Development, Optimization and Validation of an HPLC-PDA Method for Quantification of Taxifolin in the Bark Extract of Pinus pinaster

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The aqueous extract of the bark of Pinus pinaster has a high concentration of polyphenols represented by a mixture of procyanidins, besides taxifolin, phenolic acids, cinnamic acids and their glycosides. Its quality control is specified in the United States Pharmacopeia, and the assay test is performed by determination of the total procyanidins content. However, determining the individual polyphenol content may represent an additional quality parameter for this extract. In this sense, the present study aimed to develop and optimize a method of high performance liquid chromatography with photodiode array detection (HPLC-PDA) for quantification of taxifolin in the bark extract of P. pinaster, using a 3³ Box-Behnken factorial design. The proposed method was validated for specificity, linearity, limits of detection and quantification, precision, accuracy and robustness and it has shown that taxifolin may be used as a chemical marker for quality control of the bark extract of P. pinaster.

Keywords: analytical validation, high performance liquid chromatography, Pinus pinaster, quality control, taxifolin

Introduction

Over the last decades, polyphenols have had an increasing interest on understanding vital functions of biological systems because they are important antioxidants of human diet.1 Due to the health significance of these compounds, some analytical methods have been developed for their separation, identification and quantification in natural products.2-4 Polyphenols have a wide variability of chemical structures, which differ in polarity and size, from simple phenolic compounds to oligomers. Additionally, they are found at low concentration levels in natural products.5 Therefore, sample pre-treatment is normally required before instrumental analysis.6 Liquid-liquid extraction (LLE) and solid phase extraction (SPE) are the sample pre-treatment techniques most frequently used. Among analytical techniques, high performance liquid chromatography (HPLC) and capillary electrophoresis (CE) are the main methods.7,8

P. pinaster Aiton, a pine commonly known as french maritime pine and originally occurring in the Mediterranean region,9 has been investigated with regard to its chemical constituents and biological effects.10 The aqueous extract of the bark of P. pinaster has a high concentration of polyphenols represented by a mixture of procyanidins (comprising catechin and epicatechin subunits with varying chain lengths), besides phenolic acids (gallic, ferulic, caffeic and p-hydroxybenzoic acids), cinnamic acids and their glycosides forms, and taxifolin (Figure 1).11,12 It is widely used as a nutritional supplement and provides numerous health benefits due to its antioxidant, anti-inflammatory, and immuno-modulatory effects.13-15 The quality control for the bark extract of P. pinaster is specified in the United States Pharmacopeia (USP) monograph in the chapter of dietary supplements.16 A method of HPLC with ultraviolet (UV) detection is described for identification test. It analyzes four components of the extract: caffeic acid, catechin, ferulic acid, and taxifolin. The method can also be used to generate chromatographic fingerprints that provide additional qualitative information. However, for quantitative purposes,
a method of spectrophotometry is described which provides an estimation of the total procyanidins content (65-75%) and it does not give quantitative measurement of individual polyphenol content. In this regard, taxifolin, a flavanonol also known as dihydroquercetin, can be a chemical marker for quality control of the bark extract of *P. pinaster*, because it is present in quantitatively detectable amount in the the extract when compared with other individual polyphenol contents. Furthermore, it is widely found in barks of the species within the genus *Pinus*, so it is an important flavanonol for this group.

Within this context, the aim of this work was to develop, optimize and validate a method of HPLC with photodiode array detection (PDA) for quantification of taxifolin in the bark extract of *P. pinaster*. The method optimization was investigated by a multivariate approach, taking into account a $3^3$ Box-Behnken factorial design.

**Experimental**

**Reagents, standards and sample**

Acetonitrile (Panreac, Spain) and ethyl acetate (Merck, USA) were HPLC grade, and formic acid was analytical grade (Neon, Brazil). Ultrapure water obtained in an AquaMax-Ultra 370 Series (Young Lin, Korea) ($18.2 \, \text{M} \Omega \text{cm}$ resistivity at 25 °C and < 10 ppb total organic carbon) was used throughout analysis. The standards of gallic acid (purity: 102.38%), catechin (purity: 99.00%) and taxifolin (purity: 87.40%) were purchased from Sigma (USA), and ferulic acid (purity: 102.00%) was from Henrifarma (Brazil). The pine bark extract (PBE) was obtained from USP (USA). The mobile phase was filtered through a 0.45 μm filter membrane (Sartorius, Germany) and degassed by an ultrasonic apparatus (Cristófoli, Brazil) for 30 min before use.

