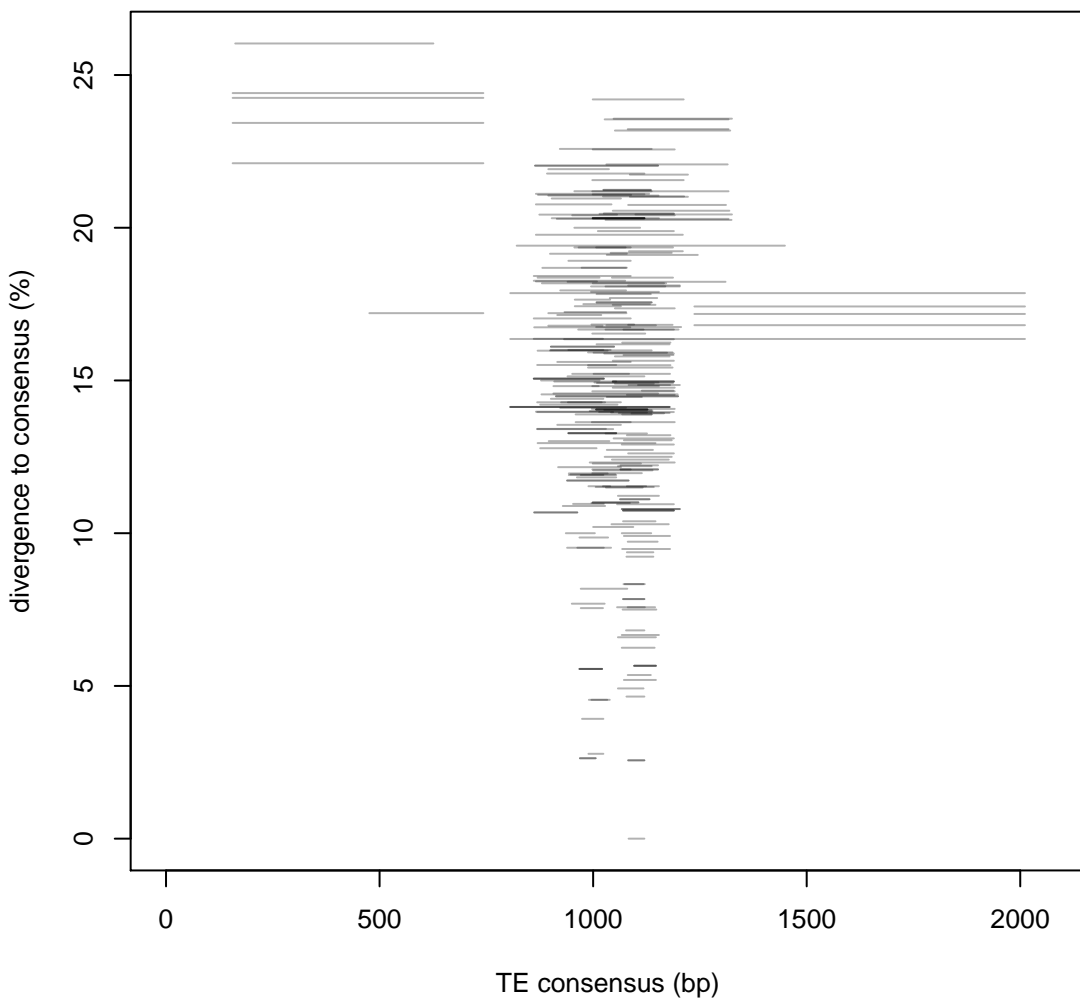
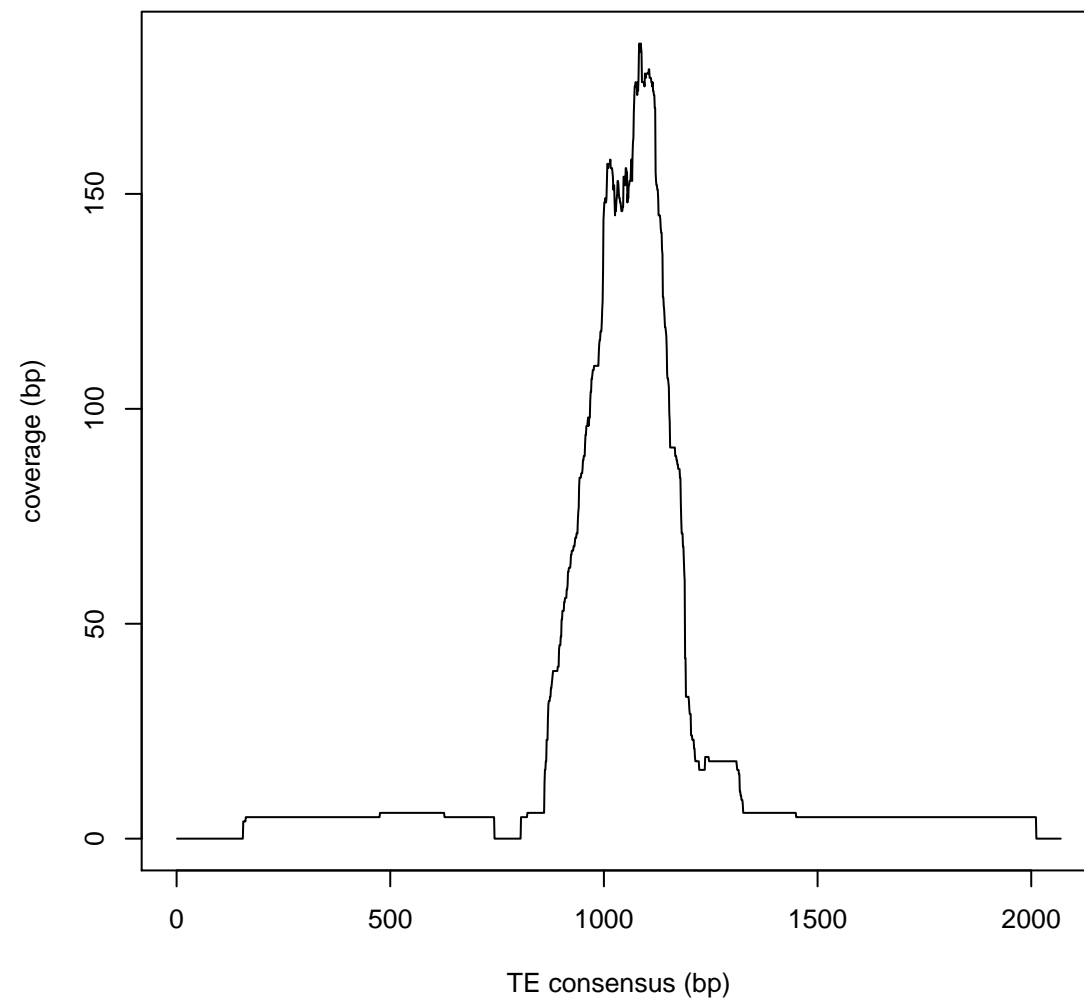


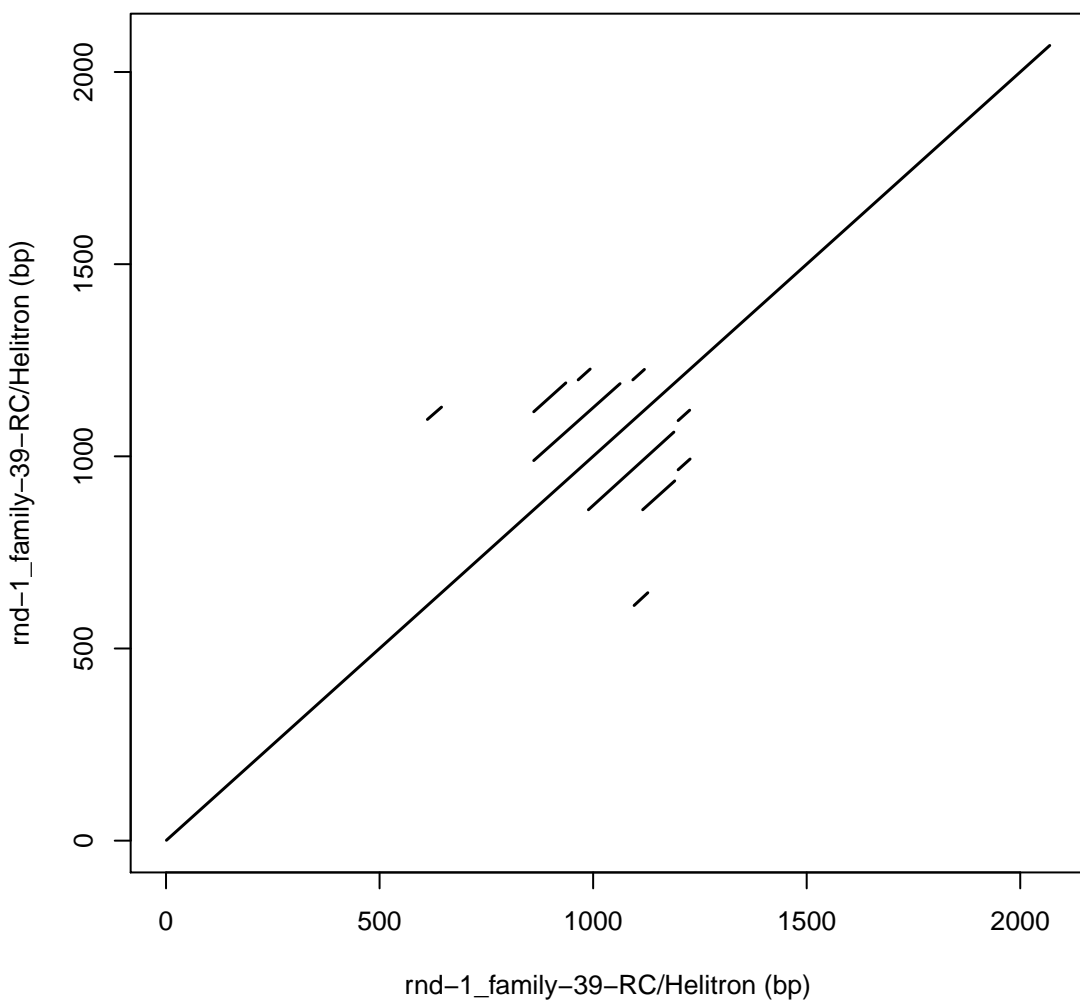
TE: rnd-1_family-39-RC/Helitron
consensus size: 2069bp; fragments: 277; full length: 0 (>=1862.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

