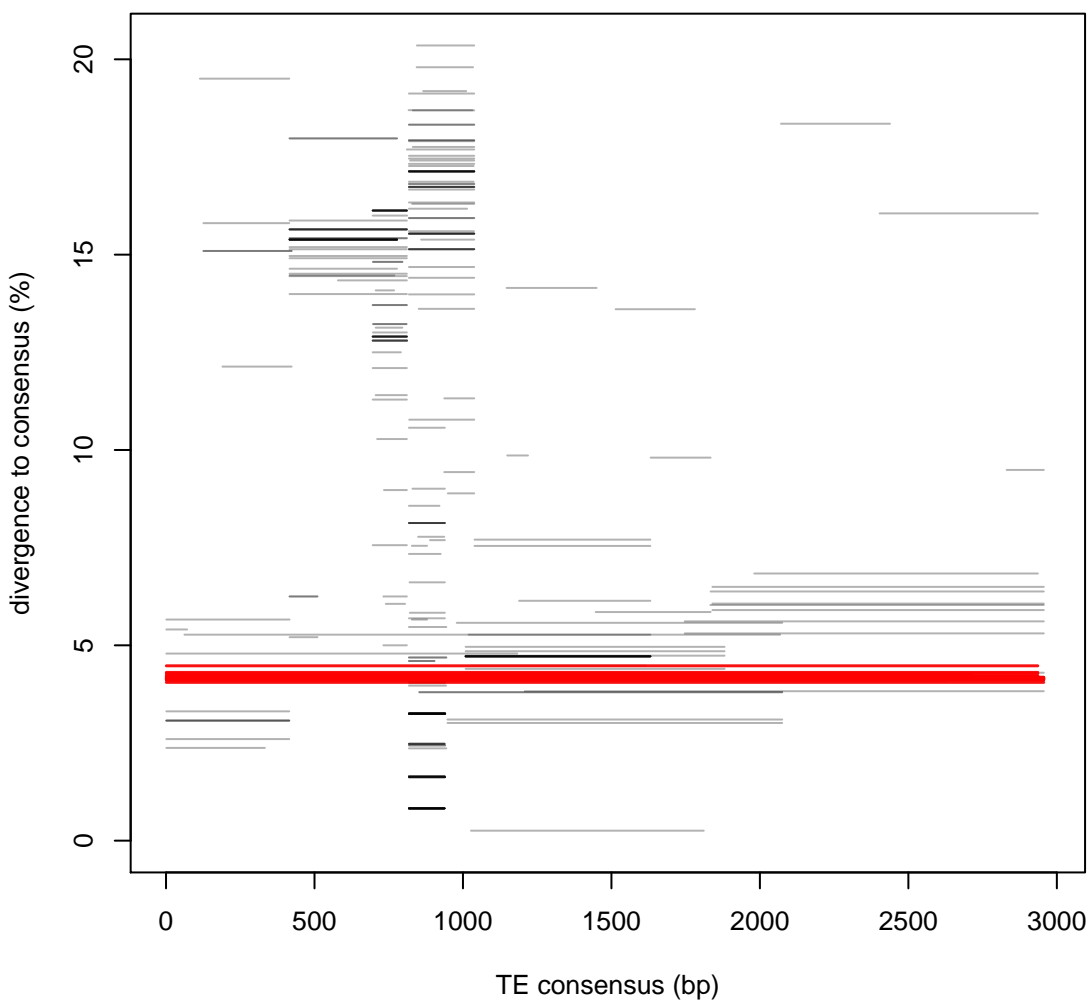
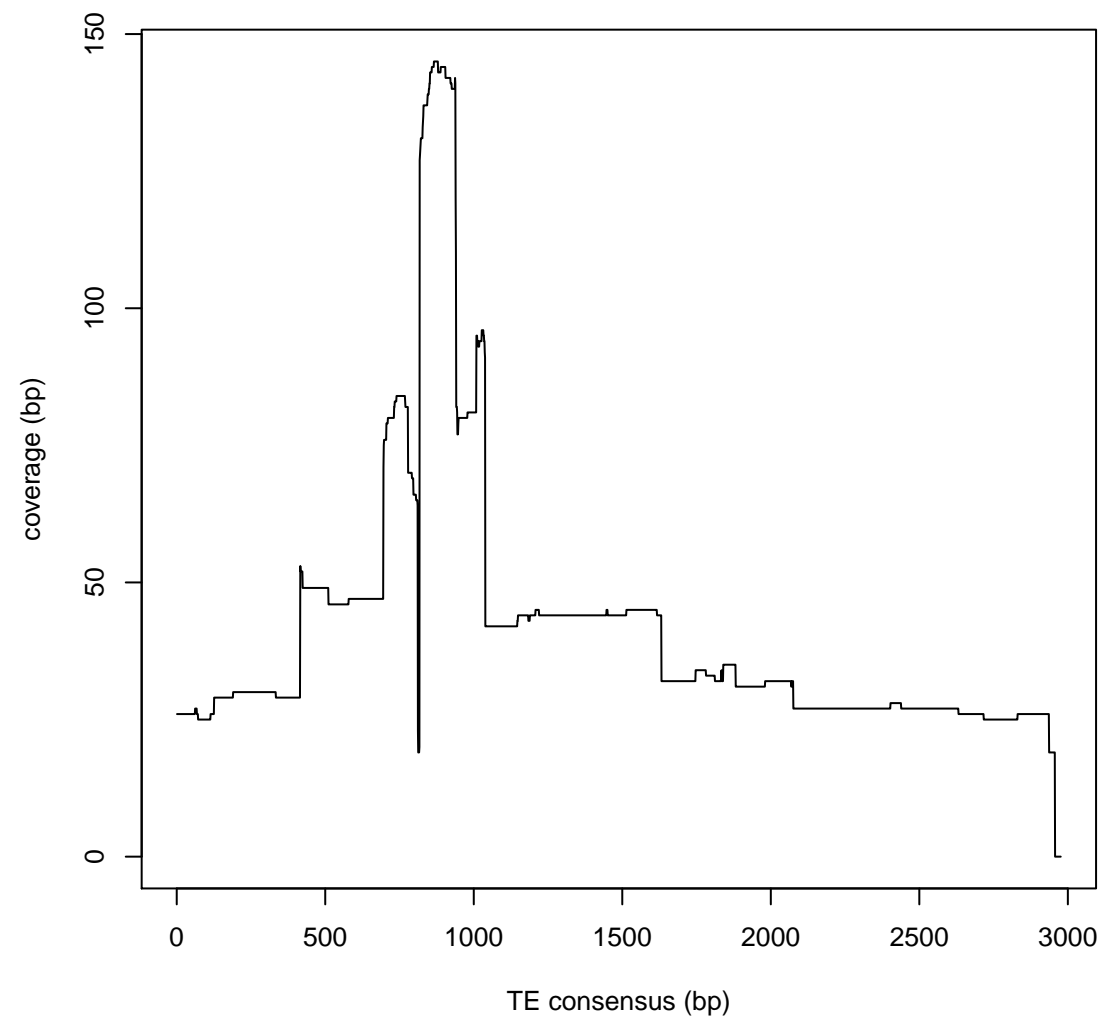


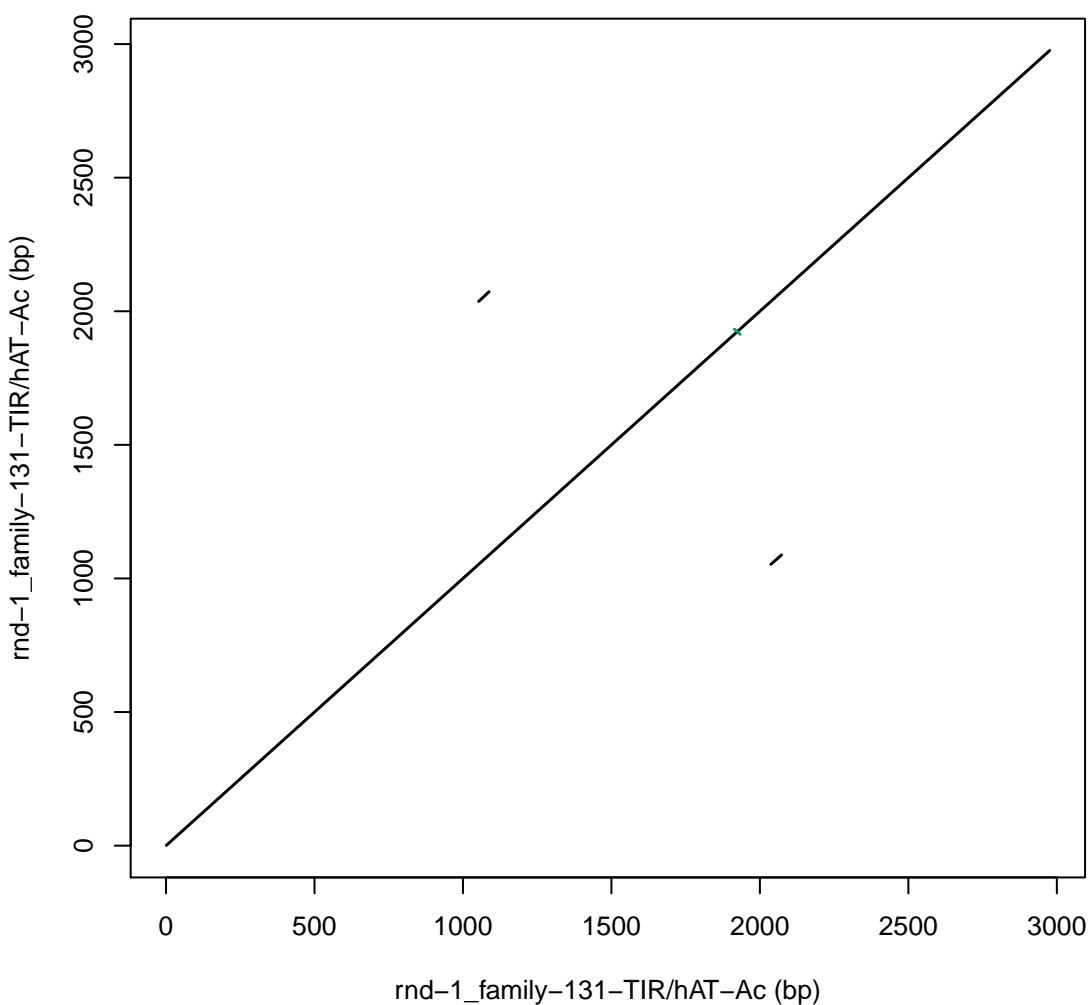
TE: rnd-1\_family-131-TIR/hAT-Ac  
consensus size: 2976bp; fragments: 272; full length: 15 (>=2678.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

