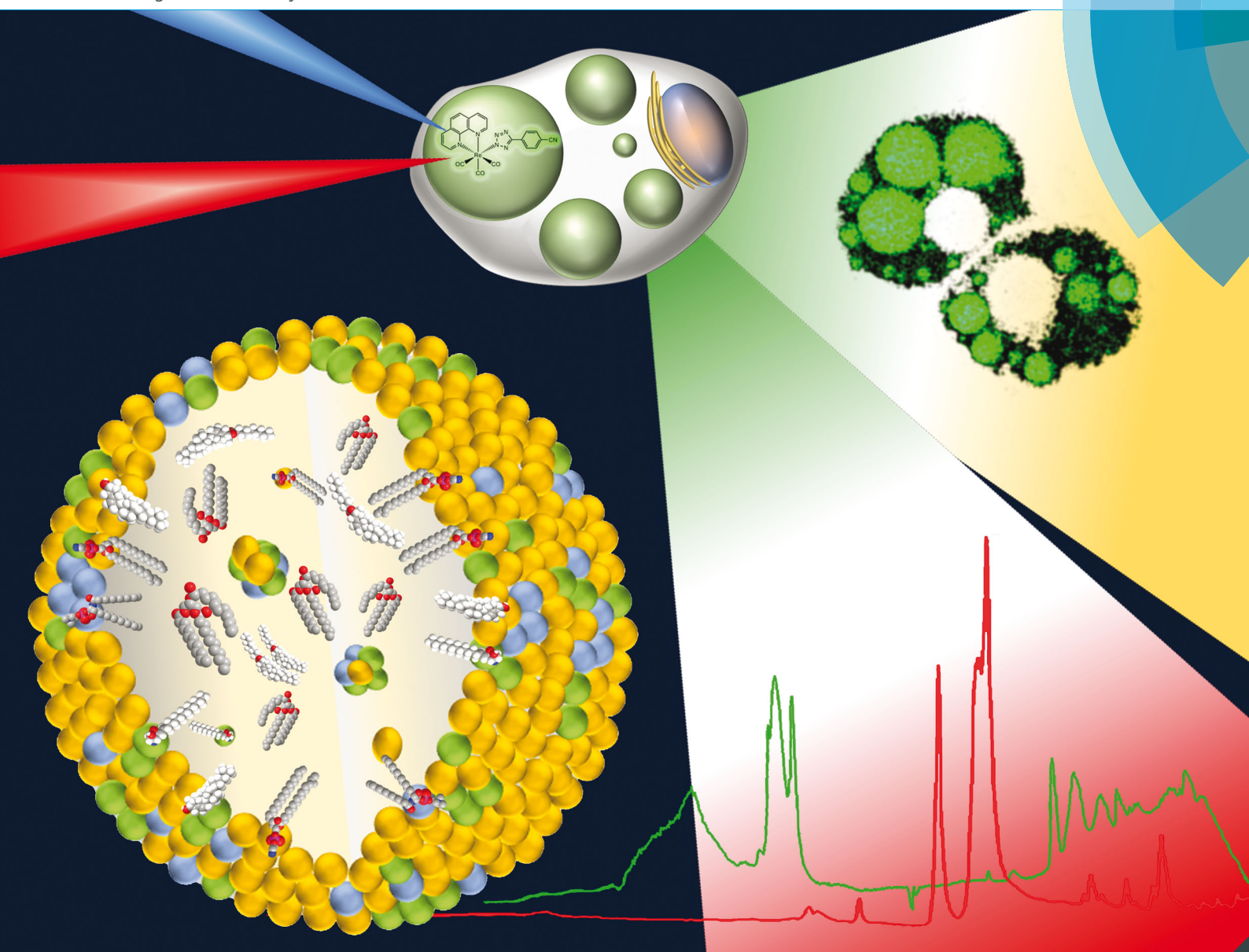


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Unprecedented staining of polar lipids by a luminescent rhenium complex revealed by FTIR microspectroscopy in adipocytes†

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Fourier transform infrared (FTIR) microspectroscopy and confocal imaging have been used to demonstrate that the neutral rhenium(I) tricarbonyl 1,10-phenanthroline complex bound to 4-cyanophenyl-tetrazolate as the ancillary ligand is able to localise in regions with high concentrations of polar lipids such as phosphatidylethanolamine (PE), sphingomyelin, sphingosine and lysophosphatidic acid (LPA) in mammalian adipocytes.

