Serine Proteases and their Inhibitors in Human Airway Epithelial Cells: Effects on Influenza Virus Replication and Airway Inflammation

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Abstract

Influenza virus replication and the production of inflammatory cytokines are associated with symptoms, including fever, and exacerbation of bronchial asthma and chronic obstructive pulmonary disease. Proteolytic activation of influenza viruses by serine proteases that are produced by airway epithelial cells is essential for viral entry and replication. Transmembrane protease serine S1 member (TMPRSS) 2, TMPRSS4 and TMPRSS11D have been detected certain cells, including the human alveolar epithelial cell line A549 and the surface epithelial cells of the human nasal mucosa, the trachea, the distal airways, and the lung. Several protease inhibitors, including aprotonin, reduce influenza virus replication. We previously demonstrated the following: (1) TMPRSSs (TMPRSS2, 4, and 11D) are present in primary cultures of human tracheal epithelial cells; (2) serine protease inhibitors, such as camostat and aprotonin, reduce the influenza virus replication and the release of the cytokines interleukin (IL)-6 and tumor necrosis factor (TNF)-α into cell supernatants; and (3) camostat reduces the cleavage of an influenza virus precursor protein, HA0, into the subunit HA1. These findings suggest that serine proteases expressed by human tracheal epithelial cells induce the proteolytic activation of influenza viruses and that serine protease inhibitors may reduce viral replication and the resultant production of inflammatory cytokines. Thus, serine protease inhibitors are potential candidates for anti-influenza virus drugs.

Here, we review the expression of serine proteases, the role of serine proteases in influenza virus activation, and the effects of serine protease inhibitors. In this review, we aim to introduce the effects of serine proteases and their inhibitors on influenza virus infection of human airway epithelial cells by discussing the findings of previous studies performed by our group and other research groups. Furthermore, the clinical features and virulence of influenza virus infection are reviewed to clarify the association of virus replication and cytokine release with disease severity.

Keywords: Airway epithelial cell; Camostat; Cytokine; Influenza virus; Proteases transmembrane protease serine S1 member; Serine protease inhibitor

Introduction

Homma and Ohuchi were the first to report that Sendai virus is activated when trypsin cleaves the viral surface glycoprotein and that the cleaved protein fuses the viral envelope to the host cell membrane and causes viral entry and replication [1,2]. Subsequent reports demonstrated that trypsin and other host proteases, by cleaving the surface glycoproteins important for membrane fusion, contribute to the spread of infection and the pathogenicity of many types of viruses, including paramyxoviruses and influenza viruses [3,4].

Serine proteases such as trypsin, transmembrane protease serine S1 member (TMPRSS) 2, TMPRSS4, and human trypsin-like protease (HAT; also known as TMPRSS11D) are important examples of host proteases that are responsible for the proteolytic cleavage of influenza virus hemagglutinin (HA), which is essential for viral genome entry to the cell and for the start of viral replication [5,6]. Furthermore, several protease inhibitors have been demonstrated to reduce the replication of influenza viruses [7-10], suggesting that these inhibitors may become candidates for anti-influenza drugs.

We previously demonstrated that serine proteases were produced by primary cultures of human tracheal epithelial cells (that retain the function of their tissue of origin [11]) and that proteases activated influenza virus HA through proteolytic cleavage. We also demonstrated that serine protease inhibitors, including camostat, reduced viral replication and inflammatory cytokine release from the cells [12].

Herein, we review the expression of serine proteases in airway epithelial cells, the role of serine proteases in influenza virus activation, and the effects of serine protease inhibitors on viral replication and cytokine release. Clinical features and treatment of influenza virus infection are also reviewed.

Clinical Features of Influenza Virus Infection and Anti-influenza Drugs

Human influenza virus infection causes a rapid onset of constitutional symptoms, including fever and lower respiratory tract symptoms. Chronic obstructive pulmonary disease (COPD) is one of
the leading causes of death worldwide, and the number of patients is increasing among older adults [13,14]. It is generally accepted that cigarette smoke is the most common identifiable risk factor for COPD [15]. Because smoking history is associated with the development of COPD, elderly people are more susceptible to COPD [14,15]. Influenza virus infection can exacerbate bronchial asthma and COPD during winter [16,17]. Therefore, vaccination is recommended to prevent exacerbation of bronchial asthma and COPD and to reduce the mortality rate of older people with COPD [18]. Furthermore, influenza vaccination is recommended for children and non-elderly adults to prevent severe illness in patients with influenza virus infection [19,20].

