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A new natural detector for irradiations with blue LED light source in photodynamic therapy measurements via UV–Vis spectroscopy

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Abstract

Photodynamic therapy has been recently studied, bringing innovations regarding the reduction of exposure time to light by the patient. This work aimed to investigate the feasibility of using *Coutarea hexandra (Jacq.) K. Schum* (CHS) as a detector in photodynamic therapy measurements. For this, an irradiator containing a blue LED bulb lamp was utilized. The CHS samples were irradiated with ten doses from 0.60 up to 6.0 kJ/cm², and six concentrations were prepared (1, 2, 3, 4, 5, and 6 mg/ml) for the CHS detector samples. After irradiation, the detector samples were evaluated using UV–Vis spectrophotometry. The results showed the behavior of the CHS detector with doses and concentrations, its sensitivity, and its linearity was also evaluated both by Wavelength Method (WM) and the Kernel Principal Component Regression (KPCR) Statistical Method. The values obtained indicate that this method can be applied to the CHS sample detector. In conclusion, the CHS is a promising detector in the field of photodynamic therapy.

Keywords New natural detector \cdot UV–Vis technique \cdot Blue LED source light

1 Introduction

Photodynamic therapy (PDT) has been investigated recently [1–4] in studies about new drugs [5], using nanoparticles [6], control microbial biofilms [7], and applications in advanced nanomaterials [8]. PDT consists of the annihilation of several cancers, through an incident visible light (VL), which sensitizes a drug called photosensitizer (PS), causing damage to living tissue (cancer cells), occurring the production of oxygen in this process [9–11]. For more details on the molecular basis, photosensitizers in medicine, the reaction mechanism, and the various modifications in PDT can be found in the literature [12–18]. The photosensitizers most used in PDT are the drug families: Porphyrin, Chlorin and Dyes [19]. Photosensitizers in recent decades have undergone modifications [20] with the main objective of reducing

concentrations and accumulation in the target areas of PS, not neglecting the effects on cancerous destruction. Some PS from plants have also been used in PDT with favorable results, such as *Rubiadin* and *Soranjidiol* [21], and *Quinones* [22].

In this context, this work focuses on the feasibility of a new photosensitizer to be used in PDT: the plant tested for its feasibility is *Coutarea hexandra (Jacq.) K. Schum* (CHS), that comes mainly from Brazil and appears in the states such as Amazonas (AM), Minas Gerais (MG), Pará (PA), extending to the state of São Paulo (SP). In each region, the *quina* popular name is modified and its applications according to the local population, being its use more common in antimalarial [23], antidiabetic, anti-contraceptive, and anti-inflammatory processes [24].

The PS feature can be performed using non-invasive and destructive techniques such as Ultraviolet–Visible spectrometry (UV–Vis) [25], which provides information on the absorbance as a function of the wavelength at which incident visible light sensitizes the PS.

Spectra from FTIR, and UV–Vis measurements, among others, can undergo pre-processing of their responses to be used in multivariate analysis [26–30]. This kind of analysis has been applied in several areas of knowledge to provide information, model, and to compare actual

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measurements with their prediction methods [31, 32]. A pre-processing method used is the Principal Component Analysis (PCA) method [33–36], which aims to resize the input data to a matrix containing the same information, but with a smaller size. This new information is associated with the number of components from the PCA called eigenvectors; usually, only two components are used for all the information described in the data system. After the PCA method, the other robust statistical methods can be applied [37, 38]. For the clustering analysis, it is necessary to apply k-means algorithm methods, consisting of partitioning the *n* observations among *k* groups. Each group is separated according to the distance of the values closest to their means (centroids). These separate groups are called clusters [39, 40].

In a physicochemical system under investigation that contains multiple degrees of freedom, e.g., several concentrations range and absorbed doses for a given sample, it is possible to apply the Kernel Principal Component Regression (KPCR) method to determine the possible linearity behavior of the system. KPCR is used to handle the multicollinearity effect among the independent variables from the regression data; this method has been used successfully for this objective [41–44].

This work aims to analyze, for the first time, the viability of CHS detector in PDT via UV–Vis measurements using multivariate statistical techniques such as PCA, k-means, and KPCR.

2 Materials and methods

2.1 Equipment and preparation of the CHS detector

The preparation of the CHS solution can be described in the schematic process as in phases a to f. In phase a, the CHS powder is measured on a semi-analytical balance; in phase b, there is the beaker containing 71.6 ml of 70% ethyl alcohol, also there is a small handle containing 126.4 ml of milli-Q water, and a vessel with 500 ml of volume, containing the CHS powder; in phase c, all reagents are added slowly to the vessel and mixed; in phase d, the solution is added to a funnel, 10 cm in diameter which contains a 9 cm diameter filter, 44 µm pore, 250 g weight, thickness of 0.5 mm connected to a 250 ml beaker which houses the filtered solution; in phase e, the solution is transferred to a 500 ml glass container, covered with aluminum foil, and kept in a refrigerator (15 °C) for 72 h; in phase f, the ready solution is inserted into standard cuvettes with dimensions of $1.0 \times 1.0 \times 4.5$ cm³ through a micropipette, thus making the CHS detector samples. Altogether there were six concentrations which are: 1, 2, 3, 4, 5, and 6 mg/ml for the CHS detector samples.

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2.2 Irradiations and evaluation of CHS detector samples

The radiation source was a blue bulb with a wavelength around 440–485 nm. This lamp was placed in front of the CHS samples (0.5 cm surface-source distance) and controlled by a circuit coupled to a microcontroller (Raspberry Computer Model B Pi 4 with 4 GB of RAM).

