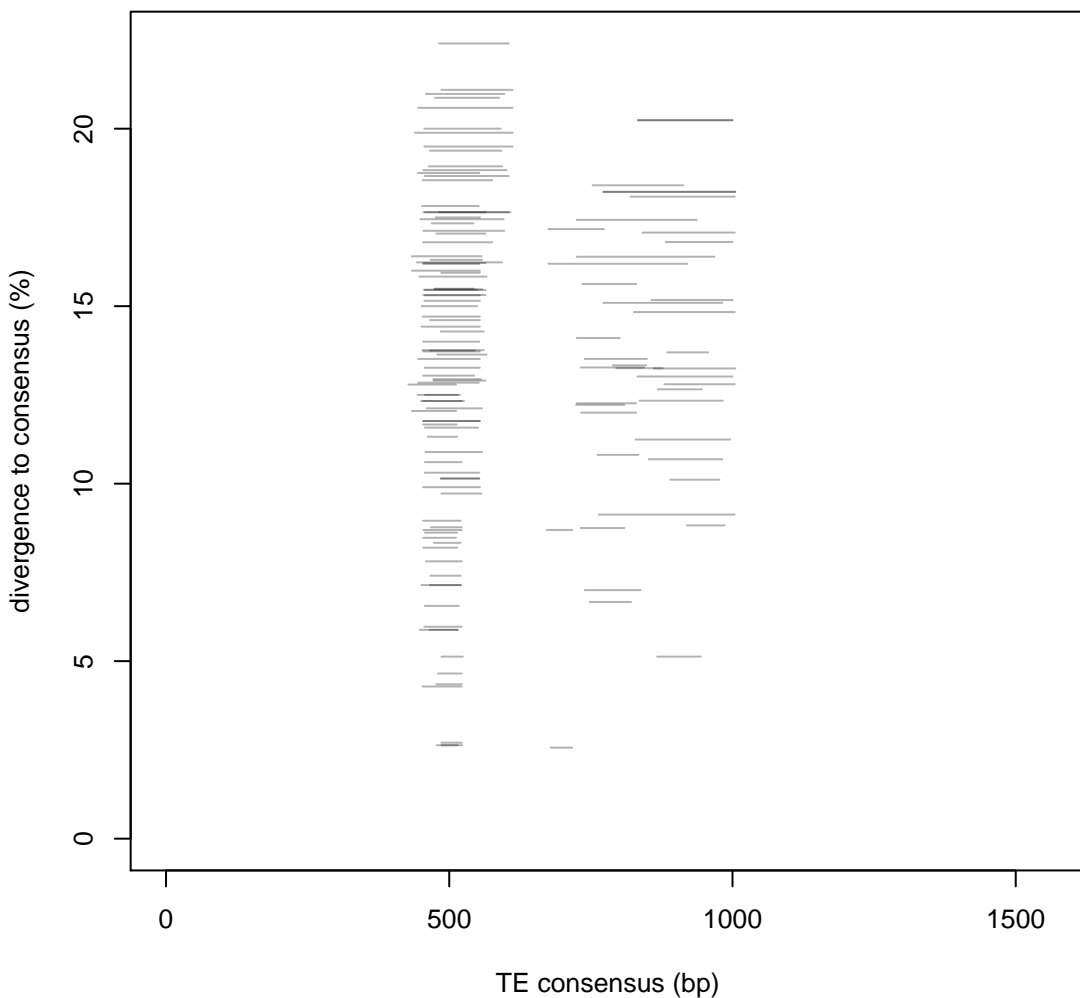
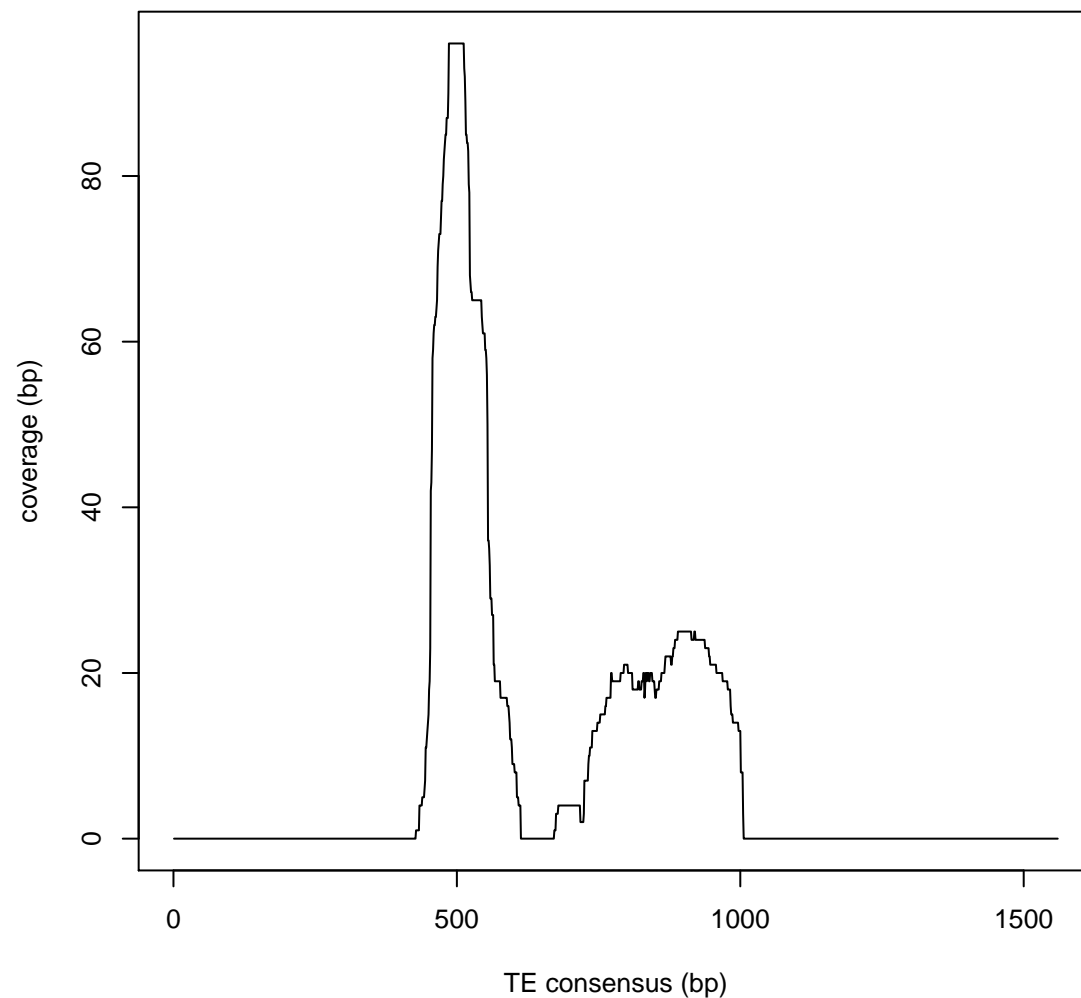


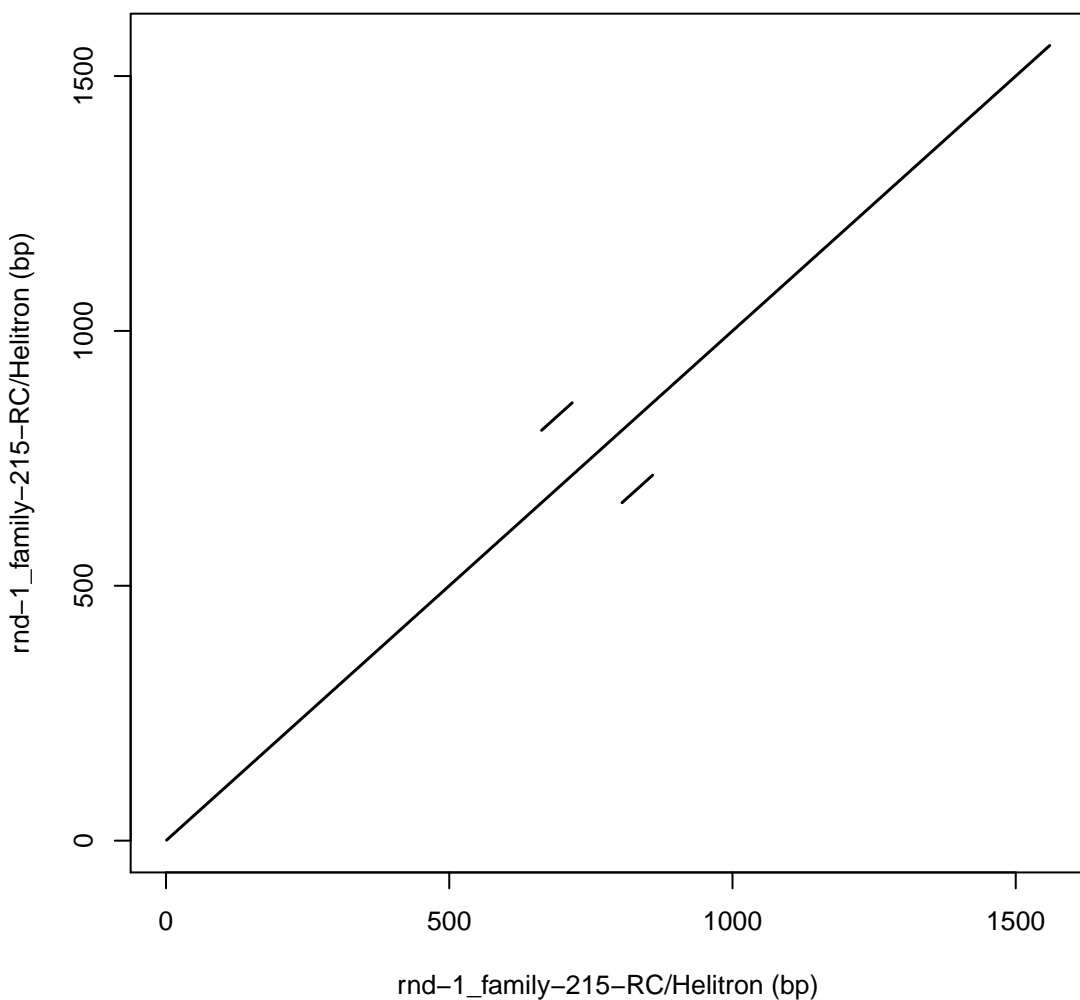
TE: rnd-1_family-215-RC/Helitron
consensus size: 1560bp; fragments: 138; full length: 0 (>=1404bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

