Regulation of Nicotine Tolerance by Quorum Sensing and High Efficiency of Quorum Quenching Under Nicotine Stress in *Pseudomonas aeruginosa* PAO1

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Quorum sensing (QS) regulates the behavior of bacterial populations and promotes their adaptation and survival under stress. As QS is responsible for the virulence of vast majority of bacteria, quorum quenching (QQ), the interruption of QS, has become an attractive therapeutic strategy. However, the role of QS in stress tolerance and the efficiency of QQ under stress in bacteria are seldom explored. In this study, we demonstrated that QS-regulated catalase (CAT) expression and biofilm formation help *Pseudomonas aeruginosa* PAO1 resist nicotine stress. CAT activity and biofilm formation in wild type (WT) and ΔlasR strains are significantly higher than those in the ΔrhlR strain. Supplementation of ΔlasI strain with 3OC12-HSL showed similar CAT activity and biofilm formation as those of the WT strain. LasIR circuit rather than RhlIR circuit is vital to nicotine tolerance. Acylase I significantly decreased the production of virulence factors, namely elastase, pyocyanin, and pyoverdine under nicotine stress compared to the levels observed in the absence of nicotine stress. Thus, QQ is more efficient under stress. To our knowledge, this is the first study to report that QS contributes to nicotine tolerance in *P. aeruginosa*. This work facilitates a better application of QQ for the treatment of bacterial infections, especially under stress.

**Keywords:** nicotine tolerance, quorum sensing, antioxidant-producing ability, biofilm formation, quorum quenching, virulence

**INTRODUCTION**

Cell density-dependent cell-to-cell communication, termed as quorum sensing (QS), regulates the behavior of bacterial populations (Waters and Bassler, 2005). Bacteria secrete and share QS signaling molecules that bind to cognate receptors, and upon reaching critical concentration induce cell density-dependent adaptive responses within the population (Albuquerque et al., 2014). QS is responsible for a number of collective behavioral properties, including virulence factor secretion, biofilm formation, and horizontal gene transfer (Antonova and Hammer, 2011; Joo and Otto, 2012; Yang et al., 2017). Compared to individuality, sociality, regulated by QS, significantly increases the bacterial fitness in various environment (Darch et al., 2012). Despite increasing recognition on
bacterial QS, the roles that they play in the response of environmental stress are far from fully understood (Garcia-contreras et al., 2015).

Quorum sensing (QS) regulates the secretion of virulence factors from a broad spectrum of bacterial pathogens, including Pseudomonas aeruginosa (De Kievit and Iglewski, 2000). QS also participates in the development of biofilms, which are responsible for resistance to antibiotics, in many infections (Hazan et al., 2016). Due to the role of QS in pathogenicity and antibiotic resistance, the different factors involved in these pathways are considered to be attractive targets for novel antimicrobial agents (Starkey et al., 2014; Wang et al., 2016; Whiteley et al., 2017). Interruption of QS, which is known as quorum quenching (QQ), has been explored to control bacterial pathogenicity (Chan et al., 2015). As QS is an active process in response to environmental changes, QQ will have to be applicable under various conditions. Therefore, analysis of the QS response under different environmental conditions is vital for developing an efficient strategy involving QQ to control pathogenicity of bacteria.

Pseudomonas aeruginosa, one of the most common pathogenic bacteria in the world, not only infects humans, but also plants (Valentini et al., 2017). Its pathogenicity is mainly regulated by QS (Girard and Bloemberg, 2008; Whiteley et al., 2017). P. aeruginosa has two acyl-homoserine lactones (AHLs) QS circuits, LasIR and RhlIR (Stover et al., 2000). In LasIR circuit, LasI catalyzes the synthesis of N-3-oxo-dodecanoyl homoserine lactone (3OC12-HSL), which binds to its cognate receptor LasR and subsequently induces the expression of elastase-encoding genes involved in the development of pathogenicity of the bacteria (Pearson et al., 1994). For RhlIR circuit, RhlI catalyzes the synthesis of butyryl-HSL (C4-HSL), which binds to RhlR and subsequently activates a series of virulence factors including pyocyanin (Mukherjee et al., 2017). The well-elucidated mechanism of QS in P. aeruginosa allows us to study the feasibility of applying QQ to reduce the pathogenicity of the bacteria.

