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Tailing miniSOG: structural bases of the complex photophysics of a flavin-binding singlet oxygen photosensitizing protein

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miniSOG is the first flavin-binding protein that has been developed with the specific aim of serving as a genetically-encodable light-induced source of singlet oxygen (¹O₂). We have determined its 1.17 Å resolution structure, which has allowed us to investigate its mechanism of photosensitization using an integrated approach combining spectroscopic and structural methods. Our results provide a structural framework to explain the ability of miniSOG to produce ¹O₂ as a competition between oxygen- and protein quenching of its triplet state. In addition, a third excited-state decay pathway has been identified that is pivotal for the performance of miniSOG as ¹O₂ photosensitizer, namely the photo-induced transformation of flavin mononucleotide (FMN) into lumichrome, which increases the accessibility of oxygen to the flavin FMN chromophore and makes protein quenching less favourable. The combination of the two effects explains the increase in the ¹O₂ quantum yield by one order of magnitude upon exposure to blue light. Besides, we have identified several surface electron-rich residues that are progressively photo-oxidized, further contributing to facilitate the production of ¹O₂. Our results help reconcile the apparent poor level of ¹O₂ generation by miniSOG and its excellent performance in correlative light and electron microscopy experiments.

miniSOG (for mini Singlet Oxygen Generator)¹ is a 106 amino acid flavin-binding protein that generates ¹O₂ under exposure to blue light. It was originally developed by Shu and coworkers for correlative light and electron microscopy (CLEM) as it both fluoresces and catalyzes the photo-oxidation of diaminobenzidine (DAB), providing high-resolution images¹. Novel applications are being actively developed since^{2–5}. miniSOG was engineered from the LOV2 (Light, Oxygen and Voltage) domain of *Arabidopsis thaliana* phototropin 2¹. Proteins based on LOV domains are blue-light photoreceptors that form a light-induced and reversible flavin-cysteine covalent adduct that consumes the energy of the excited state⁶. Replacement of the cysteine residue by an alanine or glycine avoids the formation of the covalent bond and leads to a fluorescent protein^{7,8}. miniSOG contains six mutations as compared to its precursor, two of them involving residues surrounding the chromophore. Its cofactor FMN is ubiquitously found in nature^{9,10} and generates ¹O₂ with high quantum yield (Φ_{Δ})¹¹, but also other reactive oxygen species (ROS)¹².

Close inspection of the photophysical and photosensitizing properties of miniSOG reveals a number of striking observations: (1) its Φ_{Δ} is much lower than that of FMN (0.03 vs. 0.51)^{11,13,14}; (2) the lifetime (τ_T) of triplet miniSOG (³miniSOG*) is much shorter than that of FMN in nitrogen-saturated solutions (33.6 μ s¹⁵ vs 200 μ s¹⁶); (3) oxygen quenching is less efficient than for FMN ($\tau_T^{\text{air}} = 31.3 \mu$ s¹⁵ vs 3.1 μ s in air-saturated solutions); (4) in

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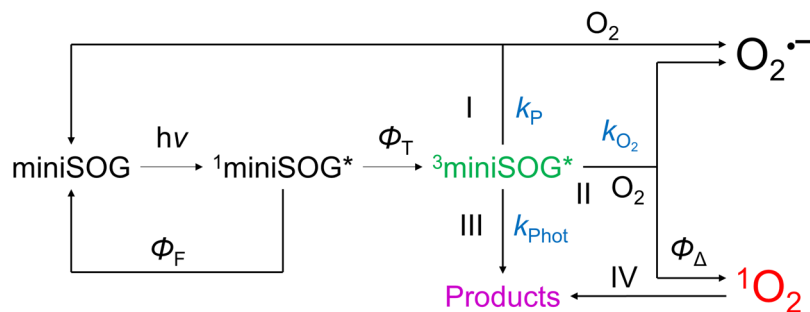


Figure 1. Model of oxygen photosensitization by miniSOG. Φ_F : fluorescence quantum yield, Φ_T : triplet state quantum yield, k_p : protein quenching rate constant, k_{Phot} : photoproduct formation rate constant, k_{O_2} : oxygen quenching rate constant, Φ_Δ : singlet oxygen quantum yield. I, II, III, and IV indicate the three deactivation and one oxidation pathways discussed in the main text.

addition to $^1\text{O}_2$ it also produces superoxide ($\text{O}_2^{\bullet-}$)^{12,14}; (5) it undergoes a remarkable transformation upon exposure to light, whereby Φ_Δ increases 10-fold (to ~ 0.3) and $\tau_{\text{T}}^{\text{air}}$ shortens by 10-fold (to $\sim 3 \mu\text{s}$)^{13,14}. The absence of a structure of miniSOG so far had prevented to rationalize these observations, which we have attempted here using a combined structural and photophysical approach.

Based on the extensive data present in the literature and the photophysical and structural results presented herein, a mechanism of excited-state deactivation of miniSOG can be proposed that involves three main pathways (Fig. 1). The shorter lifetime of $^3\text{miniSOG}^*$ compared to $^3\text{FMN}^*$ indicates that protein quenching is a major mechanism of triplet decay (pathway I). Its rate constant k_p is largely determined by electron transfer with nearby electron-rich residues¹⁷. Quenching of the singlet state can be safely ruled out since no shortening of the fluorescence lifetime or decrease in the fluorescence quantum yield are observed relative to free FMN. In the presence of oxygen, a second decay pathway (pathway II) is possible, namely oxygen quenching to produce $^1\text{O}_2$ (energy transfer) or $\text{O}_2^{\bullet-}$ (electron transfer), as observed for FMN in solution¹². It is also possible to produce $\text{O}_2^{\bullet-}$ by reaction of oxygen with a radical anion formed during protein quenching in pathway I. Finally, miniSOG undergoes a photoinduced transformation (pathway III, rate constant k_{Phot}), for which we provide here a detailed description for the first time.

