

Cite this: *Chem. Sci.*, 2015, 6, 6456

Real-time detection of histone deacetylase activity with a small molecule fluorescent and spectrophotometric probe†

Debra R. Rooker and Daniela Buccella*

Histone deacetylases (HDACs) are central players in transcription regulation and important targets in cancer treatment. Activity assays are critical tools for the study of the function and regulation of these enzymes, as well as for the screening of potential inhibitors. We report a small-molecule probe for single-step, continuous detection of deacetylase activity based on an acetyl-lysine mimic functionalized with an amine-reactive fluorophore, designed to undergo rapid intramolecular imine formation upon deacetylation. The probe exhibits a bathochromic shift in the absorption spectrum and changes in fluorescence emission intensity that enable unprecedented real-time detection of HDAC activity of purified enzymes or in cell lysates, and offers a means to evaluate HDAC inhibitors *via* simple spectrophotometric or fluorescence readings without the need of additional reagents.

Received 24th July 2015
Accepted 30th July 2015

DOI: 10.1039/c5sc02704g

www.rsc.org/chemicalscience

Introduction

Post-translational modifications are often exploited by nature to regulate a variety of protein functions in biological systems. Acetylation of the ϵ -amino group of lysine residues in the histone proteins induces changes in chromatin structure that are predominantly associated with activation of DNA transcription and replication,¹ and is one of the most prominent epigenetic mechanisms influencing human carcinogenesis.² Acetylation levels are maintained by the opposing action of two families of enzymes, the histone acetyltransferases (HAT, responsible for lysine acetylation) and the histone deacetylases (HDAC, responsible for acetyl removal). Up-regulation of HDACs has been associated with the silencing of tumor-suppressing genes, which are responsible for controlling cell proliferation, differentiation and apoptosis.³ Consequently, HDAC inhibition has emerged as a promising strategy for cancer treatment.

Activity assays are a critical tool in the study of the mechanism, biological function and regulation of enzymatic deacetylation as well as in the evaluation of HDAC-targeting inhibitors with potential therapeutic value. In early studies, deacetylase activity was estimated through the detection of [³H]acetate product from the hydrolysis of isotopically-labeled acetylated histones⁴ or by immunoblotting.⁵ Only a limited number of

non-isotopic activity assays, primarily based on the use of fluorescent substrates, have been developed to date.^{6–10} For example, a method developed by Jung and coworkers uses an acetyl-lysine functionalized with a fluorescent coumarin to quantitate deacetylation after extraction and chromatographic analysis of the product mixture.^{6,11} Alternative fluorogenic assays developed in the last decade exploit fluorophore-decorated acetylated peptides that, after deacetylation, are subjected to protease degradation leading to fluorophore release and fluorescence activation.^{7,12,13} The latter approach constitutes the basis of various commercially available activity assay kits. Nevertheless, the multistep nature of these assays, which require a separation or ‘fluorescence-developing’ step prior to sample analysis, limits the use of the probes to the assessment of activity *in vitro* in a discontinuous fashion.

More recently, a series of fluorogenic substrates amenable for single-step detection of deacetylase activity were reported by Kikuchi and coworkers.^{9,14} The peptide substrates undergo spontaneous intramolecular transesterification upon deacetylation of a lysine residue, giving rise to a fluorescence signal. The key transesterification step, however, is slow and dominates the kinetics of the fluorescence response. Under these conditions, the prospects of real-time observation of deacetylation are still limited. In this context, we report herein the design of a new small-molecule probe that serves as both HDAC substrate and optical reporter, offering a rapid response that enables the detection of enzymatic deacetylation in real time *via* spectrophotometric or fluorometric measurements. With this new probe, we demonstrate continuous detection of deacetylase activity of purified enzymes and in cell lysates, without additional reagents, which offers a convenient method for evaluation of HDAC inhibitors *via* simple optical readings.

