

Alma Mater Studiorum Università di Bologna Archivio istituzionale della ricerca

Structure illumination microscopy imaging of lipid vesicles in live bacteria with naphthalimide-appended organometallic complexes

This is the final peer-reviewed author's accepted manuscript (postprint) of the following publication:

Published Version:

Structure illumination microscopy imaging of lipid vesicles in live bacteria with naphthalimide-appended organometallic complexes / Maria Ranieri A.; Vezzelli M.; Leslie K.G.; Huang S.; Stagni S.; Jacquemin D.; Jiang H.; Hubbard A.; Rigamonti L.; Watkin E.L.J.; Ogden M.I.; New E.J.; Massi M.. - In: ANALYST. - ISSN 0003-2654. - STAMPA. - 146:12(2021), pp. 3818-3822. [10.1039/d1an00363a]

Availability:

This version is available at: https://hdl.handle.net/11585/873630 since: 2022-02-28

Published:

DOI: http://doi.org/10.1039/d1an00363a

Terms of use:

Some rights reserved. The terms and conditions for the reuse of this version of the manuscript are specified in the publishing policy. For all terms of use and more information see the publisher's website.

This item was downloaded from IRIS Università di Bologna (https://cris.unibo.it/). When citing, please refer to the published version.

(Article begins on next page)

This is the final peer-reviewed accepted manuscript of:

RANIERI, A. M.; VEZZELLI, M.; LESLIE, K. G.; HUANG, S.; STAGNI, S.; JACQUEMIN, D.; JIANG, H.; HUBBARD, A.; RIGAMONTI, L.; WATKIN, E. L. J.; OGDEN, M. I.; NEW, E. J.; MASSI, M. STRUCTURE ILLUMINATION MICROSCOPY IMAGING OF LIPID VESICLES IN LIVE BACTERIA WITH NAPHTHALIMIDE-APPENDED ORGANOMETALLIC COMPLEXES. ANALYST 2021, 146 (12), 3818–3822.

The final published version is available online at: https://doi.org/10.1039/D1AN00363A.

Terms of use:

Some rights reserved. The terms and conditions for the reuse of this version of the manuscript are specified in the publishing policy. For all terms of use and more information see the publisher's website.

This item was downloaded from IRIS Università di Bologna (https://cris.unibo.it/)

When citing, please refer to the published version.

Analyst d1an00363a

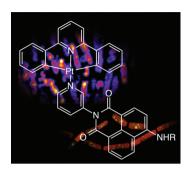
We have presented the graphical abstract image and text for your article below. This briefly summarises your work, and will be presented with your article online.

1

Structure illumination microscopy imaging of lipid vesicles in live bacteria with naphthalimide-appended organometallic complexes

Anna Maria Ranieri, Matteo Vezzelli, Kathryn G. Leslie, Song Huang, Stefano Stagni, Denis Jacquemin, Haibo Jiang, Alysia Hubbard, Luca Rigamonti, Elizabeth L. J. Watkin, Mark I. Ogden,* Elizabeth J. New* and Massimiliano Massi*

Staining bacteria is made more informative with a probe that enables imaging using super-resolution microscopy, and ion nanoscopy.



Please check this proof carefully. Our staff will not read it in detail after you have returned it.

Please send your corrections either as a copy of the proof PDF with electronic notes attached or as a list of corrections. **Do not edit the text within the PDF or send a revised manuscript** as we will not be able to apply your corrections. Corrections at this stage should be minor and not involve extensive changes.

Proof corrections must be returned as a single set of corrections, approved by all co-authors. No further corrections can be made after you have submitted your proof corrections as we will publish your article online as soon as possible after they are received.

Please ensure that:

- The spelling and format of all author names and affiliations are checked carefully. You can check how we have identified the authors' first and last names in the researcher information table on the next page. Names will be indexed and cited as shown on the proof, so these must be correct.
- Any funding bodies have been acknowledged appropriately and included both in the paper and in the funder information table on the next page.
- All of the editor's queries are answered.
- Any necessary attachments, such as updated images or ESI files, are provided.

Translation errors can occur during conversion to typesetting systems so you need to read the whole proof. In particular please check tables, equations, numerical data, figures and graphics, and references carefully.

Please return your **final** corrections, where possible within **48 hours** of receipt following the instructions in the proof notification email. If you require more time, please notify us by email to analyst@rsc.org.

Funding information

Providing accurate funding information will enable us to help you comply with your funders' reporting mandates. Clear acknowledgement of funder support is an important consideration in funding evaluation and can increase your chances of securing funding in the future.

