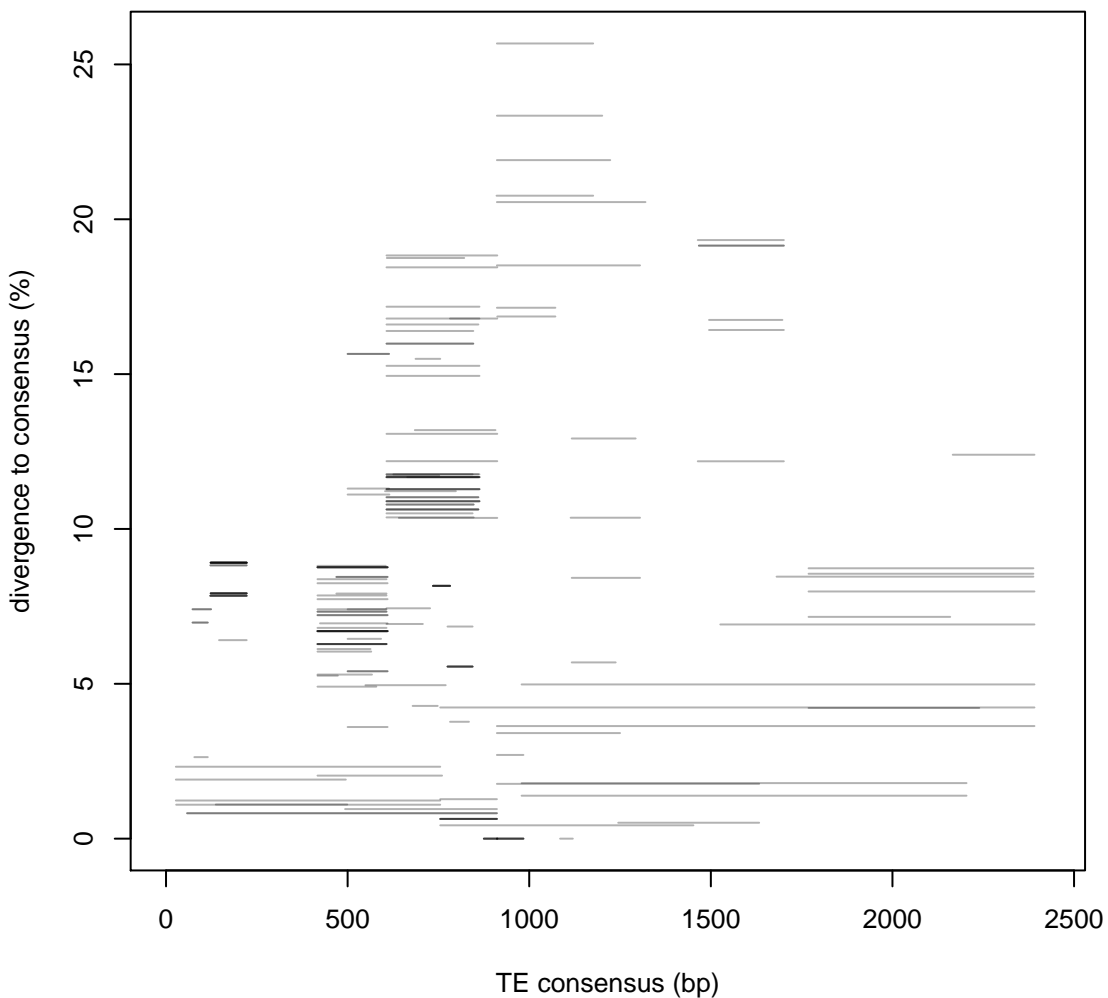
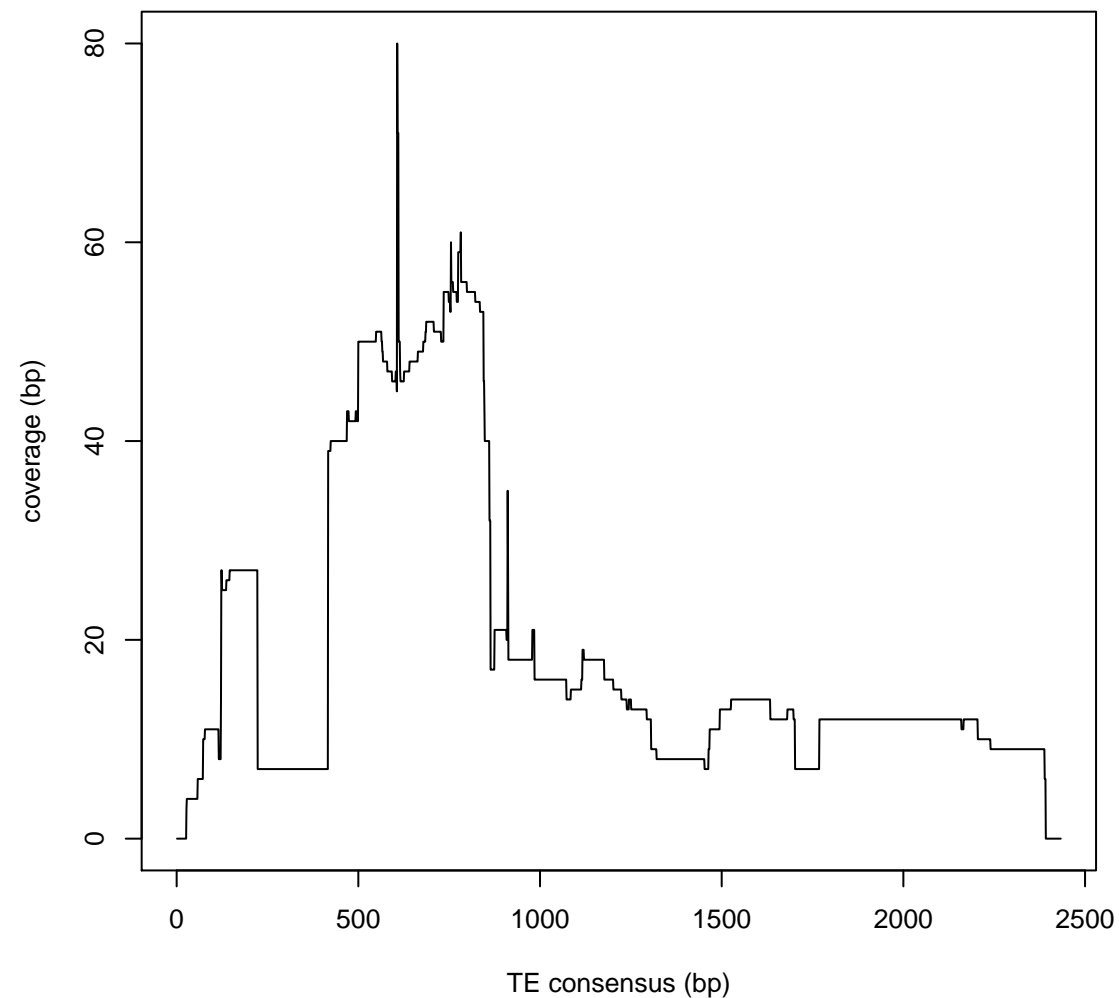


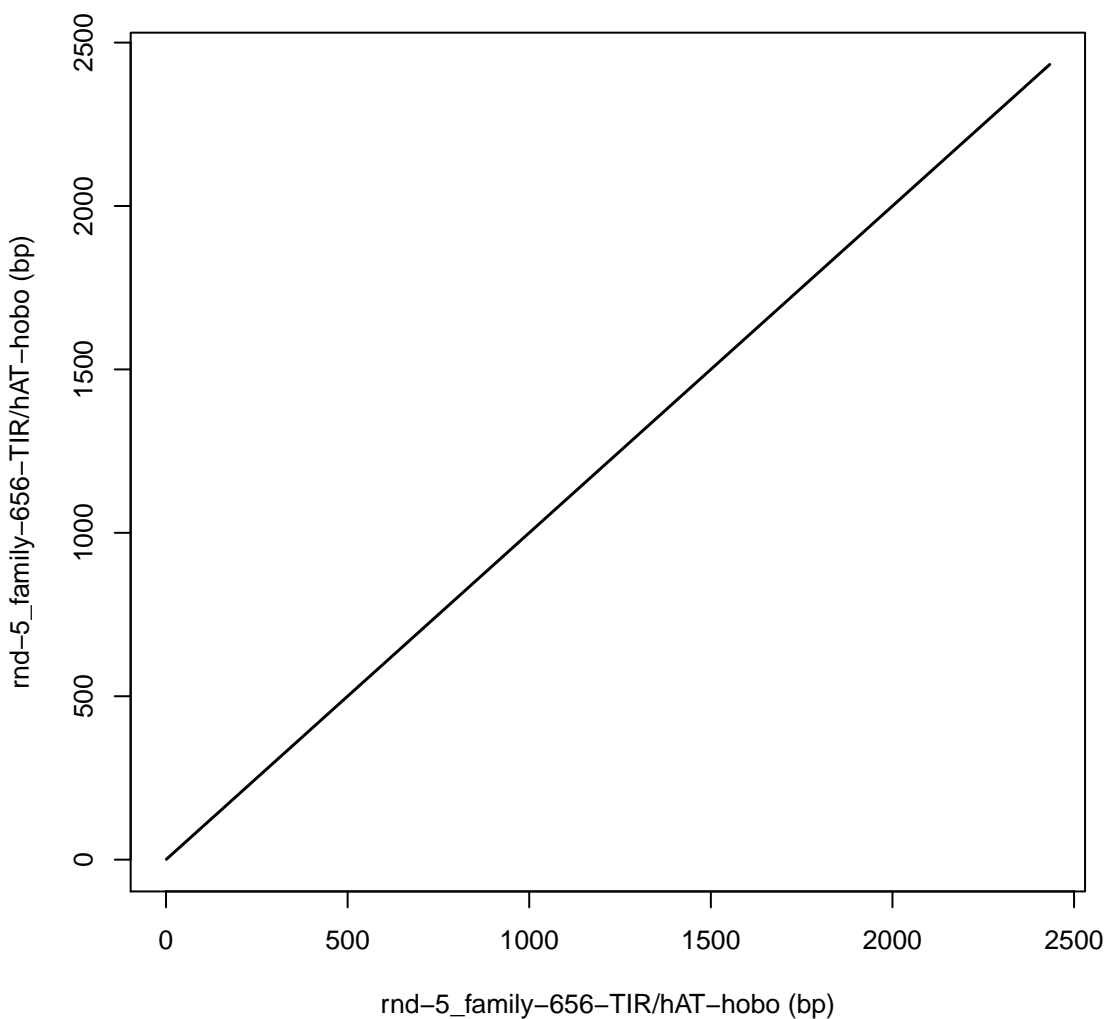
TE: rnd-5_family-656-TIR/hAT-hobo
consensus size: 2433bp; fragments: 185; full length: 0 (≥ 2189.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

