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\*CORRESPONDENCE Jichen Chen chenjichen2001@163.com

<sup>†</sup>These authors have contributed equally to this work

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# Transcriptional factor OmpR positively regulates prodigiosin biosynthesis in *Serratia marcescens* FZSF02 by binding with the promoter of the prodigiosin cluster

## Xianbo Jia<sup>1†</sup>, Ke Zhao<sup>2†</sup>, Fangchen Liu<sup>2</sup>, Junjie Lin<sup>3</sup>, Chengiang Lin<sup>1</sup> and Jichen Chen<sup>1</sup>\*

<sup>1</sup>Institute of Soil and Fertilizer, Fujian Academy of Agricultural Sciences, Fujian Key Laboratory of Plant Nutrition and Fertilizer, Fuzhou, China, <sup>2</sup>College of Resources and Environment, Fujian Agriculture and Forestry University, Fuzhou, China, <sup>3</sup>College of Life Sciences, Fujian Agriculture and Forestry University, Fuzhou, China

Prodigiosin is a promising secondary metabolite mainly produced by *Serratia marcescens*. The production of prodigiosin by *S. marcescens* is regulated by different kinds of regulatory systems, including the EnvZ/OmpR system. In this study, we demonstrated that the regulatory factor OmpR positively regulated prodigiosin production in *S. marcescens* FZSF02 by directly binding to the promoter region of the prodigiosin biosynthesis cluster with a *lacZ* reporter assay and electrophoretic mobility shift assay (EMSA). The binding sequence with the *pig* promoter was identified by a DNase I footprinting assay. We further demonstrate that OmpR regulates its own expression by directly binding to the promoter region of *envZ/ompR*. For the first time, the regulatory mechanism of prodigiosin production by the transcriptional factor OmpR was revealed.

#### KEYWORDS

Serratia marcescens, prodigiosin, OmpR, two-component system, regulatory mechanism

# Introduction

Prodigiosin, an important secondary metabolite produced by *S. marcescens* and some other bacteria, is of particular interest for its potential applications, including various pharmacological activities, food colorants, and potential sunscreens (Stankovic et al., 2014; Darshan and Manonmani, 2015). In the genus *Serratia*, biosynthesis of prodigiosin is regulated not only by external factors, including temperature, pH and medium composition (Paul et al., 2020) but also by various genes (Williamson et al., 2006). Nearly 30 genes have been reported to be involved in prodigiosin biosynthesis in *S. marcescens* (Pan et al., 2021),

and more studies should be carried out to search for new regulatory genes and uncover the complex regulatory mechanisms of this secondary metabolite.

The two-component system is a family of signal transduction proteins reported to be present in all types of life (Stock et al., 1999; Scharf, 2010; Papon and Stock, 2019). In bacteria, the classical two-component system consists of a sensor protein and a regulatory protein (Yuchuan et al., 2019). Sensor proteins respond to chemical or physical signals by phosphorylating regulatory proteins, and phosphorylated regulatory proteins can regulate the expression levels of downstream genes by binding to certain DNA sequences (Groisman, 2016). Two-component systems regulate many processes of bacteria, such as adaptation to environmental changes: osmolarity (Boyce et al., 2016), temperature (Dhiman et al., 2015; Najnin et al., 2016), oxygen (Dixon, 1998; Wright et al., 2018), regulation of developmental pathways, and behaviors, such as sporulation (Sarwar and Garza, 2015), biofilm formation (Lai et al., 2005), quorum sensing (Kruppa et al., 2004), regulation of secondary metabolite biosynthesis (Sola-Landa et al., 2003), virulence (Beier and Gross, 2006), and antibiotic resistance (Bhagirath et al., 2019; Tierney and Rather, 2019). Biosynthesis of prodigiosin was also regulated by different types of two component systems in various Serratia strains, including PigQ/W and PhoB/PhoR in Serratia 39,006 (Fineran et al., 2005; Gristwood et al., 2009); EepR/EepS in S. marcescens CMS376, S. marcescens K904, and S. marcescens Nima (Stella et al., 2015); RssB/RssA in S. marcescens CH-1 (Horng et al., 2010); and CpxR/A in S. marcescens FS14 (Qiu et al., 2021).

