

Plural assay systems derived from different cell lines and hepatitis C virus strains are required for the objective evaluation of anti-hepatitis C virus reagents

Youki Ueda, Kyoko Mori, Yasuo Ariumi, Masanori Ikeda, and Nobuyuki Kato\*

Department of Tumor Virology, Okayama University Graduate School of Medicine,  
Dentistry, and Pharmaceutical Sciences, 2-5-1 Shikata-cho, Okayama 700-8558, Japan

\*Corresponding author. Fax: +81 86 235 7392.

*E-mail address:* [nkato@md.okayama-u.ac.jp](mailto:nkato@md.okayama-u.ac.jp) (N. Kato)

3,706 words including refs and figure legends

## Abstract

Persistent hepatitis C virus (HCV) infection causes chronic liver diseases and is a global health problem. HuH-7 hepatoma-derived cells are widely used as the only cell-based HCV replication system for HCV research, including drug assays. Recently, using different hepatoma Li23-derived cells, we developed an HCV drug assay system (ORL8), in which the genome-length HCV RNA (O strain of genotype 1b) encoding renilla luciferase replicates efficiently. In this study, using the HuH-7-derived OR6 assay system that we developed previously and the ORL8 assay system, we evaluated 26 anti-HCV reagents, which other groups had reported as anti-HCV candidates using HuH-7-derived assay systems other than OR6. The results revealed that more than half of the reagents showed different anti-HCV activities from those in the previous studies, and that anti-HCV activities evaluated by OR6 and ORL8 assays were also frequently different. In further evaluation using the HuH-7-derived AH1R assay system, which was developed using the AH1 strain of genotype 1b, several reagents showed different anti-HCV activities in comparison with those evaluated by OR6 and ORL8 assays. These results suggest that the different activities of anti-HCV reagents are caused by the

differences in cell lines or HCV strains used for the development of assay systems.

Therefore, we conclude that plural HCV assay systems developed using different cell lines or HCV strains are required for the objective evaluation of anti-HCV reagents.

*Keywords:* HCV; HCV RNA replication system; Li23 cells; Reporter assay for anti-HCV reagents

## 1. Introduction

Hepatitis C virus (HCV) infection frequently causes chronic hepatitis, which often leads to liver cirrhosis and hepatocellular carcinoma. Since approximately 170 million people are infected with HCV worldwide, HCV infection is a serious global health problem [1]. Although the combination of pegylated-interferon (PEG-IFN) and ribavirin is the standard therapy worldwide, only half of the patients receiving this treatment exhibit a sustained virological response [2]. HCV is an enveloped virus with a positive single-stranded RNA virus of the *Flaviviridae* family. The HCV genome encodes a large polyprotein precursor of approximately 3,000 amino acids, which is cleaved into 10 proteins in the following order: Core, envelope 1 (E1), E2, p7, non-structural 2 (NS2), NS3, NS4A, NS4B, NS5A, and NS5B [3, 4].

To date, HuH-7 hepatoma-derived cells are used as the only cell culture system for robust HCV replication in HCV research, including drug assays. We have also developed a HuH-7-derived drug assay system (OR6), in which genome-length HCV RNA (O strain of genotype 1b derived from an HCV-positive blood donor) encoding renilla luciferase (RL) efficiently replicates [5]. Recently, we found a new human

