A Novel Compound from the Mushroom Cryptoporus volvatus Inhibits Porcine Reproductive and Respiratory Syndrome Virus (PRRSV) In Vitro

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Abstract

Porcine reproductive and respiratory syndrome (PRRS), caused by PRRS virus (PRRSV), is a serious contagious disease in the swine industry. At present, there are no effective control strategies against PRRSV. Thus, there is an urgent need for new treatment regimens that have efficacious antiviral activity to compensate for vaccines. The anti-infective effect of Cryptoporus volvatus has previously been demonstrated in Tradational Chinese Medicine. In this report, we expected to identify a new anti-PRRSV agent in the aqueous extract of C. volvatus, by employing a combination of modern chromatographic purification techniques and indirect immunofluorescence assay (IFA). Our results showed that C. volvatus extracts from every separation step differed in their inhibitory potency on PRRSV. One anti-PRRSV component designated as $C_{M-H-L-5}$ was isolated from water-soluble fraction of C. volvatus. The inhibition induced by $C_{M-H-L-5}$ occurred in a dosedependent manner. C_{M-H-L-5} appeared to be a low-molecular-weight polyol fragment with amide groups and carboxylic acid groups. Collectively, our findings imply that C_{M-H-L-5} from the aqueous extract of C. volvatus has the potential to be used for anti-PRRSV therapy.

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Introduction

Porcine reproductive and respiratory syndrome (PRRS) is a serious contagious disease in the swine industry, causing significant economical losses worldwide [1,2]. The causative agent, PRRS virus (PRRSV), can cause reproductive failure in pregnant sows, respiratory diseases in piglets, and asymptomatic infections in boars [1]. Most recently, there were devastating outbreaks of atypical PRRS in China, which is characterized by high fever, high morbidity, and high mortality in pigs of all ages [3,4]. The causative agent is a highly pathogenic PRRSV (HP-PRRSV) genotype with a discontinuous deletion of 30 amino acids in nonstructural protein 2 (nsp2) [3,4,5,6]. PRRSV belongs to the family Arteriviridae of the order Nidovirales, and its genome is a single-stranded, positive-sense RNA [7,8].

At present, vaccination is the prevailing way to control PRRS virus infection. However, commercial vaccines against PRRS virus have serious problems related to efficacy and safety. Antiviral therapeutics constitute a critical tool for combating viral infections, especially for cases in which there are no vaccines to match well with the circulating virus. Thus, an alternative measure to control PRRSV is pharmacological intervention. Previous studies have discovered a few natural compounds and compositions that have antiviral activities on PRRSV [9]. However, until now there are no effective commercial drugs available to control PRRSV infection. Currently used antiviral agents are often costly, have significant side effects [10], and lead to development of drug resistance in virus populations evolving under selective pressures [11,12]. As a result, antiviral natural products are candidates to be developed as new generations of antivirals administered either alone or, preferably, in combination with current modalities [13].

The medical use of mushrooms has a long tradition in Asian countries, and their use in the Western hemisphere has increased slightly in the past decades [14,15,16,17]. Whole extracts [18] and also isolated compounds [19,20] of medicinal mushrooms have been shown to have antiviral effects. With advances in fractionation techniques for isolating and purifying natural products and in analytical techniques for structural determination, screening of natural product mixtures is now more compatible with the expected timescale of high-throughput screening campaigns. C. volvatus belongs to Aphyllophorales [21], and grows in certain areas of China. Its fruiting body had been used for the treatment of asthma and bronchitis since the 15th century AD. [22]. Aqueous extract from the fruiting body of C. volvatus has been reported to have polysaccharose, proteins, volatile oil, and cryptoporic acids, etc and anti-tumor, anti-allergy, anti-inflammatory, and immunomodulatory activities [23,24,25].

Our research team had reported that the aqueous extract of the fruiting bodies of C. volvatus had the potential to be used for antiviral therapy [26]. However, further information about the Table 1. List of primers for real-time PCR.

^aF1: forward primer, R1: reverse primer.

^bSwine gene sequences were downloaded from GenBank.

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antiviral principles in C. volvatus is unavailable. In order to identify a new antiviral agent, we set out to isolate and purify the active compounds from the aqueous extract of C. volvatus fruiting bodies.

Materials and Methods

Material

C. volvatus was purchased from a market in the Yunnan Province of China. The mushroom was authenticated and a voucher specimen was deposited in our laboratory.

