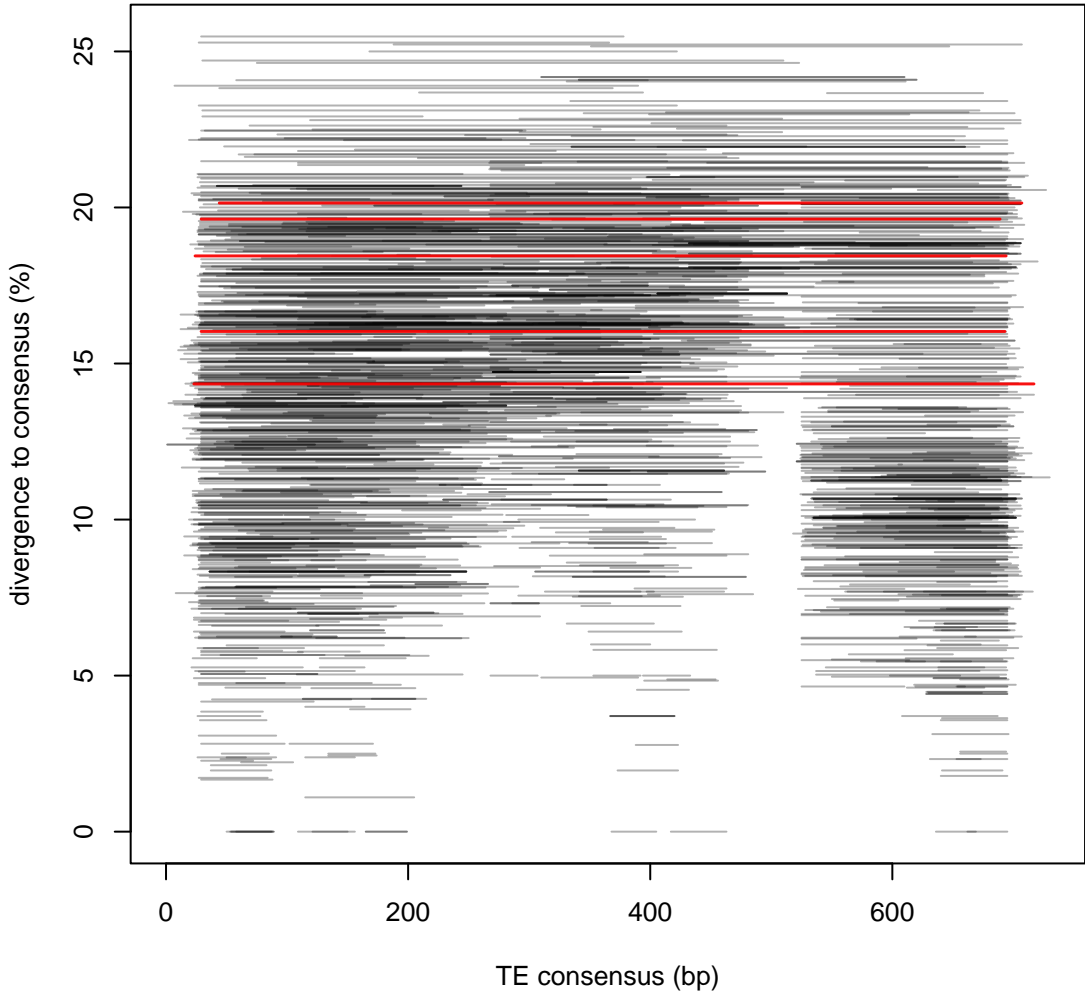
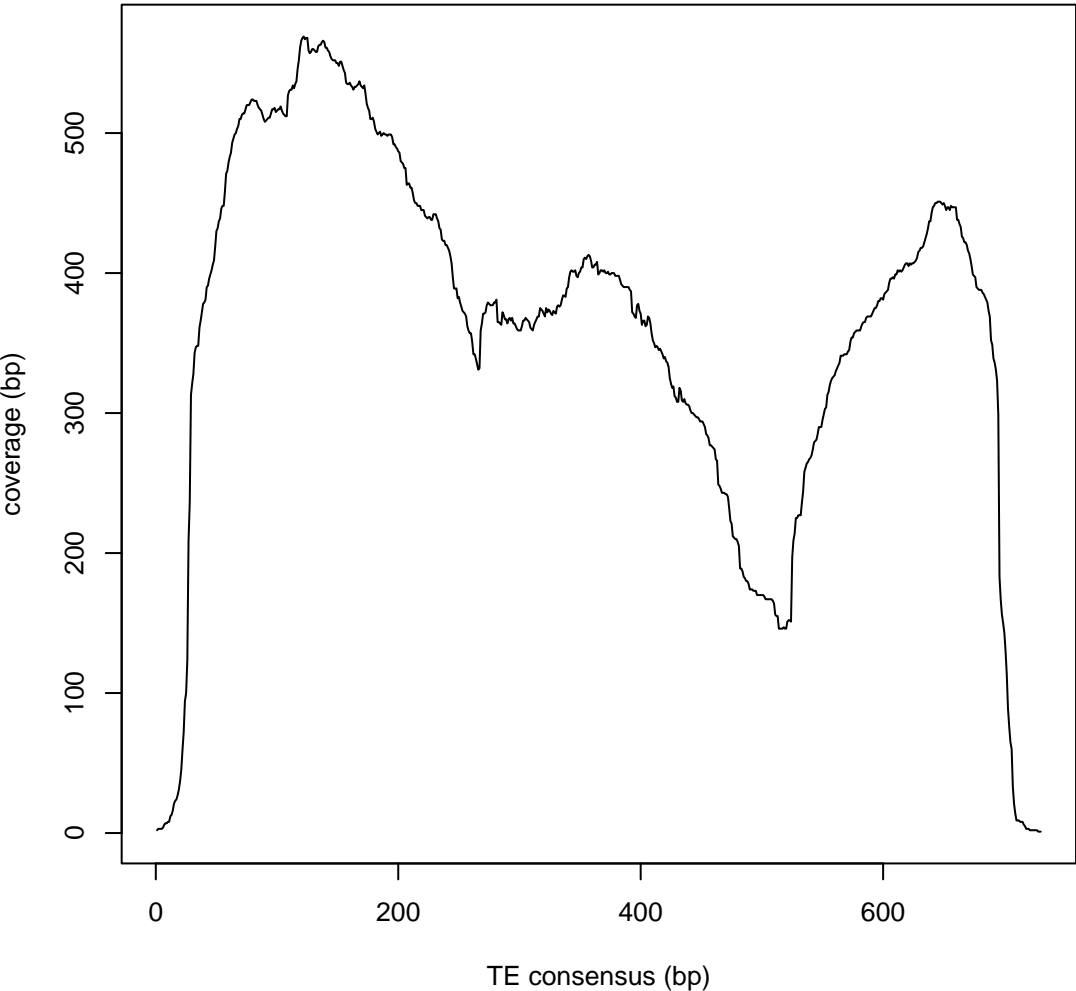


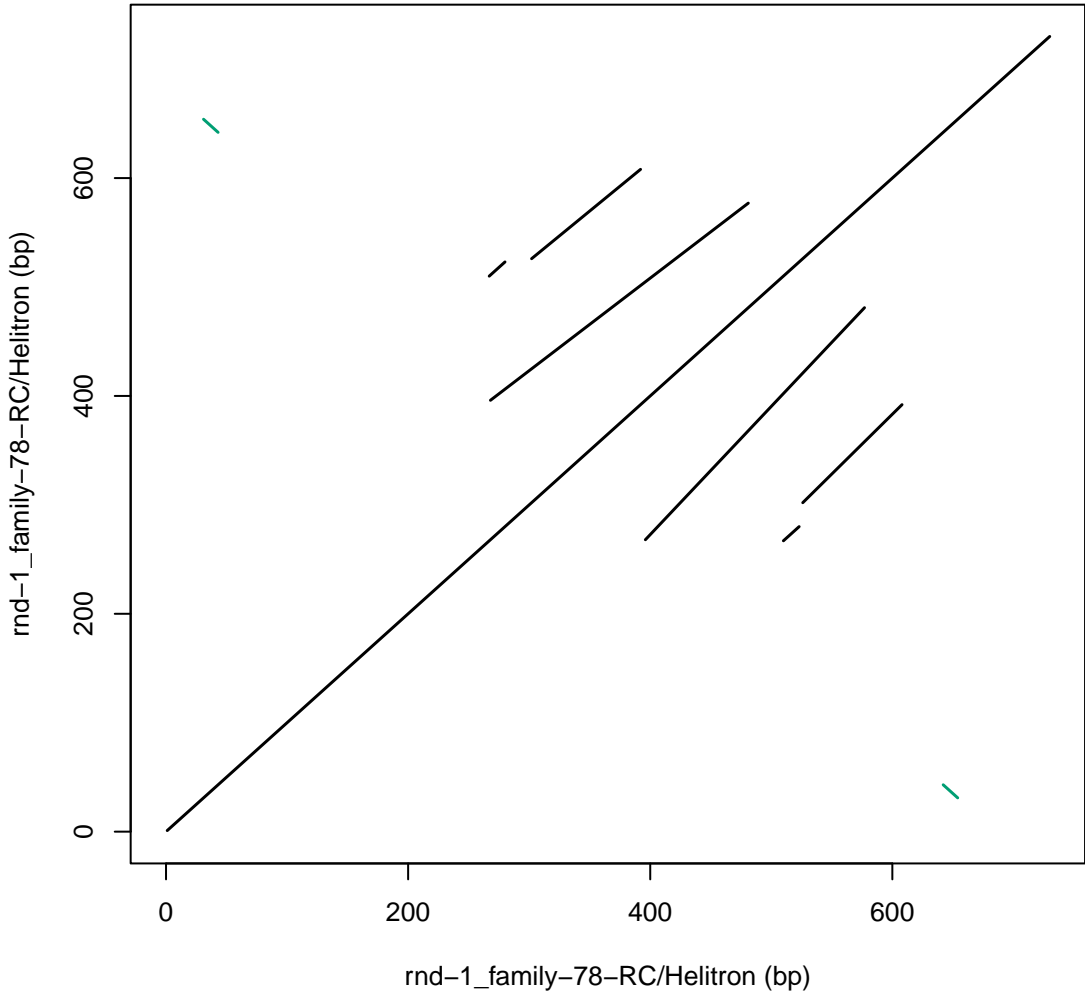
TE: rnd-1_family-78-RC/Helitron
consensus size: 730bp; fragments: 1402; full length: 5 (>=657bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

