A miniaturized in vitro release method for investigating drug-release mechanisms

E. Ahnfelt, E. Sjögren, N. Axén, H. Lennernäs *

Department of Pharmacy, Uppsala University, Box 580, 751 23 Uppsala, Sweden

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ABSTRACT

We have evaluated a miniaturized in vitro method, based on the μDISS Profiler™ technique that enables on-line monitoring of drug release from a 21 μl sample with 10 ml of release medium. Four model drugs in eight clinically used formulations, including both solid and non-solid drug delivery systems, were investigated. The acquired data were compared with historical in vitro release data from the same formulations. Use of the Weibull function to describe the in vitro drug-release profiles allowed discrimination between the selected formulations with respect to the drug-release mechanisms. Comparison of the release data from the same formulation in different in vitro set-ups showed that the methodology used can affect the mechanism of in vitro release. We also evaluated the ability of the in vitro methods to predict in vivo activity by comparing simulated plasma concentration-time profiles acquired from the application of the biopharmaceutical software GI-Sim to the in vitro observations. In summary, the simulations based on the miniaturized-method release data predicted the plasma profiles as well as or more accurately than simulations based on the historical release data in 71% of the cases and this miniaturized in vitro method appears to be applicable for both solid and non-solid formulations.

1. Introduction

Successful drug development processes benefit from the combination of in vitro methods and in silico simulations that can predict the in vivo response (Lennernäs et al., 2014a). To enable accurate in vivo predictions from in vitro drug-release data, the drug must be released from the formulation by a similar mechanism both in vivo and in vitro (Larsen et al., 2013). There are standardized in vitro quality control release methods for oral formulations, as described for example in the US Pharmacopeia (USP). The USP experimental set-ups include the paddle, rotation and reciprocating cylinder apparatuses. However, these methods might not always be accurately predictive of the in vivo situation (Kostewicz et al., 2014b). Adjustments to the stirring rate, temperature and release medium (surfactant, ionic strength and pH) could alter the release rates to varying degrees, and information on this would be of value for accelerated and biorelevant in vitro release studies of modified-release formulations (Shen and Burgess, 2012). The use of a biorelevant medium that mimics the dynamic gastrointestinal fluid has also been shown to increase the predictability of in vitro methods (Klein, 2010; Kostewicz et al., 2014b).

The μDISS profiler™, a miniaturized dissolution apparatus that uses only small amounts of active pharmaceutical ingredient (API) and release medium, was developed to measure the solubility of APIs in early drug development (Fagerberg et al., 2010; Tsinman et al., 2009). Importantly, this apparatus continuously monitors absorbance, using a fiber-optic UV dip probe located centrally in a glass vial (Avdeef and Tsinman, 2008). The mini-IDR™ disc, an additional application for the μDISS profiler™, enables measurement of the intrinsic dissolution rate (IDR). Because the exposed area of the API is constant when measuring the IDR, it has been used to characterize dissolution profiles (Avdeef and Tsinman, 2008).

Theoretical modeling is a useful tool in pharmaceutical development; it is used to investigate and characterize the release of the API from the drug delivery system (DDS) and to determine which factors influence the release rate (Siepmann and Siepmann, 2012). This type of knowledge is also useful during the innovation
and development of novel DDSs where a specifically designed drug-release profile is needed to target specific pharmacodynamic and safety properties (Johannsson et al., 2011; Selen et al., 2014; Siepmann and Siepmann, 2012; Sjögren et al., 2014).

The primary objective of this study was to improve understanding of the novel, miniaturized, in vitro, µDISSTM-based release method and to examine its potential for predicting in vivo data. The in vivo predictive abilities of release data from this novel method and of historical in vitro release data were compared using in silico simulations and the mechanisms of API release from selected DDSs were investigated using the Weibull function.

2. Materials and methods

2.1. Chemicals and drug delivery systems

Four APIs (metoprolol, diclofenac, naproxen and carbamazepine (Sigma–Aldrich, Sweden), representing biopharmaceutics classification system (BCS) classes I and II (Wu and Benet, 2005), were investigated. Their physicochemical properties are listed in Table 1. The chosen drugs have typical physicochemical properties for drugs that are formulated in modified release dosage forms. The lipophilicity (log P) covers a three-fold range but the molecular weight covered a narrow range. Stock solutions were prepared by dissolving each API in ethanol (Soelvo, Sweden). Mono- and dibasic potassium phosphate, used to prepare phosphate buffer saline (PBS), was acquired from Riedel de Haën, Germany.