A program was developed in Matlab[®] 2020a, to turn the lamp on and off; more details of the irradiation system can be seen in the literature [4]. The irradiations were carried out in the laboratory in darkness with the only window closed and with a curtain, to prevent the entry of light from outside. In this environment, the temperature was controlled through an air conditioner turned on at 20 °C throughout the experiment.

The irradiation times were: 5, 10, 15, 20, 25, 30, 35, 40, 45, and 50 min, which correspond to the doses: 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/ cm², respectively.

After irradiation, CHS samples were evaluated using the Genesys 10S/Thermo Scientific spectrophotometer, coupled to a UV–Vis Vision Life data system software.

2.3 Parameters for CHS detector samples

The absorbance of the CHS detector samples was determined by the Law of Lambert–Beer, which considers the logarithmic ratio between the non-irradiated samples and irradiated samples from the new detector. All the measurements were made in triplicate, and the average values are reported. Therefore, some characterization tests of the CHS detector samples can be performed regarding the determination of the CHS spectra for six concentrations and ten doses; calibration curves from six concentrations of CHS samples; the evaluation of the linearity considering the squared Pearson correlation coefficient (R^2); the sensitivity; and results for statistical methods. The definitions of linearity, sensitivity and Wavelength Method (WM) for spectra obtained with the UV–Vis technique can be found in the literature [4].

2.4 Multivariate dose-response analysis

Multivariate analysis is a field of mathematics that deals with multidimensional matrices. As a result, linear algebra is coupled with statistics to produce tools that find regression relationships between multiple responses and their predictor variables, regardless of the data dimensions. Specifically, in radiation physics, multivariate analysis is used as a processing technique sinking to determine a parsimonious model that fits the data.

This work used a combination of analytical methods to ascertain how the absorbance readings vary as a function of the coupled effects of the CHS powder concentration and the absorbed dose. The characterization of the absorbance distribution over the effects of concentrations and absorbed dose is essential because it is possible to assess linear relationships among the absorbance reading and the concentration factor.

The applied method to the multivariate analysis was twofold: first to find similar patterns within the absorbance distribution according to their wavelength index, while pondering the effects of the concentration and simultaneously the absorbed dose through the k-means coupled with Principal Component Analysis (PCA); second, with the Kernel Principal Component Regression (KPCR) to fit a model and then characterize the possible linear behavior between the absorbance reading as a function of the absorbed dose and the concentration.

2.4.1 *k-means algorithm coupled with* Principal Component Analysis (*PCA*)

To find similar patterns within the absorbance distribution while pondering the effects of (K=6) concentration varying from 1, 2, 3, 4, 5 to 6 mg/ml, and absorbed dose (M=10) number of total doses, 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/cm²; taken in the discrete values (N=150), the values of absorbance comprised between the region of wavelength from 400 to 550 nm. One group corresponded to 150 samples of R^2 ; thus, the sample size was 150. In total, six groups from the effect of concentration ranging from 1 to 6 ml/mg were evaluated.

The k-means algorithm clustered the absorbance data, but since the readings' dimensions were high, the k-means was coupled with the PCA. Consequently, the spectral absorbance reading for all concentrations was projected into a lower dimension space using the PCA after clustered in a two-dimensional space.

Therefore, in the PCA [45–47] method, the data is first written in a condensed matrix form as $\{x_i\}_{i=1}^N \in \mathbb{R}^N$, which is equivalent to $x_i = (x_{i1}x_{i2}\cdots x_{ip})^T$, where x_i is the regression variable, and x_{ij} corresponds to *i* th observation (*i* = 1, 2, ..., *N*), to the *j*th (*j* = 1, 2, ..., *M*) regression variables, defining the matrices $\tilde{X} = (x_1x_2\cdots x_p)^T$ and $X_c = \tilde{X}(I_N - N^{-1}1_N1'_N)$. The zero mean-centered covariance matrix is written as:

$$C = \frac{1}{M-1} X'_c X_c,$$
 (1)

where the term $(\cdot)^T$ denotes the transpose of the matrix; $(\cdot)^{-1}$ is the inverse of the matrix; \tilde{X} is a global matrix which

contents all elements; X_c is the centered matrix of \tilde{X} ; I_N is the identity matrix; 1_N is a matrix with all entries equal to 1. The covariance matrix *C* is related to the \tilde{X} matrix through the eigenspaces as a canonical basis, with the columns being the eigenvector of the observations, i.e., the principal components. Applying the eigendecomposition in Eq. 1, the results are:

$$C = V\Lambda V' = \sum_{k=1}^{p} \lambda_k v_k v'_k, \qquad (2)$$

where *C* covariance matrix *C* is related to the \tilde{X} . Such relation is made through the eigenspaces as the canonical basis; $V_{N\times N}$ is the matrix eigenvectors (orthogonal); v_k are the eigenvalues from columns of the $V_{N\times N}$; Λ is spectral matrix; diag(Λ) are composed of the eigenvalues ($\lambda_k \cdots \lambda_p$); s_{kk} is the diagonal of *S*, where $\lambda_k = s_{kk}^2$.

The first two components of *C* correspond to the most significant variances in \tilde{X} , i.e., as a corollary, with the deflation of the number of principal components represented by the eigenvectors, the data variance is maintained at high levels according to the final number of eigenvectors and its magnitudes. Thus, with the two most significant eigenvalues and eigenvectors, the \tilde{X} is projected into bi-dimensional space, which allowed the k-means to cluster the data into regions.

