Optimization and characterization of the endogenous production of protoporphyrin IX in a yeast model

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Abstract. The availability of reproducible, convenient, and inexpensive model organisms able to generate predictable levels of endogenous porphyrins, including protoporphyrin IX (PpIX), is essential in photomedicine research. *Saccharomyces cerevisiae* produces endogenous PpIX and was used as a model organism for this study with the aim to maximize endogenous PpIX fluorescence intensity. It was found that PpIX fluorescence was significantly enhanced by administration of 5-aminolevulinic acid (ALA) and 2,2′-bipyridyl. Fluorescence intensity and spectroscopy of PpIX produced endogenously were measured in diluted yeast solutions under various conditions. The optimal protocol was: 5 μM ALA and 1 mM 2,2′-bipyridyl administered synchronously at 32°C. After 3 h, PpIX in yeast demonstrated similar steady-state and time-resolved spectroscopy as that of PpIX in DMSO. Moreover, under hypoxic conditions, the reciprocal lifetime of PpIX delayed fluorescence measured in real time was correlated to the partial pressure of oxygen (pO₂) measured concomitantly with a commercially available pO₂ probe. These data show that yeast can, in optimal conditions, reproducibly generate PpIX. This is of interest in various fields such as photodiagnosis, photodynamic therapy, and photobiomodulation. Use of this model organism focuses on essential mechanisms, without the complexity of a multicellular organism. © 2016 Society of Photo-Optical Instrumentation Engineers (SPIE) [DOI: 10.1117/1.JBO.21.12.125008]

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1 Introduction

Most fundamental biological knowledge is, at least in part, based on the study of model organisms. An ideal model should not only be as relevant as possible, but it should be convenient, inexpensive, and easy to use. In addition, models able to give results as reproducibly as possible are highly desirable.

*Saccharomyces cerevisiae* is one of the most widely studied and extensively used model organisms, with fully characterized genome.1 It displays most of the features of higher eukaryotes (each yeast cell contains all the major organelles also present in the higher eukaryotes). Yet its utilization is simple, economical, and rapid in most cases. For instance, a majority of the knowledge on mitochondrial function and dysfunction were discovered by studying *Saccharomyces cerevisiae.*2 Moreover, yeast belongs to the foremost studied model organisms for nutrient sensing questions.3

5-aminolevulinic acid (ALA), as well as some of its derivatives, has an excellent clinical tolerance and good tumor selectivity in many cases. These prodrugs are clinically approved and used for diagnostic and therapeutic applications, mostly in dermatology, neurosurgery, and urology.4-9 They are precursors of protoporphyrin IX (PpIX), a fluorescent, oxygen-sensitive, and photosensitizing molecule.9,10 Topical (in the form of cream or instillation) or systemic (mostly oral) administration of ALA leads to an intracellular production of PpIX in mammalian cells.11,12 The ability of PpIX to accumulate in certain cancer cells/tissues combined with its phototoxic and fluorescent properties allow its use in photodynamic therapy (PDT) and photodiagnosis (PDD), respectively.5,9,12,13 In mammals, PpIX is produced via the heme biosynthetic pathway, which occurs partially in mitochondria. In all living organisms, heme plays a key role in oxygen sensing and utilization, as well as in regulation of various processes through similar mechanisms.14-16

In yeasts, PpIX is produced through the same biosynthetic mechanism,17 thus making them excellent models to study the modulation of endogenous PpIX production. Heme functions as the internal barometer of oxygen tension in *Saccharomyces cerevisiae* and is dependent on the levels of oxygen in the extracellular environment.18 It also impacts the expression of respiratory function genes and genes, which are responsible for the oxidative damage controlling.19-21 Furthermore, it was showed that *Saccharomyces cerevisiae* is able to produce endogenously PpIX after ALA and 2,2′-bipyridyl addition.22,23

The objective of this study was to develop a model producing endogenous levels of PpIX comparable to those observed in the abovementioned clinical applications. Such a model is of high interest to optimize or study basic mechanisms involved in PDT and PDD, such as the influence of agents or environmental factors.24-26

The study of steady-state and time-resolved spectroscopy of PpIX produced endogenously, in particular, its dependence on environmental factors [temperature and oxygen partial pressure (pO₂)]12,13,25 the development of optical phantoms mimicking the PpIX fluorescence spectroscopy and intensity observed in tumors,26,27 the study of agents modulating the production of ROS, including singlet oxygen, to optimize and/or monitor PDT.28