The in vitro release profiles of eight clinically used DDSs were investigated. Two of the DDSs were non-solid formulations (an oral suspension and a gel for dermal application), two were oral immediate-release (IR) tablets, and four were oral immediate-release (MR) tablets, and four were oral immediate-release (IR) tablets. The IR tablets contained metoprolol tartrate (Seloken™ 50 mg, Astra Zeneca, Sweden), naproxen (Pronaxen™ 250 mg, Orion Pharma, Sweden), carbamazepine (Tegretol™ 100 mg, Novartis, Sweden), and diclofenac (Voltaren™ 25 mg, Novartis, Sweden). The MR tablets contained metoprolol succinate (Seloken ZOCTM 25 mg, Astra Zeneca, Sweden), and carbamazepine (Tegretol Retard™ 200 mg, Novartis, Sweden). The non-solid DDSs were a diclofenac dermal gel (Voltaren™ gel 11.6 mg/g), and an oral suspension of carbamazepine (Tegretol Retard™ 200 mg, Novartis, Sweden). The non-solid DDSs were a diclofenac dermal gel (Voltaren™ gel 11.6 mg/g), and an oral suspension of carbamazepine (Tegretol Retard™ 200 mg, Novartis, Sweden).

2.2. Sample reservoirs for the modified µDISSTM-based release method

The dimensions of the mini-IDR™ disc were used as a starting point for developing a sample reservoir made of magnetic ferritic stainless steel, an Fe–Cr alloy (Olsson and Landolt, 2003). A small cylindrical cavity, 3 mm deep and 3 mm in diameter, with a sample volume of 21 µl was drilled centrally in the steel disc (Fig. 1). The opening was covered with a diffusion barrier (see description below) to retain the formulation sample in the reservoir. The diffusion barrier was held in place with a ring-shaped magnet placed centrally over the reservoir opening (Fig. 1).

2.3. Experimental design

A µDiss profiler™ (pION, USA) was used for all of the miniaturized in vitro release experiments (Fig. 1). The UV dip probe in the µDiss profiler™ automatically performs spectral scans at intervals of 200–720 nm and calculates the drug concentration as the area under the concentration-time curve (AUC) of the second derivative spectrum. The second derivate spectrum enhances the peaks and reduces baseline shifting of the UV spectrum, which reduces the effects of background turbidity (Avdeef et al., 2009; Bijlani et al., 2007). The drug release studies were performed in triplicate; for each experiment, a 21 µl sample was inserted into the sample reservoir and 10 ml of PBS buffer, pH 7.4, was added. A pH of 7.4 was chosen mainly because the method is intended for use with drugs delivered parenterally. However, it should be noted that the average ionization of the APIs at this pH was no different from that at the intestinal pH of 6.8. If not otherwise stated, the stirring rate was set at 400 rpm. A nylon mesh filter (pore size 150 µm), a polyvinylidene fluoride filter (pore size 0.45 µm) and a cellulose-based dialysis membrane (approximate pore size 0.003 µm) were used as drug diffusion barriers. These diffusion barriers were soaked in de-ionized water for 10 min at room temperature before use. During the drug release experiments, the temperature was kept at 37 °C using a water bath (Lauda, Germany).

2.4. In vitro release studies

In the miniaturized method, in vitro studies investigated the effect of the choice of diffusion barrier on the time taken for metoprolol and diclofenac (dissolved in de-ionized water) to reach equilibrium, and the effect of the stirring rate (100–400 rpm) on the mass transport of metoprolol and diclofenac across the diffusion barrier. The polyvinylidene fluoride filter (pore size 0.45 µm) provided appropriate mass transport values for these APIs and was use as the diffusion barrier in the stirring-rate hydrodynamic experiments.

All the solid formulations except metoprolol MR were crushed and sieved (Retsch, Germany) to prepare the material for the

Table 1

<table>
<thead>
<tr>
<th>API</th>
<th>Mw</th>
<th>S0</th>
<th>pKₐ</th>
<th>log P</th>
<th>BCS class</th>
</tr>
</thead>
<tbody>
<tr>
<td>Metoprolol</td>
<td>267.37</td>
<td>200°/700°C</td>
<td>9.7</td>
<td>1.88</td>
<td>I</td>
</tr>
<tr>
<td>Diclofenac</td>
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<td>0.375</td>
<td>3.8</td>
<td>4.51</td>
<td>II</td>
</tr>
<tr>
<td>Naproxen</td>
<td>230.27</td>
<td>0.115</td>
<td>4.4</td>
<td>3.18</td>
<td>II</td>
</tr>
<tr>
<td>Carbamazepine</td>
<td>236.28</td>
<td>0.256</td>
<td>14</td>
<td>2.45</td>
<td>II</td>
</tr>
</tbody>
</table>

BCS: biopharmaceutical classification system; log P: logarithm of the partitioning coefficient; Mw: molecular weight; pKₐ: acid dissociation constant; S0: solubility (mg/ml).

