

# Carvedilol: a new candidate for reversal of MDR1/P-glycoprotein-mediated multidrug resistance

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In 1983, carvedilol [1-[carbazolyl-(4)-oxy]-3-[(2-methoxyphenoxyethyl)amino]-2-propanol] was designed and developed as a  $\beta$ -adrenoceptor antagonist with vasodilating activity for efficacious and safe treatment of hypertension and coronary artery disease. Carvedilol belongs to the 'third generation' of  $\beta$ -adrenoceptor antagonists and shows selectivity for the  $\beta_1$ - rather than  $\beta_2$ -adrenoceptor. Carvedilol is also an  $\alpha_1$ -blocking agents, with around 2- to 3-fold more selectivity for  $\beta_1$ - than  $\alpha_1$ -adrenoceptors. This degree of  $\alpha_1$ -blockade is responsible for the moderate vasodilator properties of carvedilol, being different from other  $\beta$ -adrenoceptor antagonists. In addition, carvedilol is a potent antioxidant, with a 10-fold greater activity than vitamin E. Some carvedilol metabolites found in human plasma also exhibit antioxidative activity approximately 50- to 100-fold greater than carvedilol and other antioxidants. These unique properties of carvedilol, i.e. adrenergic ( $\beta_1$ ,  $\beta_2$  and  $\alpha_1$ ) blockade and antioxidative activity, may be important in preventing progressive deterioration of left ventricular dysfunction and chronic heart failure. Recently, carvedilol

has been demonstrated to reverse multidrug resistance (MDR) to anticancer drugs in tumor cells *in vitro* and its reversal effects were comparable with verapamil, which has been used in the first clinical trial for the reversal of MDR. This review introduces the reversal activity and usefulness against MDR, as well as an overview of the pharmacological and pharmacokinetic properties, of carvedilol. *Anti-Cancer Drugs* 15:303–309 © 2004 Lippincott Williams & Wilkins.

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## Development of carvedilol

In 1983, carvedilol [1-[carbazolyl-(4)-oxy]-3-[(2-methoxyphenoxyethyl)amino]-2-propanol] (Fig. 1) was designed and developed as a  $\beta$ -adrenoceptor antagonist with vasodilating activity for efficacious and safe treatment of hypertension and coronary artery disease. This drug contains an oxyisopropanolamine moiety with aromatic substituents linked to both the oxy and amine ends [1–3]. A common structural feature of  $\beta$ -adrenoceptor antagonists is either an aryethanolamine or an aryloxyisopropanolamine moiety, but the compounds differ in the nature of the aryl group, as well as the group(s) linked to the amine moiety. The most important differences between  $\beta$ -adrenoceptor antagonists are related to the degree of selectivity for the  $\beta_1$ - against  $\beta_2$ -adrenoceptor [4–6]. Based on these characteristics,  $\beta$ -adrenoceptor antagonists have been categorized into three classes (Table 1).

The 'first-generation' compounds, such as propranolol and timolol, are non-selective drugs with equal affinity for blocking  $\beta_1$ - and  $\beta_2$ -adrenoceptors, and no important pharmacological effects other than the  $\beta$ -blockade. The 'second-generation' compounds, such as metoprolol, atenolol, ceriprolol, acebutolol and bisoprolol, show

selectivity for the  $\beta_1$ -adrenoceptor. Carvedilol belongs to the 'third-generation' compounds including bucindolol, labetalol and nebivolol, and has selectivity (around 7-fold) for  $\beta_1$ - rather than  $\beta_2$ -adrenoceptors, but becomes non-selective at higher target doses. Carvedilol is also an  $\alpha_1$ -blocking drug, with around 2- to 3-fold more selectivity for  $\beta_1$ - than  $\alpha_1$ -adrenoceptors. This degree of  $\alpha_1$ -blockade is responsible for the moderate vasodilator properties of carvedilol, being different from other  $\beta$ -adrenoceptor antagonists.

In addition, carvedilol is a potent antioxidant, with 10-fold greater activity than the antioxidant vitamin E [7–9], and prevents depletion of vitamin E, glutathione and SH protein induced by oxidative stress. These actions are the main defense mechanisms against tissue injury caused by free radicals and their property derives from the carbazole portion of its chemical structure (Fig. 1). Also, some carvedilol metabolites found in human plasma exhibit antioxidative activity approximately 50- to 100-fold greater than that of carvedilol and other antioxidants to inhibit low-density lipoprotein oxidation by mouse macrophages [10,11]. Collectively, these unique properties of carvedilol, i.e. multiple adrenergic ( $\beta_1$ ,  $\beta_2$  and  $\alpha_1$ ) blockade and antioxidative activity, might be one of the

reasons that carvedilol was approved for the treatment of chronic heart failure, although the use of  $\beta$ -adrenoceptor antagonists for chronic heart failure had been questioned because of negative inotropic effects [12].

