



Anti-HCMV activity by an irreversible p97 inhibitor LC-1310

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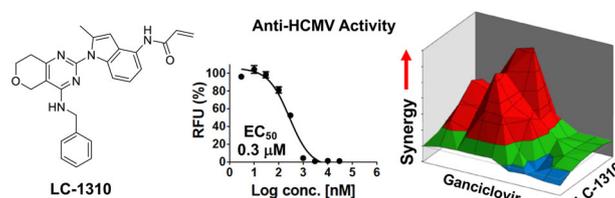
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Abstract

The AAA+ (ATPase associated with various cellular activities) protein p97, also called valosin-containing protein, is a hexameric ring ATPase and uses ATP hydrolysis to unfold or extract proteins from biological complexes. Many cellular processes are affected by p97 including ER-associated degradation, DNA damage response, cell signaling (NF- κ B), cell cycle progression, autophagy, and others. Not surprisingly, with its role in many fundamental cellular processes, p97 function is important for the replication of many viruses. We tested irreversible p97-targeting compounds for their ability to inhibit the replication of multiple viruses compared to the known p97 inhibitors NMS-873 and CB-5083. Our results indicate that overall cellular toxicity for p97 compounds provides a challenge for antivirals targeting p97. However, we identified one compound with sub-micromolar activity against human cytomegalovirus and improved cell viability to provide evidence for the potential of irreversible p97 inhibitors as antivirals.

Graphical Abstract



Keywords p97 · Valosin-containing protein · p97 inhibitors · Human cytomegalovirus · Antiviral

Introduction

The AAA+ (ATPase associated with various cellular activities) protein p97, also called valosin-containing protein, is a hexameric ring ATPase and uses ATP hydrolysis to unfold or extract proteins from biological complexes [1].

Extracted proteins are often ubiquitinated to facilitate their degradation by the 26S proteasome [2]. Many cellular processes are affected by p97 including ER-associated degradation (ERAD), DNA damage response, cell signaling (including NF- κ B), cell cycle progression, autophagy, and others. p97 is one of the most abundant proteins in cultured cells and yet upregulation of p97 levels in many cancers can signal a poor clinical outcome [3]. Therefore, p97 has been put forward as an anticancer target and a p97 inhibitor entered into a phase I clinical trial [4].

The p97 monomer has three major domains. The p97 N-terminal domain can interact with a wide range of adaptor proteins to result in different subcellular locations and substrate recognition. The two other domains, D1 and D2, have ATPase activity, however, ATP hydrolysis by D1 is relatively inactive compared to that of D2 [5]. The D1 domains promote hexamer formation and overall structural integrity. The D2 domain likely provides the major ATP

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hydrolysis activity for hexameric p97 extraction of ubiquitinated proteins.

A variety of molecular interactions have been described for p97 inhibitors. Eeyarestatin features a nitrofuranyl group thought to bind D1 and an aromatic domain that may interact with membranes to increase target specificity [6]. UPCDC30245 binds at the junction between D1 and D2 and may prevent conformational changes resulting from ATP binding [7]. NMS-873 probably also binds at the D1/D2 junction and is not competitive with ATP for inhibition [8, 9]. CB-5083 binds D2 and is competitive with ATP [10, 11]. CB5083 has been in a phase I trial with the ultimate goal of advancing it as a solid tumor or multiple myeloma treatment [4], however, the trial was halted due to adverse effects perhaps caused by unintended inhibition of phosphodiesterase-6 [12].

Not surprisingly, with its role in a myriad of fundamental cellular processes, p97 function is important for the replication of many viruses. Viruses rely upon processes such as ERAD, ribosome maintenance, mRNA transport, and autophagy to avoid host detection, facilitate viral protein expression and virus replication, including particle assembly and release [1, 12–14]. Reduction of p97 expression, mainly by siRNA knockdown, inhibits coronavirus (CoV), human cytomegalovirus (HCMV), enterovirus, Rift Valley fever virus, chikungunya virus, influenza A virus (IAV), poliovirus, and West Nile virus replication [15–22]. Published p97 inhibitors have also displayed antiviral activity against many of the aforementioned viruses and a baculovirus [16, 17, 19, 20, 22–27].