We work closely with Crossref to make your research discoverable through the Funding Data search tool (http://search.crossref.org/funding). Funding Data provides a reliable way to track the impact of the work that funders support. Accurate funder information will also help us (i) identify articles that are mandated to be deposited in **PubMed Central (PMC)** and deposit these on your behalf, and (ii) identify articles funded as part of the **CHORUS** initiative and display the Accepted Manuscript on our web site after an embargo period of 12 months.

Further information can be found on our webpage (http://rsc.li/funding-info).

What we do with funding information

We have combined the information you gave us on submission with the information in your acknowledgements. This will help ensure the funding information is as complete as possible and matches funders listed in the Crossref Funder Registry.

If a funding organisation you included in your acknowledgements or on submission of your article is not currently listed in the registry it will not appear in the table on this page. We can only deposit data if funders are already listed in the Crossref Funder Registry, but we will pass all funding information on to Crossref so that additional funders can be included in future.

Please check your funding information

The table below contains the information we will share with Crossref so that your article can be found *via* the Funding Data search tool. Please check that the funder names and grant numbers in the table are correct and indicate if any changes are necessary to the Acknowledgements text.

Funder name	Funder's main country of origin	Funder ID (for RSC use only)	Award/grant number
University of Sydney	Australia	501100001774	Unassigned
Curtin University	Australia	501100001797	Unassigned
Australian Research Council	Australia	501100000923	Unassigned

Researcher information

Please check that the researcher information in the table below is correct, including the spelling and formatting of all author names, and that the authors' first, middle and last names have been correctly identified. Names will be indexed and cited as shown on the proof, so these must be correct.

If any authors have ORCID or ResearcherID details that are not listed below, please provide these with your proof corrections. Please ensure that the ORCID and ResearcherID details listed below have been assigned to the correct author. Authors should have their own unique ORCID iD and should not use another researcher's, as errors will delay publication.

Please also update your account on our online <u>manuscript submission system</u> to add your ORCID details, which will then be automatically included in all future submissions. See <u>here</u> for step-by-step instructions and more information on author identifiers.

First (given) and middle name(s)	Last (family) name(s)	ResearcherID	ORCID iD
Anna	Maria Ranieri		
Matteo	Vezzelli		
Kathryn G.	Leslie		
Song	Huang		
Stefano	Stagni	J-4969-2012	0000-0002-7260-4845
Denis	Jacquemin	E-9020-2011	0000-0002-4217-0708

Haibo	Jiang		
Alysia	Hubbard		
Luca	Rigamonti	O-8941-2015	0000-0002-9875-9765
Elizabeth L. J.	Watkin		0000-0002-4881-7234
Mark I.	Ogden	A-1521-2008	0000-0001-5317-1637
Elizabeth J.	New	C-9934-2009	0000-0002-2310-254X
Massimiliano	Massi	G-1709-2011	0000-0001-6949-4019

Queries for the attention of the authors

Journal: Analyst Paper: d1an00363a

Title: Structure illumination microscopy imaging of lipid vesicles in live bacteria with naphthalimide-appended organometallic complexes

For your information: You can cite this article before you receive notification of the page numbers by using the following format: (authors), Analyst, (year), DOI: 10.1039/d1an00363a.

Editor's queries are marked like this $\boxed{\text{Q1}}$, $\boxed{\text{Q2}}$, and for your convenience line numbers are indicated like this 5, 10, 15, ...

Please ensure that all queries are answered when returning your proof corrections so that publication of your article is not delayed.

Query Reference	Query	Remarks
Q1	Funder details have been incorporated in the funder table using information provided in the article text. Please check that the funder information in the table is correct.	
Q2	Please confirm that the spelling and format of all author names is correct. Names will be indexed and cited as shown on the proof, so these must be correct. No late corrections can be made.	
Q3	Ref. 4e: Please check that the last name for the 2nd author is displayed correctly.	

Analyst



5

10

15

20

25

30

40

45

COMMUNICATION

10 **Cite this:** DOI: 10.1039/d1an00363a

Structure illumination microscopy imaging of lipid vesicles in live bacteria with naphthalimide-appended organometallic complexes†

Q2

45

50

Received 1st March 2021, Accepted 10th May 2021

DOI: 10.1039/d1an00363a

rsc.li/analyst

Anna Maria Ranieri, ^a Matteo Vezzelli, ^b Kathryn G. Leslie, ^c Song Huang, ^d Stefano Stagni, ^e Denis Jacquemin, ^f Haibo Jiang, ^d Alysia Hubbard, ^d Luca Rigamonti, ^e Elizabeth L. J. Watkin, ^e Mark I. Ogden, ^e * Elizabeth J. New ^f * and Massimiliano Massi ^f * ^a

