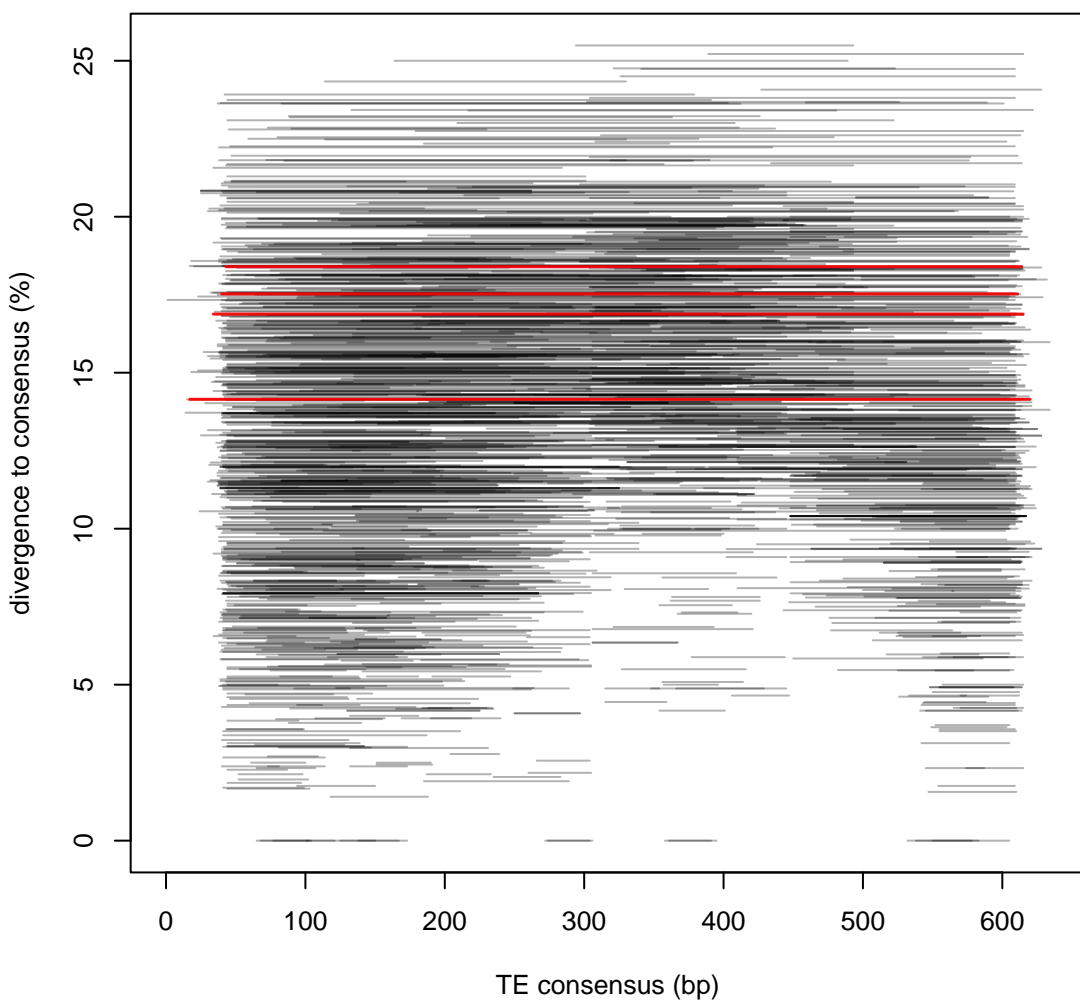
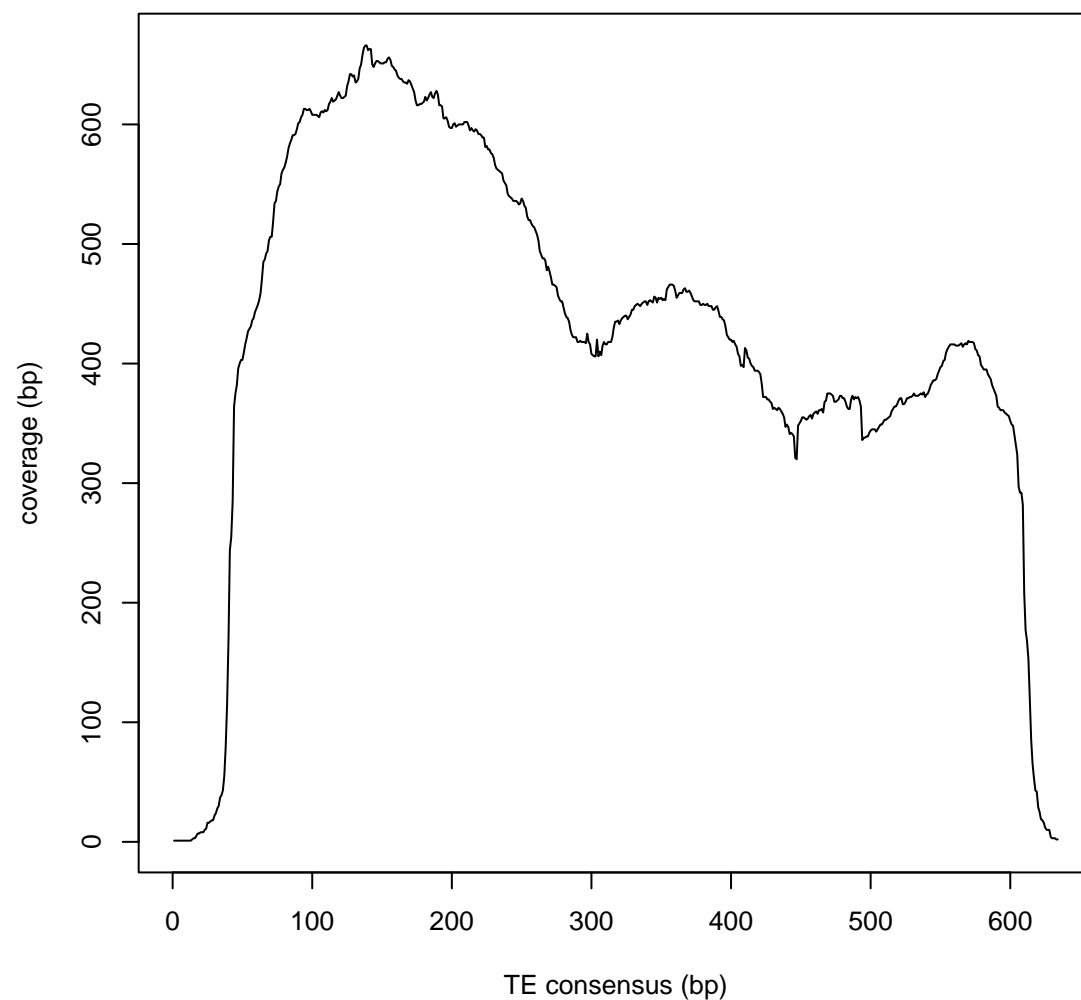


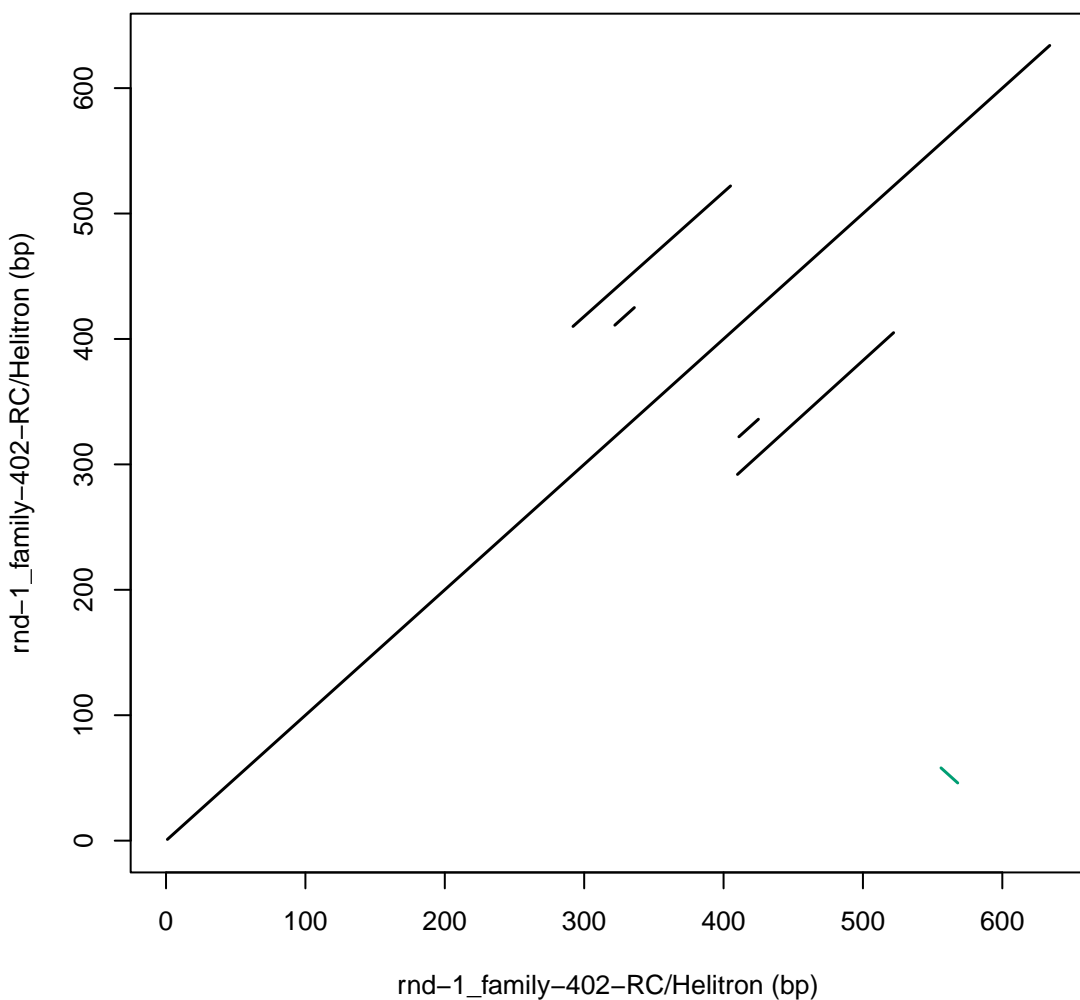
TE: rnd-1_family-402-RC/Helitron
consensus size: 634bp; fragments: 1420; full length: 4 (>=570.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