We recently described the design and synthesis of novel irreversible small molecule inhibitors targeting p97 [28]. We tested representatives of those irreversible p97-targeting compounds for their ability to inhibit the replication of multiple viruses compared to the known p97 inhibitors NMS-873 and CB-5083. Our results indicate that overall cellular toxicity for p97 compounds provides a challenge for antivirals targeting p97. However, we identified one compound that demonstrated sub-micromolar activity against human cytomegalovirus with improved cell viability to provide evidence for the antiviral potential of these irreversible p97 inhibitors.

Results/discussion

The ring ATPase p97 is involved in many important cellular processes such as ERAD, autophagy, endosomal trafficking, and regulation of multi-protein complex function. Viruses rely upon many cellular processes to establish and spread to new hosts; therefore, it is not surprising that p97 would be implicated as an important cellular factor for the replication of many viruses. We have recently published the

design and synthesis of a series of irreversible p97 inhibitors [28] and were interested in testing representatives of those and other known p97 inhibitors (Fig. 1) for possible antiviral activity using a variety of viruses.

HCMV is a human herpesvirus. Herpesviruses are enveloped viruses with large double-stranded DNA genomes (~236 kbp for HCMV). Herpesvirus genome replication, capsid assembly, and gene transcription occur in the nucleus. Gene expression involves a regulatory cascade pattern during lytic replication where genes are expressed early, midway or late in virus replication. HCMV requires p97 expression for virus replication and known p97 inhibitor, NMS-873, reduced virus replication [16]. We tested our irreversible p97 inhibitors for inhibition of HCMV replication in primary human foreskin fibroblasts (HFFs). Our approach was to inoculate HFFs with HCMV ADCREGFP, a green fluorescent protein (GFP) reporter virus [29], add compounds, and assess GFP expression 6 days later as a measure of virus replication and spread. We have used the virus-dependent GFP expression assay to evaluate many types of inhibitors [30–32]. All p97 inhibitors were evaluated in dose–response experiments for antiviral activity (EC_{50}) and cell viability (CC_{50}). Only LC-1310 showed clear antiviral activity (EC_{50} 0.3 μ M) in the absence of toxicity (CC_{50} 12 μ M) with a tissue culture therapeutic index (EC_{50}/CC_{50}) of 40 (Fig. 2 and Table 1).

The disruption of HCMV regulation of gene expression was implicated as a mechanism for the NMS-873-dependent reduction in virus replication [16]. Expression of the immediate-early protein 2 (IE2), a major viral gene expression factor, was reduced in infected and NMS-873-treated cells [16]. An interesting aspect of a possible