20 There is a lack of molecular probes for imaging bacteria, in comparison to the array of such tools available for the imaging of mammalian cells. Here, organometallic molecular probes have been developed and assessed for bacterial imaging, designed to 25 have the potential to support multiple imaging modalities. The chemical structure of the probes is designed around a metalnaphthalimide structure. The 4-amino-1,8-naphthalimide moiety, covalently appended through a pyridine ancillary ligand, acts as a luminescent probe for super-resolution microscopy. On the other 30 hand, the metal centre, rhenium(ı) or platinum(ıı) in the current study, enables techniques such as nanoSIMS. While the rhenium(i) complex was not sufficiently stable to be used as a probe, the platinum(II) analogue showed good chemical and biological stability. Structured illumination microscopy (SIM) imaging on live 35 Bacillus cereus confirmed the suitability of the probe for superresolution microscopy. NanoSIMS analysis was used to monitor the uptake of the platinum(II) complex within the bacteria and demonstrate the potential of this chemical architecture to enable multimodal imaging. The successful combination of these two moieties 40 introduces a platform that could lead to a versatile range of multi-

Sub-cellular imaging of bacteria is challenging but important. Antibiotic resistance and infection prevention are pressing health concerns, and bacteria are also used for the natural synthesis of biopolymers and are potential alternative sources for oil and fuel production. Where it has been achieved, high resolution imaging of bacterial cells has contributed to great advances in prokaryotic cell biology, enabling study of the subcellular structures that play a key role in the growth and pathogenesis of these unicellular organisms.² Given the small size of bacterial cells, super-resolution microscopy is pivotal to successful visualisation of microbial specimens with sub-cellular resolution.3 Most of the fluorescence imaging on bacterial cells has to date been achieved using genetically encoded tags or commercially available dyes. Given the lack of small-molecule probes that can readily penetrate the cell envelope of live bacteria and target cellular sub-structures, considerable recent research effort has focussed on the development of new fluorophores designed to efficiently stain live bacteria.^{2,4} While most of these fluorophores are small organic molecules, 2b,3,5 we have recently reported a sequence of phosphorescent iridium complexes that selectively stain lipid vacuoles within bacteria.4d

Here, our aim was to develop probes bearing the 4-amino-1,8-naphthalimide fluorophore that enable structured illumination microscopy (SIM) imaging, while also incorporating heavy metal ions that could support additional imaging methods,⁶ or other functionality such as antibacterial properties. In this proof of concept work, the metal ion was tracked using nano-scale secondary ion mass spectrometry (nanoSIMS).⁷ NanoSIMS is particularly well-suited to analysis of isotopes and trace elements⁸ and has been combined with confocal microscopy to monitor the cellular uptake and subcellular accumulation of platinum-based drugs in human cells and tissues.⁹ NanoSIMS protocols previously applied to bacteria have generally relied on the more traditional approach based on the stable isotope labelling of target biological substrates.¹⁰ We report the synthesis and characterisation of

School of Chemistry, The University of Sydney, 2006 NSW, Australia.

E-mail: elizabeth.new@sydney.edu.au

functional probes for bacteria.

^dCentre for Microscopy, Characterisation and Analysis, Univsersity of Western Australia, 6009 Perth, WA, Australia

^eDipartimento di Chimica Industriale "Toso Montanari", Università degli Studi di Bologna, viale del Risorgimento 4, 40136 Bologna, Italy

f_{Laboratoire} CEISAM, UMR CNRS 6230, Universit8 de Nantes, 2 Rue de la HoussiniHre. BP 92208 44322 Nantes Cedex 3. France

 g Curtin Medical School, Curtin University, Kent Street, Bentley 6102 WA, Australia \dagger Electronic supplementary information (ESI) available: Synthetic details, NMR, photophysical measurements, theoretical calculations, bacterial growth and imaging details. See DOI: 10.1039/d1an00363a

^aSchool of Molecular and Life Sciences, Curtin University, Bentley 6102, WA, Australia. E-mail: m.massi@curtin.edu.au, m.ogden@curtin.edu.au ^bDipartimento di Scienze Chimiche e Geologiche, Università degli Studi di Modena e Reggio Emilia, via G. Campi 103, 41125 Modena, Italy

Communication Analyst

rhenium(I) and platinum(II) complexes bound to naphthalimide-appended ligands (Scheme 1). Similar complexes of rhenium(I) were previously investigated by Pope *et al.* in human osteoarthritic, protistan fish parasite *Spironucleus vortens* and fission yeast *Schizosaccharomyces pombe* cells using confocal fluorescence miscroscopy, ¹¹ but never assessed for live bacterial imaging. In the present study, we determined that the stability of the rhenium(I) complex was not sufficient for synthesis and application as a biological probe, bacterial imaging was studied using exclusively the platinum(II) complex. Pt-PyNapht was tested to image the live bacterial species *B. cereus* to validate the capability of this system to enable structure illumination microscopy, with the presence of the metal confirmed by nano-scale secondary ion mass spectrometry.

