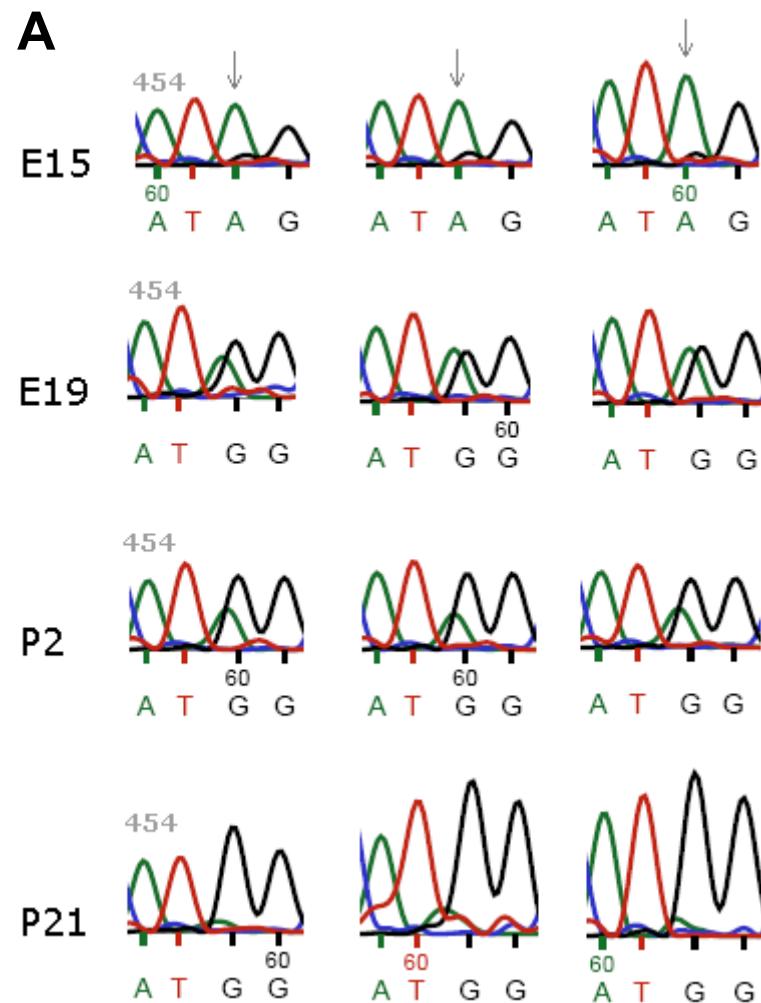


Figure S1. Validation of *Gabra3* (**A**) and *Htr2c* (**B**) editing using conventional Sanger sequencing (eurofins MWG operon). RNA from whole mouse brain (3 individuals) at 4 developmental stages (E15, E19, P2 and P21) was amplified by RT-PCR and sequenced. The edited positions are indicated by arrows. The first individual in each developmental stage was used for 454 sequencing.



B