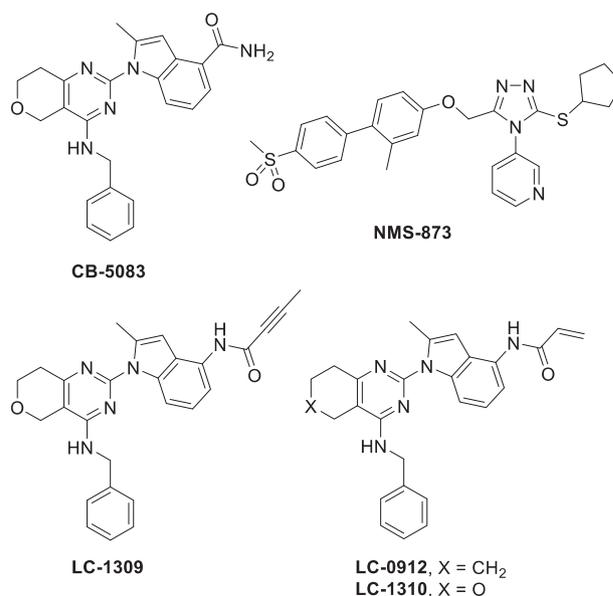


Fig. 1 Selected p97 inhibitors

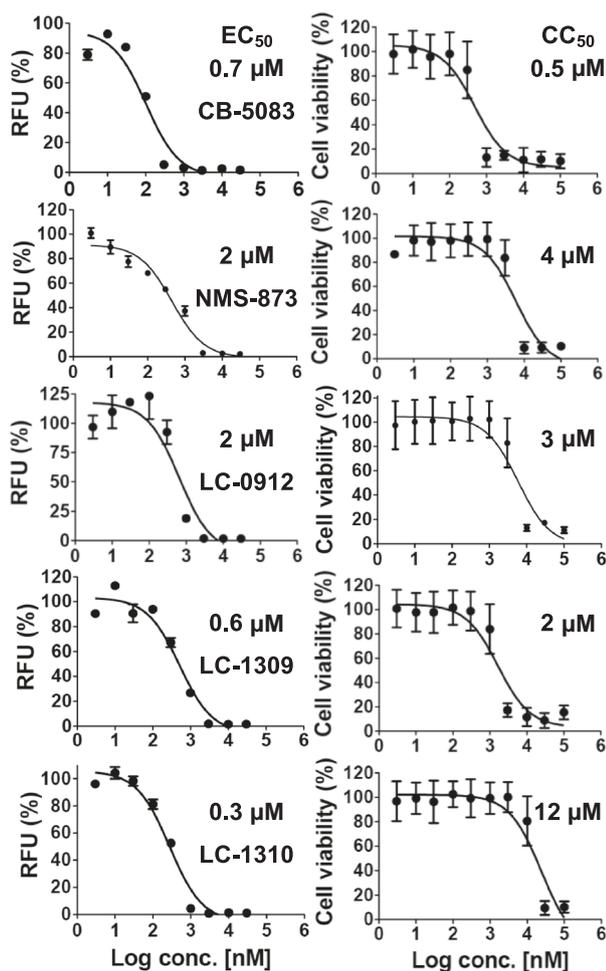


Fig. 2 Dose-response analysis of p97 inhibitors for anti-HCMV activity. Dose-response analysis of p97 inhibitors for anti-HCMV activity. HFFs were inoculated with GFP reporter HCMV and then the compound was added. Six days later, cells were lysed and GFP expression determined. (RFU relative fluorescence units). For cell viability determination, the compound was added to HFFs and 6 days later cell viability measured by an MTS assay. The EC_{50} (left graph) and CC_{50} (right graph) values and curves were determined by comparing mean activity for nine or ten serial dilutions of compound and vehicle (DMSO)-treated cells using GraphPad Prism software. The values listed are the mean of 2–3 dose–response determinations (also listed in Table 1) and representative graphs for each compound are shown

Table 1 HCMV replication assay p97 inhibitors

Compounds	EC_{50} (μ M) \pm s.d. ^a	CC_{50} (μ M) \pm s.d. ^a	TI ^b
NMS-873	2.2 ± 2.5	4.1 ± 3.3	1.9
CB-5083	0.72 ± 1.1	0.53 ± 0.1	0.7
LC-0912	2.1 ± 2.5	3.3 ± 2.3	1.6
LC-1309	0.6 ± 0.2	2 ± 0	3.3
LC-1310	0.3 ± 0	12 ± 10	40

^as.d. standard deviation

^bTissue culture therapeutic index (CC_{50}/EC_{50})

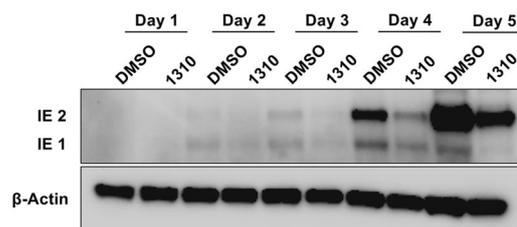


Fig. 3 HCMV IE1 and IE2 protein expression following LC-1310 treatment. Expression of IE1 and IE2 proteins during HCMV infection in the presence or absence of p97 inhibitor LC-1310. Parallel HFF cultures were inoculated, compounds added and cells processed for western blotting each day post-inoculation. The cell lysates were separated by polyacrylamide gel electrophoresis on a 10% gel and then transferred to a PVDF membrane. IE1 and 2 are detected using an anti-HCMV IE1/IE2 monoclonal antibody. The β -actin levels were detected using a rabbit anti- β -Actin monoclonal antibody and are shown as a loading and expression control