1

5

10

15

20

25

45

50

The synthetic routes for the formation of the complexes Pt-**PyNapht** and Re-**PyNapht** are described in Scheme 1, and further synthetic details are summarised in the ESI.† The dicyclometalated platinum(II) precursor Pt-**dmso** was synthesised from the corresponding dichloro-bridged dimer, following a previously reported method, ¹² whereas the cationic tricarbonyl diimine rhenium(I) precursor was prepared *in situ* by reaction

Scheme 1 Synthesis of the targeted complex Pt-PyNapht and Re-PyNapht. Reagents and conditions: (i) acetone/MeOH (9:1 v/v), reflux, overnight; (ii) CH_3CN , reflux, overnight.

of [Re(CO)₃(**phen**)Br] (**phen** = 1,10-phenanthroline) with AgBF₄ in acetonitrile at reflux. The *N*-3-pyridyl-4-butylamino-1,8-naphthalimide ligand (**PyNapht**) was prepared by reacting 4-nitro-naphthalic anhydride with 3-aminopyridine and butylamine (see ESI†). PyNapht was then made to react with the corresponding precursors to afford the target compounds *via* a ligand exchange reaction, with moderate yields for the platinum(II) complex (35%) and very low yield for the rhenium(I) complex, usually after successive flash chromatography to obtain a pure sample (7%). The complexes Pt-PyNapht and Re-PyNapht were characterised by NMR spectroscopy and elemental analysis, confirming the expected structures.

10

25

30

40

45

50

The photophysical properties of the complexes are summarised in Table 1. Due to stability issues, the data for Re-PvNapht are only reported in CH₂Cl₂. The absorption and emission spectra of Pt-PyNapht in CH₂Cl₂ are depicted in Fig. 1, whereas the corresponding spectra for the rhenium(1) complex are reported in the ESI.† The photophysical properties of the rhenium(1) complex are analogous to previously published similar complexes, indicating that the presence or absence of a methylene spacer between the pyridine ring and naphthalimide moiety in the ancillary ligand has very little effect. 11a The Pt-PyNapht complex exhibits a strong absorption band centred at 282 nm ($\varepsilon = 15.26 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$) followed by a band at 363 nm of lower intensity ($\varepsilon = 3.55 \times 10^4 \text{ M}^{-1} \text{ cm}^{-1}$) that are assigned to π - π * transitions of the triphenylpyridine ligand. ¹⁴ A further red-shifted band is observed at 433 nm (ε = 5.04 × 10⁴ M⁻¹ cm⁻¹), and is assigned to the lowest dipole-allowed transition in the 4-amino-1,8-naphthalimide, a transition presenting a partial charge-transfer (CT) character. 11a,15 These assignments are confirmed by ab initio calculations (see the ESI†). A very similar absorption band was observed by recording the absorption spectra of the uncoordinated PyNapht in the same solvent (see the ESI†), showing that the contribution of this lower energy band is mainly ascribed to the naphthalimide moiety. The absorption spectra of Pt-PyNapht in DMSO and H₂O followed the same trend (see the ESI†).

With increasing solvent polarity, a red-shift was observed for the lowest energy band, shifting from around 433 nm to 461 nm moving from CH_2Cl_2 to H_2O . This behaviour is ascribed to a strong solvatochromism that is typical of excited states of CT nature. 13,15b

Excitation of a CH₂Cl₂ solution of Pt-**PyNapht** at 430 nm resulted in a broad emission band centred at 515 nm, again

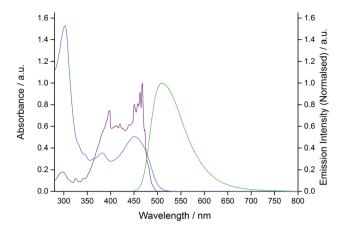
Table 1 Photophysical data of Pt-PyNapht and Re-PyNapht from dilute (10^{-5} M) solutions

Complex	$\lambda_{\rm abs}/{\rm nm} \left(10^4~arepsilon \left[{ m M}^{-1}~{ m cm}^{-1} ight] ight)$	$\lambda_{\mathrm{em}}\left[\mathrm{nm}\right]$	$ au_{ m aer}{}^a [m ns]$	$\Phi_{ m aer}^{b}$ %
Pt-PyNapht (CH ₂ Cl ₂)	282 (15.26), 363 (3.55), 433 (5.04)	515	11	28
Pt-PyNapht DMSO	287 (11.02), 348 (2.60), 448 (3.61)	542	10	5
Pt- PyNapht $(H_2O)^c$	284 (7.59), 365 (2.31), 461 (2.10)	572	3	2
Re- PyNapht (CH ₂ Cl ₂)	276 (3.17), 435 (1.05)	517	5 (57%) 10 (43%)	25

 $[^]a au_{aer}$ refers to the excited state lifetime decay obtained from air-equilibrated solution. $^b au_{aer}$ refers to the photoluminescent quatum yield obtained from air-equilibrated solution, and was measured νs . [Ru(bpy)₃]²⁺ in H₂O (φ_r = 0.028). c 0.1% DMSO was added to favour the solubilisation of the complex.

