

A Real-time Fluorescent RPA Detection Method for *Rhizoctonia Solani* Based on ITS Sequences

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ABSTRACT

Rhizoctonia solani, a pathogen with a broad host range, propagates through its hyphae in the soil, invading plant roots or seeds, and subsequently affecting surrounding plants. Given that the pathogenic fungus continues to develop within the crop population after infection, early detection of the pathogen and timely prevention measures are crucial for reducing losses and preventing the spread of the disease. In this study RPA (Recombinase polymerase amplification) specific primers and fluorescent probe were designed based on the ITS sequence of *Rhizoctonia solani* and an RPA-real time detection method for *Rhizoctonia solani* was established. By optimizing the reaction conditions of RPA, the optimal conditions for RPA detection method were 39°C for 30 min. This assay can distinguish *Rhizoctonia solani* from other related species. The sensitivity of RPA-real time detection method for *Rhizoctonia solani* can be achieved at 6×10^{-2} ng/ μ L. The method for *Rhizoctonia solani* diagnosis using RPA-real time, which is rapid convenient and sensitive can provide a new rapid assay for the detection of *Rhizoctonia solani*.

Keywords: *Rhizoctonia solani*; ITS; Real-time RPA

Introduction

Rhizoctonia solani, a significant soil-borne pathogenic fungus belonging to the genus *Rhizoctonia*, is a harmful organism with a wide range of hosts [1]. It is known to infect 263 plant species across 43 families, including important crops such as legumes, grasses, and crucifers. In the biosafety risk assessment of introducing medical plant seedlings from abroad, *R. solani* is one of the primary harmful organisms targeted for detection. This fungus can exist in the form of mycelium or sclerotia in diseased plant residues and has a broad host range. Once it spreads to planting areas through transportation or other means, it can lead to crop yield reduction or even total crop failure, causing significant economic losses to agricultural production [2-7]. Currently, the identification of *R. solani* mainly relies on mycelial fusion method, rDNA-ITS method, and isoenzyme method for identification [8-11]. Reports on the harm caused by *R. solani* have been documented in some potato-growing regions in China, where pota-

to yields have been severely affected [12]. Therefore, rapid disease detection technology as a key approach to reduce agricultural losses caused by this fungus is crucial for early detection and prevention. However, existing technologies cannot meet the practical needs for rapid detection with high sensitivity. With the development of molecular biology technology, DNA molecular marker technologies such as RAPD, RFLP, real-time PCR, LAMP, RPA-LFD, etc. have been widely applied in species identification. Recombinase polymerase amplification (RPA) technology is an isothermal nucleic acid amplification technique that is favored in many fields due to its high sensitivity, specificity, and simple operation.

In recent years, it has also been widely used in the early identification of plant pathogens [13-19]. RPA amplification products can be detected by various methods such as gel electrophoresis, fluorescent labeling, and test strips, among which the RPA real-time fluorescent detection method can observe results during the amplification pro-

cess, providing the fastest detection speed. Currently, the only molecular technology for detecting *Rhizoctonia solani* at home and abroad is the TaqMan qPCR detection technology [20]. There are few studies on the rapid detection of *Rhizoctonia solani* using RPA, and only an RPA test strip detection method for *Rhizoctonia solani* causing tobacco target spot disease has been constructed [21]. This study established an RPA real-time fluorescent detection method for *Rhizoctonia solani*, which can be applied to the rapid identification of diseases transmitted by foreign medical plant seedlings [22]. It plays an important role in the quarantine process at entry ports and provides technical support for staffs to rapidly identify *Rhizoctonia solani*.

Materials and Methods

Sampling Materials

Shaanxi Normal University provided *Rhizoctonia solani*, *Fusarium oxysporum*, *Colletotrichum gloeosporioides* penz, *Botrytis cinerea*, China Agricultural University provided *Rhizoctonia solani*, *Puccinia polysora*, *Puccinia striiformis* (Table 1).