Live cell fluorescent imaging provides a valuable tool for the visualisation of cellular structure and function. One key aspect of this research area is the design of luminescent molecules, either organic fluorophores or phosphorescent metal complexes (each of which possess unique advantages and disadvantages),^{1–7} that are able to target specific organelles in live cells or specific analytes of metabolic relevance. While significant advances have been reported in biological targeting, the specific staining of lipids (excluding those belonging to membrane layers) in live cells is an undoubtedly underdeveloped area. Lipids are critical for membrane structure, to compartmentalise cellular function, to orchestrate structural events at the membrane interface and to mediate signalling, but they also have a critical role in metabolic processes.⁸ Lipid distribution, degradation, homeostasis and metabolism have direct significance for major diseases, such as inflammation, diabetes, cancer, obesity, and heart disease.⁹ The availability of a diverse family of lipid markers, especially for visualising lipid orientated process in real time,¹⁰ therefore represents an unmet need in live cell imaging.

Advances in mass spectrometry have allowed high-sensitivity and high throughput lipid analysis for the study of lipid composition.¹¹

However, the compositional analysis of lipids by electrospray mass spectrometry relies heavily on the ability to separate cellular components, which limits the biological relevance of the information obtained when using this technology. For example, the analysis of lipids found in lipid droplet organelles requires separation of the phospholipid membrane from the hydrophobic core and from other cellular components.^{12,13} Fluorescent dyes, such as Oil Red O, Nile Red, Fillipin III, BODIPY and LipidTox, are also commonly used for the imaging of lipid biology.^{14–16} However, the need for cell fixation for effective staining with Oil Red O, Nile Red, and Fillipin III results in lipid depletion.¹⁷ BODIPY has shown promise for live cell imaging of lipids. However, dual imaging can be difficult with emission potential in the red excitation channel as well as its expected green excitation.¹⁸ BODIPY and derivatives thereof also have a small Stokes-shift, which can lead to self-quenching and can be modestly cytotoxic.¹⁹ LipidTOX is amenable to live cell imaging, however, it is prone to photo-bleaching and suffers from low staining efficiency.²⁰ Furthermore, the interaction of these dyes with lipids is non-specific by nature (*e.g.*, Fillipin III has a higher specificity for gangliosides (GM1) in contrast to cholesterol;²¹ emission from Nile Red is dependent on hydrophobicity²²), with the vast majority localising with neutral lipids or phospholipids. Consequently, there is a defined need for the development of new molecular probes that exclusively enable live cell imaging of polar lipids, particularly in conjunction with other biomarkers.²³

We have recently reported the first example of a luminescent metal complex capable of staining lipid droplets in live *Drosophila* tissue and live adipocytes.²⁴ This species is composed of the neutral Re(I) tricarbonyl 4'-cyanophenyltetrazolato complex, herein designated as ReZolve-L1™ (Fig. 1A).‡ The staining of lipid droplets in live cells is significant as cellular dysfunction involving lipid droplets is increasingly being linked to major diseases.^{25,26} Lipid droplets are intracellular organelles that have important functions in membrane formation, cellular and lipid homeostasis (storage), metabolism (hydrolysis), and trafficking (signalling and transport).^{27,28}