In addition, anti-influenza drugs, which include neuraminidase inhibitors such as Oseltamivir and Zanamivir, are beneficial in cases of uncomplicated pandemic and seasonal human influenza infection [21-23]. In contrast, certain types of patients with pandemic and seasonal influenza virus infection, including immuno-compromised patients, have died of pneumonia and multi-organ system failure despite intensive drug treatments, which include neuraminidase inhibitors and steroids [24,25]. Oseltamivir-resistant influenza A (H1N1) virus infection has been reported, and this type of seasonal influenza has caused severe disease in immuno-compromised patients [26]. In a previous study, Oseltamivir did not reduce inflammatory cytokine production by airway epithelial cells after infection with seasonal influenza viruses with an Oseltamivir-resistant genotype [27]. Moreover, Oseltamivir did not reduce viral titers or viral RNA levels. The 50% inhibitory concentration (IC50) of Oseltamivir for neuraminidase activity in the Oseltamivir-resistant seasonal virus was 300-fold higher than that observed for the pandemic influenza virus [27]. According to a report by Shobugawa et al., oral intake of Oseltamivir is feasible for children younger than 5 years old, whereas Zanamivir inhalation is prescribed for children ≥ 5 years old [28]. However, inhalation of Zanamivir may be difficult for patients with severe infections and for small children. Therefore, alternative anti-influenza virus drugs are needed.

The Virulence of Influenza Viruses

Because influenza viruses with a high virulence can cause severe illness and even fatality, precise information regarding the pathogenicity of these viruses is needed for effective treatment and to prevent confusion and fear during pandemic infection. The pathogenicity of the influenza type A (H5N1) virus has been reported to be associated with the non-structural gene segment [29], which contributes to an increase in inflammatory cytokine production. Furthermore, several studies have reported mechanisms of high pathogenicity, which include an elevation in pulmonary concentrations of inflammatory cytokines (e.g., interleukin (IL)-1, IL-6 and interferon (IFN)-γ), a decrease in anti-inflammatory cytokine production, and an elevation in viral replication [30,31].

Influenza viral replication induces cytokine production and cell damage in the airway epithelium, which is the first target of infection. Infection-induced production of inflammatory cytokines, including IL-6 and tumor necrosis factor (TNF)-α, and proteases may cause damage to airway and alveolar epithelial cells and to vascular endothelial cells [32-34]. This damage may subsequently exacerbate bronchial asthma and COPD and develop into pneumonia and acute respiratory distress syndrome (ARDS) [24,35-37] (Figure 1). IL-6 and TNF-α are associated with cell death and the activation of caspases in swine macrophages after pandemic A/H1N1 viral infection [38]. Patients infected with highly pathogenic influenza viruses experience increased viral replication and subsequent hypercytokinemia [30,39]. Influenza viral infection-induced cell damage is partly related to the magnitude of viral replication, as well as NF-κB-p65-mediated IL-6 production and caspase-3 activation [40]. Furthermore, a large number of CD8+ T cells and granzyme B+ cells have been found to be expressed in autopsy lung tissue specimens from patients infected with pandemic A/H1N1 influenza virus [32]. Granzymes are granule-contained lymphocyte serine proteases, and CD8+ cells exert their cytotoxic functions by releasing granzymes in response to influenza virus infection [41]. Because IL-6 induces granzyme production [42], hypercytokinemia during influenza virus infection may activate CD8+ cells, resulting in lung damage through the cytotoxic activities of granzymes (Figure 1).

To ameliorate the symptoms associated with the exacerbation of bronchial asthma and COPD caused by virus infection, including influenza virus infection, oral steroids and inhalation of β2 agonists have been used [43-45]. Furthermore, treatment with antibiotics is recommended when COPD patients have purulent sputum caused by a secondary bacterial infection [46].