The two component system EnvZ/OmpR is an important signal transduction system in bacteria responding to various environmental stress and growth conditions (Qin et al., 2001). We have previously demonstrated that when *envZ* or *ompR* was knocked out, *S. marcescens* FZSF02 lost its prodigiosin biosynthesis ability (Jia et al., 2021), and OmpR was also recently found to control prodigiosin biosynthesis in *S. marcescens* JNB5-1 (Pan et al., 2022). However, the regulatory mechanism of the two-component EnvZ/OmpR system on prodigiosin production is still unknown.

In this study, with *LacZ*-reporter studies and an electrophoretic mobility shift assay (EMSA), we demonstrated that OmpR positively regulated prodigiosin biosynthesis by directly binding to the promoter region of the prodigiosin biosynthesis gene cluster. We also found that OmpR can regulate its own expression level by binding the promoter of the EnvZ/OmpR genes.

# Materials and methods

# Bacterial strains, plasmids, and culture conditions

The bacterial strains and plasmids used in this study are listed in Supplementary Tables S1, S2. Serratia marcescens FZSF02 and its related mutants were incubated in lysogeny broth (LB) at 28°C and 180 rpm. *E. coli* DH5a and *E. coli* Rosetta (DE3) were cultured in LB medium at 37°C and 220 rpm. The final concentrations of antibiotics used in this study were as follows: 100 mg/L ampicillin, 100 mg/L kanamycin, and 50 mg/L chloramphenicol.

# Construction of the in-frame deletion mutant and complementary strains

FZSF02 $\Delta ompR$  and FZSF02 $\Delta envZ$  were constructed in our previous study (Jia et al., 2021). FZSF02 $\Delta envZ\Delta ompR$  was also constructed with the homologous recombination method (Jia et al., 2021). Complementary strains were constructed with the plasmid pRK415 as we have reported in our previous study (Jia et al., 2021). pRK415-ompR, pRK415-envZ and pRK415- $\Delta envZ\Delta ompR$  were transformed into FZSF02 $\Delta ompR$ , FZSF02 $\Delta envZ$ , and FZSF02 $\Delta envZ\Delta ompR$ , respectively, to construct the complementary strains. The primers used in these experiments are listed in Supplementary Table S2.

#### lacZ reporter assays

The *pig* promoter and *ompR* promoter were inserted upstream of *lacZ* in the plasmid pTOPO-*lacZ*-Cmr to construct the plasmids pTOPO-*Pig*pro-*lacZ*-Cmr and pTOPO-OmpRpro*lacZ*-Cmr. The primers used in this experiment are listed in Supplementary Table S2. The constructed plasmids with the *Pig* promoter and *ompR* promoter were transformed into the wildtype strain *S. marcescens* FZSF02 and the *ompR*-knockout strain FZSF02 $\Delta$ ompR, respectively. The plasmids pTOPO-*Pig*pro-Cmr and pTOPO-*ompR*pro-Cmr were also transformed into *S. marcescens* FZSF02 and FZSF02 $\Delta$ ompR as controls. For liquid  $\beta$ -galactosidase assays, constructed strains were cultured at 28°C and 180 rpm for 16 h, and enzyme activities were measured in sonicated extracts according to the method described by Pardee et al. (1959).