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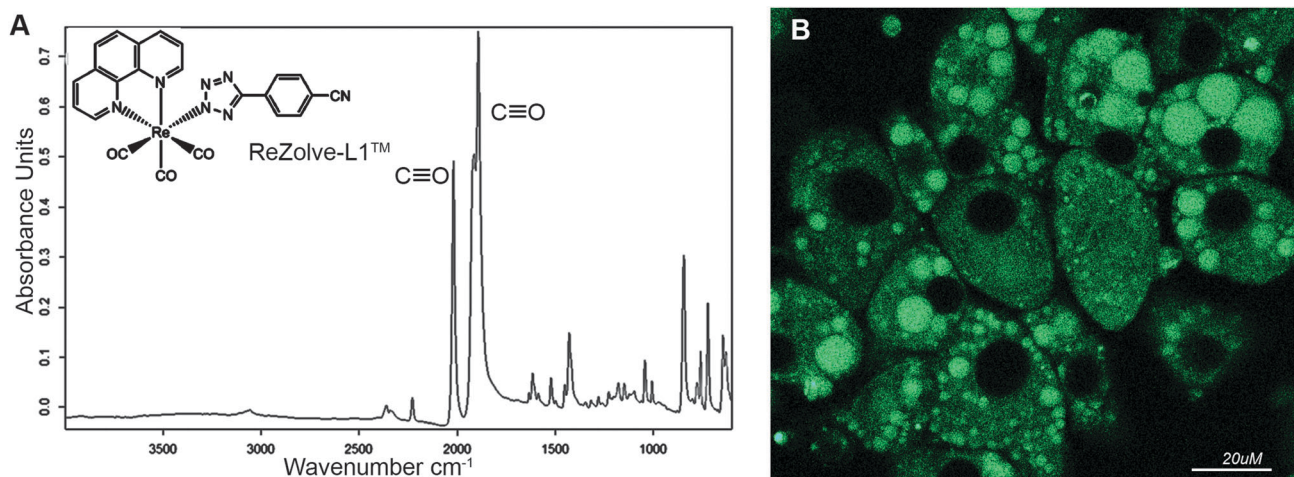


Fig. 1 (A) The structure of ReZolve-L1TM, which is a neutral *fac*-[Re(CO)₃(phen)(L)], where phen is 1,10-phenanthroline and L is 5-(4'-cyanophenyl)tetrazolate and FTIR spectrum of ReZolve-L1TM, with carbonyl ligand bands highlighted; (B) the bio-distribution of ReZolve-L1TM in live adipocytes.

The observed staining of lipid droplets by ReZolve-L1TM (Fig. 1B) suggests that this metal complex has an affinity for one of the many lipid families found within lipid droplets. It is classically accepted that the lipid droplet cores are enriched in neutral lipids, like triacylglycerides and sterol esters for energy storage and the outer limiting membrane is a lipid monolayer mainly composed of polar lipids that can include unesterified cholesterol, sphingomyelin and phosphatidylethanolamine.^{29,30} The polar lipids in the outer membrane of the lipid droplet are integrally involved in lipid droplet dynamics, intracellular signalling, and lipid compartmentalisation, as well as lipid sequestration and trafficking.^{31,32} However, the localisation of lipids, especially polar lipids, in the lipid droplet structure has yet to be fully defined.^{33–35} Therefore, it is vital that the 'selectivity' of ReZolve-L1TM for these various lipid classes be defined.

Interestingly, ReZolve-L1TM was found to also be distributed to other regions of cells (Fig. 1B). A comparison of staining patterns of ReZolve-L1TM using Fillipin III as a co-stain or Oil Red O (reported in Bader *et al.*²⁴ and reproduced as Fig. S1 in ESI†) suggests that ReZolve-L1TM may have an affinity for lipids distinct to that stained by either Fillipin III (cholesterol and GM1) or Oil Red O (neutral lipids). As there is a high demand for stains that are able to selectively discriminate polar from neutral lipids, the lipid 'specificity' of ReZolve-L1TM warranted further investigation.

Herein we demonstrate by means of IR microspectroscopy and confocal imaging^{36–38} the potential of this unique rhenium complex to localise with polar lipids in cells. Our results highlight the first example of a marker that is able to preferentially localise in areas with high polar lipid concentration, thus offering a novel biological tool for live cell imaging of lipids.

The FTIR spectrum of solid ReZolve-L1TM (Fig. 1A) is dominated by the strong C≡O stretching bands centred at ~ 2027 cm⁻¹ and the doublet $1915/1893$ cm⁻¹, characteristic of facial tricarbonyl complexes.³⁹ This is a spectral region where vibrational modes due to biochemical components are conveniently negligible.⁴⁰ There is a range of additional bands for the complex in the region between

1800 – 600 cm⁻¹ and, while these overlap with signals from cellular components, their relative intensities compared with those of the C≡O bands are such that they have not been used in the spatial analysis of the complex.