**Sample preparation**

The sample was prepared by LLE with ethyl acetate. The PBE (20 mg) was accurately weighed and transferred to a beaker in which 10 mL water/methanol (2:1, v/v) were added. The mixture was left in ultrasonic bath for 5 min. Then, the content of the beaker was poured into a separatory funnel of 100 mL and extracted with ethyl acetate (15 mL) by manual shaking for 1 min. After a 4 min rest period, the organic phase was separated from the aqueous phase. To the remaining aqueous phase, another portion of ethyl acetate was added (10 mL) and the procedure was repeated twice, making up three extractions. The combined organic phases were treated with 4 Å molecular sieves (3 g), and filtered to a round bottom flask through quantitative filter paper. The filtrate was concentrated to 1/15 of the initial volume by rotary evaporation under vacuum, transferred quantitatively to a 5 mL volumetric flask and made up to 5 mL with ethyl acetate. An aliquot of 3 mL was transferred to a 10 mL test tube and evaporated to dryness in a water bath at 50 °C. The residue was reconstituted in 2.6 mL acetonitrile, filtered in a 0.45 μm filter membrane and transferred to HPLC vials. Then, 20 μL were injected into the HPLC-PDA system.

**Instrumentation**

The HPLC analyses were performed using a qualified and calibrated Young Lin (Korea) chromatography system composed of: quaternary pump (YL 9110), photodiode array detector (YL 9160), automatic injector (YL 9150), column compartment (YL 9130) and software controller (Clarity). The HPLC-PDA method used was modeled in conformity with the method described in the USP monograph, according to the conditions available in our laboratory. Chromatographic separation was achieved using octadecylsilane (C18, 250 × 4.6 mm, 5 μm particle size) column (Agilent, Brazil) maintained at 40 °C and connected with a C18 pre-column (4.0 × 3.0 mm, 5 μm particle size) (Phenomenex, USA). The wavelength for UV detection was 288 nm. The initial HPLC gradient profile (mobile phase composition and flow rate) is shown in Table 1.

To optimize the chromatographic conditions, a $3^3$ Box-Behnken factorial design triplicated at the central
Development, Optimization and Validation of an HPLC-PDA Method


Identification of the compounds

Solutions of gallic and ferulic acids, catechin and taxifolin were prepared in acetonitrile at 50 μg mL\(^{-1}\). PBE samples (n = 3) were chromatographed and those peaks corresponding to commercially available standards were identified by UV spectra and spiking procedure.

Analytical validation

After method development and optimization, the validation tests were performed according to the International Conference on Harmonisation\(^{22}\) (ICH) and the Brazilian National Institute of Metrology, Standardization and Industrial Quality\(^{23}\) (Inmetro) guidelines, comprising the following parameters:

Specificity

The specificity of the method was determined through the comparison of standard, sample and blank chromatograms.

Linearity

The test was conducted from the plotting of three standard curves, each one constructed from five concentrations corresponding to 40, 45, 50, 55 and 60 μg mL\(^{-1}\) of taxifolin. The linear regression equation and the determination coefficient (R\(^2\)) were determined by least squares method. After regression implementation, the data for each concentration level were statistically evaluated taking into account homoscedasticity (Cochran’s test), residues’ normality (Shapiro–Wilk’s test) and lack-of-fit into the model through an \emph{a priori} test hypothesis (analysis of variance (ANOVA)). ANOVA test consists of comparing the deviations of the means between the calibration line (the residual standard deviation, s\(_{yx}\)) and the y values from their means (s\(_y\)) by using equation 1, where m\(_i\) is the number of measurements, p is the calibration point and m is the product between p and m\(_i\).