The pharmacokinetic profiles of carvedilol have been investigated in healthy volunteers and patients with hypertension [1,2,13,14]. Carvedilol is used clinically as a racemic mixture of *R*(+)- and *S*(-)-enantiomers, and demonstrates dose-linear behavior. The absolute oral bioavailability reached 24% probably due to a first-pass effect. After a single oral administration of 50 mg, maximum concentrations of about 70 ng/ml (around 0.2  $\mu$ M) were achieved within at least 2 h. The plasma concentration of carvedilol at the steady-state was around 160 ng/ml (around 0.4  $\mu$ M) after repetitive administration at a dose of 50 mg/day. In addition, carvedilol was extensively distributed to the tissues (volume of distribution: around 130 l) and eliminated primarily by hepatic metabolism (total clearance: around 600 ml/min, renal clearance: 4 ml/min). Carvedilol is metabolized by mainly cytochrome P450 2D6 (CYP2D6), with CYP2C9

and CYP1A2, but not CYP3A4, having also been clarified to participate in the metabolism in humans [15]. Detailed information on the pharmacokinetics and pharmacodynamics of carvedilol is available in the literature [1,2,16–19].

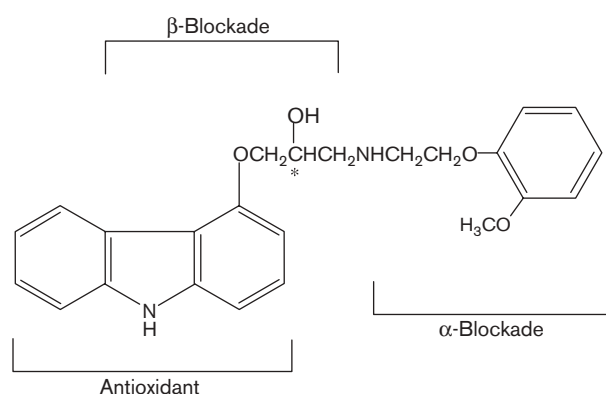
Recently, carvedilol has been demonstrated to reverse multidrug resistance (MDR) to anticancer drugs in tumor cells *in vitro* [20,21] and its reversal effects were comparable to those of verapamil, which has been used as a MDR modulator for the first time in clinical trials for the reversal of MDR. In this review, we introduce the reversal activity and usefulness against MDR of carvedilol from the viewpoint of direct interaction with MDR1,  $\beta$ -adrenoceptor selectivity, metabolic process, antioxidative activity, etc.

### MDR in cancer chemotherapy

The development of MDR remains a major hurdle to successful cancer chemotherapy [22–24]. Although the cellular basis underlying MDR is not fully understood, several factors have been identified [25–31]. These include down-regulation of uptake, induction of the efflux system (MDR1/P-glycoprotein, etc.), induction of inactivation enzymes (glutathione-S-transferase, etc.), alteration of the targeted molecule (topoisomerase, etc.), changes in DNA repair processes and alteration of the apoptotic mechanism (p53 mutation, bcl-2 overexpression, etc.). The overexpression of MDR1/P-glycoprotein in tumor cells is one of the best-characterized mechanisms of MDR [26,27,32–35]. Recently, the term MDR1 has come to be used instead of human P-glycoprotein, so MDR1 is used in this review.

In 1976, a 170-kDa glycosylated membrane protein (later called MDR1) was discovered through its ability to confer MDR in colchicine-resistant Chinese hamster ovary (CHO) cells by Juliano and Ling [36]. About 10 years later, a gene was isolated from multidrug resistant KB carcinoma cells and designated *MDR1* [37 and references therein]. *MDR1* is located on chromosome 7 at q21.1 in a 600-kb *NruI* fragment and the *MDR1* coding region is contained in a 120-kb *XhoI* fragment. This gene extends

Fig. 1



Chemical structure of carvedilol, [1-[carbazoyl-(4)-oxy]-3-[(2-methoxyphenoxyethyl)amino]-2-propanol], with postulated active sites. An asterisk indicates the point of asymmetry.

