Screening for inhibitory effects of crude drugs on furin-like enzymatic activities

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Abstract
The spike (S) protein of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) contains a cleavage motif R-X-X-R for furin-like enzymes at the boundary of the S1/S2 subunits. The cleavage of the site by cellular proteases is essential for S protein activation and virus entry. We screened the inhibitory effects of crude drugs on in vitro furin-like enzymatic activities using a fluorogenic substrate with whole-cell lysates. Of the 124 crude drugs listed in the Japanese Pharmacopeia, aqueous ethanolic extract of Cnidii Monnieris Fructus, which is the dried fruit of Cnidium monnieri Cussion, significantly inhibited the furin-like enzymatic activities. We further fractionated the plant extract and isolated the two active compounds with the inhibitory activity, namely, imperatorin and osthole, whose IC50 values were 1.45 mM and 9.45 µM, respectively. Our results indicated that Cnidii Monnieris Fructus might exert inhibitory effects on furin-like enzymatic activities, and that imperatorin and osthole of the crude drug could be potential inhibitors of the motif cleavage.

Keywords
Furin · Proprotein convertase · SARS-CoV-2 · Coumarin · Imperatorin · Osthole

Introduction
In December 2019, a novel virus, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), belonging to the human coronavirus family, was identified in Hubei Province, China [1]. It causes coronavirus disease 2019 (COVID-19), a severe respiratory disease associated with a high mortality rate. According to the World Health Organization 2019 situation report of February 16, 2021, more than 100,000,000 patients have been diagnosed with COVID-19 and 2,300,000 have died worldwide. The entry of coronavirus into host cells is mediated by the spike (S) protein [2]. Processing of the S protein by cellular proteases, such as transmembrane protease serine 2 (TMPRSS2), cathepsin, and furin is necessary for protein activation and virus entry [3]. The S protein of SARS-CoV-2 consists of the NH2-terminal S1 domain and COOH-terminal S2 domain [2, 3]. The S1 domain has a receptor-binding domain (RBD) that binds to the host angiotensin-converting enzyme 2 (ACE2) receptor and the S2 domain has an fusion peptide (FP) domain that mediates membrane fusion. The S protein cleavage at the S1/S2 boundary by host cell protease plays a key role in binding the ACE2 receptor to the S1 domain. The S protein of SARS-CoV-2 has a cleavage motif R-X-X-R for furin-like enzymes at the S1/S2 boundary, matching the consensus amino acid motif of the substrate for furin and related proprotein convertases (PCs) [2, 3]. Furin/PC inhibitors block SARS-CoV-2 S protein cleavage to suppress viral entry [2–5]. In addition, SARS-CoV-2 pseudoviruses, which have a mutated S protein at the cleavage site, showed substantially decreased efficiency of entry into host cells [2–4]. Therefore, cleavage inhibitors of the motif site are expected to be therapeutic reagents for SARS-CoV-2 infection [6–8].

Furin, a member of the proprotein convertase family, is ubiquitously expressed in mammalian cells and activates
various proprotein substrates [9–11]. Furin regulates not only pathogenic pathways but also several physiological pathways, involving hormones, growth factors, adhesion molecules, and cell surface receptors [12]. Furin is involved in calcium-dependent proteolytic cleavage at the C-terminus of a consensus amino acid motif R-X-X-R↓ (the arrow indicates the cleavage position) [9].

Peptide-based small molecules, such as hexa-D-arginine (D-6R) and chloromethylketone (CMK) have been reported to be inhibitors of furin and other PCs [13–18]. However, furin/PC-targeting therapeutic reagents for clinical application have not been identified to date. Numerous studies have evaluated furin-like (furin and other PCs) enzymatic activities using a fluorogenic substrate with whole cell lysates [19–24]. In this study, the inhibitory effects of crude drugs were evaluated using the furin-like protease assay with a fluorescent peptide substrate.