### Electrophoretic mobility shift assay

The coding sequence of *ompR* was cloned into pEASY<sup>®</sup>-Blunt E2 (TransGen Biotech, Beijing, China). The plasmid pEASY<sup>®</sup>-Blunt E2-*ompR* was transformed into *E. coli* Rosetta (DE3). The DNA fragments containing the *pig* promoter (406 bp) and *ompR* promoter (257 bp) were cloned into the pTOPO-Blunt simple vector (Aidlab, China), respectively, and then promoter probes were obtained through polymerase chain reaction (PCR) with primers M13F and M13R labeled with Cy5.5 at the 5' end. The *Pig*pro probe and *ompR*pro probe were 556 and 407 bp, respectively. Electrophoretic mobility shift assay (EMSA) was carried out with an EMSA/Gel-Shift kit (Beyotime, Shanghai, China). The purified probe and protein were mixed with EMSA/

Gel-Shift binding buffer (5×), and a total of 10µl of the reaction system was supplied with distilled water and incubated for 30 min at 28°C. 6% native PAGE was prepared as the kit protocol described, and the reaction mixture was loaded onto the PAGE. Electrophoresis was performed at 60 V for 3 h in 0.5× TBE buffer, and the gels were exposed to an Odyssey CLx (LI-COR<sup>®</sup> Biosciences).

### DNase I footprinting assay

The *pig* promoter probe was prepared by polymerase chain reaction (PCR) with primers M13F and M13R (labeled with Hex). The DNase I foot-printing assay was carried out as described by Shi et al. (2017).

### Results

# OmpR activates the transcription level of the *pig* gene cluster

We have previously demonstrated that gene deletion of ompR (GenBank: QJU42212.1) or envZ (GenBank: QJU42211.1) would result in loss of prodigiosin producing ability in S. marcescens FZSF02; transcription levels of pigA also down-regulated significantly when ompR or envZ was knocked out assayed by qPCR (Jia et al., 2021). The effect of OmpR and EnvZ on prodigiosin-producing ability was further demonstrated by knockout *ompR* and *envZ*, double knockout of envZ and ompR in this study; gene deletion strains all lost prodigiosin-producing ability, prodigiosin-producing ability restored in complementary strain (Figures 1A-C). OmpR was very conserved in amino acid sequences among many Gramnegative bacteria, such as Rouxiella aceris, Ewingella americana, Yersinia enterocolitica, Citrobacter youngae, Hafnia psychrotolerans, Escherichia coli and Klebsiella pneumoniae (Figure 1D). Many studies have demonstrated that OmpR influences a wide variety of cellular processes in E. coli, Salmonella sp. and Shigella sp., as well as pathogenic species of Yersinia sp. (Jaworska et al., 2021), but the influence of secondary metabolite biosynthesis by OmpR has rarely been reported. We have previously demonstrated that OmpR can positively regulate the production of the secondary metabolite prodigiosin in S. marcescens FZSF02, but the regulatory mechanism was unknown.

To further test whether the regulation of prodigiosin biosynthesis by OmpR is at the transcriptional level, the  $\beta$ -galactosidase activity of FZSF02 and FZSF02 $\Delta$ ompR was assayed when the *lacZ* gene was under control by the *pig* cluster promoter. The results showed that the  $\beta$ -galactosidase activity of FZSF02 $\Delta$ ompR (pTOPO-*Pig*pro-*lacZ*-Cmr) decreased by 88.5% compared with that of the wild-type strain FZSF02 WT (pTOPO-*Pig*pro-*lacZ*-Cmr; Figure 2A), and almost no  $\beta$ -galactosidase activity was tested in the control group of WT (pTOPO-*Pig*pro-Cmr) and  $\Delta ompR$  (pTOPO-*Pig*pro-Cmr; Figure 2A). This result indicates that OmpR directly or indirectly activates the transcription level of the *pig* gene cluster, and influences prodigiosin synthesis in strain FZSF02.

