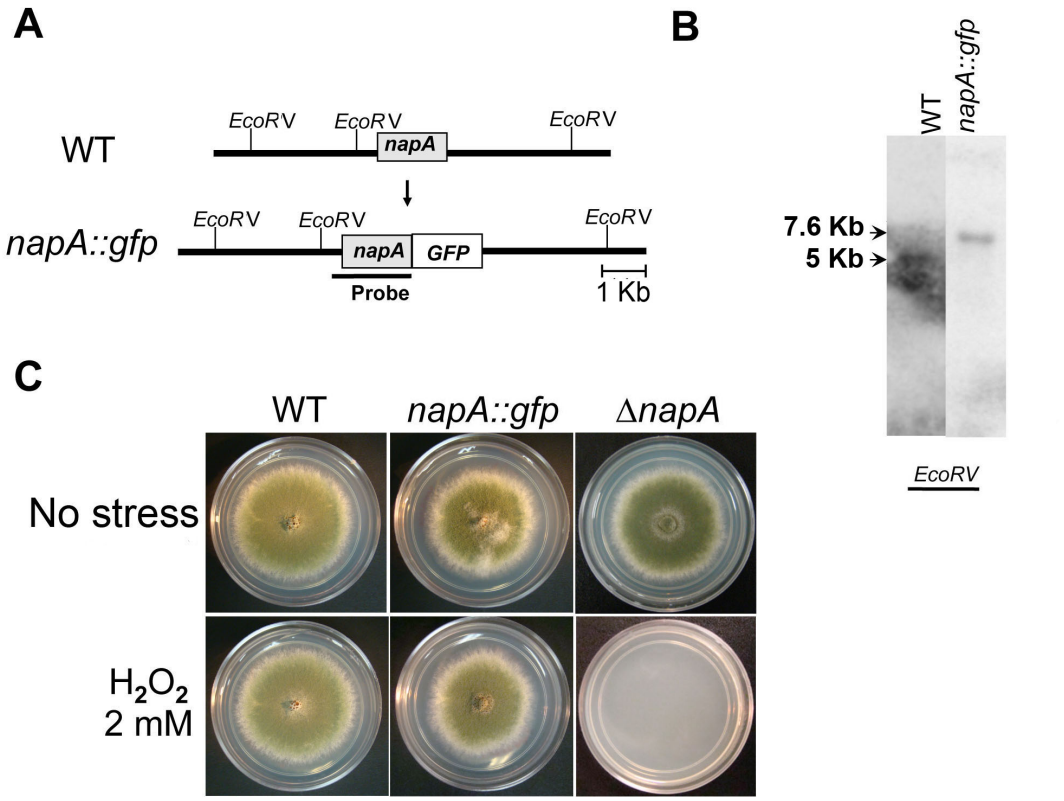

FIGURE S5. GFP tagging does not affect NapA function. **(A)** Three PCR products were used to generate an AtfA C-terminal GFP construct by double joint PCR, which was used to transform strain A1155. **(B)** Total DNA from strains A1155 (WT) and PyrG^{+} transformant TFL14 were digested with EcoRV restriction enzyme and used for Southern blot analysis using *napA* ORF as probe. The WT pattern corresponds to a 5 Kb band, while *napA::GFP* pattern corresponds to a 7.6 Kb fragment. H_2O_2 resistance of WT and *napA::GFP* strains is similar. **(C)** Conidiospores ($1X10^4$) from strains A1155 (WT), TFL9 (Δ*napA*) and TFL14 (*napA::GFP*) were inoculated on supplemented MM plates with or without 2 mM H_2O_2 and incubated at 37°C for 4 days.