The cluster was set using as a decision rule the smallest Euclidean distance between the established regions μ_j centroids (*C*) and the \tilde{X} matrix data now standardized by z-score of the *C* matrix, all of which come from the PCA, according to

$$\sum_{i=0}^{N} \min_{\mu_j \in C} (\left\| C_i - \mu_j \right\|^2)$$
(3)

New centroids were standardized by mean values, standard deviations, and *S* matrix, all of which come from the PCA.

2.4.2 Principal Kernel Component Regression—KPCR

The first step in implementing the KPCR [44, 48] was to transform the \tilde{X} matrix through a function $\psi : \mathbb{R}^g \to F$, where *F* is the feature space is a Euclidean space of higher dimension than *g*, called g_F . The ψ function is not explicitly defined but is obtained through $\kappa : \mathbb{R}^g \times \mathbb{R}^g \to \mathbb{R}$, where κ is the Kernel function. The Kernel is defined as $K = \Psi \Psi^T$ and $\Psi = (\psi(\tilde{x}_1)\psi(\tilde{x}_2)\cdots\psi(\tilde{x}_N))^T$, the dimensions of Ψ and K are $N \times g^F$ and $N \times N$, respectively.

The KPCR is fundamentally a Multiple Linear Regression (MLR) applied in the feature space F. The relationship between the feature space in the higher dimension and the Euclidean space is made through the Kernel trick.

First, it was set the Kernel Gaussian as the mapping to the feature space, given as:

$$k(x, y) = e^{-\left(\frac{\|x-y\|}{\hbar}\right)}.$$
(4)

Then, the MLR in the feature space is given as $Y = \Psi \eta + \epsilon'$, and Ψ the normalized eigenvectors v_i of $\Psi^T \Psi$ is the correspondent of $V = (v_1 v_2 \cdots v_{g^F})$ matrix, and $VV^T = I_{g^F}$, then the model above can be rewritten using the Kernel trick in the Reproducing Kernel Hilbert Space (RKHS):

$$Y_{\circ} = B\omega + \varepsilon'. \tag{5}$$

In Eq. 5, the independent variable *Y* originally in the Euclidean space should be centered by the mean values on the feature space. This process is made through Eq. 6:

$$Y_{\circ} = \left(I_N - \frac{1}{N} \mathbf{1}_N \mathbf{1}_N^T\right) Y,\tag{6}$$

where $B = \Psi V$ and $\omega = (\omega_1 \omega_2 \cdots \omega_{g^F})^T = V^T \eta$, which is equivalent to the eigendecomposition problem, where $\hat{\omega} = (\hat{\omega}_1 \hat{\omega}_2 \cdots \hat{\omega}_{g^F})^T$ is the estimator of ω , which can be written as $\hat{\omega} = (B^T B)^{-1} B^T Y$.

$$\eta = V\hat{\omega} = \sum_{i=1}^{p_f} \mu_i^{-1} v_i v_i^T \Psi^T Y$$
(7)

It is noted that the dimension of $\hat{\omega}$ depends on the number of principal components (k). Finally, the estimated model Y can be compared with the real values of doses. So basically, Eq. 7 is a projection of the mapped matrix's eigenvectors on the feature space. As a result, one may deflate the matrix Ψ , which means excluding some eigenvectors. Consequently, the deflating process handles the harmful effects associated with multicollinearity presented in the original Euclidean space. $\|\cdot\|$ is the norm of vector; Y is the predicted value from the input M; η is a vector of regression coefficients and ϵ' is a vector of random errors in the feature space.

2.5 Validation of KPCR method for dose-response

The KPCR method provides results for dose–response in association with the number of principal components (k), so it is possible to validate this method by comparing it with measurements taken with the UV–Vis spectrophotometry equipment for all doses and concentrations of this work. These validation and statistical error methods will be used with criteria established to define the method accuracy; they are Root Mean Square Percentage Error (RMSPE) defined by Eq. 8, and Mean Absolute Percentage Error (MAPE) defined by Eq. 9, as follows:

RMSPE =
$$\sqrt{\frac{1}{n} \sum_{i=1}^{n} \left(\frac{y_i - \hat{y}_i}{y_i}\right)^2} \times 100\%$$
 (8)

MAPE =
$$\sqrt{\frac{1}{n} \sum_{i=1}^{n} \left| \frac{y_i - \hat{y}_i}{y_i} \right|} \times 100\%,$$
 (9)

where y_i are the real values, \hat{y}_i are the predicted values, and n is the amount of data used in the prediction. Thus, RMSPE and MAPE methods have critical criteria characteristics of predicting values [49].

3 Results and discussion

The illuminance measurements versus exposure time, for the blue LED light source are shown in Fig. 1a. The illuminance values increase over time, until their values stabilize: for this curve, a 3rd-degree polynomial fit ($R^2 = 0.9611$) was obtained. Consequently, the CHS detector samples were irradiated after the stabilization period of the radiation source; according to the curve, this time will be obtained in > 19.7 min (in this work an adequate time for initial measurements is around 60 min).

The spectra of the CHS detector samples are shown in Fig. 1b for concentrations of 1, 2, 3, 4, 5, and 6 mg/ml, in a wavelength range from 400 to 700 nm. There is a region with higher absorbance values from 400 to 450 nm and decay of these values up to 700 nm. The absorbance values also vary with the associated concentrations, in ascending order. Thus, the UV–Vis spectroscopy technique made it possible to obtain the spectra of the CHS detector samples.

