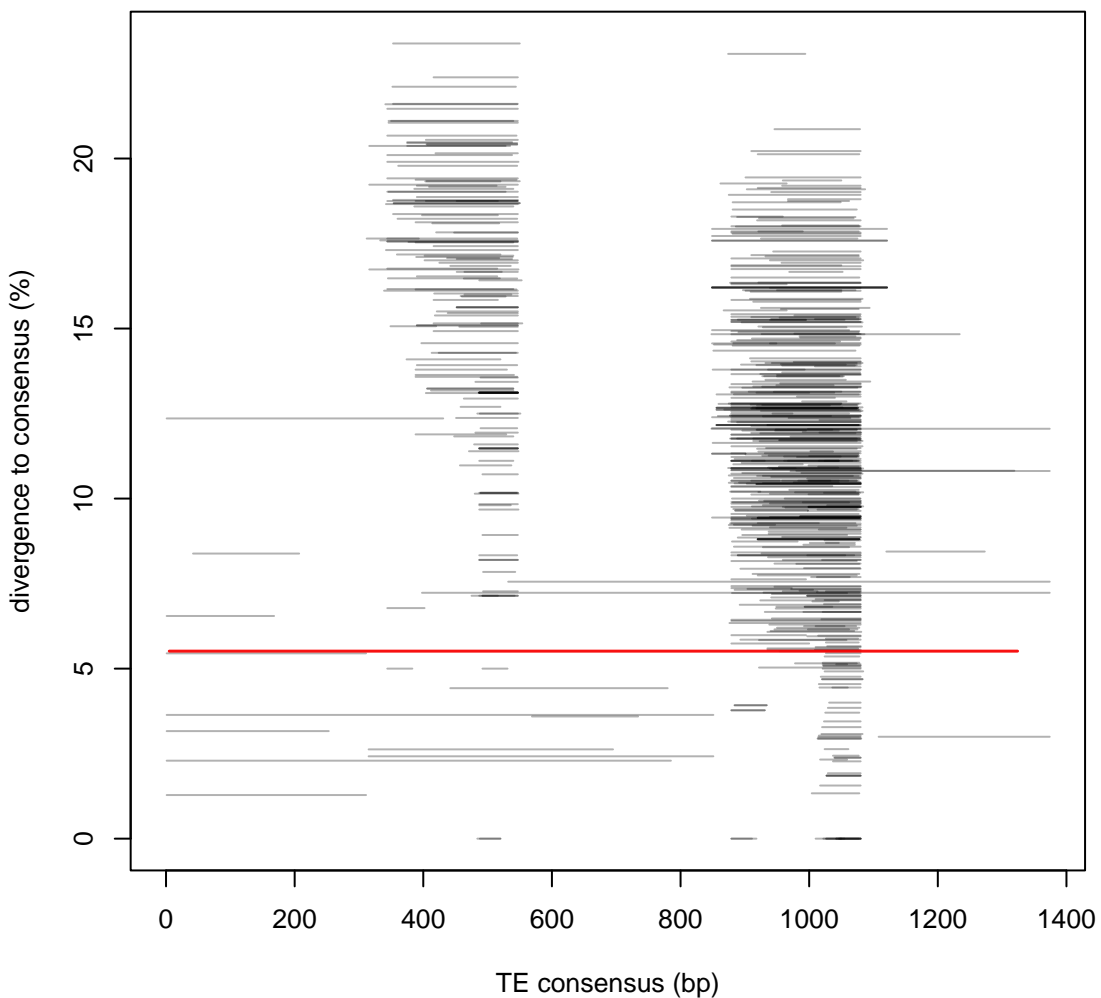
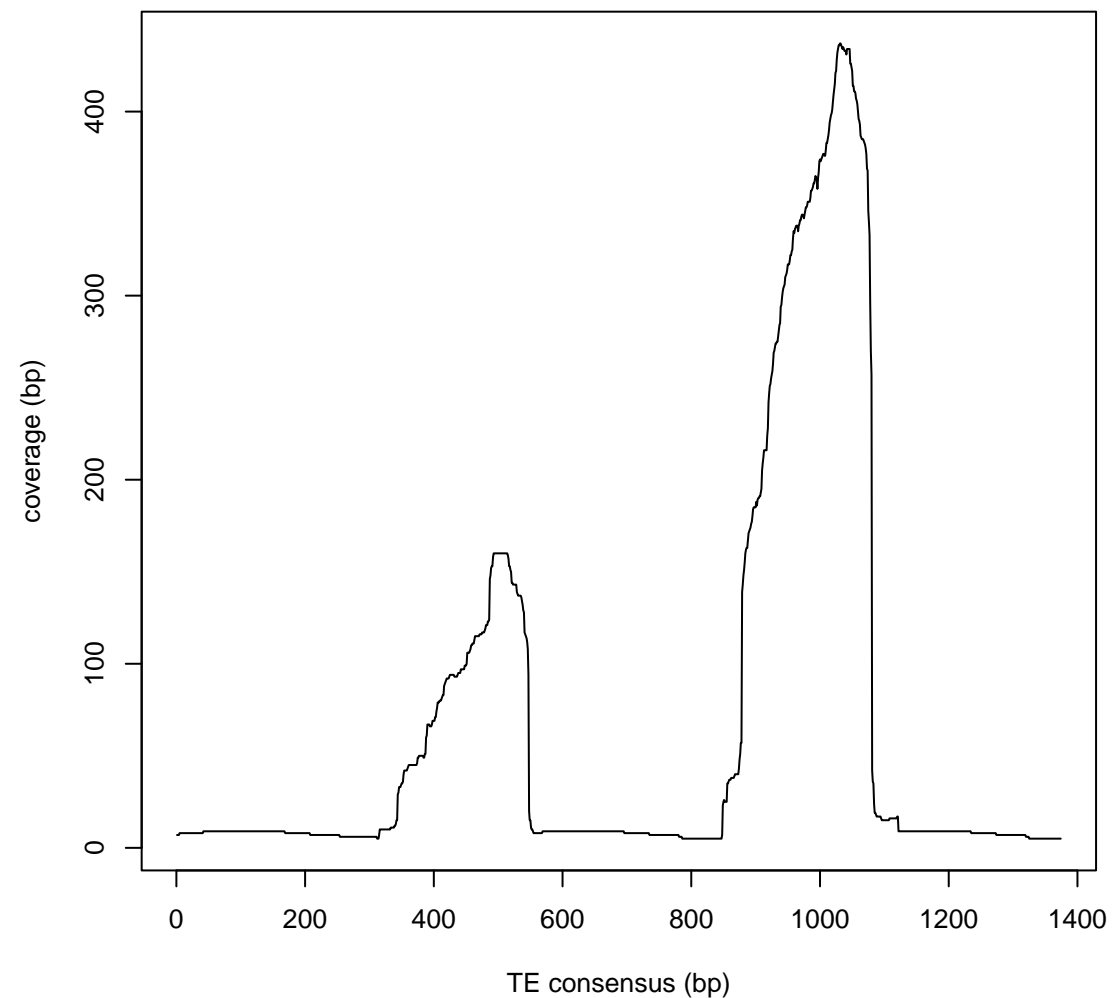


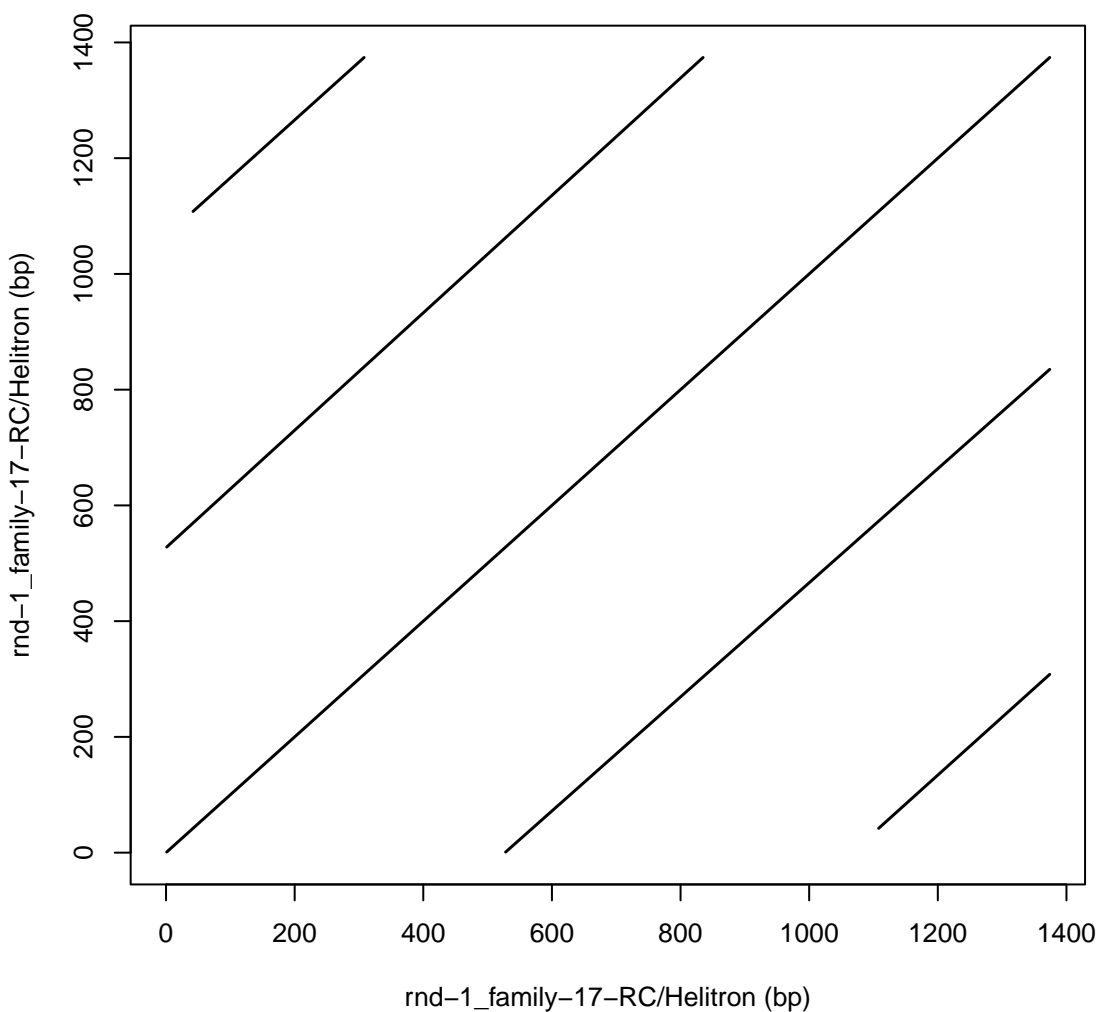
TE: rnd-1\_family-17-RC/Helitron  
consensus size: 1374bp; fragments: 655; full length: 1 ( $\geq 1236.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