mechanism of action for NMS-873 was that the viral immediate-early protein 1 (IE1) expression was not affected by NMS-873 treatment [16]. IE1 and IE2 are expressed from the same viral mRNA by differential splicing and alternative polyadenylation such that both proteins share the first three exons [33]. An NMS-873 effect on IE2 but not IE1 expression may indicate a role for p97 in splicing or polyadenylation. To examine a possible effect of LC-1310 treatment on IE1 and IE2 expression, we treated parallel infected HFFs with LC-1310 for 5 days while processing the cells for western blotting each day. Figure 3 shows that both IE1 and IE2 expression were reduced compared to the DMSO control starting at day 2 and continuing through day 5. Our result with LC-1310 differs slightly from what was published by Lin et al. using NMS-873 in that we observed a reduction in both IE1 and IE2 protein expression upon treatment with p97 inhibitor LC-1310. The simultaneous reduction in IE1 and IE2 expression could explain the increased antiviral activity and a higher therapeutic index of LC-1310 in comparison with NMS-873. Furthermore, a combined IE1/IE2 effect could be specific to LC-1310 and signal a difference in p97 inhibition or off-target effects between LC-1310 and NMS-873. LC-1310 and NMS-873 employ different mechanisms of action as p97 inhibition; NMS-873 binds in a tunnel between D1 and D2 inducing an allosteric inhibition of P97 [8, 9]. In contrast, the irreversible inhibitor LC-1310 was designed based on CB-5083, which targets the D2 ATP binding site [10], such that the extent and duration of p97 inhibition for LC-1310 could result in an enhanced anti-HCMV effect [28]. Also, in our experiments, we did not pretreat the HFFs prior to inoculation because it did not make a difference in NMS-873 HCMV inhibition [16].

Ganciclovir (GCV) is a front-line drug prescribed to treat HCMV infection and we were interested to examine how a combination of LC-1310 and GCV would impact inhibition

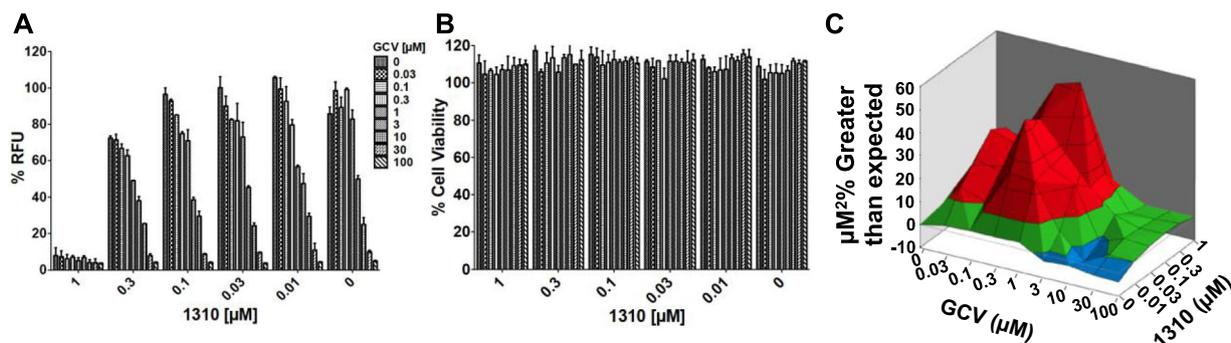


Fig. 4 LC-1310/GCV combination effects. **a** Anti-HCMV activities of the combinations. HFFs were inoculated as described in the HCMV replication assay and compound combinations applied. Six days later, GFP fluorescence was measured as an indication of virus replication and spread. Inhibitory effects of combinations were expressed as a percentage of the DMSO-alone samples. All compounds doses were performed in duplicate and mean value plus standard error depicted. **b** The viability of HFFs following compound combination treatment for 6 days was determined by the MTS viability assay and presented as a percentage of the viability of cells that were treated with DMSO

alone. **c** Three-dimensional surface plots of LC-1310 and ganciclovir combinations and anti-HCMV activities. The three-dimensional plot was generated using MacSynergy II software with the compound concentrations shown on the x and y axes. The horizontal plane represents the predicted surface if the interactions are additive. A peak above the additive plane represents synergy and a negative depression below the plane represents antagonism. Each sample was performed in duplicate and 95% confidence intervals were used to evaluate the data. Blue color = -10 to 0, green = 0–10, and red = 10–60 $\mu\text{M}^2\%$ for a given combination

of HCMV replication in cell culture. HFFs were inoculated with ADCREGFP reporter HCMV and various combinations of six LC-1310 concentrations (0–1 μM) and nine GCV concentrations (0–100 μM) were added to a 96-well plate using a six by nine matrix. After 6 days, the cultures were lysed and GFP fluorescence measured as an indication of HCMV replication and spread. The results in Fig. 4a show a clear antiviral activity at a single dose and combinations of the two compounds compared to DMSO alone and Fig. 4b indicates no significant effects on cell viability at the single dose or combination concentrations. The results were used to investigate possible synergy or antagonism by the Bliss independence model using MacSynergy II software [34]. In general, the three-dimensional surface plot of the anti-HCMV activities for the combinations will show synergy via peaks above a plane representing additive effects and antagonism via depressions below the additive plane [34]. The volumes of synergy and antagonism across all tested combinations are shown in Fig. 4c and result in a score of 261 (95% confidence interval), with anything above 40 considered synergistic. Therefore, the LC-1310 and GCV combination resulted in a synergistic inhibitory effect on HCMV replication.

