

Non-linear Mixed Effects Modeling and Simulation for Exploring Variability Sources in Dissolution Curves: A BCS Class II Case Example

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ABSTRACT

Purpose: Irbesartan is a BCS class II compound that exhibits pH- and buffer capacity-dependent dissolution behavior. The aim of this study was to apply non-linear mixed effects modelling on dissolution data of two immediate release products containing Irbesartan in order to characterize and quantify the sources of inter-dissolution profile variability.

Methods: Nonlinear mixed effects modelling was applied to describe the dissolution curves obtained for Irbesartan in three different pH-value media (1.2, 4.5, 6.8) with two different products (reference product: Aprovel® and a generic test product). Simulations performed and the impact of inter-dissolution variability was assessed.

Results: The % Irbesartan dissolved to time was found to follow a Weibull distribution. The population scale parameter was estimated 0.252 and the shape parameter was estimated 0.706. The pH-value of the dissolution medium was found to significantly affect the scale parameter, while the formulation was found to affect the shape parameter. Simulations showed that probably some discrepancies in the *in vivo* performance of the two products can be expected.

Conclusion: Through this case study the applicability and usefulness of nonlinear mixed effects modelling in oral drug formulation was highlighted and resides in its ability to identify and quantify sources of variability.

Keywords: Population modeling; Simulations; Dissolution; Irbesartan; BCS Class II

INTRODUCTION

Irbesartan is highly permeable but presents low aqueous solubility and thus belongs to BCS class II compounds. It has been claimed to present both weak acidic and weak basic properties [1-4]. In view of the complex environment of the gastrointestinal tract (GI) involving pH gradient and variation in the buffer capacity and ionic strength of GI fluids, BCS class II compounds present significantly different dissolution profile within each GI region [4,5]. As a result, dissolution studies in various pH media are highly informative of the expected *in vivo* performance [2,6], while some *in vitro* transfer models simulating the conditions and the residence time in within each region of the GI tract have also been proposed [5].

In vitro dissolution testing is a very important tool for drug development and quality control, as well as for investigation of bioequivalence [7-10]. Application of pharmacometrics, i.e., the quantitative evaluation and mathematical description of the *in vitro*

dissolution experiments, may offer significant insight regarding the underlying processes that take place, their kinetics and the sources of variability [11,12]. The main sources of variability in dissolution testing have been reported to be the analyst, the dissolution apparatus, the testing environment, the sample and naturally the product under investigation [8,9]. Various equations and mathematical approaches have been applied in order to characterize and study the dissolution profiles retrieved from *in vitro* dissolution studies either for research or for regulatory purposes [8,9,11,13-15]. Even a computer program has been developed that provides a model library for fitting dissolution data and facilitates the comparison of drug dissolution data [8].

However, in the case where there is a need to study, quantify and understand the sources of variability, nonlinear mixed effects modeling (nlme) is the more appropriate mathematical approach. Nlme consists of the identification of a model for the typical response (structural model), a model for heterogeneity which

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involves the predictable and unexplained reasons responsible for inter-individual variability and a model for uncertainty (residual error model) which describes why the previous two models do not match our observations exactly [16]. This technique has been applied, in a previous study with the aim to assess its contribution in the comparison of dissolution curves obtained for several batches of sustained release formulations of octreotide [13]. However, up to now it has not been applied clearly for the study of variability among dissolution curves.

The aim of the present study was to apply non-linear mixed effects modeling on dissolution data retrieved from two different products containing 300 mg of Irbesartan in order to describe its dissolution curves and quantify the impact of medium's pH and formulation on them. Secondly through this case study, the present work aims to highlight the advantages that nonlinear mixed effects modeling and simulation may offer in oral drug development.

