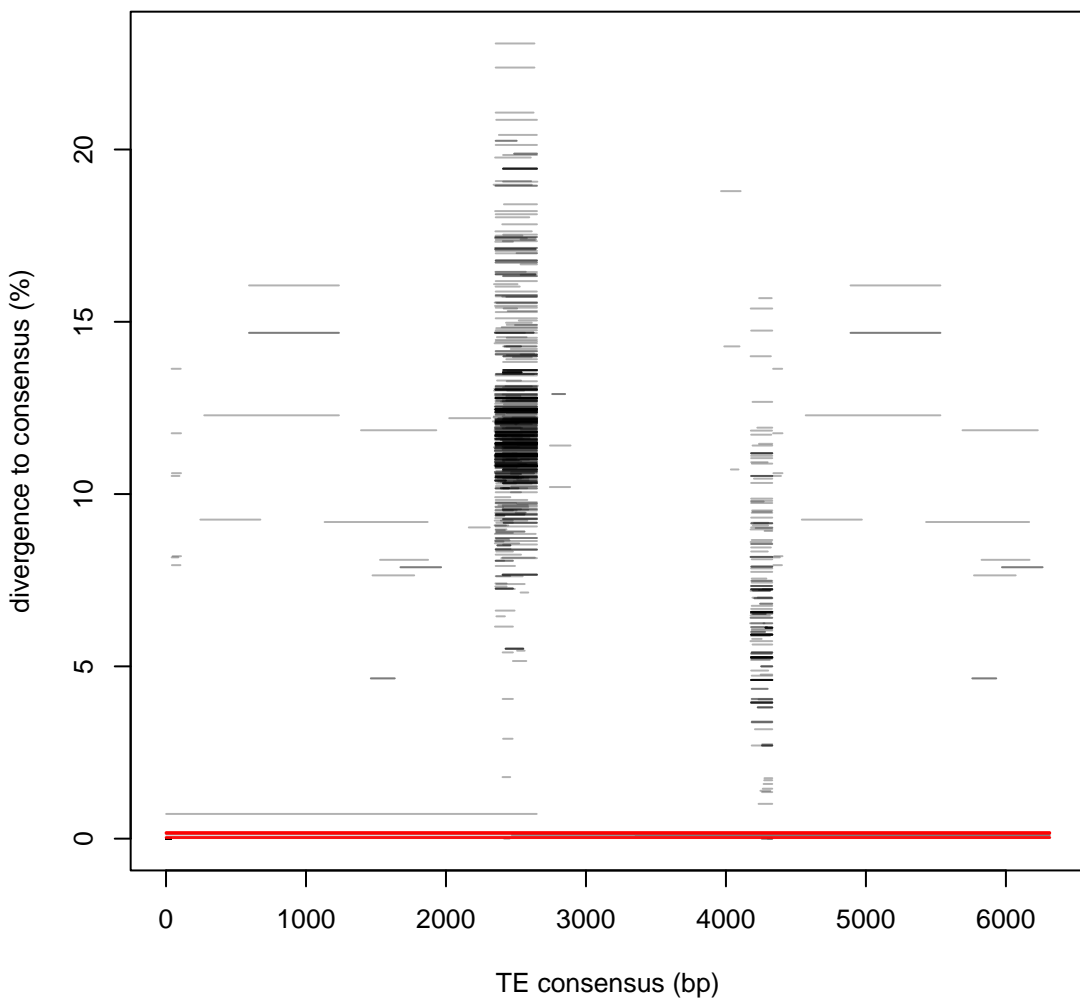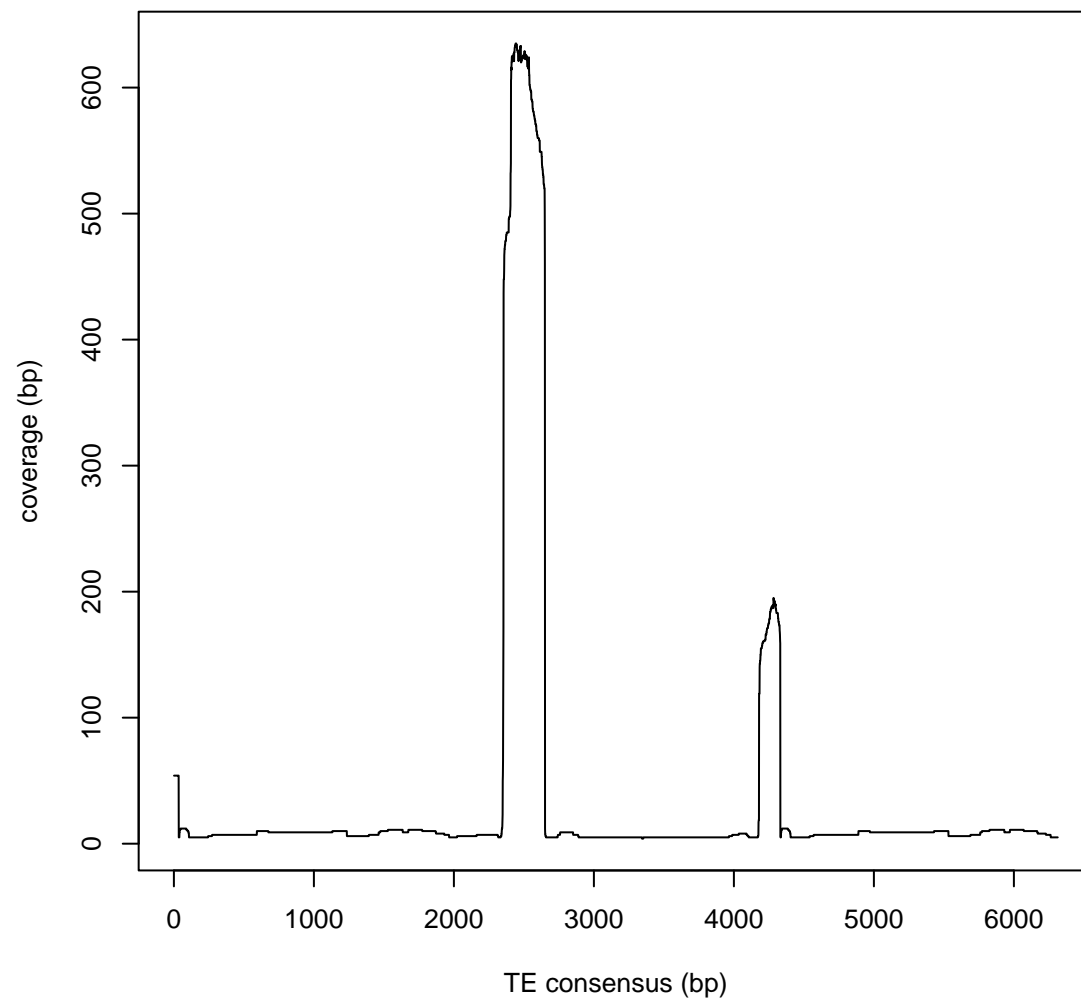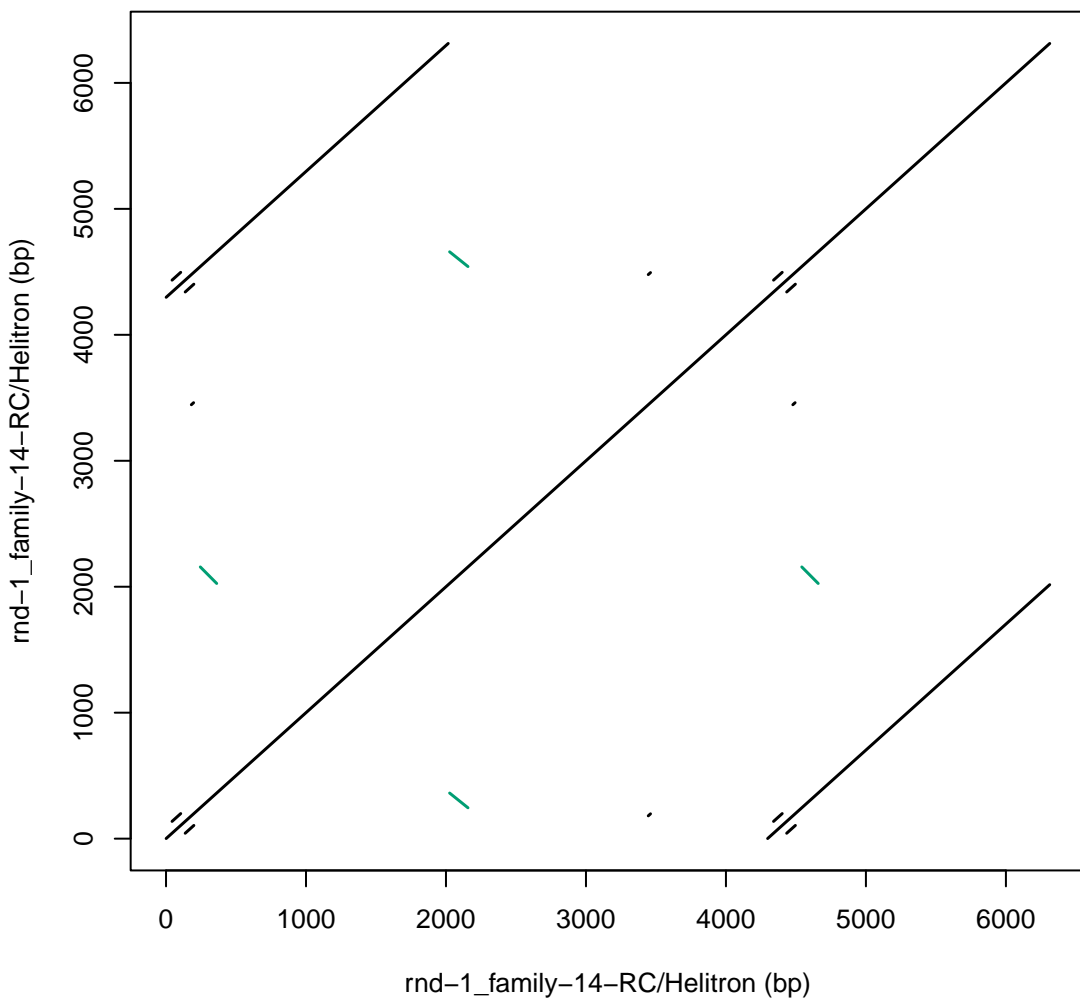**TE: rnd-1_family-14-RC/Helitron**
consensus size: 6313bp; fragments: 1002; full length: 3 (>=5681.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits