

## RESEARCH ARTICLE

# The isolation of the antagonistic strain *Bacillus australimaris* CQ07 and the exploration of the pathogenic inhibition mechanism of *Magnaporthe oryzae*

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## Abstract

Biological control as a promising method to combat plant disease has gained public attention in recent years. In the present study, we isolated 12 strains resistant to *Magnaporthe oryzae* from western Sichuan subalpine soil. Among them, CQ07 exhibited remarkable activity against *M. oryzae*. The result of 16S rRNA sequence analysis revealed that CQ07 is approximately 99% similar to *Bacillus australimaris*. The sterilized culture filtrate of CQ07 inhibited the growth of *M. oryzae*, which motivated us to deduce the influence of CQ07 on the pathogenicity of *M. oryzae*. As shown by experimentation, sterilized culture filtrate (10 µl/ml) of CQ07 can delay and even suppress the germination of conidia and prevent the formation of appressorium in vitro and in vivo. In addition, by simulative field tests, the spraying of conidia suspension diluted with sterilized culture filtrate of CQ07 reduced infection of rice blast. To better control rice blasts, understanding the infection mechanism of *M. oryzae* and inhibiting the mechanism of the antagonistic strain is of great importance.

## Introduction

Rice blast caused by the filamentous ascomycete *Magnaporthe oryzae* is one of the most destructive diseases of rice around the world[1]. Outbreaks of rice blast disease are a serious and recurrent problem in China and other rice-growing regions[2]. Therefore, many people use antifungal drugs and plant disease-resistant cultivars to reduce the loss caused by rice blasts. On the one hand, chemical antifungal drugs have many advantages, such as high efficiency and inexpensive use. On the other hand, the overuse of chemical fungicides not only pollutes the environment but also makes pathogenic fungi increasingly drug resistant. Although the introduction of resistant cultivars can prevent disease efficiently, new variations may lead to the appearance of new pathogenic bacteria[3].

With increasing focus on sustainable development in agriculture, biological control as an environmentally friendly and promising strategy has emerged to the public[4]. In recent

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studies, there has been an increasing interest in the exploitation of rice blasts for biological control by using plant beneficial microorganisms, owing to their low toxicity and a lack of pathogen resistance[5]. At present, *Bacillus* have been found to be considerable biological control candidates in agriculture, with their high production of antifungal substances and resistance to extreme conditions. Earlier studies indicated that *Bacillus* can increase plant resistance to pathogenic fungi by colonizing plants or spraying a powder made from *Bacillus* on a plant, which exerts a biological control effect. In addition, *Bacillus* can not only produce antifungal substances but also promote the growth and increase the yield of plants[6–8]. Therefore, the potential value of *Bacillus* in biological control is immeasurable.

Many studies are currently devoted to isolating antifungal components from *Bacillus* or colonizing plants with *Bacillus* to control rice blast[9–11], but little research has been performed on inhibiting pathogenic mechanisms. *M. oryzae* is a model organism for molecular biology investigations and studying the pathogenesis of filamentous fungi. To control rice blast disease effectively, understanding the infection process and mechanisms of pathogenicity are important premises. The process of infecting a rice host mainly includes contact, invasion and expansion. The spread of rice blasts is often caused by the dissemination of *M. oryzae* conidia through the air. Conidia attach to the leaf surface, and then, a dome-shaped cell with a complex cell wall structure, an appressorium, contacted by the cuticle is formed at the ends of germ tubes, which can penetrate directly through the host cuticle by its enormous turgor pressure[12, 13]. Observation of the formation process of the infection structure and real-time monitoring of the dynamic growth of rice blasts in host rice tissue will help reveal the interaction process between *M. oryzae* and rice, which is of great significance for the biological control of rice blasts. Green fluorescent protein (GFP) has been widely used as a labeling tool in life science research, so the GFP-labeled *M. oryzae* strain Guy11 was used for the following tests.

A strain of bacteria displaying strong suppression against *M. oryzae* was selected from western Sichuan subalpine soil and named CQ07. Through the examination of the morphological, biochemical and physiological properties and 16S rRNA sequence analysis, the strain was identified. Our study observed germination of conidia, formation of an appressorium, growth of mycelium, cellular permeability in vitro and infection process in plant tissues of *M. oryzae*. Furthermore, inhibition tests of sterilized CQ07 culture filtrate on infection structure development and infection process on plant tissue were conducted to confirm the potential biological control value of the strain.