Extraction and Isolation

Air-dried fruiting bodies of C. volvatus (200 g) were pulverized in a grinder and extracted overnight with distilled water (1500 ml) at 4° C. After leaving in a water bath at 65 $^{\circ}$ C for 1 h, the slurry was centrifuged at 9,000 g for 30 min. The residue was further extracted by incubation in a water bath in

Figure 1. Elution profile of crude extract C_M from HP-2MGL macroporous resino column. Fraction collected through elution with 2 l 30% ethanol, corresponding to the strand on the x axis between 0 and 2 l; fraction collected through elution with 2 l of 60% ethanol, corresponding to the strand on the x axis between 2 and 4 l; and fraction collected through sample elution with 2 l of 100% ethanol, corresponding to the strand on the x axis between 4 and 6 l. The ultraviolet absorption of C_{M-H} contained three inner peaks, C_{M-H-1} , C_{M-H-2} and C_{M-H-3} .

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500 ml distilled water for 1 h. The extraction process was repeated twice, and the combined supernatant, named C, was filtered through a Millipore ultrafiltration membrane (NMRWL: 1000, Millipore). The ultrafiltrate was evaporated to dryness to give the resulting crude extract C_M . The crude extract C_M (1 g) was dissolved in 2 ml distilled water, and then further purified sequentially by macroporous resin column chromatography, anion exchange chromatography on DEAE-cellulose, and Sephadex LH-20 chromatography.

Purification by Employing Macroporous Resino Column

The crude extract was applied to a nonpolar macroporous resin column (HP-20, 5.5×20 cm, Mitsubishi) and a polar macroporous resin column (HP-2MGL, 5.5×20 cm, Mitsubishi) successively. After unabsorbed material had been eluted with distilled water, the polar macroporous resino column was eluted with a gradient of 0 to 100% ethanol solution. Absorbance at 260 nm was measured. Active fractions were collected for lyophilization.

Purification by Employing Anion Exchange Chromatography on DEAE-cellulose

The resulting active fractions of the aqueous extract of C . *volvatus* were further dissolved in 10 mM $NH₄HCO₃$ buffer (pH 9.4) before centrifugation to remove insoluble material. The solution was chromatographed on a $2.5 \text{ cm} \times 35 \text{ cm}$ column of DEAEcellulose (Sigma) which had been equilibrated with and was then eluted with 10 mM $NH₄HCO₃$ buffer (pH 9.4). The active component was collected and evaporated to dryness.

Purification by Employing Sephadex LH-20 Chromatography

The active fractions were further fractionated by size-exclusion chromatography on Sephadex LH-20 (Pharmacia) [27]. Sephadex LH-20 was swollen in 20% ethanol overnight and then packed in a C10/20 column. Active fractions were chromatographed on the Sephadex LH-20 column and eluted successively with distilled water, 30% ethanol, 60% ethanol, 50% acetone at a flow rate of 0.2 ml/min. The eluate was monitored at UV 260 nm and 4-ml fractions were collected. Active fractions were collected and then dried by using a freeze-dryer (FD5-4, SIM). PRRSV inhibitory

Figure 2. Elution profile of crude extract C_{M-H-L} from Sephadex LH-20. Fraction collected through elution with 100 ml distilled water, 100 ml of 30% ethanol, 100 ml of 60% ethanol, 100 ml of 100% ethanol, corresponding to the strand on the x axis between 0 and 100 ml, 100 and 200 ml, 200 and 300 ml, 300 and 400 ml respectively. Sephadex LH-20 yielded five fractions at OD 260 nm, $C_{M-H-L-1}$, $C_{M-H-L-2}$, $C_{M-H-L-3}$, $C_{M-H-L-4}$ and $C_{M-H-L-5}$.

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Figure 3. C. volvatus fraction C_{M-H-L-5} inhibited PRRSV replication. (A and B) The C_{M-H-L-5} blocked PRRSV Ch1a replication in Marc-145 cells. Marc-145 cells were infected with PRRSV Ch1a at an MOI of 0.1, and then treated with IFN-a (10 units/µl) or the C_{M-H-L-5} at various concentrations. At 24 h p.i., cells were fixed and analyzed by IFA using antibody against PRRSV N protein (A), and virus yield in the supernatants was also quantified (B). Culture treated with normal saline was set up as negative control (0 mg/ml), and treated with INF-a was set up as positive control. The brightness of the fluorescence represents the level of the virus. Results are from three independent experiments, each of which was conducted in triplicate. (C) The $C_{M-H-1.5}$ potently inhibited both PRRSV HV and VR2332 replication in PAMs. A similar virus inhibition assay was performed with PAM cells infected with PRRSV strain HV or VR2332 at an MOI of 0.1 in the presence of IFN-a (10 units/µl) or the C_{M-H-L-5} at various concentrations. (D) Determination of cytotoxicity of the C_{M-H-L-5} by MTT assay. PAMs or Marc145 cells were incubated with various concentrations of the C_{M-H-L-5} or the control normal saline for 48 h prior to the MTT assay. Data are representative of three independent experiments (mean \pm SD). Statistical significance was analyzed by Student's t test. *P<0.05; **P<0.01; ***P<0.001. doi:10.1371/journal.pone.0079333.g003