Table 1 Pharmacological properties of representative  $\beta$ -adrenoceptor antagonists *in vitro*

Generation	Drug	$\beta_1$ -Blockade	$\beta_2$ -Blockade	$\alpha_1$ -Blockade	Antioxidative activity
First	propranolol	++	++	–	+
	timolol	+++	+++	–	+
Second	acebutolol	+	–	–	?
	atenolol	++	–	–	?
	bisoprolol	+	–	–	–
	celiprolol	+	–	–	–
	metoprolol	+	–	–	?
	bucindolol	++	+	–	–
Third	carvedilol	++	+	+	+++
	labetalol	+	+	+	?
	nebivolol	++	+	–	–

This table was constructed by referring to [4,5,10,11,86,87]. A question mark '?' indicates inconsistent reports concerning the antioxidative activity.

over more than 100 kb containing 28 introns, 26 of which interrupt the protein-coding sequence. MDR1 mRNA has a size of 4.6 kb, thus its coding region accounts for less than 5% of the total.

MDR1, the product of the *MDR1* gene, belongs to the ATP-binding cassette (ABC) family of transporters, currently numbering 49 members, that share sequence and structural homology (see also <http://nutrigene.4t.com/humanabc.htm>). MDR1 is a phosphorylated and glycosylated membrane protein of 1279 amino acids, and consists of two homologous halves containing six putative hydrophobic transmembrane segments and an intracellular binding site for ATP [37 and references therein]. This protein has been understood to act as an efflux pump to expel anticancer drugs from cells utilizing the energy of ATP hydrolysis, thus lowering intracellular concentrations. The anticancer drugs that are transported by MDR1 include hydrophobic and/or amphipathic drugs, such as the taxanes (paclitaxel and docetaxel), Vinca alkaloids (vinblastine, vincristine, and vinorelbine), anthracyclines (doxorubicin and daunorubicin), epipodophyllotoxins (etoposide and teniposide), topotecan, actinomycin D and mitomycin C. As a result, MDR1-overexpressing tumor cells show resistance to multiple anticancer drugs. Therefore, the development of MDR1 inhibitors, i.e. MDR modulators, has been carried out industriously with sights set on the reversal of MDR.

### Attempts to reverse MDR

The calcium channel blocker verapamil was the first drug found to inhibit MDR1-mediated transport *in vitro* [38]. Tsuruo *et al.* demonstrated a decreased efflux of vincristine in vincristine-resistant P388 leukemia cells in the presence of verapamil [38]. In 1984, Rogan *et al.* started the first clinical trial with verapamil and doxorubicin in patients with ovarian cancer [39]. However, the majority of investigations with these drugs demonstrated disappointing results because low affinities for MDR1 necessitated the use of high doses of modulators, resulting in unacceptable toxicity.

To overcome these problems, second-generation MDR modulators had been developed, such as dexverapamil, valspodar (PSC833) and biricodar (VX-710) [40–43]. The second-generation MDR modulators have been demonstrated to be more effective for reversal of MDR than the first-generation MDR modulators *in vitro* and in animal experiments. However, the second-generation MDR modulators also influence the pharmacokinetics of anticancer drug through the inhibition of MDR1-mediated biliary excretion or intestinal transport. In addition, another mechanism by which they influence the pharmacokinetics of anticancer drug was clarified to be competition for CYP3A4-mediated liver or intestinal metabolism [44,45] and thus the use of second-generation

MDR modulators has led to unacceptable toxicity of anticancer drugs in clinical trials. The most common response of clinical researchers to these pharmacokinetic interactions has been to reduce the dose of anticancer drugs. Regrettably, the pharmacokinetic interactions between MDR modulators and anticancer drugs are complicated and unpredictable, and the doses of anticancer drugs co-administered with MDR modulators could not be determined in advance. Thus, there has been no establishment of a safe and effective dose of anticancer drugs. However, it has been noted that MDR1 and CYP3A4 share significant overlap in substrate specificity [46–51] (refer to the next section).