We next combined LC-1310 and the recently approved HCMV drug Letermovir (LTV) to evaluate co-treatment potential for the p97 inhibitor. HFFs were inoculated with ADCREGFP reporter HCMV and various combinations of six LC-1310 concentrations (0–1 μM) and nine LTV concentrations (0–100 μM) were added to a 96-well plate using a six by nine matrix. After 6 days, the cultures were lysed and GFP fluorescence measured to indicate HCMV replication and spread. There were clear antiviral effects for compounds alone and in combination (Fig. 5a) without significant cellular

toxicity (Fig. 5b). When we examined the results for possible synergistic or antagonistic effects using MacSynergy, the score was 30 and fell within the additive range (-10 to 40, Fig. 5c). Therefore, the LC-1310 and LTV combination resulted in an additive inhibitory effect on HCMV replication.

IAV is an orthomyxovirus. Viral particles are enveloped and contain a segmented single-stranded negative-sense RNA genome. The IAV RNA genome segments are contained within eight ribonucleoproteins that must be transported to the infected cell nucleus to express viral proteins. IAV is another virus where p97 expression is important for virus replication and NMS-873 inhibits virus production [20]. We analyzed our p97 inhibitors for effects on IAV using MDCK cells in a viral cytopathic effects (CPE) assay. We inoculated cells for 2 hours with IAV isolate A/WS/33 (H1N1), retired the inoculum, and added compounds at 0.1, 0.25, and 0.5 μM final concentrations. Twenty-four hours later, the infected and treated cultures were analyzed in parallel to the compound-treated and uninfected cultures for cell viability. An antiviral compound will inhibit virus-induced CPE and promote cell viability. We normalized results for each compound to the uninfected DMSO vehicle control where we will see maximum cell viability and to the infected DMSO control where maximum virus-induced CPE will be observed. Clear antiviral effects and retention of cell viability were seen for the control inhibitor, anti-IAV nucleobase T-1105 (at 12.5 μM), and for NMS-873 (Table 2) in agreement with previously published studies [20, 35]. There was no detectable anti-IAV activity for the p97 inhibitors LC-0912, LC-1309, LC-1310, or CB-5083 (Table 2). It is unclear if the lack of IAV activity for the compounds that bind to the D2 ATP binding site is due to a qualitative or quantitative difference in p97 binding compared to NMS-873 or simply an issue with those compounds

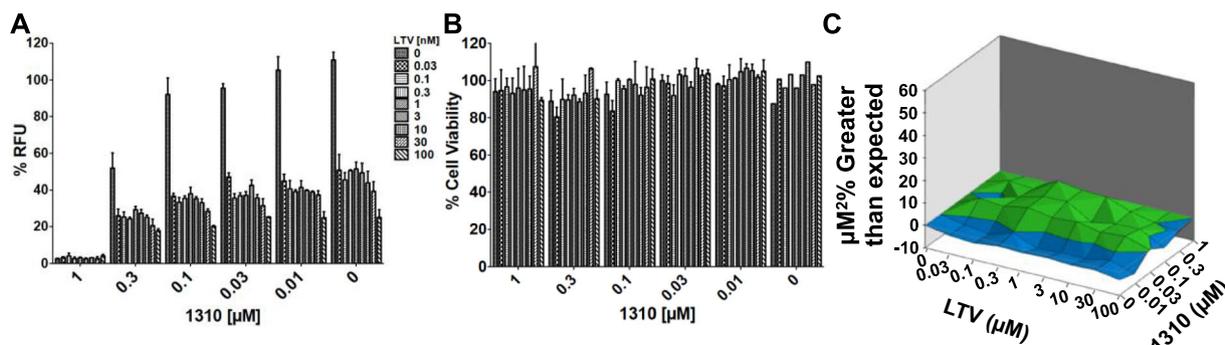


Fig. 5 LC-1310/LTV combination effects. **a** Anti-HCMV activities of the combinations. HFFs were inoculated as described in the HCMV replication assay and compound combinations applied. Six days later, GFP fluorescence was measured as an indication of virus replication and spread. Inhibitory effects of combinations were expressed as a percentage of the DMSO-alone samples. All compounds doses were performed in duplicate and the mean value plus standard error depicted. **b** The viability of HFFs following compound combination

treatment for 6 days was determined by the MTS viability assay and presented as a percentage of the viability of cells that were treated with DMSO alone. **c** Three-dimensional surface plots of LC-1310 and Letemrovir combinations and anti-HCMV activities are shown. The three-dimensional plots were generated and interpreted as described in Fig. 4 legend. Blue color = -10 to 0 and green = 0 – $10 \mu\text{M}^2$ for a given combination