hepatoma cell line, Li23, that enables robust HCV RNA replication [6], and we showed that the gene expression profile of Li23 cells was distinct from that of HuH-7 cells, although both cell lines had similar liver-specific expression profiles [7]. In that study, we identified three genes (New York esophageal squamous cell carcinoma 1,  $\beta$ -defensin-1, and galectin-3) showing Li23-specific expression profiles by a comparative analysis using several other hepatic cell lines [7]. We further developed Li23-derived drug assay systems (ORL8 and ORL11), which are relevant to the HuH-7-derived OR6 assay system [6]. During the process of evaluating ORL8 and ORL11 assay systems using anti-HCV reagents such as IFNs, we noticed that these assay systems were frequently more sensitive to anti-HCV reagents than the OR6 assay system [6]. Furthermore, we recently found that ribavirin at clinically achievable concentrations (approximately 10  $\mu$ M) effectively inhibited HCV RNA replication in both ORL8 and ORL11 assay systems, but not in the OR6 assay system [8]. This finding led to the clarification of the anti-HCV mechanism of ribavirin, and we demonstrated that ribavirin's anti-HCV activity was mediated by the inhibition of inosine monophosphate dehydrogenase, a key enzyme in the guanosine biosynthetic

pathway [8]. From these findings, we supposed that the anti-HCV reagents reported to date might show different activities among the different drug assay systems. To test this assumption, we evaluated 22 anti-HCV reagents that were reported using HuH-7-derived assay systems other than OR6, using OR6 and ORL8 assay systems. Four additional reagents predicted by antiviral activity other than HCV were also evaluated. Furthermore, a recently developed HuH-7-derived AH1R assay system (AH1 strain of genotype 1b derived from a patient with acute hepatitis) (Mori et al., in preparation) was also used for the evaluation. Here, we report that plural assay systems derived from different cell lines and different HCV strains are required for the objective evaluation of anti-HCV reagents.

## **2. Materials and methods**

### *2.1. Cell cultures*

HuH-7-derived OR6 and AH1R cells were maintained in medium containing G418 (0.3 mg/ml) as described previously [5]. Li23-derived ORL8 cells were also maintained in medium containing G418 (0.3 mg/ml) as described previously [6].

## 2.2. Reagents

Acetylsalicylic acid, cephalotaxine, clemizole, crucumin, isoliquiritigenin, nitazoxanide, and tizoxanide were purchased from Sigma-Aldrich (St. Louis, MO). Cantharidin, 2'-deoxy-5-fluorouridine, griseofulvin, guanazole, homoharringtonine, resveratrol, and Y7632 were purchased from WAKO Pure Chemical Industries, Ltd. (Osaka, Japan). Artemisinin and bisindolyl maleimide 1 were purchased from Alexis Biochemicals (San Diego, CA). Artesunate and silibinin A were purchased from Lkt Laboratories (St. Paul, MN). Esomeprazole and nelfinavir were purchased from Toronto Research Chemicals (North York, ON, Canada). Cinanserin hydrochloride and HA1077 were purchased from Tocris Bioscience (Bristol, UK). 6-Azauridine was purchased from MP Biomedicals (Solon, OH). Carvedilol was purchased from Calbiochem (San Diego, CA). Hemin was purchased from Alfa Aesar (Ward Hill, MA). Methotrexate was purchased from Tokyo Chemical Industry (Tokyo, Japan). Cinanserin hydrochloride, guanazole, HA11077, and Y27632 were dissolved in the culture medium for Li23-derived cells. Artesunate was dissolved in 0.5% NaHCO<sub>3</sub> solution. Other reagents

were dissolved in dimethyl sulfoxide.

### *2.3. RL assay*

RL assay was performed as described previously [6]. Briefly, the cells were plated onto 24-well plates ( $2 \times 10^4$  cells per well) in triplicate and then treated with each reagent at several concentrations for 72 h. After treatment, the cells were subjected to luciferase assay using the RL assay system (Promega, Madison, WI). From the assay results, the 50% effective concentration ( $EC_{50}$ ) of each reagent was determined.

### *2.4. WST-1 cell proliferation assay*

The cells were plated onto 96-well plates ( $1 \times 10^3$  cells per well) in triplicate and then treated with each reagent at several concentrations for 72 h. After treatment, the cells were subjected to the WST-1 cell proliferation assay (Takara Bio, Otsu, Japan) according to the manufacturer's protocol. From the assay results, the 50% cytotoxic concentration ( $CC_{50}$ ) of each reagent was determined.