To overcome the limitations of the second-generation MDR modulators, third-generation MDR modulators have been developed using quantitative structure–activity relationship (QSAR) analysis and combinatorial chemistry. The third-generation MDR modulators currently in clinical development include tariquidar (XR9576), zosuquidar (LY335979), R101933 and ONT-093 [52–56]. To date, the results of clinical trials show that the third-generation MDR modulators can be given with full therapeutic doses of anticancer drugs with minimal interference to the pharmacokinetics of anticancer drugs [43]. The preliminary results with the third-generation MDR modulators offer new hope that this goal might be realized. This would be mainly explained by the finding that the third-generation MDR modulators did not affect CYP3A4-mediated metabolism of anticancer drugs at clinically achievable concentrations. Therefore, a key to developing MDR modulators is understanding the overlap in substrate specificity between MDR1 and CYP3A4. Unfortunately, it will presumably take a little more time before the third-generation MDR modulators are released for clinical use. Consequently, drugs marketed for other uses have attracted attention again as candidates for MDR reversal, but the overlap in substrate specificity between MDR1 and CYP3A should be clarified in advance.

### Overlap in substrate specificity between MDR1 and CYP3A

MDR1 is expressed not only in resistant tumor cells, but also in normal tissues with an excretory function such as the biliary canalicular membrane of hepatocytes, the luminal membrane of endothelial cells in the blood–brain barrier and blood–testis barrier, the apical membrane of the syncytial trophoblasts of the placenta, the epithelial apical membrane of the intestine, and the renal proximal tubules [57]. From this evidence, MDR1 may be an important barrier to xenobiotics. In addition, pharmacokinetic studies have demonstrated the importance of MDR1 in limiting oral drug bioavailability and the distribution of drugs to tissue when the drugs are MDR1 substrates [37,58–60].

CYP3A is the CYP subfamily responsible for phase I metabolism of more than 50% of drugs administered to humans [61]. CYP3A4 is the most prominent CYP in humans, comprising about 30 and 70% of total CYP in liver and intestine, respectively. CYP3A4 has very broad substrate specificity, encompassing a wide variety of drugs [46]. Many anticancer drugs, such as epipodophyllotoxins and Vinca alkaloids, are substrates for CYP3A4 [62–64].

Like CYP3A4, MDR1 seems to have broad substrate specificity [37,45,59]. Interestingly, a striking overlap of substrates for MDR1 and CYP3A4 has been observed [46–51]. Focusing on these observations and expression sites, Benet and his collaborators proposed that proteins of MDR1 and CYP3A4 act synergistically to present a barrier to absorption from the small intestine and to distribute to the tissues [58,59]. Inhibition of one or both of these proteins can be expected to enhance the bioavailability and to change the distribution of many drugs. That is to say, MDR modulators such as the substrate for CYP3A inhibited not only the MDR1-mediated transport of anticancer drugs, but also CYP3A-mediated metabolism. Therefore, the pharmacokinetics of anticancer drugs in combination with such MDR modulators is complicated and so it is difficult to manage effectively MDR modulators in cancer chemotherapy. Although some compounds interact with MDR1 and CYP3A4 to a similar extent, for the most part, the potency of inhibition for MDR1 does not predict the potency of inhibition for CYP3A and vice versa. Moreover, not all the substrates for CYP3A are substrates for MDR1 and vice versa, e.g. ketoconazole, midazolam and digoxin, etc. [65]. Consequently, it is desirable that MDR modulators only have inhibitory activity for MDR1 not CYP3A4.

### Reversal effects of carvedilol on MDR1-mediated MDR

In 1999, Jonsson *et al.* suggested the possibility that carvedilol acts as a MDR modulator in the human breast cancer cell line Hs578T, with no measurable MDR1 expression and its MDR1-overexpressing doxorubicin-resistant subline Hs578T-Dox [20]. They demonstrated that 10  $\mu$ M of carvedilol only marginally affected the IC<sub>50</sub> value of doxorubicin, the dose which results in the death of half the number of cells, in Hs578T cells, whereas the IC<sub>50</sub> value of doxorubicin in Hs578T-Dox cells was reduced 1/20. Also, a decrease in accumulation of calcein, a fluorescence substrate for MDR1, was shown in Hs578T-Dox cells compared to the host Hs578T cells and the effect was reversed by preincubation with 1 or 10  $\mu$ M carvedilol. However, the concentration of carvedilol examined, i.e. 10  $\mu$ M, was much higher than the clinically available concentration, and an estimation of its clinical potency was complicated.