2 | Analyst, 2021, **00**, 1-5

Analyst Communication



1

10

15

25

30

35

40

45

50

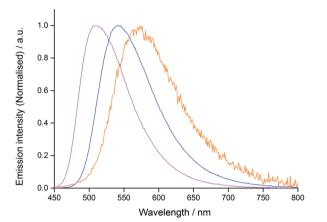


Fig. 1 Top: Absorption (blue line), normalised excitation (purple line, $\lambda_{em}=515$ nm) and normalised emission (green line, $\lambda_{exc}=430$ nm) spectra of Pt-PyNapht from a 10^{-5} M CH₂Cl₂ solution. Bottom: Normalised emission spectra (($_{\rm exc}=430$ nm) of Pt-PyNapht in CH₂Cl₂ (violet line), DMSO (blue line), H₂O (orange line).

typical of the CT emissive state of 1,8-naphthalimide fluorophores. The excitation spectrum matched the CT absorption band, suggesting that no electronic interaction existed between the platinum centre and the naphthalimide ligand. 11a,15a Moreover, the fluorescence lifetime recorded at the emission maximum exhibited a value of 11 ns, closely matching the value observed for the uncoordinated PyNapht (see the ESI†). These observations suggest that the emission observed for Pt-PyNapht is dominated by the fluorescence of the 1,8-naphthalimide, in agreement with previously reported Re(I) and Pt(II) complexes bearing related 1,8-naphthalimide ligands. 11a,15a As observed in the absorption spectra, a positive solvatochromism occurred with increasing solvent polarity, with a red-shift of 57 nm from CH₂Cl₂ to H₂O (Fig. 1). The fact that the properties of the naphthalimide are unchanged upon coordination to the rhenium(I) or platinum(II) centre is particularly advantageous, as it allows the photophysical properties of the luminescent moiety to be selected and tuned independently from the metal centre, which is acting as a "silent" carrier.

To further investigate the structural and photophysical properties of **PyNapht** and Pt-**PyNapht**, we performed time-depen-

dent density functional theory (TD-DFT) calculations, using a protocol described in the ESI.† For PyNapht, the lowest state can be ascribed to a π - π * transition, involving a CT of 0.60 e over 2.10 Å. The computed 0-0 energy is 2.77 eV, 0.19 eV off the experimental value (2.58 eV), a mismatch typical for TD-DFT (see ESI for details†).16 In Pt-PyNapht, the dihedral angle between the pyridyl ring and the 1,8-naphthalimide (PT complex) plane is 62° (54°). This leads to a near orthogonality of the two chromophores (angle of 86.2°, see the ESI†), preventing direct electronic interactions. The TD-DFT calculations reveal that the lowest singlet excited state in Pt-PvNapht is located on the 1,8-naphthalimide moiety (no lower CT-like state is found) and presents a 0-0 energy of 2.78 eV, almost identical to that determined in PyNapht. Overall, theoretical calculations therefore confirm that there is almost no interaction between the two moieties in Pt-PyNapht and that the lowest excited state is always located on the naphthalimide.

10

25

40

The use of Pt-PyNapht as a molecular probe for fluorescence microscopy in live bacteria was investigated by structured illumination microscopy (SIM). The complex was incubated with live B. cereus directly in the growth medium (Nutrient Broth, 0.1% DMSO), and the fluorescence of Pt-PyNapht was easily detected upon excitation with a 488 nm laser after only 15 min incubation (Fig. 2). To gain further spatial information about the interaction of the complex with B. cereus, bacteria were co-stained with MitoTracker Red. This commercial dye specifically stains the mitochondria in mammalian eukaryotic cells and while its use to label bacteria is quite limited, 17 we show here that MitoTracker could stain live B. cereus in the growth medium and enabled simultaneous staining with Pt-PyNapht. Notably, the super-resolution images showed that the complex exhibited sub-cellular localisation in well-defined inclusions, whose shape resembled lipid bodies. As reported previously, 4d we confirmed the lipid nature of these organelles by staining with the lipophilic dye BODIPY 493/503 (see the ESI†). The uncoordinated ligand PyNapht exhibited the same sub-cellular localisation as Pt-PyNapht (see the ESI†), in line with related 4-amino1,8-naphthalimides that

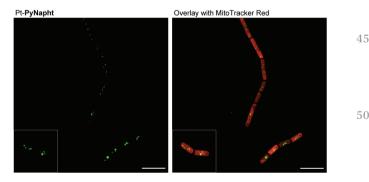


Fig. 2 SIM images of live *B. cereus* incubated with Pt-PyNapht (left) and co-stained with MitoTracker Red (right). Pt-PyNapht was excited at 488 nm and the emission collected in the 525/50 nm region. MitoTracker Red was excited at 561 nm and the emission collected in the 605/50 region. Scale bars: $5 \, \mu m$.