Table 1: Test strains used in this experiment were as follows.

Sample number	Scientific Name	Origin
1	<i>Rhizoctonia solani</i>	China Agricultural University Shaanxi Normal University
2	<i>Fusarium oxysporum</i>	Shaanxi Normal University
3	<i>Colletotrichum gloeosporioides</i> penz.	Shaanxi Normal University
4	<i>Botrytis cinerea</i>	Shaanxi Normal University
5	<i>Puccinia polysora</i>	China Agricultural University
6	<i>Puccinia striiformis</i>	China Agricultural University

Table 2: RPA primers and Probe Sequences for the ITS of *Rhizoctonia solani*.

Primers and probe (5'-3')	Sequences
RS-F	ATGTAACGCATCTAATACTAAGTTTCAACAACGG
RS-R	AGCATAACACTGAGATCCAGCTAATGAACGA
RS-P	CAATTCACATTACTTATCGCATTTCGCTGC-GTTCTTCATC

DNA Extraction

DNA was extracted from test strains using Rapid Plant Genomic DNA Isolation Kit (Sangon Biotech), the extracted DNA was dissolved in 50μL ultrapure water and then was stored at -20°C for further use.

Designing RPA Primers and Probes

The ITS region of *Rhizoctonia solani* (GenBank accession no.MF447834.1, LC384929.1, HE667746.1) were compared with those of other test strains (GenBank accession no. MT968495.1,

KC913203.1, KY031689.1, EU014048.1). The fluorescent RPA primers and probes were designed according to the variant gene site, The details of these primer pairs, along with the fluorescent probe, are summarized in Table 2.

RPA Reaction System

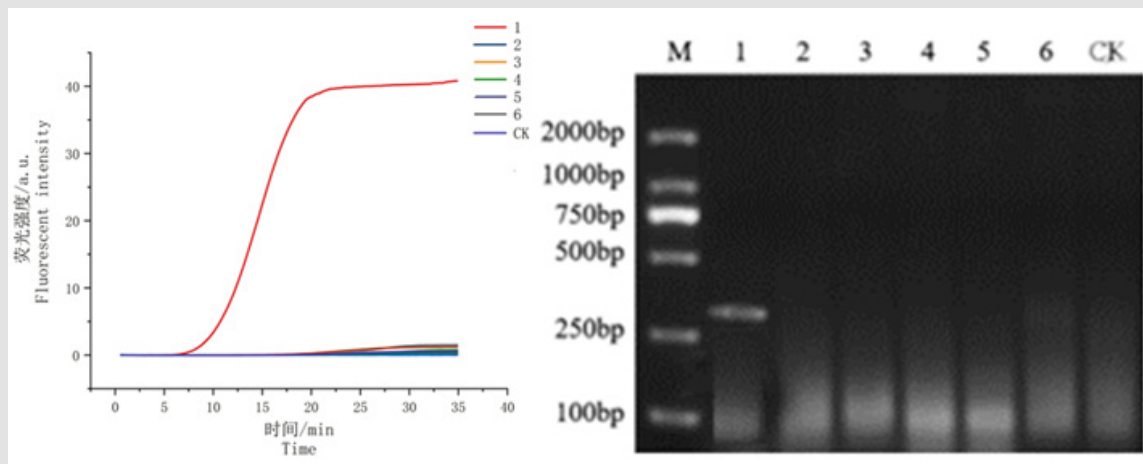
RPA reactions were performed using the TwistAmp exo Quick kit (TwistDx, Cambridge, UK) in 50 μL of reaction systems, utilizing 7500 Fast Real-Time PCR Systems (Applied Biosystems, the USA). Each reaction contained 29.5 μL of buffer, 2 μL of each forward and reverse primer (10 μM), 0.6 μL of probe (10 μM), 1 μL of template genomic DNA, 2.5 μL of magnesium acetate (280 mmol/L) and 12.4 μL of ddH₂O. The reaction was initiated by centrifugation of the pellet, followed by reaction of the tubes at 39°C for 30 minutes.

