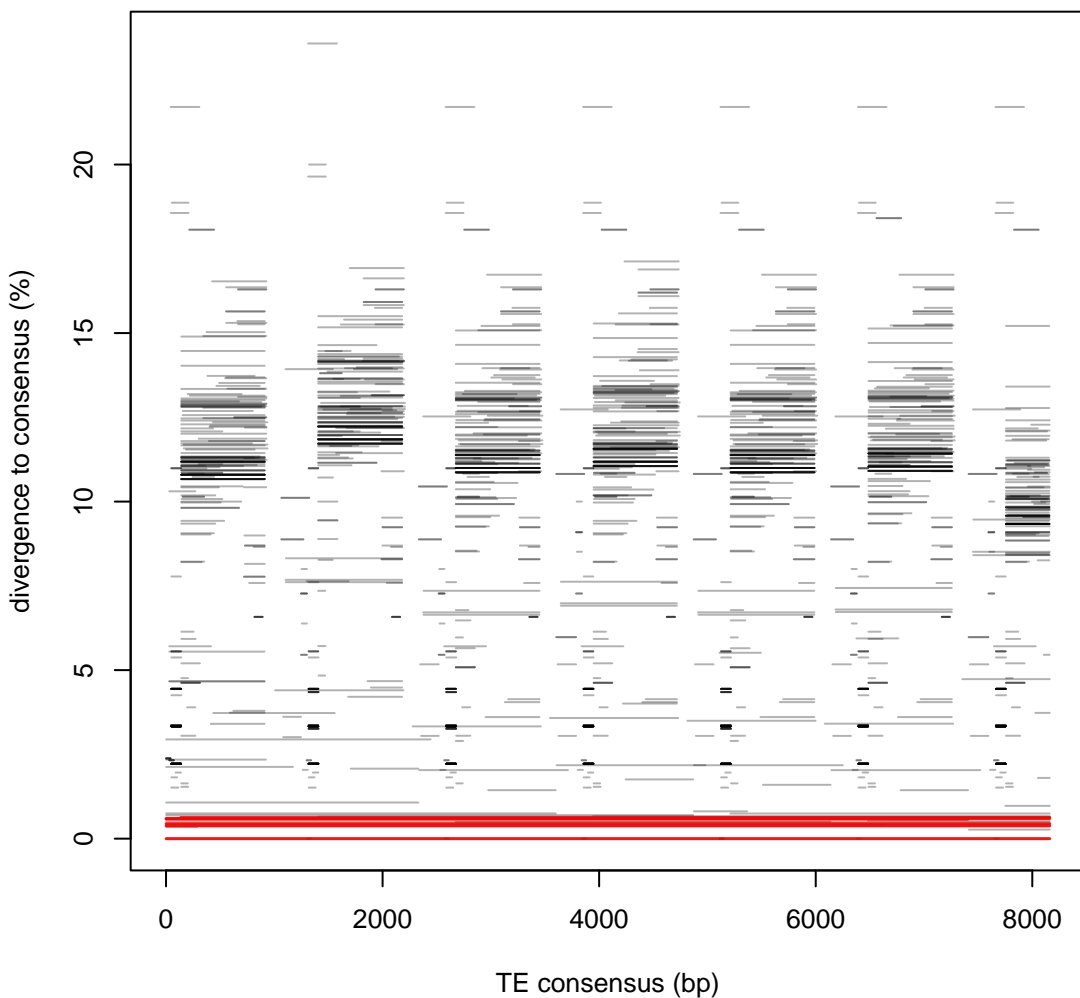
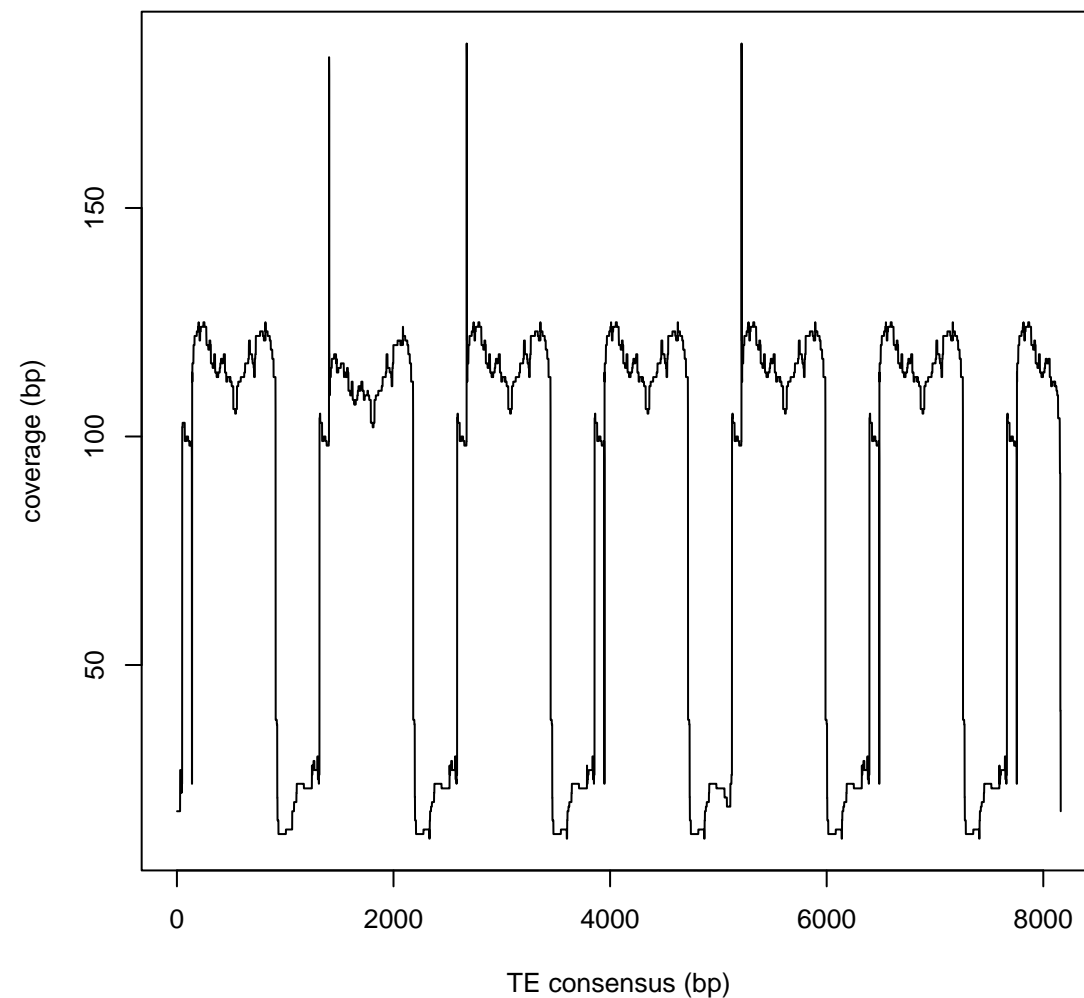


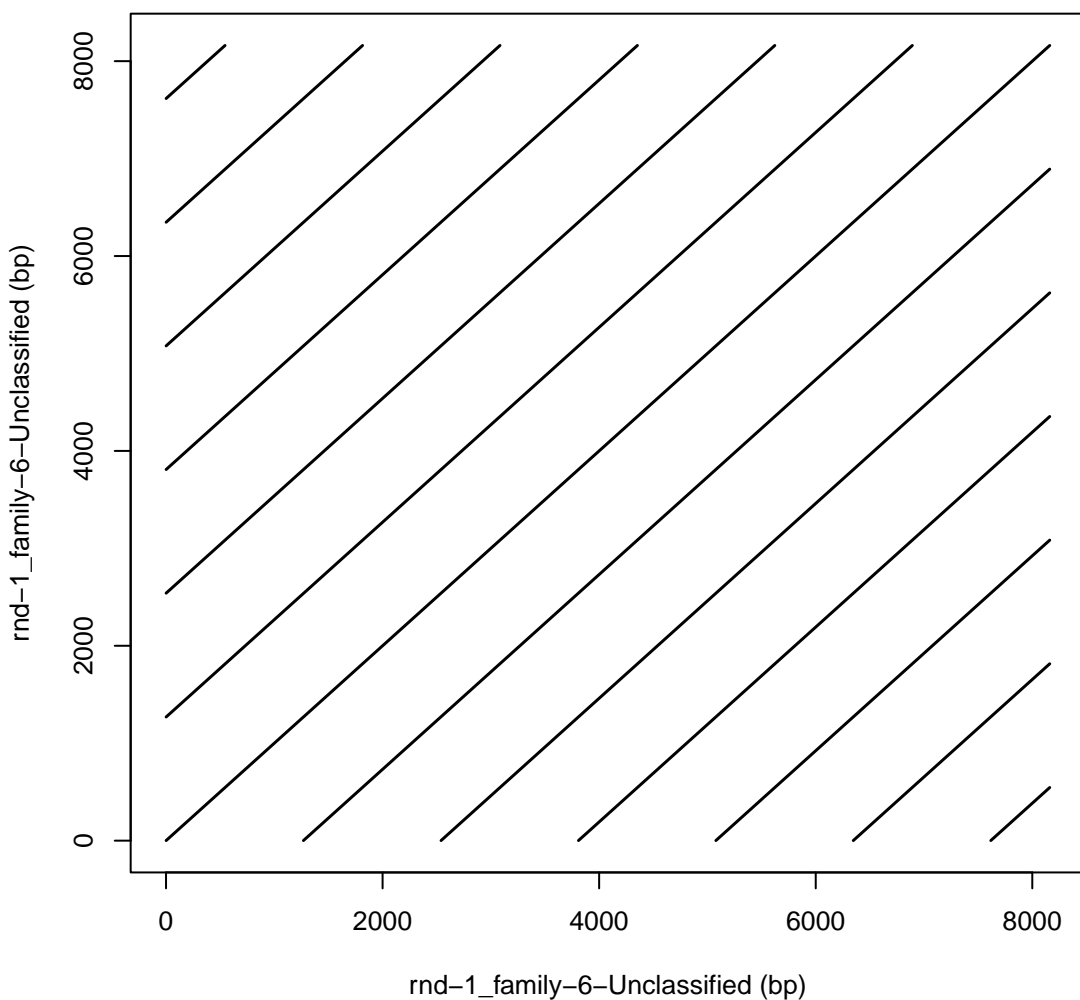
TE: rnd-1_family-6-Unclassified
consensus size: 8161bp; fragments: 1799; full length: 6 (≥ 7344.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