In Fig. 2a, the calibration curve shows the squared Pearson correlation coefficient (R^2) of 0.9895; the absorbances obtained adequately responded to the various concentrations. In Fig. 2b, linearity was plotted versus wavelength, to obtain the wavelength for the calibration curve, providing the largest R^2 ; this value was 400 nm, using all concentrations. Sensitivity versus wavelength behavior for the CHS detector samples for all concentrations are shown in Fig. 2c. The results indicate the wavelength with the highest sensitivity, which is 400 nm; afterwards, the sensitivity decreases with the increase in wavelength, and it shows that at 400 nm the maximum values for both linearity and sensitivity occur. For concentrations greater than 6 mg/ml, significant signal-tonoise ratios (SNRs) appear in the spectrum, discarded in this work. It is worth mentioning that the results of Fig. 2a-d are from non-irradiated CHS detector samples.

The absorbance versus wavelength for the CHS detector samples, UV–Vis spectra, for doses of 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/cm^2 , and for all concentrations used in this work are shown in Fig. 3.



Fig. 1 a Illuminance versus exposure time for blue LED source irradiation, in the time 0 up to 1180 s. Error bars are the standard deviations for triplicate measurements, **b** absorbance versus wavelength, UV–Vis spectra, for CHS detector samples with 1, 2, 3, 4, 5, and 6 mg/ml concentration, respectively. In this case, the non-concentration cuvette contained air. The uncertainty of the measurements obtained was lower than 1%. The uncertainty was of type C, which considers the uncertainties A (from measurements) and the uncertainties B (from equipment)

First, for all concentrations, it is possible to observe variations in their absorbance, thus proving the interaction of radiation from the blue LED source and the detector. The maximum absorbances (peaks) for the concentrations of 1, 2, 3, 4, 5, and 6 mg/ml are in the wavelength regions of $(478.0 \pm 1.2 \text{ nm})$, $(477.0 \pm 1.7 \text{ nm})$, $(508.0 \pm 1.5 \text{ nm})$, $(478.0 \pm 1.0 \text{ nm})$, $(476.0 \pm 0.7 \text{ nm})$, and $(477.0 \pm 0.3 \text{ nm})$, respectively.

These results demonstrate that most of the longest wavelengths are in the blue radiation source. These results follow the literature where the radiation wavelength is considered close to the detector wavelength [50, 51]. During the irradiations, it was also possible to observe that the CHS detector samples did not change color significantly,

so they cannot be classified as Yes/No irradiated detectors for the dosing interval analyzed in this work.

For some linearity values, with their corresponding wavelengths, they are shown in Table 1. These values are also associated with the concentrations of the CHS detector samples; this result shows that there are more regions in which the detector can be applied with adequate linearity results. It is possible to observe that for each concentration there is a value for linearity; each linear fit was carried out at the wavelength that provides the highest value of linearity (wavelength method). The maximum R^2 values were obtained, for 541, 530, 423, 443, 533, and 422 nm, and 1, 2, 3, 4, 5, and 6 mg/ml, respectively.

In Table 2 are shown the polynomial fit and linear fit for sensitivity and linearity; these results give adequate approximation mathematics for the curves from the measurements. The values for R^2 are ≥ 0.9507 and ≥ 0.9911 , for the polynomial fit and linear fit, respectively. Figure 3 shows the maximum values of absorbance and Table 1 the maximum values of R^2 . A region with higher absorbance values will not necessarily present the maximum R^2 values.

In Table 3, standard deviations are shown for all concentrations and associated with a region of 400 up to 420 nm, with irradiated and non-irradiated CHS detector samples. These results are important for the experimental response reproducibility of the detector; they can also infer that the standard deviation value decreases with the increase of the wavelength.

The results on sensitivity versus wavelength, for the CHS detector samples, are shown in Fig. 4. The results for sensitivities indicate regions and peaks where the CHS detector sample is more sensitive to non-ionizing radiation. For the concentrations of 1, 2, and 4 mg/ml the peak is at 484 nm, for 3 mg/ml the peak is at 495 nm and for concentrations 5 and 6 mg/ml the peak is at 478 nm. In Table 2 on the left, squared Pearson correlation coefficients (R^2) are shown for each concentration, which is greater than 0.9507, and these results infer that the sensitivity for the CHS detector samples can be determined through mathematical equations for the sensitivity curves.

Figure 5 presents the correlation matrix from CHS detector samples irradiated with ten doses and six concentrations. The results indicated that this matrix had highlighted a significant multicollinearity degree among the data (p < 0.01). Therefore, a multidimensional analysis was performed through the k-means algorithm seeking to evaluate the concentration effects over absorbance readings for the absorbed doses ranging from 0.60 to 6 kJ/cm². Therefore, as stated in Sect. 2.4, the whole dataset underwent the PCA transformation to assess the absorbed dose influence. This process was performed for the absorbance of the samples from to 700 nm, for each concentration ranging from 1 to 6 mg/ml.