Specificity and Sensitivity of RPA Assays

DNA templates from *Rhizoctonia solani* and its closely related species were selected for analysis of the detection specificity, Distilled water was employed as the negative control. The specificity was tested by both PCR and fluorescent RPA. In the sensitivity determination experiments, Genomic DNA from *Rhizoctonia solani* and its closely related species were serially 10-fold dilution of the initial concentration of 60 ng/μL. The results were directly examined in the UV gel system for visualization and fluorescence amplification curves.

Results and Discussion

RPA specificity validation for detection of *Rhizoctonia solani* was conducted using DNA from *Rhizoctonia solani* and its closely related species as templates. As shown in Figure 1A, for the RPA real-time fluorescent specificity assay, *Rhizoctonia solani* (sample NO.1 in Table 1) successfully generated an ‘S’-shaped trend fluorescence curve, indicating amplification. while genomic DNA fluorescence amplification curves of its closely related species overlapped with CK (ddH₂O), showing no clear amplification curve (Figure 1A). For PCR detection assay (Figure 1B), the results showed that *Rhizoctonia solani* (sample NO.1 in Table 1) could be observed with a clear amplification band, while other closely related species and negative controls CK had no amplification band (Figure 1B). These observations indicate a high degree of specificity in the well-established fluorescent RPA detection of *Rhizoctonia solani*. To further verify the sensitivity of the RPA assay, Genomic DNA from *Rhizoctonia solani* and its closely related species were serially 10-fold dilution of the initial concentration of 60 ng/μL, seven template dilutions were designed, with water as a negative control, and the experimental results showed that the fluorescence signal intensity gradually decreased with decreasing template concentration (Figure 2A). RPA gel electrophoresis demonstrated clear amplification bands when the template concentration was greater than 60×10⁻¹ ng/μL (Figure 2B). The detection limit of RPA real-time fluorescent assay was 6×10⁻² ng/μL, and that of the conventional PCR was 6×10⁻¹ ng/μL.

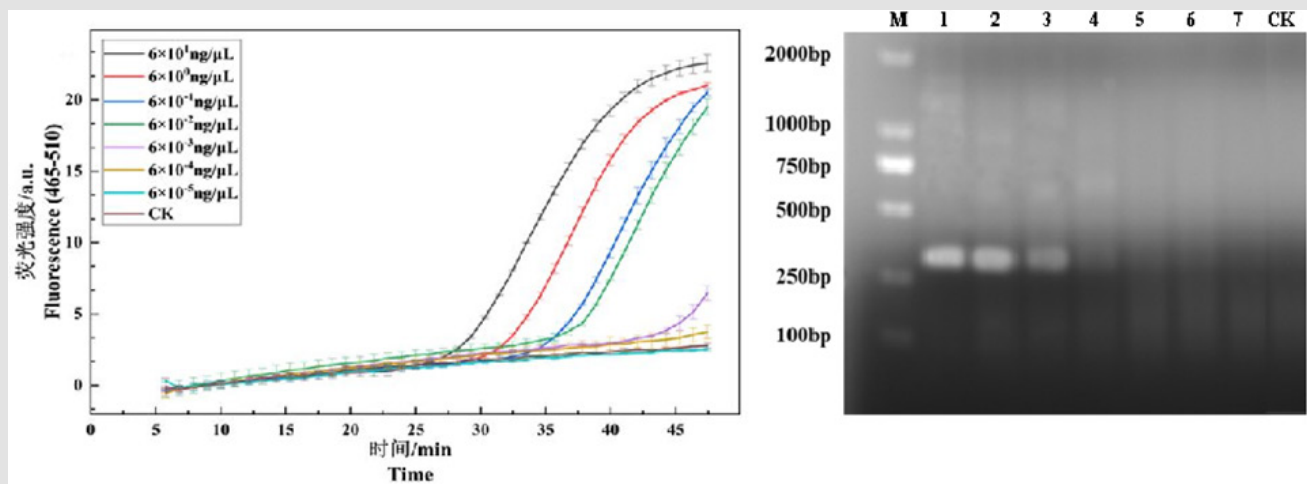


A. RPA real-time fluorescent specificity assay

B. PCR detection assay

Note: 1 to 6 are the DNA templates of corresponding samples in Table 1; CK: ddH₂O; Marker: DL2000;