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Thus, the determination of the conditions reproducibly leading to the maximal PpIX fluorescence intensity produced by *Saccharomyces cerevisiae* was studied and is reported in this paper. It is noteworthy that, although various fluorescent porphyrins with similar spectroscopy account for the typical porphyrin fluorescence spectrum, the name “PpIX” is used throughout to mean “PpIX and other fluorescent porphyrins with similar fluorescence spectrum.” PpIX accumulation in yeast after ALA and 2,2′-bipyridyl administration was visualized by fluorescence microscopy. In addition, the steady-state and time-resolved spectroscopy of PpIX produced in this model was compared to that of PpIX dissolved in DMSO. Finally, the sensitivity of PpIX delayed fluorescence (DF) lifetime on the pO\textsubscript{2} was determined in yeast and in solutions to confirm that PpIX is the main luminophore presenting a long (up to a few hundreds of microseconds) luminescence in this model.

2 Materials and Methods

2.1 Growth Conditions of the Yeast

Yeast cells from *Saccharomyces cerevisiae* YSC2 (Type II, Sigma Aldrich, Switzerland, in the order of USD 1/5 g) 2.5 mg/ml were shaken in a rotary shaker at 210 rpm and 32°C in aqueous glucose (150 mg/ml) solution. It should be noted that, under these conditions, yeast cells are not replicating. This fact does not prevent the production of PpIX, as demonstrated by our results.

2.2 Assessment of the Endogenous PpIX Fluorescence Intensity and Steady-State Fluorescence Spectroscopy of PpIX in Solution

ALA (Sigma Aldrich, Switzerland) was dissolved in distilled water with pH adjusted to pH = 5 by addition of NaOH. 2,2′-bipyridyl (Sigma Aldrich, Switzerland) was dissolved in pure ethanol. Various concentrations of ALA (5 nM to 30 mM) and/or 2,2′-bipyridyl (1 μM to 30 mM; 0.5% v/v) were added to yeast to determine the maximal PpIX fluorescence intensity. Samples were incubated in the rotary shaker for 3 h. During this incubation time, PpIX fluorescence intensity already reached half of its maximal intensity, as described below in more details. Maximal fluorescence intensity was achieved with a combination of ALA (5 μM) and 2,2′-bipyridyl (1 mM). All spectroscopic measurements were performed in 3 ml cuvettes filled with a clear diluted 2.5% solution of yeast. The fluorescence spectroscopy and intensity of the PpIX produced endogenously were measured with a fully calibrated Horiba Fluorolog\textsuperscript{3}-3 spectrophotometer operated in the front face mode and using an excitation at 405 nm with 2-nm excitation slit width and 10-nm emission slit width. These conditions were used to avoid the edge effects and spectral distortion. The temperature of the cuvette holder was controlled by a Neslab (Newington, NH) RTE refrigerated circulator and kept at 32°C (same temperature as in the incubator shaker). To prevent sedimentation, samples were stirred during the measurement. Power illuminating the samples during the spectroscopic measurements was ~400 μW. This power did not photobleach PpIX in yeast, nor did it create a PpIX photoproduct (data not shown). The identification of these conditions is based on a separate study aiming at determining the conditions that are optimal in terms of photobleaching, lack of spectral distortion due to the optical properties of the sample (inner filter effect), and temperature dependence of the PpIX spectroscopy. The intensity of the PpIX fluorescence emission peak at 635 nm was used to assess its relative concentration. All experiments were performed three times and error bars in the figures represent the standard deviation of these three independent measurements. Independent batches of yeasts were prepared for all different experimental conditions. Pure 25 nM PpIX dimethyl ester (Sigma Aldrich, Switzerland) was dissolved in DMSO (Sigma Aldrich, Switzerland). Fluorescence spectrum of PpIX dimethyl ester was measured using the same parameters as for recording fluorescence spectra of PpIX produced endogenously in yeast after 3 h incubation with ALA and 2,2′-bipyridyl.

2.3 PpIX Pharmacokinetics Measurement

PpIX fluorescence intensity in yeast was measured for 48 h after addition of 5 μM ALA and 1 mM 2,2′-bipyridyl using the set-up conditions described in Sec. 2.2. The volume of medium was checked and filled to the initial volume before each measurement to compensate the evaporation, which took place during the 48-h-long measurements.