In 2003, we examined the reversal activity of carvedilol against MDR1-mediated MDR using a realistic clinical concentration, 1  $\mu$ M [21]. In human cervical carcinoma HeLa-Ohio (HeLa) cells, the growth curves for vinblastine, paclitaxel and cisplatin were not altered by the addition of 1  $\mu$ M carvedilol. However, the cytotoxicity of vinblastine and paclitaxel in the MDR1-overexpressing vinblastine-resistant HeLa subline (Hvr100-6) was increased 3.5- and 7.1-fold by 1  $\mu$ M carvedilol treatment, respectively and this reversal occurred in a carvedilol concentration-dependent manner. In addition, carvedilol slightly modified the cytotoxicity of doxorubicin and daunorubicin in Hvr100-6 cells. In contrast, the IC<sub>50</sub> values for 5-fluorouracil and cisplatin, which were not substrates for MDR1, were not affected. These findings using a realistic clinical concentration of carvedilol suggested that carvedilol had the clinical potential to reverse MDR1-mediated MDR. We also indicated that carvedilol effectively restored the intracellular accumulation of [<sup>3</sup>H]vinblastine, [<sup>3</sup>H]daunorubicin or [<sup>3</sup>H]digoxin in MDR1-transfected cells (LLC-GA5-COL150 cells) in a concentration-dependent manner [21,66]. Carvedilol was clarified to suppress the MDR1-mediated trans-cellular transport of [<sup>3</sup>H]daunorubicin or [<sup>3</sup>H]digoxin in LLC-GA5-COL150 cells, but no alteration of transport was found in host LLC-PK<sub>1</sub> cells. Moreover, the inhibitory effects of carvedilol on the trans-cellular transport of [<sup>3</sup>H]daunorubicin, [<sup>3</sup>H]vinblastine or [<sup>3</sup>H]digoxin were comparable to those of verapamil. These findings suggest that the reversal of MDR by carvedilol was associated with the direct inhibition of MDR1 and the reversal effects of carvedilol were comparable with those of verapamil at the same concentrations *in vitro* [20,21].

Verapamil was the first drug found to inhibit MDR1-mediated transport and clinical trials using verapamil were conducted for the reversal of MDR for the first time [38,39]. In those trials, the serum concentrations of verapamil were only approximately 0.5–1.0  $\mu$ M [21]. On the other hand, the maximum concentrations of carvedilol after a single oral administration of 50 mg was about 70 ng/ml (around 0.2  $\mu$ M) and the plasma concentration at the steady state was around 160 ng/ml (around 0.4  $\mu$ M) after repetitive administration at a dose of 50 mg/day [1,2,13,14]. Since it was previously suggested that carvedilol and verapamil showed similar effects in terms of MDR reversal, carvedilol could be a new candidate modulator of MDR1-mediated MDR in clinical use. However, for controlling the pharmacokinetics and pharmacodynamics of anticancer drugs co-administered with carvedilol, details of the reversal mechanism of carvedilol should be clarified *in vitro*. Then, the effectiveness and usefulness will need to be examined in clinical studies.

## Mechanism by which carvedilol reverses MDR1-mediated MDR

Carvedilol has potent antioxidative activity, as well as  $\beta$ -adrenoceptor antagonistic activity [7–9]. Therefore, other mechanisms of reversing MDR may exist, other than the direct inhibition of MDR1.

We demonstrated that propranolol ( $\beta_1$ - and  $\beta_2$ -adrenoceptor antagonist) and metoprolol and atenolol (both  $\beta_1$ -adrenoceptor selective antagonists) did not affect MDR1-mediated MDR and transport [21], suggesting that the effects of carvedilol ( $\beta_1$ -adrenoceptor selective antagonist with  $\alpha_1$ -blockade) on MDR1 were independent of its action on  $\beta$ -adrenoceptors. However, talinolol ( $\beta_1$ -adrenoceptor selective antagonist) has been demonstrated to inhibit the MDR1-mediated transport of digoxin [67]. In addition, celiprolol ( $\beta_1$ -adrenoceptor selective antagonist with slight  $\alpha_2$ -agonist action) and acebutolol ( $\beta_1$ -adrenoceptor selective antagonist) have found to be substrates for MDR1 [68–70]. Collectively, the role of  $\beta$ -adrenoceptor antagonists in the inhibitory activity of MDR1 may be dependent on differences in receptor selectivity, not action on  $\beta$ -adrenoceptors.