activity was checked by fluorogenic assays as previously described [26]. The fractions with the highest activity were further analysed by reverse-phase HPLC.

Reverse Phase HPLC Analysis and Purification

Chromatographically pure methanol and ultrapure water were filtered with a 0.45 - μ m filter membrane before degassing. The active fraction (1 mg or 10 mg) from Sephadex LH-20 chromatography was dissolved in 1 ml of absolute methanol. Before analysis by HPLC coupled to PDA, the sample was filtered through a 0.45-um nylon filter. Separation was achieved on an Agilent reverse phase C18 column (analytical column or preparation column) at 25° C. Elution was performed with $H_2O-MeOH$ (95:5, vol:vol) using a flow rate of 1 ml/min or 3 ml/min. Detection was carried out in PDA, using 260 nm as wavelength. The combined active fractions were evaporated to dryness to give the extracts.

NMR, MS Spectra and Detection of Major Chemical Functional Groups

ESI-FT-ICR-MS spectra were obtained by using an Acuu TOF CS, and ¹H- and ¹³C-NMR spectra were acquired with a BRUKER AVANCEIII-400 spectrometer.

The chemical color reactions were carried out on TLC silica gel $(2.5 \text{ cm} \times 8 \text{ cm}, \text{EMD}$ Millipore) with Dragendorff, Ninhydrin and Bromophenol Blue (BPB) reagents.

Cells, Viruses, and Virus Preparation

Marc-145 cells are a PRRSV-permissive cell line sub-cloned from MA-104 cells [28]. Marc-145 cells were maintained in Dulbecco's minimum essential medium (DMEM) supplemented with 10% FBS and penicillin/streptomycin. Porcine alveolar macrophages (PAMs) were obtained by postmortem lung lavage of 8-week-old specific pathogen free (SPF) pigs, and maintained in RPMI 1640 supplemented with 10% FBS and penicillin/ streptomycin.

PRRSV strains, CH-1a (the first type 2 PRRSV strain isolated in China), VR2332 (the prototype of Type 2 PRRSV strain), and

Figure 4. Transcript levels of cytokines in PAMs. PAMs were infected with HV at an MOI of 0.1 in the presence or absence of C_{M-H-L-5}. Expressions of IFN- α (A), IFN- β (B), IL1- β (C)and TNF- α (D) were analyzed using Real-time RT-PCR at 12 h after treated with HV, HV plus C_{M-H-L-5} (0.8 mg/ml), or C_{M-H-L-5} only. Relative expression (fold changes) in comparison with PBS (Phosphate buffer saline) -treated cells (set up as 1) is shown. The data represent the mean \pm standard deviation from three independent experiments. Significant differences between groups treated with $C_{M+1-L-5}$ and for PBS PRRSV infected cells or uninfected cells are denoted by "*" (P<0.05), "**" (P<0.01), and "***" (P<0.001). doi:10.1371/journal.pone.0079333.g004

HV (a highly pathogenic PRRSV (HP-PRRSV) isolate, GenBank accession no. JX317648), were propagated in Marc-145 cells or PAMs. Virus preparations were titrated on Marc-145 cells or PAMs, and then stored at -80°C. Briefly, PRRSV was serially diluted 10-fold in complete DMEM or RPMI1640 to infect 5×10^4 Marc-145 cells or PAMs in 96-well plates. PRRSV infection was determined 72 h post infection using immunofluorescent staining for the PRRSV N protein. Virus titer was determined using Reed-Muench method, and expressed as tissue culture infective dose 50% (TCID₅₀). PFU was determined according to "PFU = $0.7*$ T- CID_{50} " as described before [29], and the multiplicity of infection (MOI) was calculated based on PFU.