2.4 Fluorescence Microscopy of Yeast

Yeast cells were placed onto 35-mm culture dishes with integral no. 0 glass cover slip bottoms (MatTek, Ashland, Massachusetts) covered with ECL Cell Attachment Matrix (Millipore, Bedford, Massachusetts). The optimal concentrations of ALA (5 μM) and 2,2′-bipyridyl (1 mM) were added to yeast cells, which were then incubated overnight in the rotary shaker. Cells were stained for mitochondrial membrane potential with 1 μg/ml Rhodamine 123 (ThermoFischer Scientific, Waltham, Massachusetts), green fluorescent, and cell-permeant dye. After 30 min incubation at 32°C with Rhodamine 123, cells were centrifuged (1200 g for 5 min) and visualized with fluorescence microscope (Laboratoryophot-2; Nikon, Japan) equipped with an air-cooled, slow scan 16-bit CCD camera, and mercury-vapor lamp (HB0, 100 W) and an 60X oil objective (Nikon; NA: 1.4). BV-2A Nikon filter cube (excitation: 400 to 440 nm; 455-nm dichroic mirror; emission: 470-nm high-pass filter) was used for PpIX fluorescence and FTTC Nikon filter cube (excitation: 475 to 490 nm; 500- to 540-nm dichroic mirror; emission: 520-nm long-pass filter) for Rhodamine 123 fluorescence. Images were analyzed by Image J software (National Institutes of Health, Bethesda, Maryland).

2.5 Measurement of the PpIX Delayed Fluorescence Lifetimes

DF lifetimes were measured with the optical fiber-based time-resolved setup previously described by Piffaretti et al. Briefly, PpIX molecules were excited with 5-ns light pulses at 405 nm emitted by a nitrogen-laser-pumped dye laser (MNL 100 HP, LTb) at 10 Hz. These pulses were coupled into a quartz optical fiber, which was also used to collect the PpIX emission. This emission was filtered with two high-quality band-pass filters to reject all wavelengths outside 628 ± 25 nm with an OD > 5. To discriminate the prompt fluorescence from the DF, a gated PMT (C1392, R928) was used in such a way that it was blinded during the first 1 μs after the pulse. The resulting PMT signal was sampled (LeCroy instruments LT342, 500 MHz) in temporal windows ranging between 100 μs and 10 ms. The average of
at least 10 sweeps was then fitted with a double exponential decay model. The reciprocal lifetime (inverse of the long lifetime component) was used to derive the pO₂ value according to the Stern–Volmer equation. Acquisition (LabVIEW, National Instruments, Austin, Texas) and analysis (OriginLab®, Northampton, Massachusetts) were automatically performed in real time via a homemade interface.

Yeast samples were incubated overnight with 5 μM ALA and 1 mM 2,2′-bipyridyl in the incubator shaker to reach the maximal PpIX intensity. Yeast solutions were then centrifuged at 1200g for 5 min to further increase the yeast, and hence the PpIX, concentration by a factor eight. The pellet was resuspended in the aqueous glucose (150 mg/ml) solution (5 ml) and the final solution was placed into a flask. Oxygen partial pressure and temperature of the final solution were kept constant (using air bubbling with atmospheric pO₂ and 32°C, respectively), as in the rotary shaker conditions, during all the experiment (unless otherwise specified). Yeast suspension in the flask was circulated via a peristatic pump through a flow cell cuvette (Hellma Analytics: 137-1-40). The optical path length of this cuvette was 1 mm. The total circulated volume was 4 ml and yeast suspension flow speed 5 ml/min. The tip of the measurement fiber was applied in gentle contact to the quartz window of the flow cell. In these conditions, the sensing volume was in the order of 1 μl. Ten excitation pulses (0.3 μJ/pulse, 10 Hz) were sufficient to assess the pO₂ value (∼0.5 mJ cm⁻²/measurement).