### OmpR activates the transcription level of the *pig* gene cluster by directly binding to its promoter sequence

To study whether OmpR regulates the expression of the *pig* gene cluster by binding with its promoter directly, EMSA was used to detect the binding ability between OmpR and the *pig* gene cluster promoter sequence. The results showed that OmpR can bind with the probe prepared with the prodigiosin cluster promoter sequence (Figure 2B). A DNase I footprinting assay showed that the proposed binding sequence of OmpR on the *pig* promoter was 5'CATTTATTTACATTTAC3' (Figure 3), which located on -103 bp to -86 bp relative to the A of the ATG start codon of *pigA*. *Pig* promoter sequence is between *pigA* (QJU38817.1) and *cueR* (QJU38818.1) on the genome of FZSF02.

### OMPR activates its own expression level

To test whether the autoregulation of OmpR exists in *S. marcescens* FZSF02, the  $\beta$ -galactosidase activity of FZSF02 and FZSF02 $\Delta$ ompR was assayed when the *lacZ* gene was first controlled by the ompR promoter. The results showed that the  $\beta$ -galactosidase activity of FZSF02 $\Delta$ ompR (100 U/ml) decreased by 92.3% compared with that of the wild-type strain FZSF02 (1,300 U/ml; Figure 2C). This finding demonstrates that OmpR can activate its own expression level.

### OmpR can directly bind to the envZ/ompR promoter

To test whether the activation of OmpR on its own expression is performed by binding with the *envZ/ompR* promoter, EMAS was used to examine the binding ability between OmpR and the *envZ/ompR* promoter sequence. The results showed that when the *envZ/ompR* promoter sequence was used as a probe, OmpR could bind with the labeled probe (Figure 2D). The addition of unlabeled probe can compete with the labeled probe, which further demonstrates the binding ability.

# Discussion

Prodigiosin was a kind of bacterial secondary metabolites produced mainly by many *S. marcescens* strains. Various regulating genes involved in prodigiosin biosynthesis have been found in the



past two decades, but new regulators, such as RcsB (Pan et al., 2021), CpxA/R (Qiu et al., 2021) and Fnr (Sun et al., 2021), have still been reported continuously. Research of the these genes may help to uncover the regulatory mechanism behind prodigiosin biosynthesis in *S. marcescens*.

EnvZ/OmpR is known to control motility (Prüß, 2017), intracellular survival (Du et al., 2022), antibiotic resistance (Ko and Choi, 2022), virulence (Tipton and Rather, 2017), and other characteristics of different bacterial strains, but few studies have reported the role of EnvZ/OmpR in *S. marcescens*. We have demonstrated previously that mutation of *ompR* or *envZ* would cause the loss of prodigiosin producing ability in *S. marcescens* FZSF02 and confirmed that the two-component system EnvZ/OmpR was a newly found system that can regulate prodigiosin biosynthesis (Jia et al., 2021). In this study, the regulatory function of EnvZ/OmpR on prodigiosin biosynthesis was further confirmed by gene deletion and complementation (Figure 1C). For the EnvZ/OmpR system, OmpR was reported to play the role by binding to the gene promoters and regulating the expression of other genes (Wang et al., 2021). *LacZ* reporter assays and EMSA assay in this study also showed OmpR regulate prodigiosin biosynthesis by directly binding to the promoter region of *pig* cluster (Figures 2A,B). The binding region of the OmpR was identified as 5'CATTTATTTACATTTAC3' (Figure 3) by a DNase I footprinting assay. The binding sequence showed 45% identity to the *E. coli* consensus sequence (5'TTTTACTT TTGTAACATAT3'; Maeda et al., 1991) and 55% identity to that of *Y. enterocolitica* (5'ATTTATTGATGGTAACAATT3'; Nieckarz et al., 2020).



Many two-component systems regulating proteins can autoregulate their own expression by binding to their promoters, and this kind of feedback allows the regulatory functions of the system to be more flexible (Groisman, 2016). For the EnvZ/OmpR two-component system, autoregulation differs among different strains; it exists in *Salmonella enterica* (Bang et al., 2002; Cameron and Dorman, 2012) but not in *E. coli* (Ochman and Wilson, 1987; Doolittle et al., 1996) and *Acinetobacter baumannii* (Tipton and Rather, 2017). The results of the *LacZ* reporter assay (Figure 2C) and EMSA assay (Figure 2D) indicated that OmpR can bind to the promoter region of *envZ/ompR* and promote the expression of OmpR and EnvZ.

