

ISOLATION AND STRUCTURE OF SCH 351633: A NOVEL HEPATITIS C VIRUS (HCV) NS3 PROTEASE INHIBITOR FROM THE FUNGUS *PENICILLIUM GRISEOFULVUM*

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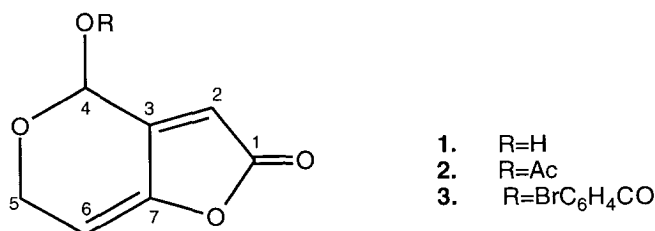
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Abstract: A new hepatitis C virus (HCV) protease inhibitor designated as Sch 351633 (**1**) was isolated from the fungus, *Penicillium griseofulvum*. Structure elucidation of **1** was accomplished by analysis of spectroscopic data, which determined compound **1** to be a bicyclic hemiketal lactone. Compound **1** exhibited inhibitory activity in the HCV protease assay with an IC_{50} value of 3.8 μ g/mL. © 1999 Elsevier Science Ltd. All rights reserved.

Hepatitis C virus (HCV) is responsible for viral infections such as non-A, non-B (NANB) hepatitis and cirrhosis, which is strongly associated with development of hepatocellular carcinoma.^{1–3} Since a significant HCV infected population in the United States and around the world has been reported, there is a great interest in developing more effective therapeutic agents against HCV to complement currently available interferon therapy.⁴ In the course of searching for novel HCV protease inhibitors,⁵ a fungal metabolite, Sch 351633 (**1**) was isolated from the fungus, *Penicillium griseofulvum* (culture Mer-0442, designated as Schering Culture Collection Fungi, SCF-1704). The microorganism was isolated from a soil sample collected from a desert terrain in the state of Arizona, U.S.⁶ Herewith we report the isolation and structure determination of **1**.

Figure 1. Structures of Sch 351633 (**1**) and Its Derivatives **2** and **3**



An ethylacetate extract (3.3 g) from a 2 L fermentation broth of *P. griseofulvum* was dissolved in 150 mL of 40% aqueous MeOH, and partitioned with 100 mL of CH₂Cl₂. The HCV active aqueous MeOH portion (2.6 g) was subjected to reverse-phase HPLC purification (YMC ODS, S-5, 20 x 250 mm semi-preparative column

with 20 x 50 mm guard column, 2-20% acetonitrile in H₂O with a linear gradient in 25 min, 12 mL/min, UV = 220 nm). A total of six replicate runs (50 mg per injection) yielded 100 mg of pure **1** which was obtained as jell-like material after lyophilization.

Table 1. ¹H and ¹³C NMR Data of **1** and **2**^a

#	1 (Acetone- <i>d</i> ₆)		2 (DMSO- <i>d</i> ₆)	
	¹³ C	¹ H	¹³ C	¹ H
1	169.4 s ^b	--	168.1 s	--
2	110.5 d	5.91 s	111.8 d	6.36 br.s
3	146.6 s	--	146.7 s	--
4	88.8 d	5.96 br.s	86.3 d	7.00 s
5	59.5 t	4.30 (dd 17.6, 3.5) ^c	60.1 d	4.51 (dd 17.0, 4.5)
		4.55 (br.d, 17.6)		4.62 (br.d 17.0)
6	108.8 d	5.96 br.s	108.5 d	6.29 (br.t 4.5)
7	151.9 s	--	144.5 s	--
COCH ₃	--	--	168.8 s	--
COCH ₃	--	--	20.6 q	2.10 s

a. Recorded on 400 MHz instrument.

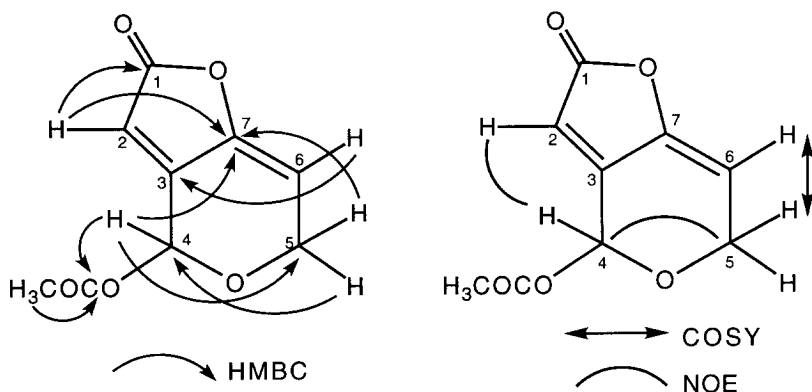
b. Multiplicity was determined by APT experiment.

c. Coupling constants in Hz.