Assay of Inhibition of Virus Infection

Marc-145 cells or PAMs in 96-well plates were inoculated with Ch-1a, HV or VR2332 at an MOI of 0.1 for 2 h at 37° C, and then the viral inoculum was removed and fresh medium containing 2% FBS and different concentrations of the Cryptoporus volvatus extract or IFN-a (10 units/ml), a known inhibitor of PRRSV replication [30], was added. Twenty-four hours later, the supernatant was collected for virus titration, and cells were fixed for indirect immunofluorescence assay. The 50% effective concentration (EC_{50}) was determined using a 4 parameter, nonlinear regression of dose response inhibition by plotting log (inhibitor(-concentration)) vs. virus titer (variable slope) using GraphPad Prism (GraphPad Software, San Diego, CA).

Indirect Immunofluorescence Assay (IFA)

Cells were fixed with cold methanol-acetone (1:1, vol:vol) for 10 min at 4° C, washed with phosphate-buffered saline (PBS), and then blocked with 5% normal goat serum for 30 min at 37° C. After blocking, cells were stained with anti-PRRSV N protein monoclonal antibody SDOW17 (1:10,000, Rural Technologies) for 60 min at room temperature. Cells were then washed and incubated with FITC-conjugated goat anti-mouse IgG (H+L) (1:2000, Jackson ImmunoResearch) for 1 h at 37° C. After three washes in PBS, cells were counter-stained with DAPI and examined by fluorescence microscopy.

MTT Assay

The MTT [3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazo-lium bromide] assay was used to examine the effect of the Cryptoporus volvatus extract on cell viability. Marc-145 cells or PAMs in 96-well plates were treated with sequential dilutions of the extract or normal saline in a total of 100 ml growth medium for 48 h. And then, 20 ml of freshly made 5 mg/ml MTT solution was added to each well, and the cells were incubated at 37° C for another 5 h before the medium was replaced with 200 ml DMSO

Figure 5. Different C. volvatus fractions blocked PRRSV replication in Marc-145 cells and PAMs. (A, B and C) Marc-145 cells were infected with PRRSV Ch1a at an MOI of 0.1, and then treated with different C. volvatus extracts at a concentration of 0.8 mg/ml or INF- α (10 units/µl). At 24 h post-infection, cells were fixed and analysed by IFA using antibody against PRRSV N protein (A). The abbreviations at the bottom of fluorescent pictures are explained as follows: C: Aqueous extract of C. volvatus; C_M: C was filtered through a Millipore ultrafiltration membrane and the ultrafiltrate gave the resulting crude extract C_M; C_{M-H}: C_M absorbed on HP-2MGL was eluted with 30% ethanol, which resulted in C_{M-H}; C_{M-H-L-5}: C_{M-H} was loaded on a DEAE and Sephadex LH-20 chromatography column sequentially, which resulted in C_{M-H-L-5}. Virus yield in the supernatant was quantified (B). The same experiment was carried out with different C. volvatus extracts at a concentration of 1.2 mg/ml or INF- α (10 units/ul) (C). (D) Different C. volvatus fractions inhibited replication of VR2332 and HV strains in PAMs at a concentration of 1.2 mg/ml or INF-a (10 units/µl). Virus yield in the supernatant was also quantified. Culture treated with normal saline was set up as negative control (0 mg/ml), and that treated with INF- α was set up as positive control. The brightness of the fluorescence represents the level of the virus. Data are representative of three independent experiments (mean \pm SD). Statistical significance was analyzed by Student's t test. *P<0.05; **P<0.01; ***P<0.001. doi:10.1371/journal.pone.0079333.g005

to dissolve the crystals. The plates were further incubated at 37° C for 5 min to dissolve any air bubbles before absorbance due to the MTT signal was measured at 550 nm. The 50% cytotoxic concentration (CC_{50}) was analyzed by GraphPad Prism (Graph-Pad Software, San Diego, CA).

Figure 6. HPLC Chromatogram of C. volvatus fraction $C_{M-H-L-5}$ at 260 nm. (mobile phase: methanol:water = 0.5:95 (vol:vol), flow rate: 1 ml/min; temperature: 25° C, retention time: 11.280 min) $C_{M-H-L-5}$ yielded a main, sharp peak when subjected to analytical HPLC, and the purity of $C_{M-H-L-5}$ was 97%. doi:10.1371/journal.pone.0079333.g006

Real-time Reverse-transcription PCR (RT-PCR)

Total RNA was extracted from PAMs using the TRIzol reagent. RNAs were converted to cDNA using Superscript III Reverse Transcriptase (Invitrogen). In replication assay, PRRSV RNA was detected using quantitative real-time RT-PCR with primers designed against PRRSV ORF7 [31]. A plasmid containing PRRSV ORF7 sequence was used to generate a standard curve [32], and then RNA copies in all samples were calculated by comparing with it. For the transcript levels of cytokines, relative expressions of IFN- α , IFN- β , IL1- β and TNF- α in C_{M-H-L-5}treated or non-treated PAMs with or without PRRSV (HV strain) infection were quantified by the $2^{-\Delta\Delta}$ Method [33]. GAPDH (glyceraldehyde-3-phosphate dehydrogenase) mRNA was set as a control. The primers used for real-time PCR amplification were listed in Table 1.

