Feasibility of Trypsin Digestion as a Sample Preparation for Daptomycin Quantification in Murine Skeletal Muscles

Yusuke Sakai,⁎ Eriko Murakami,⁎ Hideo Kato, Kaname Ohyama, Mao Hagihara, Hiroshige Mikamo, and Bunji Uno⁎⁎

⁎ The United Graduate School of Drug Discovery and Medical Information Sciences, Gifu University; 1–1 Yanagido, Gifu 501–1193, Japan; ⁎ Gifu Pharmaceutical University; 1–25–4 Daigaku- nishi, Gifu 501–1194, Japan; ⁎⁎ Department of Pharmacy, Aichi Medical University School of Hospital; 1–1 Yazakokarimata, Nagakute, Aichi 480–1195, Japan;

† Department of Pharmacy Practice, Graduate School of Biomedical Sciences, Nagasaki University; 1–7–1 Sakamoto-machi, Nagasaki 852–8501, Japan; and ⁎ Department of Infection Control and Prevention, Aichi Medical University School of Hospital; 1–1 Yazakokarimata, Nagakute, Aichi 480–1195, Japan.

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It is important to evaluate the amount of daptomycin (DAP) distributed to skeletal muscles to elucidate the mechanisms related to penetration and side effects, such as myopathies. However, no attempt has been made to measure DAP concentrations in skeletal muscles. The study’s aim is to investigate the feasibility of trypsin digestion, as a muscle sample preparation technique for the determination of DAP in murine skeletal muscle, was evaluated in conjunction with a conventional HPLC-UV analysis. Compared with trypsin digestion, DAP was less recovered from spiked skeletal muscle by the conventional extraction, including homogenization, centrifugation, and filtration, because of its incorporation into the muscle protein. On the other hand, a sample preparation technique involving enzymatic digestion employing trypsin fully recovered DAP from the spiked skeletal muscle. Based on the spike recovery assay results, we proposed an efficient muscle sample preparation method involving trypsin digestion. HPLC analysis in conjunction with the sample preparation method has successfully determined DAP concentrations of skeletal muscles collected from mice administrated subcutaneously with DAP. The proposed method is suitable for application to investigations that include animal experiments on drug migration into muscle and mechanism underlying skeletal muscle injury as a side reaction, such as myopathies, of DAP therapy.

Key words  daptomycin; murine skeletal muscle; trypsin digestion

INTRODUCTION

Daptomycin (DAP) is a Streptomyces roseosporus-derived lipopeptide antibiotic that exhibits rapid and concentration-dependent bactericidal activity against Gram-positive bacteria including methicillin-resistant Staphylococcus aureus.1–3 A few studies have identified unique bactericidal mechanisms of DAP, as well as so-called calcium-dependent antibiotic, such as permeabilization and depolarization of the bacterial cell membrane,4,5 being different from structurally similar lipopeptides such as tushimycin as well as amphomycin and laspartomycin that inhibit peptidoglycan biosynthesis.4,5 However, the activity mechanisms involved in penetration of DAP in tissues, such as skin and muscle, are not fully understood. On the other hand, it is well recognized that the characteristic toxicity of DAP therapy is caused by myopathies.6–9 Specifically appears on the skeletal muscle,8 and is induced by an overdose8 or short interval administration of DAP.9 As can be seen in a few case reports, there is also a concern about rhabdomyolysis, which is rare but serious, as a side effect of DAP therapy.10–13 However, few factors that cause side effects have been identified. Kostrominova et al. suggested that the early effects of DAP on skeletal muscles of rats are due to the loss of sarcolemmal integrity based on the results of the histopathological examination which used immunostaining technique for the muscle injury of rats that were administered with DAP.10–13 Once-daily dosing of DAP has also proved valid to minimize skeletal muscle effects.14 Therefore, a method for accurate quantitation of DAP in skeletal muscle is essential for basic research using animals to study drug penetration and distribution properties to the muscle and the mechanism of the skeletal muscle injury as a side effect of DAP therapy.