Table 2 Influenza A CPE assay p97 inhibitors

Compound	Compound + Virus ^a			Compound only ^a		
	0.5 μM	0.25 μM	0.1 μM	0.5 μM	0.25 μM	0.1 μM
DMSO	0 \pm 4	0 \pm 1	0 \pm 2	100 \pm 2	100 \pm 3	100 \pm 0.7
T-1105 12.5 μM	100 \pm 3	98 \pm 2	73 \pm 1	91 \pm 5	80 \pm 1	89 \pm 4
NMS-873	80 \pm 5	62 \pm 6	21 \pm 7	94 \pm 2	80 \pm 2	89 \pm 3
CB-5083	0 \pm 1	0 \pm 2	0 \pm 3	89 \pm 4	89 \pm 2	97 \pm 3
LC-0912	0 \pm 5	0 \pm 3	0 \pm 3	103 \pm 1	93 \pm 2	101 \pm 4
LC-1309	0 \pm 5	0 \pm 2	0 \pm 3	102 \pm 2	90 \pm 4	96 \pm 4
LC-1310	0 \pm 5	1 \pm 1	0 \pm 1	101 \pm 1	84 \pm 2	93 \pm 2

^aPercent viable cells \pm standard deviation

binding the D2 ATP site in p97 from canine MDCK cells. In general, there were reduced effects for all compounds on cell viability in the absence of virus possibly due to the use of MDCK cells or the shorter 24-h treatment compared to multiple days for the other assays.

Coronaviruses are enveloped viruses with a large single-stranded plus sense viral genomic RNA (typically 26–32 kb). Upon entry into the cytoplasm, the RNA genome is translated into two large polypeptides that are cleaved by viral proteases into their effector proteins. In addition, genes near the 3' end of the viral genome are expressed via subgenomic viral RNAs. Coronaviruses are significant human pathogens causing outcomes from a common cold-like illness to severe acute respiratory syndrome (SARS) and with the emergence of SARS-CoV-2 there is renewed interest in anti-CoV therapeutics. In addition, siRNA-mediated knockdown of p97 expression resulted in a reduction in infectious bronchitis virus (IBV) CoV replication, indicating the importance of p97 function for IBV and possibly other CoV [15]. We tested the irreversible p97 inhibitors for their effects on human coronavirus OC43 (HCoV-OC43), a betacoronavirus that can

cause a cold-like illness in humans [36]. We inoculated human hepatocellular carcinoma Huh7 cells with HCoV-OC43, retired the inoculum, and incubated the cultures with the compound for 5 days. HCoV-OC43 will cause extensive CPE in Huh-7 cultures by day 5 unless virus replication is inhibited. We included three concentrations of compound, 0.1, 0.25, and 0.5 μM . None of the compounds showed an antiviral effect against HCoV-OC43 (Table 3) whereas the broad-spectrum nucleoside inhibitor remdesivir (RDV) showed 100% protection at 1 μM as expected [37]. It could be that IBV, a gammacoronavirus of chickens, is more susceptible to p97 inhibition in cell culture than the betacoronavirus HCoV-OC43. The p97 inhibitors caused a 10–60% reduction in cell viability at 0.5 and 0.25 μM and NMS-873 showed a clear effect on cell viability additionally at 0.1 μM (Table 3).