\[
F_{\text{calculated}} = \frac{S_y^2}{S_{yx}^2} = \frac{\sum_{i=1}^{p} \sum_{j=1}^{m_i} (\bar{y}_{ij} - \bar{y})^2 / (p - 2)}{\sum_{i=1}^{p} \sum_{j=1}^{m_i} (y_{ij} - \bar{y}_{ij})^2 / (m - p)}
\]

The test is carried out by comparison between F\(_{\text{calculated}}\) and F\(_{\text{critical}}\); \(f_1 = p - 2; f_2 = m - p\) (F\(_{\text{critical}}\)). If F\(_{\text{calculated}} \geq F_{\text{critical}}\), the linear model cannot be applied.

The taxifolin quantification was performed using the calibration curve from linearity of the standard.

Limits of detection and quantification

The limit of detection (LOD) and the limit of quantification (LOQ) were determined from three standard calibration curves and were calculated as shown in equations 2 and 3, respectively:

\[
LOD = S \times \frac{2}{a}
\]

\[
LOQ = S \times \frac{10}{a}
\]

Table 1. Initial HPLC gradient profile (mobile phase composition and flow rate)

<table>
<thead>
<tr>
<th>time / min</th>
<th>0</th>
<th>15</th>
<th>20</th>
<th>25</th>
<th>35</th>
<th>40</th>
<th>50</th>
</tr>
</thead>
<tbody>
<tr>
<td>Solution A / %</td>
<td>94</td>
<td>88</td>
<td>88</td>
<td>88</td>
<td>80</td>
<td>94</td>
<td>94</td>
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<tr>
<td>Solution B / %</td>
<td>6</td>
<td>12</td>
<td>12</td>
<td>12</td>
<td>20</td>
<td>6</td>
<td>6</td>
</tr>
<tr>
<td>Mobile phase flow rate / (mL min(^{-1}))</td>
<td>1.0</td>
<td>1.2</td>
<td>1.2</td>
<td>1.0</td>
<td>1.0</td>
<td>1.0</td>
<td>1.0</td>
</tr>
</tbody>
</table>

\*0.1% formic acid in ultrapure water; \*0.1% formic acid in acetonitrile.

Table 2. Factors and levels utilized in the experimental design for chromatographic condition optimization

<table>
<thead>
<tr>
<th>Experiment</th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
<th>7</th>
<th>8</th>
<th>9</th>
<th>10</th>
<th>11</th>
<th>12</th>
<th>13</th>
<th>14</th>
<th>15</th>
</tr>
</thead>
<tbody>
<tr>
<td>X(_1)*</td>
<td>−</td>
<td>+</td>
<td>−</td>
<td>+</td>
<td>−</td>
<td>+</td>
<td>+</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>X(_2)*</td>
<td>−</td>
<td>−</td>
<td>+</td>
<td>+</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>−</td>
<td>+</td>
<td>−</td>
<td>+</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>X(_3)*</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>−</td>
<td>−</td>
<td>+</td>
<td>+</td>
<td>−</td>
<td>−</td>
<td>+</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

\*Temperature of the column oven [(−): 35 °C, (0): 40 °C, (+): 45 °C]; \*mobile phase flow rate [(−): maintained at 1.0 mL min\(^{-1}\), (0): increased until 1.2 mL min\(^{-1}\), (+): increased until 1.4 mL min\(^{-1}\)]; \*increase in the percentage of acetonitrile in mobile phase [(−): until 15%, (0): until 20%, (+): until 25%].
where \( a \) is the regression slope and \( S \) is the standard deviation of the \( y \) intercept.

**Precision**

The repeatability (intra-day precision) was determined by analyzing six replicates consecutively by a single analyst in a single day. The intermediate precision (inter-day precision) was evaluated in the same way, but with two analysts and two days. An injection precision of less than 5% relative standard deviation (RSD) was considered appropriate.

**Accuracy**

Accuracy represents the level of compliance between individual results obtained and a reference value. In this current work, the accuracy of the method was established by recovery tests at three concentration levels (40, 50 and 60 \( \mu g \) mL\(^{-1}\)) performed in triplicate. For this, aliquots of 174, 217 and 260 \( \mu L \) of the taxifolin standard solution (concentration = 1000 \( \mu g \) mL\(^{-1}\)) were added before extraction.

