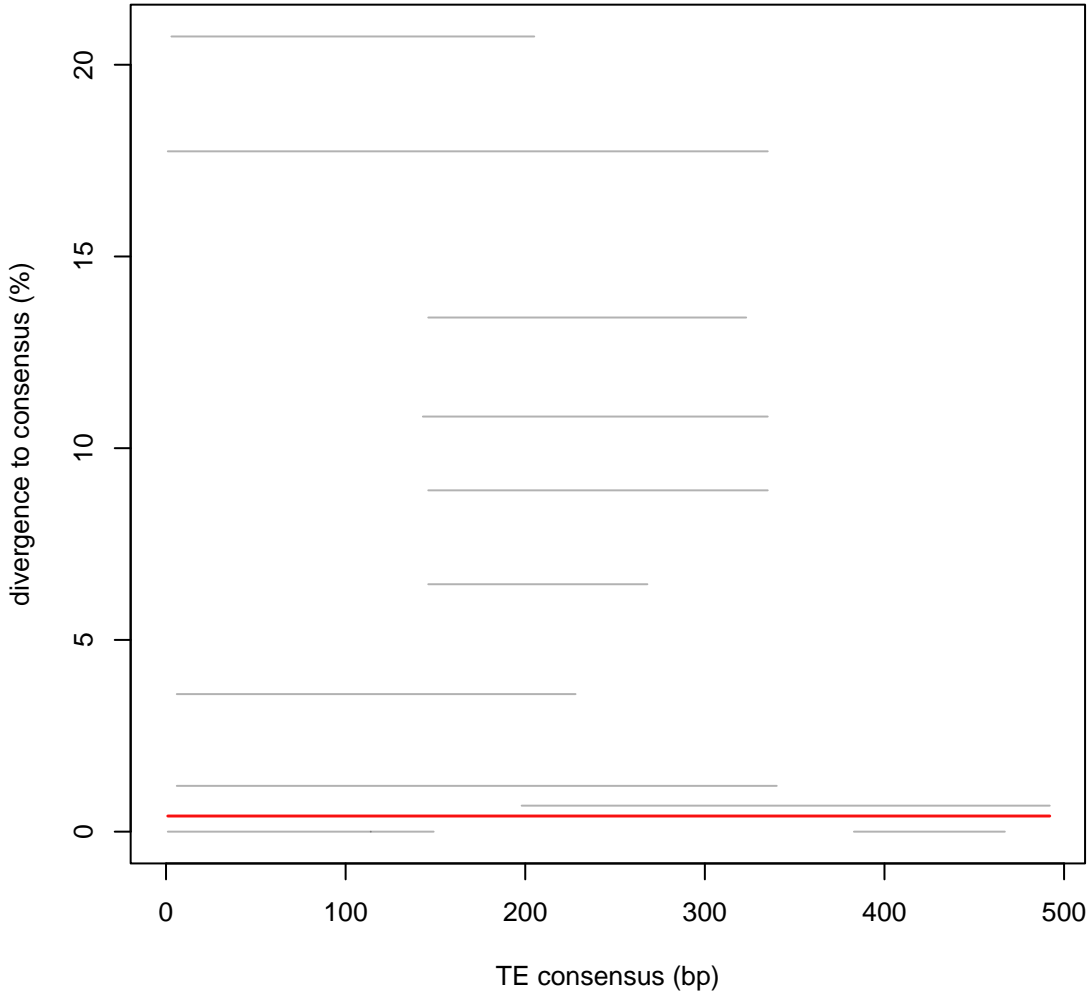
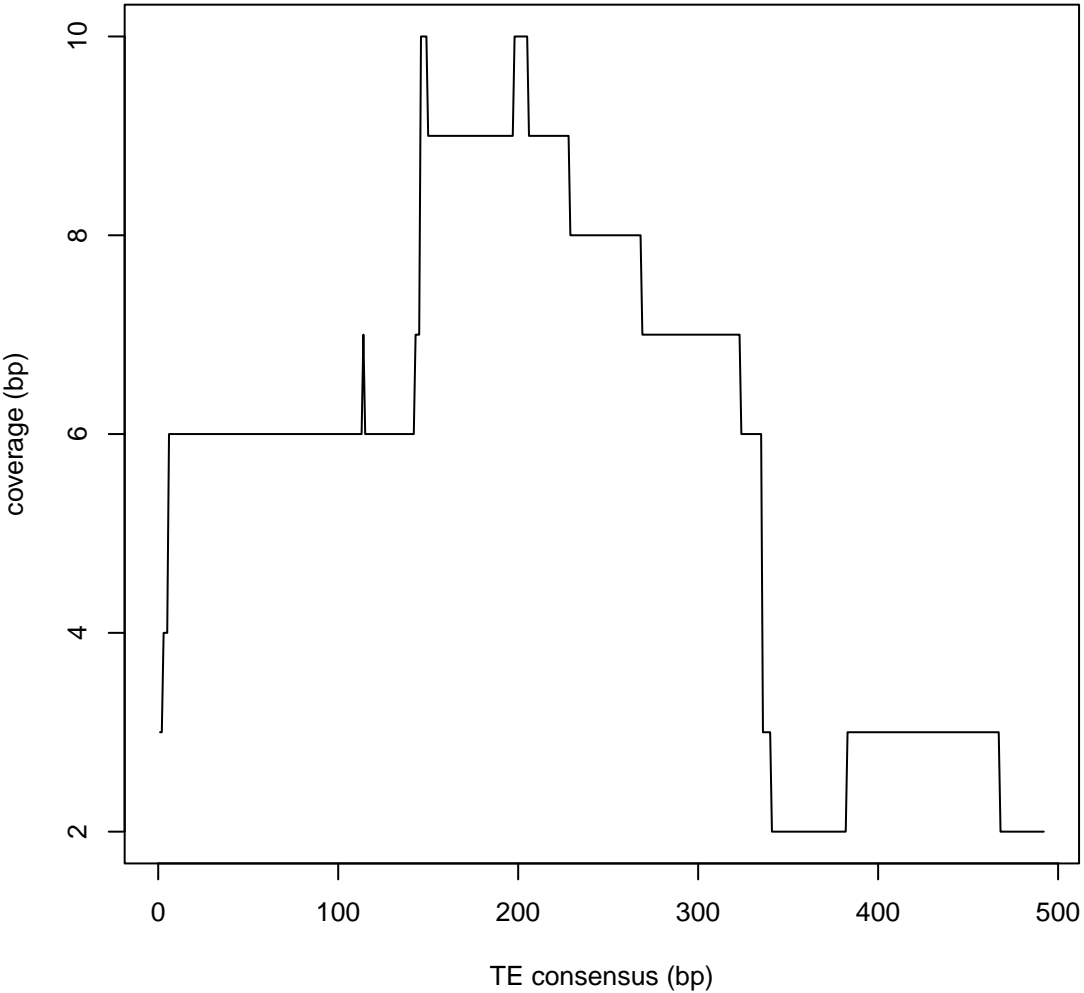


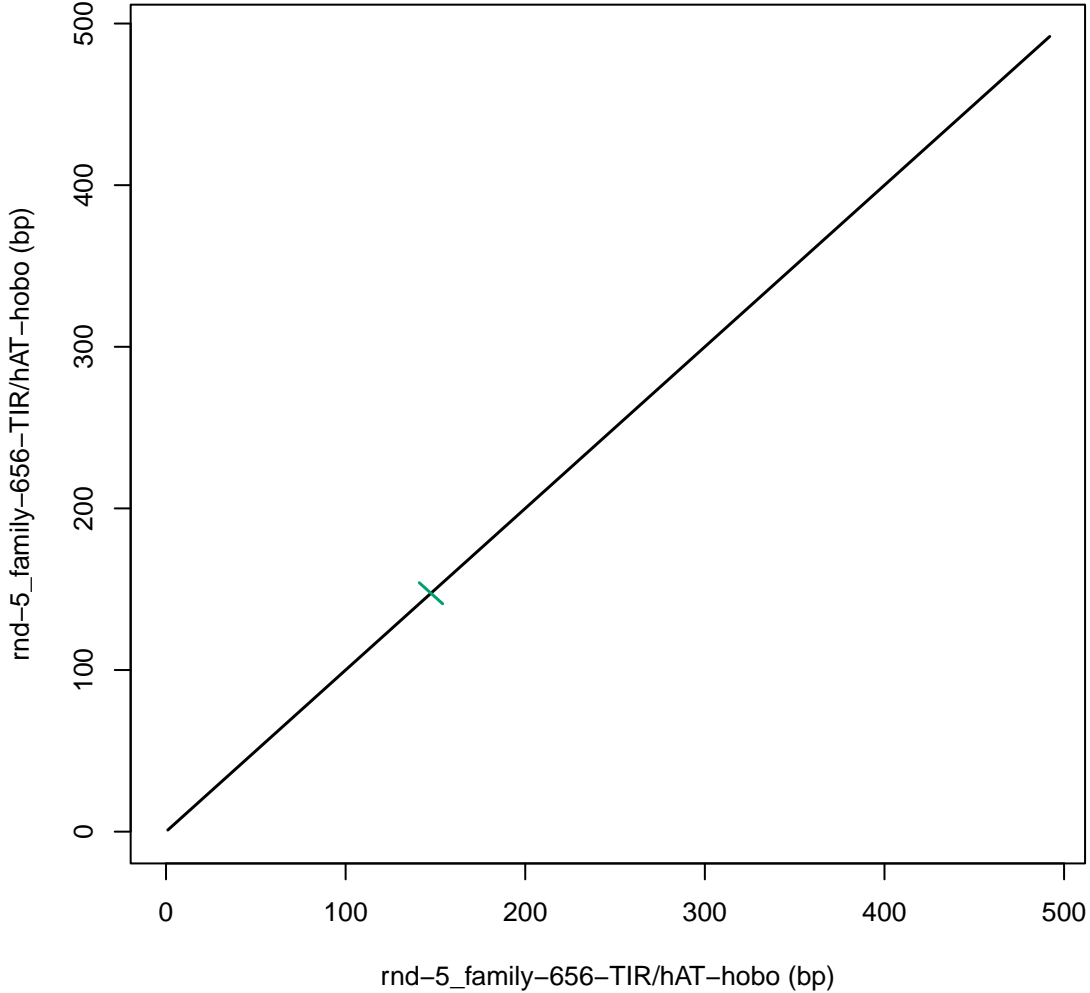
TE: rnd-5_family-656-TIR/hAT-hobo
consensus size: 492bp; fragments: 13; full length: 1 (>=442.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