Conclusion

The ring ATPase p97 is an attractive target for antiviral therapeutics. The challenge is to develop a p97 inhibitor

Table 3 HCoV-OC43 CPE assay p97 inhibitors

Compound	Compound + Virus ^a			Compound only ^a		
	0.5 μ M	0.25 μ M	0.1 μ M	0.5 μ M	0.25 μ M	0.1 μ M
DMSO	0 \pm 2	0 \pm 7	0 \pm 5	100 \pm 4	100 \pm 3	100 \pm 0.3
RDV 1 μ M	100 \pm 4	100 \pm 6	100 \pm 1	100 \pm 1	104 \pm 2	103 \pm 2
NMS-873	0 \pm 1	0 \pm 1	0 \pm 1	42 \pm 4	54 \pm 2	59 \pm 1
CB-5083	0 \pm 1	0 \pm 2	0 \pm 3	38 \pm 1	59 \pm 3	83 \pm 1
LC-0912	0 \pm 1	0 \pm 3	0 \pm 3	57 \pm 4	90 \pm 3	95 \pm 2
LC-1309	0 \pm 3	0 \pm 2	0 \pm 3	47 \pm 2	78 \pm 2	86 \pm 1
LC-1310	0 \pm 1	0 \pm 6	0 \pm 3	78 \pm 8	89 \pm 2	86 \pm 2

^aPercent viable cells \pm standard deviation

without complicating cellular toxicity. We describe here an irreversible p97 inhibitor, LC-1310, that potently inhibits HCMV replication with a better cellular toxicity profile than other known p97 inhibitors. The activity appears to be selective for HCMV and may differ mechanistically from the inhibition observed for NMS-873.

Methods

Compounds

p97 inhibitors were synthesized as previously described [28]. RDV (CAS# 1809249-37-3) was purchased from MedKoo Biosciences, Inc. T-1105 (3-hydroxypyrazine-2-carboxamide, CAS# 55321-99-8) was purchased from Alfa Aesar. All compounds were stored at -20°C as 20 mM stock solutions in dimethyl sulfoxide (DMSO) shortly before use.

Cells and viruses

The hepatocyte-derived cellular carcinoma cell line Huh-7 [38] was used for HCoV-OC43 studies. Huh-7 cells were maintained in minimum essential medium supplemented with 10% fetal bovine serum (FBS), 100 IU streptomycin/penicillin per ml, 1 mM Sodium pyruvate, 1X non-essential amino acids, and 10 $\mu\text{g}/\text{mL}$ plasmocin (InvivoGen). Human foreskin fibroblasts (HFFs, CRL-2088) and Madin-Darby Canine Kidney cells (CCL-34) were obtained from American Type Tissue Culture (ATCC). HFF and MDCK were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS) (7% for MDCK cells), 100 IU streptomycin/penicillin per ml, and 10 $\mu\text{g}/\text{mL}$ plasmocin (InvivoGen). All the cell lines were maintained at 37°C in a 5% CO_2 incubator. The GFP reporter virus derived from the lab strain HCMV AD169 strain, ADCREGFP was provided by Wade Bresnahan (University of Minnesota) [29]. HCoV-OC43 was provided by Jun Wang (University of Arizona). IAV A/WS/33 (H1N1) was obtained from ATCC (VR-825).

HCMV virus replication assay

This assay measures GFP expression from an HCMV reporter virus, ADCREGFP, and has been described previously [30–32, 39]. Briefly, $\sim 3 \times 10^4$ HFFs/well were plated into 96-well white opaque dishes and inoculated with ADCREGFP the next day at a multiplicity of infection (MOI) of 0.001 in DMEM containing 5% fetal calf serum for 2 h at 37°C and 5% CO_2 . The cells were then washed with phosphate-buffered saline (PBS) and 100 μl of DMEM containing 5% fetal calf serum with test compounds or DMSO was added and maintained at 37°C and 5% CO_2 for 6 days. The infected cells were then lysed for 10 min at 37°C with 200 μl lysis buffer (25 mM Tris [pH 7.8], 2 mM dithiothreitol (DTT), 2 mM *trans*-1,2-diaminocyclohexane-*N,N,N',N'*-tetraacetic acid, 1% Triton X-100, 10% glycerol) followed by a 30 min incubation at room temperature on a shaker [30, 40, 41]. GFP relative fluorescence units were determined at excitation/emission 495/515 nm in a Molecular Devices M5e plate reader. Mean values of triplicate wells were determined and compared to the mean value for the wells that received DMSO alone. The concentration of compound that reduced luciferase activity by 50% was defined as the 50% effective concentration (EC_{50}). The EC_{50} was determined by comparing luciferase activity for nine serial dilutions of the compound and vehicle-treated cells using GraphPad Prism software.

For the compound combination studies, cells inoculated as above were incubated with compounds alone and in various combinations in a six by nine matrix in 96-well plates. The infected and compound-treated cells were processed as above. The data were analyzed using the MacSynergy II software [34]. The synergy values were obtained at the 95% confidence interval.