The results were expressed as percentage of recovery (R%), and equation 4, where \( A_a \) is the analyte area, \( A_S \) is the standard area and \( A_{a+S} \) is the analyte area with standard addition, was used.

\[
R\% = \frac{A_{a+S} - A_a}{A_S} \tag{4}
\]

**Robustness**

The robustness parameter was evaluated by intentional minor modifications in the chromatographic conditions in the proposed method. Thus, a complete experimental design with eight experiments (2\(^3\)) and triplicate in the central point was conducted, in a total of 11 experiments performed in random order. The factors and their levels were: \( (X_1) \) temperature of the column oven [(-): 38 °C, (0): 40 °C and (+): 42 °C]; \( (X_2) \) mobile phase flow rate [(-): maintained at 1.0 mL min\(^{-1}\), (0) increased until 1.2 mL min\(^{-1}\) and (+) increased until 1.4 mL min\(^{-1}\)]; and \( (X_3) \) increase in the percentage of acetonitrile in mobile phase [(-): until 18%, (0): until 20% and (+) until 22%].

To evaluate the significance of deliberate variations in each chosen factor, their effects were calculated. From the results of each experiment, the coefficients for determining the statistical model of prediction were calculated according to equation 5:

\[
b = (X'X)^{-1}X'y \tag{5}
\]

where \( b \) is the matrix of model coefficients and \( X \) and \( y \) are the \( X \) matrix and \( y \) vector, respectively. To write the equation of the fitted model, the standard errors of the coefficients were calculated using equation 6:

\[
\varepsilon(b) = \sqrt{(X'X)^{-1}\sigma^2} \tag{6}
\]

where \( \varepsilon(b) \) is the matrix whose main diagonal represents the standard errors of the model estimators \( (b_i) \) and \( \sigma^2 \) is the population variance of the experiments, which can be estimated as \( s^2 \) using the center point replicates, from equation 7:

\[
s^2 = \frac{\sum(x_i - \bar{x})^2}{n-1} \tag{7}
\]

Effects were calculated in matrix by the product \( X'y \), where \( y \) is a column vector containing the average results of the assay.

To estimate the standard error of an effect, the square root of the value obtained in equation 8 was used, and the standard error of the mean was estimated using the square root of the value obtained in equation 9:

\[
\hat{\mu}_{\text{effect}} = \left(\frac{s^2}{2}\right)^{1/2} \tag{8}
\]

\[
\hat{\sigma}_{\text{effect}} = \left(\frac{s^2}{n}\right)^{1/2} \tag{9}
\]

With the estimated standard errors, it was possible to achieve confidence intervals for the values of effects, using Student’s \( t \)-distribution with 95% confidence, according to equation 10:

\[
\eta - t_v \times S_{\text{effect}} < \hat{\eta} < \eta + t_v \times S_{\text{effect}} \tag{10}
\]

where \( \eta \) is the true value of an effect (population value), \( \hat{\eta} \) represents the value obtained from the tests performed on the experiment, \( t_v \) is the value from the Student’s distribution, and \( S_{\text{effect}} \) is the standard error of an effect.

**Results and Discussion**

Optimization of the HPLC-PDA method

The selection of chemical markers is determinant for the quality control of herbal medicines, thus, in order to develop a method suitable for quality control of PBE, it is necessary to establish one or more chemical markers. Initially, a chromatographic fingerprint was performed using USP HPLC method, and four compounds were identified: gallic acid, catechin, taxifolin, and ferulic acid,
in this elution order. However, low signal intensity was obtained for these substances in our laboratory by UV detection, and the chromatographic profile showed low signal-to-noise ratio for these peaks. Therefore, it was evidenced which sample pre-treatment would be required. As taxifolin showed the most intensive signal, it was defined as chemical marker for method development, thus, the sample preparation was performed in order to ensure the best conditions for its extraction. After liquid-liquid extraction, the concentration found for taxifolin was 1.10 ± 0.11% at 95% confidence interval.