Analyst Communication

1

10

15

20

25

30

35

40

45

50

Fig. 3 NanoSIMS images of B. cereus incubated with Pt-PyNapht for 15 minutes. Cells were fixed, dehydrated and embedded in resin. Semithin sections were cut for NanoSIMS imaging. $^{195}\mathrm{Pt^{-}}$ image was processed with a mean filter with a radius of 3 pixels. The threshold of ¹⁹⁵Pt⁻ in the composite image is set to show the features with highest $^{195}\text{Pt}^-$ signals in B. cereus. Scale bars: 3 μm

showed preferential affinity for lipid droplets in eukaryotic cells.13 The staining experiments suggest that while the platinum(II) dicyclometalated fragment does not affect the subcellular localisation of the 4-amino-1,8-naphthalimide unit, adding the metal centre to the dye provides the additional opportunity to detect the probe with electron or isotope-based techniques not readily accessible for organic dyes. In addition, very few probes have been reported to date for use in superresolution microscopy. Hence, the detectability of Pt-PyNapht, as well as the uncoordinated PyNapht, by structured illumination microscopy opens the potential development of new classes of dyes for super-resolution microscopy.

The presence of platinum introduced using Pt-PyNapht was assessed using nanoSIMS to confirm the uptake of the platinum-based dye in the bacteria. 12C14N- secondary ions were used to show the morphology of single bacteria, and ¹⁹⁵Pt⁻ secondary ion was used to indicate the distributions of Pt-PyNapht (Fig. 3). The resulting nanoSIMS images confirmed the internalisation of Pt-PvNapht within B. cereus. The distribution of Pt-PyNapht in these cells is not homogenous, and there are some subcellular regions that yield higher ¹⁹⁵Pt⁻ signals. It should be noted that nanoSIMS images were generated from the top < 50 nm of the section whereas signals of Pt-PyNapht in structural illumination microscope images were obtained from entire bacteria. Nevertheless, the images are consistent enough with the SIM images to suggest that the platinum has been retained in the probe molecule.

This work demonstrates that the small molecule probe Pt-PyNapht efficiently stains B. cereus and enables imaging with both structured illumination microscopy (SIM), and nanoscale secondary ion mass spectrometry (nanoSIMS). The platinum(II) moiety does not change the sub-cellular localisation or photophysical properties of the 4-amino-1,8-napththalimide dye. Given the readily tailored range of known napththalimide dyes, the platform reported here can be used to develop new probes targeted to specific bacterial imaging applications.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

A. M. R. thanks Curtin University for the CIPRS Scholarship. M. V. and L. R. thank the Università degli Studi di Modena e Reggio Emilia for the More Overseas student exchange program that conferred the travel bursary to M. V. We thank the University of Sydney for funding. We acknowledge the scientific and technical assistance of the Australian Microscopy and Microanalysis Research Facility at the Australian Centre for Microscopy and Microanalysis (ACMM). This work used computational resources of the CCIPL installed in Nantes. H. J. is supported by an Australian Research Council Discovery Early Career Researcher Award. The authors thank Philip A. Gale for useful discussions.

