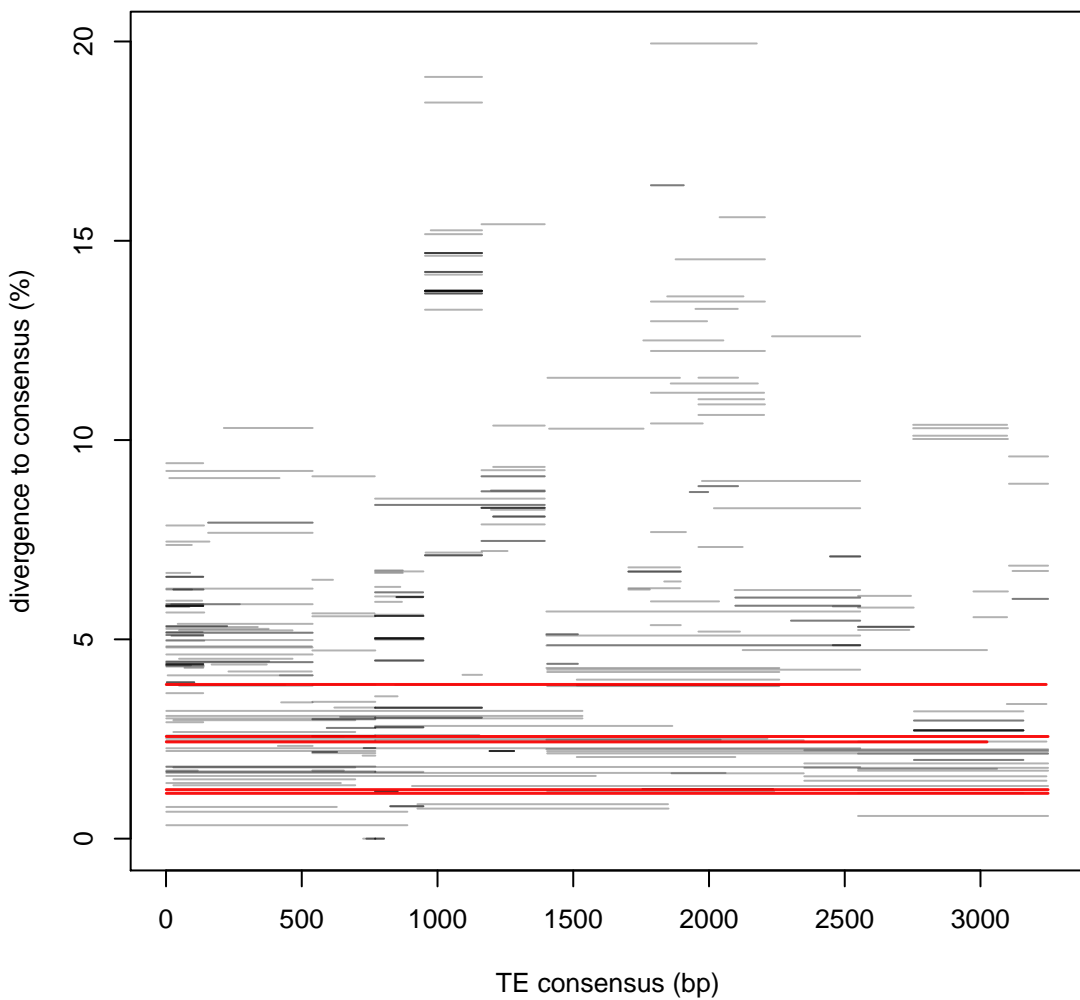
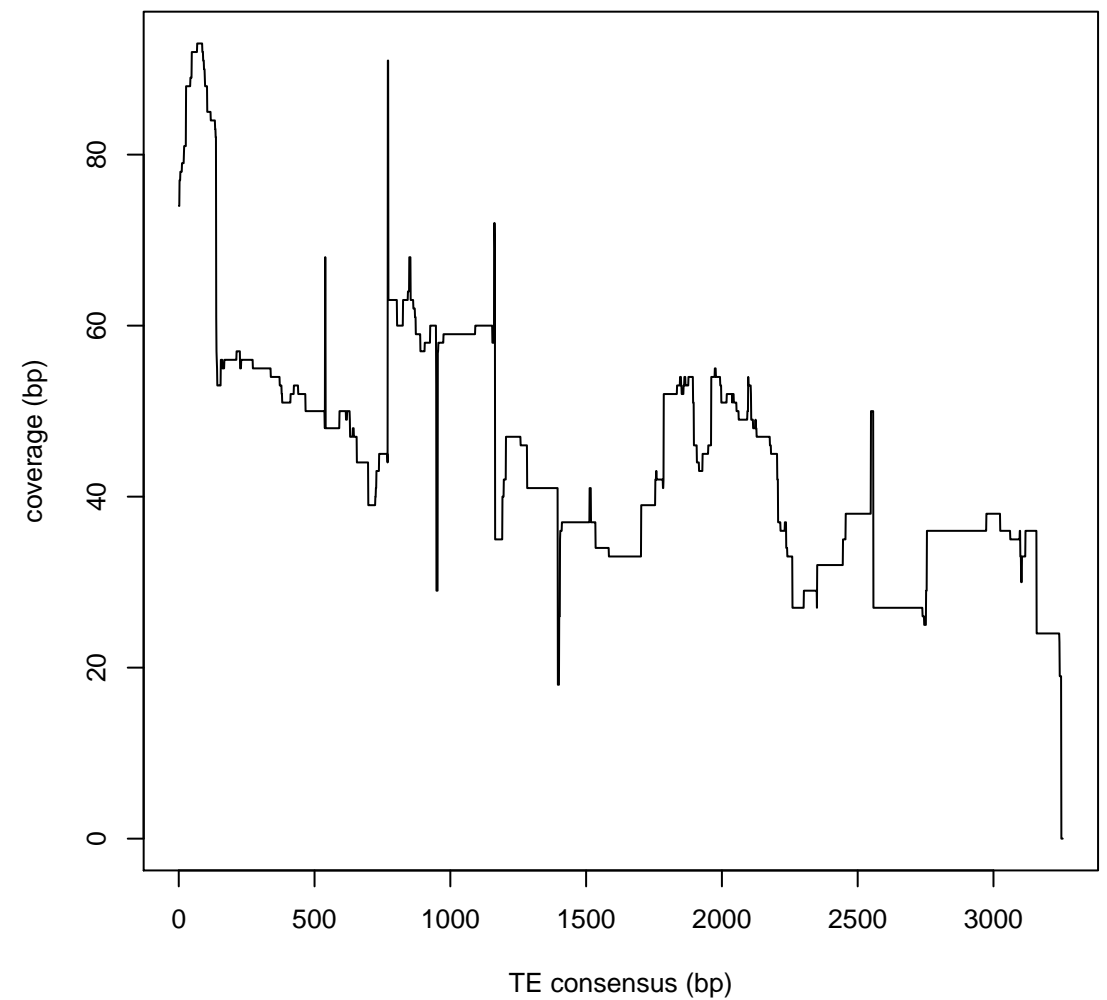


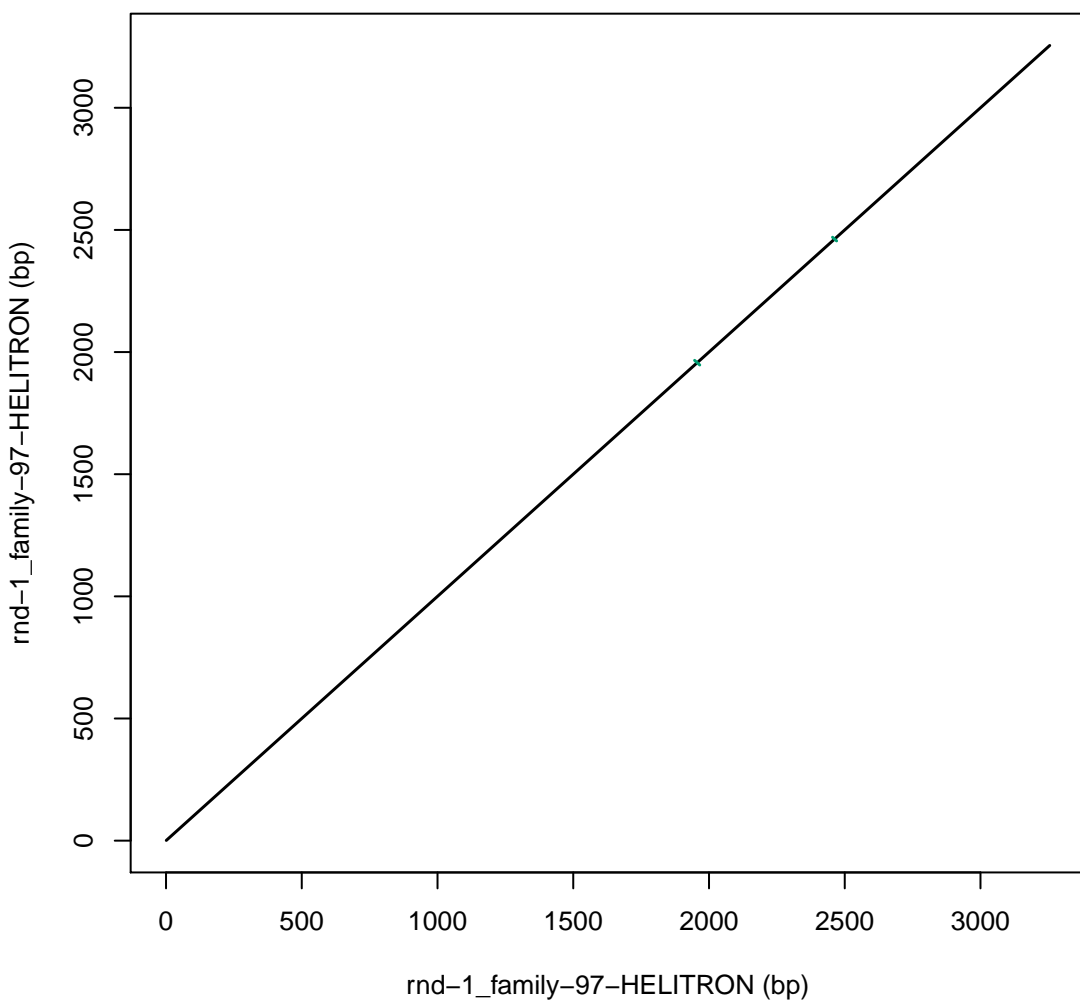
TE: rnd-1_family-97-HELITRON
consensus size: 3255bp; fragments: 367; full length: 5 (≥ 2929.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