Serine Proteases and Their Roles in Influenza Virus Replication in the Airways

The pathway of influenza virus attachment, internalization and uncoating

The hemagglutinin (HA) of human seasonal and classical H1N1 swine influenza viruses binds to the receptor Saa2, 6Gal, and the HA of most avian and equine viruses binds to Saa2, 3Gal [47]. Expression of SAA2, 6Gal was observed in epithelial cells in the nasal mucosa, pharynx, tracheae and bronchi [48-51]. SAA2, 3Gal is expressed in the human trachea, nonciliated cuboidal bronchiolar cells and type II cells lining the alveolar wall [49,52].

After the attachment of influenza virus HA to the receptor, viruses enter airway epithelial cells and are internalized by endocytic compartments via four internalization mechanisms: clathrin-coated pits; caveolae; a nonclathrin, noncaveola pathway; and macropinocytosis [53]. Of these mechanisms, the nonclathrin, noncaveola-mediated internalization pathway depends on a low pH, which triggers the fusion of the viral and endosomal membranes [54]. After binding and internalization of the virus by receptor-mediated endocytosis, the activated influenza virus HA, which is cleaved by serine proteases, induces fusion of the viral envelope to the host cell membrane in acidic endosomes [3,4]. The virus then releases its ribonucleoproteins containing viral RNA into the cytoplasm, resulting in the next steps of viral replication [53,55].

The expression of serine proteases in the airways and other organs

TMPRSSs are expressed in a broad range of human tissues (Table 1). TMPRSS2 is expressed in the human nasal and tracheal mucosa, distal airways, and lung [6,12,56,57]. TMPRSS2 is also expressed in the human Caco-2 colon cancer cell line, the human Huh7 hepatocellular carcinoma cell line [58], and swine airway epithelia [59]. TMPRSS4 is expressed in lung cancer tissue [60], the mouse lung [61] and the human trachea [12], and TMPRSS11D is expressed in human bronchi and tracheae [12,62,63] and in swine airway epithelia [59].
Furthermore, the expression of a trypsin-like protease was identified in rat bronchiolar Clara cells [64]. We identified the TMPRSS2 and TMPRSS11D protein at the cell membrane and in the cytoplasm of primary cultures of human tracheal epithelial cells using indirect immunofluorescence assay (Figure 2) [12]. We also detected mRNA encoding TMPRSS2, TMPRSS4 and TMPRSS11D in the cells [12]. Our findings pertaining to the expression of TMPRSS2, TMPRSS4 and TMPRSS11D in primary cultures of human tracheal epithelial cells [12] are consistent with the expression observed in previous reports.

**Structure of TMPRSSs and their functions other than influenza virus activation**

TMPRSS2, TMPRSS4 and TMPRSS11D belong to the type II transmembrane serine protease (TTSP) family, which was identified as a new family of S1 class serine proteases. TTSPs are characterized by an N-terminal transmembrane domain that anchors the enzymes to the plasma membrane, a stem region, and a catalytic domain of the chymotrypsin S1 type. The most prominent member of the TTSP family is the digestive enzyme enteropeptidase, which was originally isolated as a soluble protease and was later shown to be the catalytic domain of the membrane-bound protease [65]. TTSPs are synthesized as single-chainzymogens that are activated into a mature form by C-terminal cleavage of a highly conserved arginine or lysine residue.

The physiological functions of TMPRSS2, TMPRSS4 and TMPRSS11D, other than proteolytic activation of influenza virus HA, have not been completely identified. However, TMPRSS2 also activates the spike protein of the severe acute respiratory syndrome coronavirus (SARS-CoV) on the cell surface [66], and camostat inhibits TMPRSS2-dependent infection by SARS-CoV [67]. The serine protease inhibitor aprotinin inhibits mucin production by an airway epithelial cell line through its action on TMPRSS11D [68]. Camostat and aprotinin inhibit sodium channel function in human bronchial epithelial cells, which can be reversed by saturating concentrations of trypsin [69], although the specific TMPRSSs affecting the channel function have not been identified. A variety of functions are attributed to TMPRSSs,
including prostate carcinogenesis and regulation of airway surface fluid volume by proteolytic cleavage of sodium channels by TMPRSS2 [57,70]; tissue development, cell differentiation, cancer progression and metastasis by TMPRSS4 [60,71,72]; and prevention of fibrin deposition in the airway lumen by TMPRSS11D [73].