Analytical studies on the quantification of DAP have been conducted to develop a method for the determination of DAP in plasma and serum that uses HPLC and ultra-HPLC with UV detection,15–18 LC-MS,19 and LC-tandem MS.20,21 These methods rely on quite similar approaches to sample preparation for plasma and serum matrixes, such as solid-phase extraction and deproteinization with methanol or acetonitrile followed by centrifugation.15–21 However, data on the tissue distribution of DAP are sparse and are confined to a pharmacokinetics study using rat skin tissue22 and a clinical study using crushed human bone.23 The centrifuged supernatants obtained from homogenization of the tissues followed by lengthy and repeated extraction using aqueous buffer solutions were then used for HPLC measurements of DAP in the tissues.22,23 However, few details of the extractability of the drug from the homogenized skin-tissue suspension and crushed human bone before HPLC analysis are available.

We tried to determine the amount of DAP in muscle samples after administration in a preliminary experiment conducted on the use of a DAP loading dose in a murine infection model.24 However, the peak corresponding to DAP was al-

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most undetectable on the chromatogram of the muscle sample obtained by the conventional extraction with homogenization and sonication, which is widely accepted as a sample preparation method for the analysis of a drug in biological solid matrixes. Therefore, a more effective pretreatment method for a skeletal muscle sample might enable us to quantify DAP distributed in the muscle by HPLC easily and accurately. As far as we know, no attempt has been made to measure DAP concentrations in skeletal muscles.

In this study, the feasibility of trypsin digestion as an efficient sample preparation technique for the bioanalysis of DAP in murine skeletal muscles was evaluated to gain more insight into drug migration into muscles and the mechanism underlying the side effect of skeletal muscle injury. The proposed method was applied to the determination of DAP in femoral muscle samples collected from mice which were administered with single subcutaneous dose of DAP.

MATERIALS AND METHODS

Chemicals  DAP was purchased from Funakoshi Co., Ltd. (Tokyo, Japan) and was used without further purification. The DAP was subdivided into 200 µL sample tubes as a 100 µM aqueous solution and was stored in a freezer to achieve the desired concentration immediately before each experiment. Vancomycin hydrochloride from Streptomyces orientalis received from Sigma-Aldrich Co. (St. Louis, MO, U.S.A.) was used. The 100 µM stock solution of vancomycin was prepared and stored in a freezer. Analytical grade 4-nitrophenol was commercially available from Kishida Chemical Co., Ltd. (Osaka, Japan) and was used without further purification as an internal standard for HPLC analysis. HPLC-grade acetonitrile used for HPLC eluent was purchased from Kanto Chemical Co., Inc. (Tokyo, Japan) and was used without further purification. HPLC-grade methanol (500 µL) was purchased from Sigma-Aldrich Co. (St. Louis, MO, U.S.A.) was used. Methanol (600 µL) and chloroform (300 µL) were added to the tube (2 mL) containing exactly weighted muscle homogenate, which was then sealed with a cap and mixed on a vortex mixer for 10 min. Subsequently, the contents of the tube were vortex mixed with 300 µL chloroform and 400 µL water for 5 min to extract the drugs into the aqueous phase. The supernatant’s aqueous layer separated from fat-soluble components, which is known as the Bligh–Dyer extraction method. We used this method as the conventional extraction procedure, as illustrated in Fig. 1 (P1) for the thorough and detailed experimental procedures. Methanol (600 µL) and chloroform (300 µL) were added to the tube (2 mL) containing exactly weighted muscle homogenate, which was then sealed with a cap and mixed on a vortex mixer for 10 min. Subsequently, the contents of the tube were vortex mixed with 300 µL chloroform and 400 µL water for 30 s and then centrifuged at 3000 × g for 5 min to extract the drugs into the aqueous phase. The supernatant water (1.0 mL) was aspirated by using a pipette, followed by evaporation to dryness by using an evaporator. The residue dissolved in 50 µL of water containing an internal standard was applied to an Ultrafilter-MC (0.22 µm pore size) filter.