### *2.5. Western blot analysis*

The preparation of cell lysates, sodium dodecyl sulfate-polyacrylamide gel electrophoresis, and immunoblotting analysis were performed as previously described [9]. The antibodies used in this study were those against HCV Core (CP11; Institute of Immunology, Tokyo, Japan) and  $\beta$ -actin (AC-15, Sigma-Aldrich) as the control for the amount of protein loaded per lane. Immunocomplexes were detected with the Renaissance enhanced chemiluminescence assay (Perkin-Elmer Life Sciences, Boston, MA).

### *2.6. Selective index (SI)*

The SI value of each reagent was determined by dividing the  $CC_{50}$  value by the  $EC_{50}$  value.

## **3. Results**

*3.1. Evaluation of 26 reagents for anti-HCV activity using OR6 and ORL8 assay systems*

To obtain candidates for the evaluation of anti-HCV activity using OR6 and ORL8 assay systems, we first searched the literature in the PubMed database using the key words (hcv or hepatitis C) and (inhibit or antiviral or suppress or block); this yielded approximately 4,500 reports published between January 2006 and April 2010. From these results, we further selected the reports in which the  $EC_{50}$  values of reagents were determined or estimated by the HuH-7-derived HCV assay systems using the Con-1 strain (genotype 1b) [10], N strain (genotype 1b) [11], or HCV JFH-1 strain (genotype 2a) [12]. We finally chose 22 commercially available reagents for the evaluation of anti-HCV activity using OR6 and ORL8 assay systems. Four reagents predicted from the antiviral activity (hepatitis B virus, cytomegalovirus, etc.) other than HCV were also included in the evaluation study. The 26 selected reagents and their references are listed in Supplementary Table S1.

For each of the 26 reagents, we determined the  $EC_{50}$  value by RL assay and the  $CC_{50}$  value by WST-1 assay using the OR6 or ORL8 assay system, and calculated the SI value by dividing the  $CC_{50}$  value by the  $EC_{50}$  value. For each reagent, we first compared the  $EC_{50}$  value obtained from the OR6 or ORL8 assay with that of the previous study.

Consequently, we classified the 26 reagents into five classes, A to E (Table 1). Eight reagents (methotrexate, artemisinin, artesunate, clemizole, hemin, 6-azauridine, acetylsalicylic acid, and isoliquiritigenin with the order of the SI value in the ORL8 assay) belonged to class A, in which the  $EC_{50}$  value obtained by either the OR6 or ORL8 assay was less than one third of that in the previous study (Supplementary Table S1 and Table 1). Artesunate, an artemisinin-derivative possessing antiviral activity against cytomegalovirus, herpesvirus, Epstein-Barr virus etc., was included in class A by the comparison with the data on anti-cytomegalovirus activity. In this class, we especially noticed that methotrexate (an anticancer drug) showed very strong anti-HCV activity ( $EC_{50}$  0.1  $\mu$ M;  $CC_{50}$  >200  $\mu$ M; SI >2000) in the ORL8 assay (upper panel in Fig. 1A and Table 1), whereas methotrexate showed very weak anti-HCV activity ( $EC_{50}$  >200  $\mu$ M;  $CC_{50}$  >200  $\mu$ M) in the OR6 assay as well as in a previous report [13] (upper panel in Fig. 1A and Table 1). This drastic difference was confirmed by Western blot analysis (lower panels in Fig. 1A). These results indicate that only the ORL8 assay is drastically sensitive to methotrexate, and suggest that the anti-HCV activity of methotrexate depends on the types of hepatic cells. The comparison of the  $EC_{50}$  values