Fig. 1. A schematic illustration of the principles of the novel miniaturized in vitro release method. (1) The formulation sample is inserted into the sample reservoir. (2) The diffusion barrier is mounted and the ring-shaped magnet is attached to keep the diffusion barrier and the formulation sample in place. (3) The sample reservoir is placed in the glass vial and the release medium is added. The drug release is measured continuously with the µDISSTM UV-dip probe.
sample reservoir. The fraction of particles sized between 0.7 and 1.0 mm was collected and used for the in vitro release experiments. The matrix of the metoprolol MR tablet was dissolved in de-ionized water immediately prior to the experiment; within 5 min it disintegrated into small ethylcellulose-coated pellets (approximate size 0.5 mm) (Ragnarsson et al., 1987). Just before the start of the experiment, either the crushed particle fraction or the coated pellets were mixed with de-ionized water to create a slurry. 21 μl of the resulting slurry was inserted into the sample reservoir and covered with the diffusion barrier, in this case the nylon mesh filter (pore size 150 μm). The slurry was used to avoid air being trapped in the sample cavity and beneath the diffusion barrier, which could have reduced the mass transport of the API across the diffusion barrier. The non-solid DDSs (the gel and oral suspension) were inserted directly into the sample reservoir. All experiments were performed in triplicate.

Historical in vitro and in vivo data on the eight DDSs were collected from published reports (Al Ameri et al., 2012; Eldjura et al., 2004; Friedman et al., 2000; Gohel et al., 2009; Kovacevic et al., 2008; Löbenberg et al., 2005; Nan et al., 2012; Nishihata, 1987; Parsaei et al., 2002; Reiner et al., 2001; Sandberg et al., 1988; Yuksel et al., 2000; Zhang et al., 2011; Zhou et al., 1998) and obtained quantitatively with PlotDigitizer (Free Software Foundation Inc., MA, USA).

2.5. Data analysis

The nonlinear semi-log line of the software GraphPad Prism 6.04 (GraphPad Software, Inc., USA) was used to analyze the correlations between the pore size of the diffusion barrier and the time to equilibrium.

Many models are available for theoretically modeling drug-release mechanisms (Siepmann and Siepmann, 2008). However, the entire in vitro drug-release profile can be described by the Weibull function. In Eq. (1), the Weibull function has been slightly modified

\[ R(\%)_t = R(\%)_\text{max} \times (1 - \exp(-a t^b)) \]  

where \( R(\%)_t \) and \( R(\%)_\text{max} \) are the percentage of drug released at time \( t \) and the maximum percentage released, respectively; \( a \) and \( b \) are constants describing the drug-release profile, and \( t \) is the time (Papadopoulos et al., 2006). The Weibull function has been reported to be useful in comparing release profiles for the delivery of drugs from various matrices (Dash et al., 2010). The combination of Monte Carlo simulations and experimental release data has shown that the values of \( b \) in the Weibull function can provide an indication of the release mechanisms involved in the drug release (Papadopoulos et al., 2006).

In this study, Eq. (1) was fitted to the observed data using the average value of each time point for both the experimental and the historical data. The curve fitting was performed using the nonlinear least square regression in Phoenix 6 WinNonlin 6.3 software (Pharsight, USA). Estimated \( a \) and \( b \) values were then used to extrapolate the release profiles for complete drug release. The goodness of fit for each analysis was expressed as the absolute average fold error (AAFE), as described by Graham et al. (2012).

\[ \text{AAFE average folder} = 10^{\frac{\sum |\text{log}\text{predicted } \text{observed}|}{N}} \] (2)

AAFE describes the geometric mean ratio of the predicted and observed values and \( N \) describes the number of observations. An AAFE value of 1 indicates a perfect fit while a value of 2 indicates an average 2-fold difference from observed data points.

