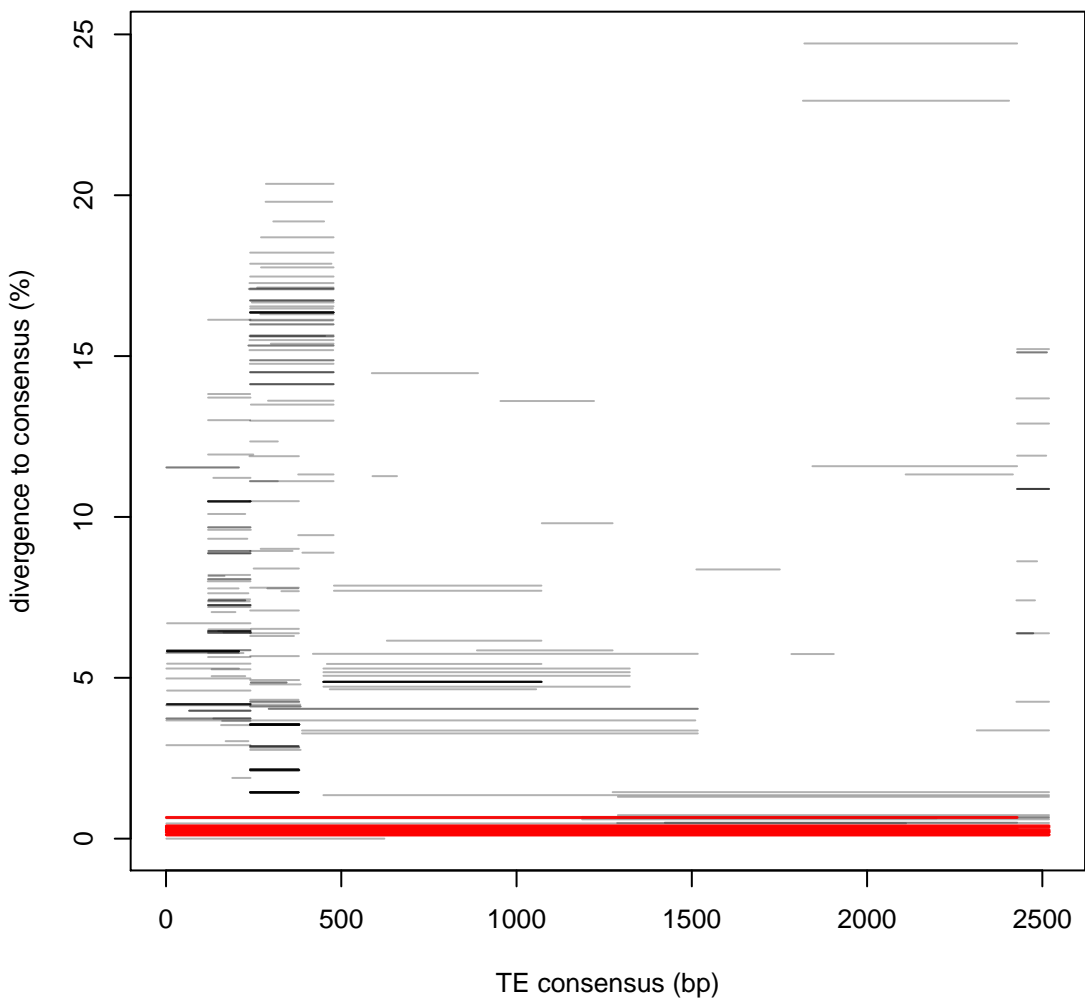
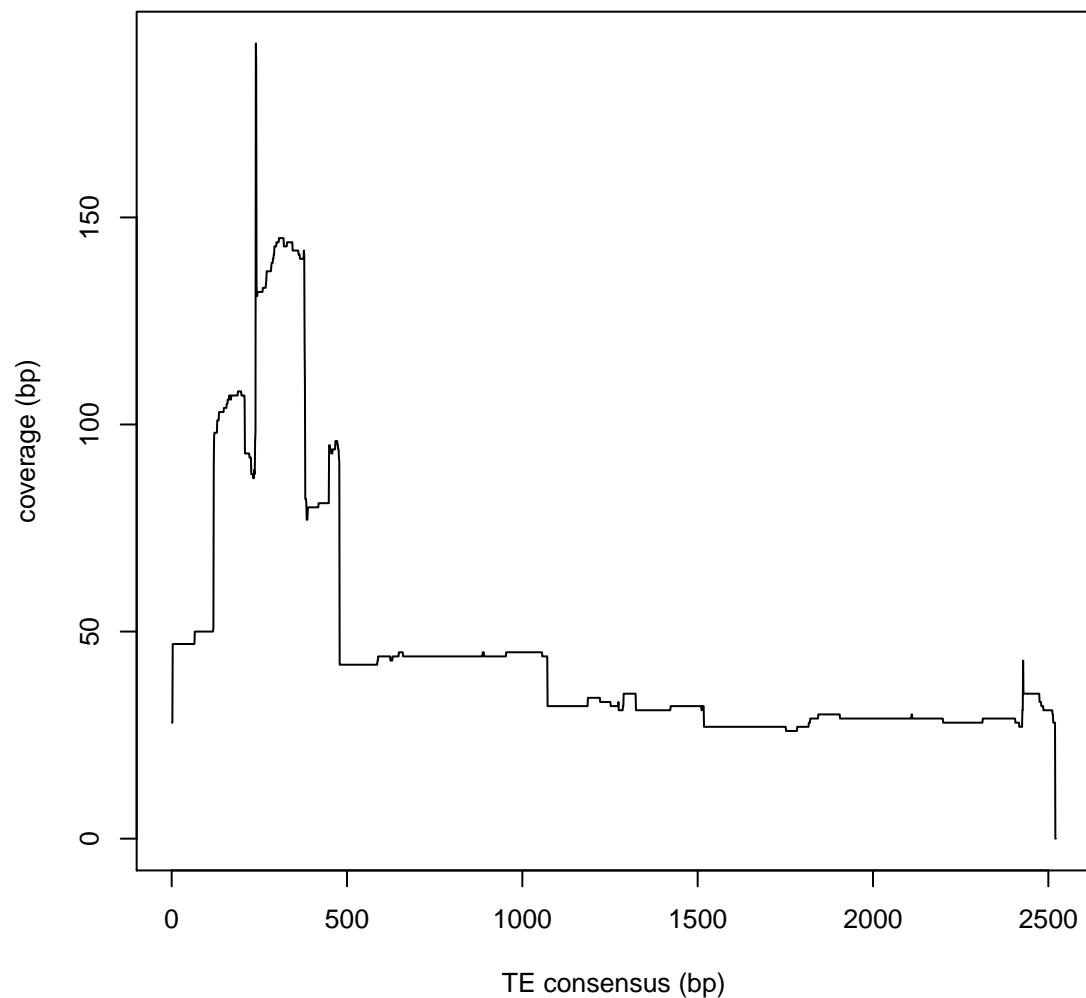


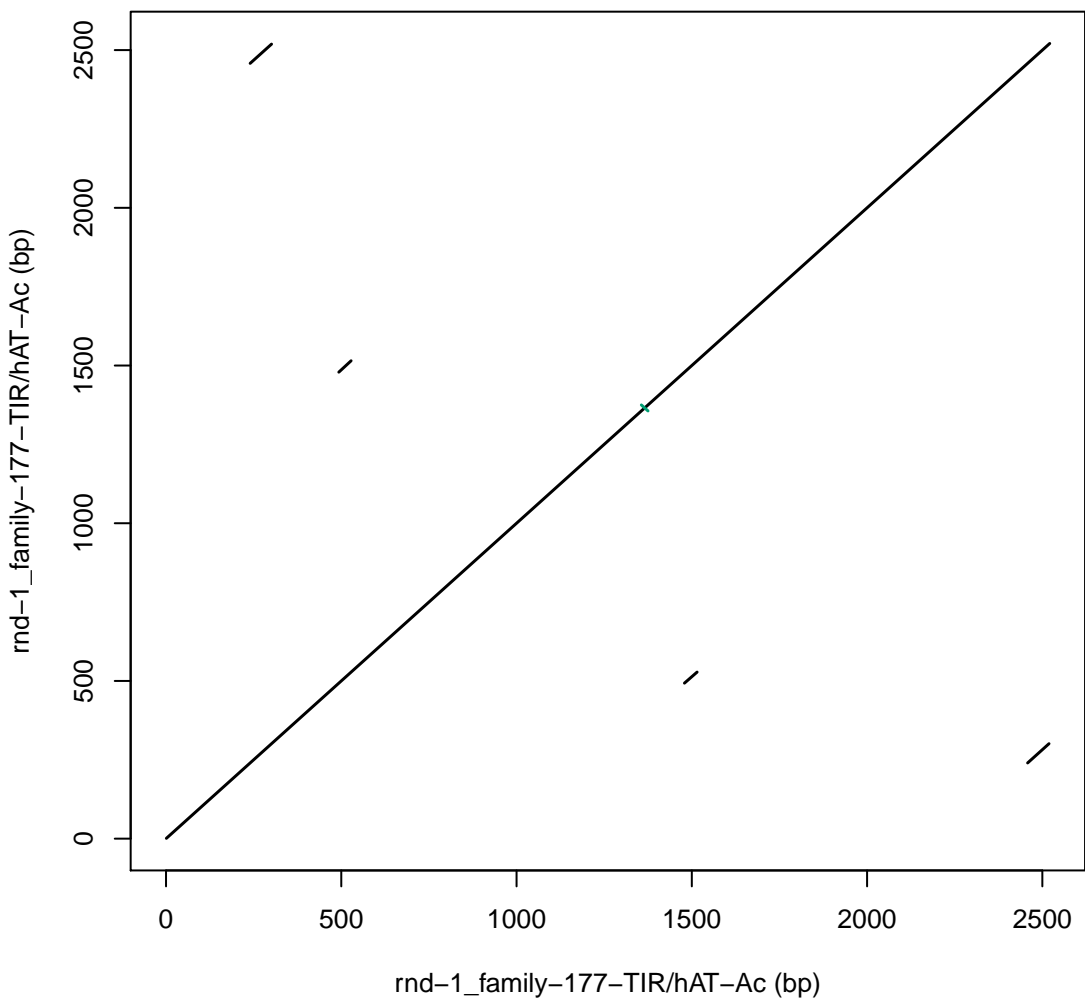
TE: rnd-1_family-177-TIR/hAT-Ac
consensus size: 2521bp; fragments: 300; full length: 14 (>=2268.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

