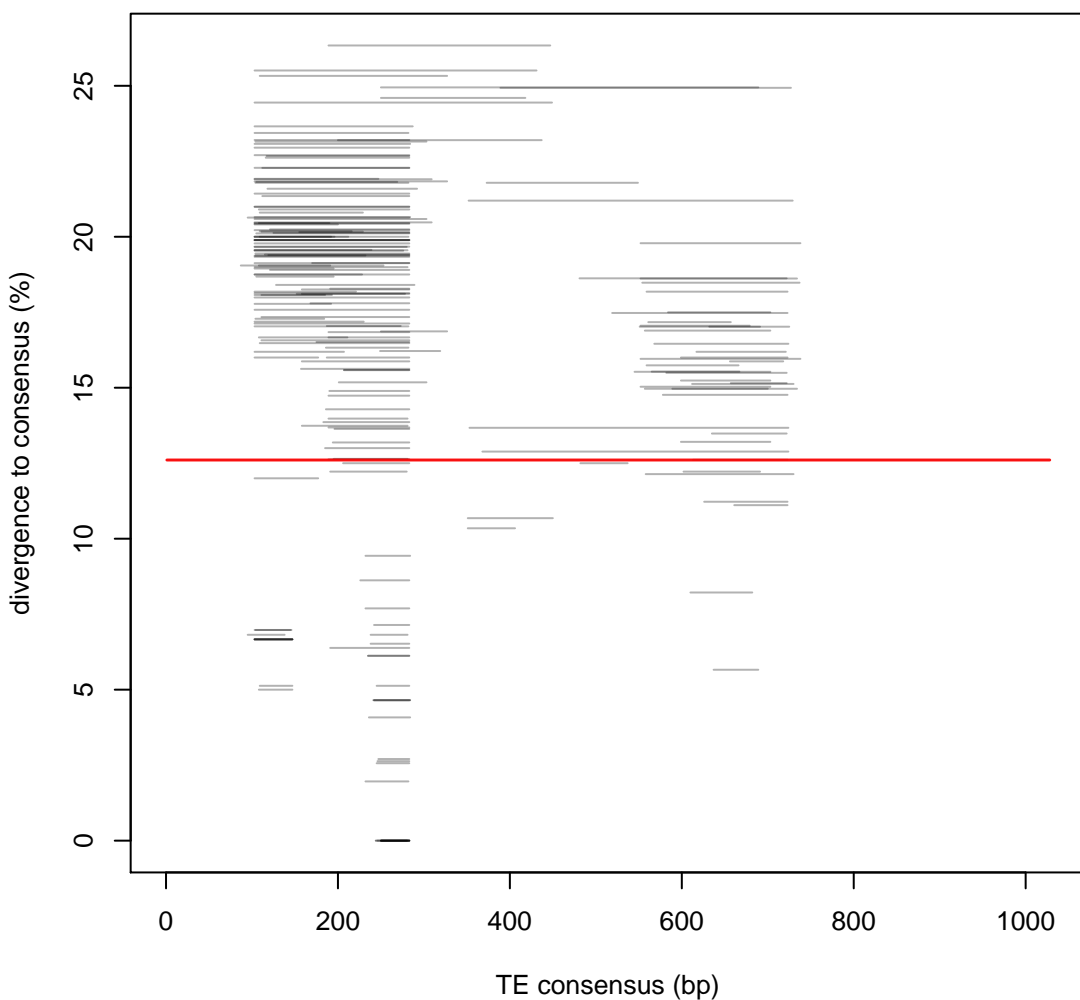
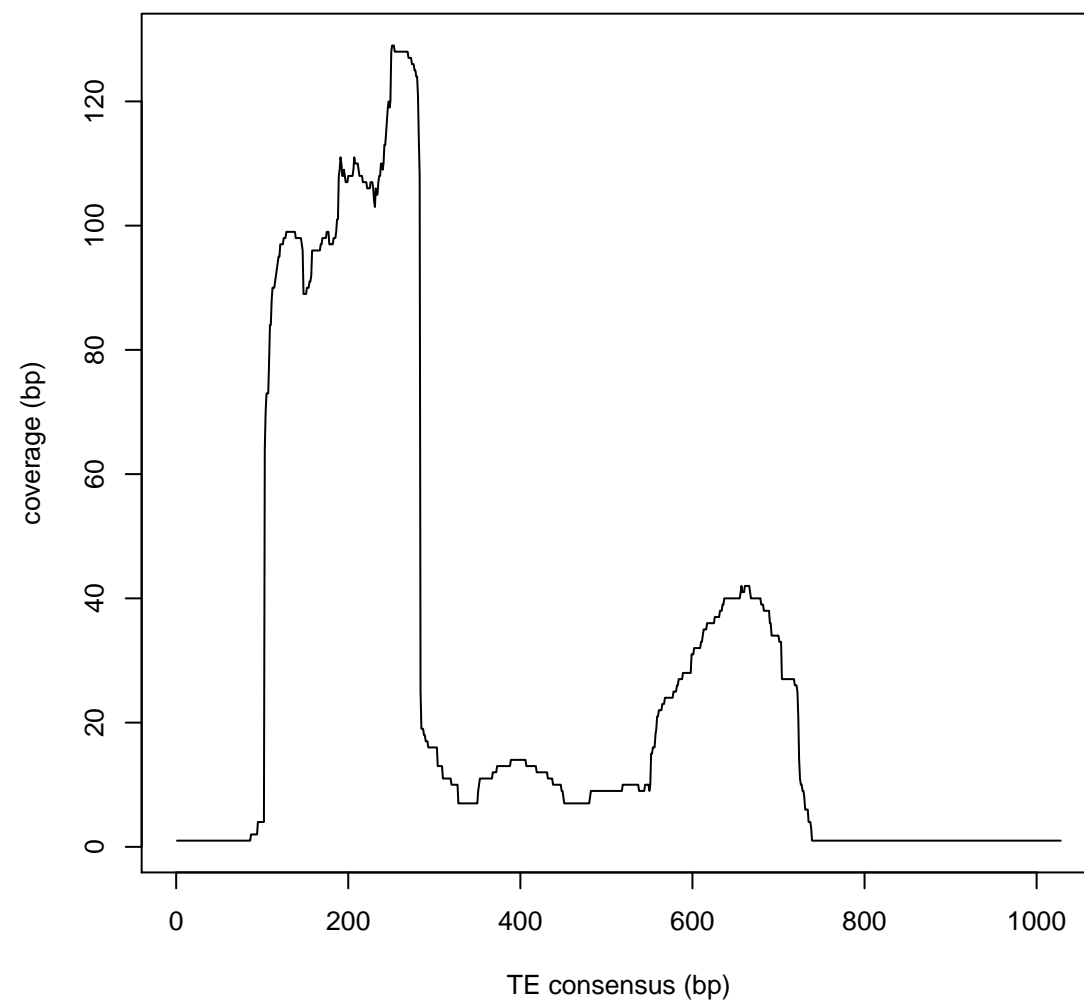


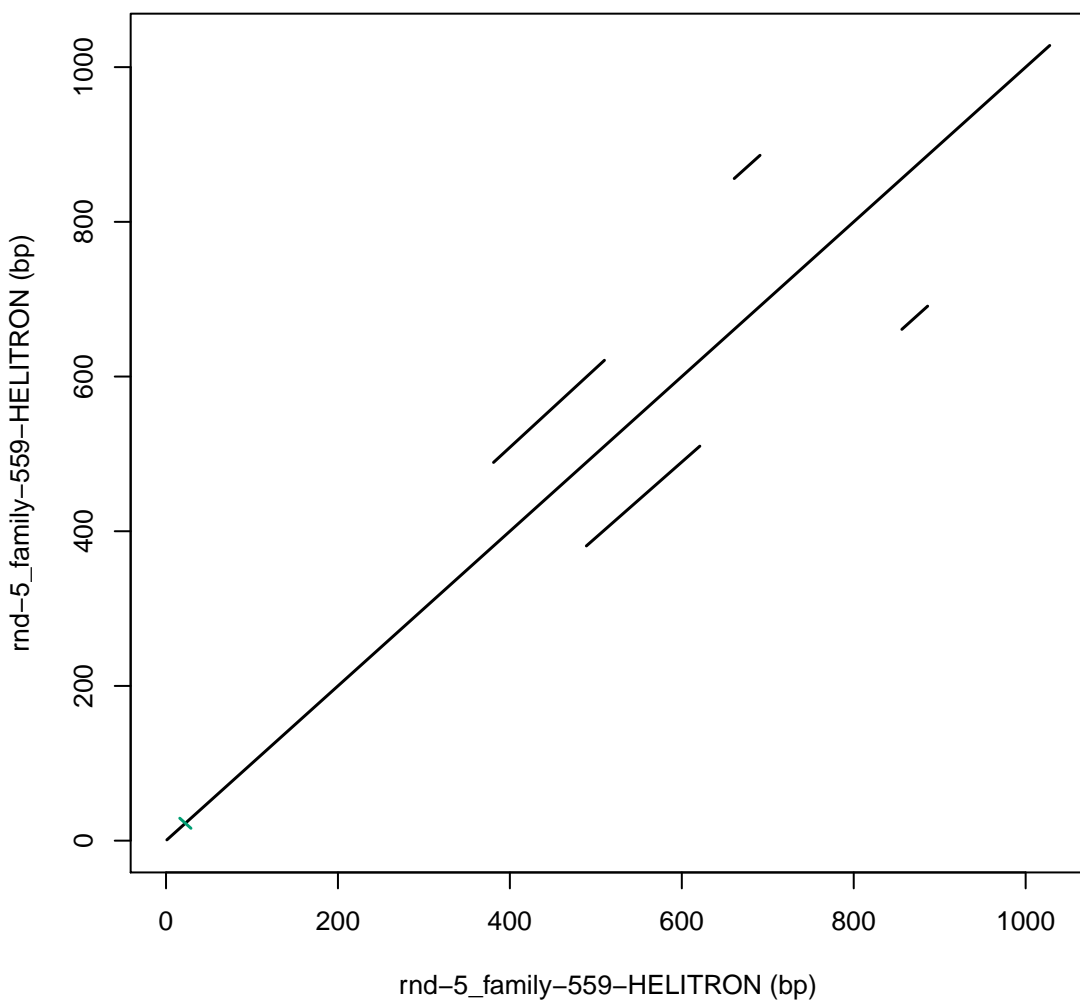
TE: rnd-5_family-559-HELITRON
consensus size: 1028bp; fragments: 209; full length: 1 (>=925.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