2.6. In silico simulations of in vivo response

In silico simulations based on the experimental and historical in vitro data were obtained using the gastrointestinal (GI) absorption model GI-Sim (Sjögren et al., 2013). Briefly, the GI-Sim simulates human GI physiology with nine simulated GI compartments. The GI compartments are connected in series to represent the stomach ( compartment 1), the small intestine ( compartments 2–7) and the colon ( compartments 8–9) (Yu and Amidon, 1999; Yu et al., 1996). Physiological parameters describing these GI compartments have been previously reported (Sjögren et al., 2013). The GI model is linked to a simulated systemic pharmacokinetic model, with up to three compartments, to enable simulations and predictions of the plasma concentration–time profiles. In GI-Sim drug particles and monomers flow from one GI compartment into the next. The particles either dissolve or grow, and the monomers partition into colloidal structures or are absorbed across the intestinal wall. The pH-dependent solubility of an ionizable compound in the lumen is calculated in GI-Sim according to the Henderson–Hasselbalch equation and the physiological pH of each GI compartment (Hasselbalch, 1916). The dissolution rate in GI-Sim is usually described by Fick’s law and the Nielsen stirring model (Nielsen, 1961); however, in this study, the in vitro experimental release profiles of the formulations were used for the appearance rate of free monomer in the GI compartments. The effective intestinal permeability for each API was used to describe the total serial transport process across the aqueous boundary layer adjacent to the intestinal wall and the small intestine membrane (Lennernäs et al., 1996; Sandberg et al., 1991; Sjögren et al., 2013; Tubic-Grozdanis et al., 2008). Absorption occurs in all intestinal compartments including the colon (there is no absorption in the stomach). The relevant physicochemical, biopharmaceutical, and pharmacokinetic properties of the investigated APIs and formulations are summarized in Table 2. The AAFE was used to analyze

| Table 2 Pharmacokinetic parameters used in in vivo simulations. |
|------------------|------------------|------------------|------------------|------------------|------------------|------------------|
| API             | CL (l/h)         | Vc (l)           | k12 (h⁻¹)        | k21 (h⁻¹)        | First pass extraction (%) | Peff (10⁻⁴ cm/s) |
| Metoprolol      | 54ᵃ             | 84ᵇ             | 6.7ᵃ             | 2.4ᵇ             | 30ᵇ             | 1.3ᵇ            |
| Diclofenac      | 16ᵇ             | 3.1ᵇ            | 4.4ᵇ             | 41ᵇ              | 60ᵇ              | 3.1ᵇ            |
| Naproxen        | 21ᵇ             | 4.0ᵇ            | 0.093ᵇ           | 0.14ᵇ            | 0ᵇ              | 8.5ᵇ            |
| Carbamazepine   | 1.1ᵇ            | 60²ᵇ            | 0.036ᵇ           | 0.24ᵇ            | 1.25ᵇ           | 4.3ᵇ            |

ᵇ Rani et al. (2004).
ᵈ Runkel et al. (1973).
ᵉ Yu and Amidon (1999).
ᶠ Sjögren et al. (2013).
ᵍ Tubic-Grozdanis et al. (2008).
the goodness of fit between the predicted and observed plasma concentrations (Graham et al., 2012).

3. Results

3.1. Effect of diffusion barrier in the miniaturized release method

There was a correlation between the pore size of the diffusion barrier and the time to reach equilibrium (time to reach plateau) in the sample reservoir and the release medium (Fig. 2). The equilibrium times for diclofenac across the nylon mesh filter, the polyvinylidene fluoride filter and cellulose-based dialysis membrane were 0.011 h, 4.3 h and 6.8 h, respectively. For metoprolol, the equilibrium times were 0.011 h, 18 h and 25 h, respectively. The correlation between equilibrium time, y (h), and pore size, x (μm) was described as $y = 3.4 - 1.5 \times \log x$, $R^2 = 0.98$, and $y = 13 - 5.4 \times \log x$, $R^2 = 0.96$, for diclofenac and metoprolol, respectively. The time to equilibrium for the nylon mesh filter (with a pore size of 150 μm) was thus more than a hundred times shorter than that for the other two diffusion barriers: the polyvinylidene fluoride filter (with a pore size of 0.45 μm) and the cellulose-based dialysis membrane (with a pore size of 0.003 μm).

The influence of stirring rate on the API mass transport as a mean value of three observations, with the polyvinylidene fluoride filter as diffusion barrier, is shown in Fig. 3. The mass transport of diclofenac across the diffusion barrier was increasing with higher stirring rate. However, there was no correlation between the stirring rate and the mass transport of metoprolol. After 1 h of stirring at 100, 200 or 400 rpm, the mass transport of metoprolol across the diffusion barrier was 76%, 86%, and 77%, respectively.