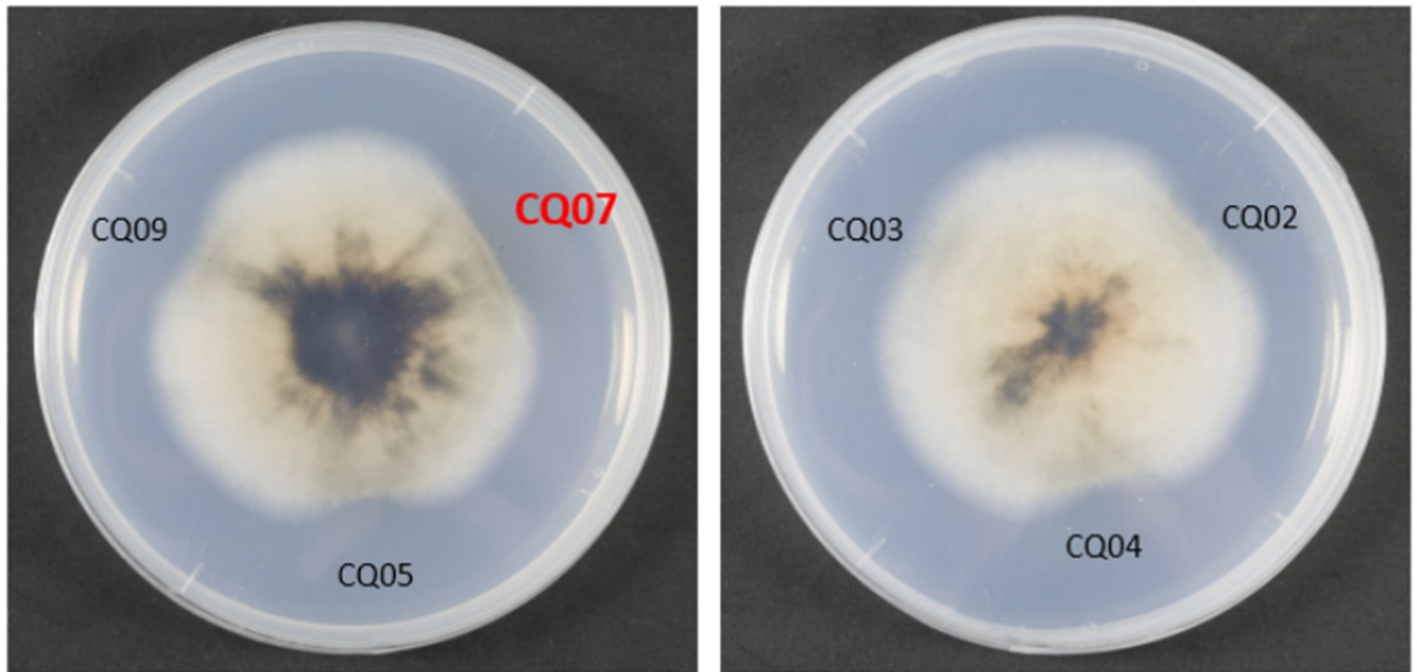
## Results

### Isolation and screening of strains antagonistic to *M. oryzae*

The soil was a mixture of soil from 800 to 3000 meters above sea level provided by the College of Resource Science and Technology of Sichuan Agricultural University. Twelve strains of bacteria antagonistic to the growth of *M. oryzae* were isolated from western Sichuan subalpine soil. Six sterilized culture filtrates of different colonies displayed obvious inhibitory effects against *M. oryzae* Guy11 by testing for mycelial growth inhibition. Among the antagonistic bacteria, the inhibition of CQ07 was particularly strong (Fig 1).

### Identification of strain CQ07

16S rRNA sequence analysis indicated that strain CQ07 displayed approximately 99% similarity to *Bacillus australimaris*. A phylogenetic tree displaying the relationship between strain CQ07 and other strains is shown in Fig 2. The morphological and physiological characteristics of CQ07 are summarized in Fig 3 and Table 1. The results showed that CQ07 was a strain with an orange-red center and feathery edges. Gram and spore staining tests were positive;



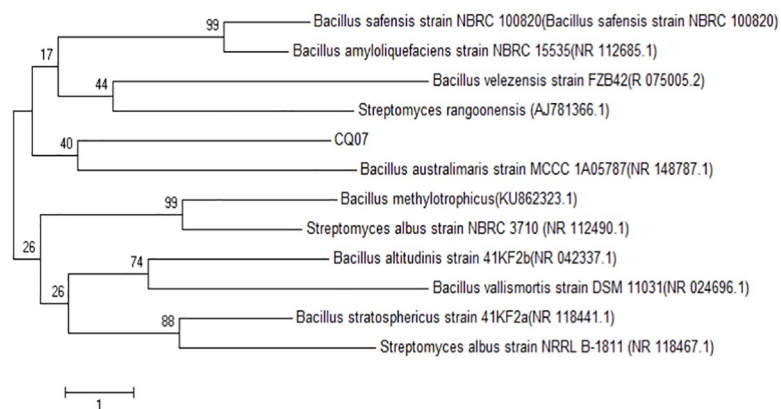
**Fig 1. CQ07 showed a strong inhibitory effect against *M. oryzae* Guy11.** CQ02, CQ03, CQ04, CQ05, CQ07, and CQ09 all showed inhibition against *M. oryzae* GUY11, but CQ07 had the strongest inhibition effect.

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therefore, CQ07 was gram-positive and spore-producing. The capsule staining was negative, which suggests that CQ07 has no capsule.

### Germination of conidia

As shown in Fig 4, the germination rate of the control group reached 93% after 2 h, while the germination rate of the treatment was 0%. At 4 hours, the treatment group germination rate reached 27%, while the control group reached 96% and formed the appressorium. The germination rate after 8 hours of treatment reached 83% and was most stable in this range. However, the control group almost always achieved 97%. On the basis of these results, we can conclude that the culture filtrate of CQ07 can delay even suppress the germination of conidia.



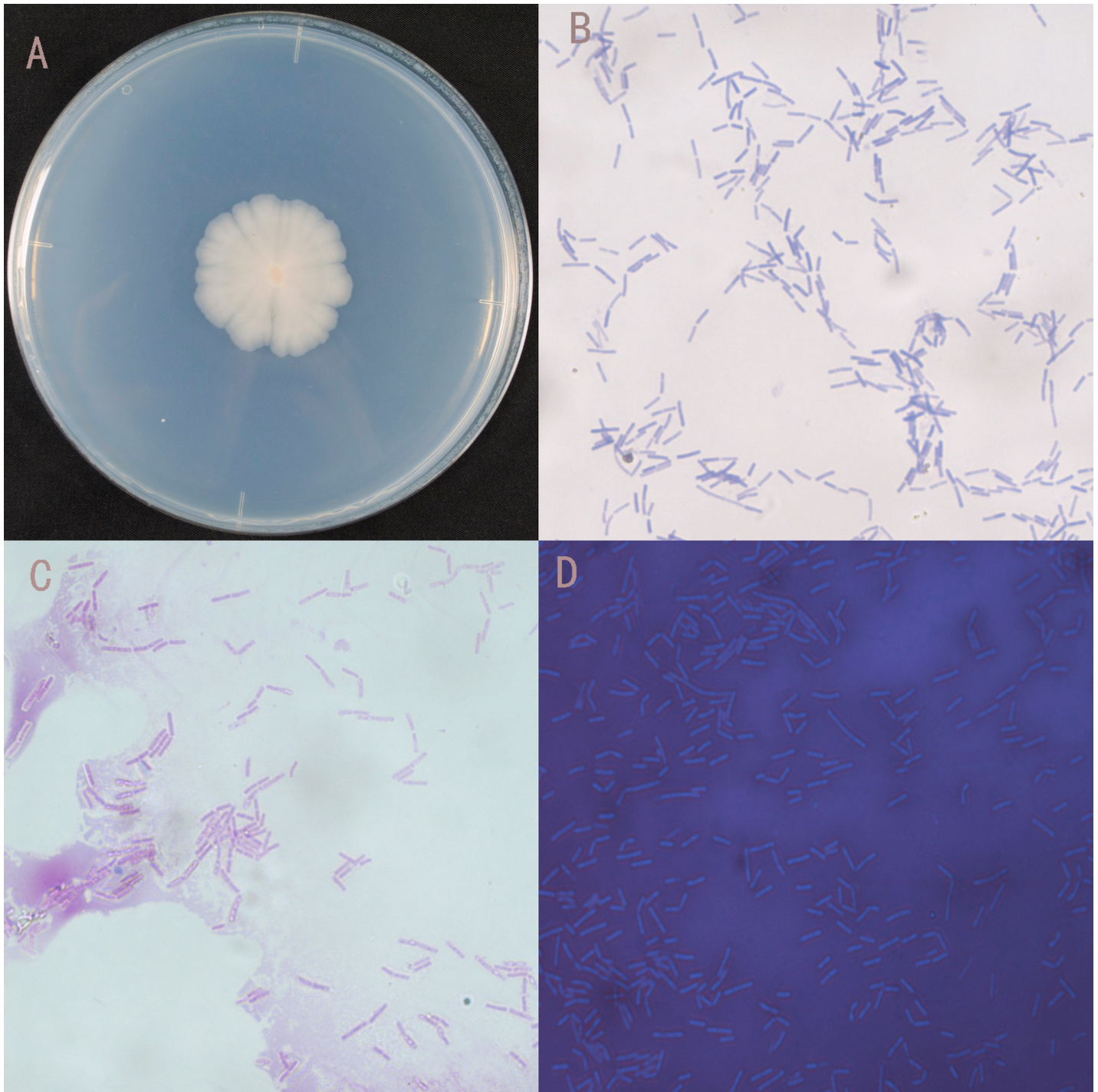
**Fig 2. Neighbor-joining phylogenetic tree of CQ07 based on 16S rRNA sequence analysis.**