To facilitate the imaging of lipid droplets (where ReZolve-L1TM has the highest intensity staining pattern), a mammalian adipocyte 3T3-L1 cell line was chosen, as these cells have large (up to 150 μm) lipid droplets when compared to other mammalian cells (0.1 – 2 μm).⁴¹ To prepare the sample for FTIR analysis, the 3T3-L1 cells were grown on a silicon nitride substrate,³⁸ then rapidly fixed using cold methanol and air dried^{38,42} before being incubated with ReZolve-L1TM (10 μM). Importantly, direct growth of the cells on the silicon nitride substrate, followed by rapid cold MeOH fixation eliminates biochemical changes that can be induced by removing adherent cells from cell culture by trypsin, so cells grow with normal morphology.²⁷ The cells prepared in this manner also retain the lipid droplets observed under optical microscopy and enable diverse biospectroscopic studies on adipocytes with minimal introduction of sampling artifacts.^{38,43}

A bright field image of an adipocyte following incubation with ReZolve-L1TM is shown in Fig. 2A and depicts large characteristic lipid droplets. IR images generated by calculating the area under the protein amide I band (1690 – 1600 cm⁻¹), the lipid ester ν (C=O) band (1712 – 1775 cm⁻¹) and the ReZolve-L1TM C≡O band (1942 – 1910 cm⁻¹) from this cell are shown in Fig. 2B–D,

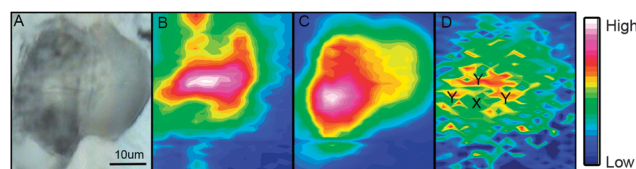


Fig. 2 (A) Bright-field image of an adipocyte incubated overnight with ReZolve-L1TM; (B) FTIR image of amide I band; (C) IR image of the lipid ester ν (C=O) band; (D) IR image of ReZolve-L1TM C≡O band. Designations Y ($n > 10$) and X ($n > 10$) indicate areas from which average spectra were taken.



respectively (see ESI,† Fig. S2 for a control cell). The FTIR image of the lipid ester, Fig. 2C, shows the highest integrated area coincides with the largest lipid droplet in the bright-field image Fig. 2A. In addition, a comparison of the highest integrated intensities from the amide I protein map against the $\nu(\text{C}=\text{O})$ lipid map showed that there was an area of high protein content adjacent to the lipid droplet location. This was assigned to the nucleus because of the high histone protein concentration in chromatin and is in close proximity to the major lipid droplet.

Lipid droplets and nuclear proteins are often found in close proximity during the nascent formation of lipid droplets as the endoplasmic reticulum is an organelle for lipid synthesis.^{34,41} ReZolve-L1™ was detected throughout the cell, which was in agreement with the localisation of lipids and confocal microscopy (Fig. 1B). Increased intensities of the bands due to ReZolve-L1™ were observed in regions of high lipid content with some overlap in the area where lipids and high protein content were in close proximity, Fig. 2D. IR spectra were then extracted from two regions within the cell and averaged (Fig. 3; see ESI,† Fig. S3 for spectra from each region with correlating 2nd derivative analysis). These regions were selected based on the ReZolve-L1™ IR map shown in Fig. 2D in combination with confocal and epifluorescence images of both fixed and live adipocytes (Fig. 1B),²⁴ which showed the highest accumulation of ReZolve-L1™ in these regions. In both extracted regions the relative absorbance from the ReZolve-L1™ $\nu(\text{C}\equiv\text{O})$ bands was relatively low in comparison with the other cellular functional groups. This is not unexpected as the cells were incubated with the same concentration of ReZolve-L1™ that is used in optical imaging experiments to ensure the same bio-distribution was obtained. Both extracted spectral averages show intense $\nu(\text{CH})$ and $\nu(\text{C}=\text{O})$ lipid bands that can be attributed to high lipid concentration, in comparison to the smaller contribution from the $\nu(\text{C}=\text{O})$ and $\nu(\text{NH})$ protein bands. This relationship suggests that the spectra were extracted from the lipid droplet (hydrophobic core and membrane) and not the nuclei or cytoplasmic regions. As expected from the integrated intensities image shown in Fig. 2D, the spectra extracted from the area surrounding the lipid droplet marked Y in Fig. 2D (spectra shown in Fig. 3), including the lipid droplet membrane, had the highest relative ratio of absorbance attributed to ReZolve-L1™ when compared to known cellular functional groups. While the spectrum extracted from the region marked X in Fig. 2D (spectra shown in Fig. 3), inside the centre of the lipid droplet, showed only a relatively small concentration of ReZolve-L1™ in comparison to cellular functional groups. This suggested that ReZolve-L1™ may be localising in higher concentration with lipids found in the lipid droplet membrane rather than the neutral lipids (*e.g.*, triglycerides).