Though P. aeruginosa causes infection in both, humans and plants, they are exposed to various conditions. P. aeruginosa is known to inhabit hypoxic mucus plugs in the lungs of cystic fibrosis (CF) patient. Nearly 30% of smokers were involved in the population of CF patient (Ortega-Garcia et al., 2012). In addition, the growth of P. aeruginosa in stems and roots leads to systemic infection and ultimately to the development of severe soft-rot symptoms in tobacco (Pfeilmeier et al., 2016). Nicotine is one of the main alkaloid in tobacco. Recent evidence has demonstrated that P. aeruginosa could grow under nicotine stress in tobacco plants or human being, but few studies regarding the role of QS in nicotine tolerance in P. aeruginosa have been performed (Hutcherson et al., 2015), limiting the development and application of strategies involving QQ to control its pathogenicity under nicotine-stress conditions.

Thus, we employed P. aeruginosa PA01 as the model bacteria and nicotine as the typical stress. First, the growth and antioxidant-producing and biofilm-formation ability of wild-type (WT) strains and their signal-blind mutants were compared to investigate the role of QS in nicotine tolerance. Second, competition assay under nicotine stress and complementation experiment using a signal-deficient mutant were performed to analyze the possible mechanism. Finally, the efficiency of a QS inhibitor was analyzed under the presence and absence of nicotine stress to evaluate the application of QQ under these conditions. To our knowledge, this is the first study to report that QS plays an important role in nicotine tolerance, and demonstrates that LasIR circuit, rather than the RhlIR circuit, is responsible for nicotine tolerance in P. aeruginosa PA01. This information will help to improve our understanding of the role of bacterial QS under stress, and to develop and apply QQ-based strategies for combating bacterial infection in the future.

**MATERIALS AND METHODS**

**Bacterial Strains, Media, and Culture**

The bacterial strains used in this study were P. aeruginosa PA01 WT strain and its QS mutants ΔlasR, ΔrhlR, and ΔlasI (Wang et al., 2015).

Luria-Bertani (LB) medium with or without nicotine was used in this study. LB medium was composed of tryptone (10 g), yeast extract (5 g), NaCl (5 g) in 1 L distilled water. Filtered-sterile nicotine (0–2.0 g/L) was replenished according to requirement.

Inocula were obtained from overnight LB cultures. The initial optical density (OD) was 0.001 (600 nm), except where noted. The culture was incubated in a shaker, at 37°C with 250 rpm.

**The Detection of Reactive Oxygen Species (ROS)**

Wildtype strain, PAO1, was inoculated into LB with initial OD_{600} of 0.01. After the growth of the cells entered the logarithmic phase (OD_{600} = 1), 0, 1.6, and 2.0 g/L nicotine was added into the culture. To measure ROS, 2′,7′-dichlorofluorescin diacetate (DCFH-DA) was added at a final concentration of 10 mM. Within 1 h of incubation, DCFH-DA was hydrolyzed into dichlorofluorescin (DCFH) in the cells. Then DCFH was oxidized by ROS into dichlorofluorescin (DCFH-DA) was added at a final concentration of 10 mM. Within 1 h of incubation, DCFH-DA was hydrolyzed into dichlorofluorescin (DCFH) in the cells. Then DCFH was oxidized by ROS into dichlorofluorescin (DCF). DCF was measured using SpectraMax® i3 plate reader at 488 nm of excitation and 525 nm of emission (Molecular Devices, Sunnyvale, CA, USA) (Yu et al., 2014). H_{2}O_{2} treatment was used as a positive control. We calculated the relative ROS level by dividing the value of the DCF level obtained for experimental samples by that for LB medium.

**The Measurement of the Activity of Catalase (CAT) and Superoxide Dismutase (SOD)**

After exposure to 0, 1.6, and 2.0 g/L of nicotine, cells in logarithmic phase were harvested to detect the activity of CAT.
and SOD, respectively. Cells were washed thrice with 0.9% NaCl and ultrasonically lysed. Subsequently, crude enzymes were obtained by centrifugation at 4°C and 12,000 rpm for 10 min. The activity of CAT and SOD was detected using the ammonium molybdate method (A007) and hydroxylamine method (A001-1-1), respectively. The total protein content was determined using a modified Bradford assay (Kit A045). All assays were performed according to manufacturer’s instructions. These kits were purchased from the Nanjing Jiancheng Bioengineering Institute (Jiangsu, China).

One unit of CAT activity was defined as the amount of lyase that catalyzes the decomposition of 1 µM of H2O2 per minute at 37°C. One unit of SOD activity was defined as the amount of lyase that inhibits the rate of xanthine/xanthine oxidase-dependent cytochrome c reduction at 25°C by 50%. The activities of both enzymes were expressed as units per mg of cellular protein.