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### Formation of appressorium

As shown in Fig 4 and Fig 5, the treatment group (Fig 5B) did not form an appressorium after 4 h, whereas the control group had an approximately 85% formed appressorium (Fig 5A). Even after 24 h, the appressorium was still not observed, except for the endlessly extended and unusual mycelium growth.



**Fig 3. Morphological characteristics of CQ07.** A. Colony morphology of CQ07. B. Gram staining of CQ07. C. Spore staining of CQ07. D. Capsule staining of CQ07.

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**Table 1. Physiological and biochemical results of strains CQ07.**

Items	CQ07
Shape	Rod
Anaerobic growth	+
Motility	+
$\beta$ -Galactoside	-
Utilization of lactose	-
Utilization of raffinose	-
Nitrate reduction to nitrite	+
Utilization of citrate	-
Mannitol	-
Indole test	-
L-Arabinose	-
Xylose	+
Oxidase activity	+
Decomposition of starch	+
Methyl red (MR) test	-
Voges-Proskauer (VP) tests	+
Growth in 7% NaCl	-
Growth in 8% NaCl	-
Growth in 9% NaCl	-

+: positive

-: negative

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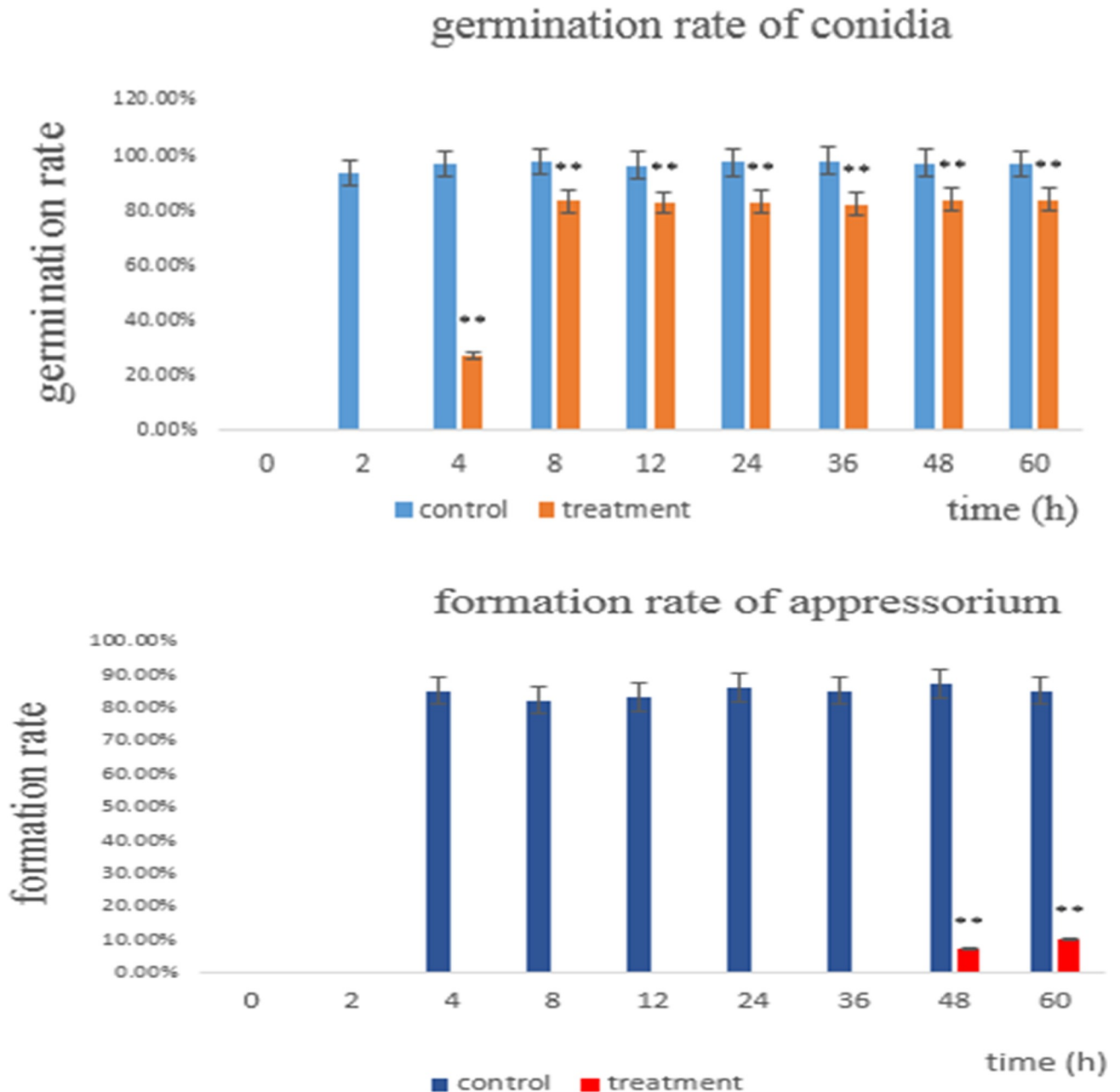
However, at 48 h and 60 h, 7% and 10% appressorium of the treatment, respectively, was found. This result also revealed why rice is still mildly infected when testing isolated leaves and living leaves and indicated that the sterilized culture filtrate of CQ07 delayed and inhibited the majority of appressorium formation, which is a necessary infection structure for the pathogenic ability of *M. oryzae*.