To investigate if ReZolve-L1™ localises with specific lipids the spectra from the extracted regions in Fig. 3 were compared with spectra obtained from pure lipids. A large number of lipid structures are known to reside in cells, hence, a representative series of lipids from each of the main lipid families were chosen for analysis (see ESI,† Fig. S4 for reference IR spectra; see ESI,† Fig. S5 for peak assignments). Peaks were assigned for the extracted regions X and Y from second derivative spectra

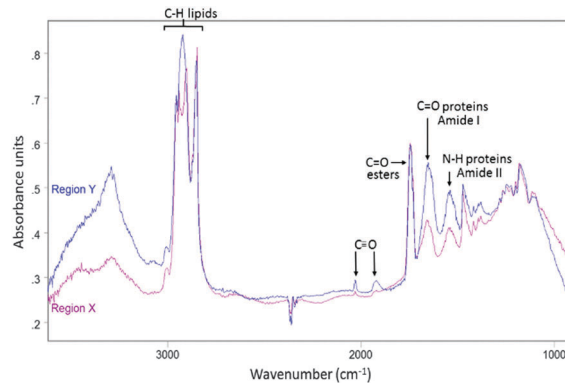


Fig. 3 Average IR spectrum collected from region Y in Fig. 2D (blue) overlaid with average IR spectrum taken from region X in Fig. 2D (purple).

which allowed for the comparison of bands in the CH stretching region ($3000\text{--}2700\text{ cm}^{-1}$) and the lipid references, see Fig. 4. Fig. 4 shows that the greatest correlation between the spectra extracted from high probe concentration (region Y) is with polar lipids. The bands at 2956 , 2921 , and 2848 cm^{-1} in the spectrum from region Y are coincident with the bands in the spectra collected from phosphatidylcholine, phosphatidylethanolamine, sphingomyelin, sphingosine and lysophosphatidic acid.

There was significantly less correlation with the non-polar lipids, such as cholesterol esters and triglycerides with the spectra from region Y. As expected the spectra extracted from region X showed high peak correlation with triglycerides and