Although taxifolin can be determined in natural products by several analytical techniques, HPLC-UV was established as the most convenient method for qualitative and quantitative analysis of phenolic compounds. Despite this, the complexity of PBE has sustained the development of HPLC methods with mass spectrometry (MS) detection such as the method proposed by Chen et al. for HPLC fingerprints analysis using the most prominent peak (taxifolin) as the reference peak. The MS detection, however, is expensive and requires more trained analysts, so a method for taxifolin determination was developed using UV detection, the most widespread detection in quality control laboratories.

PBE is a natural source of phenolic compounds. Thus, reverse chromatography with C18-stationary phases combined with binary elution systems containing an aqueous acidified solvent and an organic modifier solvent (methanol and acetonitrile, mainly) are used almost exclusively. In this study, acetonitrile was chosen because taxifolin and ferulic acid were not separated using methanol in the preliminary investigations, besides the latter provided higher pressure values than acetonitrile. Additionally, it was necessary to acidify both solvents (aqueous and organic) with formic acid to minimize peak tailing.

Within this context, the mobile phase was initially established as two solvent systems: (i) 0.1% formic acid in ultrapure water (A) and (ii) 0.1% formic acid in acetonitrile (B). The initial HPLC gradient profile (mobile phase composition and flow rate) was established after screening tests. A gradient of mobile phase flow rate in the first 20 min was also realized in order to reduce analysis time. In this regard, a Box-Behnken factorial design was performed to optimize chromatographic conditions. Figure 2 shows the representative chromatograms obtained from factorial design.

The taxifolin response (Y) took into account the resolution of the critical pair and the baseline disturbances, common to the gradient elutions. Therefore, Y was defined as the ratio between the resolution (R) of the critical pair (value obtained between the taxifolin peak and the peak indicated by an asterisk in Figure 2) and a constant (K; defined as 2 for the presence of baseline disturbances and 1 for the absence of baseline disturbances), according to equation 11:

\[ Y = \frac{R}{K} \]  

where R was calculated using retention time (t_{r1}) and peak width at baseline (t_{wb1}) as following (equation 12):

\[ R = \frac{2(t_{r2} - t_{r1})}{t_{wb1} + t_{wb2}} \]

Additionally, statistical analysis tools were used in order to identify significant effects. An independent factor has significant effect on a given response when it has a p-value smaller than 0.05, considering a 95% confidence level. The results obtained for each run are listed in Table 3.

It was found that the temperature of the column oven (X_1), and the increase of acetonitrile percentage in mobile phase (X_2) directly influence the taxifolin response since they have p-values smaller than 0.05, and also all the two-factor interactions with the exception of the interaction X_1 × X_2. The increase of the mobile phase flow rate in the first 20 min did not affect taxifolin response (p-value higher than 0.05), but it allowed the best analysis time obtained in this study (50 min). The conditions of the central point (experiments 13, 14 and 15) showed the highest response for taxifolin (Table 3) and very stable baseline. Therefore, these conditions were selected as optimal and they were used in the validation study.

### Analytical validation

Using the optimized conditions, validation studies were performed. The specificity results are shown in Figure 3. As one can observe, the matrix did not interfere with the analysis of the compound of interest. Thus, it is possible to quantify taxifolin without interference, which was confirmed by UV spectra obtained for the taxifolin peak between 200 and 400 nm (Figure 4).

For linearity, calibration plots (x = μg mL⁻¹, y = mV) of taxifolin measured by the proposed method were constructed (n = 3 for each concentration). The least squares method obtained a determination coefficient greater than 0.99 (R² = 0.9965), which indicates the existence of a significant linear relationship between two variables. The regression model diagnosis was satisfactory with no lack-of-fit because the value of F_{calculated} is lower than F_{critical} in the 95% confidence interval (F_{calculated} = 0.65 < F_{critical(0.05)} = 3.71). Moreover, the regression significance was higher, since the F_{calculated} found was 3666.92 at the 95% confidence interval.
interval. The normality in the residues (Shapiro-Wilk’s test) and homoscedasticity (Cochran’s test) were applied in order to verify if this model fits. Indeed, Shapiro-Wilk’s W value ($W_{\text{calculated}} = 0.91$) was higher than the critical value ($W_{\text{critical}(0.05)} = 0.88$) and Cochran’s C value ($C_{\text{calculated}} = 0.58$) was lower than the critical value ($C_{\text{critical}(0.05)} = 0.68$), which indicates no violation of the assumptions, thus the ANOVA is considered valid. Therefore, the method is indeed linear, over its respective range.

