Identification of a pharmacokinetic interaction between teicoplanin and sulfo-butyl ether-β-cyclodextrin, an excipient in the intravenous posaconazole

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Abstract

   Patients undergoing hematopoietic stem cell transplantation (HSCT) often receive multiple antibiotics and antifungals concurrently, making it crucial to understand their potential pharmacokinetic interactions of these agents. We report here an interaction between the glycopeptide antibiotic teicoplanin (TEIC) and sulfo-butyl ether-β-cyclodextrin (SBECD), a solubilizing excipient in the intravenous formulation of posaconazole (PSCZ). We performed a single-center retrospective analysis of patients who underwent HSCT and received oral and intravenous PSCZ during TEIC therapy. The associations between PSCZ administration and TEIC concentration-to-dose (C/D) ratios were evaluated using linear mixed-effects models. We also examined the effects of intravenous PSCZ and SBECD on TEIC pharmacokinetics in rats by assessing the area under the concentration–time curve (AUC) and urinary excretion of total TEIC and its components. In addition, molecular docking and in vitro protein-binding assays were conducted to investigate the interaction between TEIC and SBECD. In patients who underwent HSCT, TEIC C/D ratio was significantly lower following intravenous PSCZ administration than without administration. In contrast, the effect of oral PSCZ administration relative to non-administration was not statistically significant. In rats, intravenous PSCZ and SBECD decreased the AUC of TEIC and increased urinary excretion, particularly in the A_2 group. Docking simulations indicated that the hydrophobic side chain of TEIC A_{2-2} fit within the SBECD cavity, and in vitro assays confirmed SBECD concentration-dependent increases in TEIC unbound fractions. The co-administration of intravenous PSCZ containing SBECD may reduce TEIC protein binding, thereby enhancing renal excretion.

Introduction

Patients who undergo hematopoietic stem cell transplantation (HSCT) are susceptible to bacterial and fungal infections owing to immunosuppression caused by conditioning chemotherapy before transplantation and subsequent immunosuppressive therapy. Consequently, multiple antibiotics and antifungal agents are often administered concomitantly to these patients for the prophylaxis and treatment of infections. Teicoplanin (TEIC), a glycopeptide antibiotic, is one of the agents frequently used in this population. Along with vancomycin, TEIC is widely used in the treatment of methicillin-resistant *Staphylococcus aureus* infections, and is administered empirically when gram-positive bacterial infections are suspected. TEIC undergoes minimal metabolism and is eliminated primarily through the kidneys. Compared to vancomycin, TEIC exhibits a higher protein-binding rate, a larger volume of distribution, and a substantially longer half-life of approximately 83–168 h (1, 2). Target trough concentrations required for therapeutic efficacy have been established based on pharmacokinetic-pharmacodynamic analyses (3, 4); thus, maintaining appropriate plasma concentrations is critical for successful treatment. To this end, fluctuations in TEIC exposure may result in treatment failure, highlighting the need to fully elucidate inter- and intra-individual factors that contribute to pharmacokinetic variability.

Azole antifungal agents are commonly administered to patients undergoing HSCT to prevent fungal infections. Posaconazole (PSCZ), an azole antifungal agent, is known for its potent antifungal activity and inhibition of the drug-metabolizing enzyme CYP3A and efflux transporter P-glycoprotein in the liver and small intestine (5, 6). Therefore, co-administration of PSCZ with drugs that are substrates of these enzymes or transporter may reduce the clearance of the co-administered drugs, potentially leading to drug–drug interactions. In contrast, PSCZ is unlikely to significantly affect the pharmacokinetics of renally excreted drugs. Patients undergoing HSCT often receive multiple antibiotics and antifungals concurrently; however, data on the pharmacokinetic interactions among these agents in this population remain limited.

Therefore, the aim of this study was to evaluate the effect of intravenous (i.v.) PSCZ co-administration on the plasma concentration of TEIC in patients undergoing HSCT. We assessed the reproducibility of this phenomenon using an animal model, and further identified sulfo-butyl ether-β-cyclodextrin (SBECD), a pharmaceutical excipient

contained in i.v. PSCZ, as a potential contributor to the reduction in plasma TEIC

concentrations. Moreover, we predicted the formation of an inclusion complex of

SBECD with TEIC, and evaluated the effect of SBECD on TEIC protein binding.