of other reagents belonging to class A revealed that the ORL8 assay was more sensitive than the OR6 assay (1.9 ~ 15 fold) to artemisinin, artesunate, clemizole, acetylsalicylic acid, and 6-azauridine, and conversely the OR6 assay was more sensitive than the ORL8 assay (2 ~ 2.5 fold) to hemin and isoliquiritigenin (Table 1). Furthermore, the  $CC_{50}$  values of clemizole and 6-azauridine also differed more than twofold between the OR6 and OR8 assays (Table 1). These results suggest that the anti-HCV activities of these reagents are affected by the kind of assay systems used. Especially, we noticed that artemisinin and artesunate (antimalarial drugs) showed higher SI values in the ORL8 assay than previously reported [14, 15]. The anti-HCV profiles of artemisinin and artesunate in the OR6 and ORL8 assays are shown in Fig. 1B and Supplementary Fig. 1A, respectively. In addition, the comparison of SI values revealed that the OR6 assay was more sensitive to hemin and isoliquiritigenin than the HuH-7-derived assays (Con-1 and N strains) used in the previous reports (Supplementary Table S1), suggesting that the HCV strains used in the assay systems affect the evaluation of anti-HCV reagents.

Nine reagents (nelfinavir, 2'-deoxy-5-fluorouridine, resveratrol, cantharidin, homoharringtonine, curcumin, griseofulvin, cinanserin hydrochloride, and

cephalotaxine with the order of SI value in the ORL8 assay) were placed in class B, in which the  $EC_{50}$  values obtained by OR6 and ORL8 assays were similar (more than one-third to less than threefold) to those in the previous study (Table 1 and Supplementary Table S1). Cantharidin, homoharringtonine, and cephalotaxine, all of which possess anti-hepatitis B virus activity, were placed in class B by the comparison with the data on anti-hepatitis B virus activity (Supplementary Fig. 1).

Tizoxanide and nitazoxanide belonged to class C, in which the  $EC_{50}$  values obtained by both the OR6 and ORL8 assays were more than three-fold higher than in the previous study (Table 1 and Supplementary Table S1). Guanazole and HA1077 were placed in class D, in which there was no anti-HCV activity in both the OR6 and ORL8 assays (Table 1). No anti-HCV activity of guanazole and HA1077 was also confirmed by Western blot analysis (data not shown). Lastly, five reagents (Bisindoly maleimide 1, esomeprazole, Y27632, carvedilol, and silibinin A) were placed in class E, in which pro-HCV activity was exhibited in both OR6 and ORL8 assays. We unexpectedly observed that these reagents enhanced the HCV RNA replication level. As a representative reagent, pro-HCV profiles of silibinin A are shown in the upper panel of

Fig. 1C. These pro-HCV profiles were confirmed by Western blot analysis (lower panels in Fig. 1C for silibinin A and data not shown for the other reagents). Since the anti-HCV activity of silibinin A was detected by the HCV replicon assay system using the Con-1 strain [14], the converse effects obtained by our assay systems using the O strain may be due to the difference in HCV strains. In summary, the differences in anti-HCV activities observed among HuH-7- and Li23-derived assay systems used in this study and the other HuH-7-derived assay systems used in the previous studies suggest that the activities of anti-HCV reagents differ depending on which HCV strains and cell lines are used in the evaluation assays.

### *3.2. Evaluation of 18 reagents for anti-HCV activity using AH1R assay system*

We previously established a HuH-7-derived cell line (AH1), which harbors genome-length HCV RNA (AH1 strain of genotype 1b) derived from a patient with acute hepatitis [16]. To further examine the effect of the HCV strain on anti-HCV reagent activity, we developed an AH1R assay system that is based on the AH1 cell line and that corresponds to the OR6 assay system (Mori et al., in preparation).

