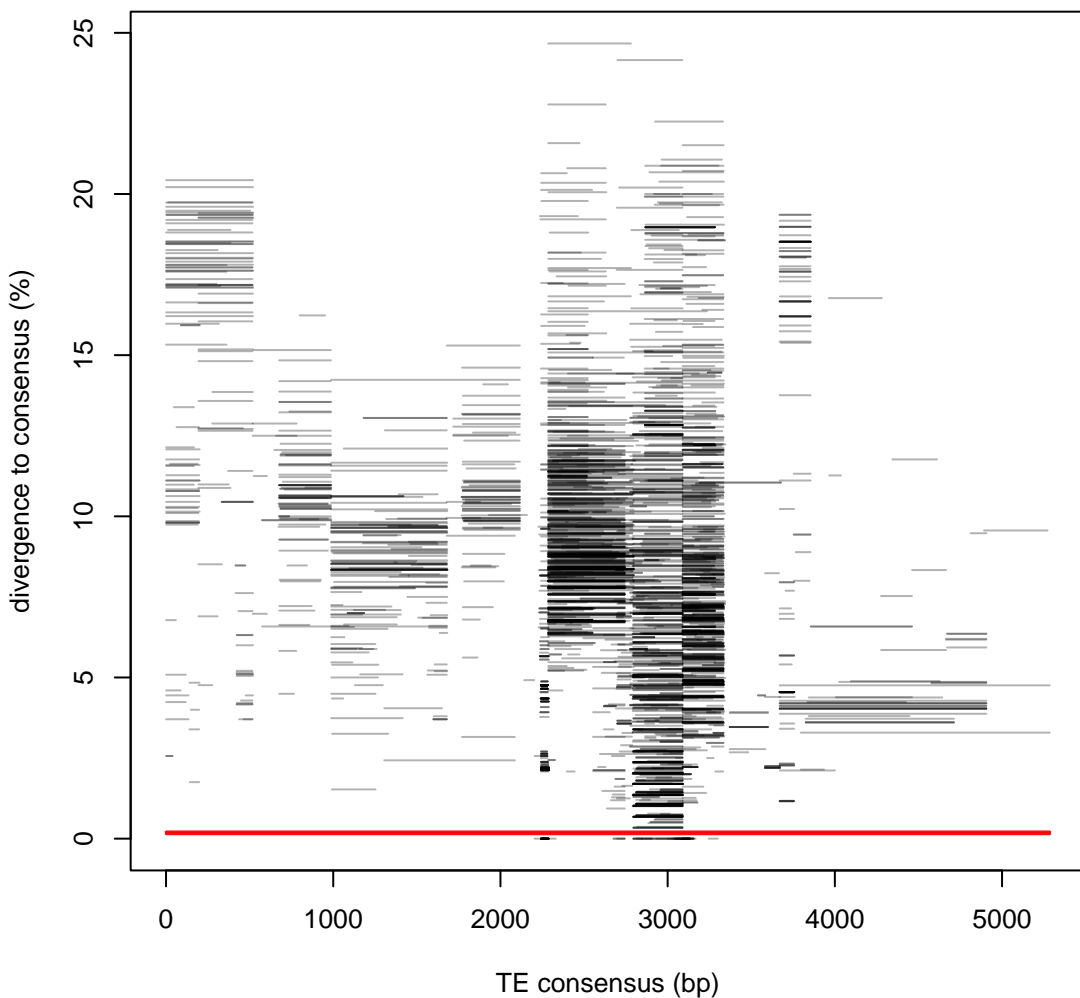
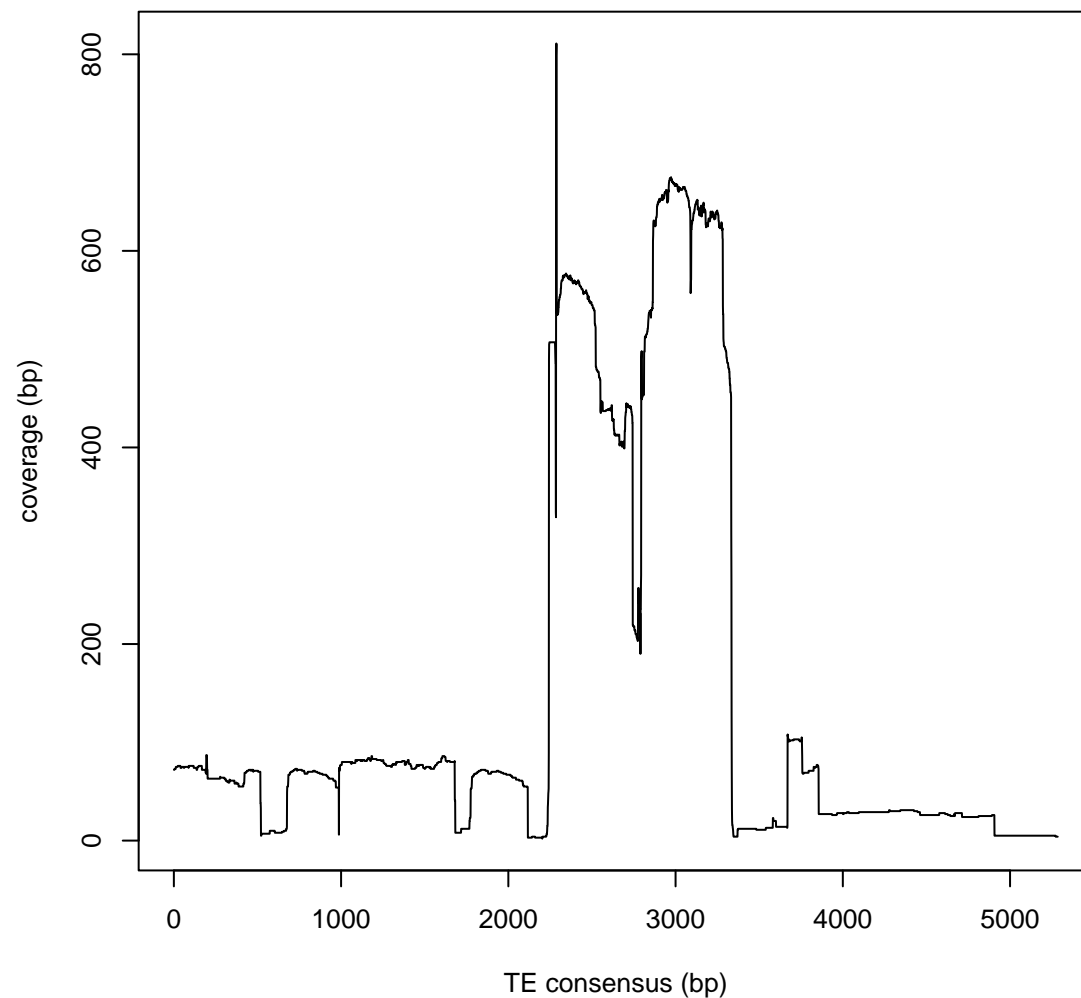


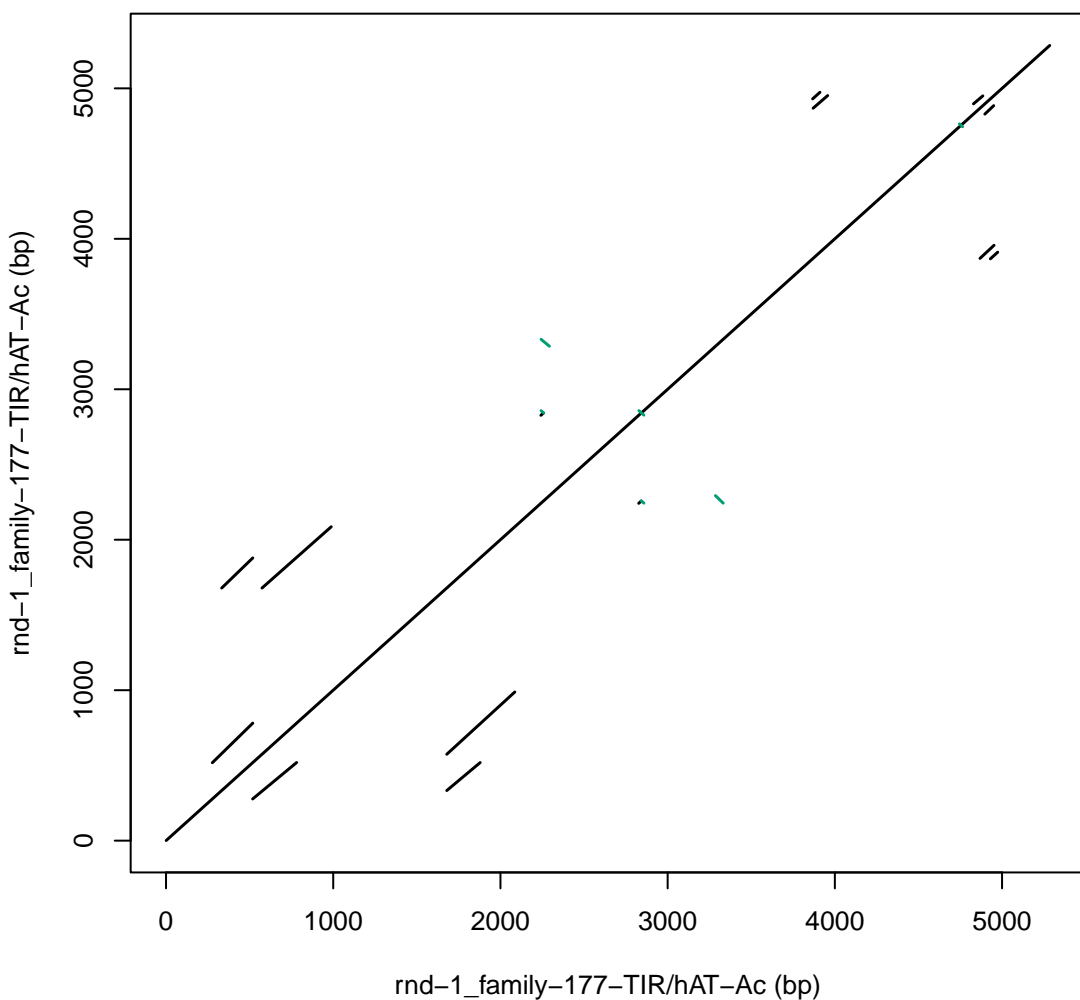
TE: rnd-1_family-177-TIR/hAT-Ac
consensus size: 5285bp; fragments: 3163; full length: 2 (>=4756.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