Roles of serine proteases in influenza virus replication

The first report by Homma et al. [1] and subsequently published reports demonstrated that trypsin cleaved the surface glycoproteins of viruses (such as the fusion protein of Sendai virus and HA of the influenza virus), and the cleaved proteins caused fusion of the viral envelope to the host cell membrane, which is required for the spread of infection and pathogenicity [1-4].

Furthermore, the role of host serine proteases, including TMPRSS2, TMPRSS4 and TMPRSS11D, in the proteolytic activation of influenza virus HA and viral replication have been demonstrated using A549, 293T and Caco-2 cell lines transiently expressing the proteases TMPRSS2, TMPRSS4, and TMPRSS11D [6,58,74]. The cleavage site of influenza HA is located in a loop that protrudes prominently from the viral surface [75]. The subunit HA1 of A/Sendai-H/108/2009/(H1N1) pdm09 influenza virus, which is produced by cleaving the influenza virus precursor protein HA0, was detected in the supernatants of primary cultures of human tracheal epithelial cells expressing TMPRSS2,TMPRSS4 and TMPRSS11D (Figure 3) [12]. TMPRSS2 and TMPRSS11D were detected at the cell membrane and in the cytoplasm of human tracheal epithelial cells (Figure 2) [12]. A serine protease inhibitor, camostat, reduced the amount of HA1 in the supernatants (Figure 3). These findings are consistent with reports that host proteases promote proteolytic activation of influenza virus HA in the trans-Golgi network and/or plasma membrane (Table 1) [76,77]. Newly synthesized influenza virus with cleaved HA may bud from the epithelial cell membrane.

Staphylococcus aureus secretes a protease that exerts a decisive influence on the outcome of influenza virus infection in mice through cleavage-mediated activation of the viral HA [78]. Extracellular proteolytic activation of HA by trypthase Clara isolated from rat lungs was also reported [79]. In contrast, Zhircov et al. demonstrated that cleavage of HA in human adenoid epithelial cells occurs intracellularly

Figure 2: A and B: Indirect immunofluorescence staining of TMPRSS2 (A) and TMPRSS11D (B) in primary cultures of human tracheal epithelial (HTE) cells. TMPRSS2 and TMPRSS11D are stained orange at the cell membrane and in the cytoplasm. Nuclei are stained blue. Magnification: x 630. C: Expression of TMPRSS2 mRNA, TMPRSS4 mRNA and TMPRSS11D mRNA in HTE cells treated with camostat (10 mg/ml) or the vehicle control (1% water). The results are expressed as the ratio of TMPRSSs (TMPRSS2, TMPRSS4 or TMPRSS11D) mRNA expression compared with β-actin mRNA and are reported as the mean ± SEM (n=3). Significant differences compared to the values of TMPRSS2 in the cells treated with vehicle alone (Vehicle) are indicated by **p<0.01. Significant differences compared to the values of TMPRSS4 in the cells treated with vehicle alone (Vehicle) are indicated by ++p<0.01. Camostat did not affect the expression of mRNA of three types of TMPRSSs [12].
by cell-associated protease and that influenza virus is also activated intracellularly in human intestinal Caco-2 cells [10,77]. Furthermore, Böttcher-Friebertshäuser et al. [80] demonstrated that TMPRSS2 and TMPRSS11D cleaved newly synthesized HA before or during the budding of newly synthesized influenza virus. TMPRSS11D can also cleave the HA of incoming viruses prior to endocytosis at the surface of Madin Darby Canine Kidney (MDCK) cells transiently transfected with these proteases (Table 1). In contrast, the authors reported low levels of enzymatic activity for soluble TMPRSS2 and TMPRSS11D in the supernatants of MDCK cells [80]. Thus, the site of influenza virus HA activation in human airway epithelial cells in vivo is still uncertain, and influenza virus HA activation may also occur outside of cells by soluble proteases released by host cells.