Using the AH1R assay system, we further evaluated the anti-HCV activities of 18 reagents, which showed differential anti-HCV activity between the OR6 and ORL8 assays, or showed either no anti-HCV activity or pro-HCV activity in both the OR6 and ORL8 assays. The results of the evaluation are shown in Table 1. The comparisons of the data obtained by OR6 and AH1R assays revealed that the difference in the  $EC_{50}$  value from reagent to reagent was held within the range of one-third to threefold. However, we noticed that the  $EC_{50}$  value (5.3  $\mu$ M) of artemisinin in the AH1R assay was remarkably lower than that (81  $\mu$ M) in the OR6 assay (Supplementary Fig. 2 and Table 1), suggesting that artemisinin's anti-HCV activity differs depending on the HCV strain. Furthermore, the results of the AH1R assay revealed that cephalotaxine, belonging to class B, would be recategorized into class D. In summary, some reagents showed differential anti-HCV activities between the HuH-7-derived OR6 (O strain) and AH1R (AH1 strain) assay systems, although most of the reagents showed similar levels of anti-HCV activity in both assays. Taking together the results of the previous and present studies, we conclude that plural assay systems derived from different cell lines and HCV strains are needed for the objective evaluation of anti-HCV reagents.

#### **4. Discussion**

In the present study, we demonstrated for the first time that a Li23-cell-derived drug assay system, not a HuH-7-derived system, was important to use for the objective evaluation of anti-HCV reagents. In addition, we demonstrated that assay systems derived from different HCV strains were also necessary for the objective evaluation of anti-HCV reagents.

Among the 28 reagents evaluated by our assay systems, methotrexate showed the most drastic differences between the HuH-7- and Li23-derived assay systems in terms of anti-HCV activity. Although methotrexate showed very weak anti-HCV activity in the HuH-7-derived assay (Con-1 strain) used in a previous study [13] as well as in our OR6 and AH1R assays (O and AH1 strains), the ORL8 assay revealed very strong anti-HCV activity (SI >2000). Such drastic differences in both assays suggest that some host factor or factors required for HCV RNA replication are different between these two cell lines, although the anti-HCV target of methotrexate is unclear. Since methotrexate is currently used as an anti-cancer drug or anti-rheumatic drug and its EC<sub>50</sub> value for



HCV RNA replication is 0.1  $\mu\text{M}$ , it may be a potential candidate for enhancing the effects of the current combination therapy of PEG-IFN and ribavirin.

The anti-HCV activities of two antimalarial drugs, artemisinin and its derivative artesunate, are interesting. Although Paeshuyse et al. [14] showed that artemisinin possessed weak or moderate anti-HCV activity using a HuH-7- or HuH-6-derived subgenomic HCV replicon system, artemisinin's anti-HCV mechanism was unclear. On the other hand, Efferth et al. [15] reported that artesunate, the most studied artemisinin-derivative for the treatment of severe malaria, possessed antiviral activity against Epstein-Barr virus, human cytomegalovirus, human herpesvirus 6A, herpes simplex virus 1, and so on, except for HCV with the low micromolar range, although artesunate's precise antiviral mechanism was ambiguous. Therefore, we supposed, and our assay systems clearly detected, that both artemisinin and artesunate possess anti-HCV activity. Especially, the AH1R assay was the most sensitive to artemisinin ( $\text{EC}_{50}$  5.3  $\mu\text{M}$ ), and the ORL8 assay was the most sensitive to artesunate ( $\text{EC}_{50}$  0.22  $\mu\text{M}$ ). Preliminary experiments for the anti-HCV mechanisms of these reagents showed that they did not activate the IFN-signaling pathway (data not shown), and that they did

not induce the oxidative stress (data not shown) as observed in the treatment with a broad range of anti-HCV reagents, including cyclosporine A [8, 17]. Further studies are needed to clarify the anti-HCV mechanisms of these reagents. Since the largest SI value of artemisinin was 58 in the AH1R assay and that of artesunate was 16 in the ORL8 assay, these reagents may be also useful for the treatment of patients with chronic hepatitis.