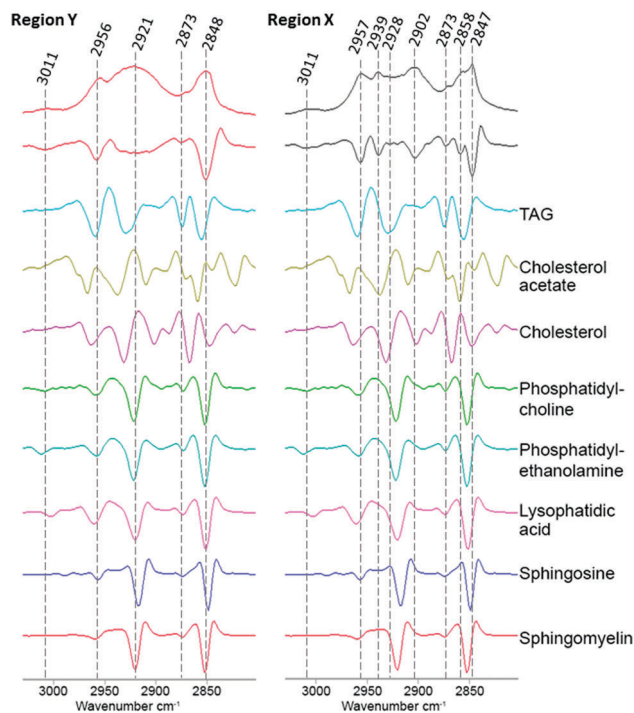


Fig. 4 Comparison of the FTIR spectra from the CH stretching region ($2750\text{--}3020\text{ cm}^{-1}$) extracted from region X and region Y (Fig. 2D) with second derivative spectra shown below followed by a range of pure lipid spectra as assigned.



cholesterol acetate (2957, 2928 and 2858 cm^{-1}). This suggests that ReZolve-L1TM mainly localised to areas directly adjacent to the hydrophobic core of the lipid droplet, corresponding to the limiting membrane. Moreover this indicates that ReZolve-L1TM is predominantly staining polar lipids and not neutral lipids in contrast to Oil Red O, Nile Red, BODIPY or LipidTOX (or derivatives thereof). Interestingly, the spectra extracted from region X did still show overlap with some of the polar lipids, albeit to a much smaller and less confident extent. It can be expected that the signal from the lipids will be more intense on the edges of the droplets as IR light is passed through many layers on the side (*i.e.* region Y) as opposed to only two layers at the top and bottom of the lipid droplet when the spectra are extracted from the middle (*i.e.* region X). Alternatively, there may be a more heterogeneous distribution of polar lipids in lipid droplets than is classically accepted. The latter correlates more with confocal microscopy Z-stack analysis which shows that while ReZolve-L1TM staining can be detected throughout the lipid droplet, it is more intense at the edges of the lipid droplet.

ReZolve-L1TM, therefore, provides new and extremely important biochemical information that has not been obtained previously with staining and fluorescence microscopies. Future work will explore how the distribution of different lipids influences the localisation of ReZolve-L1TM and which other lipids might directly interact with this molecular probe during live cell imaging.

Current commercially available imaging reagents for the live cell imaging of polar lipids are mainly based on lipid analogues conjugated to fluorescent tags, such as BODIPY or pyrene.^{44–47} Although these have assisted the investigation of lipid trafficking in cells, it is unclear if these analogue species interact with other cellular structures in the same way as their endogenous counterparts.⁴⁵ Issues with photobleaching and toxicity can also perturb time-resolved imaging using these reagents.^{19,45} Alternatively live-cell imaging of lipids can be achieved by fluorescently tagged proteins with lipid specific binding domains, which overcomes the issue of detecting endogenous lipids. However limited availability of specific proteins for this application combined with time consuming and costly transfection protocols reduces applicability.⁴⁸ To the best of our knowledge, ReZolve-L1TM is the first example of a live cell compatible molecular probe that localises to areas of high endogenous polar lipid concentration in preference to non-polar lipids. Furthermore, ReZolve-L1TM has low toxicity,²⁴ can be used in conjunction with other dyes, shows no to minimal photo-bleaching and has a high staining efficiency which are some of the drawbacks associated with other lipid dyes.^{19,20}

Conclusions

We have provided FTIR imaging evidence that ReZolve-L1TM localises to the lipid droplet region of adipocytes, which confirmed conclusions drawn from fluorescence confocal microscopy imaging. We ascertained that this molecular probe does not predominantly localise with neutral lipids, such as triglycerides, but may be more closely aligned with polar lipids such as phosphatidylcholine,

phosphatidylethanolamine, sphingomyelin, sphingosine and lysophosphatidic acid which are present in higher concentrations in lipid droplet membranes. This provided strong evidence that ReZolve-L1TM has an affinity for polar lipids and the unprecedented potential to image these lipids in live cells.

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Notes and references

‡ The authors declare that ReZolve-L1TM is the subject of a patent (PCT/AU2015/000159), which has commercial potential and has been licensed to ReZolve Scientific for commercial sales as a molecular probe. DAB, SP, MM, CAB and SS may be future shareholders in ReZolve Scientific.

