

## Antifungal Activity of *Streptomyces* sp. Against *Puccinia recondita* Causing Wheat Leaf Rust

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**Abstract** To discover a potent strain against wheat leaf rust, soil samples collected from Ilgamho, Seoul, Korea were tested *in vivo* and a strain belonging to *Streptomyces* sp. was found to show good antifungal activity when fermented in a broth. The identification of the strain was carried out based on 16S rDNA analysis, and the active compound was separated from the fermented broth. Even though its structure was not determined completely, the authors report the results obtained so far indicate that the fermented broth of the strain showed activity against wheat leaf rust. Therefore, we propose that this may be a potential novel strain showing antifungal activity against *Puccinia recondita*.

**Key words:** *In vivo* antifungal activities, *Puccinia recondita*, wheat leaf rust, *Streptomyces*

Wheat leaf rust is caused by the fungus *Puccinia triticina*. It reduces wheat yields. When the spore masses of the fungus break through the surface of the leaf, the color of the leaf becomes reddish orange like hence the disease is called leaf rust. Since the leaf rust spores are carried by wind, the disease can spread rapidly. Under favorable moisture and temperatures of 15°C to 25°C, leaf rust spores germinate and penetrate into the leaf. After 7 to 10 days the fungus can produce more spores. As a result, the disease can increase rapidly. Yield losses of wheat by leaf rust depend on its severity and the duration of infection, but severe losses occur from the seedling stage to maturity. The severity of the disease can be determined based on the amount of leaf area covered by leaf rust [2, 5, 8].

Fungicides to control wheat leaf rust include mancozeb, propiconazole, and azoxystrobin. Since there has been

a growing tendency in recent years for consumers to reject chemical synthetic pesticides, we have launched this study to discover naturally occurring biopesticides in microorganisms [3].

Soil samples were collected during 2002 at several sites of Ilgamho, Seoul, Korea. Actinomycetes were isolated by the method previously published by us [4]. Colonies showing the typical characteristics of actinomycetes were selected. They were transferred to Bennet agar medium, and the pH was adjusted to 7.2. They were then cultured for ten days at 28°C. Colonies selected from the agar were inoculated into Bennet's medium and cultured in a shaking incubator at 28°C for an additional ten days. Fermented broth was used to test antifungal activity against wheat leaf rust (*Puccinia triticina* recondita). Furthermore, 10 l of the cultured broth was prepared to perform the isolation and separation of active compounds.

The crop used for the activity assay was wheat (*Triticum aestivum* L, cv Chokwang). It was grown in vinyl pots placed in a greenhouse at 25 (±5)°C for 1–4 weeks. Potted crop seedlings were sprayed with broth dissolved in a mixture of water and methanol (95:5, v/v) containing Tween 20 (250 mg/l) as a wetting agent, and allowed to stand for 24 h [6]. Treated crop seedlings were inoculated to develop pathogenic fungi by spraying with fungal spore suspensions (10<sup>7</sup>–10<sup>9</sup> spores/l). After incubation of the crop seedlings in the dark, they were transferred to a growth chamber for one day at 25 (±2)°C and 100% RH, and were maintained at 25 (±2)°C and 70–80% RH with 12 h of light per day. The severity of the disease was determined based on a percentage of the infected leaf area. The experiments were arranged as a randomized complete block with three replicates per treatment. An assay of each pot was performed so as to visually estimate the percentage of leaf area covered by sporulating lesions, or the percentage of chlorotic presence with necrotic symptoms

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on the inoculated foliage. Data are the result of three trials. The mean value of nine estimates for each treatment was converted into a percentage figure of fungal control using the equation:

$$\% \text{ control} = 100 [(A - B)/A]$$

where A=area of infection (%) on leaves sprayed with Tween 20 solution alone, and B=area of infection (%) on treated leaves.