Cell viability assay

HFFs were plated into 96-well plates as indicated in HCMV replication assay and the next day incubated in different compound doses at 37°C for 6 days. The

3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium (MTS)-based tetrazolium reduction CellTiter 96 Aqueous Non-Radioactive cell proliferation assay (Promega G5430) was used as per manufacturer's instructions to determine cell viability. Each compound dose was tested in triplicate and the mean values of the triplicates were compared to the mean value for the wells treated with DMSO alone. Data from ten serial dilutions of the compound were used to calculate the CC_{50} using GraphPad Prism software. The CC_{50} value was the concentration of the compound resulting in a 50% viability reduction compared to DMSO.

Western blotting

HFFs were grown in 12-well tissue culture plates at $\sim 3.6 \times 10^5$ cells per well. Cells were incubated with the compound before virus inoculation. After 24 h incubation, cells were inoculated with ADCREGFP virus for 2 hours at an MOI of 0.1 in DMEM containing 5% fetal bovine serum. The inoculated cells were washed with phosphate-buffered saline and incubated with DMEM containing 5% fetal bovine serum with test compounds or DMSO at 37 °C and 5% CO_2 for 120 h. The infected cells were lysed with RIPA buffer (0.1% SDS, 1% Triton X-100, 1% deoxycholate, 5 mM EDTA, 150 mM NaCl, and 10 mM Tris at pH 7.2). Twenty micrograms of total lysate protein were separated by electrophoresis on a 10% SDS-polyacrylamide gel and transferred to a PVDF membrane (Millipore). Primary antibodies were mouse anti-CMV IE1/2 monoclonal antibody (MAB8131, Millipore) and rabbit anti- β -Actin monoclonal antibody (4970T, Cell Signaling). Blots were probed with primary antibody (1:2000) diluted in 5% BSA in Tris Buffered Saline (TBS) and subsequently the HRP-conjugated secondary antibodies (Anti-rabbit IgG, HRP-linked Antibody from Cell Signaling 7074S; Mouse IgG HRP Linked Whole Antibody from GE Healthcare NXA931) at 1:10,000. Blots were washed in TBS three times, incubated with chemiluminescent substrate (SuperSignal West Pico; Thermo Scientific) according to the manufacturer's protocol, and exposed in ChemiDoc MP (Bio-Rad) for visualization of bands.

Human betacoronavirus OC43 cytopathic effects assay

Huh-7 cells (1.25×10^4 per well) were seeded in a regular 96-well plate and the next day the cells were inoculated with HCoV-OC43 at an MOI of 0.5 for 2 hours at 37 °C 5% CO_2 . The inoculum was subsequently retired and the cells were treated with compounds or RDV 1 μ M as a positive control. Concentrations of 0.1, 0.25, and 0.5 μ M were tested for all the p97-targeting compounds. The percent viable

cells in absence of infection were analyzed in parallel cultures. Cells were incubated for 5 days after which the infected and treated cells were analyzed for cell viability using the MTS-based tetrazolium reduction CellTiter 96 Aqueous Non-Radioactive cell proliferation assay and absorbance was measured at 490 nm. The readings in infected cells were normalized to DMSO-treated cells ($\sim 0\%$ viability) and RDV-treated cells ($\sim 100\%$ viability) via GraphPad prism. Readings for uninfected cells were normalized to DMSO control ($\sim 100\%$ viability).

Influenza A virus cytopathic effects assay

MDCK cells (3×10^4 per well) were seeded in a 96-well plate. The next day the cells were inoculated with IAV A/WS/33 at an MOI 0.02 for 2 hours in an IAV-infection medium (DMEM supplemented with 0.2% BSA, 0.5 μ g/mL of TPCK-treated trypsin, 1X pen-strep, 1X glutamax, 10 mM Hepes). After 2 hours, the inoculum was retired and compounds added in an IAV-infection medium for 24 h. In parallel, cell viability in absence of infection was analyzed. Cell viability was determined using the MTS-based tetrazolium reduction CellTiter 96 Aqueous Non-Radioactive cell proliferation assay after the 24-h incubation and absorbance was measured at 490 nm. The readings in infected cells were normalized to DMSO-treated cells ($\sim 0\%$ viability) and uninfected DMSO-treated cells ($\sim 100\%$ viability) via GraphPad prism. Readings for uninfected cells were normalized to DMSO control ($\sim 100\%$ viability).

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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