- 1 K. K.-W. Lo, *Acc. Chem. Res.*, 2015, **48**, 2985–2995.
- 2 M. P. Coogan and V. Fernandez-Moreira, *Chem. Commun.*, 2014, **50**, 384–399.
- 3 H. Kobayashi, M. R. Longmire, M. Ogawa and P. L. Choyke, *Chem. Soc. Rev.*, 2011, **40**, 4626–4648.
- 4 E. J. New, D. Parker, D. G. Smith and J. W. Walton, *Curr. Opin. Chem. Biol.*, 2010, **14**, 238–246.
- 5 M. R. Gill and J. A. Thomas, *Chem. Soc. Rev.*, 2012, **41**, 3179–3192.
- 6 J. A. Thomas, *Chem. Soc. Rev.*, 2015, **44**, 4494–4500.
- 7 M. Mauro, A. Aliprandi, D. Septiadi, N. S. Kehr and L. De Cola, *Chem. Soc. Rev.*, 2014, **43**, 4144–4166.
- 8 S. Takatori, R. Mesman and T. Fujimoto, *Biochemistry*, 2014, **53**, 639–653.
- 9 K. Reue, *J. Lipid Res.*, 2011, **52**, 1865–1868.
- 10 G. van Meer, *EMBO J.*, 2005, **24**, 3159–3165.
- 11 L. Li, J. Han, Z. Wang, J. A. Liu, J. Wei, S. Xiong and Z. Zhao, *Int. J. Mol. Sci.*, 2014, **15**, 10492.
- 12 A. Reis, A. Rudnitskaya, G. J. Blackburn, N. M. Fauzi, A. R. Pitt and C. M. Spickett, *J. Lipid Res.*, 2013, **54**, 1812–1824.
- 13 A. R. Thiam, R. V. Farese, Jr. and T. C. Walther, *Nat. Rev. Mol. Cell Biol.*, 2013, **14**, 775–786.
- 14 R. N. Melo, H. D'Ávila, P. Bozza and P. Weller, in *Light Microscopy*, ed. H. Chiarini-Garcia and R. C. N. Melo, Humana Press, 2011, vol. 689, ch. 9, pp. 149–161.
- 15 P. Greenspan, E. P. Mayer and S. D. Fowler, *J. Cell Biol.*, 1985, **100**, 965–973.
- 16 R. Koopman, G. Schaart and M. K. Hesselink, *Histochem. Cell Biol.*, 2001, **116**, 63–68.
- 17 S. Fukumoto and T. Fujimoto, *Histochem. Cell Biol.*, 2002, **118**, 423–428.
- 18 Y. Ohsaki, Y. Shinohara, M. Suzuki and T. Fujimoto, *Histochem. Cell Biol.*, 2010, **133**, 477–480.



