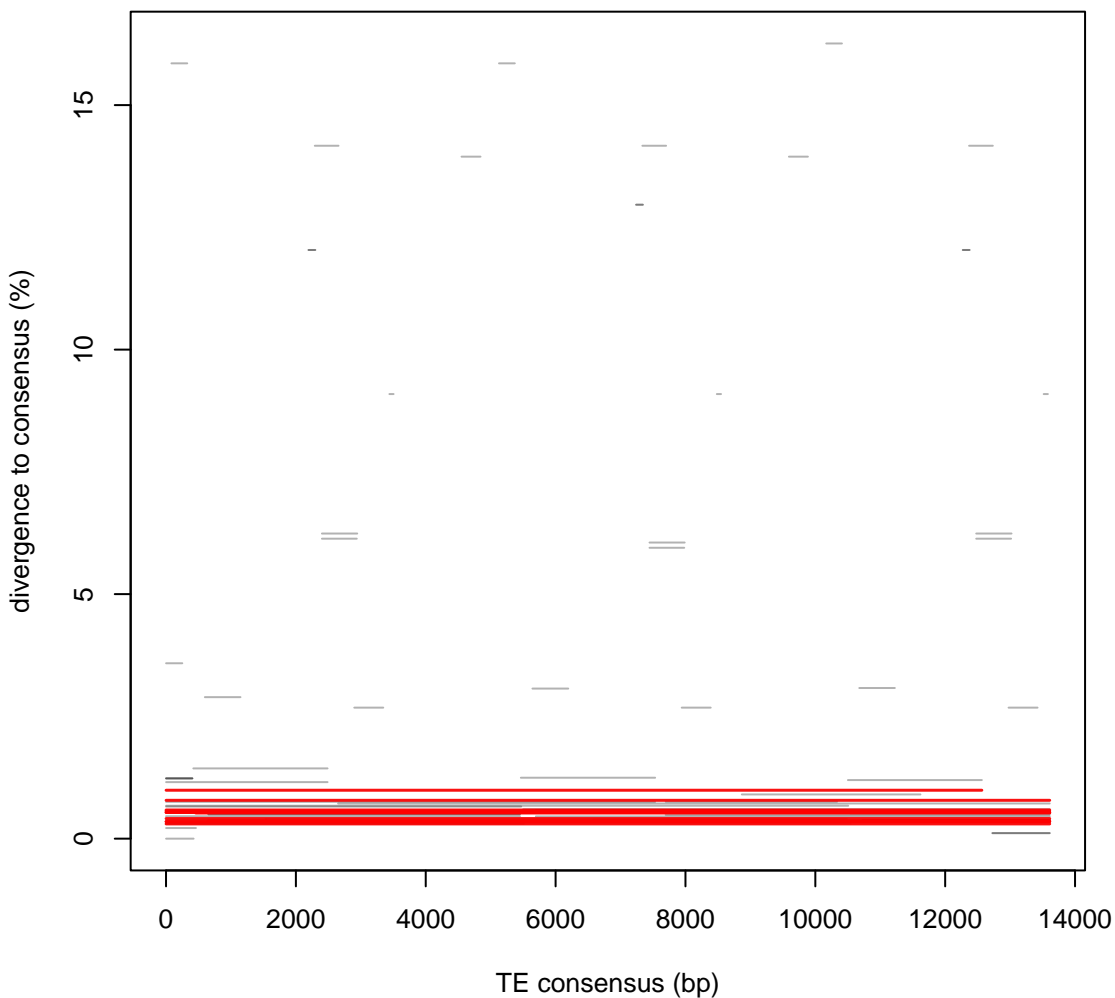
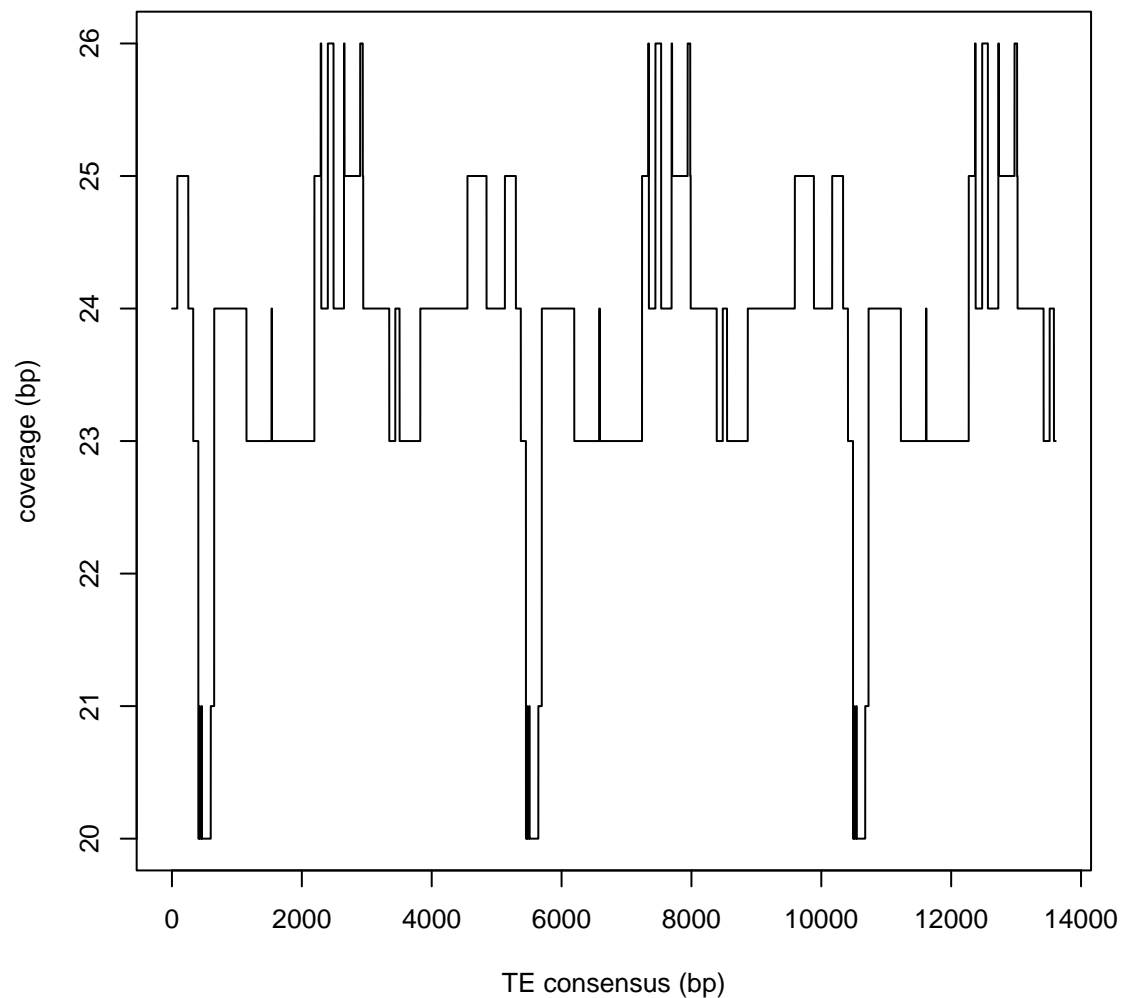


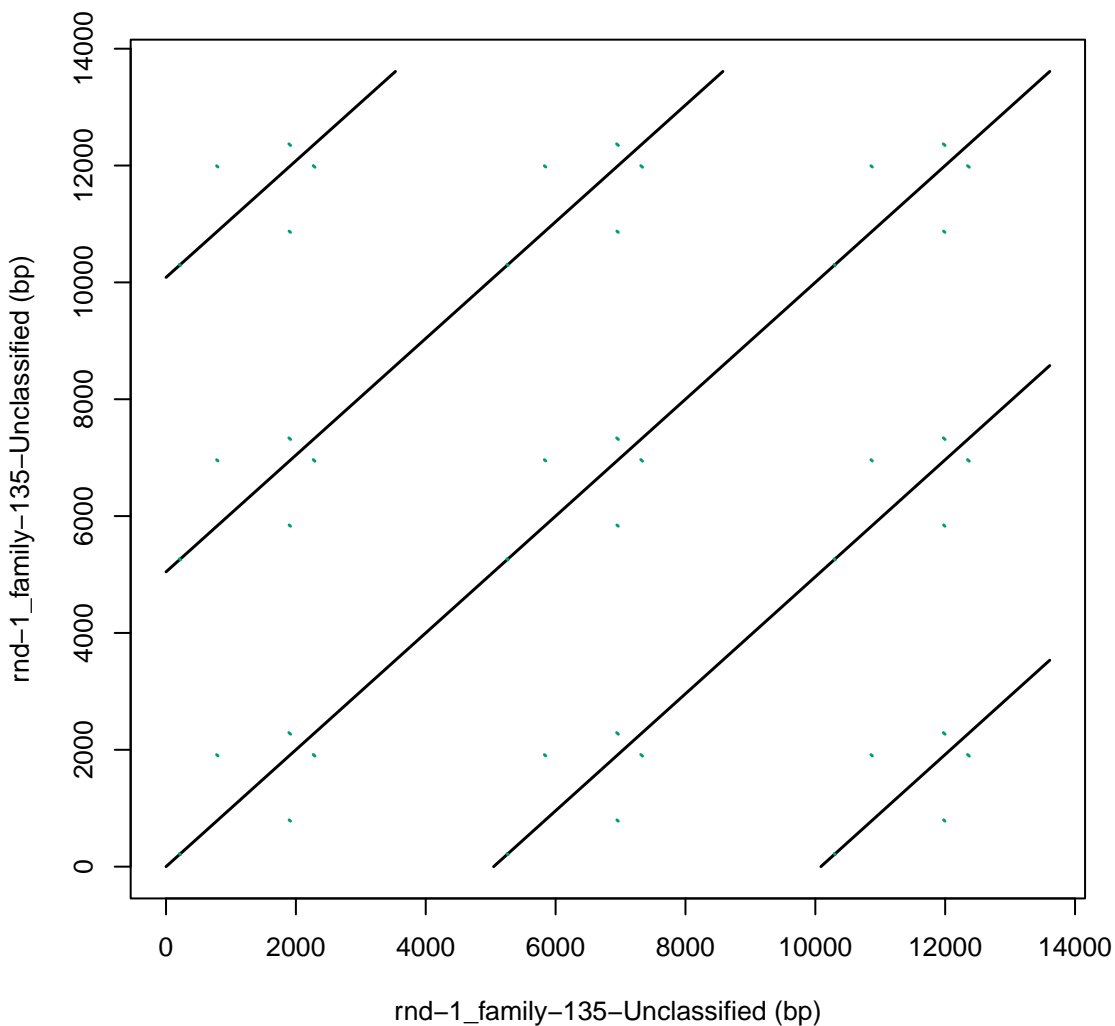
TE: rnd-1_family-135-Unclassified
consensus size: 13610bp; fragments: 79; full length: 12 (≥ 12249 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