In this study, we demonstrated that many anti-HCV reagents showed differential anti-HCV activities among different assay systems (OR6, ORL8, and AH1R) on HCV RNA replication. These results suggest that reliance on only a single assay system may lead to an incorrect evaluation of anti-HCV candidates. Therefore, we propose that plural assay systems derived from different cell lines and HCV strains should be used in order to evaluate anti-HCV candidates. Furthermore, plural assay systems derived from at least two different cell origins would be also useful for the screening of anti-HCV candidates.

## **Acknowledgments**

We thank Yusuke Wataya and Hye-Sook Kim for their helpful discussions. This work was supported by grants-in-aid for research on hepatitis from the Ministry of Health, Labor, and Welfare of Japan. K. M. was supported by a Research Fellowship for Young Scientists from the Japan Society for the Promotion of Science.

## **References**

- [1] D. L. Thomas, Hepatitis C epidemiology, *Curr. Top. Microbiol. Immunol.* 242 (2000) 25-41.
- [2] S. Chevaliez, J. M. Pawlotsky, Interferon-based therapy of hepatitis C. *Adv. Drug. Deliv. Rev.* 59 (2007) 1222-1241.
- [3] N. Kato, M. Hijikata, Y. Ootsuyama, et al., Molecular cloning of the human hepatitis C virus genome from Japanese patients with non-A, non-B hepatitis, *Proc. Natl. Acad. Sci. USA* 87 (1990) 9524-9528.
- [4] N. Kato, Molecular virology of hepatitis C virus, *Acta Med. Okayama* 55 (2001) 133-159.

- [5] M. Ikeda, K. Abe, H. Dansako, et al., Efficient replication of a full-length hepatitis C virus genome, strain O, in cell culture, and development of a luciferase reporter system, *Biochem. Biophys. Res. Commun.* 329 (2005) 1350-1359.
- [6] N. Kato, K. Mori, K. Abe, et al., Efficient replication systems for hepatitis C virus using a new human hepatoma cell line. *Virus Res.* 146 (2009) 41-50.
- [7] K. Mori, M. Ikeda, Y. Ariumi, N. Kato, Gene expression profile of Li23, a new human hepatoma cell line that enables robust hepatitis C virus replication: Comparison with HuH-7 and other hepatic cell lines. *Hepatol. Res.* 40 (2010) 1248-1253.
- [8] K. Mori, M. Ikeda, Y. Ariumi, et al., Mechanism of action of ribavirin in a novel hepatitis C virus replication cell system. *Virus Res.* 157 (2011) 61-70.
- [9] N. Kato, K. Sugiyama, K. Namba, et al., Establishment of a hepatitis C virus subgenomic replicon derived from human hepatocytes infected in vitro, *Biochem. Biophys. Res. Commun.* 306 (2003) 756-766.
- [10] V. Lohmann, F. Korner, J. Koch, et al., Replication of subgenomic hepatitis C virus RNAs in a hepatoma cell line, *Science* 285 (1999) 110-113.

- [11] M. Ikeda, M. Yi, K. Li, S. M. Lemon, Selectable subgenomic and genome-length dicistronic RNAs derived from an infectious molecular clone of the HCV-N strain of hepatitis C virus replicate efficiently in cultured Huh7 cells, *J. Virol.* 76 (2002) 2997-3006.
- [12] T. Wakita, T. Pietschmann, T. Kato, et al., Production of infectious hepatitis C virus in tissue culture from a cloned viral genome, *Nat. Med.* 11 (2005) 791-796.
- [13] L. J. Stuyver, T. R. McBrayer, P. M. Tharnish, et al., Dynamics of subgenomic hepatitis C virus replicon RNA levels in Huh-7 cells after exposure to nucleoside antimetabolites, *J. Virol.* 77 (2003) 10689-10694.
- [14] J. Paeshuyse, L. Coelmont, I. Vliegen, et al., Hemin potentiates the anti-hepatitis C virus activity of the antimalarial drug artemisinin, *Biochem. Biophys. Res. Commun.* 348 (2006) 139-144.
- [15] T. Efferth, M. R. Romero, D. G. Wolf, et al., The antiviral activities of artemisinin and artesunate, *Clin. Infect. Dis.* 47 (2008) 804-811.
- [16] K. Mori, K. Abe, H. Dansako, et al., New efficient replication system with hepatitis C virus genome derived from a patient with acute hepatitis C, *Biochem. Biophys.*

Res. Commun. 371 (2008) 104-109.

