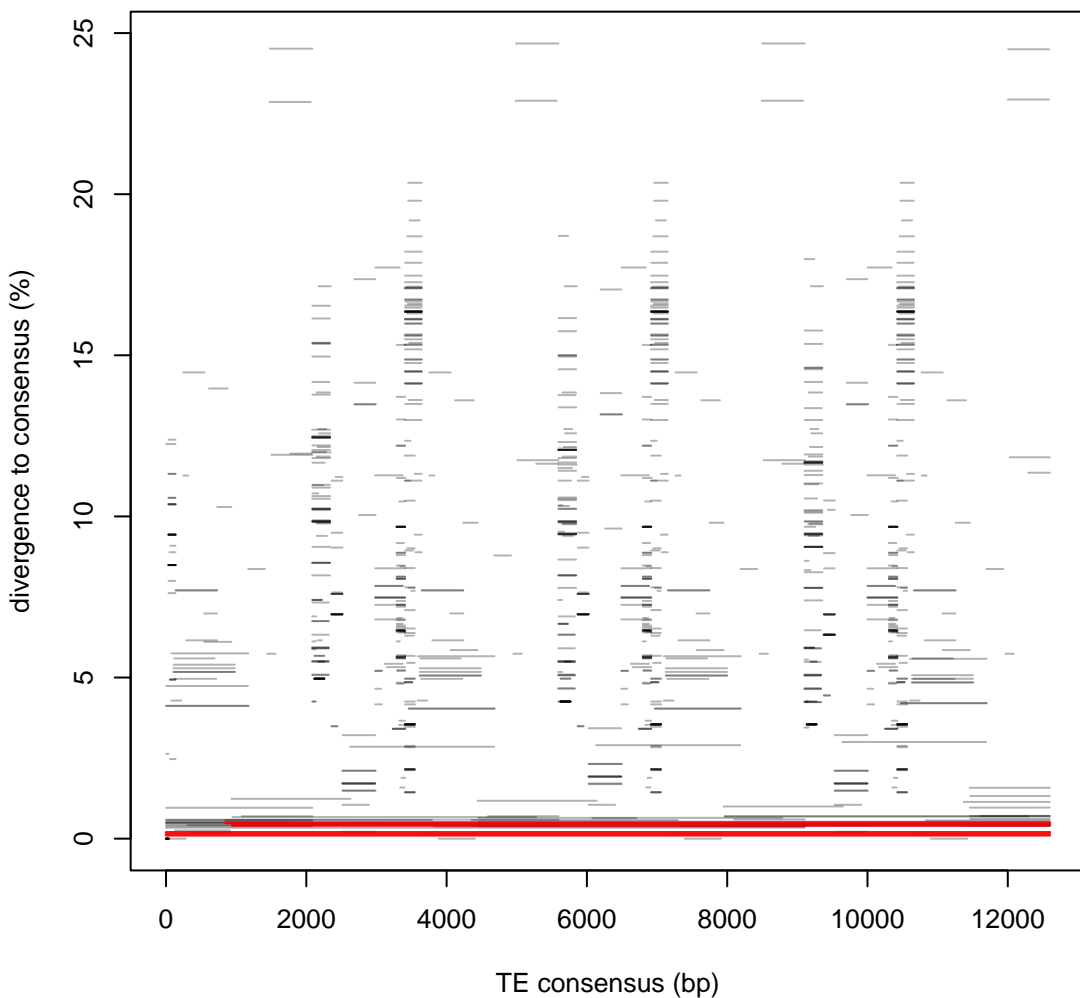
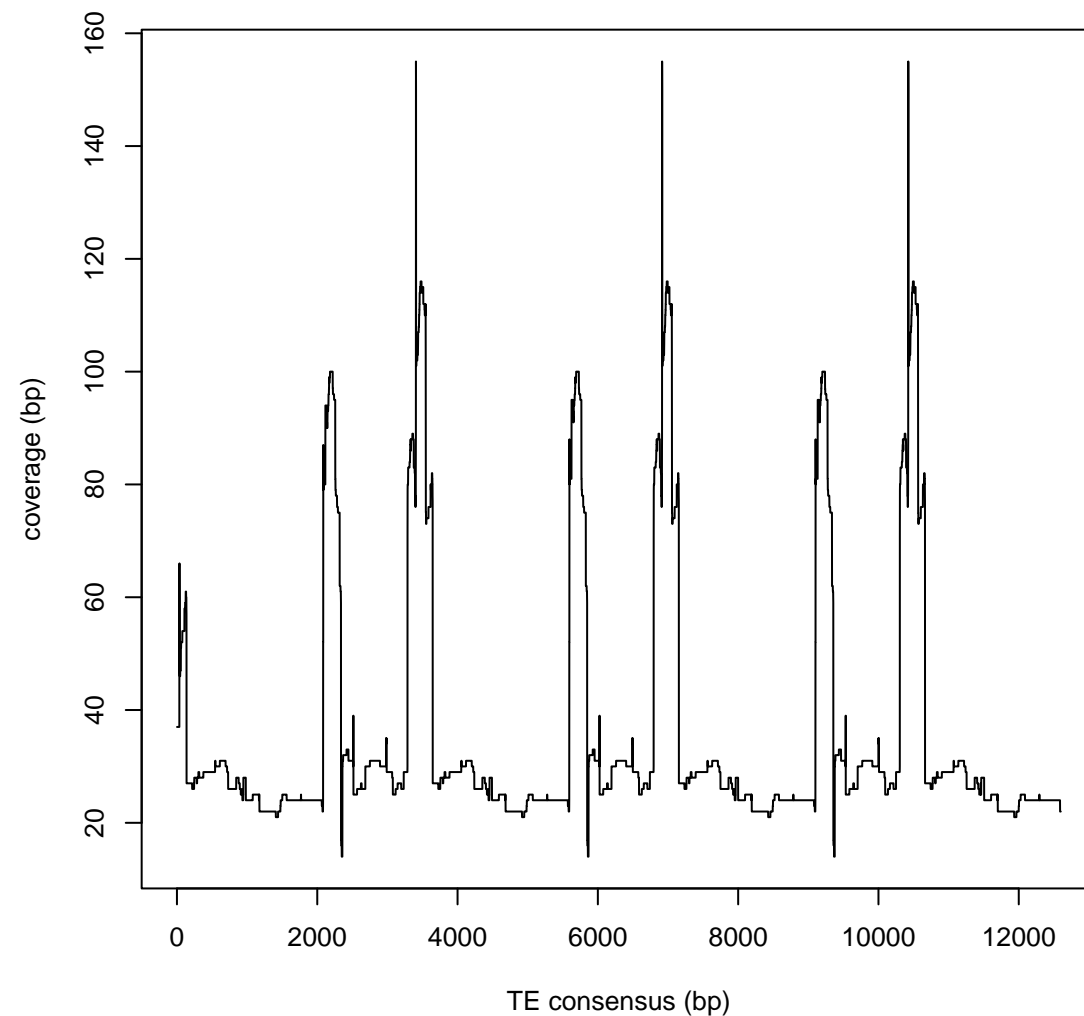


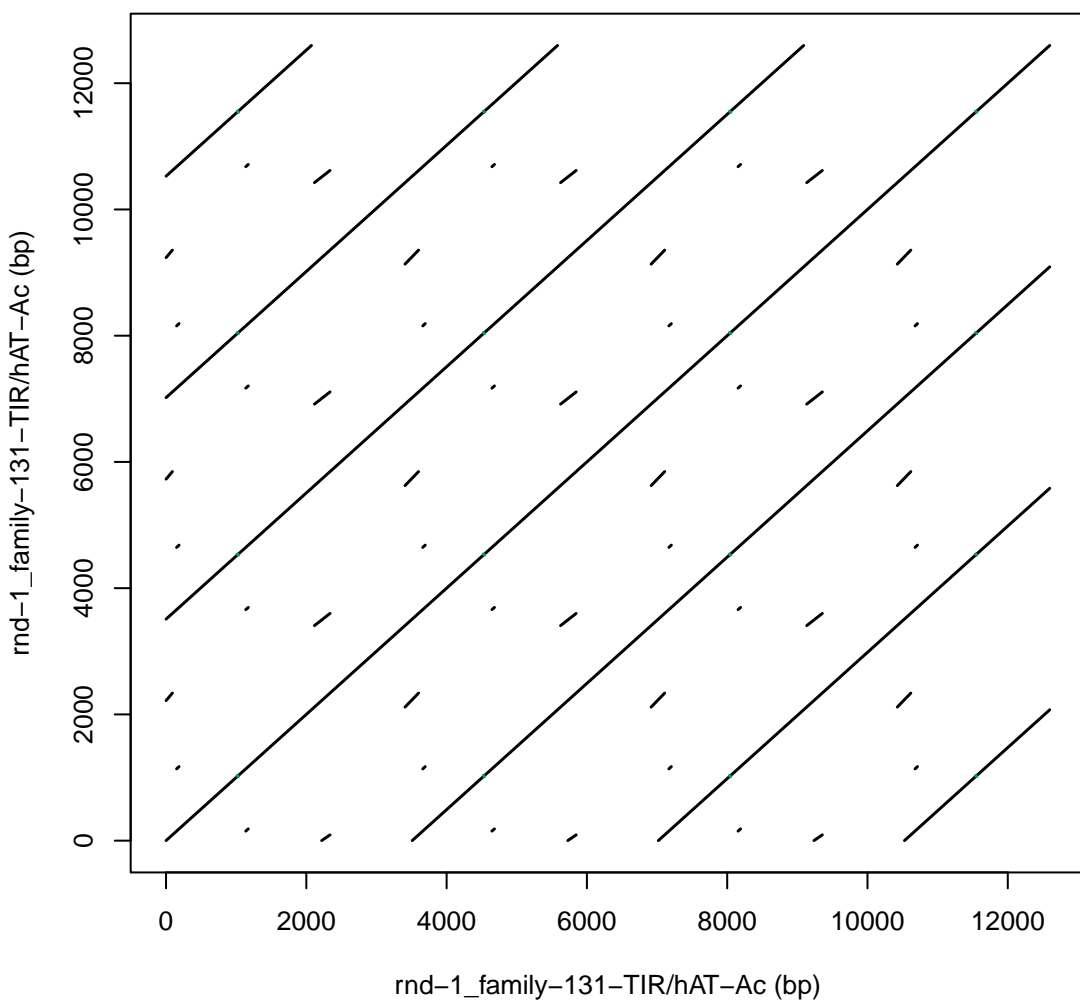
TE: rnd-1_family-131-TIR/hAT-Ac
consensus size: 12598bp; fragments: 1156; full length: 4 (>=11338.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

