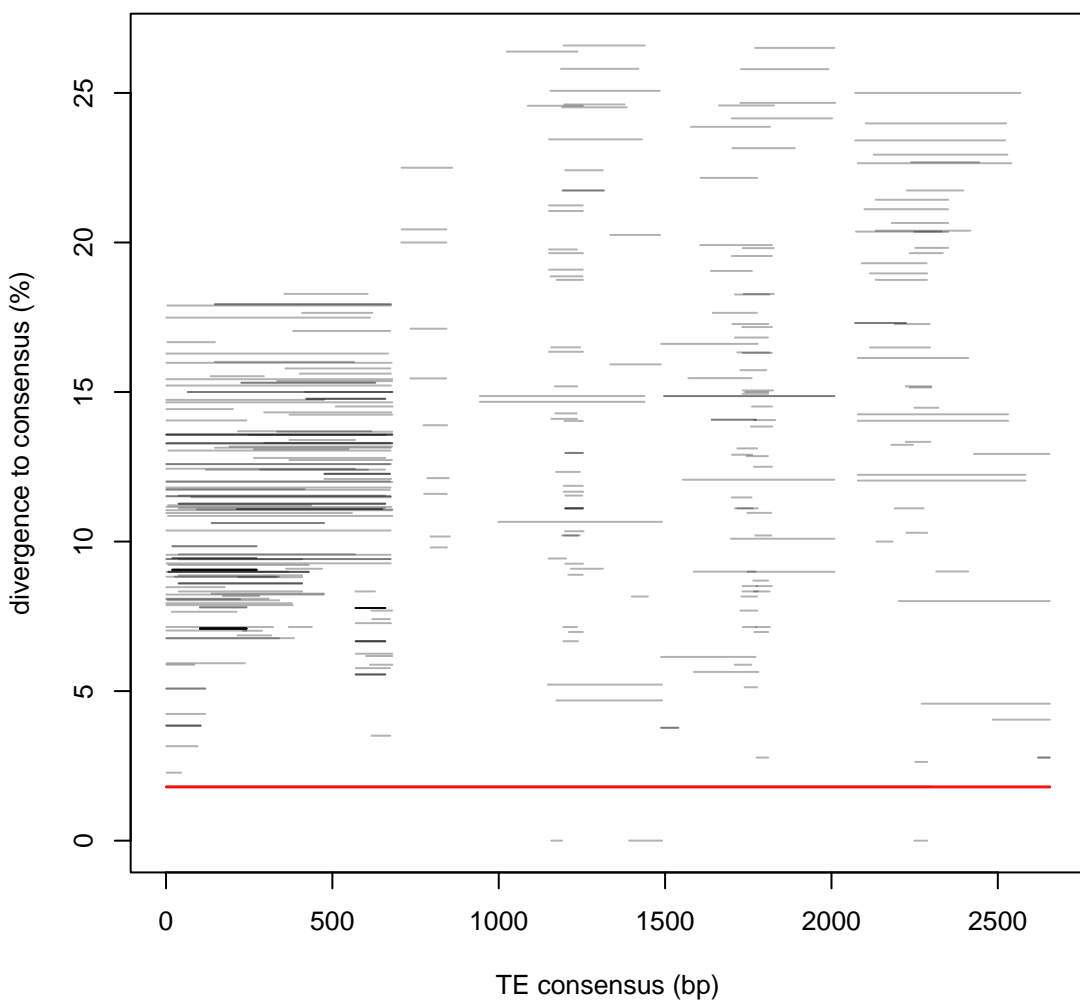
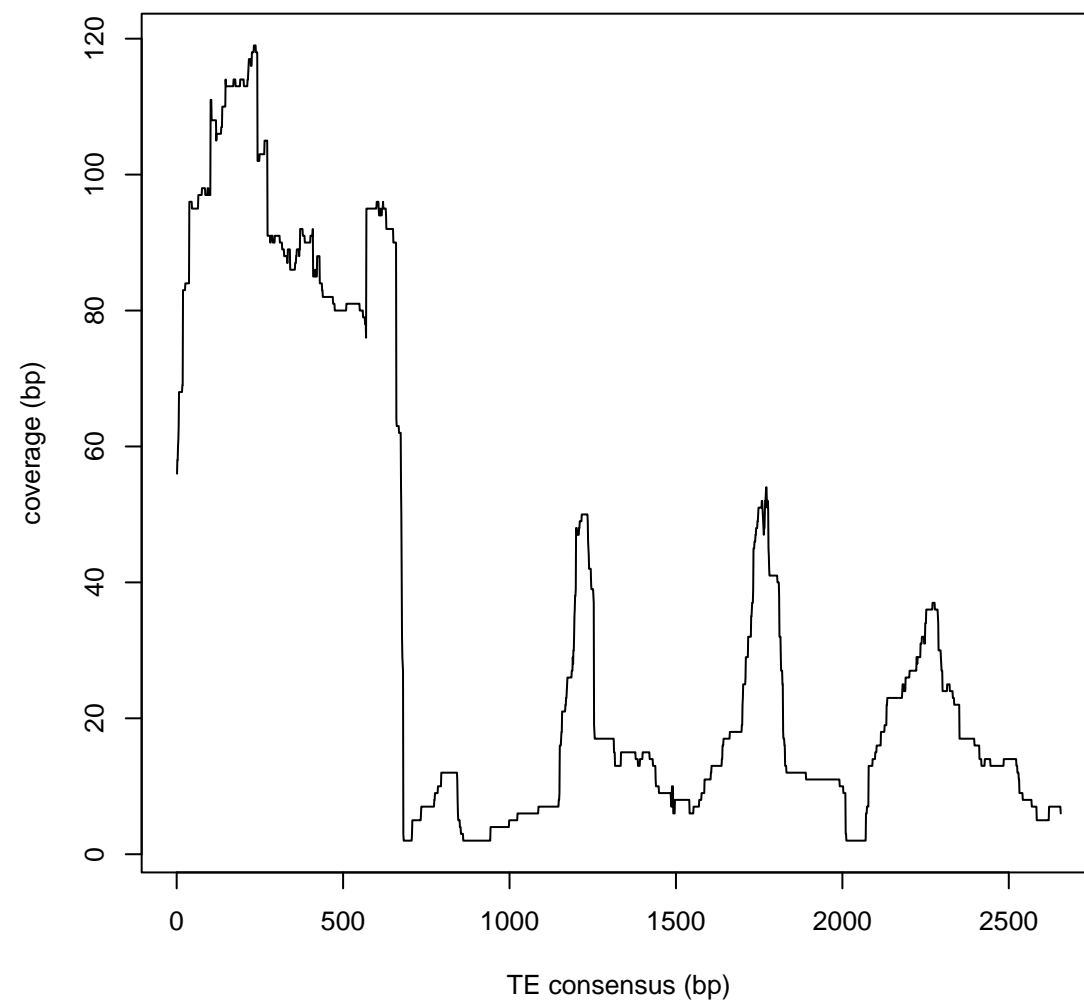


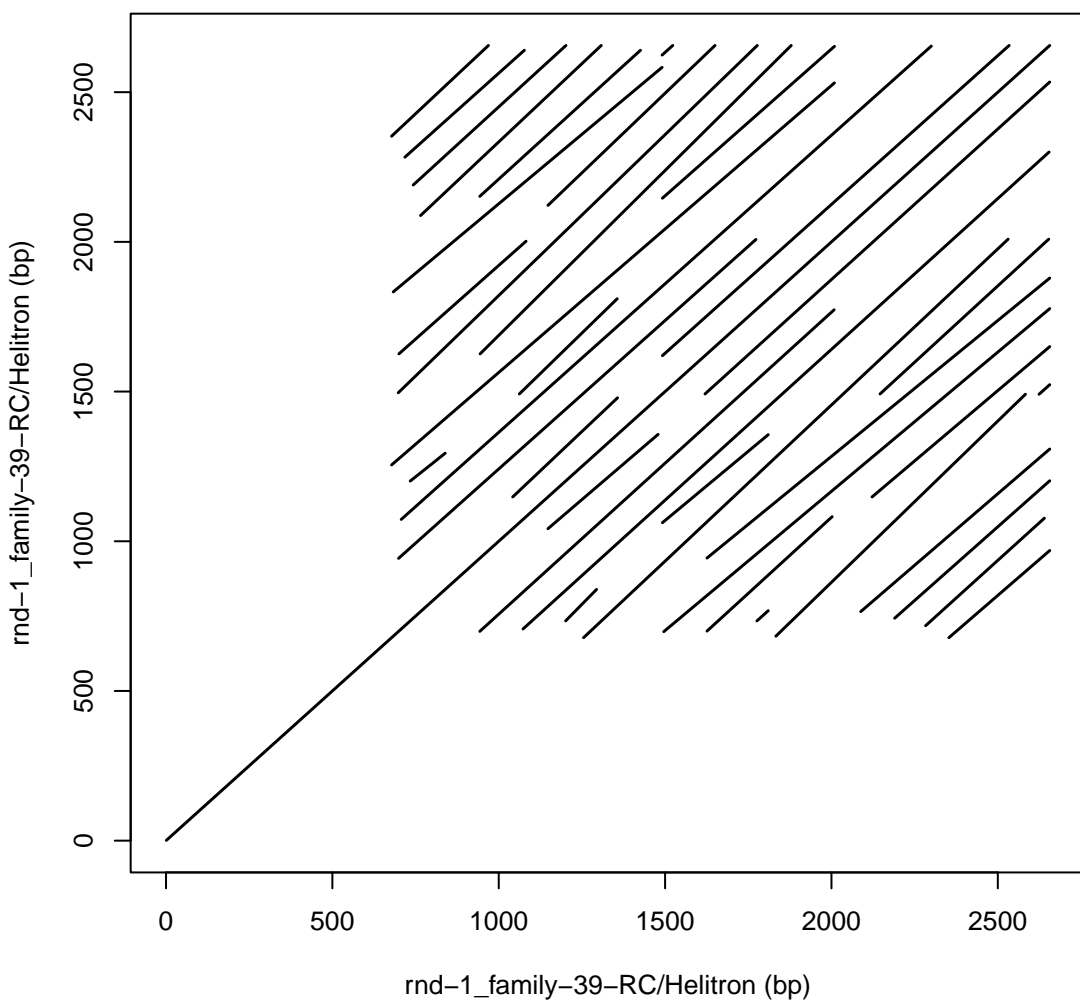
TE: rnd-1_family-39-RC/Helitron
consensus size: 2656bp; fragments: 356; full length: 1 (≥ 2390.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