| Localization | Cell type or tissue localization of TMPRSS protein and mRNA | Reference | Year |
|--------------|-------------------------------------------------------------|-----------|------|
| **TMPRSSs**  | Human lung (mRNA)                                           | [56]      | 1999 |
| **TMPRSS2**  | Human nasal, tracheal and distal airways (mRNA)             | [57]      | 2002 |
|              | Human alveolar epithelial cells (mRNA)*                     | [6]       | 2006 |
|              | Caco-2 and Huh-7 cell line (protein and mRNA)*              | [58]      | 2010 |
|              | Swine airway (protein and mRNA)                             | [59]      | 2014 |
|              | Human tracheal epithelial cells (protein and mRNA)          | [12]      | 2015 |
| **TMPRSS4**  | Human lung cancer tissue (mRNA)                             | [60]      | 2008 |
|              | Caco-2 and Huh-7 cell line (mRNA)*                          | [58]      | 2010 |
|              | Mouse lung (mRNA)                                           | [61]      | 2011 |
|              | Human tracheal epithelial cells (protein and mRNA)          | [12]      | 2015 |
| **TMPRSS11D**| Human bronchi and trachea (protein)                         | [62]      | 1997 |
|              | Human trachea (mRNA)                                        | [63]      | 1998 |
|              | Swine airway (protein and mRNA)                             | [59]      | 2014 |
|              | Human tracheal epithelial cells (protein and mRNA)          | [12]      | 2015 |

| Proteolytic activation of influenza | Cells | Reference | Year |
|------------------------------------|-------|-----------|------|
| **TMPRSS2 and 11D**               | A549 cell line | [6] | 2006 |
| **TMPRSS2 and 4**                 | 293T cell line | [74] | 2009 |
| **TMPRSS2 and 4**                 | Caco-2 cell line | [58] | 2010 |
| **TMPRSS2, 4 and 11D**            | Human tracheal epithelial cells | [12] | 2015 |

| Site of proteolytic activation of influenza | Site | Reference | Year |
|--------------------------------------------|------|-----------|------|
| **-**                                      | The late stages of intracellular transport in the trans-Golgi and plasma membrane | [77] | 2003 |
| **TMPRSS2 (1) and 11D (1 and 2)**          | (1) Before or during budding; (2) Incoming viruses prior to endocytosis at the surface of cells | [80] | 2010 |
| **TMPRSS2, 4 and 11D**                     | Before or during budding | [12] | 2015 |

| Inhibitory effects of serine protease inhibitors on influenza virus replication | Cell type or tissue | Reference | Year |
|-----------------------------------------------------------------------------|---------------------|-----------|------|
| **Aprotinin**                                                                | Chicken embryo     | [82]      | 1985 |
|                                                                             | MDCK cells         | [8]       | 1992 |
Human adenoid epithelial cells | [10] | 2002

Human tracheal epithelial cells | [12] | 2015

Camostat

MDCK cells | [8] | 1992

Mouse airway, MDCK cells | [8] | 1996

Human tracheal epithelial cells | [12] | 2015

Gabexate

MDCK cells | [8] | 1992

Human tracheal epithelial cells | [12] | 2015

Leupeptin

Mouse lung | [83] | 1987

MDCK cells | [8] | 1992

Human adenoid epithelial cells | [10] | 2002

Nafamostat

MDCK cells | [8] | 1992

Mouse lung | [61] | 2011

| Table 1: TMPRSSs: Localization, activation, cleavage site, and inhibitors of influenza virus. AEBSF: 4-(2-aminoethyl)-benzolsulfonylfluoride; A549: human type II lung cell line; Caco-2: human colon cancer cell line; Huh-7: human hepatocellular carcinoma cell line; MDCK: Madin Darby Canine Kidney; pAB: p-aminobenzidine; TMPRSS: transmembrane protease serine S1 member. *Replication of influenza viruses and/or cleavage of influenza virus HA were examined in the cells co-transfected with plasmids encoding TMPRSSs.