Materials and Methods

<u>Effects of oral and intravenous PSCZ administration on TEIC plasma concentrations in</u>

<u>patients who underwent HSCT</u>

We conducted a single-center retrospective analysis of patients who underwent

HSCT and received the tablet formulation (oral) and injectable formulation of PSCZ

(i.v. PSCZ) during TEIC treatment between April 2020 and January 2022 at the

Department of Hematology, Kyoto University Hospital. Eligible patients were required

to have available plasma TEIC concentration data under three conditions: without PSCZ

administration, following oral PSCZ administration, and following i.v. PSCZ

administration. This study was conducted in accordance with the Declaration of

Helsinki and its amendments and was approved by the Ethics Committee of Kyoto

University Graduate School and Faculty of Medicine (Approval No. R2470).

We extracted all available data on trough TEIC plasma concentrations from the

hospital electronic medical database. Plasma TEIC concentrations were measured in the

hospital's central laboratory by latex immunoturbidimetry using the Cobas 6000 system.

TEIC plasma concentration data were included if they met the following criteria: (i)

obtained at least 3 days after maintaining a constant dosage following TEIC loading, (ii)

obtained at least 3 days after dosage adjustment, or (iii) obtained at least 3 days after

switching the PSCZ formulation. We also collected data on sex and age, which are

generally assumed to influence drug pharmacokinetics, as well as serum albumin levels

and estimated glomerular filtration rate (eGFR), which are known to affect the

pharmacokinetics of TEIC (7).

To evaluate the association between PSCZ administration and changes in TEIC

pharmacokinetics, the effects of PSCZ on the concentration-to-dose (C/D) ratio of TEIC

were estimated using a linear mixed-effects model:

$$C/D\ ratio\ = \alpha + \beta_1 * I_{oral} + \beta_2 * I_{intravenous} + \gamma + \varepsilon$$

where α is the estimate of the intercept, β is the estimate of the fixed effect, I_{oral} and

$I_{intravenous}$ are the indicator variables, γ is the estimate of the random effect for the

intercept (individual-level residual with mean zero and variance σ^2_γ), and ε is the

estimated residual error (with mean zero and variance σ^2_ε). $I_{oral} = 1$ if oral PSCZ was co-

administered, and 0 otherwise; $I_{intravenous} = 1$ if i.v. PSCZ was co-administered, and 0

otherwise. The effects of PSCZ were further adjusted for age, sex, serum albumin level,

and eGFR, which were incorporated as independent covariates with fixed effects.

Analyses were performed using JMP software version 18 (SAS Institute Inc., Cary, NC,

U.S.A.)

<u>Pharmacokinetics of TEIC in rats</u>

129   Eight-week-old male Wistar/ST rats were purchased from the SLC Animal

Research Laboratories (Shizuoka, Japan) and maintained in accordance with the

guidelines for Animal Experiments of Kyoto University, Kyoto, Japan. The study was

approved by the Animal Research Committee of the Graduate School of Medicine,

Kyoto University, Kyoto, Japan (permit numbers: Medkyo 22119 and 23107). The rats

were housed in a specific pathogen-free facility, maintained in a temperature-controlled

environment with a 12-h light/dark cycle, and fed a standard diet and water ad libitum.

For the administration of TEIC and PSCZ to rats, the commercially available i.v.

formulations, Teicoplanin for Intravenous Infusion 200 mg (Towa) (Towa

Pharmaceutical Co., Ltd. Osaka, Japan) and Noxafil® for Intravenous Infusion 300 mg

(MSD K.K., Tokyo, Japan) were used. SBECD was purchased from ChemScene (NJ,

USA). All other reagents and solvents were of analytical or liquid chromatography-mass

spectrometry grade.

142   The rats were anesthetized with a combination of medetomidine, midazolam,

and butorphanol, and catheters were inserted into the left femoral vein and right femoral

artery using polyethylene tubing (SP-31; Natsume Seisakusho, Tokyo, Japan). Saline,

PSCZ (40 mg/kg PSCZ plus 890.67 mg/kg SBECD), or SBECD alone (890.67 mg/kg)

was administered through the femoral vein. These doses were set with reference to the

human equivalent dose in rats, where the dose per body weight is approximately six

times the human dose (8). Thirty minutes later, TEIC (20 mg/kg) was injected into the

femoral vein. Six rats per group were used in the experiment. PSCZ is highly

hydrophobic, making it difficult to dissolve in a solvent suitable for i.v. administration

without SBECD. Therefore, the PSCZ-alone group was not included in this study.

