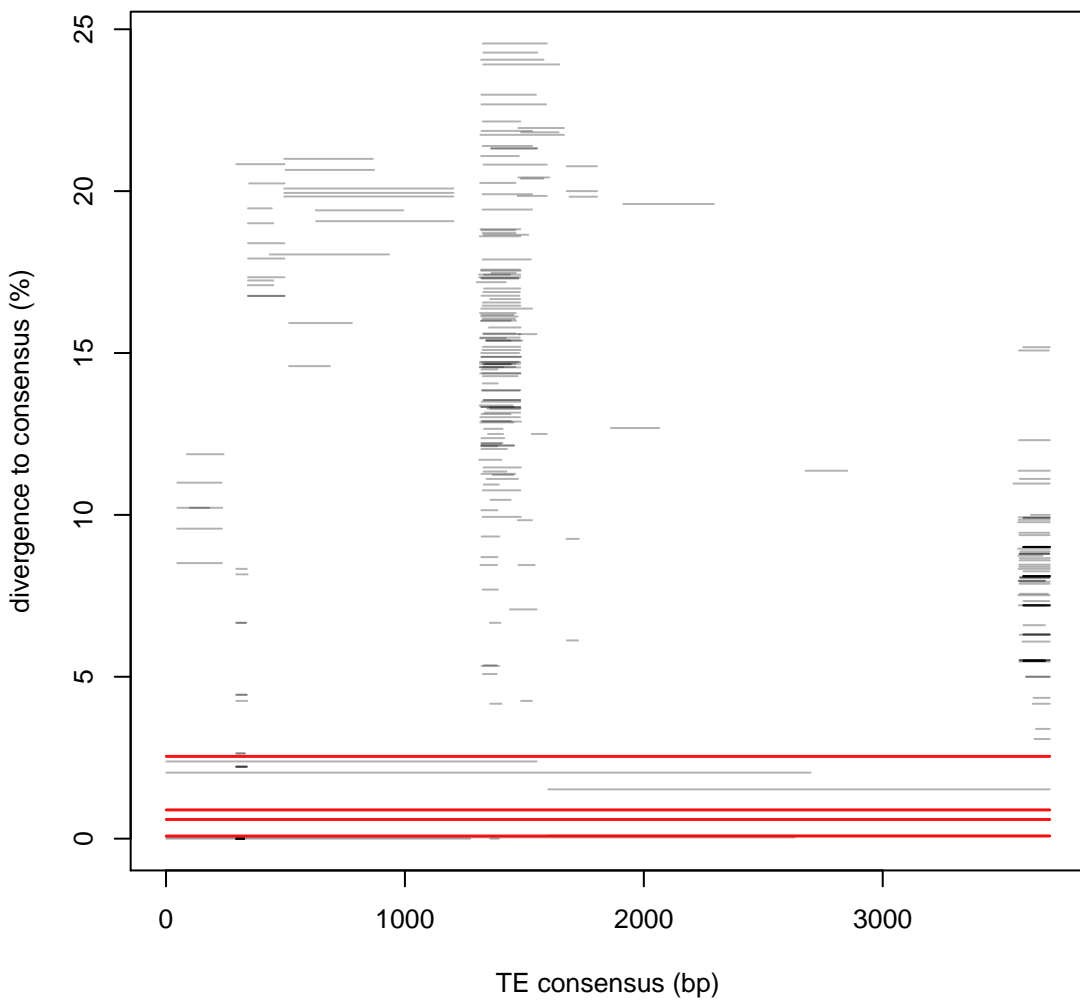
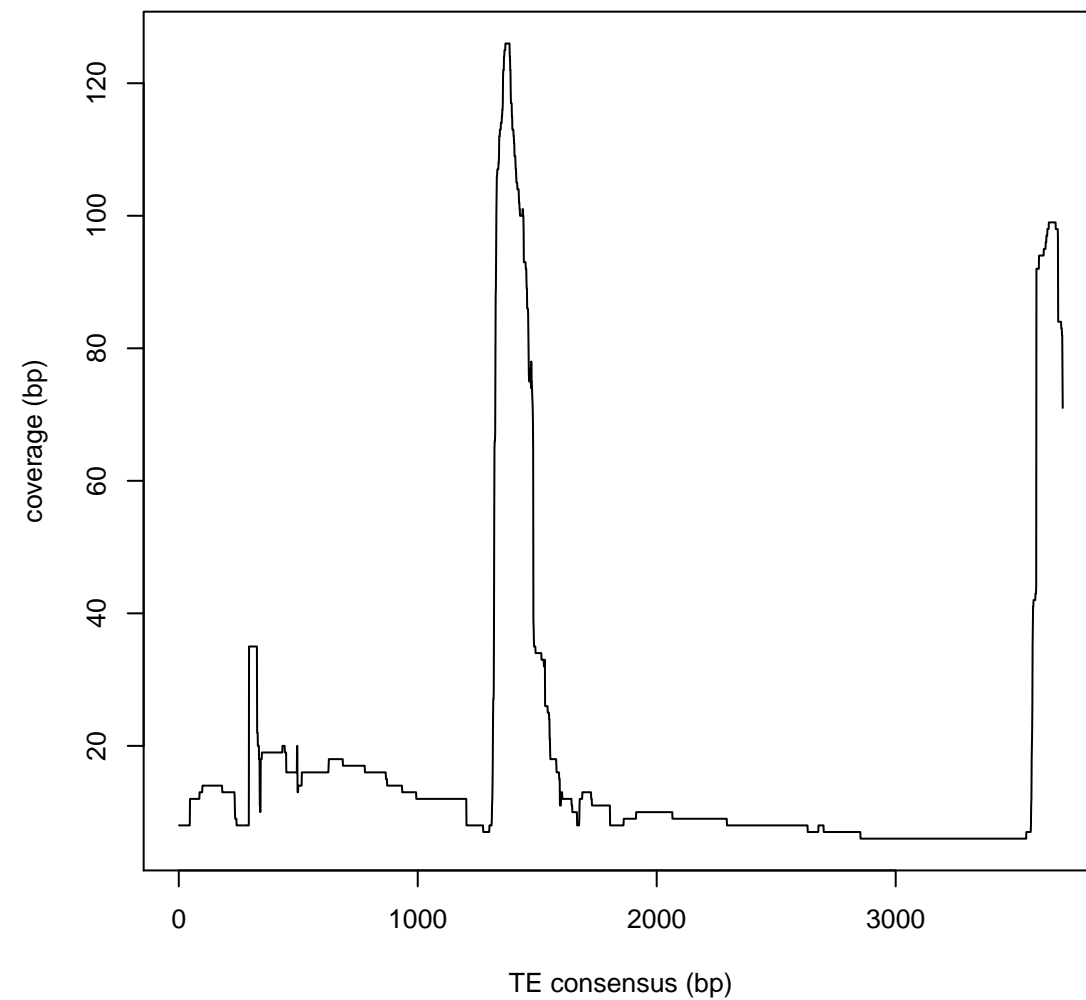


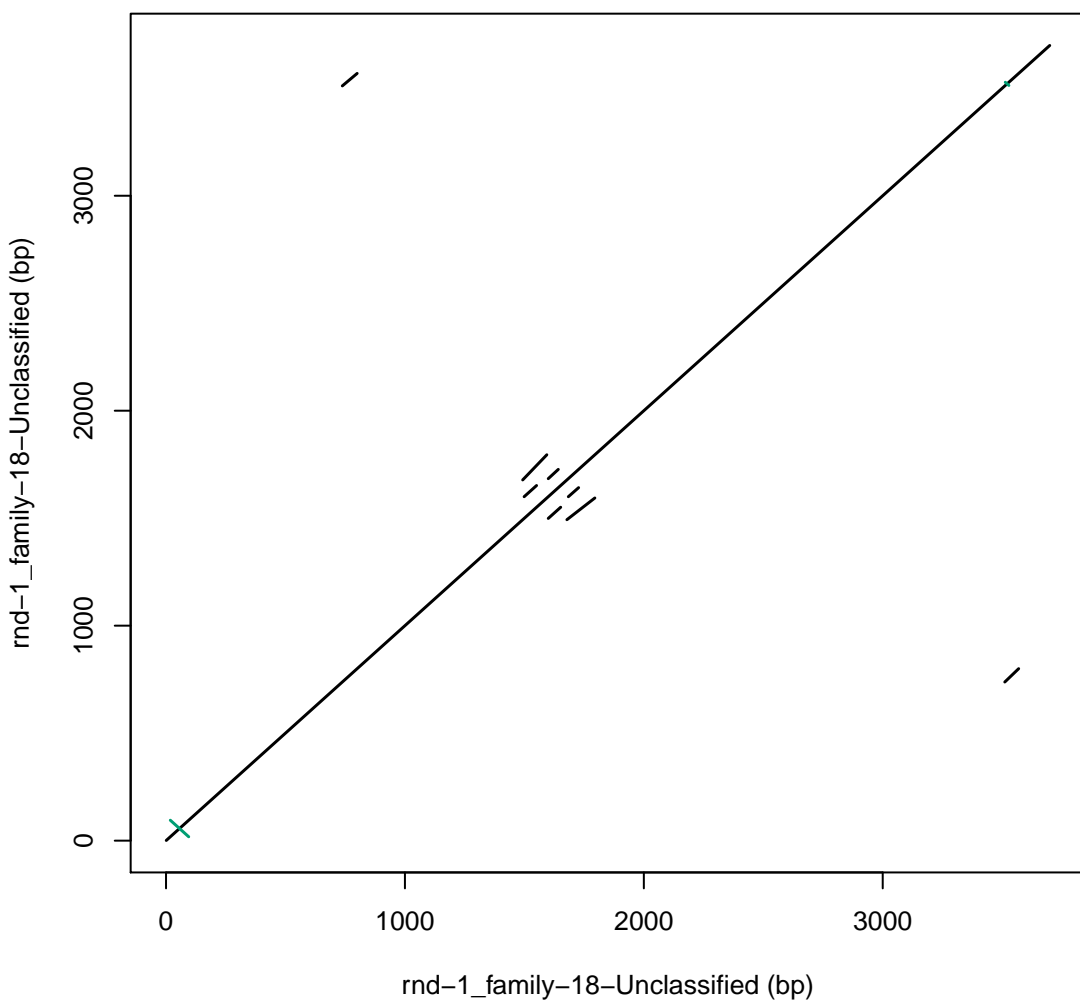
TE: rnd-1_family-18-Unclassified
consensus size: 3699bp; fragments: 294; full length: 4 (≥ 3329.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