Fig. 2 a Calibration curves of six concentrations of CHS detector samples: 1, 2, 3, 4, 5, and 6 mg/ml, with R^2 =0.9895, the standard deviations were <1% for the triplicate measurements, **b** linearity versus wavelength for all concentrations used in this work, and **c** Sen-

sitivity versus wavelength for concentrations from 1 up to 6 mg/ml. The CHS detector samples were not irradiated in these results and have a cuvette containing air as standard sample

The results on Principal Component Analysis (PCA) and k-means for all concentrations, for CHS detector samples, are shown in Fig. 6. The k-means identified cluster regions that were projected over the two-dimensional PCA component space for visualization purposes. Thus, two cluster regions were found; as a result, more than 99.97% of the variance was explained using the first principal component, which agrees with the overall squared Pearson correlation coefficient (R^2) that was above 0.9970 for the doses. These results also show the predominance of the first principal component concerning the second one while explaining the variance regarding the absorbance readings as a concentration function. Thus, within the two identified clusters, the first contained 85.50% of samples, while the second with

the remaining. Although, when weighting the dimensional effect, the cumulative variance accounted 1:45 fold ratio between the first and second eigenvalues. Thus, the second principal dimensional described only 2.22% of the first principal component, undermining the decrease in linearity as previously observed. As a result, it can be generalized that the linearity variation was inexpressive concerning the concentration variation, i.e., the overall linearity is independent of the concentration. Hence, in evaluating the concentration-effect over the linearity, it was possible to ascertain, through the cluster tail, that the second cluster manifested a decrease in the linearity as the wavenumber increased for the concentrations of 3 and 6 mg/ml.

Fig. 3 Absorbance versus wavelength, for CHS detector samples for ten doses: 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/ cm² (blue LED source). All measurements were taken with 1, 2, 3, 4, 5, and 6 mg/ml concentrations. In this case, the non-irradiated cuvette contained CHS detector solution of 1. 2. 3, 4, 5, and 6 mg/ml concentrations. The uncertainty of the measurements obtained was lower than 1%. The uncertainty was of type C, which considers the uncertainties A (from measurements) and the uncertainties

B (from equipment)



Table 1 The maximum values for squared Pearson correlation coefficient (R^2) and their appropriate locations in the spectrum (wavelength), for CHS detector sample concentrations of 1, 2, 3, 4, 5, and

6 mg/ml, and for doses: 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/cm² (blue LED source)

| 1 mg/ml | | 2 mg/ml | | 3 mg/ml | | 4 mg/ml | | 5 mg/ml | | 6 mg/ml | |
|------------------|-------------------------|------------------|-------------------------|------------------|-------------------------|------------------|-------------------------|------------------|-------------------------|------------------|-------------------------|
| $\overline{R^2}$ | Wave- length (nm) |
| 0.9944 | 541 | 0.9962 | 530 | 0.9916 | 423 | 0.9920 | 443 | 0.9968 | 533 | 0.9891 | 422 |
| 0.9941 | 545 | 0.9960 | 447 | 0.9915 | 421 | 0.9918 | 444 | 0.9968 | 535 | 0.9886 | 549 |
| 0.9939 | 542 | 0.9950 | 531 | 0.9913 | 426 | 0.9917 | 447 | 0.9968 | 541 | 0.9856 | 551 |
| 0.9937 | 446 | 0.9935 | 461 | 0.9909 | 427 | 0.9916 | 446 | 0.9967 | 537 | 0.9851 | 420 |
| 0.9936 | 445 | 0.9930 | 460 | 0.9909 | 425 | 0.9916 | 442 | 0.9967 | 540 | 0.9843 | 424 |
| 0.9934 | 443 | 0.9929 | 463 | 0.9909 | 422 | 0.9915 | 448 | 0.9965 | 532 | 0.9830 | 419 |
| 0.9931 | 436 | 0.9927 | 462 | 0.9901 | 420 | 0.9914 | 441 | 0.9964 | 441 | 0.9827 | 423 |
| 0.9930 | 444 | 0.9898 | 493 | 0.9844 | 428 | 0.9911 | 449 | 0.9963 | 442 | 0.9818 | 421 |
| 0.9927 | 440 | 0.9896 | 490 | 0.9830 | 419 | 0.9906 | 440 | 0.9963 | 542 | 0.9808 | 418 |
| 0.9927 | 439 | 0.9891 | 492 | 0.9829 | 429 | 0.9906 | 439 | 0.9962 | 536 | 0.9795 | 548 |
| 0.9925 | 543 | 0.9886 | 543 | 0.9819 | 430 | 0.9903 | 438 | 0.9961 | 547 | 0.9782 | 417 |
| 0.9924 | 547 | 0.9879 | 494 | 0.9809 | 418 | 0.9900 | 450 | 0.9961 | 544 | 0.9762 | 416 |
| 0.9924 | 536 | 0.9872 | 529 | 0.9795 | 431 | 0.9893 | 451 | 0.9960 | 440 | 0.9751 | 425 |
| 0.9922 | 550 | 0.9871 | 449 | 0.9784 | 417 | 0.9892 | 437 | 0.9960 | 539 | 0.9744 | 547 |
| 0.9921 | 441 | 0.9870 | 491 | 0.9775 | 432 | 0.9885 | 436 | 0.9959 | 531 | 0.9695 | 412 |
| 0.9920 | 449 | 0.9866 | 443 | 0.9755 | 416 | 0.9881 | 452 | 0.9959 | 534 | 0.9694 | 426 |
| 0.9919 | 448 | 0.9865 | 442 | 0.9750 | 433 | 0.9874 | 435 | 0.9959 | 530 | 0.9690 | 415 |
| 0.9920 | 549 | 0.9865 | 544 | 0.9737 | 415 | 0.9873 | 453 | 0.9958 | 550 | 0.9685 | 408 |
| 0.9916 | 442 | 0.9862 | 446 | 0.9720 | 434 | 0.9871 | 538 | 0.9958 | 543 | 0.9659 | 546 |