In our conditions, PpIX photobleaching can be neglected. This is in accordance with observations in other studies.22,23,30-32 Where 0.2 mM 2,2′-bipyridyl was shown to be the most effective for maximal endogenous PpIX fluorescence intensity in Saccharomyces cerevisiae. Since this yeast strain contains up to 1.5 mM endogenous ALA, no additional ALA was needed. On the other hand, different conditions for maintaining yeast were used in this study than in Strakhovskaya et al. and ALA addition was essential. Individual contributions to the endogenous PpIX production of ALA or 2,2′-bipyridyl are discussed later. Various concentrations of ALA and 2,2′-bipyridyl were incubated with yeast and fluorescence spectra of PpIX were recorded. Figure 2 shows the ALA and 2,2′-bipyridyl concentration dependence curve, respectively. It was found that PpIX fluorescence intensity was reproducibly the highest when it was incubated with 5 μM ALA and 1 mM 2,2′-bipyridyl. It should be noted that in the presence of 2,2′-bipyridyl, the absolute fluorescence intensity of the PpIX was ∼15 times higher than without 2,2′-bipyridyl addition. This is in accordance with observations in other studies.24-34

Optimal concentration of 2,2′-bipyridyl determined in this study (1 mM) is higher than that was reported in other papers,22,23,36 where 0.2 mM 2,2′-bipyridyl was shown to be the most effective for maximal endogenous PpIX fluorescence intensity. Higher concentration of 2,2′-bipyridyl led to inhibition of cell respiration and growth.23 In this study, this effect was observed at concentrations higher than 1 mM. Difference in the optimal 2,2′-bipyridyl concentration and essential presence of ALA is probably due to different growth conditions of the yeast. As Fig. 2 shows, there is an ALA and 2,2′-bipyridyl dose-dependent PpIX accumulation in the yeast cells. PpIX fluorescence decreases after reaching the threshold of ALA (750 μM) or 2,2′-bipyridyl concentration (30 mM). The shape of the ALA dose-dependent curve is in agreement with those reported previously for mammalian cells.13 Optimal ALA concentration at which mammalian cells produce the highest intensity of PpIX is different for various cell lines.24,35,36

![Fig. 1](https://ebooks.spiedigitallibrary.org/journals/Journal-of-Biomedical-Optics/125008-3.png)

Fig. 1: Evolution of the pO₂ in a yeast solution in the presence of high glucose (filled square) and in yeast grown in YPD (filled circle)—See text for conditions. pO₂ was measured with the Oxylab probe. The lines are drawn for visual support only.

3 Results and Discussion

3.1 Determination of the Optimal Conditions Inducing Maximal PpIX Fluorescence Intensity

The main aim of this study was to determine the optimal conditions, at which Saccharomyces cerevisiae cells reproducibly produce the maximal PpIX fluorescence intensity. It was reported that the presence of a metal-chelating agent (2,2′-bipyridyl) enhances endogenous PpIX fluorescence intensity in yeast.22,23,30-32 Same effects were observed with another iron-chelator, Desferal®, which effects were extensively studied on mammalian cells.24 2,2′-bipyridyl decreases generation of metalloporphyrins and increases accumulation of free porphyrins by complexing with a number of bivalent metals including iron,31 which would otherwise bind preferentially to porphyrins and turn them into nonfluorescent moieties. One of the most frequently used mediums for growing yeast is rich medium (YPD; containing yeast extract, peptone, and dextrose/glucose).23 However, YPD has a strong autofluorescence interfering with PpIX fluorescence (data not shown). To simplify the experiments, the absence of PpIX photobleaching can be neglected. This is confirmed by the fact that the probability that a given yeast cell be probed is in the order of 1/8 min⁻¹.

pO₂ in the medium slowly decreases (with increasing amount of yeast cells) due to the replication of yeast cells.

According to a report by Strakhovskaya et al.,23 0.2 mM 2,2′-bipyridyl induced remarkable increase of endogenous PpIX fluorescence intensity in Saccharomyces cerevisiae. Since this yeast strain contains up to 1.5 mM endogenous ALA, no additional ALA was needed. On the other hand, different conditions for maintaining yeast were used in this study than in Strakhovskaya et al. and ALA addition was essential. Individual contributions to the endogenous PpIX production of ALA or 2,2′-bipyridyl are discussed later. Various concentrations of ALA and 2,2′-bipyridyl were incubated with yeast and fluorescence spectra of PpIX were recorded. Figure 2 shows the ALA and 2,2′-bipyridyl concentration dependence curve, respectively. It was found that PpIX fluorescence intensity was reproducibly the highest when it was incubated with 5 μM ALA and 1 mM 2,2′-bipyridyl. It should be noted that in the presence of 2,2′-bipyridyl, the absolute fluorescence intensity of the PpIX was ∼15 times higher than without 2,2′-bipyridyl addition. This is in accordance with observations in other studies.24-34

