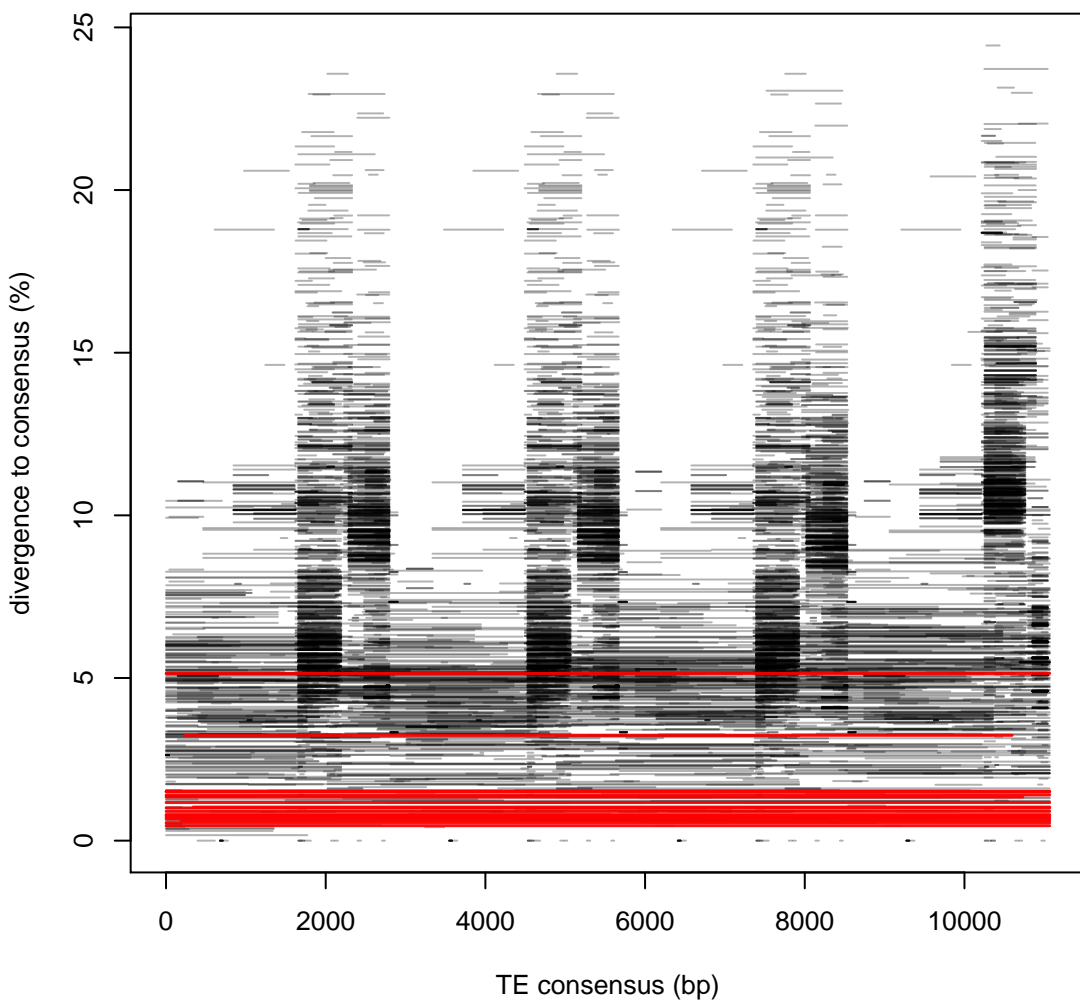
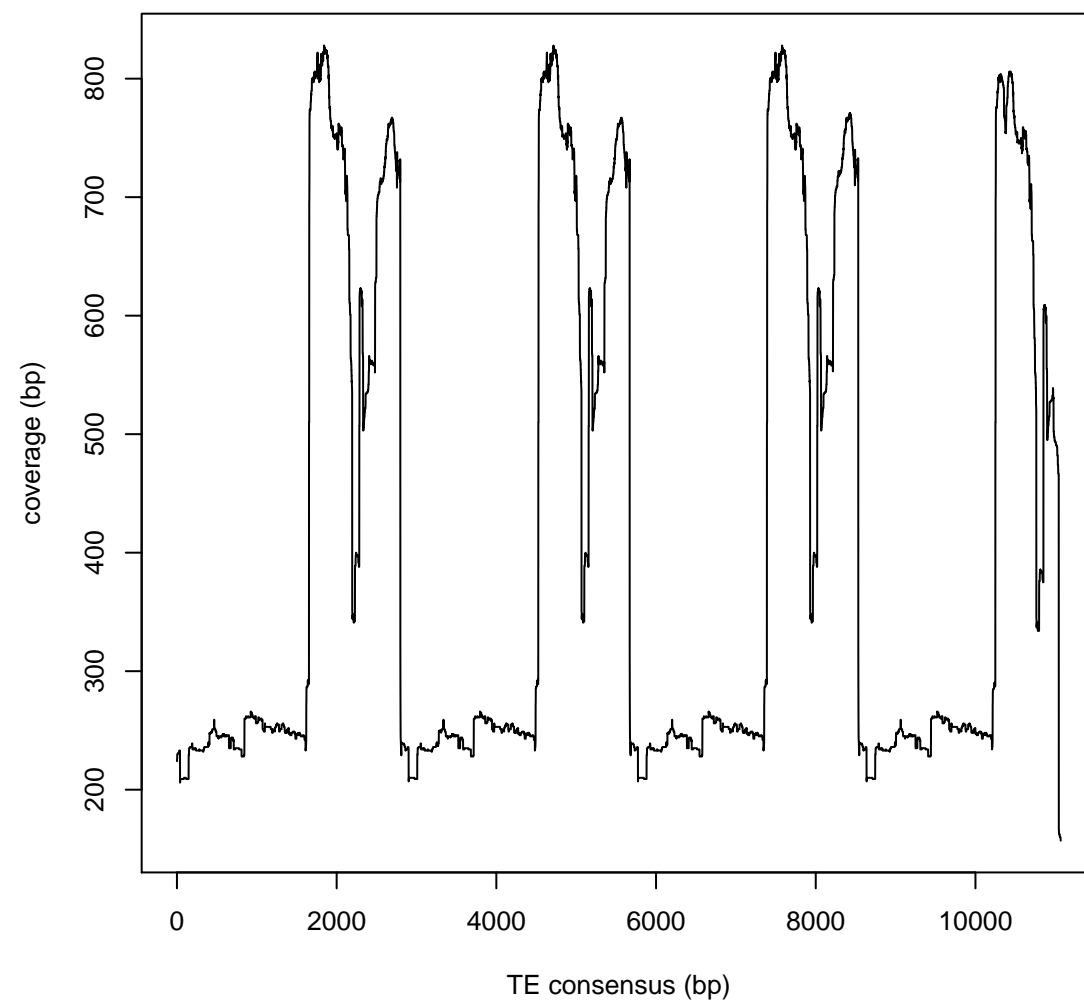


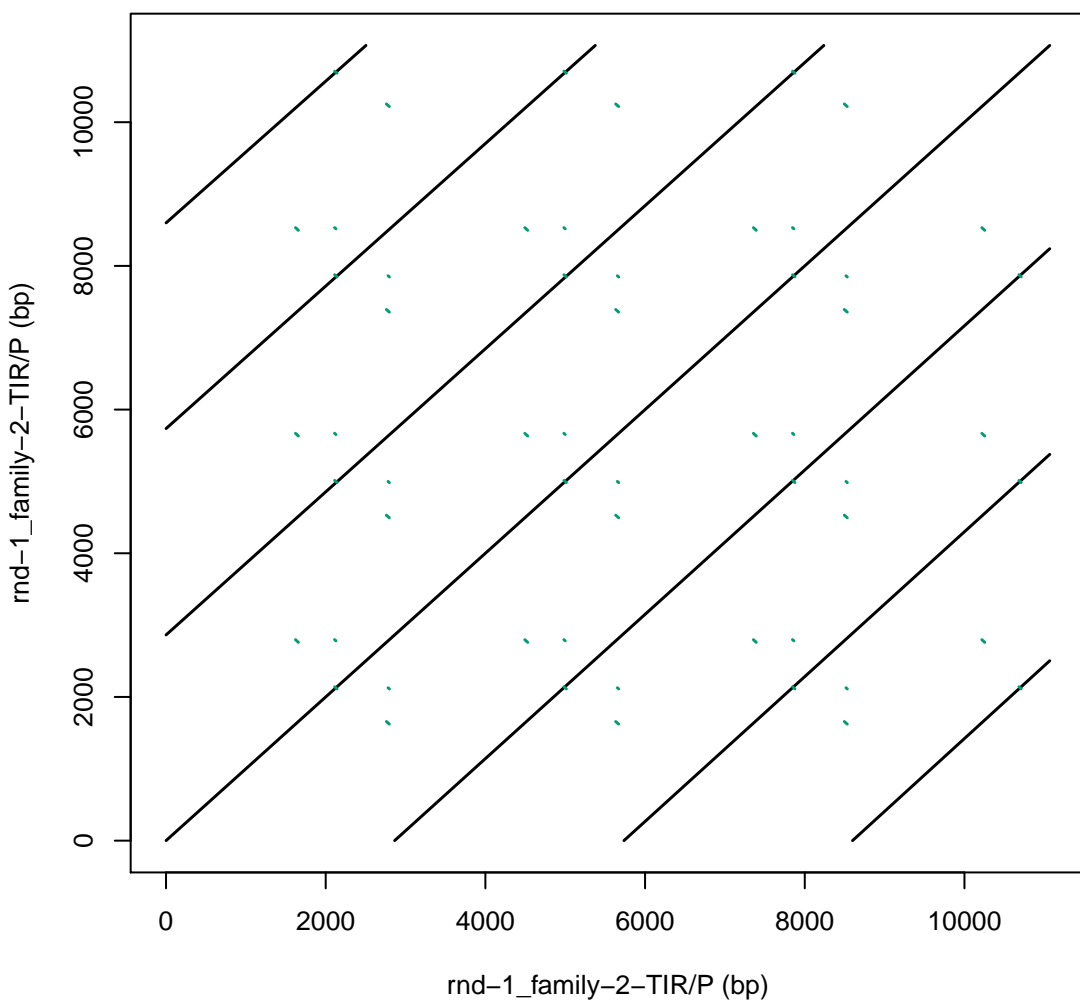
TE: rnd-1_family-2-TIR/P
consensus size: 11068bp; fragments: 7108; full length: 17 (>=9961.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

