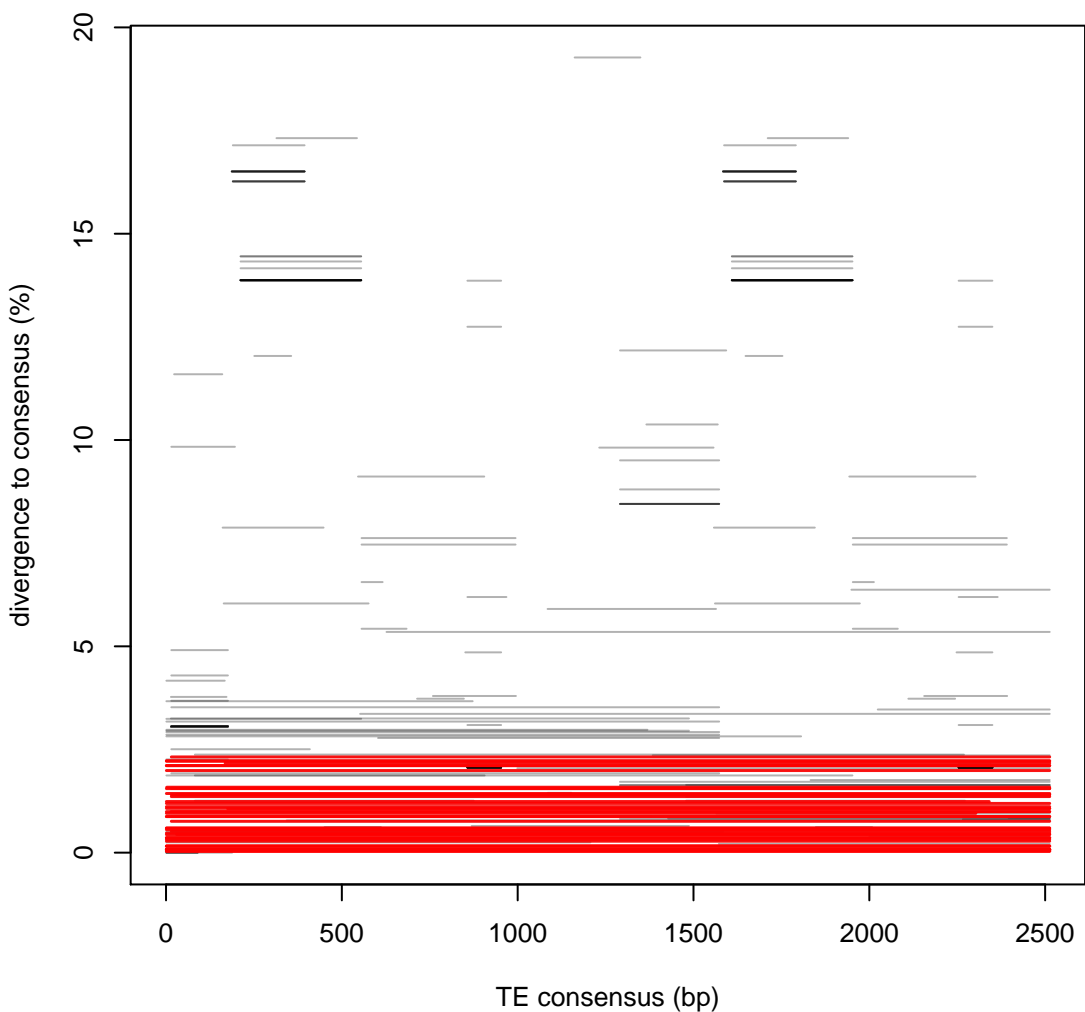
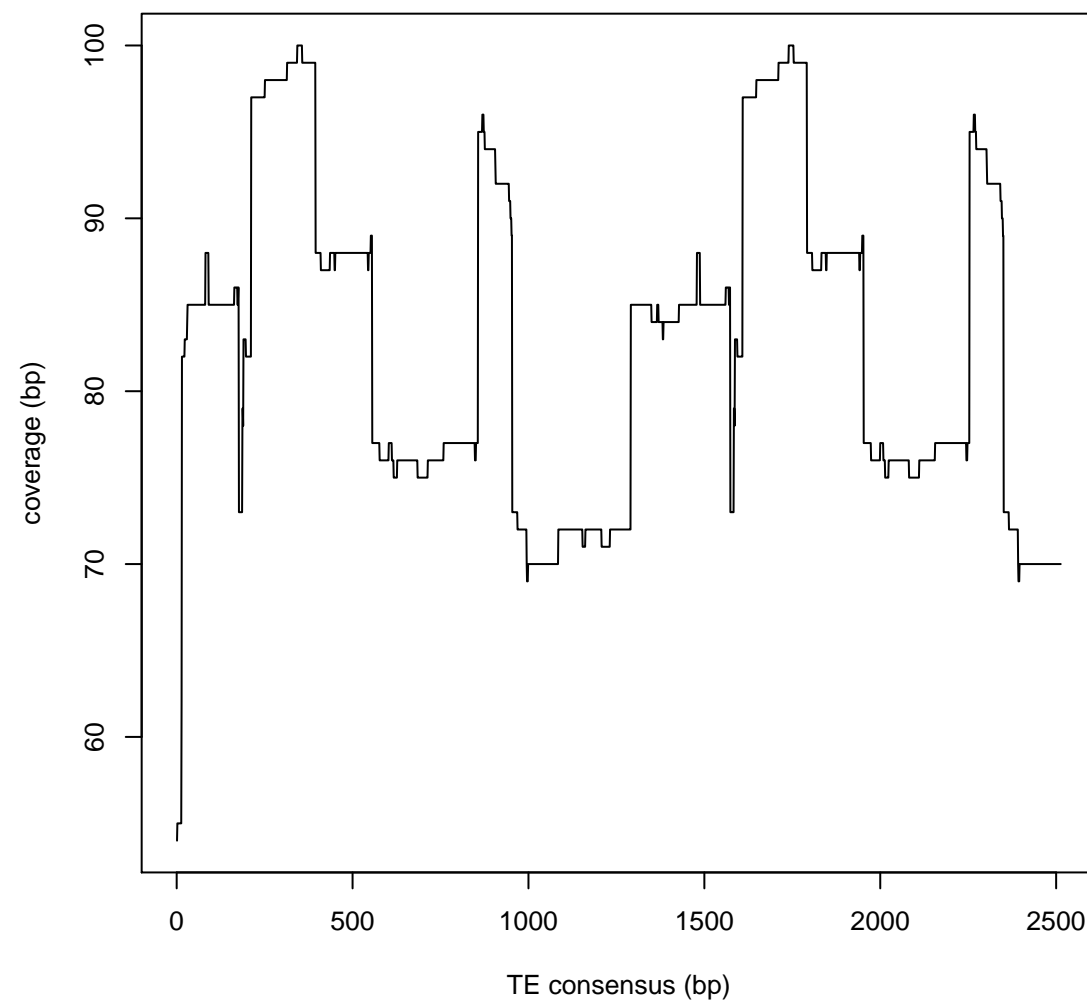


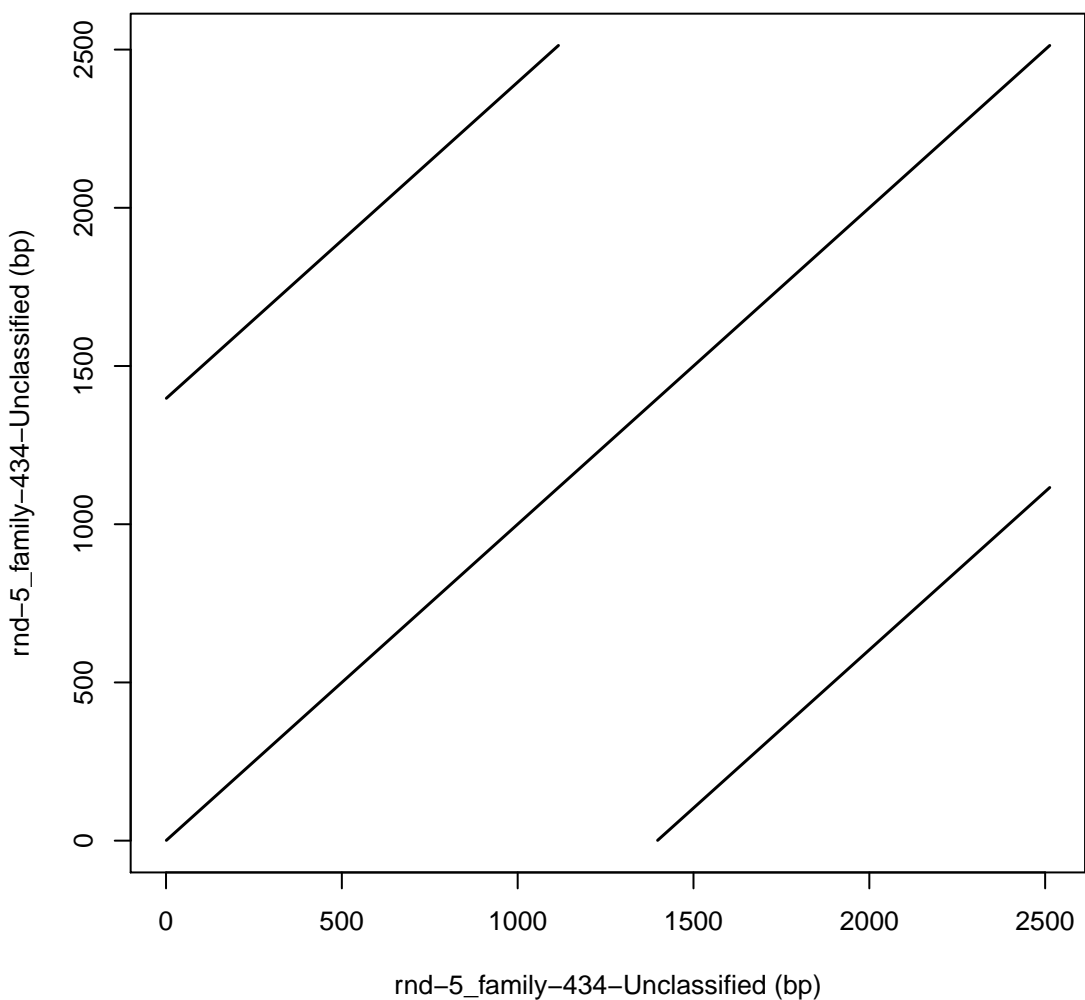
TE: rnd-5_family-434-Unclassified
consensus size: 2513bp; fragments: 243; full length: 49 (>=2261.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