Results and Discussion

High resolution crystal structure of miniSOG. We have solved the structure of miniSOG at 1.17 Å resolution (Fig. 2a and Supporting Information), which shows an increase in rigidity of the environment of the chromophore compared to that in the LOV2 domain, the location of potential quenchers of the excited states of FMN, and the phosphoribityl tail of FMN lying in a tunnel bridging the bulk solvent and the chromophore encased in the core of the protein (Fig. 2b). The latter hinders oxygen access to the isoalloxazine ring. The presence halfway through the tunnel of a chloride ion, which can be a good mimic of molecular oxygen^{18,19}, suggests that oxygen diffusion can occur.

Deactivation mechanism of miniSOG triplet excited state (Pathways I and II). The values of the relevant rate constants for pathways I and II can be inferred from the $^3\text{miniSOG}^*$ lifetime measurements. Comparison of the decay rate constant ($1/\tau_T$) of miniSOG and SOPP3, the miniSOG mutant with the longest τ_T reported so far (3.3 ms in nitrogen-saturated solutions)¹⁷ allows us to estimate the rate constant for protein quenching ($k_p = k_{\text{T}}^{\text{N}_2} - k_{\text{T}}^{\text{N}_2, \text{SOPP3}}$, Table 1). SOPP3 is a miniSOG variant, which encases the same chromophore FMN and, most importantly, lacks most of the electron-rich residues present in the vicinity of the flavin in miniSOG. Hence, protein quenching of the triplet chromophore in SOPP3 is essentially suppressed, which makes SOPP3 a convenient model for the study of the contribution of protein quenching in miniSOG. Likewise, the pseudo-first order rate constant for oxygen quenching ($k_{\text{O}_2} = k_{\text{T}}^{\text{Air}} - k_{\text{T}}^{\text{N}_2}$) can be estimated from τ_T data in air- and nitrogen-saturated solutions (Table 1).

Comparison of k_p and k_{O_2} in Table 1 reveals that protein quenching (pathway I) is the main triplet deactivation pathway, removing 93% of the triplets in air-saturated solutions $k_p/(k_p + k_{\text{O}_2})$. Oxygen only quenches 7% of the triplets, which limits Φ_Δ to $0.6 \times 0.07 = 0.042$ (Eq. 1), in excellent agreement with the experimental value. It can therefore be concluded that the modest Φ_Δ of miniSOG is due to an unfavorable combination of low k_{O_2} and high k_p values, as proposed previously¹⁷.

$$\Phi_\Delta = \Phi_T \times \frac{k_{\text{O}_2}}{k_p + k_{\text{Phot}} + k_{\text{O}_2}} \approx \Phi_T \times \frac{k_{\text{O}_2}}{k_p + k_{\text{O}_2}} \quad (1)$$

Our structural results above suggest that the low value of k_{O_2} is due to the steric hindrance of the ribityl tail within the tunnel which provides oxygen access to the FMN. Regarding k_p , the miniSOG structure shows that six electron-rich residues are positioned within 8.2 to 10.2 Å from the isoalloxazine ring, namely Tyr30, Tyr73, Trp81, His85, Met89 and Tyr98, and are thus close enough to the chromophore to act as electron-transfer quenchers of $^3\text{miniSOG}^*$ ²⁰. In addition, four hydrophilic residues, Glu44, Asp72, Asp82 and Glu103, form hydrogen bonds

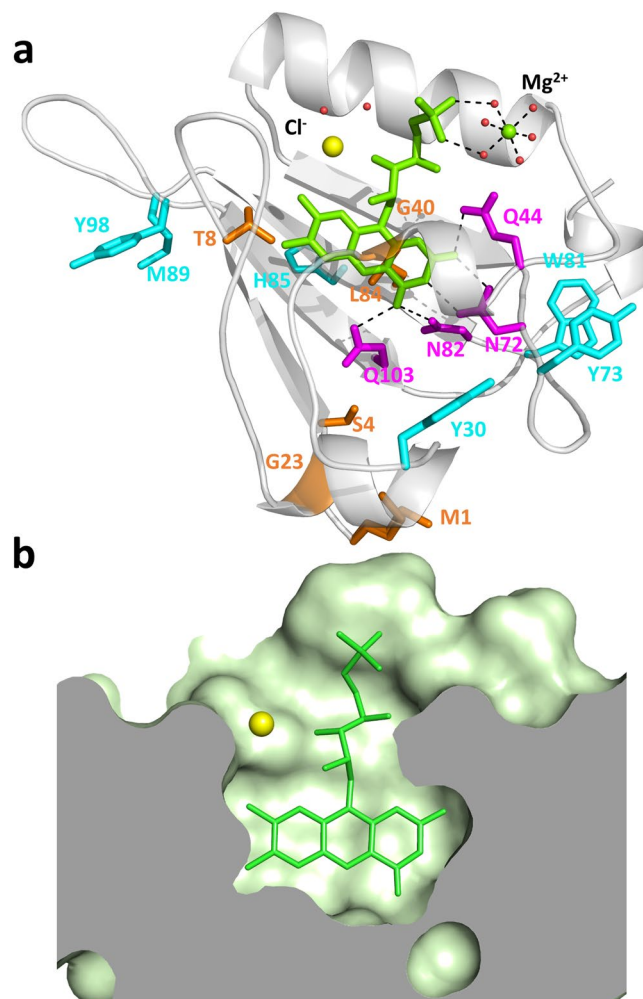


Figure 2. High-resolution crystallographic structure of miniSOG. **(a)** Secondary structure (white) represented with FMN (green), chloride (yellow), magnesium (green) and water molecules (red). Represented residues: mutations from LOV2 (orange), residues hydrogen-bonded to the FMN ring (magenta), and potential quenchers of $^3\text{FMN}^*$ (cyan). **(b)** Topology of the FMN-binding site.