Biofilm Formation Analysis
After exposure to 0, 1.6, and 2.0 g/L of nicotine, the biofilm formation in 10-mL tubes was evaluated. Biofilm biomass was analyzed by crystal violet (CV) staining method described by Wang et al. (2012). After 24 h of incubation, the tubes were carefully washed twice with phosphate-buffered saline (PBS) to remove planktonic cells. After air drying for 5 min, biofilms were stained with 1 mL of 0.1% CV for 10 min, then the tubes were rinsed thoroughly thrice with distilled water to remove the unabsorbed CV. Finally, adhered CV was solubilized with 3 mL of alcohol acetone (4:1, v/v) and measured at 570 nm using a Nano-drop 2000 spectrophotometer after quantification using a Nano-drop 2000 spectrophotometer.

The polysaccharides, protein and DNA component of biofilm was analyzed according to Wang et al. (2012). In brief, the biofilm was washed thrice and resuspended in PBS. Subsequently, the suspension was heated to 80°C for 45 min, and the mixture was centrifuged at 13,000 rpm for 20 min to remove solid residues. The extracellular polysaccharides (EPS) and extracellular protein as the two main components of biofilm were determined using the phenol/sulfuric acid method (Dubois et al., 1956) and Coomassie brilliant blue assay (Bradford, 1976), respectively. The content of extracellular DNA as the other component of biofilm was quantified using a Nano-drop 2000 spectrophotometer after purification with a phenol/chloroform/isoamyl reagent.

The morphology of biofilm was observed by confocal laser scanning microscopy (CLSM, Leica, Germany). For ease of observation, crude glass slides were placed in flasks containing 0, 1.6, and 2.0 g/L of nicotine, and biofilms formed on these slides. The cell viability in biofilm was determined using a double live/dead staining kit containing nucleic acid stains SYTO 9 and propidium iodide (PI). After biofilm formation, the glass slides were gently rinsed by immersing them in PBS, removing all unadhered cells, and subsequently, stained for 15 min. Viable bacteria with intact cell membrane were stained with green, whereas dead bacteria with damaged membrane were stained with red. Stained samples were visualized with the following excitation/emission detectors and filter sets: for SYTO 9, 480/500 and for PI, 490/635 (Shi et al., 2016).

Coculture Assay
WT, ΔlasR, and ΔrhlR strains were grown to mid-logarithmic phase, respectively. WT vs. ΔlasR, and ΔrhlR vs. ΔlasR with the ratio of 1:1 (cell number) were separately cocultured in LB media with 0, 0.4, 0.8, 1.2, 1.6, and 2.0 g/L nicotine under 37°C for 24 h. The initial OD600 was 0.05. Then, skim milk agars were used to differentiate the ΔlasR strains from WT or ΔrhlR strains, where a clear zone appeared around WT and ΔrhlR colonies but not around ΔlasR colonies (Wang et al., 2015). Skim milk agar was prepared as follows (L): 1.25 g NaCl, 1.25 g yeast extract, 2.5 g trypptone, 80 g skim milk powder, and 15 g agar. For each value reported, at least 300 colonies were screened.

QQ Assay
Acylase I (Kit A8376-1G, Sigma, Germany) was used for QQ (Yeon et al., 2008) Overnight culture of the WT strain was inoculated into LB with 0, 1.6, and 2.0 g/L of nicotine. After 12 h of incubation, 0.25 mg/L acylase I was replenished to interrupt both, 3OC12-HSL and C4-HSL-mediated QS circuits. After another 12 h of incubation, the production of QS-regulated products including elastase, pyocyanin, and pyoverdine was compared among different culture conditions.

Elastase was detected by Pierce Fluorescent Protease Assay kit (Thermo). In brief, the culture was centrifuged at 12,000 rpm for 15 min. Subsequently, 100 µL of the supernatant was mixed with 100 µL of succinylated-casein solution (1:500 mixture of 2 g/L lyophilized succinylated casein and trinitrobenzene sulfonic acid, pH = 8.5) and incubated for 45 min in the dark at room temperature. The fluorescence was detected at 450 nm using a plate reader (SpectraMax® i3, Molecular Devices, Sunnyvale, CA, USA).