Based on the above results, we proposed that the regulatory mechanism of the two-component EnvZ/OmpR system on prodigisin biosynthesis was probably as follows (Figure 4): Some unknown factors induced the expression of EnvZ and OmpR, and OmpR was then phosphorylated by EnvZ (Cai and Inouye, 2002). Phosphorylated OmpR activated the expression of more EnvZ and OmpR by binding with the *envZ/ompR* 

promoter. When the concentration of OmpR reached a certain level, the *pig* gene cluster promoter persistently bound with OmpR, and genes involved in prodigiosin biosynthesis were highly expressed at the transcriptional level.

Although we have previously found that when ompR was knocked out, *S. marcescens* FZSF02 lost the prodigiosin biosynthesis ability (Jia et al., 2021), in this study, the proposed regulatory mechanism of OmpR on prodigiosin biosynthesis was demonstrated. EnvZ/OmpR was demonstrated to be a new two-compound system that can directly positively regulate prodigiosin production in *S. marcescens* FZSF02.

# Data availability statement

The original contributions presented in the study are included in the article/Supplementary material, further inquiries can be directed to the corresponding author.



indicates that the promoter probe was not incubated with OmpR protein before treatment with DNase I, and the OmpR (+) group indicates that the promoter probe was incubated with OmpR protein before treatment with DNase I.

# Author contributions

XJ and JC designed the study, wrote the manuscript, and analyzed the results. XJ, FL, JL, CL, and KZ performed the experiments. All authors contributed to the article and approved the submitted version.

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# Conflict of interest

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.

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### References

Bang, I. S., Audia, J. P., Park, Y. K., and Foster, J. W. (2002). Autoinduction of the ompR response regulator by acid shock and control of the *Salmonella enterica* acid tolerance response. *Mol. Microbiol.* 44, 1235–1250. doi: 10.1046/j.1365-2958. 2002.02937.x

Beier, D., and Gross, R. (2006). Regulation of bacterial virulence by twocomponent systems. *Curr. Opin. Microbiol.* 9, 143–152. doi: 10.1016/j.mib. 2006.01.005

Bhagirath, A. Y., Li, Y., Patidar, R., Yerex, K., Ma, X., Kumar, A., et al. (2019). Two component regulatory systems and antibiotic resistance in gram-negative pathogens. *Int. J. Mol. Sci.* 20:1781. doi: 10.3390/ijms20071781

Boyce, K. J., Cao, C., and Andrianopoulos, A. (2016). Two-component signaling regulates osmotic stress adaptation via SskA and the high-osmolarity glycerol MAPK pathway in the human pathogen *Talaromyces marneffei. mSphere* 1, e00086–e00015. doi: 10.1128/mSphere.00086-15

Cai, S. J., and Inouye, M. (2002). EnvZ-OmpR interaction and osmoregulation in *Escherichia coli. J. Biol. Chem.* 277, 24155–24161. doi: 10.1074/jbc.M110715200

Cameron, A. D., and Dorman, C. J. (2012). A fundamental regulatory mechanism operating through OmpR and DNA topology controls expression of *salmonella* pathogenicity islands SPI-1 and SPI-2. *PLoS Genet.* 8:e1002615. doi: 10.1371/journal.pgen.1002615

Darshan, N., and Manonmani, H. K. (2015). Prodigiosin and its potential applications. J. Food Sci. Technol. 52, 5393–5407. doi: 10.1007/s13197-015-1740-4

Dhiman, A., Gopalani, M., and Bhatnagar, R. (2015). WalRK two component system of *Bacillus anthracis* responds to temperature and antibiotic stress. *Biochem. Biophys. Res. Commun.* 459, 623–628. doi: 10.1016/j.bbrc.2015.02.159