Figure 7 presents the Cumulative Distribution Function (CDF) versus squared Pearson correlation coefficient (R^2) for all concentrations. The order in the legend is ascending,

but the results do not follow the order shown in the legend. These results show that the organization of the different spectra from the concentrations provide distinct and

| Concentra- tion (mg/ml) | Equation (Fig. 6) | Fit | R^2 | Equation (Table 1) | Fit | R^2 |
|----------------------------|--|------------|--------|-------------------------------------|--------|--------|
| 1 | $y = 4.5 \times 10^{-9} x^3 - 9.4 \times 10^{-6} x^2 + 0.006x - 1.218$ | Polynomial | 0.9860 | $y = -1.1 \times 10^{-2} x - 0.001$ | Linear | 0.9945 |
| 2 | $y = 6.0 \times 10^{-9} x^3 - 1.2 \times 10^{-5} x^2 + 0.008 x - 1.516$ | Polynomial | 0.9793 | $y = -1.4 \times 10^{-2} x + 0.013$ | Linear | 0.9967 |
| 3 | $y = -2.0 \times 10^{-8} x^3 + 2.5 \times 10^{-5} x^2 - 0.010x - 1.384$ | Polynomial | 0.9770 | $y = -1.5 \times 10^{-2} x - 0.001$ | Linear | 0.9918 |
| 4 | $y = -2.5 \times 10^{-8} x^3 + 3.4 \times 10^{-5} x^2 - 0.014x + 2.064$ | Polynomial | 0.9627 | $y = -1.2 \times 10^{-2} x + 0.013$ | Linear | 0.9923 |
| 5 | $y = 6.9 \times 10^{-9} x^3 - 1.3 \times 10^{-6} x^2 + 0.008x - 1.522$ | Polynomial | 0.9678 | $y = -1.1 \times 10^{-2} x + 0.003$ | Linear | 0.9969 |
| 6 | $y = -1.6 \times 10^{-8} x^3 + 2.1 \times 10^{-5} x^2 - 0.009 x + 1.193$ | Polynomial | 0.9507 | $y = -4.1 \times 10^{-3} x - 0.030$ | Linear | 0.9911 |

Table 2 Linear fit equations with six concentrations and ten doses. The respective squared Pearson correlation coefficients (R^2) correspond to the absorbance values with wavelengths at 541, 530, 423, 443, 533, and 422 nm for 1, 2, 3, 4, 5, and 6 mg/ml, respectively.

A polynomial fit was obtained for the CHS detector samples to all concentrations, spectra, and doses. The values of y and x are the absorbance and dose, respectively

independent R^2 and CDF values, as can be seen in Fig. 7. These results corroborate those obtained by the previous analysis for the PCA and k-means methods. This means that most of the data from the CHS detector samples are in the region of strong linearity, specifically with values of 80% for the CDF provides $R^2 \ge 0.9706$, for all concentrations and dose data of the CHS detector samples.

Figure 8 shows the values predicted from the KPCR method (k=2) in comparison with the measured values of the CHS detector samples with all concentrations. As a corollary of the above observation, the results show KPCR regression lines employed to quantify the statistical significance of the linearity variation to the concentration effects. The hypothesis test through the ANOVA, in terms of the difference of mean values, showed that the R^2 for all concentrations is not different (p < 0.01). The result for ANOVA indicates that the null hypothesis is rejected, thus the mean values for all six groups are different and statistically significant; therefore, the ANOVA method may be applied here. Thus, the overall linearity of the adjusted multivariate KPCR model indicates invariance of linearity taking simultaneously the effects of the concentration and the spectral (wavenumbers) variations. Table 4 shows the values for the squared Pearson correlation coefficient (R^2) associated with the concentrations and the number of principal components (k) from the KPCR method. These results indicate the evolution of the number of principal components from k=1up to k=9. With the increase in these values, R^2 will also increase until their maximum value is obtained, for example, for the concentration of 6 mg/ml, the maximum values were already reached in k=7. This fact demonstrates that the KPCR method fitted the entire concentrations made in the measurements and not just a concentration in its analysis, that is, for global analysis of the system an appropriate method is the KPCR, which provided adequate results for comparisons with the measurements of the CHS detector samples.

Statistical error methods as the Root Mean Square Percentage Error (RMSPE) and the Mean Absolute Percentage Error (MAPE) versus number of principal component (*k*) for 1, 2, 3, 4, 5, and 6 mg/ml concentrations are presented in Fig. 9a–f, respectively. For the initial analysis, the best criterion for comparisons between the RMSPE and MAPE methods was chosen in which the error < 10% is considered highly accurate. In this criterion, the concentration of 1 mg/ ml requires k=1 for both MAPE and RMSPE methods, the concentration of 2 mg/ml requires k=1 (MAPE) and k=2(RMSPE), the other concentrations of 3, 4, 5, and 6 mg/ ml, for MAPE and RMSPE require k=2. This work will assign a criterion with more accuracy than observed in the literature, that is, an analysis for an error < 5% was considered here as excellent accuracy. For this second analysis,

