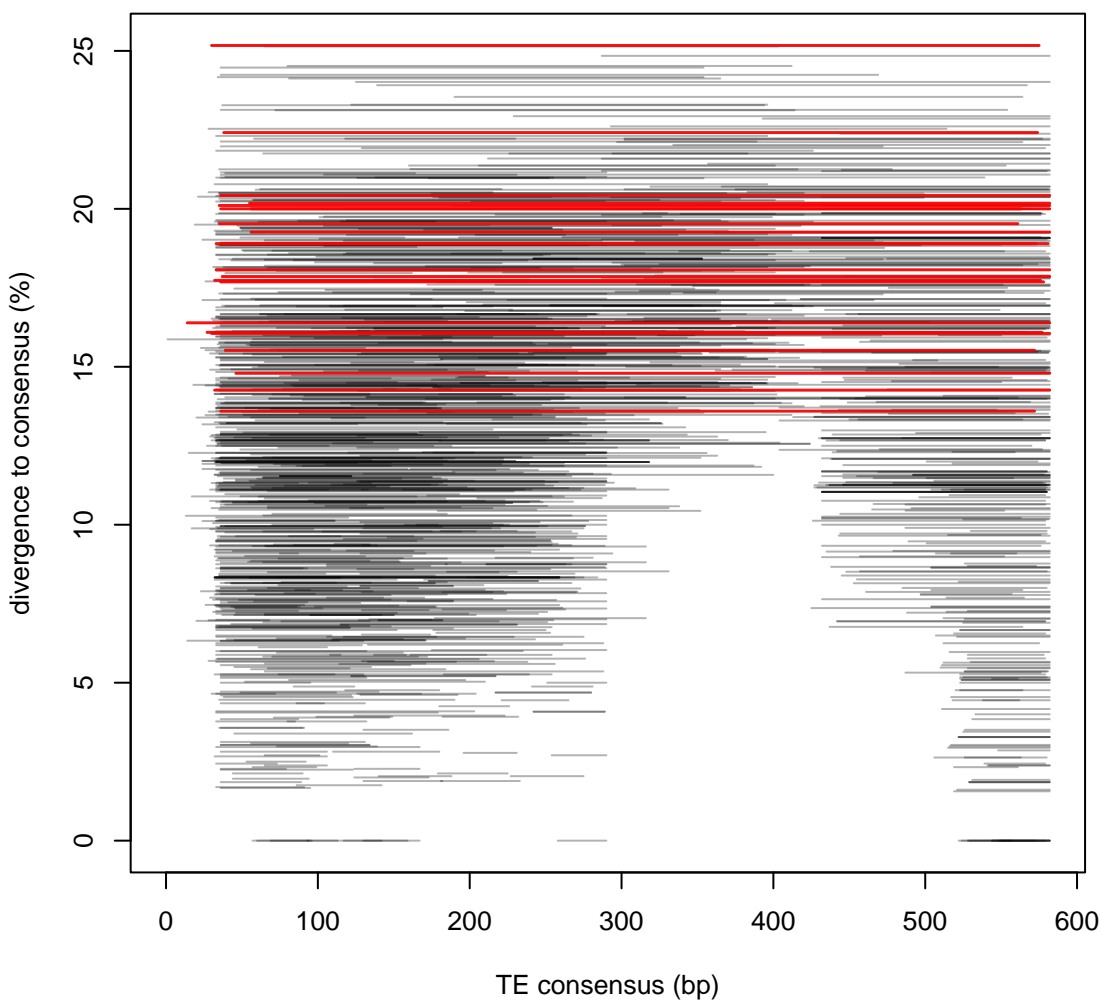
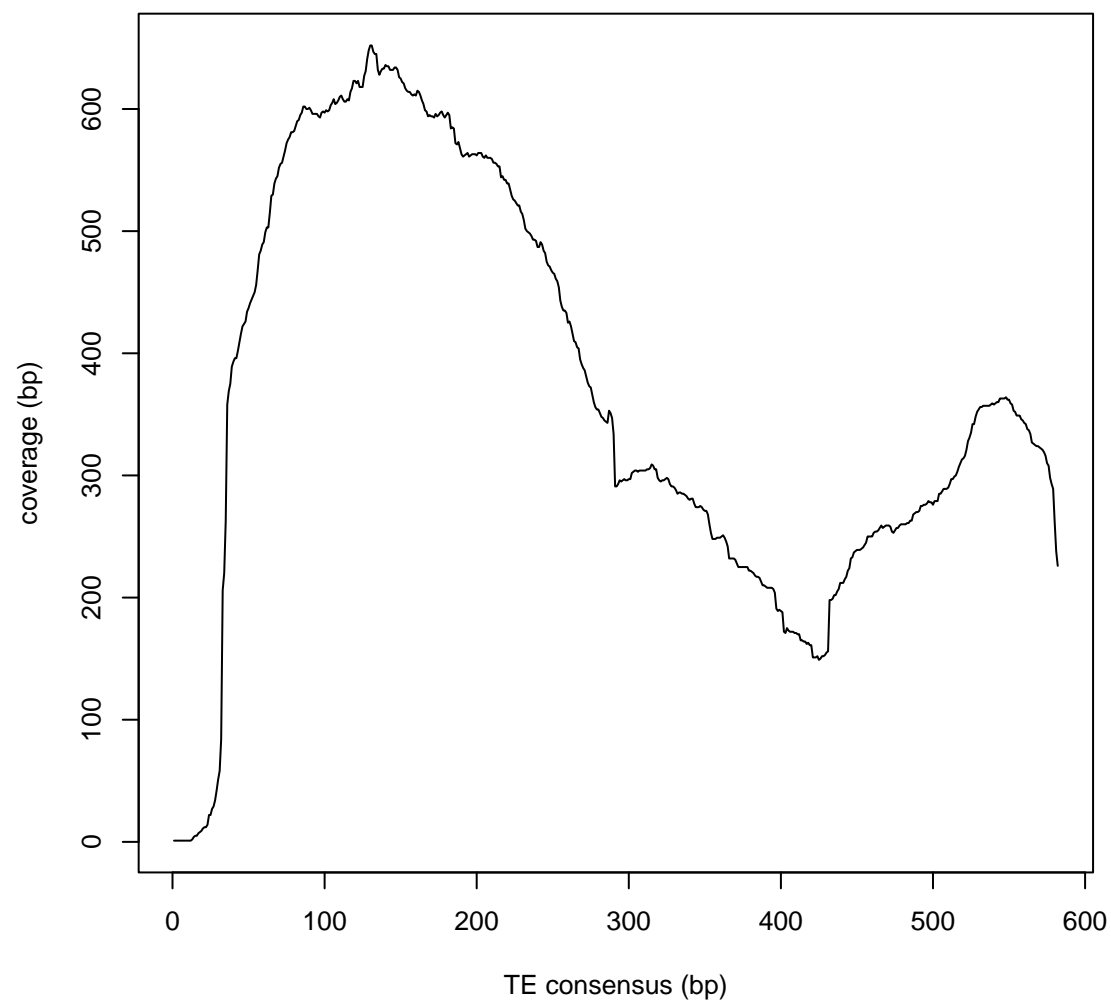


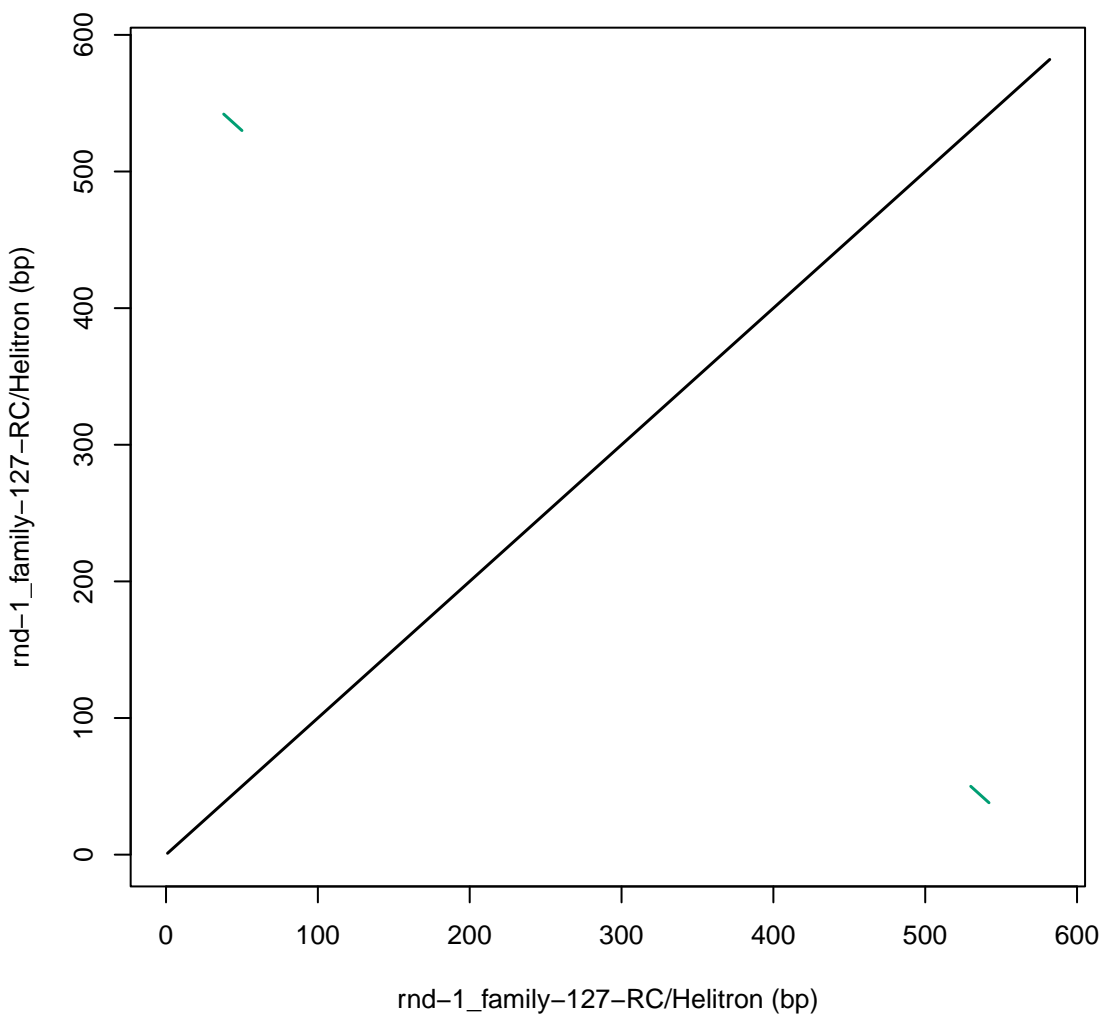
TE: rnd-1_family-127-RC/Helitron
consensus size: 582bp; fragments: 1184; full length: 22 (≥ 523.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