Parameter	in D ₂ O	in H ₂ O	Pathway	References
τ_S	5.0 ns	4.9–5.5 ns		13,15,43
Φ_F	0.43	0.37–0.44		1,14,15,43,44
Φ_T	0.6	0.6		15
Φ_Δ	0.03–0.04	0.03–0.05		13,14,23,31,44
$k_T^{\text{N}_2}$	$2.41 \times 10^4 \text{ s}^{-1}$	$2.98 \times 10^4 \text{ s}^{-1}$		15
k_p	$2.38 \times 10^4 \text{ s}^{-1}$	$2.95 \times 10^4 \text{ s}^{-1\text{b}}$	I	This work
k_T^{Air}	$2.59 \times 10^4 \text{ s}^{-1}$	$3.19 \times 10^4 \text{ s}^{-1}$		15
k_{O_2}	$1.8 \times 10^3 \text{ s}^{-1}$	$2.3 \times 10^3 \text{ s}^{-1}$	II	This work
k_{Phot}	6.0 s^{-1}	—	III	This work

Table 1. Photophysical properties of miniSOG in D₂O- and H₂O-based phosphate buffer. D₂O was used to increase the singlet oxygen lifetime, thus boosting the reactions and processes in which singlet oxygen is involved and facilitating its detection⁴⁵. ^aAssuming the same value of $k_T^{\text{N}_2}$,_{SOPP3} in H₂O and D₂O.

with FMN, and may thus enhance protein quenching and $\text{O}_2^{\bullet-}$ formation²¹. Replacing selectively these residues should lead to a lengthening of the triplet lifetime of miniSOG²² and hence to a higher fraction of triplets being trapped by oxygen, thus to a higher Φ_Δ value. In fact, some of these positions have already been mutated in light of their capacity of direct electron transfer from the FMN: such miniSOG mutants show considerably longer τ_T values (e.g., 196 μs for miniSOG Q103L (SOPP)²³, 1.1 ms for miniSOG W81F (Supplementary Fig. S2), and 3.3 ms for SOPP3¹⁷ in oxygen-free solutions) and larger Φ_Δ values (0.25, 0.33 and 0.6, respectively), in agreement with

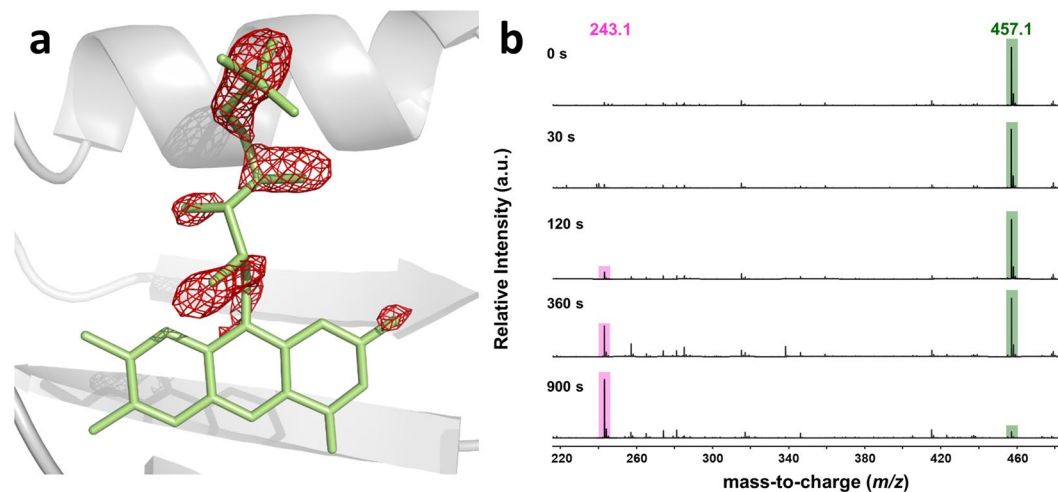


Figure 3. Blue-light induced structural changes on the chromophore of miniSOG. (a) 2.0 Å resolution difference Fourier map calculated between non-irradiated and irradiated parts of a miniSOG crystal contoured at a -3.0σ level (magenta) superimposed on the FMN molecule (green). (b) ESI-TOF mass spectra acquired in the low mass range ($m/z < 500$) of miniSOG progressively irradiated with blue-light.

Eq. 1. It is worth noting also that miniSOG produces more $O_2^{\bullet-}$ than free FMN¹², which indicates that the radical anion pathway contributes to the production of $O_2^{\bullet-}$. Indeed, SOPP shows an 8-fold higher Φ_{Δ} value than miniSOG but only a 1.3 higher yield of $O_2^{\bullet-}$ ²³. Thus, removal of hydrophilic side chains in the vicinity of the chromophore should strongly reduce the relative formation of $O_2^{\bullet-}$ vs. 1O_2 .