Blood samples were collected from the femoral artery at the end of TEIC administration

and 5, 15, 30, 60, 90, and 120 min thereafter. Plasma was separated by centrifugation.

Urine was collected from the bladder into a polyethylene tube for 120 min after TEIC

administration. Plasma and urine samples were deproteinized by adding four volumes of

methanol, followed by centrifugation at $10,000 \times g$ for 5 min. A 50 µL aliquot of the

supernatant was mixed with 200 µL of methanol. Urine samples were further diluted

five-fold with methanol. Lastly, all samples were filtered through a 0.45 µm membrane

filter prior to analysis.

TEIC was quantified using a triple-quadrupole mass spectrometry system

(LCMS-8040; Shimadzu, Japan) equipped with an electrospray ionization source. The

mobile phases were 0.1% formic acid in water (phase A) and 0.1% formic acid in

acetonitrile (phase B). The gradient elution program was as follows: 0–1.0 min, 5% B;

1.0–3.0 min, 5–100% B; 3.0–3.5 min, 100% B; 3.5–4.0 min, 100–5% B; 4.0–6.5 min,

5% B. The flow rate was 0.25 mL/min, the column oven temperature was 60 °C, and the

injection volume was 5 µL. Chromatographic separation was achieved using an

ACQUITY UPLC BEH C18 column (50 × 2.1 mm, 1.7 µm particle size; Waters, USA).

Detection was performed in positive ion mode using multiple reaction monitoring.

Since the clinically used i.v. formulation of TEIC comprises five major components ($A_{2\text{-}1}$,

$A_{2\text{-}2}$, $A_{2\text{-}3}$, $A_{2\text{-}4}$, and $A_{2\text{-}5}$), and one hydrolysis component ($A_{3\text{-}1}$) (Fig. 1), the following

ion transitions were monitored to detect each component: $A_{2\text{-}1}$, from m/z 939.7 to 314.2;

$A_{2\text{-}2}$ and $A_{2\text{-}3}$, from m/z 940.7 to 316.2; $A_{2\text{-}4}$ and $A_{2\text{-}5}$, from m/z 947.8 to 330.2; $A_{3\text{-}1}$,

from m/z 782.4 to 203.7 (9).

The area under the plasma concentration–time curve (AUC) and the urinary

excretion of TEIC were determined over 120 min after administration. The AUC and

urinary excretion were calculated for total TEIC, and each individual component.

Statistical analyses were performed using GraphPad Prism version 8.0 software

(GraphPad, San Diego, CA, USA). The data were analyzed using one-way analysis of

variance; a P-value < 0.05 was considered statistically significant.

<u>Molecular docking simulation</u>

Since TEIC $A_{2\text{-}2}$ is the major component of the clinical formulation and its

conformation is crystallographically characterized, only this dominant analog was