Statistical Analysis

All experiments were performed with at least three independent replicates. Results were analyzed using Student's t test. Differences were considered to be statistically significant if the P value is less than 0.05. *P<0.05; **P<0.01; ***P<0.001.

Figure 7. Mass spectrometric detection of C. volvatus chromatographic fraction C_{M-H-L-5}. The HPLC-MS of C_{M-H-L-5} gave m/z data as the protonated molecular ion [M+H]⁺. The data of m/z of protonated molecular ion was deducted one, and then presented a molecular mass. doi:10.1371/journal.pone.0079333.g007

Results

Chromatographic Action of Aqueous Extract from C. Volvatus

The crude extract C_M was first chromatographed on HP-20 to remove impurities, and then the unadsorbed components were loaded onto HP-2MGL. The components adsorbed on HP-2MGL were eluted with an ethanol gradient (30%, 60% and 100% ethanol). The elution profile of the fractions from HP-2MGL is shown in Fig. 1. Three fractions were obtained from the HP-2MGL chromatography, corresponding to 30%, 60% and 100% ethanol eluate respectively. The fraction eluted with 30% ethanol was designated as C_{M-H} . The fractions eluted with 60% and 100% ethanol were not named (Fig. 1). Besides, the ultraviolet absorption of C_{M-H} was higher than any other fraction (fraction eluted with 60% or 100% ethanol), and showed several inner peaks, C_{M-H-1} ,

Figure 8. Detection of major chemical functional groups by color reactions. (A) Organic acid color reaction (0.05% BPB - ethanol solution): The BPB was used to detect the carboxylic acid group, along with yellow color change. (B) Ninhydrin color reaction: The existence of amide groups can induce mulberry color change in Ninhydrin color reaction. (C) Dragendorff color reaction: The compound, containing nitrogen atoms, can cause orange or yellow color change in Dragendorff color reaction. However, there are not any color changes when the nitrogen atoms are in the form of amide groups or not in existence.

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C_{M-H-2} and C_{M-H-3} , as indicated by OD 260 nm.

The fraction C_{M-H} was loaded on a DEAE-cellulose ion exchange chromatography column, and the unadsorbed fraction was collected. This fraction was selected for size exclusion chromatography on Sephadex LH-20 (Fig. 2). Five fractions resulted, $C_{M-H-L-1}$, $C_{M-H-L-2}$, $C_{M-H-L-3}$, $C_{M-H-L-4}$, and $C_{M-H-L-5}$. $C_{M-H-I-3}$ was a filemot powder eluted from Sephadex LH-20 as a single, sharp peak. $C_{M-H-L-5}$ was a white powder eluted from Sephadex LH-20 as a relative broad, small single peak.

We isolated and purified several fractions, C, C_M , C_{M-H} , and $C_{M-H-I-5}$, from the fruiting body of C. volvatus.

$C_{M-H-L-5}$ Inhibits PRRSV Replication

To explore the antiviral activity of $C_{M-H-L-5}$ against viruses, we first investigated its antiviral effect on PRRSV infection. The antiviral activity of related fraction was tested by IFA and the brightness of the fluorescence which represented the level of the virus. As shown in Fig. 3A and B, $C_{M-H-L-5}$ significantly inhibited PRRSV (CH-1a strain) replication in Marc-145cells. The extract at the concentration of 1.2 mg/ml reduced the virus yields by about 3000-fold when compared to normal saline control, and this inhibition was in a dose-dependent manner. To further verify its anti-PRRSV activity, we examined whether $C_{M-H-L-5}$ could inhibit replication of more than one PRRSV strain in PAMs. As illustrated in Fig. 3C, $C_{M\text{-}H\text{-}L\text{-}5}$ potently inhibited the replication of both the prototype of Type 2 PRRSV strain (VR2332) and HP-PRRSV strain (HV) in PAMs, which could reach a 300-fold suppression at the concentration of 1.2 mg/ml. The extract inhibited PRRSV infection with a 50% effective concentration (EC_{50}) value of 0.29 mg/ml for CH-1a strain in Marc-145 cells,

Figure 9. UV-VIS absorption spectrum of C_{M-H-L-5}. (start wavelength: 190 nm, end wavelength: 500 nm). UV-VIS absorption spectrum scan showed that the value of ultraviolet absorbance of $C_{M-H-L-5}$ at 260 nm was the highest. doi:10.1371/journal.pone.0079333.g009

0.29 mg/ml for HV strain, and 0.31 mg/ml for VR2332 in PAMs.