Consequences of blue-light irradiation of miniSOG on its FMN chromophore (Pathway III). In light of Eq. 1, the observed 10-fold decrease in τ_T and similar increase in Φ_{Δ} upon extended photolysis suggest severe changes in both k_p and k_{O_2} . Blue-light (440 nm) irradiation of a miniSOG crystal at $10 \text{ W}\cdot\text{cm}^{-2}$ led to a five-fold decrease of the fluorescence signal over a 30 min course (Supplementary Fig. S3) and was gentle enough to keep diffraction around 2.0 Å resolution while affecting a sufficient fraction of molecules so that structural alterations could be visualized in electron density maps. A difference Fourier map calculated from non-irradiated and irradiated parts of a crystal revealed the loss of electron density all along the ribityl tail of the FMN (Fig. 3a), strongly suggesting its cleavage. Besides, Electrospray ionisation time-of-flight (ESI-TOF) mass spectrometry performed on irradiated protein samples show (Fig. 3b) the progressive disappearance of the FMN peak at $m/z = 457.1$ in favor of a peak at $m/z = 243.1$.

To get further insights into the photoconversion, we performed additional photophysical investigations. Besides the already-known shortening of τ_T and increase in Φ_{Δ} , exposure of miniSOG samples to light induces photobleaching of the FMN chromophore and appearance of new absorption and fluorescence bands (Fig. 4a,b). The leaching out of FMN from miniSOG was routinely checked and could be safely ruled out. The quantum yield and rate constant of pathway III could be estimated (Table 1, Supplementary Fig. S1). Noteworthy, the Φ_{Δ} value increases when the photoconverted miniSOG is excited at 355 nm, but remains essentially constant when probed at 473 nm (Fig. 4c,d).

Phototransformation of FMN to lumichrome (LC) is consistent with all of the above observations: (1) LC is a photodegradation product of flavins in aqueous solutions²⁴; (2) the observed mass loss upon irradiation matches the molar mass difference between FMN (456.3 Da) and LC (242.2 Da); (3) LC absorbs and fluoresces at shorter wavelengths than FMN, (Fig. 5); (4) LC lacks the phosphoribityl tail of FMN, which facilitates the access of molecular oxygen to the isoalloxazine ring, resulting in the increase of k_{O_2} and the decrease of τ_T ; (5) LC is a worse electron acceptor than FMN, hence protein quenching is less favored. The $\Delta_r G^{\circ}$ value for quenching of $^3\text{riboflavin}^*$ by tryptophan is $-86.5 \text{ kJ}\cdot\text{mol}^{-1}$ (riboflavin is analogous to FMN except for the phosphate group) while is more positive for $^3\text{LC}^*$, $-67.2 \text{ kJ}\cdot\text{mol}^{-1}$ ²⁵; (6) finally, LC is also an excellent 1O_2 photosensitizer^{25–27}, hence the combination of a higher k_{O_2} and a lower k_p yield a higher Φ_{Δ} value (Eq. 1) when excited at 355 nm but not at 473 nm, where LC barely absorbs.

Consequences of blue-light irradiation of miniSOG on its amino acid residues (Pathway IV).

We investigated if our structural data could also support a decrease in k_p . Indeed, the $2F_{\text{obs}} - F_{\text{calc}}$ electron density map of blue-light irradiated miniSOG reveals the unambiguous oxidation of three surface residues during irradiation (Fig. 6a, Supplementary Fig. S4). Tyr73 has been partially converted to a γ -peroxytyrosine. The loss of electron density on Trp81 is compatible with the formation of *N*-formylkynurenine (NFK), a well-known tryptophan oxidation product^{28,29}. Finally, His85 can be modeled by either a singly, or a doubly oxidized histidine, namely 2-oxo-histidine and 2,4-dioxo-histidine. Mass spectrometry analysis of blue-light irradiated miniSOG samples reveals sequential additions of +16 mass units to the native protein mass of 13882.0 Da, consistent with increasing oxidation steps of the protein (Fig. 6b). All three structural modifications account for six of the eight additional