The molecular weight of **1** was determined to be 154 by EI and CI-MS data that showed the molecular ion M⁺ = 154 and protonated molecular ion (M+H)⁺ at *m/z* 155, respectively. Molecular formula of C₇H₆O₄ for **1** was deduced based on elemental analysis (Calcd: C 54.55%, H 3.90%. Measured: C 54.32%, H 3.73%) as well as carbon NMR data. IR spectral data showed maximum absorption of 1741 cm⁻¹ suggesting the presence of a conjugated lactone or ester functional group. UV maximal absorption of 285 nm indicated an extended conjugation system of **1**. ¹H NMR spectrum of **1** (Table 1) showed an AB quartet spin system at δ 4.30 and 4.55, in which each doublet had very small coupling with an adjacent proton reflecting an oxygenated allylic CH₂ group. A singlet at δ 5.91 and a broad singlet at δ 5.96 suggested two olefinic protons. It should be noted that the integration of broad singlet at δ 5.96 indicated two protons associated with this signal. ¹³C NMR spectral data were consistent with ¹H NMR data showing one methylene carbon at δ 59.5, two olefinic methines at δ 108.8 and 110.5 and one dioxygenated methine at δ 88.8. In addition, one carbonyl at δ 169.4 and two olefinic quarternary carbons at δ 146.6 and 151.9 were observed the ¹³C NMR experiments of **1**. Interestingly, the tautomeric (aldehyde-alcohol, ring cleavage) form of **1** was not observed in NMR spectra by using different solvents (MeOH-*d*₆ and DMSO-*d*₆). The hemiacetal form presents predominantly perhaps due to the more favorable stabilization of forming a six ring in the molecule.

In order to obtain additional information and better quality of spectrum for assignments, compound **1** was acetylated with acetic anhydride-pyridine. Extensive 2-D NMR experiments and all detailed assignments, therefore, were performed on acetylation product **2**. Deuterated DMSO solvent was chosen instead of acetone for NMR experiments to avoid an overlap of the acetyl-CH₃ signal at ~2 ppm. COSY experiment clearly indicated direct connection of 5-CH₂ to 6-H. This connection was confirmed by observation of proton coupling and difference NOE experiments. Difference NOE experiments further revealed correlation of 2-H to 4-H, and 4-H to 5-CH₂. The carbon and proton connections were finally established based on HMBC experiments. The structure of **1** was proposed by analysis of long range carbon-proton coupling data from HMBC experiments, which are summarized and as shown in Figure 2. It should be noted that the stereochemistry of chiral center at 4-position was not determined by NMR spectroscopic data. A *p*-bromobenzoate derivative product **3** was made by esterification with *p*-bromobenzyl chloride/triethylamine.⁷ However, several attempts of crystallization of **3** failed to provide suitable crystals for X-ray analysis.

Figure 2. 2-D NMR Data of Acetylated Derivative of **1**



Compound **1** demonstrated antiviral activity in the *in vitro* HCV protease scintillation proximity assay with the IC₅₀ = 3.8 µg/mL.⁸ *In vitro* antiviral activity for the acetylated product **2** and *p*-bromobenzoate derivative **3** were also evaluated. The IC₅₀ values of **2** and **3** were shown to be 7.2 and 12.6 µg/mL, respectively.

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6. The fungus was supplied by Mr. G. M. Luedemann.
7. To a mixture of Sch 351633 (100 mg, 0.65 mmol) and pyridine (0.5 mL) in dry CH_2Cl_2 (10 mL) and acetonitrile (1 mL) solution was added *p*-bromobenzoyl chloride (150 mg, 0.68 mmol), and then stirred at room temperature overnight. The reaction was quenched by adding water (30 mL) followed by additional 15 mL of CH_2Cl_2 . The CH_2Cl_2 layer was separated, dried over NaSO_4 and evaporated under nitrogen. The residue was redissolved in acetonitrile and yielded 65 mg of crude acetonitrile soluble material. The crude material was purified on semi-preparative HPLC (YMC ODS 20 x 250 mm column with guard column, S-5, 20–100% ACN in H_2O with linear gradient in 30 min, 12 mL/min, UV = 220 nm) to afford 11 mg of pure **3**. CI-MS m/z 339 ($\text{M}+2$)⁺; 337 (M)⁺; ¹H NMR (400 MHz, CDCl_3) δ 4.53 (dd, J = 4.7, 17.5 Hz, 1H), 4.75 (br.d, J = 17.5 Hz, 1H), 5.97 (br.s, 1H), 6.08 (s, 1H), 7.16 (s, 1H), 7.56 (d, J = 8.7 Hz, 2H) 7.34 (d, J = 8.7 Hz, 2H); ¹³C NMR (100 MHz, CDCl_3) δ 168.1, 164.1, 146.3, 145.8, 132.1, 132.1, 131.5, 131.5, 129.4, 127.7, 112.7, 106.7, 88.1, 60.9.
8. Scintillation Proximity Assay (SPA) for HCV NS3 Protease.
Inhibition of proteolytic activity of HCV NS3 protease was determined by scintillation proximity assay (SPA). NS3 631/4A protease, purified to 95% homogeneity from a baculovirus expression system, was incubated at 10 nM concentration with the peptide substrate containing the HCV NS5A/5B cleavage site. This peptide substrate was biotinylated at the N-terminus and ³H-labeled at the C-terminus. After incubation at 22 °C for 2 h, the reaction was stopped, and cleavage of the Cys-Ser bond was detected by the addition of streptavidin-coated SPA beads (Amersham), which can bind to the biotin-labeled peptide. Signal is detected as cpm using a Packard Topcount Model NXT instrument. Cleavage is indicated by reduction in signal caused by separation of the radiolabeled fragment from the bound biotinylated fragment, which is proportional to proteolytic activity.