considered for molecular modeling. The 3D structure of TEIC A_{2-2} was obtained from
the Protein Data Bank (ID: 4PJZ) (10). The structure of SBECD was constructed using
Avogadro software (ver. 1.2.0) (11) on the basis of the 3D structures of native β-CD that
was retrieved from the Protein Data Bank (ID: 3CGT). Based on previous reports (12),
the structure of SBECD was prepared by assuming that seven glucopyranose units were
substituted with sulfobutyl ether, namely four at position C2 on the secondary face
(residues 1, 3, 5, and 7) and three at position C6 on the primary face (residues 2, 4, and
6), in accordance with the average degree of substitution of the SBECD reagent (6.0-
7.1). The molecular geometries of TEIC A_{2-2} and SBECD were fed into GAMESS (ver.
2023 R1) (13) for geometry optimization using the semi-empirical quantum chemistry
method PM3. By treating the geometry-optimized structures of TEIC A_{2-2} and SBECD
as ligands and receptors, respectively, molecular docking to obtain complexes of TEIC
A_{2-2} with SBECD was performed with 100 runs using the Lamarckian genetic algorithm
in AutoDock 4.2 (14). The receptor and ligand were assigned Gasteiger partial charges
and the rotatable bonds were not defined. All other parameters were set to the default
values for AutoDock 4.2.
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<u>Effect of SBECD on TEIC protein binding in human serum</u>
SBECD (40 mg/mL in distilled water) was diluted with human serum (Kojin
Bio, Saitama, Japan) to prepare serum samples containing 0, 100, 300, and 1000 µg/mL
of SBECD. These concentrations were within the range observed in clinical settings (15,
16). Aliquots of 380 µL of each serum sample were incubated at 37 °C for 30 min.
Subsequently, 20 µL of TEIC injection solution, prepared with distilled water to a
concentration of 0.8 mg/mL, was added to achieve a final concentration of 40 µg/mL.
This corresponded to the upper limit of the clinically targeted trough concentration of
TEIC. The samples were then incubated at 37 °C for 60 min, with gentle mixing for 10 s
every 10 min during incubation. After incubation, 300 µL of each sample was
immediately collected and filtered through an ultrafiltration membrane (30 kDa,
Millipore, Tokyo, Japan). TEIC calibration curves were generated using serum
subjected to the same ultrafiltration procedure used for the matrix. The concentration of
TEIC in the filtrate was quantified using the method described above, and the unbound
fraction was calculated relative to the concentration of 40 µg/mL. Data were analyzed

using one-way analysis of variance, and a P-value < 0.05 was considered statistically

significant.

Results

<u>Decreased TEIC plasma concentrations following i.v. PSCZ administration in patients</u>

<u>who underwent HSCT</u>

This study included eight patients who underwent HSCT and received both

TEIC and PSCZ during the observation period. The plasma TEIC concentrations were

measured under three conditions: without PSCZ administration, following oral PSCZ

administration, and following i.v. PSCZ administration. Among these patients, TEIC

was administered prior to PSCZ in five, while PSCZ was administered prior to TEIC in

three. A total of 83 plasma concentration measurements were obtained: 23 without

PSCZ administration, 27 following oral administration, and 33 following i.v.

administration. The data points per person ranged from a minimum of five to a

maximum of 18. The patient characteristics are summarized in Table 1. The average

TEIC C/D ratio was 3.53 (standard deviation [SD], 1.29) in the absence of PSCZ, 3.10

(SD, 1.26) following oral PSCZ administration, and 2.23 (SD, 0.95) following i.v.

administration (Fig. 2). A linear mixed-effects model demonstrated that the C/D ratio

was significantly lower following i.v. PSCZ administration than without administration

(estimate: -1.24; 95% confidence interval [CI], -1.74 to -0.74). After adjusting for serum

albumin, eGFR, age, and sex as covariates, the reduction in the C/D ratio associated

with i.v. PSCZ remained significant (estimate: -1.22; 95% CI, -1.84 to -0.60). This

corresponds to a decrease of approximately 35% compared to the non-PSCZ condition.

In contrast, the effect of oral PSCZ administration on the TEIC C/D ratio relative to

non-administration was not statistically significant (unadjusted: -0.36; 95% CI, -0.92 to

0.20; adjusted: -0.21; 95% CI, -0.88 to 0.45).

<u>Changes in TEIC pharmacokinetics following SBECD administration in rats</u>

The above results suggest that the i.v. formulation of PSCZ is associated with

reduced plasma TEIC concentrations. Because the dosage of PSCZ was identical

regardless of the route of administration (300 mg/day), we focused on SBECD, which is

present only in the i.v. formulation, as a potential factor influencing TEIC

pharmacokinetics. To test this hypothesis, rats were pretreated with either an i.v. PSCZ

formulation or SBECD followed by TEIC administration, and the AUC and urinary

excretion of TEIC were evaluated (Fig. 3). Compared to rats pretreated with saline,

those pretreated with either i.v. PSCZ or SBECD showed a significant decrease in TEIC

AUC and a significant increase in urinary excretion (Fig. 3A–C). No significant

differences were observed in the plasma concentrations immediately after

administration (Figure 3A) or significant differences in urine volume (Fig. 3D) between

the groups. Furthermore, the relative abundance of the individual components of the

TEIC formulation, A_{2-1}, $A_{2-2/2-3}$, $A_{2-4/2-5}$, and A_{3-1}, was determined. Following the i.v.

administration of PSCZ or SBECD, the AUC of the A_2 group significantly decreased,

whereas the urinary excretion significantly increased (Fig. 3E–G, 3I–K). In contrast, no

significant effects were observed for A_{3-1} (Fig. 3H and 3L).