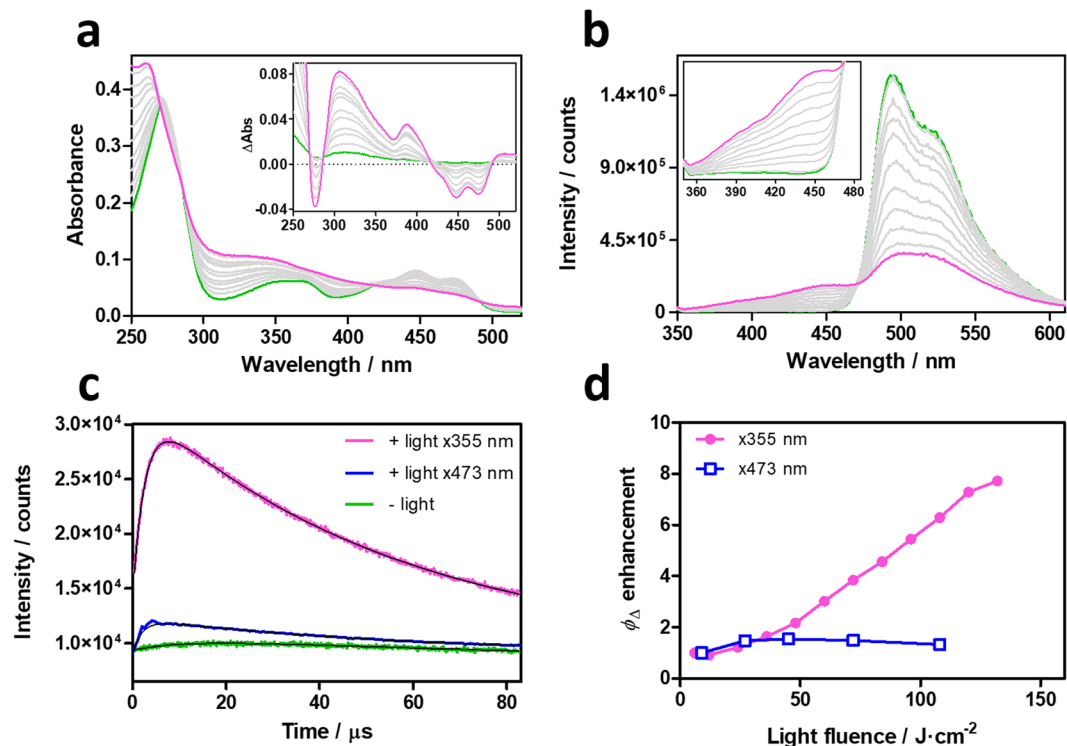


Figure 4. Spectroscopic characterization of photoconverted miniSOG. Evolution of miniSOG's absorbance (a) and fluorescence (b) upon laser irradiation at 355 nm. Insets show the difference absorption spectra before and after irradiation, and a zoomed-in image of the new fluorescence bands. (c) Time-resolved NIR $^1\text{O}_2$ phosphorescence decays of native (green) and photoconverted miniSOG excited at 473 nm (blue) or 355 nm (magenta). (d) Observed Φ_{Δ} enhancement at 473 nm (blue) and 355 nm (magenta).

oxygen atoms evidenced in the mass spectrometry analysis. The two non-assigned additions could correspond to oxidation of Tyr30, Met89 or Tyr98, although we did not observe unambiguous oxidation of these residues. Oxidation of Tyr73, His85 and Trp81 eliminates potential quenchers of $^3\text{miniSOG}^*$, thereby decreasing the value of k_p . According to Eq. 1, this should contribute to an increase in Φ_{Δ} . However, since protein oxidation (pathway IV) occurs simultaneously to FMN \rightarrow LC transformation, which also increases Φ_{Δ} , it is not possible to ascertain the individual contribution of both effects.

Finally, oxidation of tryptophan into NFK could contribute to the increased Φ_{Δ} value observed at 355 nm since NFK is a potent singlet oxygen photosensitizer ($\Phi_{\Delta} = 0.17$)³⁰. However, the W81F mutant shows a doubled Φ_{Δ} ($=0.33$) already before photolysis on account of its lower k_p value (Eq. 1), indicating that the potential benefits of producing NFK as secondary photosensitizer are of minor value as compared to the effect of eliminating a protein quencher.

Conclusion

We have performed an extensive structural characterization of miniSOG in the dark and its photoproduct formed in the presence of molecular oxygen, which led us to explain in structural terms the details of its complex photo-physical behavior. miniSOG is initially moderately efficient towards $^1\text{O}_2$ generation because of a combination of limited oxygen accessibility and ^3FMN quenching by electron-rich side chains. Prolonged irradiation to blue light leads to several structural alterations of miniSOG, which include photodegradation of FMN into LC and oxidation of the quenching side chains. All this results in an increase of Φ_{Δ} when photoconverted miniSOG is excited at the wavelengths where the formed LC absorbs. Formation of LC liberates the access of molecular oxygen to the alloxazine ring and reduces protein quenching of the triplet state, while oxidized electron-rich side chains cannot quench the triplet state of the chromophore. The competition between oxygen quenching and protein quenching of flavin triplet state seems to be a general feature of flavin-binding proteins³¹, hence our results will be useful to guide the evolution of such a protein towards retaining or gaining a specific function. Finally, our results explain the apparent discrepancy between the poor level of singlet oxygen generation by miniSOG, which had been consistently measured at low light fluences, and its efficiency in CLEM experiments, in which the singlet oxygen generation capability of miniSOG is exploited over its whole lifetime.

Methods

Chemical compounds. Riboflavin-5'-monophosphate sodium salt hydrate (FMN) (Chemochroma), Lumichrome (Santa Cruz Biotechnology), tris(hydroxymethyl)aminomethane (Merck), sodium chloride (Panreac Applichem), imidazole (Merck), L-arabinose (Sigma Aldrich) and ampicillin (Sigma Aldrich) were used as received. Phosphate-buffered saline (PBS) or deuterated dPBS solutions were prepared by dissolving the required amount of a PBS tablet (Sigma Aldrich) in milliQ water or deuterium oxide (Sigma-Aldrich).