### The integrity of the cell membrane was destroyed by CQ07 culture filtrate

FDA and PI are two types of biological compound dyes that stain dead and alive cells, displaying different fluorescence through the cellular membrane permeability principle [14]. As shown in Fig 6, the cell membrane of *M. oryzae* was destroyed by CQ07 culture filtrate. Combining this test with the appressorium formation test, the mycelium displayed unusual growth in both these two tests. Furthermore, the cell membrane permeability test suggested that the permeability of swell parts was also destroyed.

### Observation of the infection process

In agreement with germination and appressorium formation tests on hydrophobic slides in vitro, the conidia on onion epidermis did not germinate in the expected amount of time or form appressorium at the end of the growth period. As shown in Fig 7, the appressorium of the control group had already formed, and secondary mycelium extended into the onion cells at 24 h, whereas the mycelium infected the neighboring cells at 36 h. As shown in the treatment group shown in Fig 7, the treatment group only extended the malformed mycelium on the surface of the onion cells. From above, we can speculate that the filtrate of CQ07 still inhibits the activity of conidia in vivo.



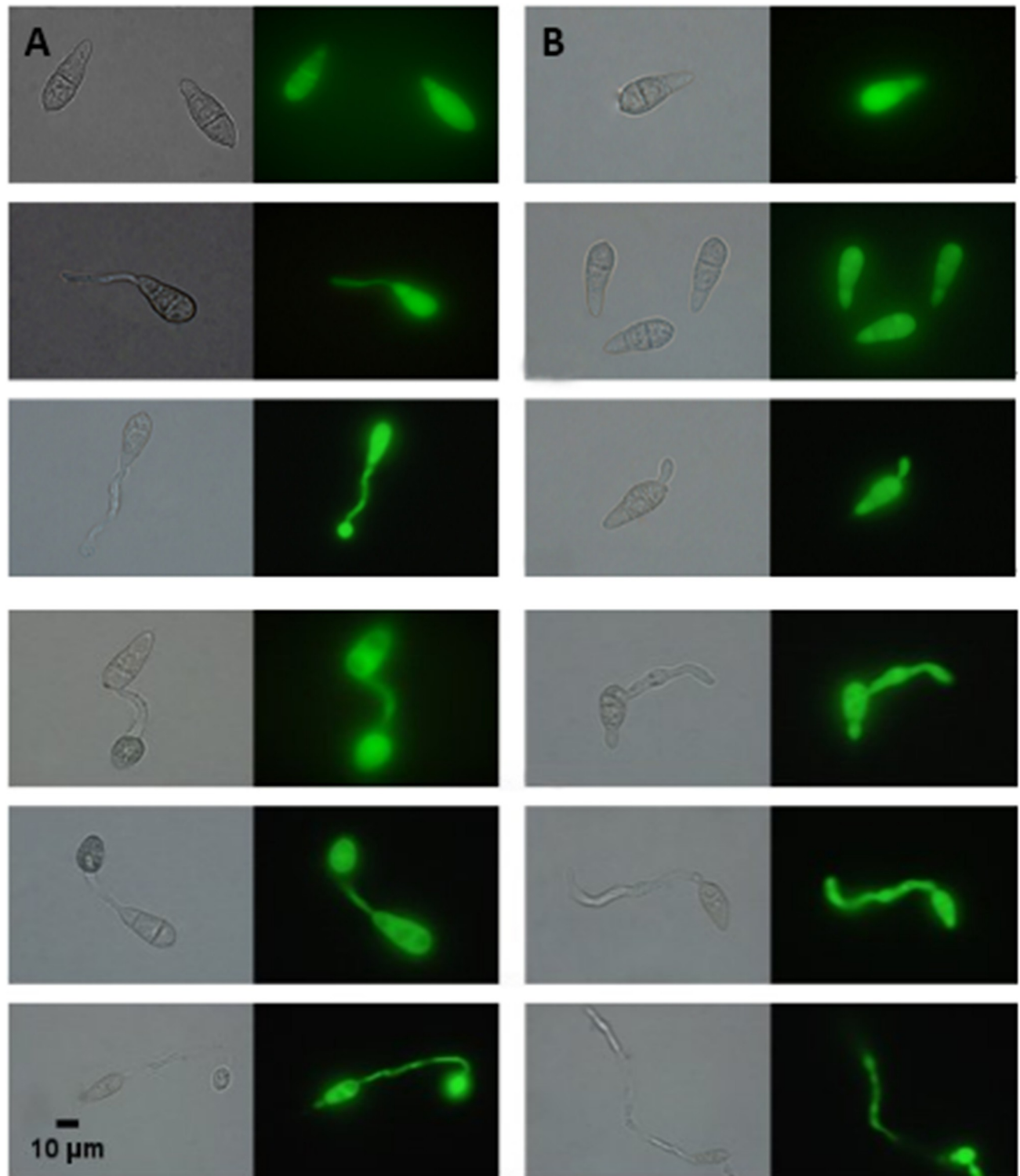
**Fig 4. Germination rate of conidia at different times.** The control group had already germinated after 2 hours, while the treatment group had not yet germinated. At 4 hours, the treatment group began to germinate, but the rate of germinated conidia was only 27%. After 8 hours, the germination rate of the treatment group was almost always stable at 83%, while the germination rate of the control group was 97%.