Figure 1: RPA specificity validation for detection of *Rhizoctonia solani*.



A. RPA real-time fluorescent sensitivity assay

B. PCR detection assay

Note: 1 to 7 are *Rhizoctonia solani* DNA samples with pre-set concentrations of 60×10^1 ng/μL, 60×10^0 ng/μL, 60×10^{-1} ng/μL, 60×10^{-2} ng/μL, 60×10^{-3} ng/μL, 60×10^{-4} ng/μL, or 60×10^{-5} ng/μL; CK: ddH₂O; Marker: DL2000;

Figure 2: RPA sensitivity validation for detection of *Rhizoctonia solani*.

Rhizoctonia solani is an important plant pathogenic fungus that poses severe threats to various crops and plants, such as tomatoes, beans, potatoes, tobacco, strawberries, soybeans, and tulips. Traditional morphological identification methods and commonly used molecular detection methods such as PCR and qPCR for *Rhizoctonia solani* are cumbersome and time-consuming. In this study, we developed a rapid detection method for *Rhizoctonia solani* based on RPA real-time fluorescence detection technology. This method can accurately real-time monitor the progress and results of the amplification reaction

and achieve results in only 40 minutes. Therefore, the method established in this study is suitable for rapid detection of diseases carried by imported Chinese herbal medicine seedlings at entry ports and other links, thereby effectively preventing and controlling the spread and diffusion of diseases and ensuring the safety of agricultural crops and plant resources. In the future, this technology can be further optimized and promoted, expanding its application in other plant disease monitoring fields and providing more reliable and efficient technical support for agricultural production and plant protection.

Conclusion

In summary, the rapid detection method for *Rhizoctonia solani* based on real-time fluorescent RPA established in this study is characterized by high sensitivity, high specificity, and a rapid operation procedure, making it highly practical for on-site plant disease monitoring. This method has wide application potential and great development prospects in the field of agricultural production and plant protection.

Conflicts of Interest

The authors declare no competing interests.