To exclude the possibility that nonspecific toxicity induced by the extract could affect PRRSV replication, we evaluated PAM and Marc-145 cell viability under various concentrations of $C_{M-H-L-5}$ using the MTT assay (Fig. 3D). The 50% cytotoxic concentrations (CC_{50}) of the $C_{M-H-L-5}$ for PAMs and Marc-145 cells were 55 mg/ml and 28 mg/ml, respectively, which greatly exceeded its EC_{50} value. The therapeutic index (CC_{50}/EC_{50}) was 97 for CH-1a strain in Marc-145 cells, 190 for HV strain, and 177 for VR2332 in PAMs.

These initial studies confirmed that the C. volvatus extract $C_{M-H-I,5}$ could inhibit PRRSV infection. Therefore, it is necessary for us to compare antiviral cytokine gene expression and different C. volvatus extracts on inhibition of PRRSV in subsequent studies.

Antiviral Cytokine Gene Expression in $C_{M-H-L-5}$ -treated Porcine Alveolar Macrophages

Cytokines are able to interfere with viral infection. Thus, we postulated that $C_{M\text{-}H\text{-}L\text{-}5}$ might induce antiviral cytokine expressions. To investigate this possibility, the expressions of four cytokines including IFN- α , IFN- β , IL1- β and TNF- α , known to be involved in antiviral response and inflammation, were analyzed in the presence or absence of C_{M-H-L-5}. Porcine alveolar macrophages (PAMs) were incubated with HV, HV plus $C_{M-H-L-5}$ (0.8 mg/ml) , or $C_{M\text{-}H\text{-}L\text{-}5}$ only, and real-time RT-PCR was performed to assess the relative mRNA level in PAMs after cultured for 12 h. $\rm C_{M\text{-}H\text{-}L\text{-}5}$ did not significantly induce IFN- α (Fig. 4A) and IFN- β (Fig. 4B) expression in both infected and noninfected PAMs. However, $C_{M\text{-}H\text{-}L\text{-}5}$ treatment could elevate the levels of IL-1 β (Fig. 4C) and TNF- α (Fig. 4D) expression in PAMs. The expressions of IL1- β and TNF- α exhibited 8-fold and 25-fold increase in $\rm C_{M\text{-}H\text{-}L\text{-}5}$ -treated infected cells respectively. Taken together, these data suggested that $C_{M-H-L-5}$ -impaired PRRSV infection could be partially due to the up-regulation of certain cytokines.

Effect of Different Extracts on Inhibition of PRRSV

Next, we investigated the effect of C, C_M , $C_{M\text{-H}}$, and $C_{M\text{-H-L-5}}$ on PRRSV infection to compare the antiviral activity of different C. volvatus extracts against PRRSV viruses. As shown in Fig. 5A and B, C, C_M , C_{M-H} and $C_{M-H-L-5}$ significantly inhibited PRRSV (CH-1a strain) replication in Marc-145 cells which produced different extents of inhibition of PRRSV at a concentration of 0.8 mg/ml. As the purification proceeded, the virus titration decreased to the lowest in C_{M-H} treatment, and then significantly increased following $C_{M\text{-}H\text{-}L\text{-}5}$ treatment. $C_{M\text{-}H}$ elicited the strongest inhibition compared with other components. The virus titration values of C_{M-H} and $C_{M-H-L-5}$ at a concentration of 0.8 mg/ml were $10^{2.5}$ TCID₅₀/ml and $10^{3.25}$ TCID₅₀/ml, respectively. The same trend occurred in treatment at 1.2 mg/ ml, and the virus titration values of $C_{M\text{-}H}$ and $C_{M\text{-}H\text{-}L\text{-}5}$ were $10^{2.1}$ $TCID_{50}/ml$ and $10^{2.75}$ $TCID_{50}/ml$, respectively (Fig. 5C). We further verified different C. volvatus extracts could inhibit replication of VR2332 and HV strains in PAMs in the same trends at a concentration of 1.2 mg/ml (Fig. 5D).