<u>Molecular modeling of TEIC A_{2-2} inclusion complex with SBECD</u>

The abovementioned findings indicate that SBECD contributes to a reduction

in TEIC plasma concentration, with a greater impact on the A_2 group than A_{3-1}.

Structural differences exist between these groups because only the A_2 group possesses a

long carbon chain (Fig. 1). Furthermore, a previous study showed that a β-cyclodextrin

derivative, 2-hydroxypropyl-β-cyclodextrin (HP-β-CD), potentially prevents the

hydrolysis of the glycosyl group with a hydrophobic side chain of dalbavancin, a

semisynthetic glycopeptide with an analogous molecular scaffold of TEIC (17).

Therefore, we hypothesized that SBECD accommodates the side chain of TEIC, thereby

inhibiting its binding to serum proteins. To test this postulation, we performed docking

simulations for TEIC A_{2-2}, a major analog of TEIC, against SBECD using AutoDock

software. Through 100 docking runs, we obtained two probable binding modes for the

TEIC A_{2-2} complex with SBECD: the dimethyl terminus of the TEIC A_{2-2} side chain was

directed towards the primary face of the SBECD molecule for Type I (Fig. 4A), and the

dimethyl terminus was directed towards the secondary face of the SBECD molecule for

Type II (Fig. 4B). Regardless of the binding mode, most of the TEIC A_{2-2} hydrophobic

side chain, which contributes to an increased protein-binding rate, was accommodated

within the hydrophobic inner cavity of SBECD.

<u>Effect of SBECD on TEIC protein binding</u>

The above findings suggest that SBECD exerts a strong influence on TEIC

protein binding, particularly on the A_2 group. Therefore, we evaluated the unbound

fraction of TEIC in human serum containing clinically relevant concentrations of

SBECD (Fig. 5). The results demonstrated that the unbound fractions of both the A_2

group and A_{3-1} increased in a concentration-dependent manner with increasing SBECD

concentration. Compared with the condition without SBECD, the unbound fractions of

A_{2-1}, $A_{2-2/2-3}$, $A_{2-4/2-5}$, and A_{3-1} increased by 3.4-fold, 9.6-fold, 25.9-fold, and 1.3-fold,

respectively, at an SBECD concentration of 300 µg/mL.

Discussion

Multiple antibiotics and antifungal agents are often used concurrently in

patients undergoing HSCT, making it essential to fully understand the potential

interactions between these drugs. In this study, we demonstrated, for the first time, that

the co-administration of an i.v. formulation of PSCZ with TEIC in patients undergoing

HSCT resulted in reduced TEIC plasma concentration. A key strength of this study is

that the reproducibility of this phenomenon was confirmed by integrating clinical

observations with an animal model. Moreover, by focusing on the pharmaceutical

excipient SBECD, we further elucidated its impact on TEIC protein binding using in

vitro assays and the mode of inclusion complex formation through molecular modeling.

SBECD is a derivative of β-cyclodextrin, composed of seven glucose units

linked in a ring, with a hydrophobic inner cavity and a hydrophilic outer surface. This

structure enables the encapsulation of hydrophobic drugs, thereby improving their

aqueous solubility (18-20). Each vial of the i.v. PSCZ formulation (Noxafil® 300 mg)

contains 6,680 mg of SBECD. In general, cyclodextrin–drug inclusion complexes are

thought to have minimal effects on pharmacokinetics, as they dissociate rapidly after i.v.

administration (21-27). However, exceptionally stable complexes are known to

markedly alter drug elimination pathways. For instance, the rocuronium–sugammadex

complex exhibits a high stability constant, whereas rocuronium alone is excreted

through bile and rapidly eliminated in urine in the presence of sugammadex (21).

