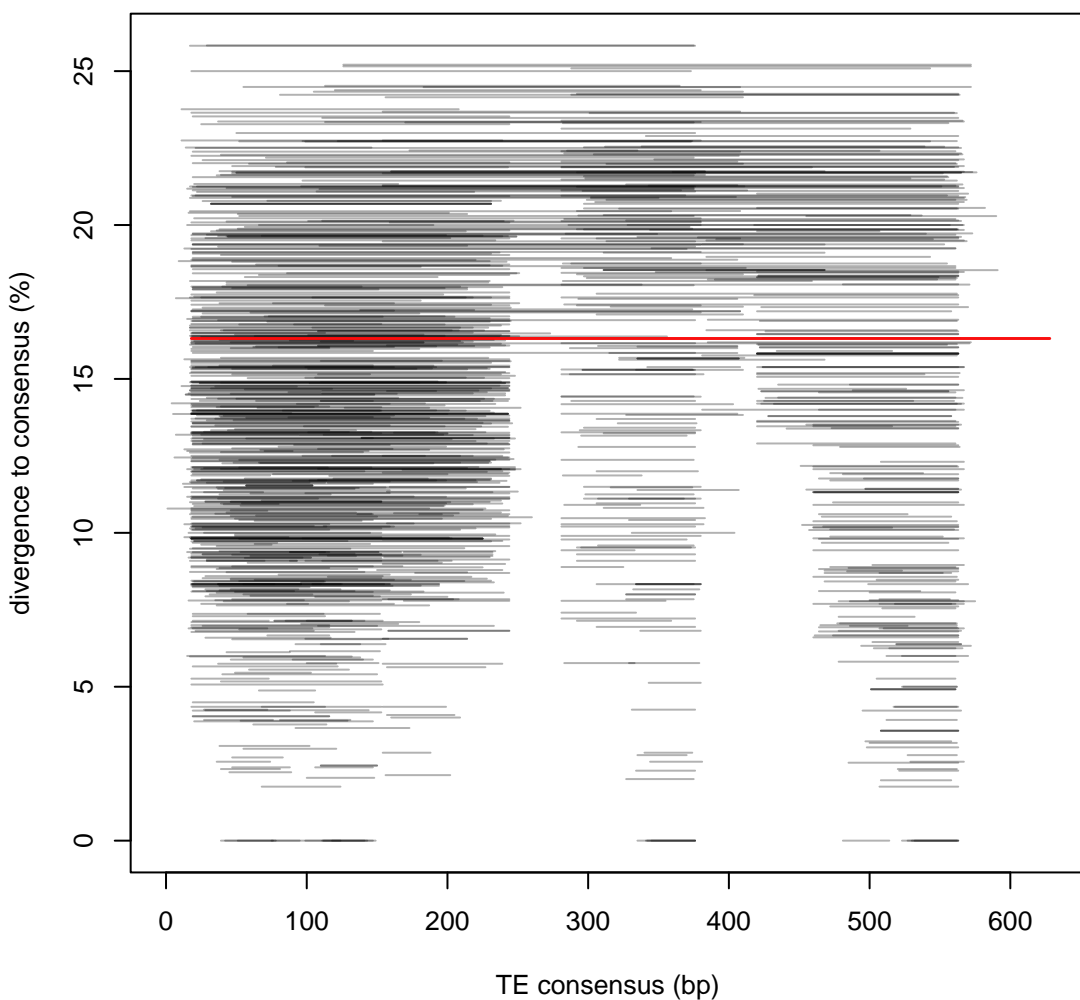
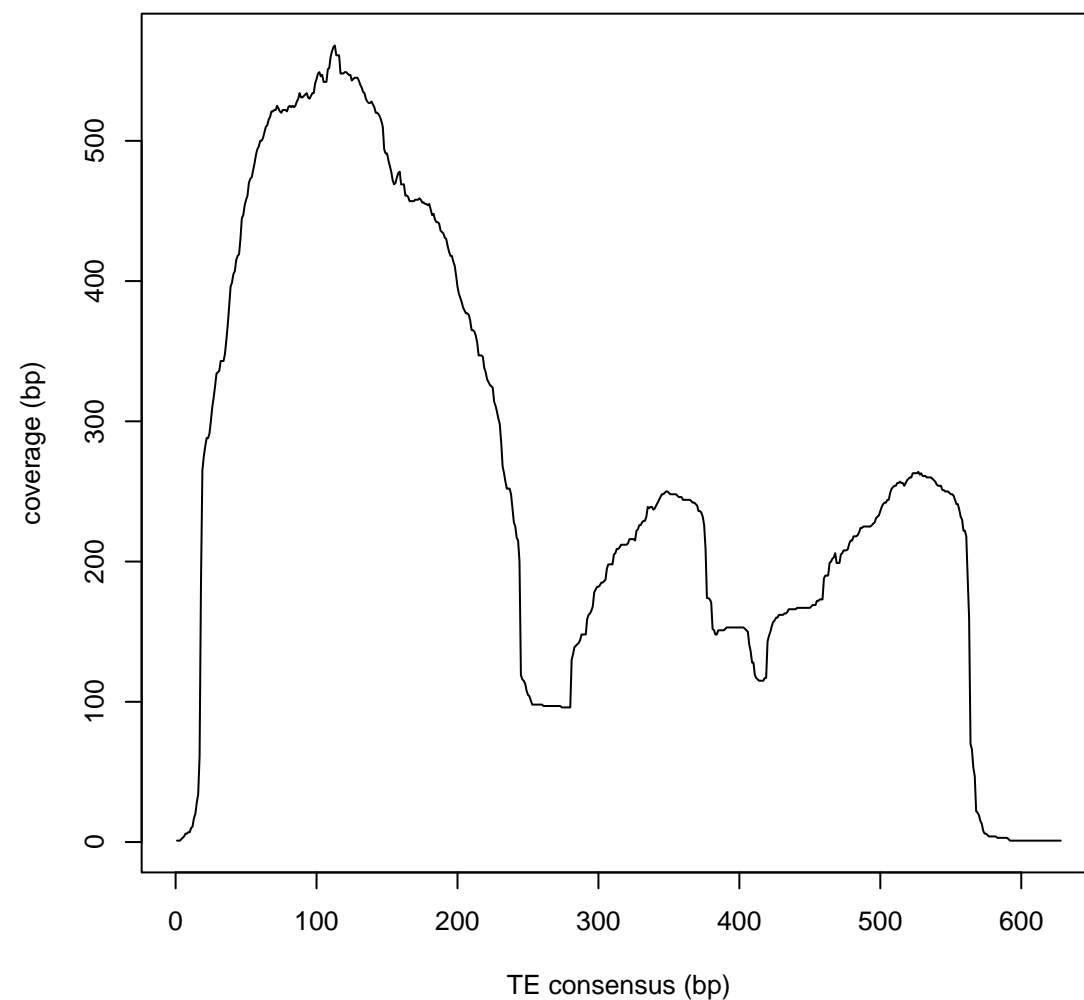


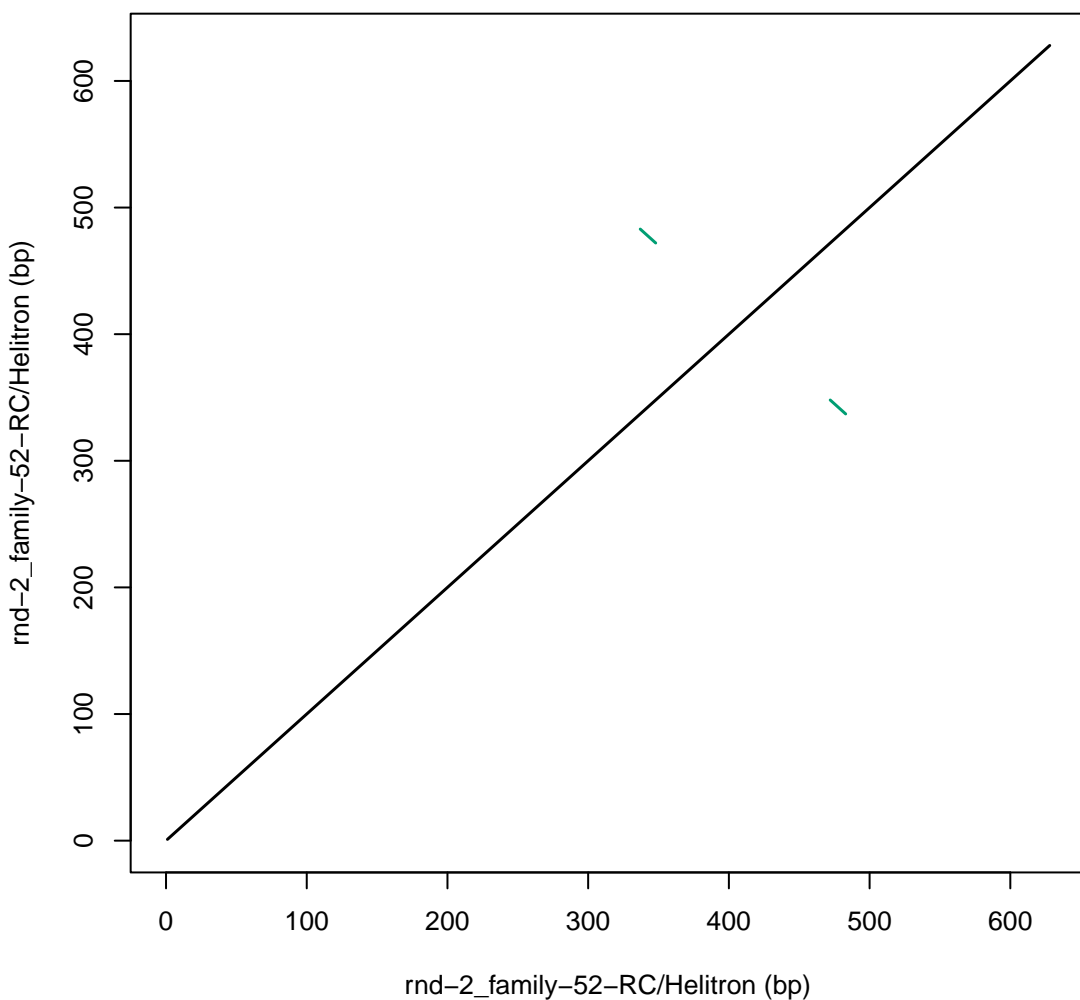
TE: rnd-2\_family-52-RC/Helitron  
consensus size: 628bp; fragments: 1035; full length: 1 (>=565.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