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### Evaluation of inhibition in vivo

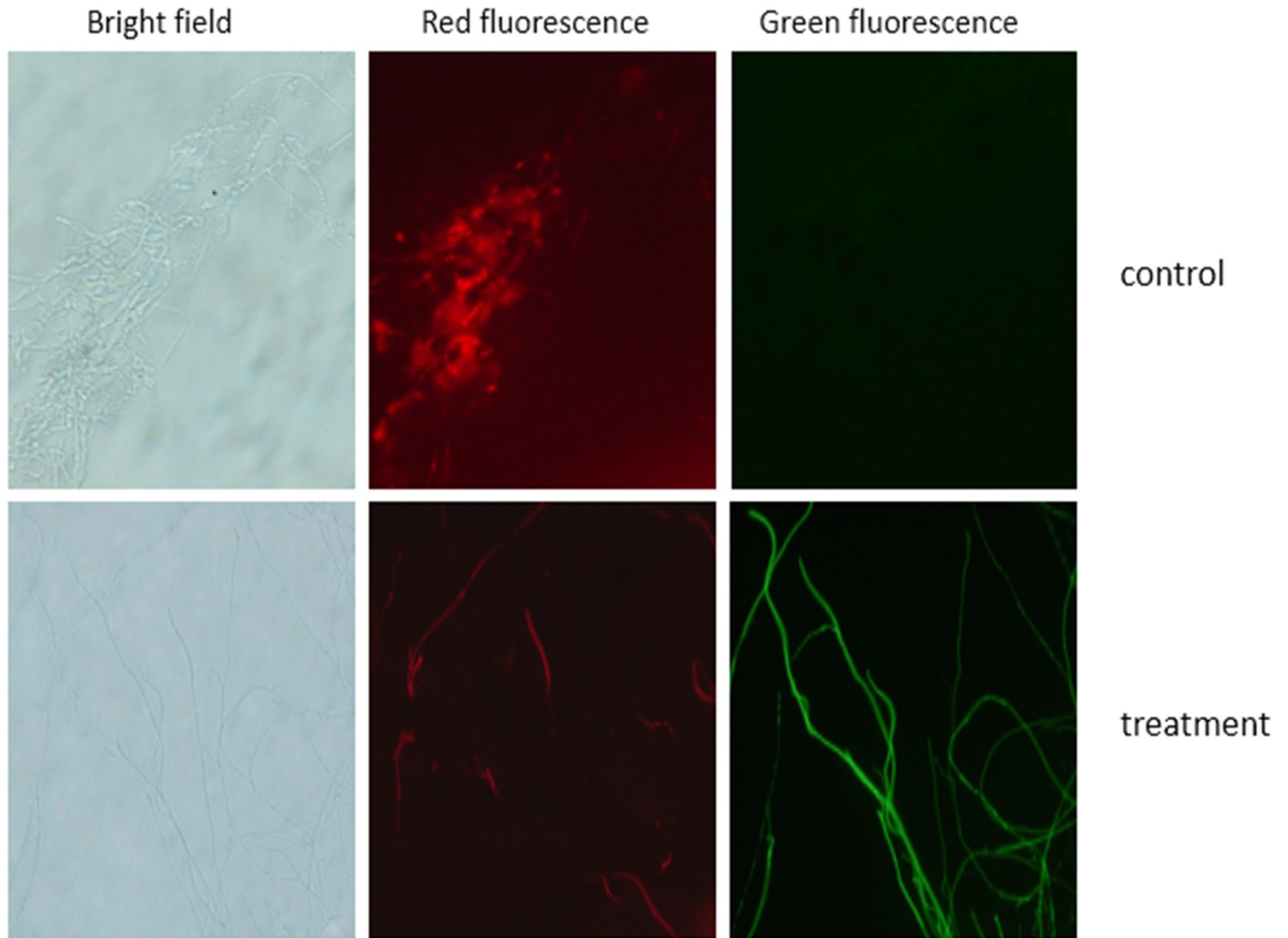
The inhibitory pathogenicity of CQ07 against rice blasts is shown in Fig 8. In Fig 8, compared with the control group, the lesion diameter on isolated leaves of the treatment group was obviously smaller than that of the control group. The average lesion diameter of the control group in isolated leaves reached 10.3 mm, while the diameter of the treatment group was only 2.8 mm. The lesion quantity on living leaves of the control group was 24 lesions per leaf on average, while the lesion quantity of treatment was 6 lesions per leaf on average. This phenomenon





**Fig 5. A. Germination of conidia at different time points in the control group. B. Germination of conidia at different time points in the treatment group.** At 4 hours, 85% of appressorium of the control group was formed. After 8 hours, endlessly extended and unusual mycelium growth were observed in the treatment group. After 24 hours, the appressorium of the treatment group still did not form. However, after 48 hours, 7% of appressorium was observed in the treatment group. After 60 hours, the appressorium formation rate of the treatment group reached only 10%, while that of the control group reached 85%.

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**Fig 6.** The cell membrane of the treatment group was destroyed by the culture filtrate of CQ07. Through the FDA-PI staining test, the mycelium of the control group showed the most green fluorescence. The mycelia of the treatment group all showed red fluorescence.

