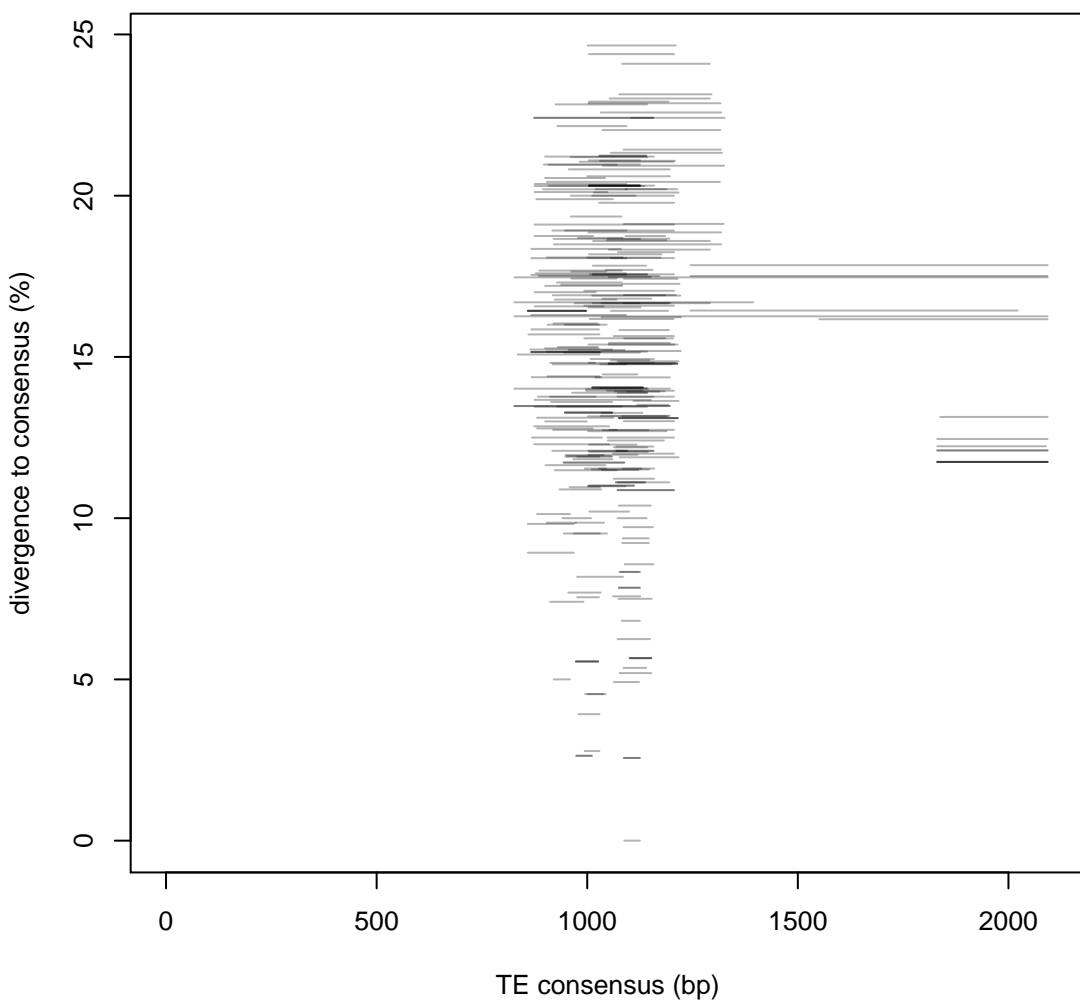
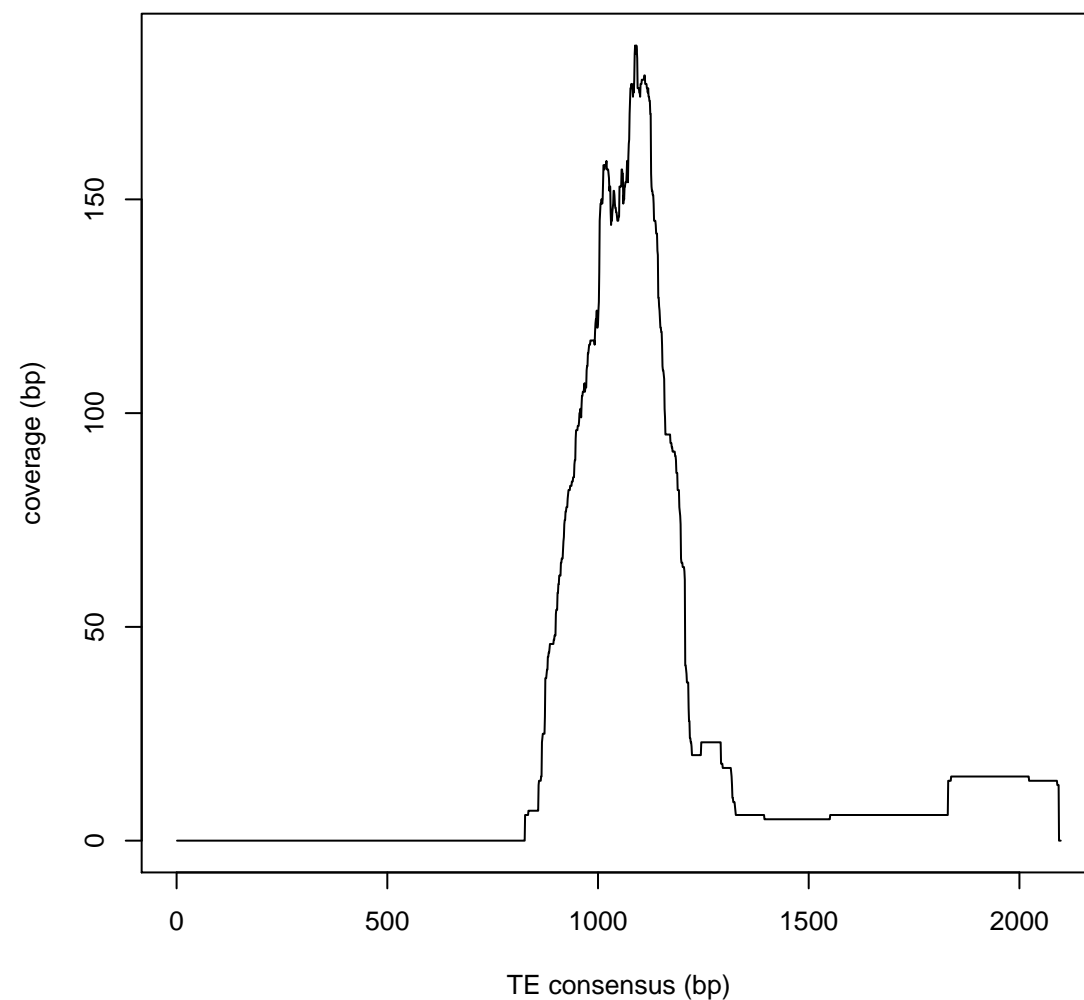


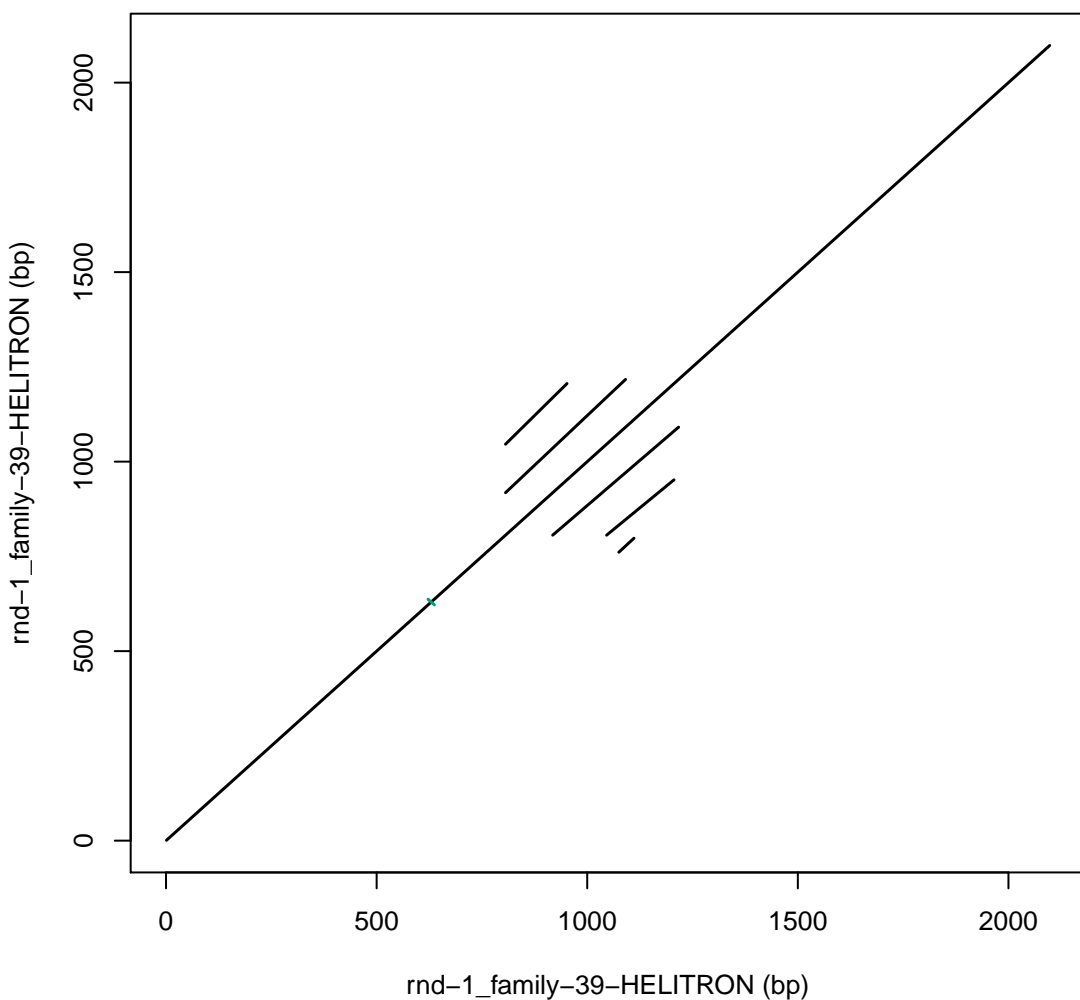
TE: rnd-1\_family-39-HELITRON  
consensus size: 2098bp; fragments: 291; full length: 0 (>=1888.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