Taken together, these data showed that C, C_M , C_{M-H} , and $C_{M-H-L-5}$ produced different extents of inhibition of PRRSV.

Reverse Phase HPLC Analysis, Purification and Identification

Fraction $C_{M-H-L-5}$ was analysed or purified on an HPLC preparative column followed by an analytical column or a

preparative column with 5% methanol. $C_{M-H-L-5}$ yielded a single, sharp peak when subjected to analytical HPLC, and the purity of $C_{M-H-L-5}$ was 97%. Currently, the retention time of $C_{M-H-L-5}$ was 11.28 min (Fig. 6). Moreover, mass spectrometric analyses tested with electrospray ionization mass spectrometry (ESI-MS) showed that the m/z $[M+H]^+$ of $C_{M-H-L-5}$ was 404.0955 Da which presented a molecular mass as 403.1 Da (Fig. 7).

¹H-NMR and ¹³C-NMR Spectra were Acquired with the Use of a BRUKER

AVANCEIII-400 spectrometer. The NMR spectrum showed that there were very little impurities in the sample (please refer to supplementary data). Based on nuclear magnetic resonance spectra data and spectrum information of structure analysis, it was deduced that the sample structure contained amide groups and carboxylic acid groups ; The peak at δ 90-110 ppm in the ¹³C-NMR spectrum of $C_{M-H-L-5}$ indicated the absence of sugar fragments. However, the peaks at $\delta 60-70$ ppm showed the presence of polyol fragment containing 11 or 12 carbon atoms (please refer to supplementary data).

The amide groups and carboxylic acid groups were further indicated by results of chemical color reactions (Dragendorff, Ninhydrin and Organic Acid Reaction). As shown in Fig. 8, the existence of amide groups induced mulberry color change in Ninhydrin color reaction and not any color changes in Dragendorff color reaction. Moreover, Bromophenol blue detected the carboxylic acid group, along with yellow color change.

Integrating the above structural information, it was concluded that $C_{M-H-L-5}$ was a polyol fragment with amide groups and carboxylic acid groups.

Discussion

C. volvatus is commonly used as an anti-infective agent in traditional Chinese medicine. The inhibitory effect of C. volvatus has been previously demonstrated. Gao et al. (2013) found that the aqueous extract from the fruiting bodies of C. volvatus has the potential to be used for antiviral therapy [26]. In the present study, we further isolated and purified the antiviral compounds in the aqueous extract prepared from the fruiting bodies of C. volvatus.

Three kinds of chromatographic techniques were used in this study to isolate $C_{M-H-L-5}$ from *C. volvatus* in an efficient way [34]. Macroporous resin polymer contains a permanent network of pores, and the network is independent of the swelling state of the resin. The weak polar resin HP-2MGL with highly cross-linked structure is a good carrier which is appropriate for $C_{M-H-L-5}$ with weak polarity. DEAE-cellulose is a weakly basic anion exchanger, which can adsorb the undesired electronegative components. Sephadex is a gel filtration medium prepared by cross-linking dextran with epichlorohydrin. Different types of Sephadex differ in their degree of cross-linking and modifying groups, and hence the degree of swelling and the molecular fractionation range. Sephadex LH-20 is a well-known gel filtration medium for removing contaminants for small biomolecules including $C_{M-H-L-5}$ in a single step. Moreover, the sample was purified by using only distilled water and methanol which were safer than organic solvents such as chloroform, ethyl acetate and ether.

We also showed that C_{M-H-L-5} not only significantly inhibited CH-1a strain replication in Marc-145cells, but also VR2332 and HV strains in PAMs, excluding the possibility of nonspecific toxicity. All initial studies confirmed that $C_{M\text{-}H\text{-}L\text{-}5}$ could inhibit PRRSV infection (Fig. 3). Besides, the decline in brightness of fluorescence represents the increasing antiviral effect of the fraction and a decline in the fluorescence brightness was associated with a lower virus titration. In our studies, the decline in fluorescence brightness and virus titration were observed with the increase of concentration, indicating that the inhibition of $C_{M-H-L-5}$ was in a dose-dependent manner.