Similarly, drugs with an adamantane structure show particularly high affinity because of

their close fit with the cyclodextrin cavity (28), and urinary excretion rates have been

reported to increase from 0.1% to 37% in the presence of SBECD (29). In the current

study, rats administered with either the i.v. PSCZ formulation or SBECD alone

exhibited a significant increase in TEIC urinary excretion, supporting the interaction

between TEIC and SBECD in vivo.

Semisynthetic lipoglycopeptides, such as the TEIC A_2 group, telavancin,

dalbavancin, and oritavancin, exhibit high protein binding and consequently much

longer half-lives than vancomycin owing to their lipophilic side chains (30). Although

the binding affinity of TEIC for human serum albumin (HSA), the predominant serum

protein, has been reported (31) , its binding conformation remains unclear. Dalbavancin,

which shares a large structural similarity with TEIC, binds to HSA by inserting its

hydrocarbon chain into the hydrophobic pocket within subdomain IA of HSA,

composed entirely of nonpolar amino acids (32). Given that TEIC, despite isoform

differences, also possesses a long hydrocarbon chain and a conserved cyclic peptide

core, it is likely to bind albumin in a similar manner. This hypothesis is consistent with

our in vitro findings that showed a higher proportion of the unbound fraction of A_{3-1}

than that of the A_2 group. Furthermore, HP-β-CD has been shown to slowly hydrolyze

the glycosidic linkage between the mannosyl aglycone and hydrophobic side chain of

dalbavancin (17), suggesting that HP-β-CD interacts with and shields this region. The

use of HP-β-CD as a pharmaceutical excipient to enhance the solubility of telavancin

(Vibativ®) (33) and oritavancin (Kimyrsa®) (34) further supports our findings

regarding TEIC–CD interactions. A_{2-2}, A_{2-3}, A_{2-4}, and A_{2-5} are the major components of

clinical TEIC formulations. In this study, SBECD significantly increased the unbound

fraction of TEIC, particularly in the A_2 group. These results suggest that the

hydrophobic cavity of SBECD accommodates much of the hydrophobic side chain of

TEIC A_2, thereby reducing protein binding and leading to decreased plasma

concentrations through enhanced renal excretion.

In patients who underwent HSCT, the C/D ratio of TEIC decreased by an

average of 1.2 when the i.v. formulation of PSCZ was administered, compared with no

administration. In other words, when TEIC was administered at 10 mg/kg/day, co-

administration of i.v. PSCZ resulted in an approximate 12 µg/mL reduction in TEIC

trough levels. However, this study had certain limitations. The clinical sample size in

this study was limited, and large-scale prospective studies are needed to evaluate the

magnitude of this effect more accurately. In addition, an assessment of how these

plasma concentration changes affect the efficacy of TEIC should be conducted through

pharmacokinetic-pharmacodynamic analyses. Moreover, since SBECD is widely used

in injectable formulations beyond PSCZ, future studies should investigate potential

interactions between TEIC and these SBECD-containing products, as well as with drugs

possessing similar hydrophobic side chains.

Taken together, our findings suggest that the co-administration of i.v. PSCZ

containing SBECD and TEIC may reduce the protein binding of the latter, thereby

promoting its renal excretion. The findings of this study provide a new perspective for

assessing the risk of drug–drug interactions when using SBECD-containing

formulations.

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Figure legends

Fig. 1. Chemical structures of teicoplanin (TEIC, left) and sulfo-butyl ether-β-

cyclodextrin (SBECD, right).

Fig. 2. Trough plasma concentrations of TEIC in patients who underwent hematopoietic

stem cell transplantation (HSCT) during periods without posaconazole (PSCZ)

administration, following PSCZ oral administration, and following PSCZ intravenous

administration. The TEIC concentration/dose (C/D) ratios were obtained from 83

plasma concentration measurements across eight patients: 23 who did not receive

PSCZ, 27 who administered oral PSCZ, and 33 who received intravenous (i.v.) PSCZ.

Bars indicate the mean C/D ratio, and error bars represent standard deviations.

Fig. 3. Pharmacokinetics of TEIC in rats. Male Wistar/ST rats (8 weeks old) received an

intravenous injection via the femoral vein of saline (Control), PSCZ intravenous

formulation (i.v. PSCZ), or SBECD alone (SBECD). Thirty minutes later, TEIC was

administered intravenously. (A–D) The area under the plasma concentration–time curve

(AUC) and urinary excretion of TEIC were evaluated over 120 min after TEIC

administration. (E–L) The AUC and urinary excretion of individual TEIC components

were evaluated and expressed as relative values to the Control group. *$P < 0.05$, **$P <$

0.01, significantly different from the Control group (n = 6 per group).