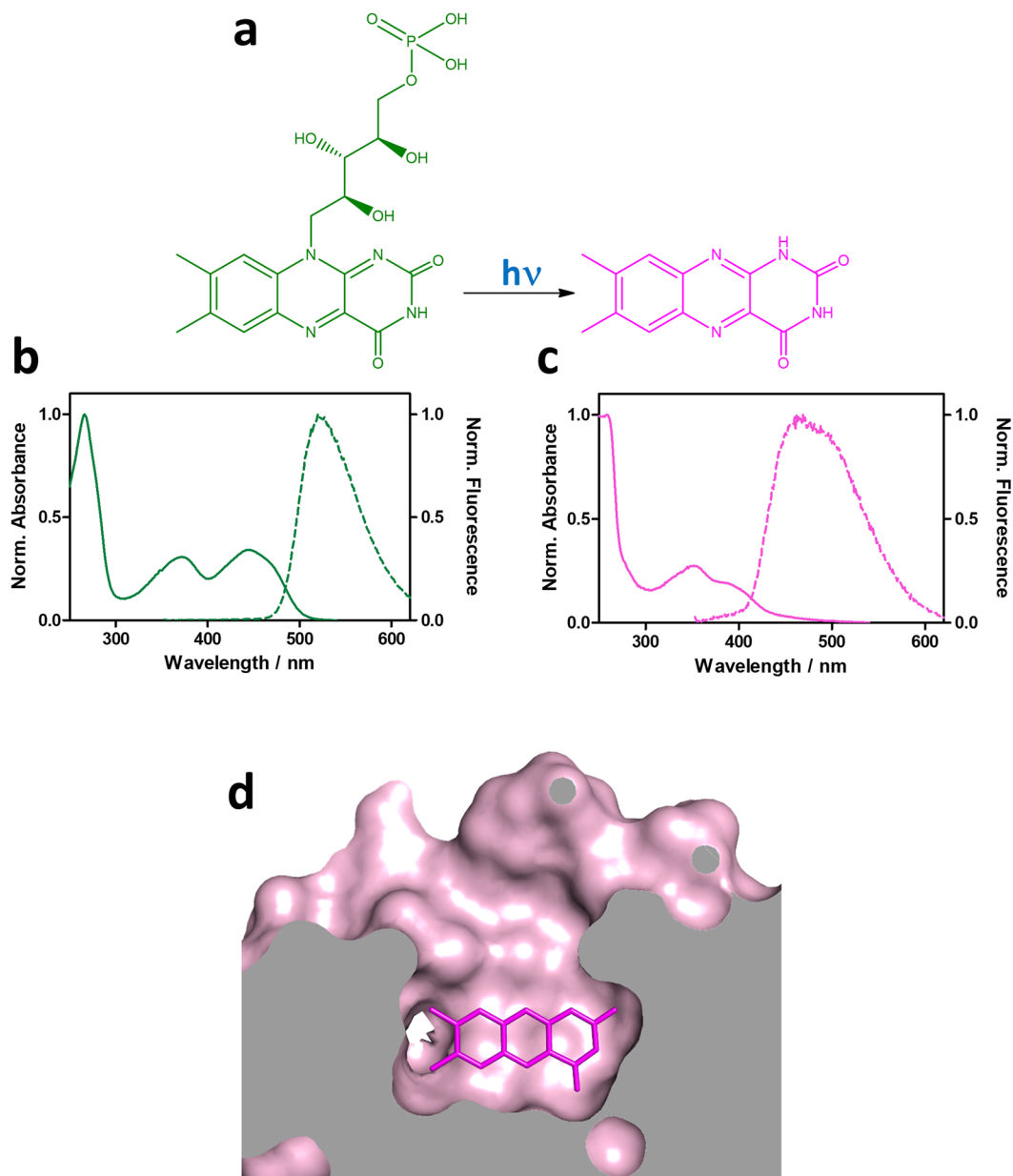


Figure 5. FMN is photoconverted to lumichrome. (a) Photoconversion of FMN (green) to LC (magenta). Normalized absorption and fluorescence spectra of (b) FMN and (c) LC. (d) Topology of blue-light irradiated miniSOG showing the increased access to the alloxazine ring.

Expression and purification. Genes coding for a C-terminal 6xHis-tagged recombinant miniSOG and miniSOG W81F were inserted in a pBad expression vector and over-expressed in *Escherichia coli* CodonPlus (DE3) RIL Cells (Stratagene) or in TOP10 cells (Invitrogen). Bacterial cells were grown in LB broth medium containing 1 mM Ampicillin. At an OD_{600} of approximately 0.6, expression of recombinant protein was induced by the addition of L-arabinose and cells were grown for an additional 24 h at 25 °C. Cells were pelleted by centrifugation (4000 g, 4 °C, 30 min), re-suspended in *buffer A* (20 mM Tris-HCl pH 8.0, 500 mM NaCl), complemented with complete protease inhibitors-EDTA (Roche) and disrupted using a micro-fluidizer. The soluble fraction was recovered by centrifugation (40,000 g, 4 °C, 30 min), and loaded on a 1 mL Ni-NTA superflow column (Qiagen) pre-equilibrated with *buffer A*. The His-tagged protein was eluted with 150 mM imidazole in *buffer A*. Fractions containing purified proteins were pooled and concentrated to a volume of 0.5 mL using Centricon devices (Amicon 10 kDa cut-off) and loaded onto a size-exclusion chromatography column (Hiload Superdex75 10/300, GE Healthcare) for the final step of the purification procedure. The column was equilibrated with 20 mM Tris-HCl pH 8.0 and the pooled peak fractions were concentrated to 4 mg·mL⁻¹. Protein expression and purification was always performed in the dark or under red light. The purity of the protein solutions was confirmed by SDS-PAGE. The final concentration was determined by UV-vis absorption spectroscopy using a molar absorption coefficient of 14 mM⁻¹·cm⁻¹ at 448 nm.