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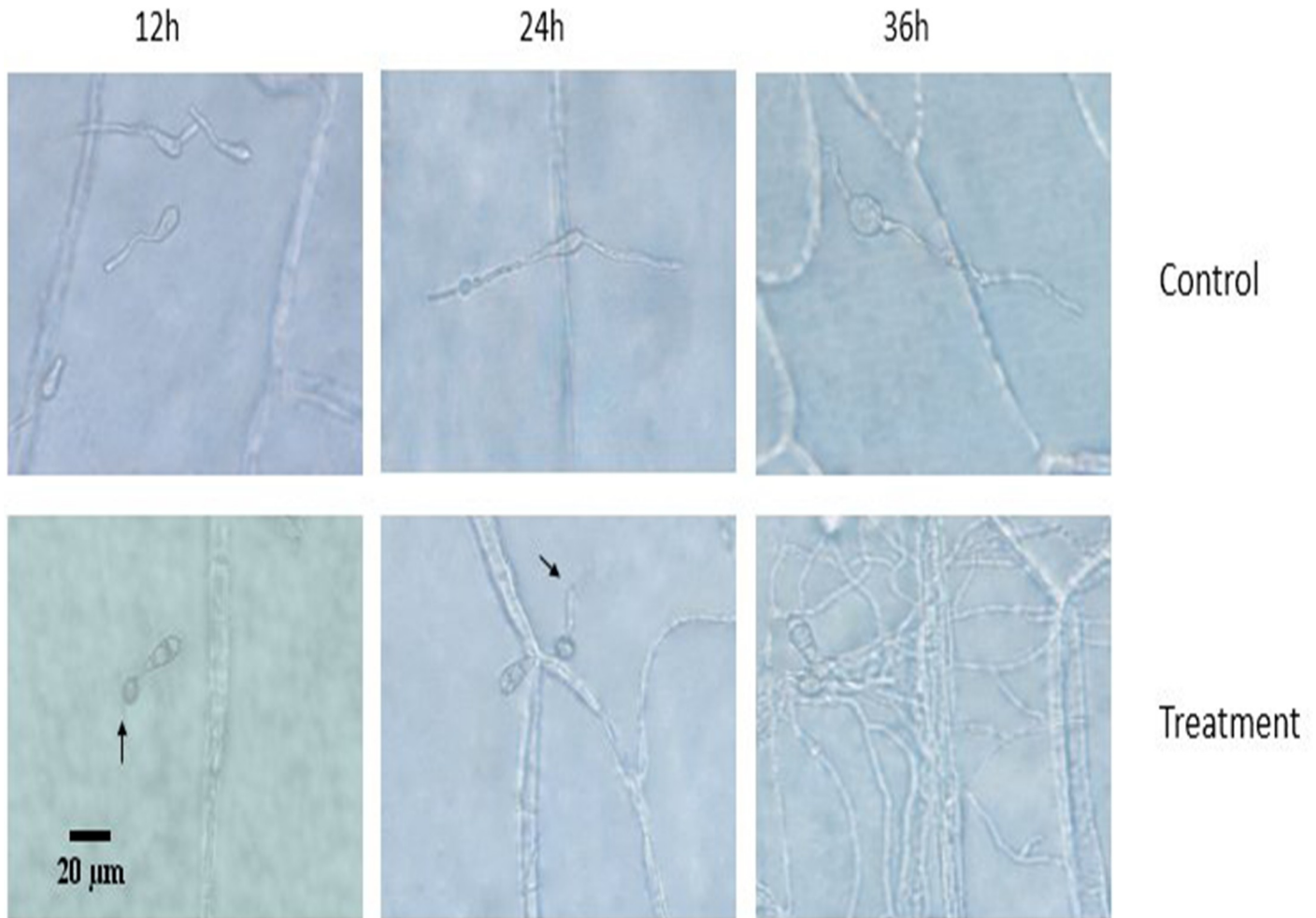
indicated that the sterilized culture filtrate of CQ07 had a strong ability to inhibit the pathogenicity of *M. oryzae*, and the number of lesions decreased by 75% compared with that in the control group.

## Discussion

In nature, antagonism among different species is ubiquitous. For a long time, interest has focused on taking full advantage of antagonism, applying it to many fields such as agricultural defense or the medical domain. Plant fungal diseases are difficult to control, which leads to very large losses in economic crops. The leading strategy applied to control plant fungal disease is chemical pesticides, which cause serious environmental pollution[15]. Thus, strategies to protect plants from *M. oryzae* infection should be developed, especially environmentally friendly biological control methods.

Rice blasts have received considerable attention because of the increasing importance of food security worldwide, especially in East Asia. *M. oryzae* is one of the causal pathogenic fungi of rice blast disease. Host plants of *M. oryzae* are not only rice but also other Poaceae crops and their wild relatives, such as wheat, barley, finger millet, foxtail and wild grasses. The losses of rice harvest caused by rice blast reached 10–30% each year[16–18].