| tor samples | | | | I | | 1 | | | | | | |
|---------------|-------------------------|--------------------------|-------------------------|--------------------------|---------------------------|--------------------------|---------------------------|--------------------------|-------------------------|----------------------------|-------------------------|--------------------------|
| Wavelength nm | 1 mg/ml | | 2 mg/ml | | 3 mg/ml | | 4 mg/ml | | 5 mg/ml | | 6 mg/ml | |
| | Standard dev | viation | Standard dev | iation | Standard devi | ation | Standard devi | iation | Standard devi | iation | Standard devi | ation |
| | (0.0 kJ/cm^2) | (0.60 kJ/cm^2) | (0.0 kJ/cm^2) | (0.60 kJ/cm^2) | (0.0 kJ/cm ²) | (0.60 kJ/cm^2) | (0.0 kJ/cm ²) | (0.60 kJ/cm^2) | (0.0 kJ/cm^2) | (0.60 kJ/cm ²) | (0.0 kJ/cm^2) | (0.60 kJ/cm^2) |
| 400 | 0.0013 | 0.0034 | 0.0211 | 0.0056 | 0.0384 | 0.0083 | 0.0112 | 0.0042 | 0.0012 | 0.0095 | 0.0067 | 0.0050 |
| 401 | 0.0013 | 0.0033 | 0.0205 | 0.0052 | 0.0370 | 0.0072 | 0.0094 | 0.0037 | 0.0014 | 0.0116 | 0.0029 | 0.0132 |
| 402 | 0.0015 | 0.0031 | 0.0193 | 0.0052 | 0.0354 | 0.0068 | 0.0091 | 0.0029 | 0.0022 | 0.0087 | 0.0095 | 0.0028 |
| 403 | 0.0014 | 0.0032 | 0.0175 | 0.0048 | 0.0342 | 0.0067 | 0.0108 | 0.0029 | 0.0033 | 0.0051 | 0.0023 | 0.0050 |
| 404 | 0.0013 | 0.0030 | 0.0152 | 0.0045 | 0.0328 | 0.0061 | 0.0096 | 0.0041 | 0.0013 | 0.0083 | 0.0049 | 0.0060 |
| 405 | 0.0012 | 0.0027 | 0.0145 | 0.0039 | 0.0313 | 0.0059 | 0.0086 | 0.0028 | 0.0020 | 0.0071 | 0.0063 | 0.0075 |
| 406 | 0.0011 | 0.0027 | 0.0137 | 0.0040 | 0.0303 | 0.0055 | 0.0082 | 0.0034 | 0.0041 | 0.0083 | 0.0056 | 0.0033 |
| 407 | 0.0014 | 0.0024 | 0.0131 | 0.0040 | 0.0289 | 0.0052 | 0.0081 | 0.0030 | 0.0028 | 0.0057 | 0.0033 | 0.0038 |
| 408 | 0.0013 | 0.0024 | 0.0128 | 0.0042 | 0.0276 | 0.0051 | 0.0079 | 0.0025 | 0.0034 | 0.0057 | 0.0058 | 0.0077 |
| 409 | 0.0011 | 0.0021 | 0.0119 | 0.0039 | 0.0262 | 0.0047 | 0.0077 | 0.0024 | 0.0022 | 0.0062 | 0.0032 | 0.0054 |
| 410 | 0.0011 | 0.0023 | 0.0110 | 0.0037 | 0.0249 | 0.0048 | 0.0068 | 0.0018 | 0.0020 | 0.0053 | 0.0028 | 0.0026 |
| 411 | 0.0011 | 0.0020 | 0.0101 | 0.0033 | 0.0237 | 0.0043 | 0.0066 | 0.0019 | 0.0023 | 0.0056 | 0.0030 | 0.0034 |
| 412 | 0.0011 | 0.0021 | 0.0102 | 0.0035 | 0.0227 | 0.0043 | 0.0064 | 0.0017 | 0.0022 | 0.0050 | 0.0027 | 0.0050 |
| 413 | 0.0009 | 0.0020 | 0.0096 | 0.0033 | 0.0214 | 0.0043 | 0.0060 | 0.0017 | 0.0020 | 0.0046 | 0.0037 | 0.0027 |
| 414 | 0.0010 | 0.0019 | 0.0092 | 0.0033 | 0.0205 | 0.0038 | 0.0058 | 0.0018 | 0.0017 | 0.0041 | 0.0019 | 0.0042 |
| 415 | 0.0010 | 0.0018 | 0.0085 | 0.0031 | 0.0194 | 0.0035 | 0.0055 | 0.0018 | 0.0018 | 0.0042 | 0.0030 | 0.0045 |
| 416 | 0.0008 | 0.0015 | 0.0076 | 0.0028 | 0.0186 | 0.0032 | 0.0053 | 0.0015 | 0.0019 | 0.0040 | 0.0023 | 0.0042 |
| 417 | 0.0008 | 0.0014 | 0.0067 | 0.0026 | 0.0178 | 0.0032 | 0.0047 | 0.0014 | 0.0017 | 0.0039 | 0.0029 | 0.0040 |
| 418 | 0.0007 | 0.0013 | 0.0060 | 0.0025 | 0.0165 | 0.0029 | 0.0047 | 0.0015 | 0.0018 | 0.0035 | 0.0026 | 0.0033 |
| 419 | 0.0007 | 0.0012 | 0.0023 | 0.0023 | 0.0157 | 0.0028 | 0.0043 | 0.0013 | 0.0017 | 0.0036 | 0.0023 | 0.0037 |
| 420 | 0.0006 | 0.0011 | 0.0021 | 0.0020 | 0.0151 | 0.0024 | 0.0041 | 0.0013 | 0.0015 | 0.0034 | 0.0021 | 0.0033 |

Fig. 4 Sensitivity versus wavelength, for CHS detector samples and ten doses: 0.60, 1.20, 1.80, 2.40, 3.00, 3.60, 4.20, 4.80, 5.40, and 6.00 kJ/ cm^2 (blue LED source), and six concentrations: 1, 2, 3, 4, 5, and 6 mg/ml