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5 AGAGTTTGATCCTGGCTCAGGACGAAACGCTGGCGCGTCTTAACACATGCAAGTCGAAC 60
6 AGAGTTTGATCCTGGCTCAGGACGAAACGCTGGCGCGTCTTAACACATGCAAGTCGAAC 60
8 GATGAAGCCCTTCGGGTGGATTAGTGGCGAACGGGTAGTAACACGTTGGCAATCTGCC 120
9 GATGAAGCCCTTCGGGTGGATTAGTGGCGAACGGGTAGTAACACGTTGGCAATCTGCC 120
11 CTTCACTCTGGGACAAAGCCCTGGAAACGGGGTCTAATACCGGATACGACCGCTCGGGCA 180
12 CTTCACTCTGGGACAAAGCCCTGGAAACGGGGTCTAATACCGGATACACCTGCCCGGGCA 180
14 TCCGATGTGCGTGGAAAGCTCCGGCGGTGAAGGATGAGCCGCGGCTATCAGCTTGTGG 240
15 -----GTGGGTAAAGCTCCGGCGGTGAAGGATGAGCCGCGGCTATCAGCTTGTGG 240
17 GTGAGTAAACGGCTCAACAGCGCAGCAGCGGTAGCCGCGCTGAGAGGGCAGCCGCGCAC 300
18 GTGAGTAAACGGCTCAACAGCGCAGCAGCGGTAGCCGCGCTGAGAGGGCAGCCGCGCAC 294
20 ACTGGACTGAGACAGCGCCAGACTCTACCGGAGGCGAGCAGTGGGGAATATTGCACAA 360
21 ACTGGACTGAGACAGCGCCAGACTCTACCGGAGGCGAGCAGTGGGGAATATTGCACAA 354
23 TGGCGAAAGCCTGATGACGCGACCGCCGCTGAGGGATGACGGCCCTCGGGTGTAAACC 420
24 TGGCGAAAGCCTGATGACGCGACCGCCGCTGAGGGATGACGGCCCTCGGGTGTAAACC 414
26 TCTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAAGAAGCGCGCTTAACATAC 480
27 TCTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAAGAAGCGCGCTTAACATAC 474
29 GTGCCAGCAGCCGCGTAATACCTAGGCGCGAGCGTTGTCCGGAATATTGGCGGTAAA 540
30 GTGCCAGCAGCCGCGTAATACCTAGGCGCGAGCGTTGTCCGGAATATTGGCGGTAAA 534
32 GAGCTCGTAGGCGCTGTGTCGCTCGGATGTAAAGCCCGGGGCTTAACCCCGGCTGTC 600
33 GAGCTCGTAGGCGCTGTGTCGCTCGGATGTAAAGCCCGGGGCTTAACCCCGGCTGTC 596
35 ATTCGATACCGGCTAGCTAGAGTGTGGTAGGGAGATCGGAATTCCTGCTAGCGGTGA 660
36 ATTCGATACCGGCTAGCTAGAGTGTGGTAGGGAGATCGGAATTCCTGCTAGCGGTGA 653
38 AATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGCGGATCTCTGGGCCATTACTGAC 720
39 AATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGCGGATCTCTGGGCCATTACTGAC 713
41 GCTGAGGAGCGAAACGCTGGGAGCGAACAGGATTAGATACCCCGTAGTCCACGCCGTA 780
42 GCTGAGGAGCGAAACGCTGGGAGCGAACAGGATTAGATACCCCGTAGTCCACGCCGTA 773
44 AACGGTGGGAACCTAGGTGTGGCGACATCCACGCTCGTGGTCCGCGAGCTAACGCATTA 840
45 AACGGTGGGAACCTAGGTGTGGCGACATCCACGCTCGTGGTCCGCGAGCTAACGCATTA 833
47 AGTTCGCCCTCGGGAGTACGGCCGCAAGGCTAAAACCTCAAGGAATTGACGGGGGCC 900
48 AGTTCGCCCTCGGGAGTACGGCCGCAAGGCTAAAACCTCAAGGAATTGACGGGGGCC 893
50 GCACAAGCAGCGGACATGTGGCTTAATTCGACGCAACCGGAAGAACCTTACCAAGGCTT 960
51 GCACAAGCAGCGGACATGTGGCTTAATTCGACGCAACCGGAAGAACCTTACCAAGGCTT 953
53 GACATACCGGAAACACCCAGAGATGGGTGCCCCCTTGTGGTTCGGTGTACAGGTGGTGC 1020
54 GACATACCGGAAACCGCCAGAGATGGGTGCCCCCTTGTGGTTCGGTGTACAGGTGGTGC 1013
56 ATGGCTGTGCTCAGCTCGTGTGCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACC 1080
57 ATGGCTGTGCTCAGCTCGTGTGCTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACC 1073
59 TTGTTCTGTGTTGCCAGCATGCCCTTCGGGCTGATGGGACTCACAGGAGACTGCCGGG 1140
60 TTGTTCTGTGTTGCCAGCATGCCCTTCGGGCTGATGGGACTCACAGGAGACTGCCGGG 1133
62 TCAACTCGGAGGAAGTGGGGACGACGTCAGTCAATCATGCCCTTATGTCTTGGGCTGC 1200
63 TCAACTCGGAGGAAGTGGGGACGACGTCAGTCAATCATGCCCTTATGTCTTGGGCTGC 1193
65 ACACGTGTACAATGGCAGGTACAATGAGTGCGAAGCCGTGAGGCGGAGCGAATCTCAA 1260
66 ACACGTGTACAATGGCAGGTACAATGAGTGCGAAGCCGTGAGGCGGAGCGAATCTCAA 1253
68 AAAGCCTGTCTCAGTTCGGATTGGGGTCTGCAACTGACCCCATGAAGTCGGAGTTGCTA 1320
69 AAAGCCTGTCTCAGTTCGGATTGGGGTCTGCAACTGACCCCATGAAGTCGGAGTTGCTA 1313
71 GTAATCGCAGATCAGCAGTGTGCTCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCC 1380
72 GTAATCGCAGATCAGCAGTGTGCTCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCC 1373
74 TCACGTACCGAAAGTCGGTAAACCCGAAGCCGGTGGCCCAACCCCTTGTGGGAGGGAGC 1440
75 TCACGTACCGAAAGTCGGTAAACCCGAAGCCGGTGGCCCAACCCCTTGTGGGAGGGAGC 1433
77 TGTGCAAGGTGGGACTGGCGATTGGGACGAAGTCGTAACAGGTAGCCGTA 1491
78 TGTGCAAGGTGGGACTGGCGATTGGGACGAAGTCGTAACAGGTAGCCGTA 1484
    
