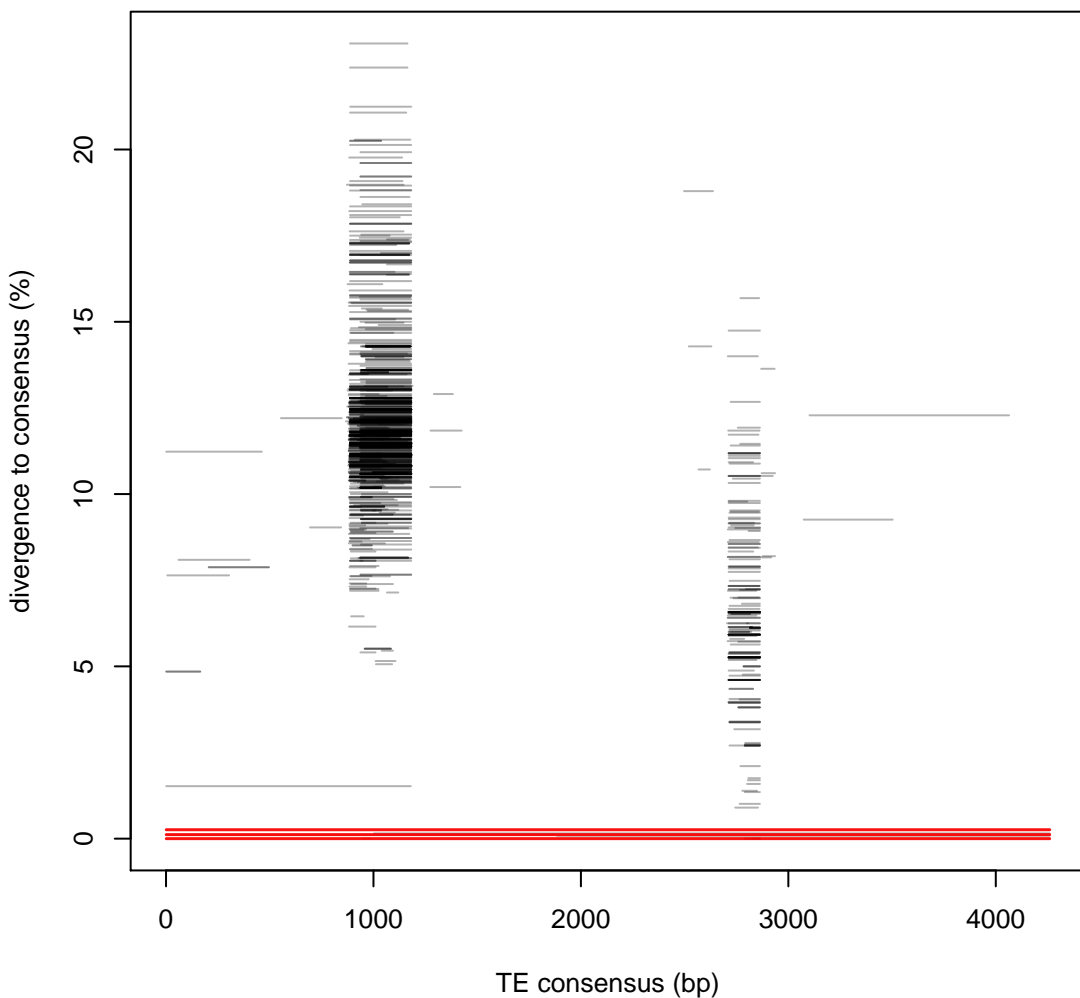
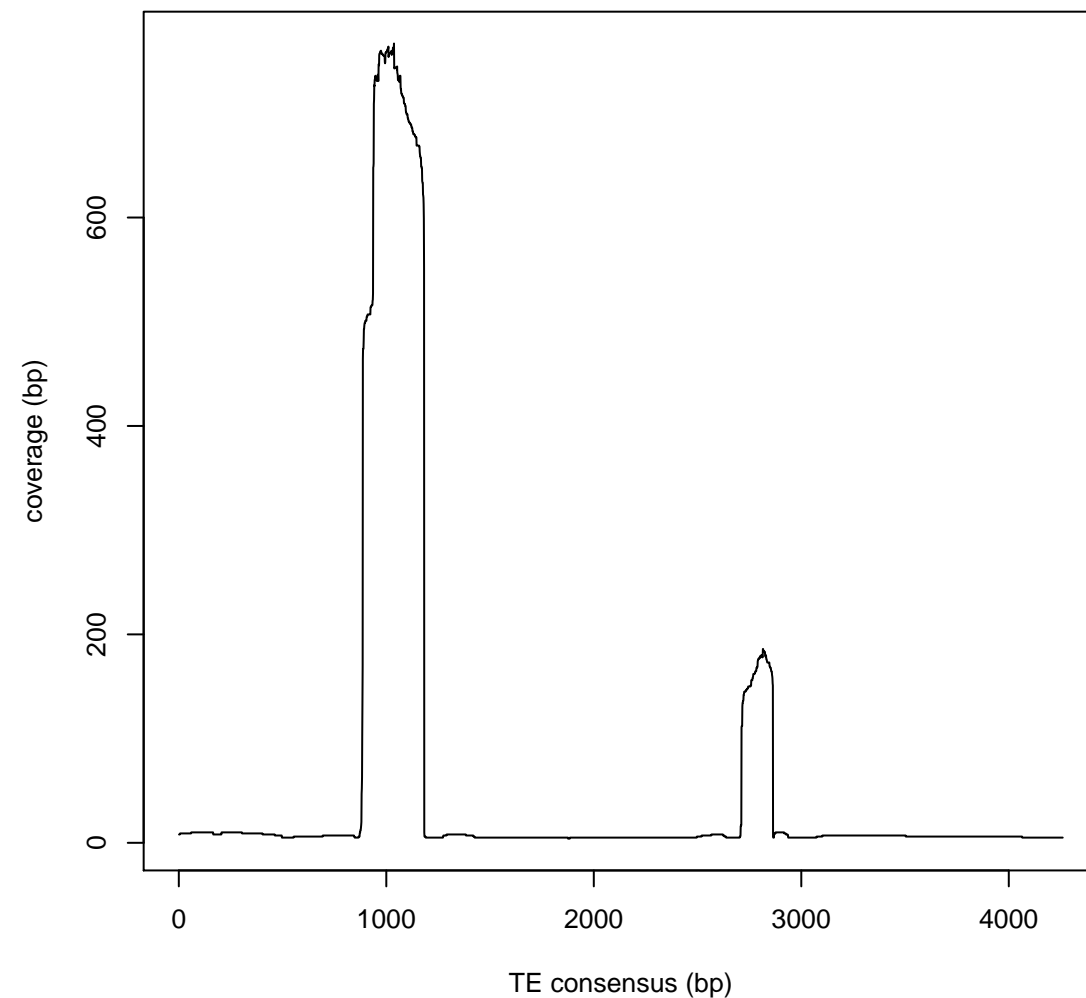


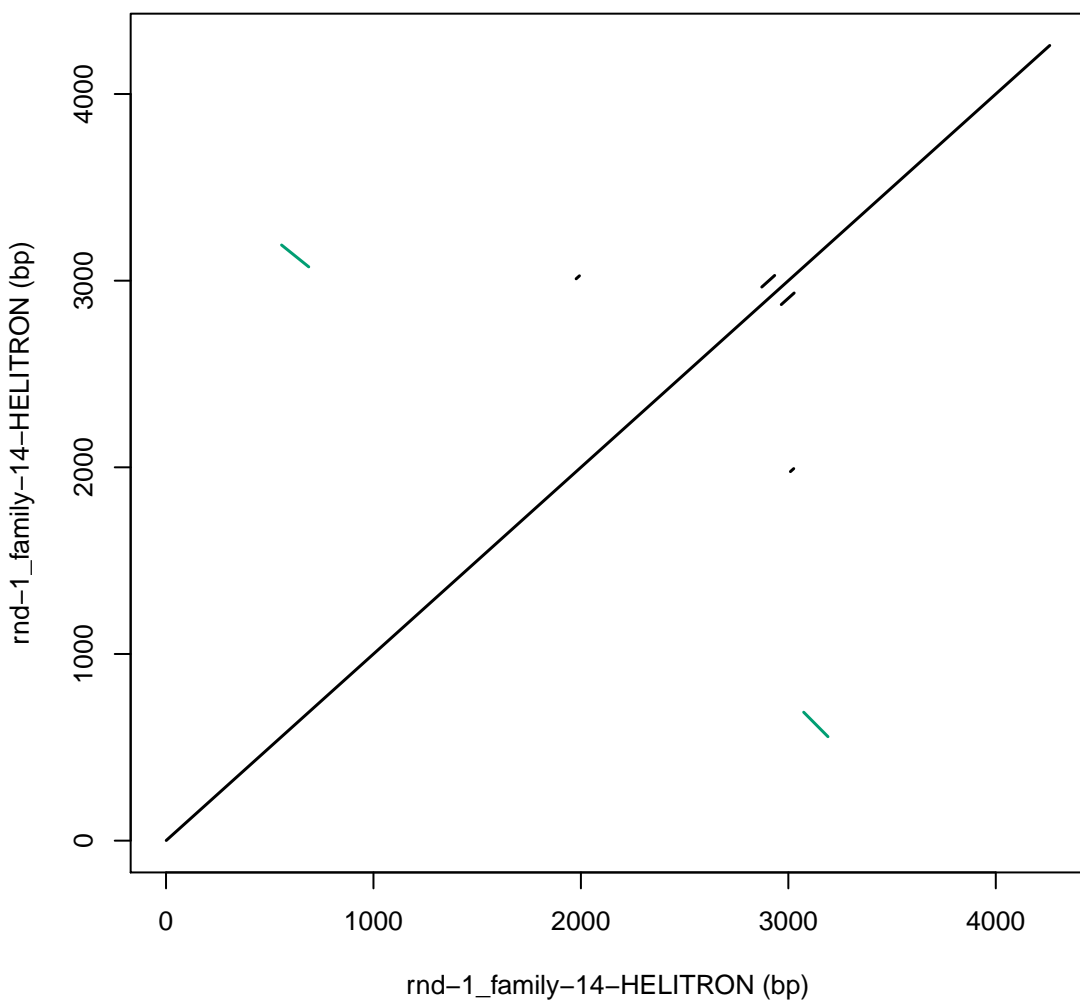
TE: rnd-1_family-14-HELITRON
consensus size: 4260bp; fragments: 1042; full length: 3 (>=3834bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