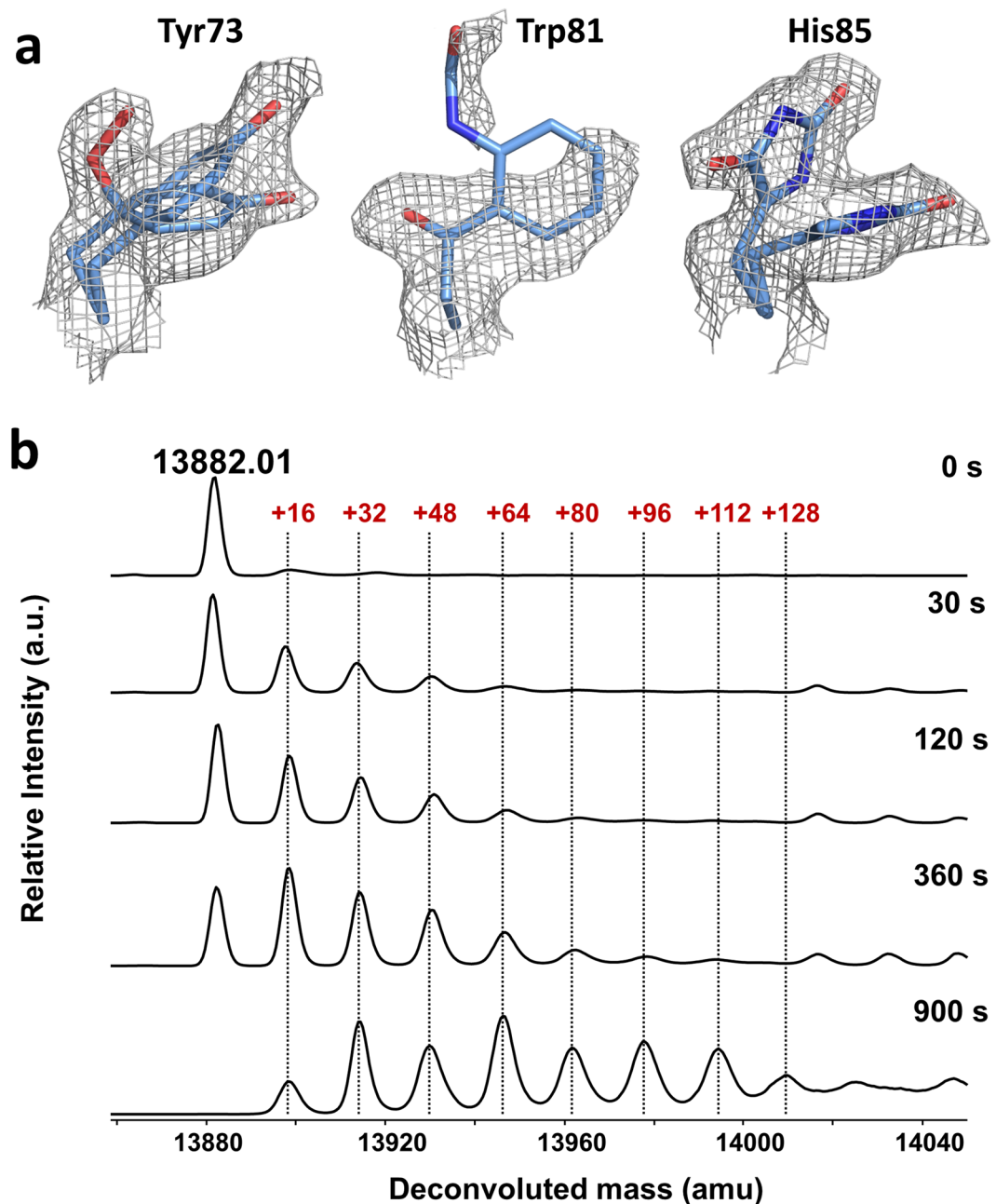


Figure 6. Identification of protein residue oxidation in blue-light irradiated miniSOG. (a) $2F_{\text{obs}} - F_{\text{calc}}$ electron density maps contoured at a 1.0, 0.6 and 1.0 σ level superimposed on the refined model of residues Tyr73, Trp81, and His85, respectively. (b) Deconvoluted ESI-TOF mass spectra of miniSOG progressively irradiated with blue-light. The peak at 13882.0 Da corresponds to the native protein.

Spectroscopic measurements. All spectroscopic measurements were performed using quartz cuvettes (Hellma) under magnetic stirring and at room temperature. Absorption spectra were recorded on a double beam Cary 6000i spectrophotometer (Varian). Fluorescence spectra were measured on Fluoromax-4 spectrofluorometer (Horiba). Time-resolved near-infrared (NIR) phosphorescence signals at 1275 nm were measured using a customized PicoQuant Fluotime 200 lifetime system. Briefly, an AO-Z-473 solid state AOM Q-switched laser (Changchun New Industries Optoelectronics Technology Co., China) was used for excitation at 473 nm, working at 1.0 kHz repetition rate at 473 nm. The average power that reached the sample was conveniently modulated by neutral density filters. For excitation at 355 nm, the frequency-tripled output of a diode-pumped pulsed Nd:YAG laser (FTSS355-Q, Crystal Laser, Berlin, Germany) was used, working at 1 kHz repetition (0.5 mW, or 5 mW, 1 ns pulse width). An uncoated SKG-5 filter (CVI Laser Corporation, Albuquerque, U.S.A.) was placed at the exit port of the laser to remove any NIR component. The luminescence exiting from the sample was filtered by a 1100 nm long-pass filter (Edmund Optics, York, U.K.) and a narrow bandpass filter at 1275 nm (bk-1270-70-B, bk Interferenzoptik, Germany) to remove any scattered laser radiation and isolate the $^1\text{O}_2$ emission. A

TE-cooled near-IR sensitive photo multiplier tube assembly (H9170-45, Hamamatsu Photonics Hamamatsu City, Japan) in combination with a multichannel scaler (NanoHarp 250, PicoQuant GmbH, Germany) was used as photon-counting detector. The time-resolved $^1\text{O}_2$ emission decays were analyzed by fitting Eq. 2³² to the data using GraphPad Prism 5.

$$S_{(t)} = S_{(0)} \frac{\tau_{\Delta}}{\tau_{\Delta} - \tau_T} \left(e^{-\frac{t}{\tau_{\Delta}}} - e^{-\frac{t}{\tau_T}} \right) \quad (2)$$

τ_T and τ_{Δ} are the lifetimes of the photosensitizer triplet state and of $^1\text{O}_2$, respectively, and $S_{(0)}$ is a quantity proportional to Φ_{Δ} . Φ_{Δ} values were determined by comparing the $S_{(0)}$ values of optically-matched solutions of the corresponding flavoprotein and FMN at 473 nm (Eq. 3)³².

$$\phi_{\Delta, \text{protein}} = \frac{S_{(0)\text{protein}}}{S_{(0)\text{FMN}}} \phi_{\Delta, \text{FMN}} \quad (3)$$

FMN was taken as reference photosensitizer with $\Phi_{\Delta} = 0.51$ in PBS¹¹ and 0.57 in dPBS³³.