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Fig. 1. The result obtained by BLASTN. K: KACC91021; S: *Streptomyces neyagawaensis*.

In order to identify the strain, a 16S rDNA sequence analysis was carried out. To extract DNA, the strain was fermented in Bennet's medium for 7 days and the fermented broth was centrifuged. The extraction of DNA was carried out using a Genomic DNA Extraction Kit (Intronbiotechnology, Kyungkido, Korea). Amplification of the DNA was performed using a PCR kit using primers (fD1 5'-AGA GTT TGA TCC TGG CTC AG-3', rP2 5'-ACG GCT ACC TTG TTA CGA CTT-3'). The conditions used for the thermal cycles were as follows: denaturation at 94°C for 5 min prior to cycling; 35 cycles of denaturation at 94°C for 1 min, annealing at 58°C for 1 min, extension at 72°C for 2 min, further extension at 72°C for 10 min, and cooled to 4°C. The PCR products were cloned using a pGEM-T cloning kit (Promega, Madison, WI, U.S.A.) according to the manufacture's instructions. The sequences of the PCR products were determined by four internal primers which were : p510r 5'-TAT TAC CGC GGC TGC TG-3', p364f 5'-GGC AGC AGT GGG GAA TAT TG-3', p783f 5'-TAG ATA CCC TGG TAG TCC AC-3', and p1037f 5'-TCG TCA GCT CGT GTC GTG AG-3'. The determined sequences were analyzed by a BLAST program at the National Center for Biotechnology Information (Fig. 1). The analysis shows that the strain has a 98% homology (1484/1491) with *Streptomyces neyagawaensis*. The strain was deposited in the Korean Agricultural Culture Collection (KACC) at Suwon, Korea. Its registration number is KACC91021. The evolutionary distance calculated by the Jukes and Cantor method [1], and a phylogenic tree created using the neighbor-joining method are shown in Fig. 2.

Morphological observations of strain KACC91021 were made using scanning electron microscopy (SEM, JSM 5410LV, JEOL, Peabody, U.S.A.). Strain KACC91021 for the SEM observation was cultured in a small Petri dish containing Bennet's medium supplemented with agar at 28°C for seven days. Samples were fixed in an osmotically balanced solution of 2% glutaraldehyde in 0.05 M sodium cacodylate buffer (pH 7.2) at 25°C for 3 h and were then washed three times at 4°C for ten min in 0.05 M sodium

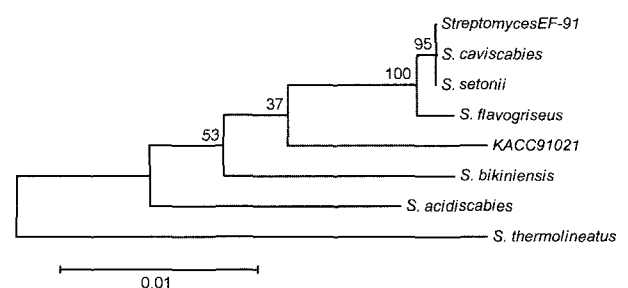


Fig. 2. The evolutionary distance calculated by the Jukes and Cantor method and a phylogenic tree created using the neighbor-joining method.