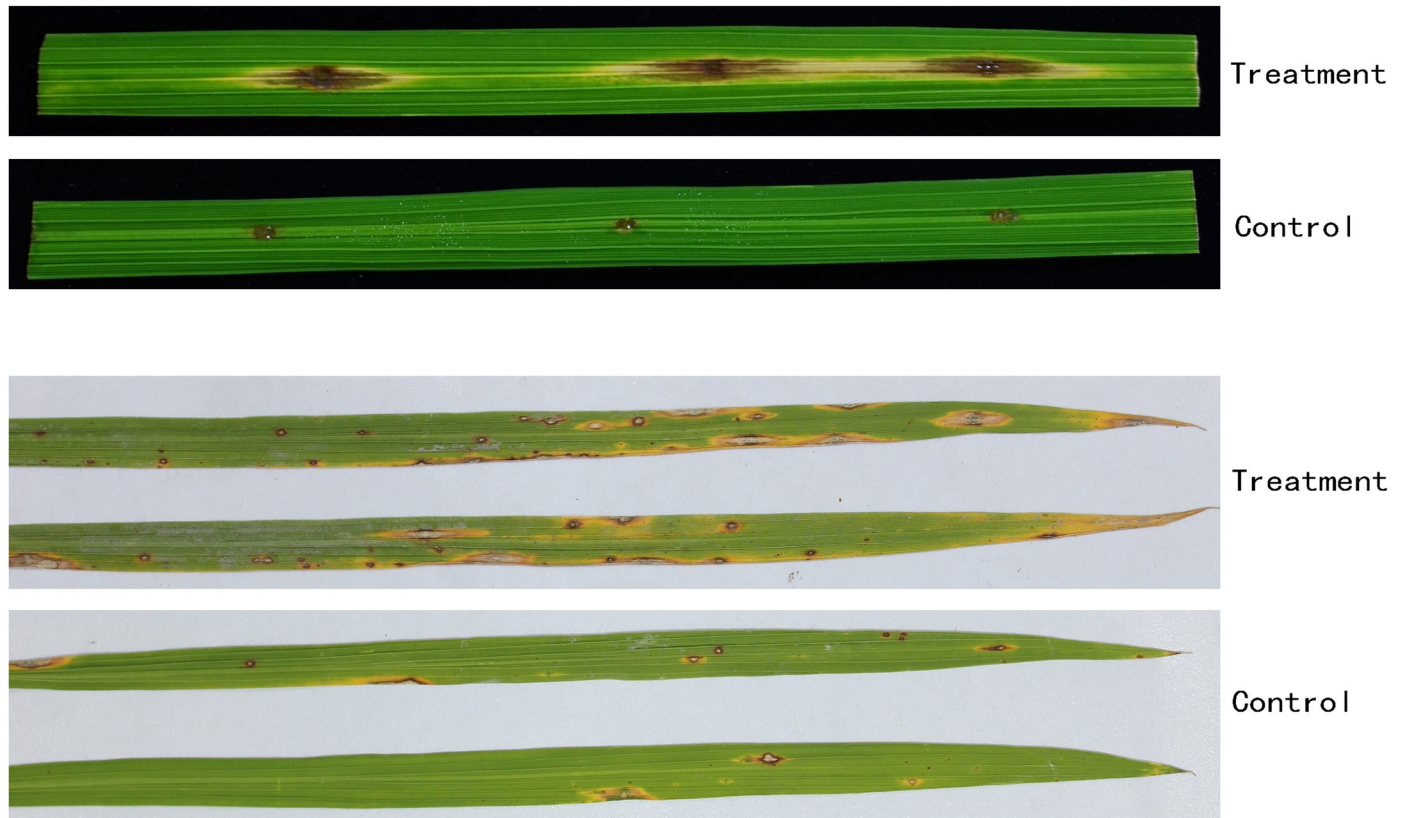


**Fig 7. Infection status of conidia at different times.** At 12 hours, appressorium of the control group was observed. At 24 hours, infection pegs of the control group were observed. At 36 hours, the mycelium of the control group infected the neighboring cells. At 24 and 36 hours, the treatment group only extended the malformed mycelium on the surface of the onion cells.

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Many species of bacteria, especially *Bacillus*, are known as biological control agents that inhibit several phytopathogenic fungi due to the production of cell wall-degrading enzymes and other antifungal metabolic products [19–21]. Many *Bacillus* strains antagonistic to fungi have been isolated from soil [10, 22–24]. In our current study, several antagonism strains were isolated from western Sichuan subalpine soil. CQ07 was one of the most active strains that inhibited the growth of *M. oryzae*. The sterilized culture filtrate of CQ07 also significantly inhibited the growth of *M. oryzae* on nutrient medium. In the present study, based on the physiological and biochemical characteristics and 16S rRNA sequence analysis, CQ07 was identified as *B. australimaris*. To the best of our knowledge, this is the first isolation of *B. australimaris* from western Sichuan subalpine soil.

Germination of conidia and formation of appressorium are two crucial processes of *M. oryzae* during plant infection. However, through the results of the conidia germination test, the process of germination was obviously delayed and even completely suppressed. The conidia treated with sterilized culture filtrate did not germinate after 2 hours compared with those in the control group. After 4 hours, the control group had already formed a dome-shaped appressorium, while the treated group reached a germination rate of only 27% and exhibited a short



**Fig 8. Pathogenicity of conidia on isolated and living leaves.** The experiment on isolated leaves showed that the average lesion diameter of the control group was 10.3 mm, while the diameter of the treatment group was only 2.8 mm. The experiment on living leaves showed 24 lesions on average per leaf in the control group and 6 lesions on average per leaf in the treatment group.

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germ tube. At 8 hours and afterwards, the germ tube of the treated group began to extend without limits, and unusual growths emerged on the mycelium. Only after 48 hours, there was 7% appressorium began to form. The results shown above definitely illustrate that the germination of conidia was delayed and inhibited and the formation of appressorium was hindered by the crude fermentation filtrate of CQ07. Therefore, we can deduce that the filtrate of CQ07 can inhibit the formation of infectious structures of *M. oryzae* in vitro.

Cellular activity is observed by cell permeability, and the permeability of *M. oryzae* was destroyed by the culture filtrate of CQ07 through the FDA-PI dyeing experiment. Dynamic observation of the infection process and pathogenicity test in vivo can verify the impact of the CQ07 filtrate against the pathogenicity of *M. oryzae*. Onion epidermis is an ideal material to observe the infection process because of its large cells, easy access and convenience to observe. Researchers have already utilized onion epidermis to examine the infection of *M. oryzae* [25, 26]. Therefore, we used onion epidermis to perform pathogenicity test observations. Through the test results, we observed that the conidia handled with the culture filtrate of CQ07 lost pathogenicity. From the image, we can see that the appressorium and infection peg were only observed in the control group at 12 h and 24 h. Only long and unusual mycelia were observed on the surface of onion cells in the treatment group.

As shown in Fig, the lesion of the control group was obviously larger than that of the treatment group. Furthermore, this result was consistent with the phenomenon in the living leaf experiment. The number of lesions in the treatment group was much lower than that in the

control group. By evaluating the infection situation, we found that the pathogenicity of *M. oryzae* was significantly inhibited.

From the above, we can deduce that CQ07 can inhibit not only the growth, germination, and formation of appressorium in vitro but also the pathogenicity of *M. oryzae* in vivo. Therefore, we can conclude that CQ07 might effectively inhibit infection by influencing the formation of the infectious structure of *M. oryzae*. However, the formation of infectious structures is regulated by various genes, and the process is complicated. In future studies, we will explore the molecular mechanism related to our present studies.

In summary, *B. australimaris* CQ07 is not only a candidate biological control agent, but its ability to inhibit pathogenicity has furthered the progress of pathogenesis research.

## Methods

### Isolation of antagonistic strains from soil

The western Sichuan subalpine soil used to isolate the bacteria was provided by the College of Resource Science and Technology of Sichuan Agricultural University. Then, 1 g of soil was suspended in 10 ml of sterilized water and vortexed for 1 min before centrifugation at 2000 rpm for 10 min. Then, 0.1 mL of supernatant was diluted and plated onto potato dextrose agar [27]. After cultivation at 37°C for 48 h in a homothermal incubator, strains that appeared on nutrient agar plates were isolated as single colonies on potato dextrose medium.

### Screening of the antagonistic bacteria against *M. oryzae* in vitro

The rice blast pathogenic fungus *M. oryzae* Guy11 was provided by the Plant Pathogenic Laboratory of Sichuan Agricultural University. To select the antagonistic strains, every strain was inoculated in homologous fluid nutrient medium and incubated at 28°C for 48 h in a rotary shaker at 180 rpm. One milliliter of sterilized culture filtrate from every strain was removed and stored in a small EP tube. Ten microliters of sterilized culture filtrate of each strain was removed from every small EP tube and injected into the edge of a *M. oryzae* colony, which had been deposited in the center of potato dextrose agar medium. After the sterilized culture filtrate of the antagonistic strain was injected, the *M. oryzae* medium was incubated at 28°C in an incubator. After 2 or 3 days, the inhibition of growth of *M. oryzae* was compared by observing the degree of diminished growth. The strongest antagonistic bacteria were named CQ07 and taken for further study.