Concerning the relation between MDR1 and reactive oxygen species, the induction of MDR1 has been reported to be associated with the generation of reactive oxygen species [71–73]. Ziemann *et al.* [72] indicated that the addition of  $H_2O_2$  or a catalase inhibitor induced the expression of *mdr1b*, the MDR1 ortholog in rats, and its mRNA, and antioxidants markedly suppressed them in primary rat hepatocyte cultures. However, we found no effects of antioxidants, i.e. baicalein, genistein, quercetin, ascorbic acid, hydroquinone, superoxide dismutase and catalase, on MDR1 mRNA expression in HeLa cells and MDR1-overexpressing Hvr100-6 cells [74]. Moreover, the cytotoxicity of anticancer drugs in HeLa and Hvr100-6 cells was not altered by addition of these antioxidants [74]. Jonsson *et al.* [20] also reported that carvedilol did not affect pyrogallol cytotoxicity, which was a superoxide radical generator, and pyrogallol was without any effect on calcein accumulation of MDR1-overexpressing Hs578T-Dox cells, indicating a lack of antioxidative properties affecting MDR1 activity and associated toxicity of the drug [21]. Anyway, the action of antioxidants against MDR1 remains unclear.

Various types of drugs except for anticancer drugs have been clarified to regulate MDR1 expression [75–78]. In addition, some factors affecting MDR1 expression have been identified [79–81], although the transcriptional regulation of the *MDR1* gene was unexpectedly complex and is far from being completely understood. The *MDR1* promoter was found to contain a GC-box for Sp1, an inverted CCAAT element (Y-box) for YB-1 and NF-Y, a p53 element, an AP-1 element, a CAAT element for a

complex of NF- $\kappa$ B and *c-fos* proteins, a C/EBP element for NF-IL-6, a heat-shock element (HSE) for heat-shock transcription factor (HSF), an inverted MED-1 for an unknown nuclear protein of about 150–160 kDa, and a steroid xenobiotic receptor (SXR) element for the orphan nuclear receptor SXR (also known as PXR, PAR, PRR or NR1I2). In the latest investigations, some drugs have been found to interact with these transcription factor(s) [82–84]. Moreover, it was reported that the development of MDR1-mediated MDR was a two-step post-transcriptional process mediated by changes in both MDR1 mRNA stability and translation [85]. Consequently, the understanding of these transcriptional processes of MDR1 has led to new possibilities for the reversal of MDR in cancer chemotherapy.

## Future directions

From the standpoint of the reversal of MDR using  $\beta$ -adrenoceptor antagonists, the role of  $\beta$ -adrenoceptor antagonists in the inhibitory activity of MDR1 may be dependent on receptor selectivity, not action on  $\beta$ -adrenoceptors. Although the evidence is not complete, the utilization of differences in receptor selectivity for  $\beta$ -adrenoceptor antagonists may lead to a novel course for the development of MDR modulators.

In addition, it is desirable that MDR modulators have inhibitory activity for MDR1 but not CYP3A4, because MDR1 and CYP3A4 share significant overlap in substrate specificity. So, it will be important to design and develop MDR modulators with such specificity. To accomplish this,  $\beta$ -adrenoceptor antagonists including carvedilol, which are mainly metabolized by CYP2D6, will be attractive as novel candidates for the lead compound of a MDR modulator.

A link between MDR1 and reactive oxygen species certainly exists based on the past evidence. Carvedilol has the strongest antioxidative activity among the  $\beta$ -adrenoceptor antagonists (Table 1). Therefore, it is speculated that the antioxidative activity of carvedilol participates in the reversal of MDR. From the evidence of this, the use of antioxidants may achieve a breakthrough in the reversal of MDR.

On the other hand, the transcriptional processes of the *MDR1* gene have gradually been clarified with the advancement of genome science. Moreover, it has been demonstrated that some drugs interacted with factors participating in the transcriptional processes of the *MDR1* gene and thus this transcriptional process must be a hot target for the reversal of MDR. In the near future, these understandings will lead to new possibilities for the reversal of MDR in cancer chemotherapy. Therefore, the effects of  $\beta$ -adrenoceptor antagonists including carvedilol on the transcriptional processes should be elucidated.

## Conclusion

It is very important that MDR modulators have a strong and specific inhibitory potency against MDR1, are clinically available drugs at present, and are non-substrates/inhibitors for CYP3A4. So, carvedilol could be a novel candidate modulator of MDR1-mediated MDR. However, its usefulness and effectiveness cannot be satisfied yet for the reversal of MDR in the clinical setting. To be successful in reversing MDR, it will be also necessary to carry out the development of MDR modulators from the viewpoint, which has not been done up to this point.

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