METHODOLOGY

Dissolution data

Dissolution data were retrieved from tests designed for the *in vitro* comparison of pilot batches of a generic film coated tablet formulation containing 300 mg of Irbesartan with the reference product Aprovel® 300 mg. A dissolution USP II paddle apparatus was used, while the experiments were performed in three different mediums with pH values 1.2, 4.5 and 6.8. The dissolution experiments were performed in accordance to the EMA guideline on the investigation of bioequivalence [7]. Samples were collected at pre-defined time points and the % amount dissolved was calculated. In view of the fact that Irbesartan presents a pH-dependent dissolution behavior [1] a different sampling scheme was applied for studying the dissolution kinetics in each medium. Each experiment was replicated 12 times; as a result data from a total of 72 dissolution experiments were retrieved. Each dissolution experiment was considered as a separate individual, in order to perform a population analysis.

Nonlinear mixed effects modelling

Monolix Suite® 2019R2 (Lixoft, Orsay France) was used as software in order to apply non-linear mixed effects modeling. All model parameters were assumed to follow a lognormal distribution, and an exponential model was used to describe inter individual variability.

Structural model

The initial dose was set at 100, as the % dissolved was used as dependent variable and a nonlinear mixed effect model was sought for % dissolved (t). The models explored were the simple first order with only one parameter standing for a first order dissolution rate constant (kd), the Weibull model with a scale and shape parameter, Korsmeyer-Peppas (Power Law) with a kinetic constant and an exponent characterizing the diffusion mechanism and the logistic function with a scale, a shape and a location parameter. These models were selected as they are models that have been found to efficiently describe dissolution kinetics [8,9,14,15].

Modeling inter-individual and residual variability

The impact of formulation type (reference or test) and pH value of

the medium (1.2, 4.5 or 6.8) were explored as potential categorical covariates on model parameters. Statistical significance of these categorical was assessed by a one-way ANOVA. A Wald test and a Likelihood ratio test (LRT) were used in order to decide their inclusion in the model. A significance level of 5% was considered in all cases. Different error models of residual variability (constant, proportional, combined) were assessed based on the OFV but also on the model evaluation and validation criteria as analyzed below.

Model evaluation and validation

The models were evaluated both graphically (goodness of fit plots) and statistically (-2LL, Akaike and Bayesian information criteria). More specifically the goodness of fit plots inspected were primarily the individual fittings of the predicted profile given by the estimated individual model versus the observed data and the population and individual predictions overlaid on observations versus time. The predictive performance was assessed by prediction corrected visual predictive checks (VPCs) generated using Monte Carlo simulations of 1,000 datasets and 90% prediction intervals.

Simulations

Simulations were performed the expected range of dissolution profiles based on the estimated variability and have a more robust assessment of the dissolution behavior. Taking into account the inter-dissolution variability, a total of 500 dissolution profiles were simulated in each case and the effect of pH and formulation on the dissolution curves was explored more in depth without the need to perform any additional experiments.

The R function 'Simulx' included in the 'mlxR' package was used to perform the simulations.

RESULTS AND DISCUSSION

All dissolution data included in this analysis are presented in Figure 1. A population model able to describe the dissolution profiles of Irbesartan was successfully developed. A Weibull model was found to most adequately fit the data. The parameters estimated for the model are presented in Table 1. The accuracy and reliability of the parameter estimates obtained, was confirmed by the low % RSE values. The model described the data adequately as shown by the individual fittings of the predicted profile given by the estimated individual model versus the observed data (Figure 2) and by the individual predictions overlaid on observations (Figure 3A). The robustness and predictive capacity of the model can be noted by the VPC and the observed versus predicted plot (Figure 3B).

The Weibull model even though is an empirical equation, it can be applied to almost all kinds of dissolution curves [14]. It expresses the % cumulative amount dissolved to time by:

$$\% \text{dissolved (t)} = 100 * \left(1 - e^{-\left(\frac{t}{a}\right)^k} \right),$$

Where a, the scale parameter that defines the time scale of the process and k the shape parameter that characterizes the curve as either exponential (k=1), sigmoid (k>1) or parabolic (k<1) [14].