- 19 R. D. Moriarty, A. Martin, K. Adamson, E. O'Reilly, P. Mollard, R. J. Forster and T. E. Keyes, *J. Microsc.*, 2014, **253**, 204–218.
- 20 E. J. O'Rourke, A. A. Soukas, C. E. Carr and G. Ruvkun, *Cell Metab.*, 2009, **10**, 430–435.
- 21 J. R. Arthur, K. A. Heinecke and T. N. Seyfried, *J. Lipid Res.*, 2011, **52**, 1345–1351.
- 22 D. L. Sackett and J. Wolff, *Anal. Biochem.*, 1987, **167**, 228–234.
- 23 J. Spandl, D. J. White, J. Peychl and C. Thiele, *Traffic*, 2009, **10**, 1579–1584.
- 24 C. A. Bader, R. D. Brooks, Y. S. Ng, A. Sorvina, M. V. Werrett, P. J. Wright, A. G. Anwer, D. A. Brooks, S. Stagni, S. Muzzioli, M. Silberstein, B. W. Skelton, E. M. Goldys, S. E. Plush, T. Shandala and M. Massi, *RSC Adv.*, 2014, **4**, 16345–16351.
- 25 N. Kraemer, R. V. Farese, Jr. and T. C. Walther, *EMBO Mol. Med.*, 2013, **5**, 905–915.
- 26 D. A. Gross and D. L. Silver, *Crit. Rev. Biochem. Mol. Biol.*, 2014, **49**, 304–326.
- 27 S. D. Kohlwein, M. Veenhuis and I. J. van der Klei, *Genetics*, 2013, **193**, 1–50.
- 28 M. Beller, K. Thiel, P. J. Thul and H. Jäckle, *FEBS Lett.*, 2010, **584**, 2176–2182.
- 29 H. Abramczyk, J. Surmacki, M. Kopec, A. K. Olejnik, K. Lubecka-Pietruszewska and K. Fabianowska-Majewska, *Analyst*, 2015, **140**, 2224–2235.
- 30 D. J. Murphy, *Prog. Lipid Res.*, 2001, **40**, 325–438.
- 31 I. Dugail and E. Hajdouch, *Cell. Mol. Life Sci.*, 2007, **64**, 2452–2458.
- 32 Z. Sun, J. Gong, L. Wu and P. Li, in *Methods in Cell Biology*, ed. Y. Hongyuan and L. Peng, Academic Press, 2013, vol. 116, pp. 253–268.
- 33 E. Muro, G. E. Atilla-Gokcumen and U. S. Eggert, *Mol. Biol. Cell*, 2014, **25**, 1819–1823.
- 34 A. Pol, S. P. Gross and R. G. Parton, *J. Cell Biol.*, 2014, **204**, 635–646.
- 35 F. Wilfling, J. T. Haas, T. C. Walther and R. V. F. Jr, *Curr. Opin. Cell Biol.*, 2014, **29**, 39–45.
- 36 E. A. Carter, K. K. Tam, R. S. Armstrong and P. A. Lay, *Biophys. Rev.*, 2009, **1**, 95–103.
- 37 M. Diem, M. Romeo, S. Boydston-White, M. Miljkovic and C. Matthaus, *Analyst*, 2004, **129**, 880–885.
- 38 E. A. Carter, B. S. Rayner, A. I. McLeod, L. E. Wu, C. P. Marshall, A. Levina, J. B. Aitken, P. K. Witting, B. Lai, Z. Cai, S. Vogt, Y.-C. Lee, C.-I. Chen, M. J. Tobin, H. H. Harris and P. A. Lay, *Mol. BioSyst.*, 2010, **6**, 1316–1322.
- 39 M. V. Werrett, D. Chartrand, J. D. Gale, G. S. Hanan, J. G. MacLellan, M. Massi, S. Muzzioli, P. Raiteri, B. W. Skelton, M. Silberstein and S. Stagni, *Inorg. Chem.*, 2011, **50**, 1229–1241.
- 40 K. Meister, J. Niesel, U. Schatzschneider, N. Metzler-Nolte, D. A. Schmidt and M. Havenith, *Angew. Chem., Int. Ed.*, 2010, **49**, 3310–3312.
- 41 M. Digel, R. Ehehalt and J. Füllekrug, *FEBS Lett.*, 2010, **584**, 2168–2175.
- 42 V. Zohdi, D. R. Whelan, B. R. Wood, J. T. Pearson, K. R. Bamberg and M. J. Black, *PLoS One*, 2015, **10**, e0116491.
- 43 L. E. Wu, A. Levina, H. H. Harris, Z. Cai, B. Lai, S. Vogt, D. E. James and P. A. Lay, *Angew. Chem., Int. Ed.*, 2016, **55**, 1742–1745.
- 44 K. Tanhuanpää, J. Virtanen and P. Somerharju, *Biochim. Biophys. Acta, Mol. Cell Res.*, 2000, **1497**, 308–320.
- 45 P. Somerharju, *Chem. Phys. Lipids*, 2002, **116**, 57–74.
- 46 M. Holtta-Vuori, R. L. Uronen, J. Repakova, E. Salonen, I. Vattulainen, P. Panula, Z. Li, R. Bittman and E. Ikonen, *Traffic*, 2008, **9**, 1839–1849.
- 47 D. L. Marks, R. Bittman and R. E. Pagano, *Histochem. Cell Biol.*, 2008, **130**, 819–832.
- 48 Y. Mély and G. Duportail, *Fluorescent Methods to Study Biological Membranes*, Springer, Berlin, Heidelberg, 2012.

