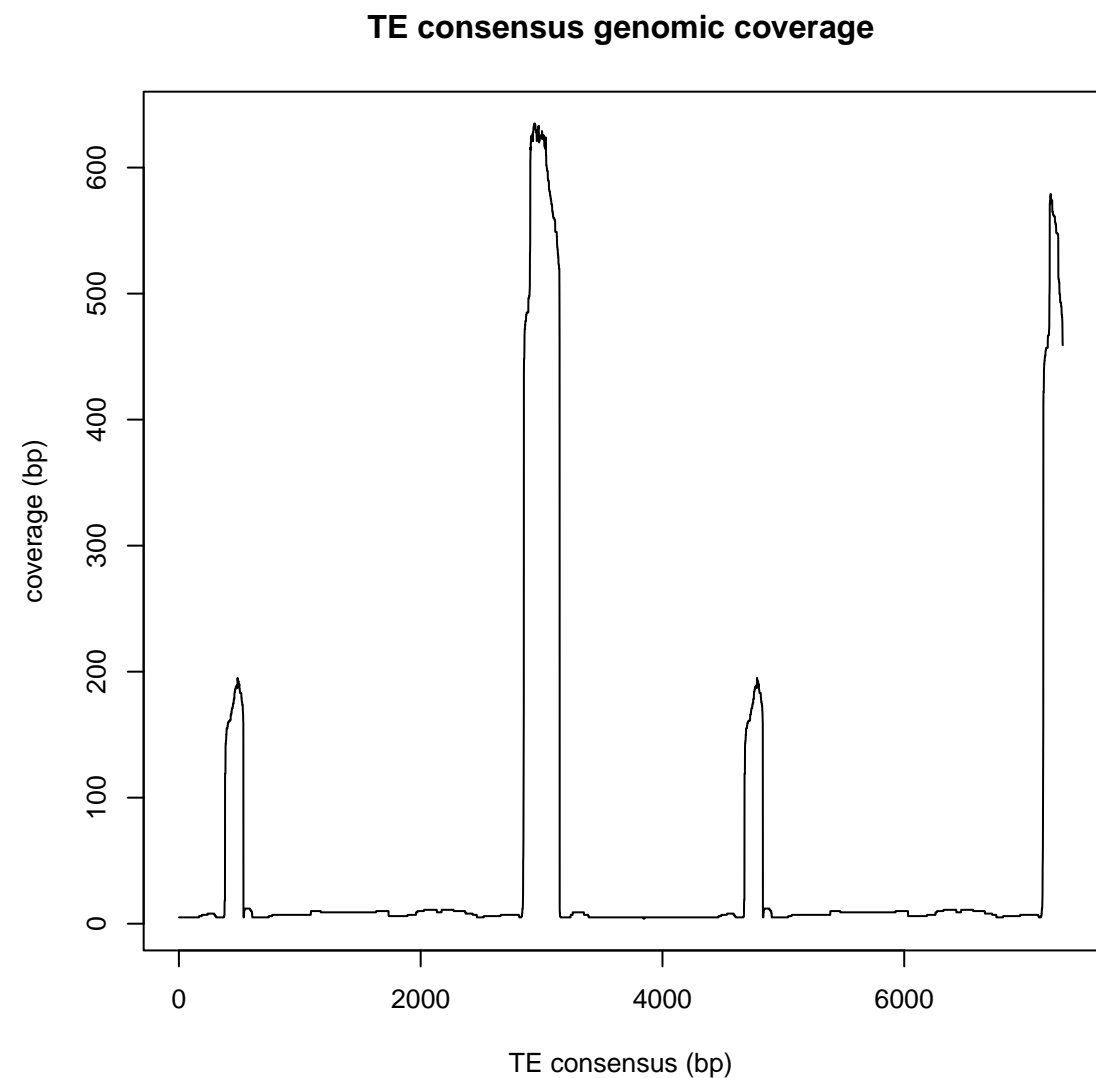
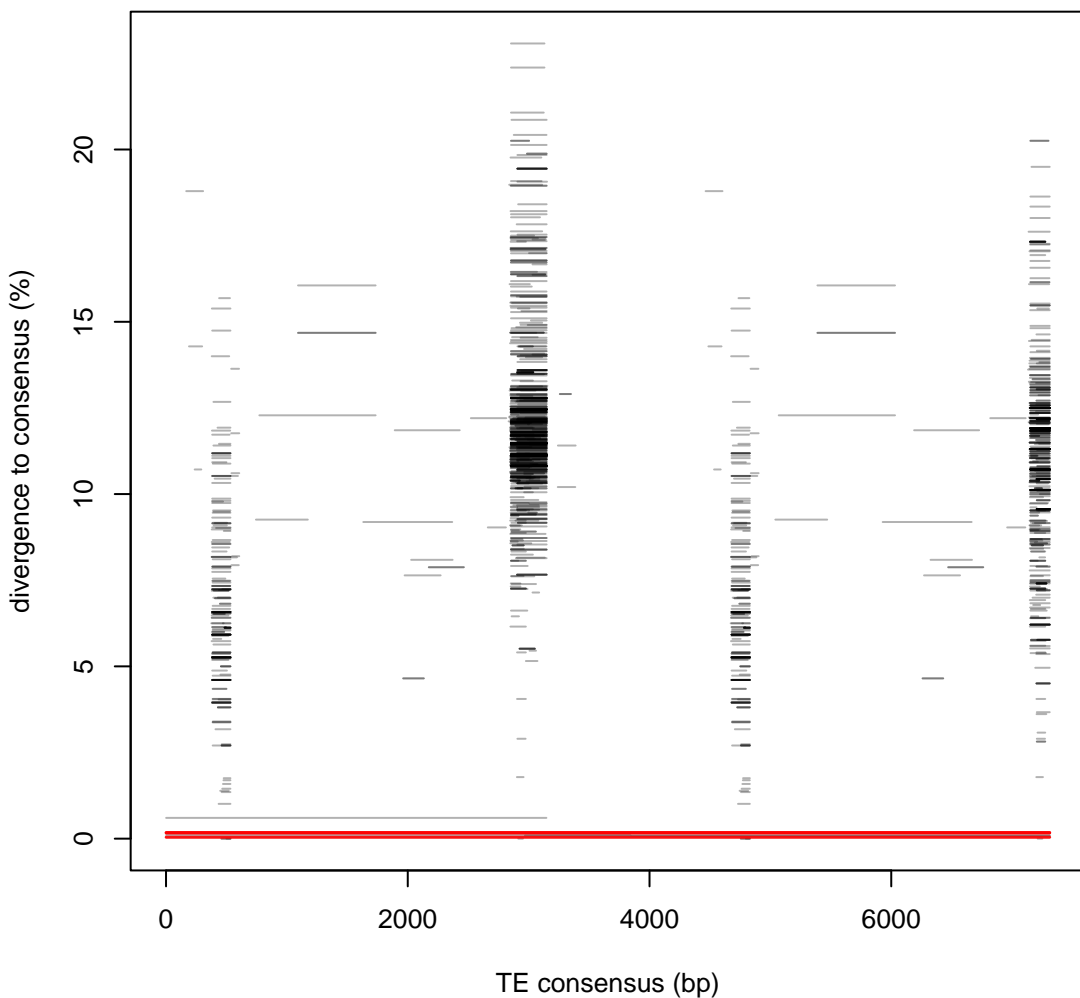
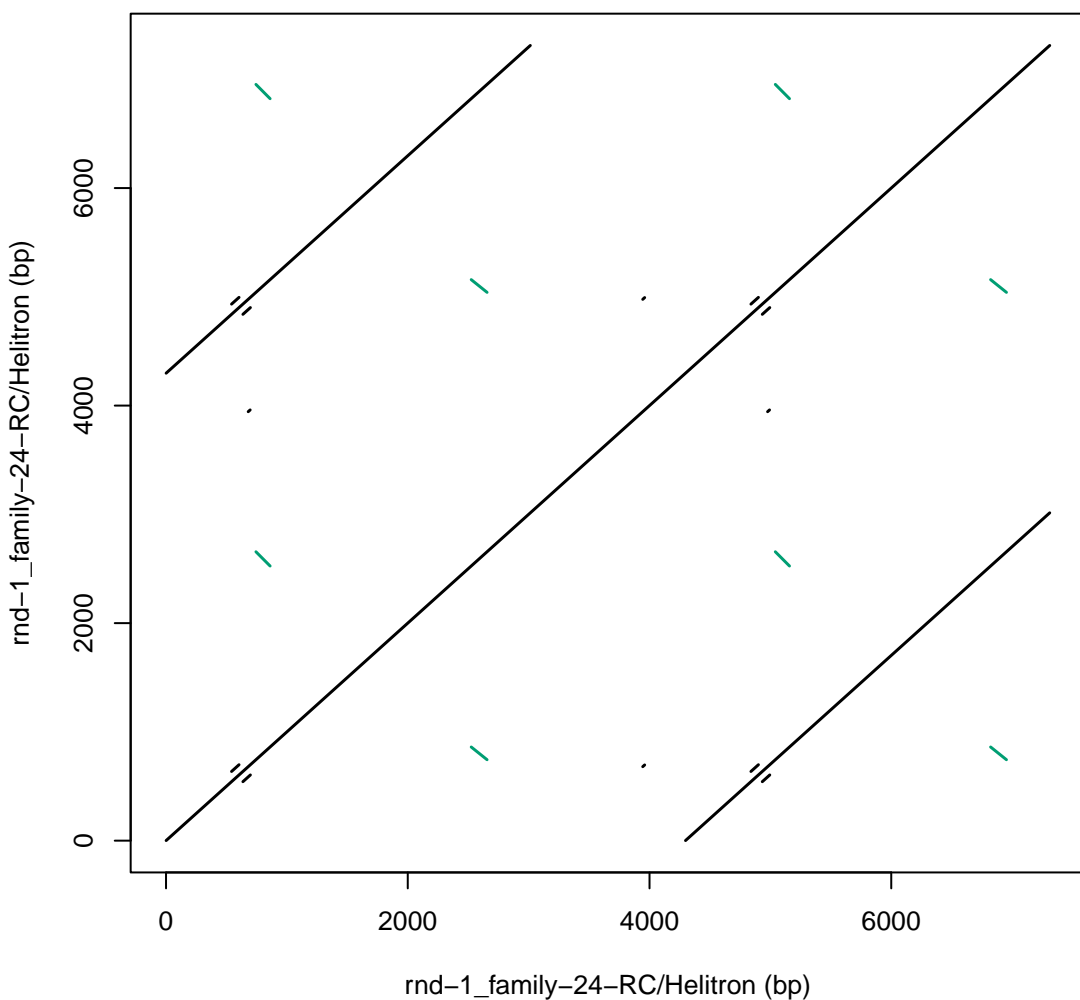


TE: rnd-1_family-24-RC/Helitron
consensus size: 7311bp; fragments: 1738; full length: 3 (>=6579.9bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