In the model developed in this study the pH value of the medium was found to significantly affect the scale parameter as described below:

$$a = a_{\text{pop}} \times \exp(-0.268)^{\text{cat}1} \times \exp(-0.773)^{\text{cat}2} \times \exp(\eta_a)$$

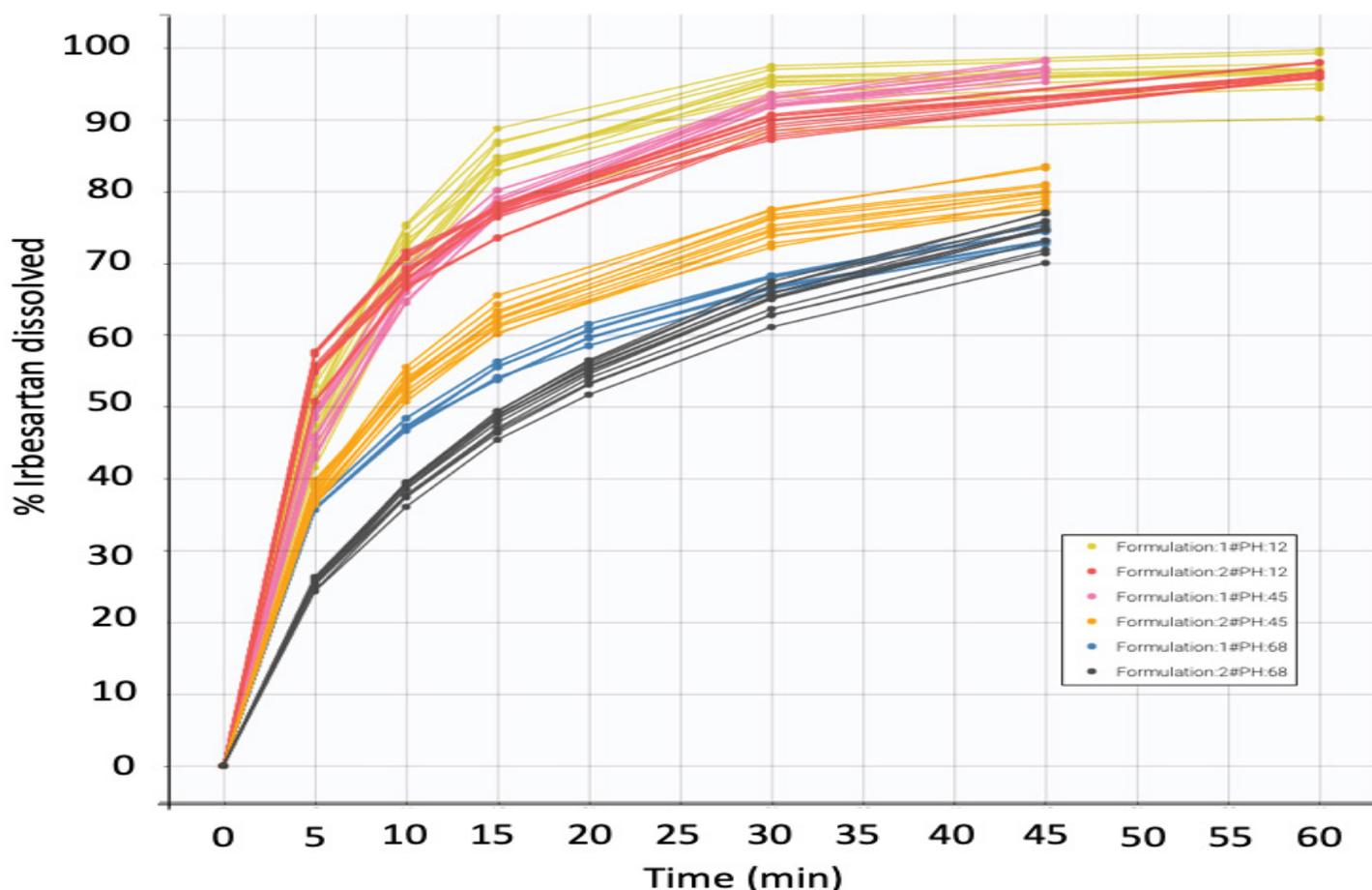


Figure 1: Spaghetti plot of the observed dissolution profiles per formulation and per pH of the medium used for the experiment. Therefore yellow represents the dissolution profiles retrieved with the reference product in a medium of 1.2 pH value, red with the test product in a medium of 1.2 pH value, pink with the reference product in a medium of 4.5 pH value, orange the test product in a medium of 4.5 pH value, blue with the reference product in a medium of 6.8 pH value and black with the test product in a medium of 6.8 pH value.

Table 1: Population estimates identified for the description of irbesartan’s dissolution kinetics following a Weibull distribution.

Fixed effects				
Variables	Estimate	SE	RSE (%)	p-value
a_pop	0.252	0.0111	4.41	~
beta_PH_45	-0.268	0.0273	10.2	0.00001