The values obtained for LOD and LOQ were 6.52 and 21.70 $\mu$g mL$^{-1}$, respectively. These values showed adequate sensitivity for the analytical assay.

For precision, the RSD found were 3.46, 3.62 and 3.48% from intra-day, first day ($n = 12$); intra-day, second day ($n = 12$); and inter-days ($n = 24$), respectively. All RSD were within specification limits recommended ($\leq 5\%$). For accuracy, the average recovery percentage of the standard added was calculated, and the value found was 98.23% (RSD = 1.36%), which indicates that this method has an acceptable accuracy and suitability to quantify taxifolin without interferences.

Table 3. $3^1$ Box-Behnken design results for the method optimization

<table>
<thead>
<tr>
<th>Experiment</th>
<th>Taxifolin response (Y)</th>
<th>Factor</th>
<th>Coefficient</th>
<th>Error</th>
<th>p-Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.00</td>
<td>mean</td>
<td>2.24</td>
<td>0.07</td>
<td>0.01$^b$</td>
</tr>
<tr>
<td>2</td>
<td>2.10</td>
<td>$X_1$</td>
<td>0.32</td>
<td>0.04</td>
<td>0.02$^a$</td>
</tr>
<tr>
<td>3</td>
<td>0.82</td>
<td>$X_2$</td>
<td>0.17</td>
<td>0.04</td>
<td>0.06</td>
</tr>
<tr>
<td>4</td>
<td>1.71</td>
<td>$X_3$</td>
<td>-0.31</td>
<td>0.04</td>
<td>0.02$^a$</td>
</tr>
<tr>
<td>5</td>
<td>1.79</td>
<td>$X_1 \times X_1$</td>
<td>-0.79</td>
<td>0.07</td>
<td>0.01$^a$</td>
</tr>
<tr>
<td>6</td>
<td>0.00</td>
<td>$X_2 \times X_2$</td>
<td>-0.29</td>
<td>0.07</td>
<td>0.05</td>
</tr>
<tr>
<td>7</td>
<td>0.00</td>
<td>$X_3 \times X_3$</td>
<td>-0.68</td>
<td>0.07</td>
<td>0.01$^b$</td>
</tr>
<tr>
<td>8</td>
<td>1.32</td>
<td>$X_1 \times X_2$</td>
<td>-0.30</td>
<td>0.06</td>
<td>0.04$^b$</td>
</tr>
<tr>
<td>9</td>
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<td>0.78</td>
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<td>10</td>
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<tr>
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</tbody>
</table>

$^a$The best results for taxifolin response; $^b$statistically significant effects ($p < 0.05$).
For robustness analysis, a $2^3$ complete experimental design with triplicate in the central point was realized. The calculated values for main effects and interactions in a 95% level of confidence are shown in Table 4. From the analysis of the results, it is possible to see that small variations in these chromatographic parameters did not show significant changes in the overall assay values. Therefore, it is possible to conclude that the method is robust for the parameters analyzed within the intervals investigated.
Conclusions

Under the conditions described, the method for quantitative determination of taxifolin in the bark extract of *P. pinaster* is in accordance with the validation parameters required by the ICH and INMETRO guidelines. The method of HPLC-UV is available in quality control laboratories, and it is adequate for taxifolin quantification without matrix interferences. Therefore, the quantification of taxifolin may be an additional quality parameter for this extract in association with the spectrophotometric determination of the total procyanidins content. However, for this purpose, future investigations are necessary in order to establish limits for taxifolin content.

Acknowledgments

This work was supported by CAPES, CNPq, FAPEMIG and PROPESQ/UFJF. The authors would also like to thank Sharlene Loures for her help throughout the work.

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Submitted: September 30, 2015
Published online: February 11, 2016