Fig. 5 Correlation matrix for CHS detector samples with 0.60, 1.20, 1.80, 2.40, 3.00,3.60, 4.20, 4.80, 5.40, and 6.00 kJ/cm^2 doses, and 1, 2, 3, $4, 5, and 6 \text{ mg/ml concentra$ $tions}$

the concentration of 1 mg/ml requires k=2 for both MAPE and RMSPE methods, the concentration of 2 mg/ml requires k=3 for both MAPE and RMSPE methods, the concentration of 3 mg/ml requires k=2 (MAPE) and k=3 (RMSPE), the concentration of 4 mg/ml requires k=3 (MAPE) and k=4 (RMSPE), the concentrations of 5 and 6 mg/ml require

k=4 and k=5 for both, MAPE and RMSPE methods, respectively. For all analyzed concentrations: (i) the RMSPE values are greater than the MAPE values, (ii) the errors are greater when k=1, (iii) they converge, when the predictive data are equal to the real ones, for k=9.

Fig. 6 Principal Component Analysis (PCA) for 1, 2, 3, 4, 5, and 6 mg/ml concentrations. The first principal component is represented by the *x* axis, while the second main component is represented by the *y* axis. The k-means clusters can also be observed, on the left side the first cluster and its centroid, and to the right the second cluster and its centroid





4 Conclusions

The CHS detector presents preliminary results which can be useful in measurements for photodynamic therapy, such as: (i) the preparation of the CHS samples is simple, following the steps they can be easily reproduced; (ii) the application of the UV–Vis spectrophotometric technique to the CHS detector samples could be performed with low standard deviations, thus guaranteeing the reproducibility of the results obtained for both irradiated and non-irradiated samples; (iii) the irradiation system was effective in irradiating the samples and the blue LED light sensitized the photosensitizer in the region of the detector wavelength; (iv) the results for linearity through the best wavelength were adequate, thus providing a quick analysis of the detector at a specific wavelength and visually determined on the graph (peaks) of absorbance versus wavelength; (v) the correlation matrix, PCA together with k-means provided information regarding the clusters and the main components; these were determined and were accurate for the Fig. 8 Predicted dose versus measured dose, through the KPCR method, using k=2 as number principal of component, for CHS detector samples with 1, 2, 3, 4, 5, and 6 mg/ml concentrations



Table 4 Squared Pearson correlation coefficient (R^2) versus number of principal component (k) for 1, 2, 3, 4, 5, and 6 mg/ml concentrations for CHS detector samples irradiated with blue LED source

| Squared Pearson correlation coefficient (R^2) | | | | | | | | | |
|---|-------------|---------------|---------|---------|---------|---------|--|--|--|
| Number of principal | Concentrati | Concentration | | | | | | | |
| component (k) | 1 mg/ml | 2 mg/ml | 3 mg/ml | 4 mg/ml | 5 mg/ml | 6 mg/ml | | | |
| 1 | 0.9870 | 0.9764 | 0.9509 | 0.9822 | 0.9868 | 0.8523 | | | |
| 2 | 0.9873 | 0.9936 | 0.9625 | 0.9921 | 0.9948 | 0.9920 | | | |
| 3 | 0.9934 | 0.9986 | 0.9972 | 0.9924 | 0.9962 | 0.9920 | | | |
| 4 | 0.9944 | 0.9986 | 0.9989 | 0.9924 | 0.9986 | 0.9968 | | | |
| 5 | 0.9990 | 0.9986 | 0.9989 | 0.9989 | 0.9987 | 0.9968 | | | |
| 6 | 0.9993 | 0.9991 | 0.9990 | 0.9989 | 0.9989 | 0.9980 | | | |
| 7 | 0.9998 | 0.9999 | 0.9993 | 0.9998 | 0.9996 | 1.0000 | | | |
| 8 | 0.9999 | 1.0000 | 0.9999 | 1.0000 | 1.0000 | 1.0000 | | | |
| 9 | 1.0000 | 1.0000 | 1.0000 | 1.0000 | 1.0000 | 1.0000 | | | |

most robust employed analyses; (vi) the sensitivity of the CHS detector was also determined with irradiated and non-irradiated samples, the results showing that in both situations the greatest sensitivity is observed in the region of greatest linearity; (vii) the KPCR method had excellent results when applied to the CHS detector spectra; with the control of the number of main components, it was possible to determine how close the R^2 values were to the experimental values; (viii) the results for linearity (both for the method of choosing the wavelength where R^2 occurs, and for KPCR method) can be obtained through mathematical methods and functions; (ix) the RSMPE and MAPE methods for statistical errors were used, and their results showed that the KPCR method is confident and determinant for the maximum agreement between the predictive and measured values, through the number of main components (k). Therefore, it is concluded that the feasibility of employing the CHS detector samples was confirmed in measurements involving photodynamic therapy measurements. Fig. 9 Root Mean Square Percentage Error (RMSPE) and Mean Absolute Percentage Error (MAPE) versus number of principal components from k=1 up to k=9, for **a** 1 mg/ml. **b** 2 mg/ml, **c** 3 mg/ml, **d** 4 mg/ ml, e 5 mg/ml, and f 6 mg/ml concentrations. The standard deviations of the RMSPE and MAPE methods obtained were lower than 1%



Number of principal component (k)

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Declarations

Conflict of interest There are no conflicts to declare.

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