beta_PH_68	-0.773	0.0286	3.71	~
k_pop	0.706	0.017	2.41	~
beta_Formulation_Test	-0.134	0.0125	9.34	0.00175
Standard deviation of the random effects				
Variables	Estimate	SE	RSE (%)	
omega_a	0.344	0.0296	8.62	
omega_k	0.194	0.0167	8.6	
Correlations				
Variables	Pearson correlation coefficient	SE	RSE (%)	~~~~~
corr_k_a	-0.964	0.00861	0.893	
Proportional error model parameters				
Variables	Estimate	SE	RSE (%)	
Prop. (%)	1.80%	0.000826	4.4	

Where a_{pop} equals to the population parameter estimate, η_a represents the random effect, cat1 takes the value 1 when the medium has pH=4.5 and cat2 takes the value 1 when the medium has pH=6.8, otherwise they take the value 0. Evidently, the scale parameter equals to $0.252 \cdot \exp(\eta_a)$, $0.192 \cdot \exp(\eta_a)$ and $0.116 \cdot$

$\exp(\eta_a)$ for a pH medium of 1.2, 4.5 and 6.8, respectively. These findings are in line with the pH-dependent dissolution behavior of Irbesartan [1-4]. Also, it is evident from the faster dissolution rate, i.e., higher time scale parameter that Irbesartan’s basic properties prevail in aqueous media [3].