**Fig. 3.** Morphological observation of strain KACC91021 obtained using scanning electron microscopy after growth on Bennet's medium at 28°C for 7 days.

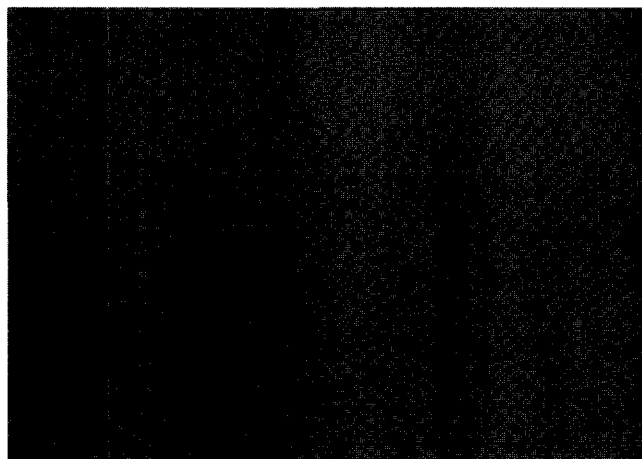
cacodylate buffer (pH 7.2). Samples were post-fixed in 1% aqueous osmium tetroxide in 0.05 M sodium cacodylate buffer (pH 7.2) for 2 h at 4°C, and were then washed twice with distilled water. Samples were dehydrated in a graded ethanol series (30, 50, 70, 80, 90, 100, 100, and 100), and were dried in liquid CO<sub>2</sub> using the Balzers CPD 010 (Balzers Instruments, Liechtenstein) mounted on aluminum stubs, and sputter-coated with gold/palladium using the Polaron SEM Coating Unit E5100 (Thermo VG Scientific, Beverly, MA, U.S.A.). The image obtained by SEM is shown in Fig. 3.

Figure 4 shows antifungal effects of the broth produced by strain KACC91021 against *Puccinia triticina*. Ten ml of the broth showed 80% of control. The broth was concentrated using a rotary evaporator under reduced pressure. Solvents such as ethylacetate, n-hexane, and butanol were used for extraction employing a separation funnel. The fraction separated from the aqueous layer showed the best activity. Column chromatography with



**Fig. 4.** Effects of strain KACC91021 against *Puccinia triticina* recondita.

Left: wheat leaves infected by *Puccinia triticina* recondite; center: wheat leaves with the treatment of Flusilazole; right: wheat leaves with the treatment of the broth of the strain.



**Fig. 5.** The chromatogram of the fraction 21-4-1 obtained from HPLC with photodiode array detector.

Silica Gel was performed on a solvent system of methanol: chloroform (3:7, v/v). Seven fractions were collected using an auto-collector. They were named from 21-1 to 21-7. Among these, the fraction 21-4 showed the best activity. Fractions were separated again under a different solvent condition (methylenechloride:ethanol=10:1). They were named 21-4-1 to 21-4-6. Of the six fractions, 21-4-1 showed the best activity, and fortunately it did not need further separation. As shown in Fig. 5, the chromatogram obtained by HPLC (Waters, Milford, MA, U.S.A.) with a two-dimensional photodiode array detector gives a single spot. The HPLC analysis was performed on a Waters HPLC system with a flow rate of 1 ml/min, a Vydak C8 column of 4.5 mm×300 mm, and eluent of methylenechloride: ethanol=10:1 [9]. The collected sample was freeze-dried. To identify the structure of 21-4-1, several NMR experiments were carried out on a Bruker DRX400 (Karlsruhe, Germany) [7].

The structural determination of the active compound requires further work. Even though the compound has not been identified, it is meaningful that the fermented broth of strain KACC91021 shows an activity against wheat leaf rust. Therefore, we propose that this may be a potential novel strain showing antifungal activity against *Puccinia recondita*.

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