Fig. 4. Estimated binding modes of TEIC A_{2-2} with SBECD. The lowest TEIC A_{2-2}

binding energy conformations for SBECD in the most probable docking clusters from

100 runs. The dimethyl terminus of the TEIC A_{2-2} side chain is directed towards the

primary face of the SBECD molecule in Type I (A) and is directed towards the

secondary face of the SBECD molecule in Type II (B). The left, middle, and right

panels show the front, side, and back views as the chemical structure of TEIC in Figure

1, respectively.

Fig. 5. Unbound fraction of TEIC in human serum. Human serum containing various

concentrations of SBECD was supplemented with TEIC. The unbound fraction of each

TEIC component was determined. *$P < 0.05$, **$P < 0.01$, significantly different from

the group without SBECD ("0") (n = 3 per group).

Table 1. Characteristics of patients treated with teicoplanin and posaconazole

Dose of teicoplanin, mg/kg/day (mean ± S.D.)	9.95 ± 3.64
Male/female, n	4/4
Age, years (mean ± S.D.)	45.9 ± 11.0
Serum albumin, g/dL (mean ± S.D.)	2.92 ± 0.45
Estimated glomerular filtration rate, mL/min/1.73 m^2 (mean ± S.D.)	101.5 ± 46.3
Number of data points, n	
Without posaconazole administration	23
Administered oral posaconazole	27
Administered intravenous posaconazole	33

505 S.D., standard deviation. Posaconazole was administered at a dose of 300 mg/day

506 regardless of body weight or route of administration.

Fig. 1

Teicoplanin (TEIC)   Sulfo-butyl ether-β-
                                  cyclodextrin (SBECD)

Fig. 1. Chemical structures of teicoplanin (TEIC, left) and sulfo-butyl ether-β-cyclodextrin (SBECD, right).

Fig. 2

Fig. 2. Trough plasma concentrations of TEIC in patients who underwent hematopoietic stem cell transplantation (HSCT) during periods without posaconazole (PSCZ) administration, following PSCZ oral administration, and following PSCZ intravenous administration. The TEIC concentration/dose (C/D) ratios were obtained from 83 plasma concentration measurements across eight patients: 23 who did not receive PSCZ, 27 who administered oral PSCZ, and 33 who received intravenous (i.v.) PSCZ. Bars indicate the mean C/D ratio, and error bars represent standard deviations.

Fig. 3

Fig. 3. Pharmacokinetics of TEIC in rats. Male Wistar/ST rats (8 weeks old) received an intravenous injection via the femoral vein of saline (Control), PSCZ intravenous formulation (i.v. PSCZ), or SBECD alone (SBECD). Thirty minutes later, TEIC was administered intravenously. (A–D) The area under the plasma concentration–time curve (AUC) and urinary excretion of TEIC were evaluated over 120 min after TEIC administration. (E–L) The AUC and urinary excretion of individual TEIC components were evaluated and expressed as relative values to the Control group. *$P < 0.05$, **$P < 0.01$, significantly different from the Control group (n = 6 per group).

Fig. 4

Front Side Back

A

Type I

B

Type II

TEIC A$_{2-2}$ SBECD

Fig. 4. Estimated binding modes of TEIC A$_{2-2}$ with SBECD. The lowest TEIC A$_{2-2}$ binding energy conformations for SBECD in the most probable docking clusters from 100 runs. The dimethyl terminus of the TEIC A$_{2-2}$ side chain is directed towards the primary face of the SBECD molecule in Type I (A) and is directed towards the secondary face of the SBECD molecule in Type II (B). The left, middle, and right panels show the front, side, and back views as the chemical structure of TEIC in Figure 1, respectively.

Fig. 5

Fig. 5. Unbound fraction of TEIC in human serum. Human serum containing various concentrations of SBECD was supplemented with TEIC. The unbound fraction of each TEIC component was determined. *$P < 0.05$, **$P < 0.01$, significantly different from the group without SBECD ("0") (n = 3 per group).