Pyocyanin was measured by chloroform and hydrochloric acid extraction (Pearson et al., 1994). A total of 1.5 mL of chloroform was used to extract 2.5 mL of the supernatant. The pyocyanin was re-extracted from the chloroform using 1 mL of 0.2 M hydrochloric acid. Finally, the absorbance of the supernatant was measured at 520 nm. The concentration of pyocyanin was equal to the absorbance multiplied by 12.8 mg/L.

Pyoverdine was detected using the method described by Wurst et al. (2014). In brief, the cultures were centrifuged at 12,000 rpm for 15 min. The absorbance of the supernatant was measured at 405 nm.

The level of elastase, pyocyanin, and pyoverdine were expressed as units per OD600 unit in order to avoid the interference of cell density. All experiments were in triplicate.

Statistical Analysis
GraphPad Prism 6.0 software was used for statistical analyses. Two-way ANOVA and t-test were performed. Differences with a value of p < 0.05 were considered to be statistically significant.

RESULTS
QS Plays an Important Role in Nicotine Tolerance
QS is involved in the regulation of the behavior of a bacterial population, whereby the cells secrete diffusible substances that
generate phenotypic responses in the living group. Compared to individuality, sociality confers a 100–1,000-fold increase in resistance to stress (Hazan et al., 2016). Thus, our hypothesis is that QS possibly plays an important role in nicotine tolerance. To confirm this hypothesis, a simple experiment comparing the growth of the WT strain with complete QS circuits and the signal-blind mutants under nicotine stress, was performed. Signal-blind mutants cannot respond to their cognate signals, and therefore, the expression of their corresponding regulons is inhibited.

As shown in Figure 1, there was no difference of bacterial growth between the WT and signal-blind mutant ΔlasR and ΔrhlR strains in the absence of nicotine. Under a 1.6 g/L-nicotine treatment, the growth of the WT, ΔlasR, and ΔrhlR strains was inhibited. However, the growth of the ΔlasR strain was significantly lower than that of the WT and ΔrhlR strains. Similar to the result of the 1.6 g/L-nicotine treatment, the growth of all three strains was inhibited under a 2.0 g/L-nicotine treatment. The lowest growth was observed in ΔlasR culture. Though other mechanisms possibly exist, the results indicated that QS played an important role in nicotine tolerance by P. aeruginosa PAO1.

Antioxidant Ability Regulated by QS Benefit for Nicotine Tolerance

Nicotine is a carcinogenic, teratogenic, and mutagenic substance, which can induce the production of a large number of free radicals, resulting in oxidative damage to cells (Haussmann and Fariss, 2016). The comparison of bacterial growth indicated that QS played an important role in nicotine tolerance. According to García-contreras et al. (2015), QS is able to exert a robust anti-oxidative response. Thus, one possibility could be that the role of QS in anti-oxidative response was beneficial for nicotine tolerance.

In order to validate this assumption, we first evaluated the ROS generation under nicotine exposure. As shown in Figure 2A, the level of intracellular ROS in WT cells increased significantly with the increase in nicotine. Nicotine-treated WT cells exhibited a higher level of ROS compared to the untreated

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**Figure 1** Comparison of bacterial growth among different strains (WT, blue bars; ΔlasR, red bars; ΔrhlR, yellow bars) under exposure to different concentrations of nicotine. Different letters indicate significant difference at p < 0.05 and the same letter indicates no significant difference.

**Figure 2** Relative ROS levels (A) in WT on exposure to different concentrations of nicotine (left) and H2O2 (right); CAT activity (B) and SOD activity (C) among different strains (WT, blue bars; ΔlasR, red bars; ΔrhlR, yellow bars) under exposure to different concentrations of nicotine. Different letters indicate significant difference at p < 0.05 and the same letter indicates no significant difference.
WT cells. Especially a 2.0 g/L-nicotine treatment led to the increase in the level of ROS in nicotine-treated cells, and this level was 24.4 times higher than that in untreated cells. Using H$_2$O$_2$ as positive control, it was observed that the level of ROS produced by 2.0 g/L-nicotine treatment, is higher than that produced by 2 mM-H$_2$O$_2$ treatment. Therefore, it can be inferred that the
higher the concentration of nicotine, the stronger the oxidative stress induced.

To confirm that QS could contribute to nicotine tolerance by activating antioxidant defense system, the activity of antioxidant enzymes were measured among WT, ΔlasR, and ΔrhlR strains. As shown in Figure 2B, there was no difference in the activity of CAT among the WT and mutant strains without nicotine stress. The activity of CAT significantly increased on exposure to 1.6 g/L of nicotine in the WT and ΔrhlR strains compared to that in the ΔlasR strain. Though the CAT activity decreased under a 2.0 g/L-nicotine treatment due to toxicity, the WT strain showed a significantly higher activity of CAT than that observed in ΔlasR, and this activity had no significant difference with that observed in ΔrhlR strain.