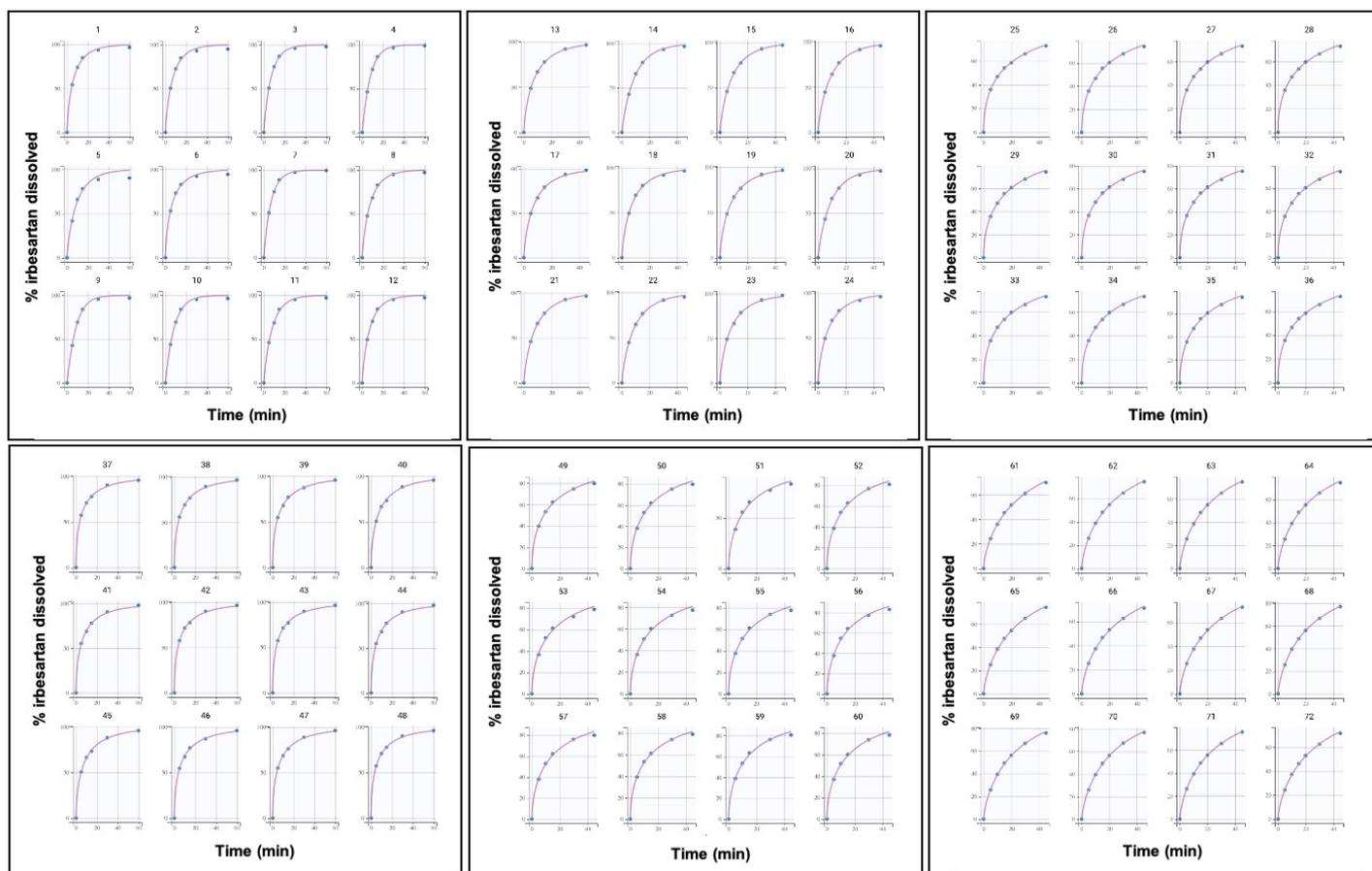


Figure 2: Fittings of the predicted profile given by the estimated individual model versus the observed data. The purple line represents the individual predicted profiles and the black closed circles the observed data.