Inhibitory effects of serine protease inhibitors on influenza virus replication

The serine protease inhibitor aprotinin and similar agents, such as leupeptin and camostat, suppress virus HA cleavage and reduce the replication of influenza viruses with a single arginine in the HA cleavage site [7]. Protease inhibitors, including e-aminocaproic acid, can substantially reduce fatality in mice after influenza virus infection [81]. Similarly, viral replication in human adenoid epithelial cells was inhibited by aprotinin [10]. The protease inhibitors 1-1, 1-2 and 1-3 inhibit influenza HA cleavage in MDCK cells co-transfected with plasmids encoding TMPRSS2 and TMPRSS11D [80]. Similarly, the protease inhibitors 4-(2-aminoethyl)-benzolsulfonylfluoride (AEBSF) and p-aminobenzidine (pAB) decrease influenza virus replication in mouse lung cells expressing proteases, including TMPRSS4 [61]. Furthermore, serine protease inhibitors, including camostat and aprotinin, inhibit HA cleavage and influenza virus replication in human tracheal epithelial cells that express TMPRSS2, 4 and 11D [12]. The inhibitory effects of protease inhibitors, such as aprotinin, camostat, gabexate, leupeptin and nafamostat, on influenza virus replication have also been reported in MDCK cells, human airway epithelial cells, chicken embryos and mice (Table 1) [7-10,12,82,83].

Camostat, a serine protease inhibitor used to treat patients with pancreatitis [84], reduced the replication of the influenza A/H1N1 pdm 2009 and A/H3N2 viruses in primary human tracheal epithelial cells as evaluated by viral titers and viral RNA levels (Table 2) [12]. Other types of serine protease inhibitors, such as aprotinin, gabexate and sivelestat, also reduced viral titers and/or RNA levels in the cells.

Thus, a variety of serine protease inhibitors reduce influenza virus replication in human and mouse airway and alveolar epithelial cells (Table 2). Serine protease inhibitors may prevent the exacerbation of bronchial asthma and COPD and the development of pneumonia and ARDS by reducing viral replication (Figure 1).

The effects of serine protease inhibitors on inflammatory cytokine release

IL-6 and TNF-α are associated with disease symptoms and severity in influenza-infected patients [39,85] and with cell damage [40]. Camostat and gabexate inhibit lipopolysaccharide-induced TNF-α release from macrophages [86] and the development of influenza-associated pneumonia in mice by inhibiting the production of cytokine, including IL-6 [87]. Aprotinin reduces ICAM-1 expression in human umbilical vein endothelial cells [88] and myocardial IL-6 gene expression [89].
expression after cardiac ischemia and reperfusion in rats [89]. Serine protease inhibitors, including camostat, reduce production of IL-6 and TNF-α in human tracheal epithelial cells (Table 2) [12]. Furthermore, treatment of epithelial cells with an anti-IL-6 antibody attenuated cell damage in this model [40]. These findings suggest that the serine protease inhibitors may have anti-inflammatory effects in the lung, airways and other organs of patients with an influenza virus infection.

|               | Aprotinin | Camostat | Gabexate | Sivelestat |
|---------------|-----------|----------|----------|------------|
| Viral release | ++        | ***      | +        | NS         |
| Viral RNA replication | ++        | ***      | +        | +          |
| IL-6          | +         | ++       | NS       | NS         |
| TNF-α         | +         | ++       | NS       | NS         |

Table 2: Inhibitory effects of serine protease inhibitors on viral release, RNA replication, and cytokine release after influenza A/H1N1 pdm 2009 virus infection. A/H1N1 pdm 2009: Influenza A/H1N1 pdm 2009 virus; Viral release and RNA replication: The viral titers in the supernatants and viral RNA replication in the primary cultures of human tracheal epithelial cells 5 days after exposure to the influenza A/H1N1 pdm 2009 virus in the presence of camostat (10 mg/ml), sivelestat (10 mg/ml), gabexate (10 mg/ml), aprotinin (1000 KIU/mL) or the vehicle control; Cytokine release: The concentration of IL-6 and TNF-α in the supernatants before or 5 days after exposure to the influenza A/H1N1 pdm 2009 virus in the presence of camostat (10 mg/ml), sivelestat (10 mg/ml), gabexate (10 mg/ml), aprotinin (1000 KIU/ml) or the vehicle control. Significant differences from the viral infection alone are indicated by +p<0.05, ++p<0.01 and +++p<0.001. NS: Not significant [12].