After 10 mice were euthanized by CO2 asphyxiation without administering drugs, the thighs were immediately excised, followed by homogenization with 2 mL of sterile iced saline (0.9% (w/v)), and was used as analyte-free femoral muscle for drug recovery experiments. The homogenized analyte-free muscle was cut into pieces weighing approximately 0.2 g (wet weight); one piece was added to a pre-weighed 2 mL tube (Watson), and the exact weight was determined. The tube was stored at −80°C until use. Another eight mice were injected subcutaneously with 100 mg/kg DAP in 200 µL of saline, and five of them and the other three were sacrificed at 2 and 1 h, respectively, after drug administration to collect the femoral muscles. The muscles were harvested in the same manner as the analyte-free muscles and were frozen at −80°C until analysis.

Conventional Extraction Procedure Conventional liquid–solid extraction coupled with homogenization or sonication is still widely accepted as the most practical sample preparation method for the analysis of organic compounds, such as drugs, in biological solid matrixes. It is well recognized that the use of a mixed solution consisting of chloroform and water efficiently extract water-soluble drugs, such as DAP and vancomycin hydrochloride, into the supernatant’s aqueous layer separated from fat-soluble components, which is known as the Bligh–Dyer extraction method. We used this method as the conventional extraction procedure, as illustrated in Fig. 1 (P1) for the thorough and detailed experimental procedures. Methanol (600 µL) and chloroform (300 µL) were added to the tube (2 mL) containing exactly weighted muscle homogenate, which was then sealed with a cap and mixed on a vortex mixer for 10 min. Subsequently, the contents of the tube were vortex mixed with 300 µL chloroform and 400 µL water for 30 s and then centrifuged at 3000 × g for 5 min to extract the drugs into the aqueous phase. The supernatant water (1.0 mL) was aspirated by using a pipette, followed by evaporation to dryness by using an evaporator. The residue dissolved in 50 µL of water containing an internal standard was applied to an Ultrafilter-MC (0.22 µm pore size) filter.

Fig. 1. Experimental Protocol for HPLC Analysis of DAP in Skeletal Muscle Samples

Sample preparation procedures of the conventional extraction (P1) and those involving trypsin digestion (P2). (Color figure can be accessed in the online version.)
available from Merck Millipore by using a centrifuge set at 6000 \times g \text{ at } 4^\circ C \text{ for } 10 \text{ min. A } 20 \mu L \text{ aliquot of the filtrate was injected onto the HPLC column.}

**Extraction Procedure Coupled with Trypsin Digestion**

Trypsin-ethylendiaminotetraacetic acid (EDTA) solution without phenol red, containing 0.5 wt% trypsin and 5.3 mM EDTA-2Na, was purchased from FUJIFILM Wako Pure Chemical Corporation (Osaka, Japan) and used as a trypsin standard solution without further treatment. The optimal conditions for trypsin digestion of the murine skeletal muscles are shown in Fig. 1 (P2). The trypsin standard solution (150 \mu L) and 0.1 M Tris–HCl buffer (pH 8) solution (200 \mu L) were added to the tube (2 mL) containing exactly weighed muscle homogenate, which was then sealed with a cap and incubated at 37°C for 30 min. Subsequently, the contents of the tube were vortexed to yield a homogeneous suspension, followed by the conventional extraction, as illustrated in Fig. 1.

**HPLC Analyses**

A Model 10AD VP HPLC system (Shimadzu, Kyoto, Japan) equipped with a DGU-12A degasser, a CLASS M10A (SPD-MXA) photodiode array detector, and a DGU-12A degasser, a Class M10A system (Shimadzu) with mobile phases of 0.01 M phosphate buffer (pH 2.1) containing 10% acetonitrile and 0.01 M phosphate buffer (pH 7.2) containing 10% acetonitrile for DAP and vancomycin, respectively, at a flow rate of 1.0 mL/min at 40°C. The column effluent was monitored at 221 and 280 nm for DPA and vancomycin, respectively, and the peak analyses were performed by using a Class M10A system (Shimadzu).

**Calibration Curve**

A calibration curve was created by plotting the peak height ratios relative to an internal standard against the various concentrations of the standard DAP solutions. The limit of detection, defined as the lowest concentration that gave a signal-to-noise ratio of at least 3, was found to be 0.3 \mu M for DAP. The limit of quantification defined as the drug sample concentration meeting prespecified requirements for precision within 20% was found to be 0.75 \mu M. The calibration curve was linear across the examined dynamic range from 0.75 to 10 \mu M (n = 6, r = 0.992). The relative standard deviation (RSD) values of the UV responses of DAP in 6 trials were 12.86, 6.76, and 2.18% for standard samples containing 0.75, 2.5, and 7.5 \mu M DAP, respectively.