Dixon, R. (1998). The oxygen-responsive NIFL-NIFA complex: a novel twocomponent regulatory system controlling nitrogenase synthesis in gammaproteobacteria. *Arch. Microbiol.* 169, 371–380. doi: 10.1007/s002030050585

Doolittle, R. F., Feng, D. F., Tsang, S., Cho, G., and Little, E. (1996). Determining divergence times of the major kingdoms of living organisms with a protein clock. *Science* 271, 470–477. doi: 10.1126/science.271.5248.470

Du, Z., Zhang, M., Qin, Y., Zhao, L., Huang, L., Xu, X., et al. (2022). The role and mechanisms of the two-component system EnvZ/OmpR on the intracellular survival of *Aeromonas hydrophila*. J. Fish Dis. 45, 1609–1621. doi: 10.1111/jfd.13684

Fineran, P. C., Slater, H., Everson, L., Hughes, K., and Salmond, G. P. (2005). Biosynthesis of tripyrrole and beta-lactam secondary metabolites in *Serratia*: integration of quorum sensing with multiple new regulatory components in the control of prodigiosin and carbapenem antibiotic production. *Mol. Microbiol.* 56, 1495–1517. doi: 10.1111/j.1365-2958.2005.04660.x

Gristwood, T., Fineran, P. C., Everson, L., Williamson, N. R., and Salmond, G. P. (2009). The PhoBR two-component system regulates antibiotic biosynthesis in *Serratia* in response to phosphate. *BMC Microbiol.* 9:112. doi: 10.1186/1471-2180-9-112

Groisman, E. A. (2016). Feedback control of two-component regulatory systems. Annu. Rev. Microbiol. 70, 103–124. doi: 10.1146/annurev-micro102215-095331

Horng, Y. T., Chang, K. C., Liu, Y. N., Lai, H. C., and Soo, P. C. (2010). The RssB/ RssA two-component system regulates biosynthesis of the tripyrrole antibiotic, prodigiosin, in *Serratia marcescens. Int. J. Med. Microbiol.* 300, 304–312. doi: 10.1016/j.ijmm.2010.01.003

Jaworska, K., Ludwiczak, M., Murawska, E., Raczkowska, A., and Brzostek, K. (2021). The regulator OmpR in *Yersinia enterocolitica* participates in iron homeostasis by modulating fur level and affecting the expression of genes involved in iron uptake. *Int. J. Mol. Sci.* 22:1475. doi: 10.3390/ijms22031475

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# Supplementary material

The Supplementary material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fmicb.2022.1041146/ full#supplementary-material

Jia, X., Liu, F., Zhao, K., Lin, J., Fang, Y., Cai, S., et al. (2021). Identification of essential genes associated with prodigiosin production in *Serratia marcescens* FZSF02. *Front. Microbiol.* 12:705853. doi: 10.3389/fmicb.2021.705853

Ko, D., and Choi, S. H. (2022). Mechanistic understanding of antibiotic resistance mediated by EnvZ/OmpR two-component system in *salmonella enterica* serovar Enteritidis. *J. Antimicrob. Chemother.* 77, 2419–2428. doi: 10.1093/jac/dkac223

Kruppa, M., Krom, B. P., Chauhan, N., Bambach, A. V., Cihlar, R. L., and Calderone, R. A. (2004). The two-component signal transduction protein Chk1p regulates quorum sensing in *Candida albicans. Eukaryot. Cell* 3, 1062–1065. doi: 10.1128/EC.3.4.1062-1065.2004

Lai, H. C., Soo, P. C., Wei, J. R., Yi, W. C., Liaw, S. J., Horng, Y. T., et al. (2005). The RssAB two-component signal transduction system in *Serratia marcescens* regulates swarming motility and cell envelope architecture in response to exogenous saturated fatty acids. *J. Bacteriol.* 187, 3407–3414. doi: 10.1128/JB.187.10.3407-3414.2005

