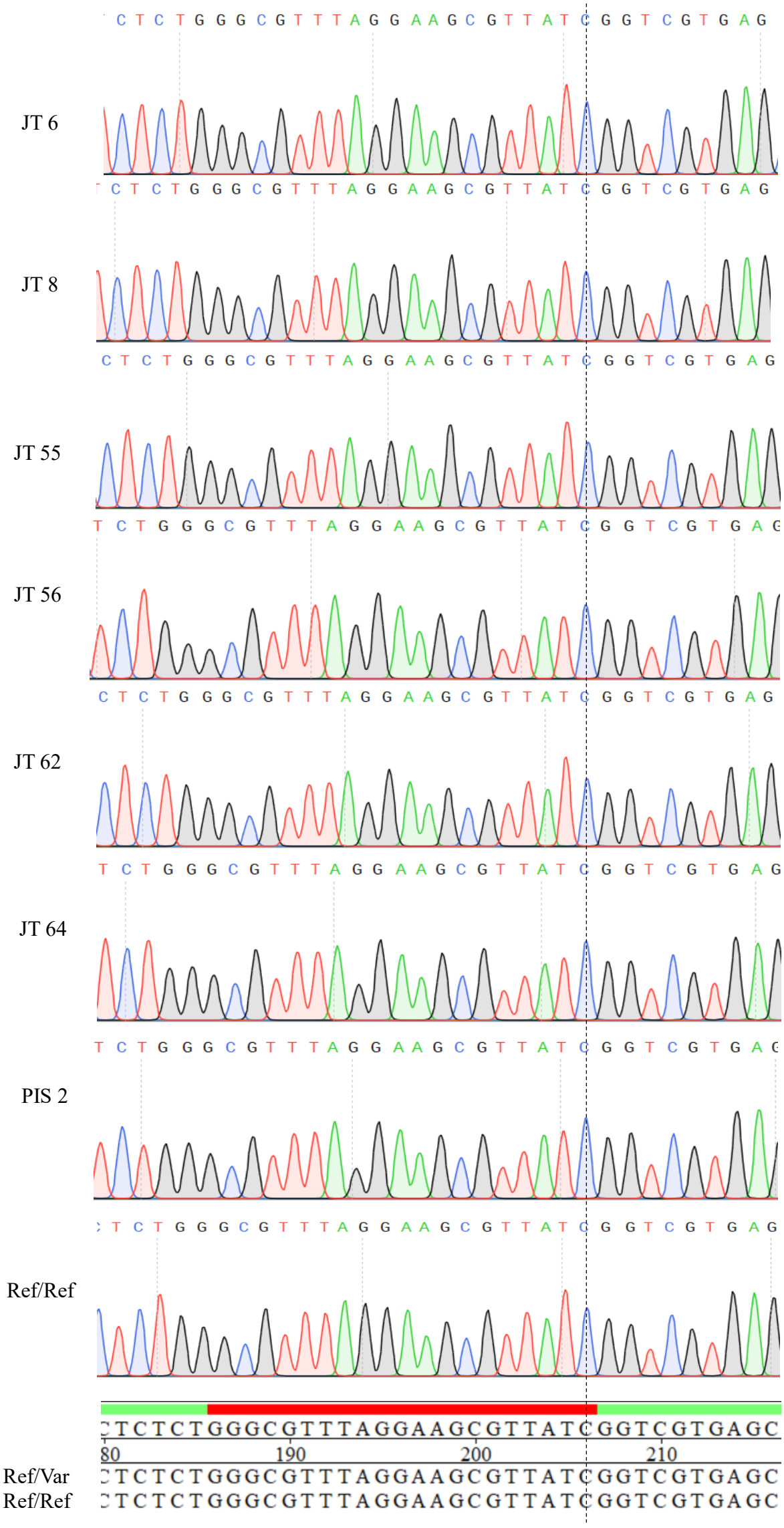

Figure S2. Validation of a mutation within the *ERG* gene through PCR and Sanger sequencing