**RESULTS AND DISCUSSION**

**Recovery Using the Conventional Extraction Method from Spiked Murine Skeletal Muscles**

Water-soluble DAP and vancomycin hydrochloride were expected to be extracted into the aqueous layer of an extract solution from femoral muscle homogenates. We conducted the additional recovery assay of the muscle homogenates spiked with DAP and vancomycin as preliminary experiments. Figure 2 shows the dependence of recovery of the drugs upon time after spiking the muscles. Recovery was evaluated by measuring three replicates of the samples. The mean recoveries for vancomycin were 96–104% at all the times observed, whereas those for DAP exponentially reduced with the passage of time and converged to approximately 25%. This implied that DAP can be incorporated into skeletal muscle in the manner characterized by somewhat specific binding unrecoverable by solvent extraction ascribed to partition equilibrium. Although approximately 92% DAP binds to plasma protein,7 DAP is easily recovered from plasma by using the conventional pretreatments comprising deproteinization by methanol or acetonitrile, centrifugation, and extraction without the interference of the protein binding.15–20 The physicochemistry underlying the strong incorporation of DAP into the muscle remains currently unknown; however, it is really interesting from the perspective of clinical analytical chemistry.

**Enzymatic Skeletal Muscle Digestion with Trypsin**

To recover DAP incorporated into the skeletal muscles, we considered enzymatic digestion as an alternate tissue preparation technique, taking a hint from the efficient extraction of some lipophilic drugs from fat tissue digested with lipase.26 As trypsin specifically cleaves peptide chains at the carboxyl side of lysine or arginine, except when either is followed by proline,27 it is conceivable that trypsin does not inactivate lipopeptide DAP. It has been demonstrated that DAP is not susceptible to hydrolysis by trypsin despite that diverse proteases such as chymotrypsin and type I proteases hydrolyze it.28 Additionally, it is well recognized that trypsin is inhibited by calcium and magnesium ions. Therefore, we employed a trypsin solution containing EDTA as the most suitable enzyme for digestion of skeletal muscles.
Trypsin digestion conditions were optimized by using the homogenized samples of skeletal muscle left for 3 h at a refrigerator after infusion of a DAP solution. Figure 3a shows recoveries of DAP from the muscle samples plotted against amounts of the trypsin standard solution used for the muscle digestion at 37°C for 10 min. The use of a trypsin standard solution of >100 µL produces an exponential increase in the recovery of DAP. The reason for this phenomenon is still unknown, but it suggests that DAP is instantaneously released from the collapsed structure of some proteins through cleavage of specific peptide bonds. On the other hand, the dependence of recovery of DAP from the skeletal muscle upon digestion time is illustrated in Fig. 3b. Complete dissolution of 0.2 g of murine muscle was obtained by digestion with a 150 µL trypsin standard solution at 37°C for 30 min. We note that DAP responses start to decrease after 1 h of digestion, as shown in Fig. 3b. As a result, digestion of 0.2 g of murine skeletal muscle at 37°C for a period of 30 min was determined to be the optimal digestion conditions for a 150 µL trypsin standard solution, as shown in Fig. 1 (P2). In subsequent investigations, these conditions were employed.