Materials and methods

Materials

We selected 124 crude drugs listed in the Japanese Pharmacopeia, 17th Edition, and purchased them from several distributors (Supplementary Material, Table S1) [25]. Crude drugs (10 g) were refluxed with 300 mL of 70% EtOH for 1 h, and the resultant extracts were dried by evaporation. The samples were dissolved in dimethyl sulfoxide (DMSO) to a concentration of 10 mg/mL and stored at 4 °C until use. Imperatorin and osthole were obtained from Tokyo Chemical Industry Co., Ltd. (Tokyo, Japan) and FUJIFILM Wako Pure Chemical Corporation (Osaka, Japan), respectively.

Furin-like enzyme assay

A549 cells, human lung carcinoma epithelial cells, were obtained from RIKEN BioResource Center (Tsukuba, Japan) and cultured in Dulbecco’s modified Eagle’s medium containing 10% fetal bovine serum, 100 µg/mL streptomycin, and 100 units/mL penicillin. A549 cells were seeded in 100-mm-diameter dishes (1.0 × 10⁶ cells/plate) and cultured in staining 10% fetal bovine serum, 100 µg/mL streptomycin, and cultured in Dulbecco's modified Eagle’s medium containing 10% fetal bovine serum, 100 µg/mL streptomycin, and 100 units/mL penicillin. A549 cells were seeded in 100-mm-diameter dishes (1.0 × 10⁶ cells/plate) and cultured for 24 h at 37 °C with 5% CO₂. After 24 h, the cells were counted and treated with 1 mL of 2× lysis buffer (20 mM HEPES–KOH [pH 7.4], 0.5% Triton X-100, 1 mM CaCl₂) per 1.0 × 10⁶ cells. The cell lysates were vortexed for 1 min and centrifuged at 13,000 × g for 10 min at 4 °C. The supernatants were transferred to 1.5-mL tubes and stored at −80 °C until use. Supernatants (10 µL), crude drug extracts (10 µL), and H₂O (70 µL) were added to a 96-well black microplate and incubated at 37 °C for 30 min. Drug extracts were diluted and adjusted to a final concentration of 20 µg/mL for screening. To the mixture, 10 µL of 1 mM Pyr-Arg-Thr-Lys-Arg-methyl-coumaryl-7-amide (pyr-RTKR-MCA) was added (PEPTIDE INSTITUTE, Inc., Osaka, Japan). The mixture was incubated at 37 °C for 30 min, and fluorescence intensity of the sample was measured with excitation at 380 nm and emission at 460 nm using SpectraMax M2 (Molecular Devices, LLC, CA, USA). The 124 samples were subjected to screening using the furin-like enzyme assay, and the results are presented as mean ± standard deviation of at least three independent experiments. Ethylenediaminetetraacetic acid (EDTA, final conc. 50 mM) was used as the control in the assay. Half-maximal inhibitory concentration (IC₅₀) was obtained by logistic regression analysis using the drc package for R [26].

Extraction and isolation of the bioactive compounds

The dried fruits of Cnidii monnieri (100 g) were extracted three times with 70% aqueous EtOH (1 h, each) under reflux, and the solvent was evaporated in vacuo to obtain the corresponding extract (55 g). The extract was suspended in water and fractionated with ethyl acetate three times to obtain an ethyl acetate layer. The water-soluble portion was partitioned with n-BuOH three times. The yield of ethyl acetate soluble extract and n-BuOH soluble extract were 4.7 and 1.2 g, respectively. The ethyl acetate soluble extract (0.3 g) was subjected to chromatography on an ODS column (ODS-SM 50C; Yamazen Corporation, Osaka, Japan) with MeOH–H₂O (4:1, v/v) as a solvent to yield 16 fractions. Fraction 4 (12 mg) was chromatographed on a preparative HPLC column (Senshu Pak ODS-4151-N; 10 mm × 150 mm) eluted with MeOH–H₂O (2:1, v/v) and monitored at 254 nm to obtain 1 (5.2 mg). Fraction 6 (15 mg) was purified by HPLC (Senshu Pak ODS-4151-N; 10 mm × 150 mm) with MeOH–H₂O (2.8:1, v/v) as a solvent, and monitored at 254 nm to obtain 2 (12 mg).

Identification

Compounds 1 and 2 were identified as imperatorin and osthole, respectively. Their structures were confirmed by comparing their spectroscopic data, such as NMR and MS, with those of authentic compounds.

