Data in Brief 11 (2017) 617-627



Contents lists available at ScienceDirect

Data in Brief

journal homepage: www.elsevier.com/locate/dib

Data Article



Fluorescence spectroscopy of roGFP2-based redox probes responding to various physiologically relevant oxidant species *in vitro*

Alexandra Müller^{a,*}, Jannis F. Schneider^a, Adriana Degrossoli^a, Nataliya Lupilova^a, Tobias P. Dick^b, Lars I. Leichert^a

 ^a Institute of Biochemistry and Pathobiochemistry – Microbial Biochemistry, Ruhr-University Bochum, 44780 Bochum, Germany
^b Division of Redox Regulation, DKFZ-ZMBH Alliance, German Cancer Research Center (DKFZ).

Im Neuenheimer Feld 280, 69120 Heidelberg, Germany

A R T I C L E I N F O

Article history: Received 28 February 2017 Accepted 7 March 2017 Available online 11 March 2017

Keywords: Genetically encoded redox probes Polysulfides HOCI Peroxynitrite glutathione H₂O₂ Nitric oxide

ABSTRACT

This article contains representative fluorescence excitation spectra of roGFP2-based probes used for ratiometric analysis of redox changes as presented in the article "Systematic *in vitro* assessment of responses of roGFP2-based probes to physiologically relevant oxidant species" [1]. The recombinant probes roGFP2, roGFP2-Orp1, and Grx1-roGFP2 were exposed to various oxidative and nitrosative species, including hydrogen peroxide (H_2O_2), aldrithiol-2 (AT-2), glutathione disulfide (GSSG), hypochlorous acid (HOCI), S-nitrosoglutathione (GSNO), peroxynitrite (ONOO⁻), potassium polysulfide (K_2S_x), spermine NONOate (SperNO), and diethyl amino NONOate (DeaNO) at different molar ratios. Fluorescence excitation spectra of the probes were recorded in the excitation wavelength range between 350 and 500 nm and for a total of 60 min. Analysis and interpretation of the data is presented in an associated article [1].

© 2017 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

DOI of original article: http://dx.doi.org/10.1016/j.freeradbiomed.2017.02.044

* Corresponding author.

http://dx.doi.org/10.1016/j.dib.2017.03.015

E-mail address: alexandra.mueller-2@ruhr-uni-bochum.de (A. Müller).

^{2352-3409/© 2017} The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

618

Subject area	Biology
More specific subject area	Redox Biology
Type of data	figure, raw data file (.csv)
How data was acquired	JASCO FP-8500 fluorescence spectrometer equipped with a Peltier thermo-holder 'EHC-813' (JASCO, Darmstadt, Germany)
Data format	Raw
Experimental factors	Reduction of the probes with DTT
Experimental features	Assessment of changes in the fluorescence excitation characteristics of roGFP2- based probes upon oxidant treatment
Data source location	Bochum, Germany, Latitude 51.4445974, Longitude 7.258836
Data accessibility	Spectral data are displayed in Figs. 1–10. Associated raw data can be accessed as . csv text files in the supplementary data section

Specifications Table

Value of the data

• The influence of diverse oxidative and nitrosative species on roGFP2-based probes is compared side-by-side.

• Spectral responses of roGFP2, roGFP2-Orp1 and Grx1-roGFP2 are compared side-by-side

- Full spectra are recorded every minute for 60 min.
- The data delineate probe redox responses under strictly controlled *in vitro* conditions.

1. Data

Each experiment represents a time series of 60 fluorescence excitation spectra with one spectrum recorded per minute. The first spectrum shows the fluorescence excitation spectrum prior to treatment. Following the addition of oxidative or nitrosative species, spectral changes of roGFP2, roGFP2-Orp1, and Grx1-roGFP2 were recorded for a total of 60 min. Autoxidation of roGFP2-Orp1 under aerobic conditions is shown in Fig. 1. Experiments involving roGFP2-Orp1 were thus performed under anaerobic conditions, while measurements with roGFP2 and Grx1-roGFP2 were performed under aerobic conditions. Reference spectra were recorded using 2 μ M of control reductant (DTT or GSH), or $2 \mu M$ of control oxidant (AT-2) (Fig. 2). The specificity of the two fusion probes roGFP2-Orp1 and Grx1-roGFP2 was tested using 2 μ M H₂O₂ and GSSG (Fig. 3). Spectral responses to 2 μ M of various other oxidative and nitrosative species were recorded (Fig. 4). Unfused roGFP2 was treated with increasing concentrations of H₂O₂, HOCl, and ONOO⁻ to determine the minimal oxidant concentration eliciting a spectral response (Figs. 5–7). The response of the three probes to treatment with $100 \,\mu$ M of the above tested oxidants was recorded (Figs. 8, 9). To rule out artifacts, non-redoxsensitive eGFP was treated with all compounds used in this study (Fig. 10). The raw data for all figures (excitation wavelength vs. fluorescence intensity values) is made available as comma separated value (.csv) text files.



Fig. 1. Response of the three probes to air oxygen. Each figure displays the dynamics of 60 measured emission spectra (one spectrum per minute; E_m =510 nm) in the excitation wavelength range of 350–500 nm. First (0 min) and last spectra (60 min) are displayed in green and red, respectively. The probes were added to PBS buffer at a final concentration of 0.2 μ M (A–E) or expressed in *E. coli* (F). Measurements were either done under aerobic conditions (A–C) or under fully anaerobic conditions (D). One volume of aerobic buffer corresponding to 100 μ M O₂ was added after measurement of the first spectrum (E).