In addition to the reduction of viral replication, serine protease inhibitors may provide clinical benefits by preventing the exacerbation of bronchial asthma and COPD and the development of pneumonia and ARDS, through reducing production of cytokines, granzymes and other types of cytotoxic factors (Figure 1).

Both neuraminidase inhibitors and serine protease inhibitors reduce the production of inflammatory cytokines in vitro [12,27]. Therefore, the combined use of the two inhibitors may enhance the inhibitory effects on cytokine production and may serve as a more efficient antiviral therapy. However, because physicians usually prescribed only one type of neuraminidase inhibitor for treatment, these serine protease inhibitors may be used alone at first, even in cases when these inhibitors can be clinically used. The effects of combination therapies involving serine protease inhibitors and other types of drugs, such as neuraminidase inhibitors and a polymerase inhibitor, should be elucidated.

The effects of granzymes and their inhibitors on inflammation and influenza virus infection

Because granzymes are granule-stored lymphocyte serine proteases and are produced in CD8+ cells in response to influenza virus replication [41], it could be speculated that serine protease inhibitors inhibit the cytotoxic activities of granzymes themselves and reduce CD8+ cell activation and granzyme production in response to virus replication. Serine protease inhibitors reduce the production of cytokines, including IL-6 [12], that induce granzyme production [42]. Thus, serine protease inhibitors could inhibit the cytotoxic activities of granzymes produced by immune cells during influenza virus replication through these mechanisms. However, the effects of granzymes and other serine proteases and their inhibitors on inflammation in response to influenza virus infection are unclear.

Aprotinin has been shown to inhibit the L-glutamyl-2-naphthylamide activity of granzyme B in a human cytolytic lymphocyte line, Q31, whereas leupeptin does not have a significant inhibitory effect [90]. In contrast, Zhong et al. reported that granzyme K inhibited replication of influenza virus by cleaving the nuclear transport complex (the importin α1/β dimer) of infected host cells [91]. However, the effects of granzymes and other serine proteases and their inhibitors on virus entry, including the proteolytic activities of influenza HA, are still unclear. Further studies are required to clarify these effects.

Potency of serine protease inhibitors

Of all the protease inhibitors studied (aprotinin, camostat, gabexate and sivelestat) in human tracheal epithelial cells, camostat was the most potent inhibitor of influenza virus replication and inflammatory cytokine production (Table 2) [12]. This is supported by the similar levels of potency observed in a report by Hosoya et al. using MDCK cells [8]. Observations of the MDCK influenza infection model suggest that nafamostat may be more potent than camostat and that leupeptin and pepstatin are less effective than aprotinin [8]. Thus, of the protease inhibitors studied, camostat and nafamostat may have the highest potential for reducing influenza virus replication and/or inflammatory cytokine production (Table 2) [8,12]. However, the exact difference in the potency between camostat and nafamostat has not been examined in human airway epithelial cells.

Conclusion

The serine proteases of host cells induce the proteolytic activation of influenza virus HA, and serine protease inhibitors reduce the replication of the influenza virus within, and the production of inflammatory cytokines by, human airway epithelial cells. In vitro models suggest that serine protease inhibitors are a potential treatment for influenza patients, but animal experiments are required to confirm their anti-viral effects and low toxicity. Ultimately, clinical trials in influenza-infected patients will be needed to assess the clinical benefits of these inhibitors.

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Conflicts of Interest

All authors have no conflict of interest. Yamaya is a professor in the Department of Advanced Preventive Medicine for Infectious Disease, Tohoku University Graduate School of Medicine. This department had been funded by eleven pharmaceutical companies until March 31, 2014, including Ono Yakuhin Co., Ltd. which provided camostat mesilate, sivelestat and gabexate mesilate. From April 1, 2014, this department is funded by eight pharmaceutical companies, which are as follows: Kyorin Pharmaceutical Co. Ltd., Abbott Japan Co., Ltd., Taisho Toyama Pharmaceutical Co., Ltd., AstraZeneca Co. Ltd, Otsuka

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