Results and discussion

We screened 124 crude drug extracts for inhibitory effects on furin-like activities. The furin-like activity was evaluated using pyr-RTKR-MCA as a fluorogenic substrate and cell lysates as whole proteolytic enzyme. Of the 124 crude
drug extracts, three extracts, Cnidii Monnieris Fructus (dried fruits of *C. monnieri*), Hydrangeae Dulcis Folium [dried leaves of *Hydrangea macrophylla* (Thunb.) Ser. var. *thunbergii* (Siebold) Makino], and Forsythiae Fructus [dried fruit of *Forsythia suspensa* (Thunb.) Vahl] suppressed furin-like activities by more than 40% (activity: 6.2% ± 0.3%, 56.5% ± 1.8%, and 42.9% ± 2.3%, respectively) (Table 1). We then evaluated the IC₅₀ of the three samples and Cnidii Rhizome (the dried rhizome of *C. officinale*) as the control. The IC₅₀ values of Cnidii Monnieris Fructus, Hydrangeae Dulcis Folium, and Forsythiae Fructus were 1.10, 7.12, and 6.52 µg/mL, respectively (Table 2). Cnidii Monnieris Fructus showed stronger inhibitory effects on furin-like activity than Cnidii rhizome (IC₅₀ > 50 µg/mL). Cnidii Monnieri Fructus (*Jashoshi* in Japanese) has been traditionally used to treat osteoporosis, sexual dysfunction, asthma, and skin ailments [27]. *Cnidium monnieri* Cusson contains several compounds, such as bergapten, imperatorin, osthole, and xanthotoxin [28]. Here, we fractionated and isolated bioactive compounds from Cnidii Monnieris Fructus contributing to the inhibitory effects on furin-like enzymatic activity. We isolated and identified two coumarin compounds, imperatorin and osthole, with inhibitory activity (Fig. 1). Osthole (IC₅₀ = 9.45 µM) showed significant inhibitory effects on furin-like enzymatic activity when compared with imperatorin (IC₅₀ = 1.45 mM). The autofluorescence of two coumarins (imperatorin and osthole) did not occur because reaction mixture (compounds and substrates) without cell lysates did not show fluorescence signal. These results indicate that Cnidii Monnieris Fructus might inhibit furin-like enzymatic activities, and that imperatorin and osthole of the crude drug could be candidates for inhibitors of motif cleavage.

In the present study, we screened the anti-furin-like activity of crude drugs using an in vitro furin-like assay with a fluorogenic substrate. Since furin is a Ca²⁺-dependent serine protease, EDTA, a popular chelating agent was used as positive control in this screening. However, a high concentration (IC₅₀ 50 mM) was required to exert its inhibitory activities. Although polyphenols, such as tannin is known to show chelating activities, our medicinal plant extracts containing polyphenols did not show inhibitory effects on furin-like activities. It is considered that the concentration of polyphenols in our medicinal extracts was not sufficient to exhibit inhibitory activity. Of the 124 crude drugs, Cnidii Monnieris Fructus showed strong inhibitory effects on furin-like activity, and two coumarin compounds (imperatorin and osthole) exerted inhibitory activity. Further studies are required to understand if Cnidii Monnieris Fructus and its bioactive compounds block S protein processing. For example, the inhibitory effect on S protein processing could be proven if the S protein expressed in *Escherichia coli* is used as a cleavage substrate instead of pyr-RTKR-MCA [29]. When the S protein gene was transfected into mammalian cells, the S