[17] M. Yano, M. Ikeda, K. Abe, et al., Comprehensive analysis of the effects of ordinary nutrients on hepatitis C virus RNA replication in cell culture, *Antimicrob. Agents Chemother.* 51 (2007) 2016-2027.

### Figure legends

**Fig. 1.** Anti-HCV profiles of representative reagents in OR6 and ORL8 assay systems.

(A) Methotrexate sensitivities on genome-length HCV RNA replication in OR6 and ORL8 assay systems. OR6 and ORL8 cells were treated with methotrexate for 72 h, followed by RL assay (black circle in the upper panel) and WST-1 assay (open triangle in the upper panel). The relative value (%) calculated at each point, when the level in nontreated cells was assigned to 100%, is presented here. Western blot analysis of the treated cells for the HCV Core was also performed (lower panel). (B) Artemisinin sensitivities on genome-length HCV RNA replication in OR6 and ORL8 assay systems. RL assay, WST-1 assay, and Western blot analysis were performed as described in (A).

(C) Silibinin A sensitivities on genome-length HCV RNA replication in OR6 and ORL8 assay systems. RL assay, WST-1 assay, and Western blot analysis were performed as described in (A).

Table  
[Click here to download high resolution image](#)

**Table 1**  
 Anti-HCV activities of 26 reagents evaluated in this study

Assay		*1		OR6		ORL8		AHIR	
Cell origin		HuH-7		HuH-7		Li23		HuH-7	
HCV strain		Con-1, N, JFH-1 etc.		O		O		AH1	
Class	Reagent	$\frac{CC_{50}}{EC_{50}}$	SI	$\frac{CC_{50}}{EC_{50}}$	SI	$\frac{CC_{50}}{EC_{50}}$	SI	$\frac{CC_{50}}{EC_{50}}$	SI
A	Methotrexate	$\frac{>100}{>100}$	-	$\frac{>200}{>200}$	-	$\frac{>200}{0.1}$	>2000	$\frac{170}{>200}$	<0.9
A	Artemisinin	$\frac{>177}{78}$	>2.3	$\frac{380}{81}$	4.7	$\frac{370}{23}$	16	$\frac{310}{5.3}$	58
A	Artesunate*2	$\frac{>15}{3.9}$	>3.8	$\frac{6.1}{2.3}$	2.7	$\frac{3.4}{0.22}$	15	$\frac{4}{0.81}$	4.9
A	Clemizole	$\frac{>20}{8}$	>2.5	$\frac{11}{22}$	0.5	$\frac{22}{2.0}$	11	$\frac{7.3}{>25}$	<0.3
A	Hemin	$\frac{>52}{22}$	>2.4	$\frac{10}{1.2}$	8.3	$\frac{18}{2.4}$	7.5	$\frac{7.2}{1.1}$	6.5
A	6-Azauridine	$\frac{>100}{100}$	>1.0	$\frac{10}{5.7}$	1.8	$\frac{1.5}{0.37}$	4.1	$\frac{14}{3.3}$	4.2