Maeda, S., Takayanagi, K., Nishimura, Y., Maruyama, T., Sato, K., and Mizuno, T. (1991). Activation of the osmoregulated *ompC* gene by the OmpR protein in *Escherichia coli*: a study involving synthetic OmpR-binding sequences. J. Biochem. 110, 324–327. doi: 10.1093/oxfordjournals.jbchem.a123579

Najnin, T., Siddiqui, K. S., Taha, T., Elkaid, N., Kornfeld, G., Curmi, P. M., et al. (2016). Characterization of a temperature-responsive two component regulatory system from the *Antarctic archaeon, Methanococcoides burtonii. Sci. Rep.* 6:24278. doi: 10.1038/srep24278

Nieckarz, M., Kaczor, P., Jaworska, K., Raczkowska, A., and Brzostek, K. (2020). Urease expression in pathogenic *Yersinia enterocolitica* strains of bio-serotypes 2/O:9 and 1B/O:8 is differentially regulated by the OmpR regulator. *Front. Microbiol.* 11:607. doi: 10.3389/fmicb.2020.00607

Ochman, H., and Wilson, A. C. (1987). Evolution in bacteria: evidence for a universal substitution rate in cellular genomes. *J. Mol. Evol.* 26, 74–86. doi: 10.1007/BF02111283

Pan, X., Tang, M., You, J., Liu, F., Sun, C., Osire, T., et al. (2021). Regulator RcsB controls prodigiosin synthesis and various cellular processes in *Serratia marcescens* JNB5-1. *Appl. Environ. Microbiol.* 87, e02052–e02020. doi: 10.1128/AEM.02052-20

Pan, X., You, J., Tang, M., Zhang, X., Xu, M., Yang, T., et al. (2022). Improving prodigiosin production by transcription factor engineering and promoter engineering in *Serratia marcescens. Front. Microbiol.* 13:977337. doi: 10.3389/fmicb.2022.977337

Papon, N., and Stock, A. M. (2019). Two-component systems. Curr. Biol. 29, R724–R725. doi: 10.1016/j.cub.2019.06.010

Pardee, A. B., Jacob, F., and Monod, J. (1959). The genetic control and cytoplasmic expression of "Inducibility" in the synthesis of  $\beta$ -galactosidase by *E. coli. J. Mol. Biol.* 1, 165–178. doi: 10.1016/S0022-2836(59)80045-0

Paul, T., Bandyopadhyay, T. K., Mondal, A., Tiwari, O. N., and Bhunia, B. (2020). A comprehensive review on recent trends in production, purification, and applications of prodigiosin. *Biomass Conv. Bioref.* 12, 1409–1431. doi: 10.1007/s13399-020-00928-2

Prüß, B. M. (2017). Involvement of two-component signaling on bacterial motility and biofilm development. J. Bacteriol. 199, e00259–e00217. doi: 10.1128/JB.00259-17

Qin, L., Yoshida, T., and Inouye, M. (2001). The critical role of DNA in the equilibrium between OmpR and phosphorylated OmpR mediated by EnvZ in *Escherichia coli. Proc. Natl. Acad. Sci. U. S. A.* 98, 908–913. doi: 10.1073/pnas.98.3.908

Qiu, S., Jia, S., Zhang, F., Liu, X., Ran, T., Wang, W., et al. (2021). Two component system CpxR/a regulates the prodigiosin biosynthesis by negative control in *Serratia*  marcescens FS14. Biochem. Biophys. Res. Commun. 579, 136-140. doi: 10.1016/j. bbrc.2021.09.050

Sarwar, Z., and Garza, A. G. (2015). Two-component signal transduction systems that regulate the temporal and spatial expression of *Myxococcus xanthus* sporulation genes. *J. Bacteriol.* 198, 377–385. doi: 10.1128/JB.00474-15