Optimal concentration of 2,2′-bipyridyl determined in this study (1 mM) is higher than that was reported in other papers,22,23,36 where 0.2 mM 2,2′-bipyridyl was shown to be the most effective for maximal endogenous PpIX fluorescence intensity. Higher concentration of 2,2′-bipyridyl led to inhibition of cell respiration and growth.23 In this study, this effect was observed at concentrations higher than 1 mM. Difference in the optimal 2,2′-bipyridyl concentration and essential presence of ALA is probably due to different growth conditions of the yeast. As Fig. 2 shows, there is an ALA and 2,2′-bipyridyl dose-dependent PpIX accumulation in the yeast cells. PpIX fluorescence decreases after reaching the threshold of ALA (750 μM) or 2,2′-bipyridyl concentration (30 mM). The shape of the ALA dose-dependent curve is in agreement with those reported previously for mammalian cells.13 Optimal ALA concentration at which mammalian cells produce the highest intensity of PpIX is different for various cell lines.24,35,36
As discussed above when yeast cells are grown in the presence of YPD, addition of 2,2′-bipyridyl alone leads to a small endogenous PpIX fluorescence intensity [Fig. 3(a)] albeit much smaller than after incubation with ALA and 2,2′-bipyridyl. It is also interesting to note that in the presence of 2,2′-bipyridyl alone there is significantly higher PpIX fluorescence intensity than when in the presence of ALA alone. Finally, although very weak, a detectable signal corresponding to PpIX is observed in the absence of both ALA and 2,2′-bipyridyl [dashed dotted line in Fig. 3(a)].

The results presented in Fig. 3(a) demonstrate the strong synergy between ALA and 2,2′-bipyridyl to potentiate the PpIX fluorescence intensity in yeast in the presence of high glucose concentration. Since it cannot be assumed that the internalization and metabolism kinetics of these two compounds are identical, a separate preliminary study was conducted to assess influence of different ALAs and 2,2′-bipyridyl administration on the PpIX fluorescence. Figure 3(b) shows the emission spectra of PpIX fluorescence intensity at various ALA and 2,2′-bipyridyl administration conditions. Surprisingly, it was found that a synchronous administration of ALA (5 μM) and 2,2′-bipyridyl (1 mM) led to a maximal PpIX fluorescence [Fig. 3(b)]. Incubation of yeast with ALA alone for 1 h with additional 2,2′-bipyridyl administration led to lower PpIX fluorescence. This effect was even lower when yeast cells were first incubated with 2,2′-bipyridyl with following ALA addition.

Based on all these results, it was concluded that the optimal conditions leading to maximal endogenous PpIX fluorescence intensity were an incubation of yeast cells for 3 h after synchronous administration of 5 μM ALA and 1 mM 2,2′-bipyridyl.

### 3.2 Pharmacokinetics of the PpIX Fluorescence

A pharmacokinetics study was carried out to determine how long was PpIX fluorescence present in yeast cells and at what time the highest PpIX fluorescence intensity was reached after ALA and 2,2′-bipyridyl application. Figure 4 shows the curve of the endogenous PpIX fluorescence over time.

Half the maximal PpIX fluorescence intensity was reached 3 h after ALA and 2,2′-bipyridyl application. The maximal intensity was reached 12 h after ALA and 2,2′-bipyridyl administration and plateaued for ~4 h. The fluorescence of PpIX then decreased slowly with almost no PpIX present 44 h after ALA and 2,2′-bipyridyl application. This slow decrease may be due to the depletion of glucose in the culture medium and subsequent possible disruption of the yeast cells, which resulted in the leak of PpIX into the solution. Since PpIX is lipophilic, its fluorescence was quenched in the aqueous solution.

### 3.3 Steady-State and Time-Resolved Fluorescence Spectroscopy of PpIX in Solution or Produced Endogenously

Figure 5 shows the normalized fluorescence spectrum of 25 nM PpIX dimethyl ester, which was used as a reference to the fluorescence spectrum of PpIX in yeast after 3 h incubation with 5 μM ALA and 1 mM 2,2′-bipyridyl.