10

15

20

25

30

40

45

Notes and references

- 1 H. W. Boucher, G. H. Talbot, J. S. Bradley, J. E. Edwards, D. Gilbert, L. B. Rice, M. Scheld, B. Spellberg and J. Bartlett, Clin. Infect. Dis., 2009, 48, 1-12.
- 2 (a) O. Kocaoglu and E. E. Carlson, Nat. Chem. Biol., 2016, 12, 472-478; (b) Z. Yao and R. Carballido-López, Annu. Rev. Microbiol., 2014, 68, 459-476.
- 3 (a) C. Coltharp and J. Xiao, Cell. Microbiol., 2012, 14, 1808-1818; (b) J. P. Schneider and M. Basler, Philos. Trans. R. Soc., B, 2016, 371, 20150499.
- 4 (a) L. Li, Y. Li, T. M. Lim and S. Q. Pan, FEMS Microbiol. Lett., 1999, 179, 141-146; (b) X. Ning, S. Lee, Z. Wang, D. Kim, B. Stubblefield, E. Gilbert and N. Murthy, Nat. Mater., 2011, 10, 602-607; (c) L. C. Ong, L. Y. Ang, S. Alonso and Y. Zhang, Biomaterials, 2014, 35, 2987-2998; (d) A. M. Ranieri, C. Caporale, V. Fiorini, P. Rigby, S. Stagni, E. Watkin, M. I. Ogden, M. J. Hackett and M. Massi, Chem. - Eur. J., 2019, 25, 10566-15070; (e) C. Reimold, H. J. Defeu Soufo, F. Dempwolff and P. L. Graumann, Mol. Biol. Cell, 2013, 24, 2340–2349; (f) P. Shieh, M. S. Siegrist, A. J. Cullen Q3 and C. R. Bertozzi, Proc. Natl. Acad. Sci. U. S. A., 2014, 111, 5456-5461; (g) P. Stiefel, S. Schmidt-Emrich, K. Maniura-Weber and Q. Ren, BMC Microbiol., 2015, 15, 1-9; (h) A. G. White, N. Fu, W. M. Leevy, J.-J. Lee, M. A. Blasco and B. D. Smith, Bioconjugate Chem., 2010, 21, 1297-1304; (i) C. Zhang, L. Yang, Y. Ding, Y. Wang, L. Lan, Q. Ma, X. Chi, P. Wei, Y. Zhao, A. Steinbchel, H. Zhang and P. Liu, Nat. Commun., 2017, 8, 15979.
- 5 K. Ritchie, Y. Lill, C. Sood, H. Lee and S. Zhang, Philos. Trans. R. Soc., B, 2013, 368, 1-8.
- 6 (a) R. T. Schirra Jr. and P. Zhang, Curr. Protoc. Cytom., 2014, **70**, 12.36.11–12.36.10; (b) E. Brown, J. Mantell, D. Carter, G. Tilly and P. Verkade, Semin. Cell Dev. Biol., 2009, 20, 910-919; (c) J. Caplan, M. Niethammer, R. M. Taylor and K. J. Czymmek, Curr. Opin. Struct. Biol., 2011, 21, 686-693; (d) K. Cortese, A. Diaspro and C. Tacchetti, *I. Histochem*. Cytochem., 2009, 57, 1103-1112; (e) S. Halary, S. Duperron and T. Boudier, Appl. Environ. Microbiol., 2011, 77, 4172-4179; (f) S. G. Li, G. Ji, Y. Shi, L. H. Klausen, T. X. Niu,

Analyst Communication

S. L. Wang, X. J. Huang, W. Ding, X. Zhang, M. D. Dong, W. Xu and F. Sun, *J. Struct. Biol.*, 2018, 201, 63–75; (g) J. M. Plitzko, A. Rigort and A. Leis, *Curr. Opin. Biotechnol.*, 2009, 20, 83–89; (h) P. Schellenberger, R. Kaufmann, C. A. Siebert, C. Hagen, H. Wodrich and K. Grnewald, *Ultramicroscopy*, 2014, 143, 41–51; (i) E. Brown and P. Verkade, *Protoplasma*, 2010, 244, 91–97.

10

15

20

25

30

35

40

45

50

- 7 (a) P. Agüi-Gonzalez, F. Opazo, K.-A. Saal, S. Jähne, S. Kabatas, N. T. N. Phan and S. O. Rizzoli, Angew. Chem., Int. Ed., 2019, 1–7; (b) C. Höschen, F. Opazo, S. O. Rizzoli, U. Diederichsen, I. C. Vreja, K. Kröhnert, S. Kabatas and S. K. Saka, Angew. Chem., Int. Ed., 2015, 54, 5784–5788; (c) S. K. Saka, A. Vogts, K. Kröhnert, F. Hillion, S. O. Rizzoli and J. T. Wessels, Nat. Commun., 2014, 5, 1–8.
- 8 (a) A. C. Komor and J. K. Barton, *Chem. Commun.*, 2013, 49, 3617–3630; (b) L. E. Wedlock, M. R. Kilburn, J. B. Cliff, L. Filgueira, M. Saunders and S. J. Berners-Price, *Metallomics*, 2011, 3, 917–925.
- (a) B. K. Keppler, M. Galanski, A. A. Legin, I. Lichtscheidl,
 A. Schintlmeister, M. Wagner and M. A. Jakupec, *Chem. Sci.*, 2014, 5, 3135–3143; (b) M. T. Proetto, C. E. Callmann,
 J. Cliff, C. J. Szymanski, D. Hu, S. B. Howell, J. E. Evans,
 G. Orr and N. C. Gianneschi, *ACS Cent. Sci.*, 2018, 4, 1477–1484.
- 10 (a) D. M. Doughty, M. Dieterle, A. L. Sessions, W. W. Fischer and D. K. Newman, *PLoS One*, 2014, **9**, 1–8; (b) N. Musat, R. Foster, T. Vagner, B. Adam and M. M. M. Kuypers, *FEMS Microbiol. Rev.*, 2012, **36**, 486–511.
- 11 (a) E. E. Langdon-Jones, C. F. Williams, A. J. Hayes, D. Lloyd, S. J. Coles, P. N. Horton, L. M. Groves and S. J. A. Pope, *Eur. J. Inorg. Chem.*, 2017, 2017, 5279–5287;
 (b) E. E. Langdon-Jones, N. O. Symonds, S. E. Yates, A. J. Hayes, D. Lloyd, R. Williams, S. J. Coles, P. N. Horton and S. J. A. Pope, *Inorg. Chem.*, 2014, 53, 3788–3797.
- 12 (a) G. W. V. Cave, N. W. Alcock and J. P. Rourke, Organometallics, 1999, 18, 1801–1803; (b) G. W. V. Cave,