Scharf, B. E. (2010). Summary of useful methods for two-component system research. *Curr. Opin. Microbiol.* 13, 246–252. doi: 10.1016/j.mib.2010.01.006

Shi, K., Fan, X., Qiao, Z., Han, Y., McDermott, T. R., Wang, Q., et al. (2017). Arsenite oxidation regulator AioR regulates bacterial chemotaxis toward arsenite in *Agrobacterium tumefaciens* GW4. *Sci. Rep.* 7:43252. doi: 10.1038/srep43252

Sola-Landa, A., Moura, R. S., and Martín, J. F. (2003). The two-component PhoR-PhoP system controls both primary metabolism and secondary metabolite biosynthesis in *Streptomyces lividans*. *Proc. Natl. Acad. Sci. U. S. A.* 100, 6133–6138. doi: 10.1073/pnas.0931429100

Stankovic, N., Senerovic, L., Ilic-Tomic, T., Vasiljevic, B., and Nikodinovic-Runic, J. (2014). Properties and applications of undecylprodigiosin and other bacterial prodigiosins. *Appl. Microbiol. Biotechnol.* 98, 3841–3858. doi: 10.1007/s00253-014-5590-1

Stella, N. A., Lahr, R. M., Brothers, K. M., Kalivoda, E. J., Hunt, K. M., Kwak, D. H., et al. (2015). *Serratia marcescens* cyclic AMP receptor protein controls transcription of EepR, a novel regulator of antimicrobial secondary metabolites. *J. Bacteriol.* 197, 2468–2478. doi: 10.1128/JB.00136-15

Stock, A. M., Robinson, V. L., and Goudreau, P. N. (1999). Two-component signal transduction. Annu. Rev. Biochem. 69, 183–215. doi: 10.1146/annurev.biochem.69.1.183

Sun, D., Zhou, X., Liu, C., Zhu, J., Ru, Y., Liu, W., et al. (2021). Fnr negatively regulates prodigiosin synthesis in *Serratia* sp. ATCC 39006 during aerobic fermentation. *Front. Microbiol.* 12:734854. doi: 10.3389/fmicb.2021.734854

Tierney, A. R., and Rather, P. N. (2019). Roles of two-component regulatory systems in antibiotic resistance. *Future Microbiol.* 14, 533–552. doi: 10.2217/fmb-2019-0002

Tipton, K. A., and Rather, P. N. (2017). An ompR-envZ two-component system ortholog regulates phase variation, osmotic tolerance, motility, and virulence in *Acinetobacter baumannii* strain AB5075. *J. Bacteriol.* 199, e00705–e00716. doi: 10.1128/JB.00705-16

Wang, S. T., Kuo, C. J., Huang, C. W., Lee, T. M., Chen, J. W., and Chen, C. S. (2021). OmpR coordinates the expression of virulence factors of Enterohemorrhagic *Escherichia coli* in the alimentary tract of *Caenorhabditis elegans. Mol. Microbiol.* 116, 168–183. doi: 10.1111/mmi.14698

Williamson, N. R., Fineran, P. C., Leeper, F. J., and Salmond, G. P. (2006). The biosynthesis and regulation of bacterial prodiginines. *Nat. Rev. Microbiol.* 4, 887–899. doi: 10.1038/nrmicro1531

Wright, G., Saeki, A., Hikima, T., Nishizono, Y., Hisano, T., Kamaya, M., et al. (2018). Architecture of the complete oxygen-sensing FixL-FixJ two-component signal transduction system. *Sci. Signal.* 11:eaaq0825. doi: 10.1126/scisignal.aaq0825

Yuchuan, M., Sayak, B., Tatsat, , and Banerjee, (2019). Wave patterns organize cellular protrusions and control cortical dynamics. *Mol. Syst. Biol.* 15:e8585. doi: 10.15252/msb.20188585