Fluorescence spectra of PpIX in yeast and PpIX dimethyl ester dissolved in DMSO are similar, with maximum fluorescence intensity at 635 nm for PpIX present in yeast and 632 nm for PpIX dimethyl ester dissolved in DMSO (excitation: 405 nm). The small shift in the fluorescence maximum could be due to environmental/derivation effects. It should also be noted that PpIX is not the only porphyrin produced by yeast and that in the presence of ALA and 2,2′-bipyridyl application: coproporphyrin contributes to the fluorescence spectrum with fluorescence maximum at 620 nm. Presence of uroporphyrin can be expected as well with very similar fluorescence properties to those of coproporphyrin. Nevertheless, the fluorescence lifetime measured in yeast was very similar (16.58 ± 0.06611 ns) to the fluorescence lifetime of PpIX dimethyl ester in DMSO (16.9 ± 0.0436 ns). This observation suggests that PpIX produces the most important contribution to the yeast fluorescence in our conditions.
The fluorescence spectrum of PpIX in yeast presents a peak at 590 nm. It has been reported that, when ALA is applied to yeast in certain conditions, in particular in combination with iron chelators, it can induce the generation of Zn- and Mg-porphyrins, which present fluorescence maxima at 580 to 605 nm. This effect was consistently observed when iron chelators were coadministered. Moreover, it was found that PpIX can be converted to Zn-PpIX. However, 2,2'-bipyridyl induces accumulation mainly of PpIX, Zn-PpIX is synthetized enzymatically through Zn-chelatase activity of ferrochelatase. Therefore, it can reasonably be hypothesized that the peak which can be observed at 590 nm is due to the endogenous conversion of PpIX to metallo-porphyrins (Zn- and Mg-) and preferential binding of 2,2'-bipyridyl to the bivalent iron. The fluorescence lifetime of Zn-PpIX is in the order of 2.1 ns, as reported in Ref. 40. This value is significantly different than ours (5.37 ± 0.07061 ns) when an excitation wavelength of 405 nm was used.

### 3.4 Fluorescent Microscopy of the Yeast

Fluorescent microscopy was used to study the mitochondrial membrane potential of yeast in the presence or absence of ALA and 2,2'-bipyridyl (Fig. 6). Yeast cells were stained with Rhodamine 123, a mitochondrial potential-dependent probe. Depolarization of mitochondria induces red spectral shift and intensity quenching of the probe. Fifteen-hour incubation with 5 \( \mu \)M ALA and 1 mM 2,2'-bipyridyl did not induce any significant difference in the Rhodamine 123 intensity. In addition, distribution of PpIX in the yeast cells could be observed.
PpIX was localized in mitochondria and cytoplasm, which is in agreement with previous descriptions of the PpIX biodistribution.\textsuperscript{44} The presence of PpIX in mitochondria was confirmed by partial colocalization of PpIX and Rhodamine 123 [Fig. 6(b)]. This observation is of interest since it suggests that the pO$_2$ can be determined specifically in this organelle by measuring the PpIX DF lifetime, as described in the following chapter.

### 3.5 Influence of the pO$_2$ on the PpIX Delayed Fluorescence Lifetime in Yeasts and in Solution.

The triplet state of the most porphyrin photosensitizers, including endogenous PpIX is quenched by molecular oxygen. Unfortunately, certain porphyrins, such as PpIX, do not emit detectable phosphorescence \textit{in vitro} or \textit{in vivo}. Since the inverse intersystem crossing transition takes place between PpIX triplet and singlet states, the measurement of its triplet state’s lifetime can be obtained via the time-resolved detection of its DF. The lifetime of this DF yields information on the surrounding pO$_2$ via the Stern–Volmer relationship.\textsuperscript{12} Endogenous PpIX, which is produced in the mitochondria, can be therefore used as a mitochondrial-pO$_2$ probe.\textsuperscript{25}

In order to verify whether these concepts that are well established in mammalian cells are also valid in yeast, the pO$_2$ was also measured synchronously with a commercially available thermo-compensated optical oxygen device (Optronix; Oxylab pO$_2$E\textsuperscript{TM}; probe: BF/OT/E) where the probe (with 350-μm tip

![Fig. 6](https://example.com/fig6.png)

**Fig. 6** Fluorescence images of yeast stained with rhodamine 123 for mitochondrial potential. (a) Control yeast; (b) yeast incubated for 15 h with 5 μM ALA and 1 mM 2,2′-bipyridyl. See text for further conditions.