### Morphological characteristics and 16S rRNA analysis of CQ07

The method of extracting total genomic DNA was based on the manufacturer's instructions of a commercial DNA extraction kit. The 16S rRNA of the strain was amplified using forward and reverse primers in a small EP tube. The polymerase chain reaction program proceeded in a PCR thermocycle instrument. The PCR products were purified using a PCR Purification Kit, and the PCR products were identified by horizontal electrophoresis on a 1% agarose gel. The fragments of amplified 16S rRNA were transformed into *Escherichia coli*. Sequences were sequenced by TSINGKE (Chengdu, China). The similarity of 16S rRNA sequences was compared using the BLAST search program in the NCBI database (<http://www.ncbi.nlm.nih.gov>).

A colony of CQ07 was observed on PDA medium. Bacteriological, biochemical and physiological properties were tested according to Bergey's Manual.

### Germination test of conidia

The sterilized culture filtrate and sterilized water were used to scrape the conidia from 10-day-old *M. oryzae* conidia-producing medium, which was adjusted to a concentration of  $1 \times 10^5$

conidia/ml after filtration with two layers of lens paper[28]. Fifty microliters of conidia suspension was dropped on the hydrophobic cover slide. Then, the hydrophobic cover slides were placed in a moist petri dish to germinate at room temperature. After 0, 2, 4, 8, 12, 24, 36, 48, and 60 hours, the germination degree of conidia was observed under a ZEISS microscope, and 100 conidia were selected to calculate the germination rate. The experiment was repeated three times.

The formula for the germination rate was as follows:

$$\text{Germination rate (\%)} = [R1/R2] \times 100\%$$

where R1 = the number of germinated conidia and

R2 = the number of total conidia.

### Formation of appressorium

Conidia were scraped with sterilized culture filtrate and sterilized water from 10-day-old *M. oryzae* conidia-producing medium, then the conidia suspension was diluted to  $1 \times 10^5$  conidia/ml. Fifty microliters of conidia per droplet were dripped on a hydrophobic cover slide. After incubation at room temperature in a moist petri dish for 0, 2, 4, 8, 12, 24, 36, 48, and 60 hours, appressorium formation was observed under a ZEISS fluorescence microscope. The experiment was repeated three times.

The formula for appressorium formation rate was as follows:

$$\text{Formation rate of appressorium(\%)} = [X_1/X_2] \times 100\%$$

X<sub>1</sub>: the number of conidia that formed appressorium;

X<sub>2</sub>: the number of total conidia.

### Experiment of cell activity

*M. oryzae* was cultivated with 99 ml of fluid medium at 28°C for 24 h, and then 1 ml of sterilized culture filtrate was added to the fluid medium. After the *M. oryzae* continued cultivation in the mixed medium for 24 h, the mycelia of *M. oryzae* were cut off and dyed with FDA-PI composite dyes[14] for 5~10 min under dark conditions before being observed under a ZEISS fluorescence microscope at 20 times magnification. The experiment was repeated three times.

### Experiment of the infection process

The isolated strain that demonstrated the greatest suppression activity against *M. oryzae* in vitro was observed with respect to its ability to inhibit rice blast in plant tissue. The onion epidermis was divided into two groups, and each group had 3 epidermis samples. The epidermis was injected with a 50 µl of conidia suspension (concentration of  $1 \times 10^5$  conidia/ml) that was treated with two kinds of liquid as detailed below: sterilized water and sterilized culture filtrate of CQ07. The infection status was observed under a ZEISS microscope at the corresponding time (12, 24, and 36 h). The experiment was repeated three times.

### Experiment on isolated leaves

Rice seedling were grown at artificial climate chamber (30°C) for 30 days. Leaves of rice without obvious symptoms of disease were placed in petri dishes with a solution of 6-benzylamino-purine. Prior to that, every leaf was poked slightly with a needle, which made leaves more susceptible to *M. oryzae* infection.

Treatment group: a 5 µL volume of *M. oryzae* conidia suspension ( $1 \times 10^5$  conidia/mL) droplets was diluted with a sterilized culture filtrate of CQ07 and applied to the slightly punctured sites of the leaves. Then, the leaves were incubated at 25°C in the dark for 24 h.



Subsequently, all leaves were incubated in the light at 25°C[29]. In the control group, sterile water was used instead of the sterile culture filtrate, and other steps were unchanged. After 5 days, lesion length was compared.

The experiment was repeated three times.

### Experiment on living leaves

Considering that experiments in the isolated leaves do not accurately reflect inhibitory abilities under field conditions, we carried out simulate field tests aiming to confirm the biological control effects of CQ07. The experiment was finished under growth climate chamber conditions. The bioassay was conducted in accordance with previous methods[30]. First, 30-day-old rice seedlings were transferred to an inoculation chamber(30°C). In the treatment experiment, the rice seedlings were sprayed to runoff with the conidia suspension of *M. oryzae* ( $1 \times 10^5$  conidia/mL) diluted by sterilized culture filtrate of CQ07 containing Tween 20. The control group was treated in the same way, but the sterile water was in place of the culture filtrate. After 5 days, lesion quantity was compared (both 100 leaves of control group and treatment group were counted; the experiment was repeated three times).

The formula for the inhibition rate was as follows:

$$\text{Inhibition rate (\%)} = [N_1 - N_2 / N_1] \times 100\%$$

$N_1$ : lesion quantity before treatment;

$N_2$ : lesion quantity after treatment

### Supporting information

**S1 Data. Raw data for Fig 4.**

(XLSX)

**S2 Data. Raw data of Fig 8.**

(XLSX)

**S1 File. 16S rRNA sequence of CQ07.**

(DOCX)

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### Author Contributions

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**Formal analysis:** Rongjun Chen, Zhengjun Xu.

**Investigation:** Wenqian Chen, Ying Guan.

**Methodology:** Lu Zhao, Hong Xu, Ying Guan.

**Project administration:** Lu Zhao.

**Resources:** Yilun Dong.

**Software:** Lihua Li.

**Supervision:** Yilun Dong, Hong Xu, Songhao Rong.

**Validation:** Hui Li.

**Visualization:** Hui Li.

**Writing – original draft:** Wenqian Chen.

**Writing – review & editing:** Zhengjun Xu.

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