3.2. Indicated mechanisms of drug release

Interpretations of the $b$ values in the Weibull function are summarized in Table 3 (Papadopoulou et al., 2006). The drug release profiles for each of the investigated DDSs are shown in Fig. 4 and the corresponding $a$ and $b$ estimates are listed in Table 4.
**Fig. 4.** *In vitro* drug release using data acquired with the novel, miniaturized, μDISS™ *in vitro* release method and historical *in vitro* data. Symbols represent the observed data and lines represent WinNonlin curve fitting of the Weibull function. (A) Release profile for metoprolol from a slurry of modified-release (MR) pellets using data from the miniaturized method and historical data (Gohel et al., 2009; Sandberg et al., 1988). (B) Release profile for metoprolol from a slurry of immediate-release (IR) metoprolol using miniaturized-method data and historical data (Löbenberg et al., 2005). (C) Release profile for carbamazepine from a slurry of MR carbamazepine using miniaturized-method data and historical data (Friedman et al., 2000). (D) Release profile for carbamazepine from a slurry of IR carbamazepine using miniaturized-method data and historical data (Elqidra et al., 2004; Kovacevic et al., 2008; Nan et al., 2012). (E) Release profile for carbamazepine from a carbamazepine oral suspension using miniaturized-method data and...
The indicated types of drug release varied according to the methodology used. Drug release through a highly disordered space was indicated for one data set (IR carbamazepine) (Elqidra et al., 2004). Drug release governed by diffusion in a fractal or disordered substrate was indicated only by data for IR formulations, including both the miniaturized-method experimental data for IR naproxen, IR diclofenac and IR metoprolol and historical data for IR carbamazepine and IR metoprolol (Kovacevic et al., 2008; Löbenberg et al., 2005). Diffusion through a normal Euclidian space was the most plausible drug release mechanism for one historical set of MR carbamazepine data (Friedman et al., 2000). Diffusion through a normal Euclidian space with the contribution of another release mechanism was suggested by historical data for MR metoprolol (Gohel et al., 2009) and diclofenac gel (Gohel et al., 2009; Parsaeae et al., 2002), and by the miniaturized-method experimental results for MR carbamazepine, the oral carbamazepine suspension, and diclofenac gel. A first-order release rate was indicated by both historical data (Nan et al., 2012) and the experimental results for IR carbamazepine. A sigmoidal release curve, interpreted as indicating a complex release mechanism, was suggested by historical data sets for IR naproxen (Yuksel et al., 2000), IR diclofenac (two data sets) (Al Ameri et al., 2012; Nishihata, 1987) and MR metoprolol Sandberg et al. (1988), as well as by experimental data for MR metoprolol. The historical data for the oral carbamazepine suspension had too few reported data points (first data point at 80% released) to be analyzed in this manner.

### 3.3. In silico simulations of in vivo response

The simulations of the plasma profiles obtained with GI-Sim were based on the miniaturized-method (μDISS™) or historical in vitro release profiles (Al Ameri et al., 2012; Elqidra et al., 2004; Friedman et al., 2000; Gohel et al., 2009; Kovacevic et al., 2008; Löbenberg et al., 2005; Nan et al., 2012; Nishihata, 1987; Parsaeae et al., 2002; Reiner et al., 2001; Sandberg et al., 1988; Yuksel et al., 2000; Zhang et al., 2011; Zhou et al., 1998). The plasma-time concentration profiles and the AUCs, maximum concentrations \( C_{\text{max}} \) and times to \( C_{\text{max}} \) from the experimental and simulated data are shown in Fig. 5 and Table 5, respectively. The agreement between the predicted concentration-time plasma profiles and the observations was assessed using the AAFE with the respective values at the times of the observations. In addition, the predictive ability was evaluated from the agreement between the observed and simulated PK parameters \( AUC, T_{\text{max}} \) and \( C_{\text{max}} \). In general, the simulations based on the data from the miniaturized in vitro method were in good agreement with the observed plasma profiles, with AAFE values in the range of 1.03–1.53. For five of the seven investigated DDSs (71%), the in vivo predictions (i.e., the accuracy of \( AUC, T_{\text{max}} \) and \( C_{\text{max}} \) predictions) based on the miniaturized-method experimental data were equally or more accurate than the predictions based on historical in vitro release data. For the remaining two DDSs, MR metoprolol and IR carbamazepine, the in vivo prediction based on the miniaturized-method experimental data was less accurate than that based on two of the historical release data, with AAFE values of 1.53 and 1.31, respectively.