A	Acetylsalicylic acid	$\frac{8^{**}}{4^{**}}$	2.0	$\frac{2.6^{**}}{1.6^{**}}$	1.6	$\frac{2.4^{**}}{0.83^{**}}$	2.9	ND	-
A	Isoliquiritigenin	$\frac{<24}{24}$	<1.0	$\frac{12}{3.9}$	3.1	$\frac{15}{9.8}$	1.5	ND	-
B	Nelfinavir	$\frac{>10}{9.9}$	>1.0	$\frac{26}{11}$	2.4	$\frac{68}{12}$	5.7	ND	-
B	2'-Deoxy-5-fluorouridine	$\frac{<15}{15}$	<1.0	$\frac{31}{32}$	1.0	$\frac{36}{14}$	2.6	$\frac{13}{86}$	0.2
B	Resveratrol	$\frac{>10}{10}$	>1.0	$\frac{35}{4.3}$	8.1	$\frac{42}{16}$	2.6	$\frac{76}{9.9}$	7.7
B	Cantharidin*3	$\frac{3.5}{0.3}$	12	$\frac{1.5}{0.28}$	5.4	$\frac{1.8}{0.69}$	2.6	ND	-
B	Homoharringtonine*3	$\frac{0.5}{30^{**}}$	17	$\frac{38^{**}}{18^{**}}$	2.1	$\frac{0.11}{45^{**}}$	2.4	$\frac{22^{**}}{19^{**}}$	1.2
B	Curcumin	$\frac{>15}{15}$	>1.0	$\frac{18}{14}$	1.3	$\frac{19}{11}$	1.7	ND	-
B	Griseofulvin	$\frac{207}{6.1}$	34	$\frac{16}{4.4}$	3.6	$\frac{14}{8.6}$	1.6	ND	-
B	Cinanserin hydrochloride	$\frac{>10}{>10}$	-	$\frac{33}{25}$	1.3	$\frac{39}{35}$	1.1	ND	-
B	Cephalotaxine*1	$\frac{>100}{60}$	>1.7	$\frac{35}{29}$	1.2	$\frac{38}{47}$	0.8	$\frac{4.8}{41}$	0.1
C	Tizoxanide	$\frac{15}{0.15}$	100	$\frac{11}{2.4}$	4.6	$\frac{24}{9.6}$	2.5	ND	-
C	Nitazoxanide	$\frac{38}{0.21}$	181	$\frac{11}{2.8}$	3.9	$\frac{17}{9.2}$	1.8	$\frac{7.2}{2.2}$	3.3
D	Guanazole	$\frac{<100}{>100}$	<1.0	$\frac{200}{>200}$	<1.0	$\frac{170}{>200}$	<0.9	$\frac{173}{>200}$	<0.9
D	HA1077	$\frac{50}{15}$	3.3	$\frac{>50}{>50}$	-	$\frac{>50}{>50}$	-	$\frac{>50}{>50}$	-
E	Bisindolyl maleimide 1	$\frac{ND}{5}$	-	$\frac{8.1}{6.2}$	1.3	$\frac{15}{15}$	1.0	$\frac{14}{9.1}$	1.5
E	Esomeprazole	$\frac{ND}{>10}$	-	$\frac{67}{67}$	1.0	$\frac{27}{27}$	1.0	$\frac{20}{25}$	0.8
E	Y27632	$\frac{>50}{50}$	>1.0	$\frac{>80}{>80}$	-	$\frac{>80}{>80}$	-	$\frac{39}{>80}$	<0.5
E	Carvedilol	$\frac{17}{4.5}$	3.8	$\frac{4.4}{3.7}$	1.2	$\frac{6.6}{8.8}$	0.8	$\frac{6.3}{6.2}$	1.0
E	Silibinin A	$\frac{ND}{23}$	-	$\frac{12}{85}$	0.1	$\frac{26}{89}$	0.3	$\frac{28}{96}$	0.3

\*1 Assay used in previous reports. \*2 Reported as anti-cytomegalovirus reagent. \*3 Reported as anti-hepatitis B virus reagent. EC<sub>50</sub> and CC<sub>50</sub> values are indicated by the order of μM except \*4 (mM) and \*5 (nM). ND, not determined



Figure  
[Click here to download high resolution image](#)