- F. P. Fanizzi, R. J. Deeth, W. Errington and J. P. Rourke, *Organometallics*, 2000, **19**, 1355–1364; (*c*) A. M. Ranieri, L. K. Burt, S. Stagni, S. Zacchini, B. W. Skelton, M. I. Ogden, A. C. Bissember and M. Massi, *Organometallics*, 2019, **38**, 1108–1117.
- 13 K. G. Leslie, D. Jacquemin, E. J. New and K. A. Jolliffe, *Chem. Eur. J.*, 2018, **24**, 5569–5573.

10

2.5

30

35

40

45

50

55

- 14 (a) J. R. Berenguer, E. Lalinde and J. Torroba, *Inorg. Chem.*, 2007, 46, 9919–9930; (b) S. Fuertes, S. K. Brayshaw, P. R. Raithby, S. Schiffers and M. R. Warren, *Organometallics*, 2012, 31, 105–119; (c) M. Krause, D. Kourkoulos, D. Gonzlez-Abradelo, K. Meerholz, C. A. Strassert and A. Klein, *Eur. J. Inorg. Chem.*, 2017, 5215–5223; (d) S. C. F. Kui, S. S. Y. Chui, C. M. Che and N. Zhu, *J. Am. Chem. Soc.*, 2006, 128, 8297–8309; (e) S. C. F. Kui, F. F. Hung, S. L. Lai, M. Y. Yuen, C. C. Kwok, K. H. Low, S. S. Y. Chui and C. M. Che, *Chem. Eur. J.*, 2012, 18, 96–109; (f) W. Lu, M. W. C. Chan, K.-K. Cheung and C.-M. Che, *Organometallics*, 2001, 2477–2486
- (a) S. Banerjee, J. A. Kitchen, S. A. Bright, J. E. O'Brien, D. C. Williams, J. M. Kelly and T. Gunnlaugsson, *Chem. Commun.*, 2013, 49, 8522–8524; (b) R. M. Duke, E. B. Veale, F. M. Pfeffer, P. E. Kruger and T. Gunnlaugsson, *Chem. Soc. Rev.*, 2010, 39, 3936–3953; (c) H. Guo, M. L. Muro-Small, S. Ji, J. Zhao and F. N. Castellano, *Inorg. Chem.*, 2010, 49, 6802–6804; (d) F. Zhong and J. Zhao, *Eur. J. Inorg. Chem.*, 2017, 2017, 5196–5204.
- 16 (a) D. Jacquemin, B. Moore, C. Adamo and J. Autschbach,
 J. Chem. Theory Comput., 2014, 10, 1677–1685;
 (b) D. Jacquemin, A. Planchat, C. Adamo and B. Mennucci,
 J. Chem. Theory Comput., 2012, 8, 2359–2372.
- I. Maslov, A. Bogorodskiy, A. Mishin, I. Okhrimenko,
 I. Gushchin, S. Kalenov, N. A. Dencher, C. Fahlke, G. Bldt,
 V. Gordeliy, T. Gensch and V. Borshchevskiy, *Sci. Rep.*,
 2018, 8, 1–12.

Proof Central

Dear Author

Please use this PDF proof to check the layout of your article. If you would like any changes to be made to the layout, you can leave instructions in the online proofing interface. First, return to the online proofing interface by clicking "Edit" at the top page, then insert a Comment in the relevant location. Making your changes directly in the online proofing interface is the quickest, easiest way to correct and submit your proof.

Please note that changes made to the article in the online proofing interface will be added to the article before publication, but are not reflected in this PDF proof.