| Latin Name                  | Furin-like activity (%) |
|-----------------------------|-------------------------|
| ACHYRANTHIS RADIX           | 95.2 ± 1.8              |
| ACONITI RADIX PROCESSA      | 72.5 ± 3.6              |
| AKEBIAE CAULIS              | 92.8 ± 1.8              |
| ALISMATIS TUBER             | 101.9 ± 0.7             |
| ALOE                        | 83.1 ± 7.9              |
| ALPINIAE OFFICINARHIZOMA    | 92.2 ± 1.5              |
| AMOMI SEMEN                 | 92.6 ± 2.5              |
| ANEMARRHENAE RHIZOMA        | 95.4 ± 3.4              |
| ANGELICAE ACUTILOBAE RADIX | 101.8 ± 2.3             |
| ANGELICAE DAHURICAE RADIX  | 99.8 ± 3.1              |
| ARALIAE CORDATAE RHIZOMA   | 60.7 ± 4.2              |
| ARCTII FRUCTUS              | 97.2 ± 4.1              |
| ARECAE SEMEN                | 99.2 ± 2.3              |
| ARMENIACAE SEMEN           | 93.0 ± 9.7              |
| ARTEMISIAE CAPILLARIS FLOS | 80.0 ± 8.3              |
| ARTEMISIAE FOLIUM          | 79.0 ± 3.8              |
| ASIASARI RADIX              | 78.8 ± 2.7              |
| ASPARAGI RADIX              | 94.2 ± 1.1              |
| ASTRAGALI RADIX             | 92.4 ± 4.6              |
| ATRACTYLODIS LANCEAE RHIZOMA| 94.4 ± 3.1              |
| ATRACTYLODIS RHIZOMA        | 84.6 ± 3.1              |
| AURANTII FRUCTUS IMMATUS    | 86.3 ± 2.8              |
| AURANTII PERICARPII         | 100.3 ± 3.8             |
| BENINCASAE SEMEN           | 84.7 ± 0.3              |
| BUPLEURI RADIX              | 95.0 ± 2.0              |
| CANNABIS FRUCTUS            | 89.2 ± 1.7              |
| CARTHAMI FLOS               | 93.7 ± 5.3              |
| CASSIAE SEMEN               | 83.6 ± 4.9              |
| CATALPAE FRUCTUS            | 84.8 ± 2.8              |
| CHRYSANTHEMI FLOS           | 85.3 ± 4.9              |
| CIMICIFUGAE RHIZOMA         | 96.4 ± 2.6              |
| CINNAMOMI CORTEX            | 86.9 ± 5.7              |
| CITRI UNSHIU PERICARPII     | 80.8 ± 2.7              |
| CLEMATIDIS RADIX            | 85.9 ± 5.6              |
| CNIDII MONNIERIS FRUCTUS    | 6.2 ± 0.3               |
| CNIDII RHIZOMA              | 103.0 ± 3.3             |
| COICIS SEMEN                | 93.0 ± 2.3              |
| COPTIDIS RHIZOMA            | 82.9 ± 7.5              |
| CORNI FRUCTUS               | 89.8 ± 1.8              |
| CORYDALYS TUBER             | 86.5 ± 4.4              |
| CRATAEGI FRUCTUS            | 98.1 ± 2.4              |
| CURCUMAE RHIZOMA            | 78.2 ± 1.5              |
| CYPERI RHIZOMA              | 90.0 ± 0.9              |
| DIGENEIA                    | 101.1 ± 3.6             |
| DIOSCOREA RHIZOMA           | 92.1 ± 3.0              |
| EPHEDRAE HERBA              | 86.8 ± 4.7              |
| EPIPEMEDH HERBA             | 73.4 ± 11.3             |
| ERIOBOTRYAE FOLIUM          | 79.7 ± 3.1              |
| EUODIAE FRUCTUS             | 83.1 ± 2.5              |
protein was processed by furin/PC, and syncytial phenotype was observed [5, 30]. Evaluation of S protein processing by western blotting and syncytial formation by microscopy would provide direct evidence that the samples affect S protein processing and virus entry.