### 4. Discussion

Eight DDSs were investigated using a novel, miniaturized, in vitro drug-release method and in silico modeling. The benefits of this method include the ability to collect data continuously, the use of only small amounts of sample and release medium, and the historical data (Zhang et al., 2011) (too few data points to perform curve fitting). (F) Release profile for diclofenac from diclofenac gel using miniaturized-method data and historical data (Parsaeae et al., 2002). (G) Release profile for naproxen from a slurry of IR naproxen using miniaturized-method data and historical data (Yuksel et al., 2000). (H) Release profile for diclofenac from a slurry of IR diclofenac using miniaturized-method data and historical data (Al Ameri et al., 2012; Nishihata, 1987). In vitro drug release was measured with a 150 μm nylon mesh filter as diffusion barrier and 10 ml of PBS buffer, at a temperature of 37 °C, with a stirring rate of 400 rpm.
Fig. 5. Historical in vivo drug concentrations in plasma over time (CP OBS) and simulated (GI-Sim) in vivo plasma concentration-time profiles based on in vitro data obtained from the miniaturized method (µDISSSTM) (data acquired with a nylon mesh, pore size 150 μm) and historical sources. Symbols represent historical CP OBS, lines represent simulated plasma concentrations: (A) observed data from Sandberg et al. (1988), simulated red line data from Gohel et al. (2009), simulated green line data from Sandberg et al. (1988), and simulated blue line data from the miniaturized method. (B) observed data from Sandberg et al. (1988), red line data from Löbenberg et al. (2005) and blue line data from the miniaturized method. (C) observed data from Zhang et al. (2011), red line data from Friedman et al. (2000), and blue line data from the miniaturized method. (D)
Table 5
Pharmacokinetic parameters (AUC, $C_{\text{max}}, t_{\text{max}}$) from clinical observations (in bold) and in vitro-based in silico simulations (in italics).

<table>
<thead>
<tr>
<th>API</th>
<th>DDS</th>
<th>AUC (ng*h/ml)</th>
<th>$t_{\text{max}}$ (h)</th>
<th>$C_{\text{max}}$ (ng/ml)</th>
<th>AAFE</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>Naproxen</td>
<td>IR</td>
<td>1200000</td>
<td>3</td>
<td>68000</td>
<td>1.26</td>
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<td></td>
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<td>1300000</td>
<td>1.1</td>
<td>110000</td>
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</tr>
<tr>
<td></td>
<td></td>
<td>1300000</td>
<td>0.96</td>
<td>110000</td>
<td></td>
<td>Yuksel et al. (2000)</td>
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<tr>
<td>Metoprolol</td>
<td>IR</td>
<td>1300</td>
<td>1</td>
<td>170</td>
<td>1.11</td>
<td>Sandberg et al. (1988)</td>
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<tr>
<td></td>
<td></td>
<td>1300</td>
<td>1.9</td>
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<tr>
<td></td>
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<td>Diclofenac</td>
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<td></td>
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<td>3700</td>
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<td>4900</td>
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<td>Friedman et al. (2000)</td>
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<td>MR</td>
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<td>6</td>
<td>40</td>
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<td></td>
<td>800</td>
<td>12</td>
<td>47</td>
<td>1.30</td>
<td>Gohel et al. (2009)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>820</td>
<td>8.2</td>
<td>47</td>
<td>1.70</td>
<td>Sandberg et al. (1988)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>750</td>
<td>12</td>
<td>43</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Carbamazepine</td>
<td>Susp</td>
<td>31000</td>
<td>0.99</td>
<td>3100</td>
<td>1.03</td>
<td>Zhang et al. (2011)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>33000</td>
<td>1.4</td>
<td>3100</td>
<td>1.08</td>
<td>Miniaturized method</td>
</tr>
<tr>
<td></td>
<td></td>
<td>33000</td>
<td>1.6</td>
<td>3200</td>
<td></td>
<td>Zhang et al. (2011)</td>
</tr>
</tbody>
</table>

AAFE: absolute average fold error; API: active pharmaceutical ingredient; AUC: area under the concentration-time curve; $C_{\text{max}}$: maximum concentration; DDS: drug delivery system; IR: immediate-release tablet; MR: modified-release tablet; Susp: oral suspension; $t_{\text{max}}$: time to $C_{\text{max}}$.

The diffusion barrier investigation showed that the nylon mesh filter with 150 $\mu$m pores had only a minor impact on the equilibrium time (0.011 h), while the filters with smaller pores significantly increased the equilibrium time. There are several possible explanations for this observation. The pore size in the nylon mesh filter was presumably sufficiently large to facilitate the flow of fluids. Since the smaller pores reduced this fast fluid flow, the dominating mass transport mechanism across the dialysis membrane and the polyvinylidene filter may thus have been the slower molecular diffusion (Kostewicz et al., 2014b). The total available pore area determines the rate of mass transport when molecular diffusion is the main mechanism of transport. The two selected APIs, diclofenac and metoprolol, showed slightly different equilibrium times for the dialysis membrane which might be explained by different API affinity to the same dialysis membrane. Possibly, the positively charged metoprolol ($pK_a = 9.7$) is attracted to the negatively charged cellulose membrane while the negatively charged diclofenac is not ($pK_a = 3.8$). In all, care should be taken when choosing an appropriate diffusion barrier since the material may affect the equilibrium time. However, for DDSs such as emulsions, a dialysis membrane could prevent the formulation from spreading outside the sample reservoir, thereby reducing the pharmaceutical effect of the excipients on the UV measurements as well as better control of the available total surface area for release (Morais and Burgess, 2014). If the diffusion barrier influences the API mass transport, this effect needs to be evaluated and corrected for in future data analyses (Levy and Benita, 1990; Washington, 1989).

