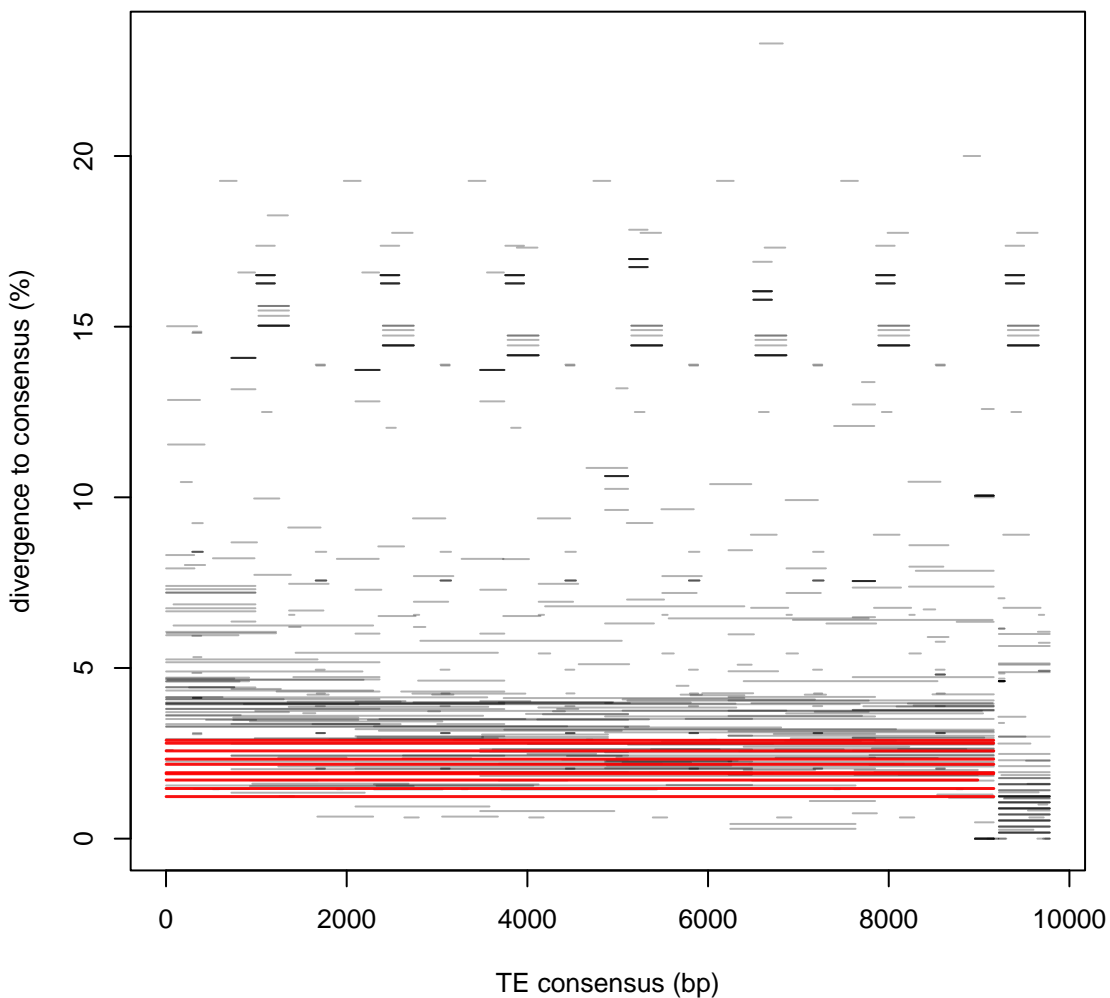
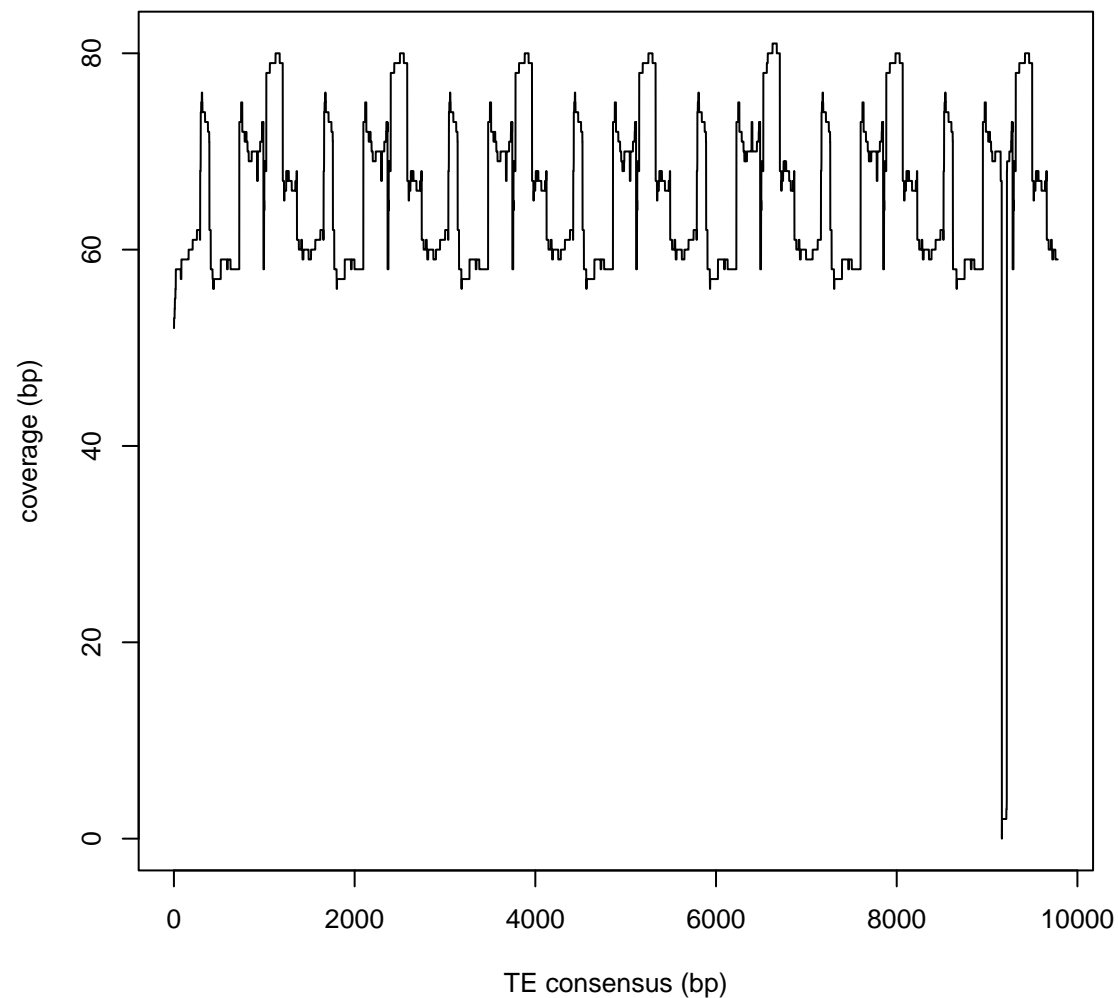


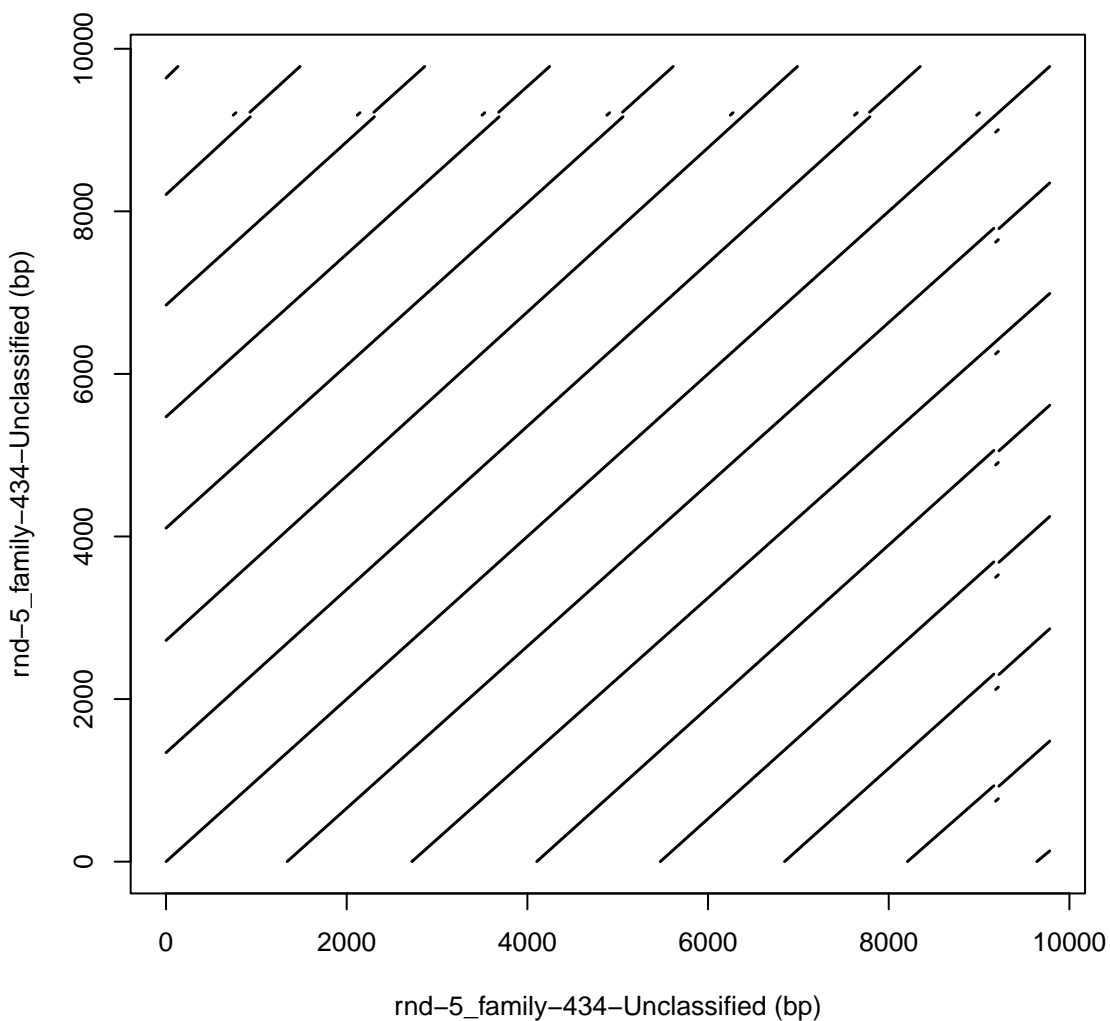
TE: rnd-5\_family-434-Unclassified  
consensus size: 9782bp; fragments: 730; full length: 10 ( $\geq 8803.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