Osthole is a multifunctional compound with antioxidative, antiproliferative, anti-inflammatory, and antiallergic properties [31]. A recent study indicated that osthole suppressed TGF-β1-induced epithelial-mesenchymal transition (EMT) in lung cancer A549 cells [32]. Because TGF-β1 activates furin expression in several cell lines [33, 34], and

| Latin Name | Furin-like activity (%) |
|------------|-------------------------|
| FOENICULI FRUCTUS | 79.9 ± 11.7 |
| FORSYTHIAE FRUCTUS | 42.9 ± 2.3 |
| FRTILLARIAE BULBUS | 90.3 ± 3.0 |
| GARDENIAE FRUCTUS | 89.0 ± 4.2 |
| GASTRODIA TUBER | 98.9 ± 2.7 |
| GENTIANAE RADIX | 101.1 ± 8.1 |
| GENTIANAE SCABRAE RADIX | 99.6 ± 3.3 |
| GERANI HERBA | 91.2 ± 12.4 |
| GINSENG RADIX | 99.5 ± 4.4 |
| GINSENG RADIX RUBRA | 97.0 ± 1.4 |
| GLYCRRHIZAE RADIX | 91.3 ± 4.7 |
| GLYCRRHIZAE RADIX PRAEPARATA | 89.3 ± 1.4 |
| HOUTTUYNIAE HERBA | 92.9 ± 11.7 |
| HYDRANGEAE DULCIS FOLIUM | 56.5 ± 1.8 |
| KOI | 95.4 ± 5.3 |
| LONICERAE FOLIUM CUM CAULIS | 94.6 ± 2.2 |
| LONICERAE FOLIUM CUM CAULIS | 94.6 ± 2.2 |
| LCHRUS CUM CAULIS | 90.8 ± 4.6 |
| TRIBULI FRUCTUS | 95.1 ± 1.8 |
| TRICHOANTHIS RADIX | 87.1 ± 9.2 |
| UNCARIAE UNCIS CUM RAMULUS | 104.1 ± 5.8 |
| UVAE URSI FOLIUM | 89 ± 6.5 |
| VALERIANAE FAURIEI RADIX | 90.7 ± 6.2 |
| ZANTHOXYLII PIPERITI PERICARPUM | 82.4 ± 3.9 |
| ZEDOARIAE RHIZOMA | 77.1 ± 3.7 |
| ZINGIBERIS RHIZOMA | 96.8 ± 0.5 |
| ZINGIBERIS RHIZOMA PROCESSUM | 79.4 ± 6.1 |
| ZIZYPHI FRUCTUS | 97.7 ± 3.9 |
| ZIZYPHI SEMEN | 99.7 ± 1.7 |

Ethanol extracts of crude drugs (20 µg/mL) were pre-incubated with cell lysates and added to fluorogenic substrates (pry-RTKR-MCA). The data are presented as mean ± standard deviation of at least three independent experiments.

### Table 2 IC50 of different crud drugs

| Sample | IC50 (µg/mL) |
|--------|--------------|
| Cnidii Monnieris Fructus | 1.10 |
| Cnidii Rhizoma | > 50 |
| Hydrangeae Dulcis Foliu | 7.12 |
| Forsythiae Fructus | 6.52 |

protein was processed by furin/PC, and syncytial phenotype was observed [5, 30]. Evaluation of S protein processing by western blotting and syncytial formation by microscopy would provide direct evidence that the samples affect S protein processing and virus entry.

Osthole is a multifunctional compound with antioxidative, antiproliferative, anti-inflammatory, and antiallergic properties [31]. A recent study indicated that osthole suppressed TGF-β1-induced epithelial-mesenchymal transition (EMT) in lung cancer A549 cells [32]. Because TGF-β1 activates furin expression in several cell lines [33, 34], and
proteolytic processing of the TGF-β1 precursor by furin is an essential step in the formation of biologically active TGF-β1 [35], osthole might suppress TGF-β1-induced autocrine effects by blocking furin-like activities.

In conclusion, we screened the inhibitory effects of 124 crude drugs listed in the Japanese pharmacopoeia on in vitro furin-like enzymatic activities. Of these drugs, Cnidii Monnieris Fructus, which is the dried fruit of C. monnieri (Japanese name Jashoshi), strongly inhibited furin-like activity. We further isolated and identified two bioactive coumarins, imperatorin and osthole, from Cnidii Monnieris Fructus.

**Supplementary Information** The online version contains supplementary material available at [https://doi.org/10.1007/s11418-021-01519-9](https://doi.org/10.1007/s11418-021-01519-9).

**Declarations**

**Conflict of interest** The authors declare no conflict of interest.

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