A stirring rate of 400 rpm allowed sufficiently fast mass transport across the nylon mesh to minimize any influence on the drug-release profiles. However, further hydrodynamic investigation of the influence of the stirring rate on mass transport in the miniaturized release method is needed and is currently on-going.

4.2. Interpretation of the in vitro drug-release mechanism from drug delivery systems

The use of the Weibull function, suggested by Papadopoulou et al. (2006) facilitates straightforward analysis of drug-release mechanisms and is applicable for a wide range of formulations and types of mechanism. The release of the drug in fractal or disordered matrices appears to be driven only by diffusion. Diffusion in irregular fractal spaces is slower than in a homogeneous Euclidian space (Kosmidis et al., 2003a). Diffusion in a homogeneous space
when another release mechanism also is involved indicates diffusion-driven release with additional input from case II transport. In case II transport, the relaxation of the tablet matrix polymers influences the movement of the APIs through the matrix (Kosmidis et al., 2003c). First-order drug release follows Fick’s law of diffusion (Papadopoulou et al., 2006). A sigmoidal release curve is indicative of a complex release mechanism, involving erosion-based release or the release of a relatively insoluble API (Kosmidis et al., 2003b; Papadopoulou et al., 2006).

The drug-release mechanisms for seven of the IR formulation data sets (see Table 4) were, in accordance with theory and formulation strategy, based on diffusion through a fractal/disordered matrix or diffusion according to Fick’s law (first-order release). In contrast, the data sets for IR naproxen and IR diclofenac suggested complex release mechanisms involving erosion or influence of the relative insolubility of the APIs (Kosmidis et al., 2003b).

The release kinetics for the individually coated pellets of MR metoprolol indicated almost zero-order release over 24 h (Sandberg et al., 1988). To provide a drug release mechanism that follows zero-order kinetics, the DDS must be designed to overcome Fick’s law of diffusion, in which the diffusion is non-linear over time due to the changes in surface area and diffusion length (Chidambaram et al., 1988). Therefore, a more complex mechanism is required; this can be achieved, for example, by coating the pellets, as seen with the ethylcellulose coating used in MR metoprolol (Sandberg et al., 1988; Varelas et al., 1995). For the three MR metoprolol data sets, the indicated release mechanisms were either a complex mechanism or diffusion-controlled in combination with case II transport. Hence, the indicated release mechanisms for MR metoprolol are in line with the reported approximate zero-order release (Sandberg et al., 1988).

For MR carbamazepine, the release mechanism indicated by the miniaturized-method data was different from that indicated by the historical data (see Table 4) (Friedman et al., 2000). API release was driven only by diffusion in the historical data set, while the miniaturized-method data set indicated the presence of diffusion with contribution from case II transport. When MR carbamazepine is used clinically, the patient is instructed not to crush the tablets. However, the prolonged in vitro release from the crushed tablet indicated that the individually coated pellets of the MR tablet remained intact (Kesarwani et al., 2007).

The difference in the rate of drug release from diclofenac gel between data obtained using the miniaturized method and historical data was probably due to the large difference in pore size (more than a 50,000-fold difference) between the diffusion barriers. Nylon mesh (150 μm pores) was used in the miniaturized method, whereas a cellulose acetate dialysis membrane with a molecular weight cut-off point of 1000 was used for collection of the historical data (Parsaei et al., 2002).

In the novel, miniaturized, μDISSTM in vitro method, only a representative proportion of each DDS was investigated whereas, in the historical release methods, an intact formulation unit such as a tablet was examined. While the surface area of the diffusion barrier was constant during the miniaturized-method experiments, the surface area of the particles was not constant. The historical in vitro release experiments differed in the applied release equipment and set-ups, such as stirring rate, release medium (e.g., differing in pH and/or the addition of surfactant), as well as in the volume of release medium used (Al Ameri et al., 2012; Elqidra et al., 2004; Friedman et al., 2000; Gohel et al., 2009; Kovacevic et al., 2008; Löbenberg et al., 2005; Nan et al., 2012; Nishihata, 1987; Parsaei et al., 2002; Sandberg et al., 1988; Yulkes et al., 2000; Zhang et al., 2011). Therefore, the in vitro release profile and consequently the indicated release mechanism varied for the investigated DDS. As an example, the four in vitro data sets for IR carbamazepine differed with regard to the release method and the release medium used, and consequently the b values ranged between 0.27 and 1.0. The DDS for IR metoprolol was the exception; the same release mechanism was indicated in both the miniaturized-method and historical data sets. As expected, the release mechanism differed between the IR, MR and non-solid drug delivery systems. In conclusion, there was wide variation in the indicated release mechanisms between the data sets. The differences in method set-ups seemed to affect the drug release to a greater extent than the differences caused by using a representative part of the DDS or an intact unit.