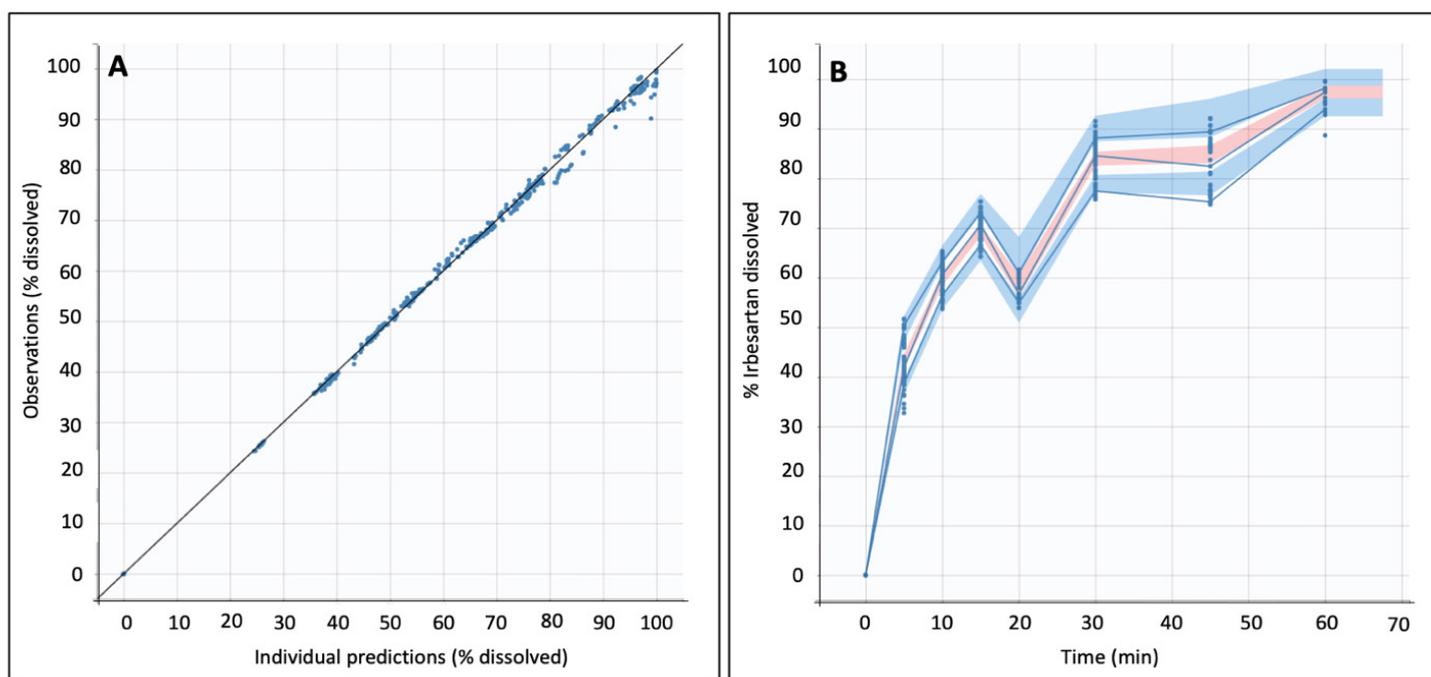


Figure 3: Goodness of fit plots of the final model developed for the % amount of irbesartan dissolved. On the left: Individual predictions overlaid on observations, On the right: Prediction corrected visual predictive check (VPC) of the Weibull model developed using 1000 Monte Carlo simulations. Median (solid line), 10th, 50th and 90th percentiles (blue line) of the observed data overlaid to the 90 % confidence intervals (colored areas) for the median, 10th, 50th and 90th percentiles of the simulated data.

The formulation type was found to significantly affect the shape parameter as described below:

$$k = k_{pop} \times \exp(-0.134)^{cat} \times \exp(\eta_k)$$

Where k_{pop} equals to the population parameter estimate, η_k

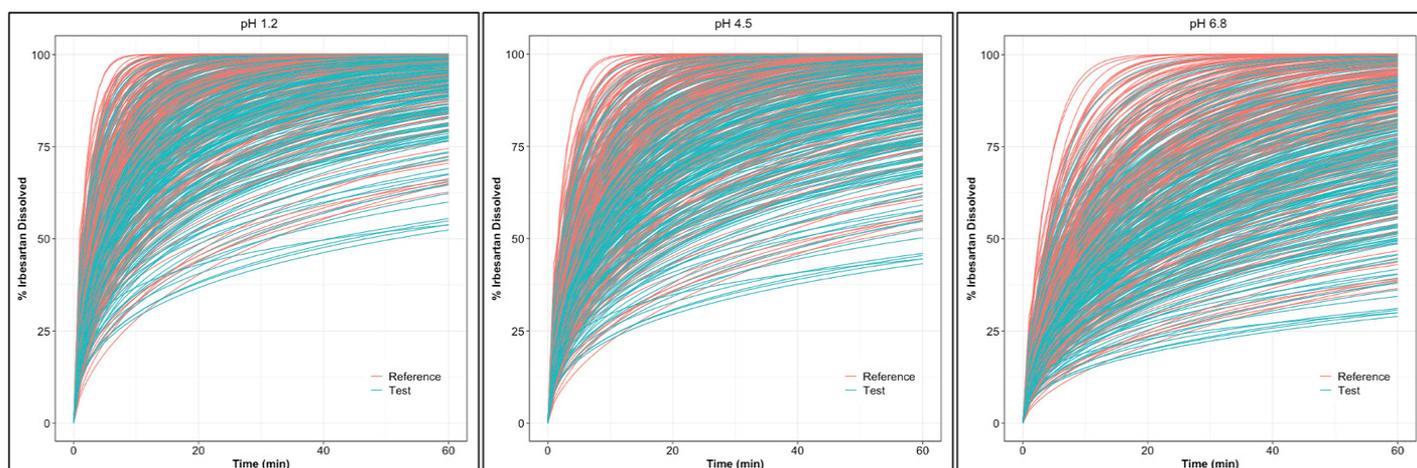


Figure 4: Simulated dissolution profiles ($n=500$) in three pH media, namely 1.2, 4.5 and 6.8. Red lines indicate profiles expected with the reference product (Aprovel®) and blue lines with the test product.

represents the random effect, cat takes the value 1 for the test formulation and 0 for the reference formulation. Therefore, for the reference product the shape parameter equals to $0.706 \cdot \exp(\eta_k)$ and for the test product to $0.617 \cdot \exp(\eta_k)$. This finding indicates that the curves present the same shape as expected.

A statistically significant inverse correlation between the random effects of the shape and scale parameter was also identified (Table 1), indicating that probably when the one grows the other lessens. This fact may lead to the assumptions that from a kinetic point of view there are some restrictions regarding the shape of the curve in relation to the time scale of the phenomenon.

Even though the Weibull model effectively characterized the dissolution profiles retrieved, this model doesn't include any kinetic fundament able to characterize the dissolution kinetic properties of the drug [14-16]. However, the combination of this equation to fit the data and nonlinear mixed effects modeling that is able to identify and quantify the sources of inter-profile variability, the effect of the formulation and the pH medium on the dissolution profiles were quantified.

Through simulations, it was noted that in all three pH the reference formulation presented a rather faster dissolution rate than the test formulation (Figure 4), indicating that probably *in vivo* some discrepancies of product performance should be expected. In addition, profiles obtained in media with higher pH values present a significantly higher variability, probably because Irbesartan's dissolution in acidic pH is faster. The significant impact of pH on dissolution of Irbesartan was noted both in simulated profiles of the reference and of the test product. Profiles obtained in media with pH 6.8 showed a much slower solubility than in pH 1.2 confirming the pH-dependent dissolution behavior of the compound and its basic properties.

CONCLUSIONS AND FUTURE WORK

Nonlinear mixed effects modeling and simulation help to gain a deeper understanding of the dissolution process as well as how various factors impact the dissolution curves. In the case example presented herein the impact of pH and formulation on dissolution curves of Irbesartan (BCS class II) were addressed. In fact, using these techniques, it was noted that due to inter-dissolution variability some differences in the *in vivo* dissolution and by

extension *in vivo* absorption should be expected. In this vein, it would be interesting to explore the impact of various excipients and/or their percentage in the formulation on dissolution kinetics. Nonlinear mixed effects modeling may constitute a valuable tool for formulation development that could provide guidance to the R&D department for formulation optimization.

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CONFLICTS OF INTEREST/COMPETING INTERESTS

The authors declare that they have no conflict of interest.

AVAILABILITY OF DATA AND MATERIALS

The data may be available upon request, even though that can be easily retrieved from Figure 1.

CODE AVAILABILITY

Monolix Suite® 2019R2 (Lixoft, Orsay France) was used.

AUTHORS' CONTRIBUTIONS

E.K. performed the modeling exercise, the simulations and contributed to the writing and editing of the manuscript. V.K. conceptualized the study, supervised the computational work and contributed to the writing and editing of the manuscript.

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