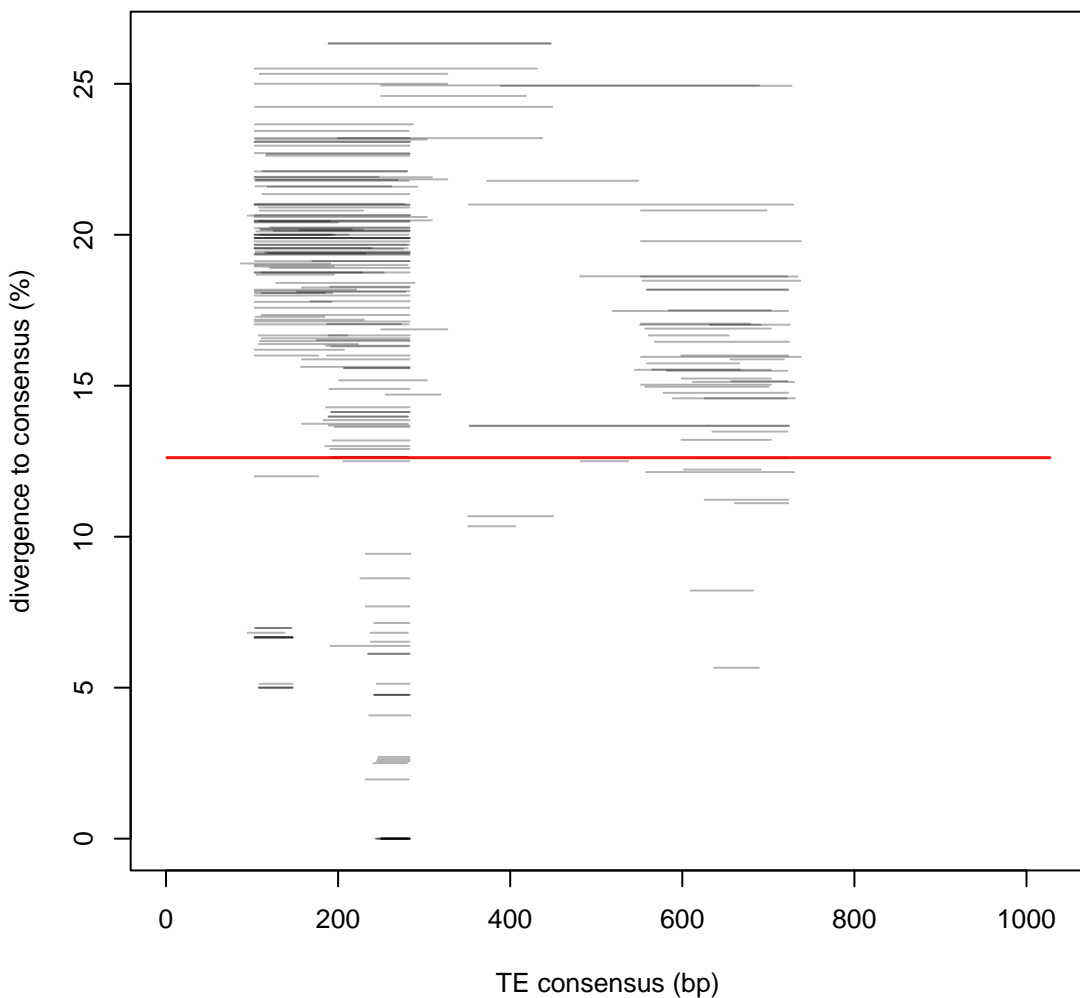
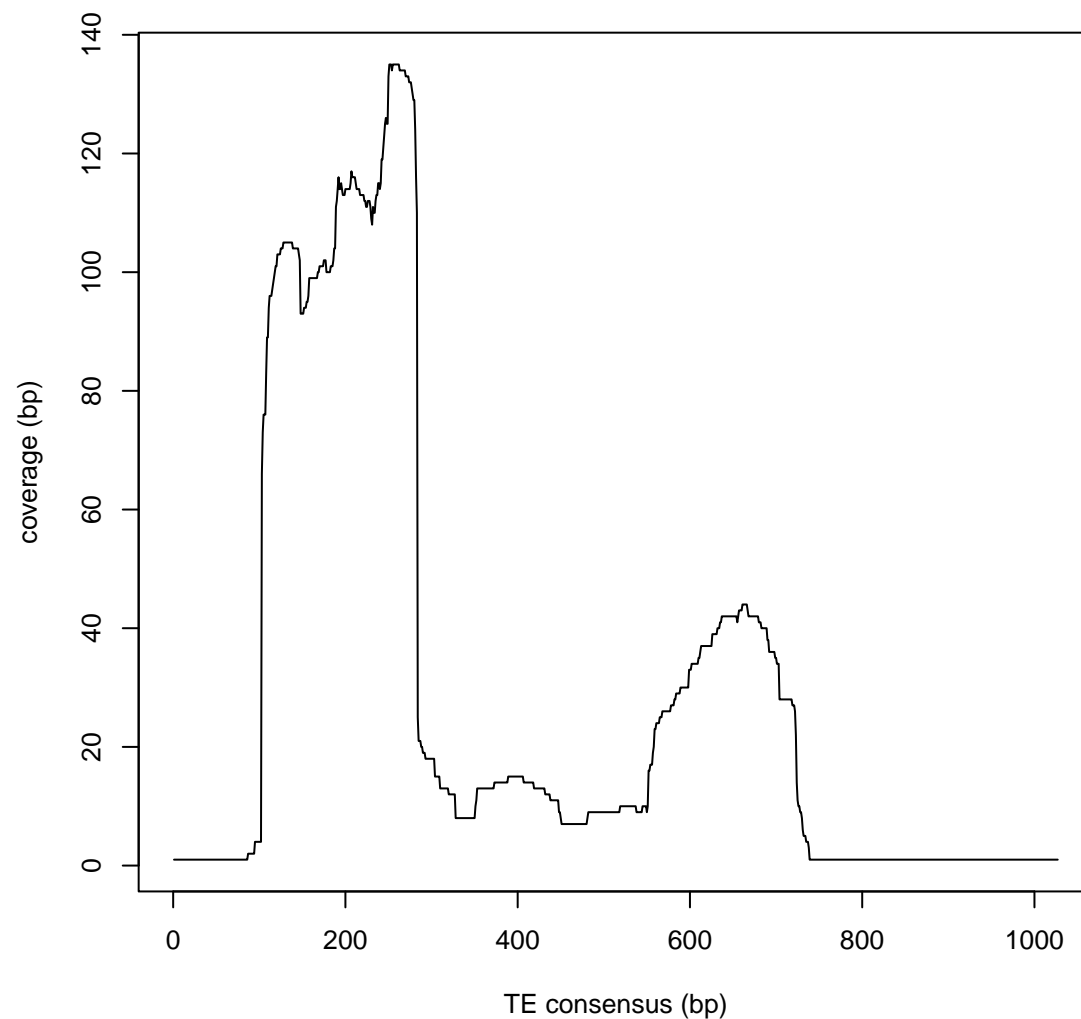


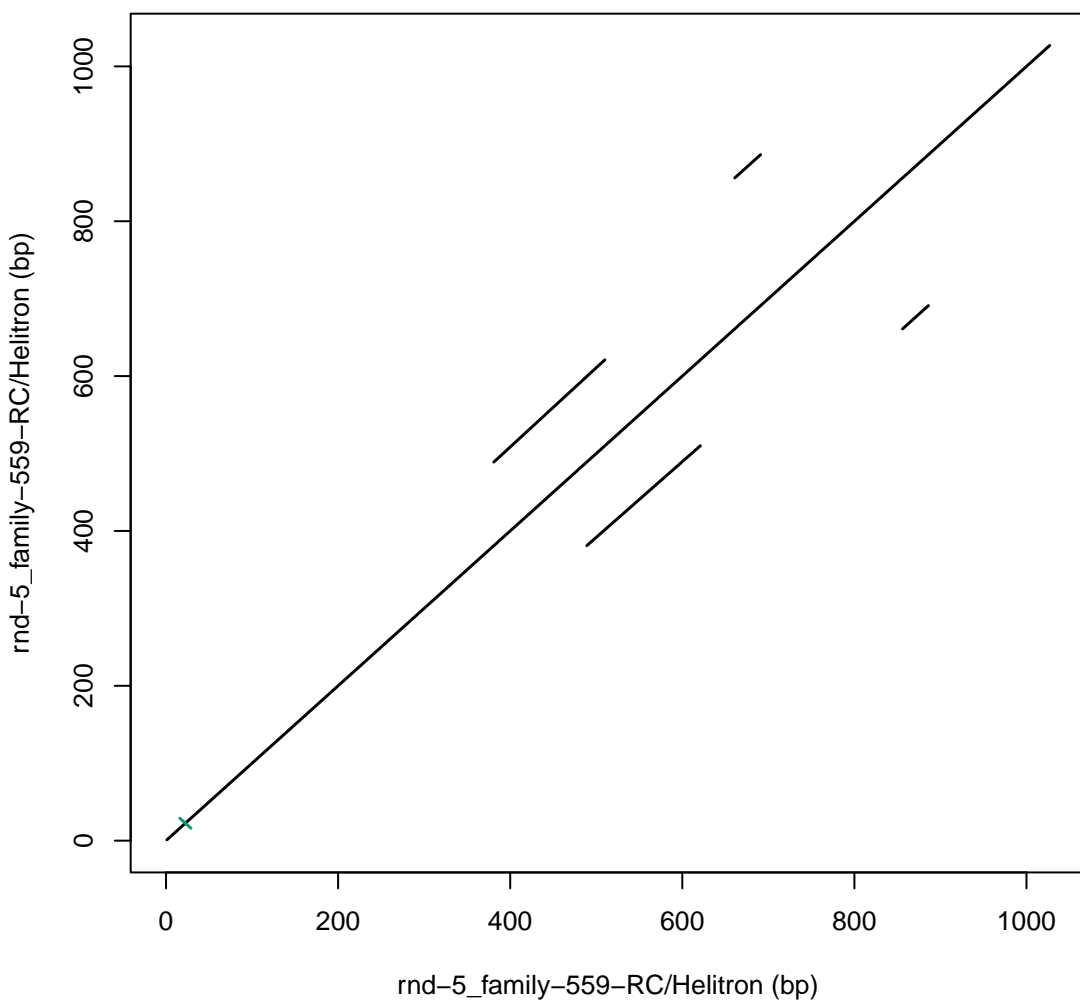
TE: rnd-5_family-559-RC/Helitron
consensus size: 1027bp; fragments: 221; full length: 1 (>=924.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

