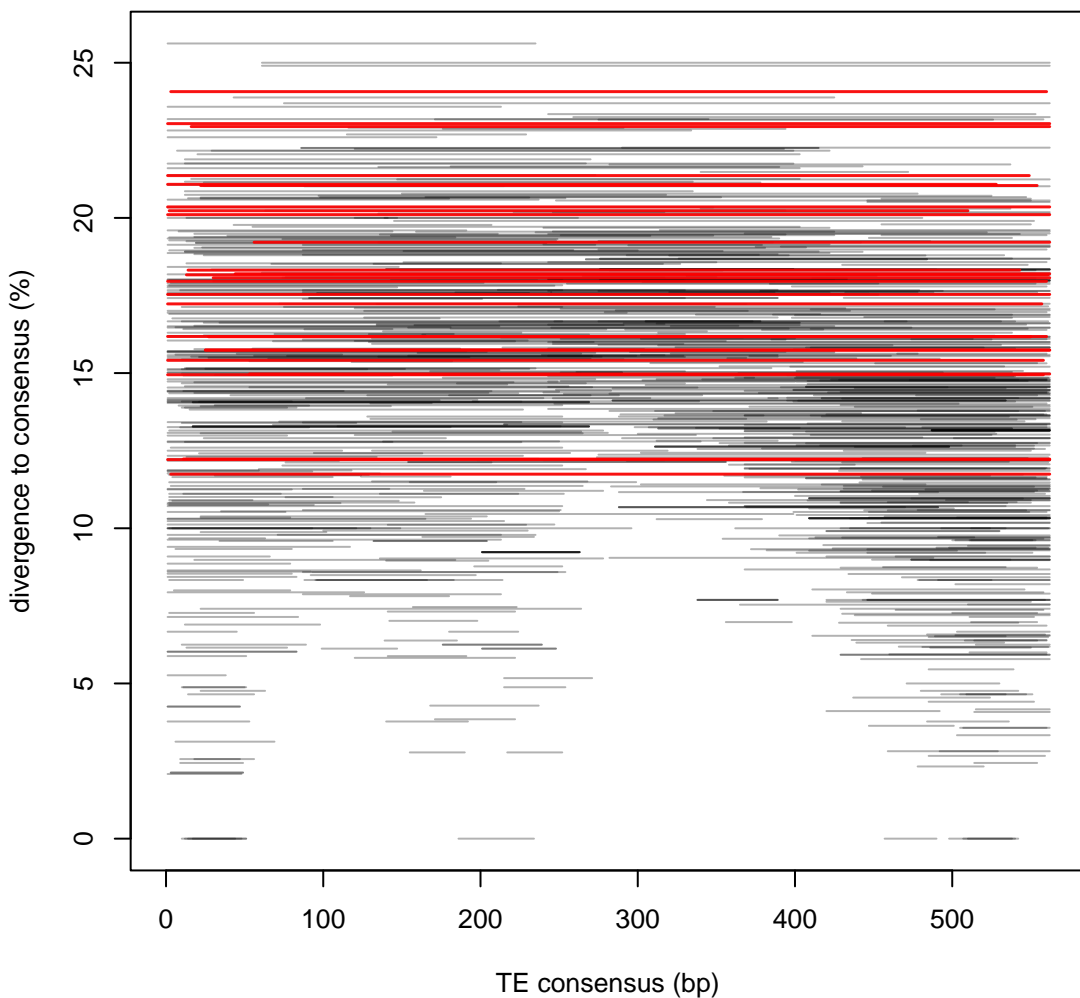
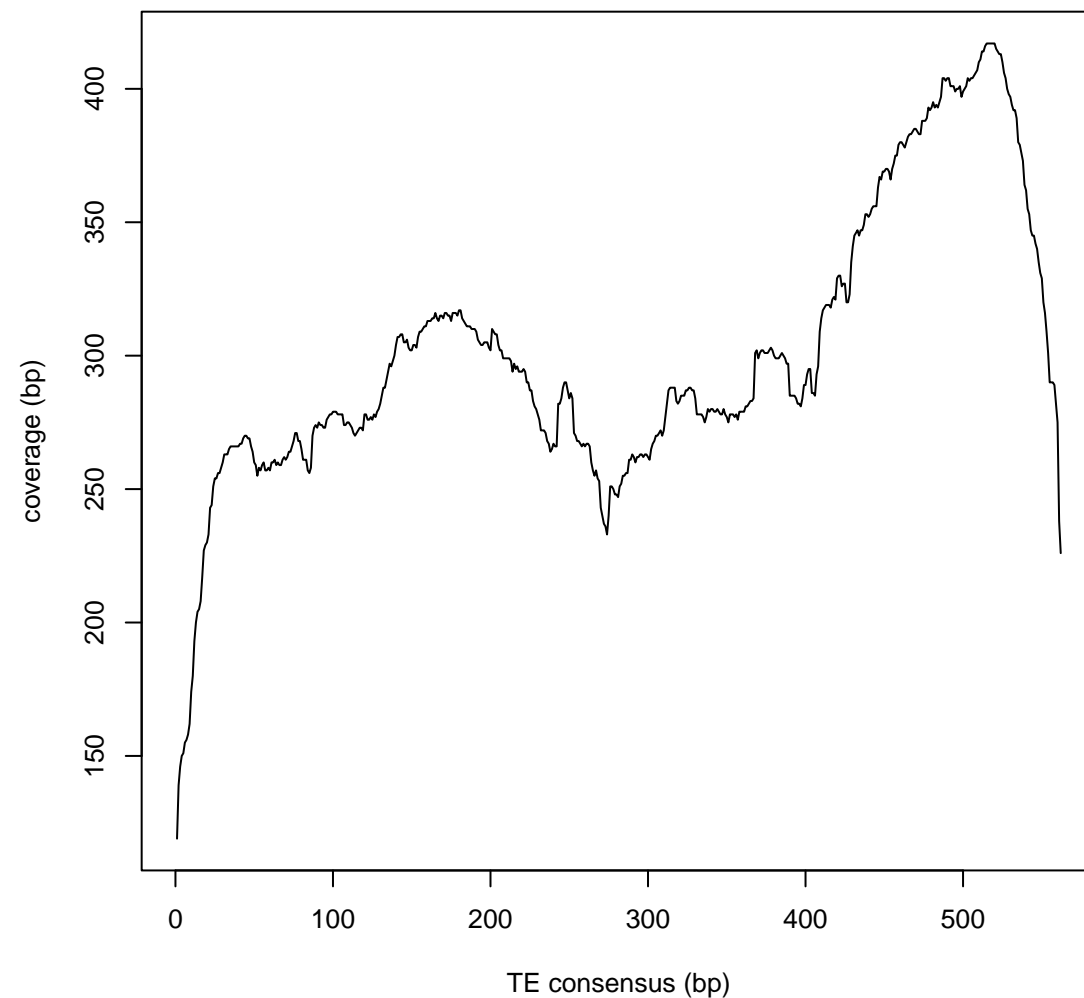


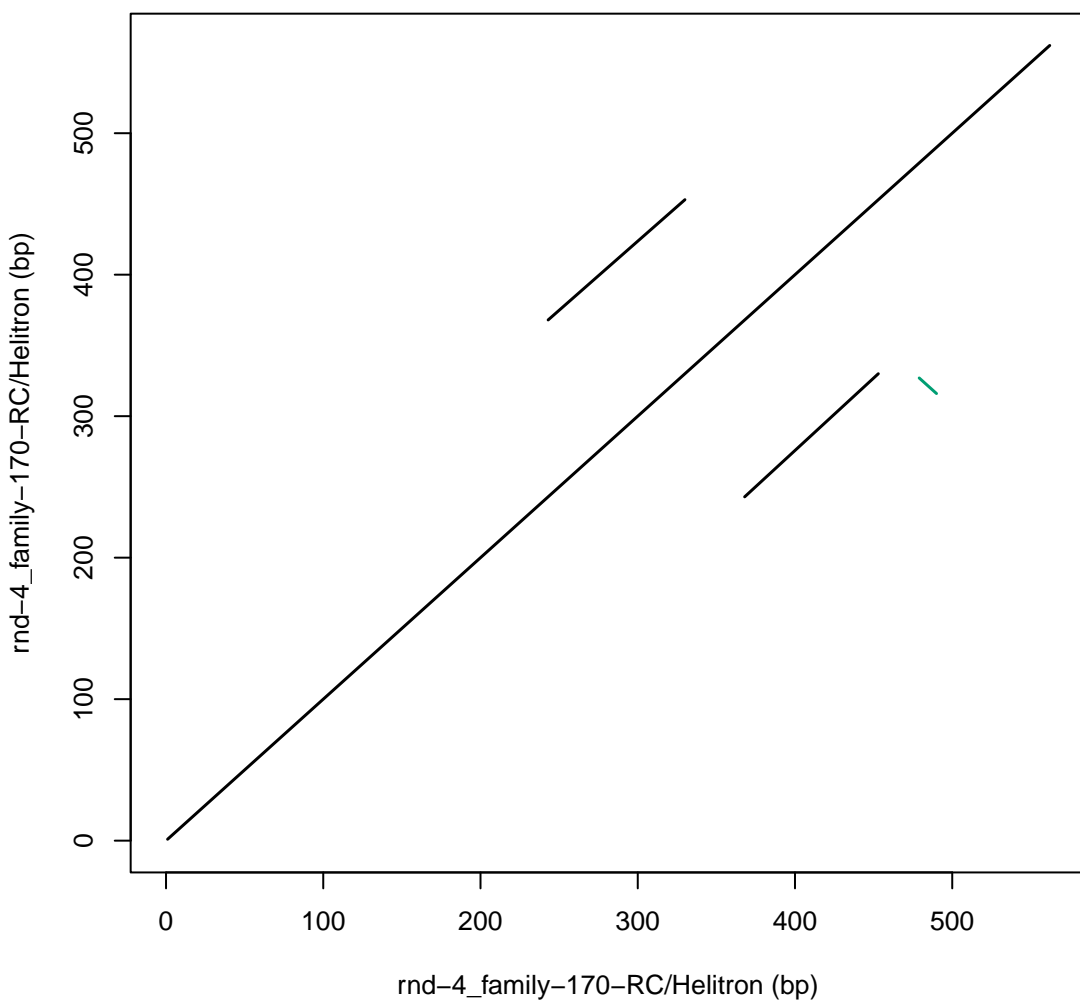
TE: rnd-4\_family-170-RC/Helitron  
consensus size: 562bp; fragments: 881; full length: 27 ( $\geq 505.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