4.3. Gastrointestinal (GI) absorption simulation

In vitro data originating from both the miniaturized method and the historical investigations for four model drugs were inserted into the in silico model GI-Sim to predict the plasma concentration-time profiles in vivo. The in vitro release profiles as well as selected parameters listed in Tables 1 and 2 determined the predicted in vivo response. As shown in Fig. 4, the in vitro release method had an impact on the release profiles and consequently on the predicted plasma-time profiles. Based on the in vitro drug-release profiles, GI drug absorption was predicted to occur mainly from the proximal small intestine or throughout the whole intestine including the colon. The potential for colonic absorption will be relevant for DDSs with prolonged drug-release profiles (>3 h), such as MR metoprolol, MR carbamazepine, and IR carbamazepine, as the specified mean transit time to reach the colon in GI-Sim is 3.5 h (Sjögren et al., 2013). Overall, the experimental data acquired with the novel, miniaturized, in vitro μDISSTM method gave more accurate predictions of the plasma concentration profiles than the historical in vitro data. The predictions based on the miniaturized method data were equally or more accurate than the predictions based on historical in vitro release data for 71% of the investigated DDSs. This was the case for both IR and MR formulations, indicating the overall accuracy of the modified miniaturized method. However, for IR carbamazepine and MR metoprolol one historical in vitro release data set for each formulation gave more accurate predictions. The in vivo profile for MR carbamazepine was very well described by the in vitro data, while the in vitro data for IR carbamazepine were somewhat less predictive. This appears contradictory, as it could be assumed that the in vitro drug-release profile for an IR formulation would be more likely to reflect the in vivo release profile. The inaccurate prediction for the IR formulation was also observed for two historical data sets (Elqidra et al., 2004; Nan et al., 2012). The in vivo predictions based on the in vitro data set from Kovacevic et al. were, however, accurate (Kovacevic et al., 2008). It is noteworthy that, in this investigation of the IR formulation, only 50% of the carbamazepine was released in 2 h (Kovacevic et al., 2008). The administered dose was predicted to be incompletely absorbed throughout the small and large intestine as a consequence of this slow release. For MR metoprolol, the in vitro release of the three data sets are quite similar (see Fig. 4), however, the predicted plasma-time profiles (see Fig. 5) are not. One in vitro release data set (Gohel et al., 2009) have a slightly higher drug release compared to the miniaturized method which lead to a higher absorption of metoprolol before the colon compartment of GI-Sim. This might be explained by a delay in the predicted tmax when based on release data from the miniaturized method.

In this study, the combination of in vitro and in silico tools was a valuable asset for the evaluation of the in vivo predictive performance of the in vitro results. This advantageous combination has been shown previously, and there is an ongoing effort to further improve these synergistic methodologies (Kostewicz et al., 2014a; Lennernäs et al., 2014b).
5. Conclusion

The developed in vitro release method based on the miniaturized µDiss™ technique enables continuous sampling with small volumes of both DDS (21 µl) and release medium (10 ml). In this experimental study, in vitro release data were collected from eight selected DDSs with four APIs. Collection of API release data was possible for both solid and non-solid formulations, as well as for charged and uncharged APIs. Both the miniaturized-method and the historical release data were used to estimate the release mechanisms and to predict the plasma concentration-time profiles in a biopharmaceutical GI absorption model (Al Ameri et al., 2012; Elqidra et al., 2004; Friedman et al., 2000; Gohel et al., 2009; Kovacevic et al., 2008; Löbengrin et al., 2005; Nan et al., 2012; Nishihata, 1987; Parsae et al., 2002; Reiner et al., 2001; Sandberg et al., 1988; Sjögren et al., 2013; Yuksel et al., 2000; Zhang et al., 2011; Zhou et al., 1998). The indicated release mechanisms for the APIs varied according to the source of the data. This discrepancy most likely originated from differences in the methods used, such as the stirring rate, the release medium used, and the in vitro apparatus used. For five of the seven evaluable drug formulations, predictions of the in vivo plasma concentration-time profiles obtained using the novel, miniaturized, µDiss™ in vitro data were at least as accurate as those obtained from the historical in vitro data. These promising results could reduce the amounts of the formulations and release media required in the collection of in vitro data, while still generating accurate in vivo predictions.

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