Our work provided evidences that $C_{M-H-L-5}$ was able to inhibit viral replication in vitro. In subsequent studies, two cytokines, IL1- β and TNF- α , were confirmed to be induced by $\rm C_{M\text{-}H\text{-}L\text{-}5}$ treatment. This provided a possibility that $C_{M-H-L-5}$ could indirectly inhibit the PRRSV replication by regulating some antiviral cytokines. However, more study is required to verify if the induced cytokines by $C_{M-H-L-5}$ play a critical role in the $C_{M-H-L-5}$ -induced inhibition of PRRSV replication. Nevertheless, our data suggested that $C_{M-H-L-5}$ is probably regulating the host immune response.

The aqueous extract from the fruiting bodies of C. volvatus is a crude extract, which has many components. The antiviral effects of the extract might result from a mixture of active compounds rather than from a single chemical entity, the different compounds in the mixture mediate a synergistic antiviral effect [26]. The efficacy of Traditional Chinese Medicine (TCM) is a characteristic of a complex mixture of chemical compounds present in the various herbs. The concept of combinatorial medicines has been exemplified by the drug cocktail used in the treatment of acquired immunodeficiency syndrome. C. volvatus fractions from every separation step differ in their PRRSV inhibitory potency. With the increase in purification, antiviral activity component first increased up to C_{M-H} , and then decreased with $C_{M-H-L-5}$ obviously. However, the antiviral activity of $C_{M-H-L-5}$ was still much higher than that of the crude extract C, and $C_{M-H-L-5}$ could achieve a 102 -fold suppression compared to crude extract C (Fig. 5). So we conclude that there are more than one active component exist in C. volvatus.

As we know, the optimal wavelength of nucleoside absorbance is about 260 nm and the value of ultraviolet absorbance of $C_{M-H-I-5}$ at 260 nm was the highest, which matched with the HPLC chromatogram and UV-VIS absorption spectrum scan (Figures 6 and 9). The amide groups not only exist in peptides, but also in several nucleosides, such as adenosine, guanosine, and uridine etc. Many antiviral nucleoside analogues derived from the aforementioned nucleosides have amide groups. Moreover, most antiviral drugs are nucleoside analogues which interfere with reverse transcriptase competitively [10]. Gao et al. proved that the crude extract of *C. volvatus* could interfere with reverse transcriptase for PRRSV inhibition [26]. As the most important antiviral component of C. volvatus, $\text{C}_{\text{M-H-L-5}}$ may play an important role on reverse transcriptase inhibition. At last, we had isolated guanosine and uridine in similar separation methods with $C_{M-H-L-5}$ for their similar physical properties. Those three substances may have

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similar structural features. Integrating the above information, it was concluded that $C_{M-H-L-5}$ could be a nucleoside analogue or a polypeptide with a polyol fragment, and the nucleoside analogue or polypeptide component was composed of amino acids by condensation.

C. volvatus exerts a range of generally beneficial effects on respiratory disease and cancer. Cryptoporic polysaccharides and cryptoporic acids have been isolated from C. volvatus as significant active ingredients [35,36,37,38]. Nowadays, most of the clinical antiviral drugs are nucleoside analogues [39]. In our study, $C_{M-H-L-5}$ was first extracted from the edible mushroom C. volvatus. Besides, the structural nature of $C_{M\text{-}H\text{-}L\text{-}5}$ was similar to both polysaccharides and nucleoside drugs, suggesting $C_{M-H-L-5}$ might have a potential for medicinal use.

Above all, although the chemical composition of $C_{M-H-L-5}$ of C. volvatus has been characterized, $C_{M-H-L-5}$ is a novel antiviral compound with a very complex structure.

Supporting Information

Figure S1 ¹H-NMR spectrum. ¹H-NMR spectra were acquired with the use of a BRUKER AVANCEIII-400 spectrometer. (TIF)

Figure S2 13 C-NMR spectrum (1). ¹³C-NMR spectra were acquired with the use of a BRUKER AVANCEIII-600 spectrometer. The peaks at $\delta 60$ –70 ppm showed the presence of polyol fragment containing 11 or 12 carbon atoms. (TIF)

Figure S3 13 C-NMR spectrum (2). ¹³C-NMR spectra were acquired with the use of a BRUKER AVANCEIII-600 spectrometer. The peak at $\delta 90$ –110 ppm in the ¹³C-NMR spectrum of $C_{M-H-L-5}$ indicated the absence of sugar fragments. (TIF)

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Author Contributions

Conceived and designed the experiments: ZM HW WHF TBN. Performed the experiments: ZM WZ LW MZ. Analyzed the data: ZM HW WHF TBN. Contributed reagents/materials/analysis tools: ZM WZ. Wrote the paper: ZM HW WHF TBN.

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A Novel Compound Inhibits PRRSV In Vitro

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