Additionally, we measured the SOD activity among these three strains. However, no significant increase was observed for this parameter (Figure 2C). Taking the above-mentioned data into account, bacterial QS involving the LasIR and RhlIR circuits, regulate the anti-oxidative response to nicotine stress in WT strain. Further studies are required to explain why QS promotes CAT activity, and not SOD activity.

QS-Regulated Biofilm Formation Favored of Nicotine Tolerance

Biofilm formation, mainly regulated by QS, could be another reason for stress tolerance (Hammer and Bassler, 2003; Daniels et al., 2004; Shrout and Nerenberg, 2012). Compared to planktonic cells, biofilm formation increases stress tolerance up by 10–1,000 folds (Hazan et al., 2016). Another parallel assumption is that QS-regulated biofilm formation is beneficial for nicotine tolerance. Therefore, to clearly understand the effect from QS-regulated biofilm formation on nicotine tolerance, we compared the biofilm formation of WT and ΔlasR and ΔrhlR strains on exposure to nicotine.

As shown in Figure 3A, there was no significant difference in the biofilm formation of WT and ΔlasR and ΔrhlR strains in absence of nicotine. On treating with 1.6 and 2.0 g/L of nicotine, the biofilm biomass of WT and ΔrhlR increased significantly. There was no difference of biofilm biomass between WT and ΔrhlR. However, the biofilm biomass of ΔlasR was significantly lower than that of the other two strains.

In addition, the amount of certain biofilm components was analyzed. As shown in Figures 3B–D, the level of EPS and extracellular proteins in the biofilms of the WT and ΔrhlR strains was significantly higher than that of the ΔlasR strains.
under a 1.6 g/L-nicotine treatment. After exposure to 2.0 g/L of nicotine, no significant difference in the level of EPS between the biofilms of ΔlasR and ΔrhlR was observed. The level of EPS and extracellular protein in the biofilm of the WT strain was significantly higher than that in the biofilm of ΔlasR under a 2.0 g/L-nicotine treatment. The extracellular DNA content was almost equivalent among three strains, indicated by an extremely small amount of extracellular DNA in the biofilm.

Moreover, we used the CLSM to observe the structure of biofilm and employed a double live/dead staining to determine cell viability in biofilm. As shown in Figure 4, the biofilm thickness of WT and ΔrhlR strains increased under nicotine stress. However, the biofilm formation of ΔlasR was significantly inhibited under nicotine stress. Compared to WT and ΔrhlR biofilm, the number of dead cells dramatically increased in the ΔlasR biofilm. All above data demonstrated that QS-regulated biofilm formation was also involved in enhancement of nicotine tolerance.

LasIR Being Responsible for Nicotine Tolerance

As seen in Figures 2B, 3A, the CAT activity and biofilm biomass in the ΔlasR strain was significantly lower than the WT and ΔrhlR strain. Meanwhile there were no significant differences for the same parameters between the WT and ΔrhlR strains. It suggested that the LasIR circuit played more important role in nicotine tolerance than the RhlIR circuit. Bacteria lacking a functional LasIR circuit, are sensitive to nicotine. To confirm these, competition experiments between the WT and ΔlasR strains or between the ΔrhlR and ΔlasR strains were conducted.

As shown in Figure 5, without nicotine stress, ΔlasR growth was higher than that of the WT or ΔrhlR strains. After 24 h, 79.1 and 86.1% of the total population in the WT competition system and the ΔrhlR competition system, respectively, were ΔlasR cells. With the increase in nicotine concentration, the proportion of ΔlasR population significantly decreased. It was reduced to 16.7% in WT competition system under 2.0 g/L-nicotine stress. The decrease of ΔlasR fitness advantage with the increase of nicotine is consistent with the above hypothesis.

For the competition experiment, other factors except the nicotine tolerance could affect the advantageous fitness. Thus, ΔlasI supplementation with 3OC12-HSL was implemented in further experiments. ΔlasI is a signal-deficient mutant, without the ability to synthesize 3OC12-HSL, but with the functional signal receptors, LasR. According to the mechanism of QS, exogenous additional of 3OC12-HSL also could bind to LasR and trigger the expression of the corresponding regulon (Wang et al., 2015). As shown in Figure 6, the CAT activity and biofilm formation in the ΔlasI strain was similar to those in the ΔlasR strain. However, addition of 3OC12-HSL significantly increased the CAT activity and biofilm formation in the ΔlasI strain, and they were nearly identical with those in the WT strain. Both competition systems in coculture and signal complementary assays for ΔlasI confirm that LasIR circuit is important for nicotine tolerance in P. aeruginosa.