2. Experimental design, materials and methods

Purification and reduction of roGFP2, roGFP2-Orp1, and Grx1-roGFP2, and preparation of oxidants is described in [1]. Purification of eGFP, heterologously expressed in *E. coli* BL21 from plasmid pET16b_eGFP SC+ [2] was essentially performed as described for the roGFP2-base probes [1]. roGFP2 or Grx1-roGFP2 was added to a 1,500 μ L QS fluorescence cuvette with a stirring bar containing 1 mL of buffer (PBS, 5 mM EDTA, pH 7.4) at a final concentration of 0.2 μ M. Measurements of roGFP2-Orp1 were performed under anaerobic conditions as described in [1]. Measurements were done at 20 °C in a JASCO FP-8500 fluorescence spectrometer equipped with a Peltier thermo-holder 'EHC-813' at 20 °C for 60 min under continuous stirring. Measurement parameters were as follows: 510 nm (Em), 350–500 nm (Ex), 5 nm slit width (Ex/Em), medium sensitivity. After a single recorded spectrum, oxidants were added at the indicated final concentrations and a time series of 59 additional spectra was recorded. Expression of roGFP2-Orp1 in *E. coli* MG1655 (Fig. 1F) was performed as described in [1].



Fig. 2. Response of the three probes to DTT, AT-2 and GSH. The probes were added to PBS buffer to a final concentration of 0.2 μ M. A single spectrum (green) was recorded prior to addition of DTT (A), AT-2 (B), or GSH (C) to a final concentration of 2 μ M. The responses of the probes were measured for a total of 60 min. Each figure displays 60 single spectra with one spectrum per minute. The last recorded spectrum is shown in red.



Fig. 3. Response of the three probes to H_2O_2 and GSSG. A single spectrum (green) of roGFP2, roGFP2-Orp1 and Grx1-roGFP2 at a final concentration of 0.2 μ M was measured prior to the addition of 2 μ M H₂O₂ (A) or 2 μ M GSSG (B). Subsequently, another spectrum was recorded every minute for a total of 60 min. The last recorded spectrum is shown in red.



Fig. 4. Response of the three probes to various oxidative and nitrosative agents. 0.2μ M of the probes were added to PBS. After the measurement of one spectrum (green), the indicated compounds were added to a final concentration of 2μ M. Subsequently, a single spectrum was measured every minute for a total of 60 min. The last recorded spectrum is shown in red.



Fig. 5. Response of unfused roGFP2 to increasing concentrations of H_2O_2 . A single spectrum of roGFP2 (0.2 μ M) was recorded (green). Oxidation was measured for a total of 60 min after addition of the indicated H_2O_2 concentrations (one spectrum each minute). The last recorded spectrum is shown in red.



Fig. 6. Response of unfused roGFP2 to increasing concentrations of HOCI. HOCI concentrations between 2 μ M and 2 mM were applied to roGFP2 after measurement of the first spectrum (green). Subsequently, a total of 60 spectra (one spectrum each minute) were recorded. The last recorded spectrum is shown in red.



Fig. 7. Response of unfused roGFP2 to peroxynitrite. Peroxynitrite concentrations between 2 μ M and 200 μ M were applied to 0.2 μ M roGFP2 after measurement of the first spectrum (green). Subsequently, a total of 60 spectra (one spectrum each minute) were recorded. The last recorded spectrum is shown in red.



Fig. 8. Response of the three probes to a 500-fold molar excess of H_2O_2 , GSSG and HOCI. The probes were measured at a concentration of 0.2 μ M. 100 μ M of oxidant were added after measurement of a single spectrum (green). Each figure displays the dynamics of a total of 60 recorded spectra (one spectrum per minute). The last recorded spectrum is shown in red.



Fig. 9. Response of the three probes to various nitrosative agents. The indicated compounds were added at a final concentration of 100 μ M after measurement of the first spectrum (green). Subsequently, a total of 60 spectra were recorded. The last recorded spectrum is shown in red.



Fig. 10. Response of eGFP to the compounds used in this study. The response of eGFP to the compounds used in this study was measured to rule out general effects of these compounds on GFP fluorescence characteristics. eGFP was added to PBS to a final concentration of 0.2 μ M. After the measurement of one spectrum (green), 2 μ M of the compounds were added and a total of 60 spectra were recorded. The last recorded spectrum is shown in red.

Acknowledgments

We thank Franz Narberhaus and Johanna Rosmanith for providing plasmid pET16b_eGFP SC+. A.D. acknowledges support from CAPES/CNPq (Ciência sem Fronteiras postdoctoral fellowship) (248836/2013-7). L.I.L. and T.P.D. are supported by grants LE 2905/1-1 and DI 731/3-1 under the DFG Priority Program "Dynamics of thiol-based redox switches in cellular physiology" (SPP1710).

Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at http://dx.doi. org/10.1016/j.dib.2017.03.015.

References

- A. Müller, J.F. Schneider, A. Degrossoli, N. Lupilova, T.P. Dick, L.I. Leichert, Systematic *in vitro* assessment of responses of roGFP2-based probes to physiologically relevant oxidant species, Free Radic. Biol. Med. 106 (2017) 329–338.
- [2] M. Wieland, J.S. Hartig, Improved aptazyme design and in vivo screening enable riboswitching in bacteria, Angew. Chem. Int. Ed. Engl. 47 (2008) 2604–2607.