![Fig. 7](https://example.com/fig7.png)

**Fig. 7** DF-pO$_2$ dependence of the PpIX produced endogenously in yeast; comparison between the pO$_2$ measured with the Oxylab probe (gray line) and the reciprocal lifetime of the PpIX delayed fluorescence (black line). Inset: correlation between this reciprocal lifetime and the pO$_2$ measured with the Oxylab probe. The lines are drawn for visual support only.
diameter) was plunged directly into the medium. Measurements were compared with the PpIX DF reciprocal long lifetime obtained in yeast. In Fig. 7, a time series of pO2 measurements based on these two approaches were performed during the air bubbling (environmental pO2 ~90 mmHg) phases. These results indicate that, in our conditions, the consumption of oxygen by yeast is more important than the oxygen delivery due to its limited diffusion and solubilization in the solution.

After stopping the air bubbling, yeast depleted the oxygen, which led to hypoxic condition (0 mmHg). As expected, the evolution of the PpIX DF reciprocal lifetime is nicely correlated with the pO2 measured with the Oxylab probe. Plotting reciprocal lifetime against values obtained with the Oxylab probe showed a linear correlation (Fig. 7, inset).

These measurements enabled to derive the Stern–Volmer quenching constant, as described by Piffaretti et al. This derivation resulted in: $K_{SV} \approx 1500 \text{ mmHg}^{-1} \text{s}^{-1}$. The lifetime without oxygen ($\tau_o$) was determined by purging the residual oxygen with nitrogen bubbling during 5 min. Remaining oxygen traces were removed by the irradiation of the sample (1 J/cm² at 405 nm). $\tau_o$ was taken when the DF lifetime reached a constant value ($\tau_0 = 388 \pm 40 \mu$s).

PpIX Stern–Volmer constants reported in the literature differ depending on the calibration protocol and microenvironments. $K_{SV}$ was found to be $\approx 600 \text{ mmHg}^{-1} \text{s}^{-1}$ for HeLa cells and $\approx 1000 \text{ mmHg}^{-1} \text{s}^{-1}$ for lung fibroblast cells with $\tau_o$ around 500 and 300 $\mu$s, respectively. The concordance between the values reported by Mik et al. in cells and these present values in yeast support the fact that yeast cells produce PpIX and that its microenvironment is similar to that of mammalian cells.

A temporal shift of about 10 s is observed between the two curves presented in Fig. 7. This shift can reasonably be assigned to the time response of the Oxylab probe setup since this time is indicated as being “<10 s” according to the specifications. This illustrates the advantage of using the PpIX DF to measure the pO2 in quasi real time. This approach opens the way to dynamic measurements of the pO2 in different tissular and cellular compartments, including mitochondria, to obtain information of fundamental interest regarding their metabolism and respiration.

4 Conclusion

Overall, we have established conditions to use Saccharomyces cerevisiae as a reproducible, convenient, and inexpensive model organism to produce PpIX. Because our conditions are characterized by the absence of yeast replication, thus constant pO2, experiments can be performed over several dozen hours. Fluorescence spectroscopy of the endogenous PpIX in yeast resembles that of PpIX in DMSO. In addition, fluorescence lifetime of PpIX in yeast is very similar to that of PpIX in DMSO. This suggests that PpIX is the main fluorescent component produced by yeast after the administration of its precursor in our conditions. The values we have obtained for the Stern–Volmer constants ($K_{SV}$ and $\tau_o$) also support the fact that PpIX is endogenously produced in yeast and is responsible for the DF we have observed.

Using a yeast model enables focusing on essential mechanisms, without the complexity of a multicellular organism. In addition, such a model is of high interest in PDT and PDD research for optimization and basic mechanistic studies. In particular, a version of this model loaded with appropriate amounts of light absorbing and scattering particles could be used as phantom to mimic tumors containing PpIX, a useful tool to optimize certain cancer photodetection apparatus.

Disclosures

All the authors have no conflict of interest to declare.

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References

18. J. Mattoon et al., “Oxygen and catabolite regulation of hemoprotein biosynthesis in the yeast Saccharomyces cerevisiae,” in Biosynthesis of...
Joniová et al.: Optimization and characterization of the endogenous production of...


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Georges Wagnières obtained his PhD in physics from the Swiss Federal Institute of Technology at Lausanne (EPFL) in 1992 and did a postdoc in the Wellman Laboratories of Photomedicine, Boston, from 1993 to 1994. He obtained a master’s degree in management of technology in 2001. He has led a group active in biomedical optics at EPFL since 1994. He has authored more than 220 papers (130 peer-reviewed) and filed 15 patents.