**Recoveries from the Skeletal Muscles Spiked with DAP**

We found that trypsin treatment is very effective for the extraction of DAP incorporated strongly into skeletal muscles. We conducted experiments for recovery of DAP from femoral muscles spiked with DAP. Figure 4 shows the experimental procedures for pretreatments of the muscle samples. A 20 µL aliquot of 50 µM DAP solution was added to the homogenate of analyte-free femoral muscle (0.2 g), which was stored in a refrigerator for 3 h. The four samples were used for the conventional extraction experiments (Fig. 4A) and those involving trypsin digestion (Fig. 4C), respectively. In addition to this, the residual muscle after the conventional extraction was treated with trypsin followed by the conventional extraction again (Fig. 4B). The mean recovery data are also shown in Fig. 4. It is interesting that the sum of the collected amounts of A and B is comparable to the recovery amount of C, which implies that more than two-thirds of DAP present in skeletal muscle

![Fig. 3. Dependences of Recovery of DAP from the Muscle Spiked with DAP on the Amounts of Trypsin Standard Solution Used for Digestion (a) and on Incubation Time for Muscle Digestion (b)](image)

*The additional recovery samples were prepared by using analyte-free muscle (0.2 g) spiked with a 20 µL aliquot of 50 µM DAP standard solution, and treated with trypsin followed by the conventional extraction, as shown in Fig. 1. (a) Trypsin was used for muscle digestion involving incubation for 10 min at 37°C. (b) Incubation time was measured for muscle digestion with 150 µL trypsin standard solution at 37°C.*

![Fig. 4. Experimental Scheme for Recovery Assays from Skeletal Muscle Samples](image)

*The muscle samples were analyte-free femoral muscles spiked with 1.62 µg DAP. The measured mean values (n = 4) are shown in gray.*
strongly interacts with some muscle proteins and is difficult to extract without the digestive power of trypsin.

Table 1 shows the results of recovery assays for the repeated experiments performed using the homogenized femoral muscle samples infused at three different concentration levels in quadruplicate. The obtained recovery data, except for the case of the 0.064 µg addition, satisfy the acceptance criteria for biochemical assays. The results show that the enzyme digestion employed in the extraction process contributed to the high and stable recovery of DAP >0.8 µg/g (muscle) in murine skeletal muscle. Alternatively, a spike with 0.064 µg DAP gave an unacceptable deviation, as presented in Table 1. The amount of addition converted into the final analyte concentration to be injected into the HPLC was <0.8 µM, which is close to the quantification limit of the present HPLC-UV method (0.75 µM). Therefore, pretreatment involving trypsin digestion appeared to contribute to precise quantification of DAP in murine skeletal muscle under the assumption that DAP transferred to the muscle through drug administration is in the same state as that spiked into the muscle directly.

**Table 1. Recovery Data and the RSD of DAP from Murine Skeletal Muscles Spiked with 0.064, 0.16, and 1.62 µg DAP**

<table>
<thead>
<tr>
<th>Addition amounts (µg)</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>Mean</th>
<th>S.D.</th>
<th>RSD (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.064&lt;sup&gt;b)&lt;/sup&gt;</td>
<td>75.2</td>
<td>42.3</td>
<td>87.3</td>
<td>52.8</td>
<td>64.4</td>
<td>20.52</td>
<td>31.87</td>
</tr>
<tr>
<td>0.16&lt;sup&gt;c)&lt;/sup&gt;</td>
<td>86.3</td>
<td>84.4</td>
<td>103.1</td>
<td>94.3</td>
<td>92.0</td>
<td>8.54</td>
<td>9.28</td>
</tr>
<tr>
<td>1.62&lt;sup&gt;d)&lt;/sup&gt;</td>
<td>97.5</td>
<td>99.5</td>
<td>101.1</td>
<td>100.3</td>
<td>99.6</td>
<td>1.54</td>
<td>1.55</td>
</tr>
</tbody>
</table>

<sup>a</sup>) Data obtained from HPLC analysis after sample pretreatment P2 shown in Fig. 1. <sup>b)</sup>) Prepared by spiking a 20 µL aliquot of a 2.0 µM DAP standard solution. The concentration of this muscle sample is calculated as 0.32 µg/g (muscle). <sup>c)</sup>) Prepared by spiking a 20 µL aliquot of a 5.0 µM DAP standard solution. The concentration of this muscle sample is calculated as 0.80 µg/g (muscle). <sup>d)</sup>) Prepared by spiking a 20 µL aliquot of a 50 µM DAP standard solution. The concentration of this muscle sample is calculated as 8.0 µg/g (muscle). The mean value shown in Fig. 4C was obtained from these experiments.