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References

- Li KM, Guo QY, Zhao Li, et al. (2009) Study on anastomosis groups and their pathogenicity of *Rhizoctonia solani* isolated from alfalfa in Xinjiang. *Pratacultural Science* 26(05): 151-154.
- Liu Ya-ya, Chen Yuan, Guo Feng-xia, Shi You-tai, Li Zeng-xuan, et al. (2011) Isolation and identification of the pathogens causing *Rheum palmatum* root rot. *Acta Prataculturae Sinica* 20(01): 199-205.
- Li BJ, Chen QH, Lan CZ, Wang NN, Wang YC, et al. (2011) Identification and Pathogenicity Test of the Pathogens Causing Soybean Root Rot in Fujian. *Fujian Journal of Agricultural Sciences* 26(05): 798-803.
- Wang D, Xie XW, Chai AL, Shi YX, Li BJ (2019) Biological characteristics on *Rhizoctonia solani* isolated from Cruciferous-vegetables in North China. *Acta Phytopathologica Sinica* 49(05): 590-601.
- Tang CR, Chen J, Ji MS, et al. (2000) Study on the Pathogens of Corn Sheath Blight in Liaoning Province. *Acta Phytopathologica Sinica* (04): 319-326.
- Liang JY, Zhao C, Cai YB, Wu X (2024) Identification of the anastomosis groups of *Rhizoctonia* spp. causing diseases on watermelon seedling. *Acta Phytopathologica Sinica* 54(3): 658-663.
- Cheng R, Dong Z, Li W, Dai L, Liu S (2016) Research Progress of Soybean Root Rot. *Chinese Agricultural Science Bulletin* 32(08): 58-62.
- Chand T, Logan C (1983) Cultural and pathogenic variation in potato isolates of *Rhizoctonia solani* in Northern Ireland[J]. *Transactions of the British Mycological Society* 81(3): 585-589.
- Luo D, Tian H, Zhang CX (2020) Advances in the research on plant root rot caused by *Rhizoctonia solani*. *China Plant Protection* 40(03): 23-31.
- Carling DE, Kuninaga S, Brainard KA (2002) Hyphal Anastomosis Reactions, rDNA-Internal Transcribed Spacer Sequences, and Virulence Levels Among Subsets of *Rhizoctonia solani* Anastomosis Group-2 (AG-2) and AG-BI. *Phytopathology* 92(1): 43-50.
- Xiao Y, Liu MW, Li G (2008) Genetic Diversity and Pathogenicity Variation of Different *Rhizoctonia solani* Isolates in Rice from Sichuan Province China. *Chinese Journal of Rice Science* 15(2): 137-144.
- Gao YH, Meng LQ, Liu ZT, JIANG Chao, DIAO Zhuo, et al. (2025) Research progress on pathogenicity and integrated control of *Rhizoctonia solani* Kühn to potato. *Agricultural Engineering* 15(1): 85-92.
- Zhao ZX, Fan QX, Feng LX, et al. (2022) Establishment of a rapid detection method for maize dwarf mosaic virus based on recombinase polymerase amplification technology. *Plant Protection* 48(9): 83-89.
- Bier F F, Kersting S, Rausch V (2014) Multiplex isothermal solid-phase recombinase polymerase amplification for the specific and fast DNA-based detection of three bacterial pathogens. *Mikrochimica Acta: An International Journal for Physical and Chemical Methods of Analysis* 181: 1715-1723.
- Li C, Ju Y, Wu X, Le Cao, Benguo Zhou, et al. (2021) Development of recombinase polymerase amplification combined with lateral flow detection assay for rapid and visual detection of *Ralstonia solanacearum* in tobacco. *Plant Disease* 105(12): 3985-3989.
- Miles T D, Martin F N, Coffey M D (2015) Development of rapid isothermal amplification assays for detection of *Phytophthora* spp. in plant tissue. *Phytopathology* 105(2): 265-278.
- Wei MS, Tian Q, Zhao WJ, et al. (2016) Rapid detection of *Pseudomonas syringae* pv tomato by RPA method. *Plant Protection* 42(1): 150-153.
- Ma ZM, Duan Y, Xu JJ, et al. (2021) The Rapid Detection of *Xanthomonas citri* ssp. *Citri* (Xcc) Based on Recombinase Polymerase Amplification (RPA) Assay. *Acta Horticulturae Sinica* 48 (3): 590-599.
- Wu JC, Wang YZ, Zhang PP, et al. (2024) Development of RPA assay for detection of the corn rust pathogens *Puccinia polysora*. *Plant Quarantine* 38(2): 43-49.
- Luo SH, Luo YZ, Zhao YY, et al. (2022) Quantitative real-time PCR for rapid detection of *Rhizoctonia solani* and *Sclerotium rolfsii* in urban green space. *Journal of Zhejiang A&F University* 39(5): 1087-1095.
- Li YY, Qiu MJ, Li XH, et al. (2023) Establishment of LFD-RPA Rapid Detection Technique for Tobacco Target Spot. *Chinese Tobacco Science* 44(5): 62-69.
- Rush CM, Carling DE, Harveson RM, et al. (1994) Prevalence and Pathogenicity of Anastomosis Groups of *Rhizoctonia solani* from Wheat and Sugar Beet in Texas. *Plant Disease* 78(4): 349-352.

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