QQ Acting Even Better Under Nicotine Stress

Quorum quenching (QQ) was widely used for controlling pathogenicity in P. aeruginosa, and reducing the level of virulence factors such as elastase, pyocyanin, and pyoverdine (Lee and
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the ΔrhlR and ΔlasR strains. In LB media without nicotine, the ΔlasR strain had a significant fitness than the ΔrhlR strain. However, with the increase in nicotine concentration in LB media, the growth of the ΔrhlR strain increased significantly compared to that of the ΔlasR strain (Figure 5). From Figure 6, supplementation of the ΔlasI strain with 3OC12-HSL led to the culture showing similar CAT activity and biofilm formation to those of the WT strain, under nicotine stress. Both competition in coculture and signal complementary assays for ΔlasI confirmed that LasIR circuit was more important than the RhlIR circuit in the response to nicotine stress.

The members of the QS pathway are promising targets for treatment of pathogenic infection (Köhler et al., 2010). Several QQ reagents have been developed (O’Loughlin et al., 2013). As shown in Figure 7, the inhibition efficiencies of acylase I are different for various of virulence factors. According to the genetic network of the PAO1 strain, lasR, rhlR, and pqsE have been reported to be involved in the production of pyocyanin (O’Loughlin et al., 2013; Rampioni et al., 2016), while ampR, ppyR, mexT, and lasR are involved in the production of elastase (Van Delden et al., 1998; Maseda et al., 2004; Kong et al., 2005; Attila et al., 2008). There are much more genes contributing to elastase production than those contributing to pyocyanin production. Thus, the inhibition efficiency for pyocyanin was higher, while less elastase production was inhibited. The production of pyoverdine was regulated by PQS, a type of a QS pathway that is not mediated by AHLs, in P. aeruginosa (Lee and Zhang, 2015). Acylase I can only interrupt AHLs-mediated QS (Zhang et al., 2015). Thus, acylase I did not inhibit the production of pyoverdine under no nicotine treatment conditions. Different QS circuits regulate the secretion of different virulence factors (Chugani et al., 2001). One virulence factor is regulated by completely or partially regulated by QS (O’Loughlin et al., 2013; Husain et al., 2017). QQ was successful in reducing the production of certain, but not all, kinds of tested virulence factors in P. aeruginosa.

Various conditions, such as pH and temperature, possibly affect the application of QQ in pathogenicity control. pH and temperature could affect the existence of QS signal in the environment (Yates et al., 2002). Few studies have focused on the efficiency of QQ under stress. In this study, the QQ showed a higher efficiency in decreasing the production of virulence factors, including elastase, pyocyanin, and pyoverdine under nicotine stress compared to no stress. Nicotine is toxic to most kinds of bacteria. QS contributes to nicotine tolerance (Figure 8A). Interruption of QS led to the decrease in both, nicotine tolerance and virulence (Figures 8B,C). After loss of nicotine tolerance, the bacterial population possibly reduces their virulence in order to survive as a trade-off. Though we can not apply of QQ under nicotine stress due to its addiction, it gives us an explanation that the combination of QQ with antibiotics is higher efficient than only one treatment (Wang et al., 2018). Therefore, this study not only improves our understanding regarding the role of QS in environmental stress tolerance, but also provides a foundation for the development of QQ-based strategies to control or reduce the pathogenicity of bacteria (Figure 8).

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**AUTHOR CONTRIBUTIONS**

MW, HT, and DS conceived and designed the experiments. HT, YZ, YM, and MT performed the experiments. HT and MW analyzed the data. MW and DS contributed reagents, materials, and analysis tools. MW and HT wrote the paper.

**ACKNOWLEDGMENTS**

This study was supported by the National Science Foundation of China (Grant no. 31570490, 51478432, 41403080) and National Undergraduate Training Program for Innovation and Entrepreneurship (Grant no. LY20171035302). We also thank Professor E. Peter Greenberg and Professor Ajai A. Dandekar for providing us with the strains and for their helpful assistance.
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**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The reviewer MC-Y and handling Editor declared their shared affiliation.

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