**Table 2. DAP Concentrations in Murine Femoral Muscles Collected 2h after the Single Subcutaneous Administration of DAP (100 mg/kg)**

<table>
<thead>
<tr>
<th>Procedure&lt;sup&gt;a)&lt;/sup&gt;</th>
<th>Measured values of DAP (µg/g)&lt;sup&gt;b)&lt;/sup&gt;</th>
<th>S.D.</th>
<th>RSD (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1</td>
<td>2</td>
<td>3</td>
</tr>
<tr>
<td>A</td>
<td>1.1</td>
<td>1.3</td>
<td>1.4</td>
</tr>
<tr>
<td>B</td>
<td>1.5</td>
<td>1.9</td>
<td>2.2</td>
</tr>
<tr>
<td>C</td>
<td>2.5</td>
<td>3.0</td>
<td>3.5</td>
</tr>
<tr>
<td>(1 h)&lt;sup&gt;c)&lt;/sup&gt;</td>
<td>3.0</td>
<td>2.8</td>
<td>3.8</td>
</tr>
</tbody>
</table>

<sup>a)</sup>) Sample preparation procedures shown in Fig. 5. <sup>b)</sup>) Measured values are indicated by the amount (µg) of DAP contained in the 1g of muscle. <sup>c)</sup>) Data obtained from femoral muscle collected 1 h after administration of DAP.

**Fig. 5. Experimental Scheme for the HPLC Analyses of the Determination of DAP in Murine Skeletal Muscles Collected from Mice (n = 5) Administered a Single Subcutaneous Dose of 100 mg/kg DAP**
DAP determination in murine skeletal muscles, as is denoted P2 in Fig. 1. We conducted quantification experiments of DAP in the femoral muscles collected from mice administered with DAP according to the proposed method. Figure 5 shows the experimental scheme for pretreatments of the muscle samples collected 2 h after single subcutaneous administration of 100 mg/kg DAP, and the measured values are presented in Table 2.

Since all of the samples have individual differences and issues based on the localization of analytes in the muscle tissue, variability in the analysis results of in vivo muscle tissue samples is both expected and acceptable. The incurred DAP levels were found to be higher with the present method (C) than the values obtained by the muscle preparation using the conventional extraction (A). It was also found that the relationship in which the sum of A and B is equal to C is established for all the samples, respectively. This finding implies that trypsin treatment is essential to determine the precise concentration of DAP involving incorporation through interaction with some muscle proteins. On the other hand, the large changes in muscle concentration 1 to 2 h after administration may be expected as it is well recognized that DAP concentrations in plasma and serum show a clear decrease within a few hours.21,24) However, both concentrations of DAP appeared to be comparable, as listed in Table 2. These are only the results of a pilot experiment; further studies are needed using this proposed method to investigate the relationship between the drug concentration variances in plasma and muscle over time after administration. Although the dosage is more than amounts generally used in the administration experiments, the feasibility of DAP determination in the skeletal muscle is well illustrated.

CONCLUSION

The feasibility of enzymatic digestion as a sample preparation technique for bioanalysis of DAP in murine skeletal muscles was evaluated. Our results showed that trypsin digestion was an appropriate sample preparation method for analysis of DAP incorporated into skeletal muscle through interaction with some proteins. This sample preparation method involving trypsin digestion could be used to identify factors that might predispose some patients to develop DAP-related myopathies with the aid of a highly-sensitive analytical method for DAP. The strong incorporation into the muscle may be possibly associated with skeletal muscle injury of DAP therapy.

There are only a few attempts to employ enzymatic digestion in tissue sample preparation of drug molecules.26,29,30) Yu et al. have proposed that enzymatic digestion as an alternate tissue preparation technique for bioanalysis of drugs has extraction efficiency comparable to that of homogenization.29) However, as far as we know, there have been no reports evaluating the feasibility of enzymatic digestion as a release technique from some muscle proteins interacting with the drug. This is the first report on the pretreatment of muscle sample using trypsin digestion. Further work to estimate the relationship between DAP concentrations in the serum and the skeletal muscle, and to identify the muscle proteins strongly interacting with DAP by using this muscle sample preparation technique is ongoing.

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Conflict of Interest The authors declare no conflict of interest.

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