Transient absorption spectra were monitored by nanosecond laser flash photolysis using a Q-switched Nd-YAG laser (Surelite I-10, Continuum) operating at the 3rd harmonic. The luminescence exiting from the sample was filtered by a 610 nm long-pass filter (CVI Laser Corporation, NM, USA). Changes in the sample absorbance were detected at 715 nm using a Hamamatsu R928 photomultiplier to monitor the intensity variations of an analysis beam produced by a 75 W short arc Xe lamp (USHIO) and spectral discrimination was obtained using a PTI 101 monochromator. The signal was fed to a Lecroy Wavesurfer 454 oscilloscope for digitizing and averaging (typically 10 shots) and finally transferred to a PC for data storage and analysis. The system was controlled using the in-house-developed LKS software (LabView, National Instruments).

Determination of k_{phot} . The rate constant for the photoproduct formation has been determined measuring the progressive loss of miniSOG in solution as a function of the absorbed light dose at 473 nm using Eq. 4. The slope of the resulting plot yielded the photobleaching quantum yield Φ_{phot} , from which the rate constant for photobleaching was calculated as:

$$k_{\text{phot}} = \frac{\Phi_{\text{phot}}}{\tau_{\text{Air}}} \quad (4)$$

X-ray crystallography. *Crystallization procedures.* miniSOG was concentrated to 4 mg·mL⁻¹. The crystallization condition consisted of 100 mM Tris-HCl pH 8.0, 20 mM MgCl₂, 28% PEG 4000, 0 or 15 mM CoCl₂ at 20 °C. Crystals appeared and grew to final size after 1–2 days.

Data collection and processing. X-ray data were collected on beamlines ID23-1³⁴ and ID29³⁵ of the ESRF and were indexed, integrated, merged and scaled using the XDS software package³⁶. Molecular replacement was carried out using the model structure of LOV2 (PDB ID: 4eep) with the program Phaser MR³⁷. Structure refinement was performed using Refmac5³⁸ and manual improvement of the model with Coot³⁹. The native structure of miniSOG was used as a starting model for model building of bleached-miniSOG. Data collection and refinement statistics are presented in Supplementary Table S1. Structure analysis and representation were performed with Pymol⁴⁰.

Preparation of photobleached miniSOG samples. *miniSOG crystals.* A single miniSOG crystal was soaked in a cryoprotectant solution containing 20% of glycerol then harvested with a nylon loop. The crystal was exposed to 440 nm laser (10 W·cm⁻²) on the ID29S-Cryobench setup⁴¹ at room temperature using a HCl1 humidity control device⁴². Spectra were recorded at a 1 Hz rate. After 30 min of total exposure, the crystal was flashcooled in liquid nitrogen.

miniSOG solutions. Fresh miniSOG or miniSOG W81F solutions in air-saturated deuterated PBS were illuminated at 355 nm (~5 mW·cm⁻²) or 473 nm (~15 mW·cm⁻²) for elapsed irradiation times. Absorption and fluorescence spectra as well as time-resolved $^1\text{O}_2$ phosphorescence decays were recorded at different time intervals of cumulative irradiation.

Liquid chromatography-mass spectrometry (LC-MS). Liquid Chromatography Electrospray Ionization Mass Spectrometry (LC/ESI-MS) was carried out on a 6210 LC/ESI-TOF mass spectrometer interfaced with a binary HPLC pump system (Agilent Technologies). The mass spectrometer was calibrated in the positive ion mode with ESI-L (low concentration tuning mix, Agilent Technologies) before each series of measurements, the calibration providing mass accuracy <1 ppm in the 100–3200 *m/z* range. All solvents used were HPLC grade: water and acetonitrile (LC-MS Chromasolv, Sigma-Aldrich); formic acid was from Acros Organics (puriss., p.a.).

Data acquisition was carried out in the positive ion mode with spectra in the profile mode and mass spectra were recorded in the 130–2000 *m/z* range. The mass spectrometer was operated with the following experimental settings: ESI source temperature was set at 325 °C; nitrogen was used as drying gas (5 L/min) and as nebulizer gas (30 psi); the capillary needle voltage was set at 3500 V. Fragmentor value was of 250 V and skimmer of 65 V. The instrument was operated in the 2 GHz (extended dynamic range) mode and spectra acquisition rate was of 1 spectrum/s.

Before analysis, miniSOG samples were diluted to a final concentration of 10 μM in acetonitrile/water/formic acid (50:50:0.1, *v/v/v*) and infused directly in the mass spectrometer by a syringe pump at a flow rate of 10 $\mu\text{l}/\text{min}$. A blank run was carried out infusing only protein buffer diluted at the same ratio as the protein sample in the same solvent system.

The MS data were acquired and processed with the MassHunter workstation software (Data acquisition v.B.04.00, Qualitative analysis with Bioconfirm v.B.07.00, Agilent Technologies).

Protein Data Bank accession codes. The structures of non-irradiated and blue-light irradiated miniSOG have been deposited in the Protein Data Bank under entry codes 6GPU and 6GPV, respectively.

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Author Contributions

J.T. performed and analyzed the spectroscopic and photophysical investigations. C.L., S.A., G.G. and A.R. performed and analyzed the structural work. L.S. performed and analyzed the mass spectrometry experiments. C.F. and X.S. participated to the discussion of the data. S.N., G.G. and A.R. designed the study, discussed the data and wrote the manuscript with J.T. and C.F.

Additional Information

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