Department of Chemistry, New York University, New York, New York 10003, USA.
E-mail: dbuccella@nyu.edu

† Electronic supplementary information (ESI) available: Experimental details, additional absorption and fluorescence spectra, kinetic and thermodynamic parameters of intramolecular imine formation and characterization data for all compounds. See DOI: 10.1039/c5sc02704g

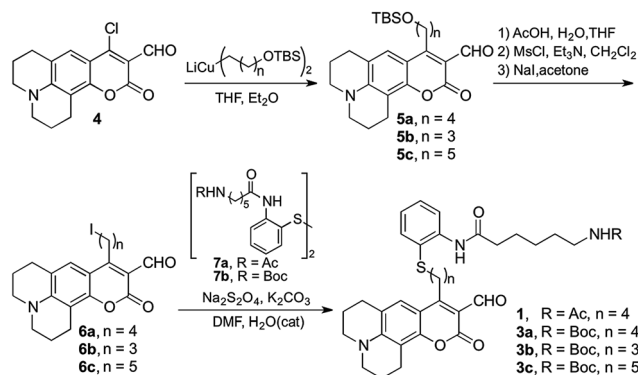


Results and discussion

We envisioned a small-molecule deacetylase activity probe consisting of an acetyl-lysine mimic functionalized with an amine-reactive coumarin fluorophore, designed to undergo rapid intramolecular imine formation that ensued optical changes upon enzymatic deacetylation (Fig. 1). Based on crystal structures of inhibitor-bound HDACs^{15,16} and insight from design of HDAC inhibitors,¹⁷ we chose to introduce an aromatic moiety as part of the linker between the reporter and reactive moieties of the probe, to aid its interaction with residues surrounding the catalytic pocket of the enzymes.

Probe **1** was prepared following the protocol depicted in Scheme 1. Amine-reactive 4-chloro-3-formylcoumarin **4** was treated with an appropriate alkyl cuprate yielding *tert*-butyldimethylsilyl (TBS) ether **5**. Following deprotection under acidic conditions, the alcohol was mesylated and converted to alkyl iodide **6**.¹⁸ Final probe **1** was obtained *via* reductive alkylation¹⁹ of aryl disulfide **7a** bearing an amide of 6-acetamidohexanoic as an acetyl lysine mimic. A similar synthetic route was employed to prepare Boc-protected analogue **3a**, which provides convenient access to the product of deacetylation (compound **2a**, Fig. 1) *via* chemical deprotection.

With probe in hand, we focused our initial studies on the intramolecular imine formation step. Boc-protected compound **3a** was deprotected quantitatively with trifluoroacetic acid and the product amine was diluted and allowed to equilibrate in aqueous buffer at pH 8.0 and 37 °C. A bathochromic shift from 471 nm to 499 nm was observed in the absorption spectrum upon equilibration, consistent with the formation of an imine product (Fig. 2A and Table 1 and S1, ESI[†]). In this regard, Glass



Scheme 1 Synthetic route to HDAC activity probe and acid-cleavable Boc-protected analogues.

and coworkers have reported similar spectroscopic changes induced by the intermolecular formation of imines between 7-diethylamino-3-formyl-coumarins and amino acids^{20,21} or catecholamine neurotransmitters.^{18,22} On the basis of NMR and optical spectroscopic studies, it has been suggested that the *s-trans* conformation of the imine favors its protonation and intramolecular hydrogen bonding with the coumarin carbonyl,²⁰ leading to a red shift in the absorption maximum.

In addition to the observed changes in absorption, compound **2a** exhibited changes in fluorescence emission intensity upon imine formation (Fig. S2[†]). An increase in emission was observed upon excitation at 499 nm, whereas a modest decrease resulted upon excitation at 477 nm (isobestic point). Measurements of the quantum efficiency of the equilibrated sample indicate that the mixture of amine and imine

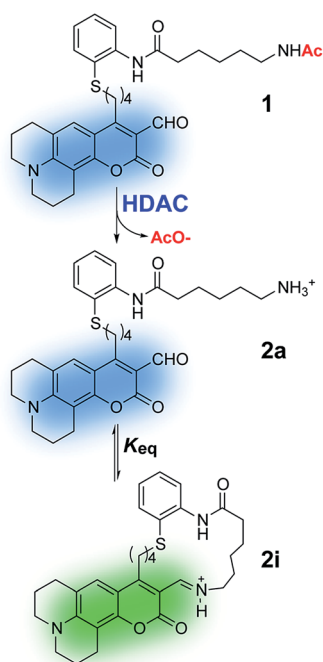


Fig. 1 Probe for single-step spectrophotometric and fluorescence-based detection of HDAC activity.

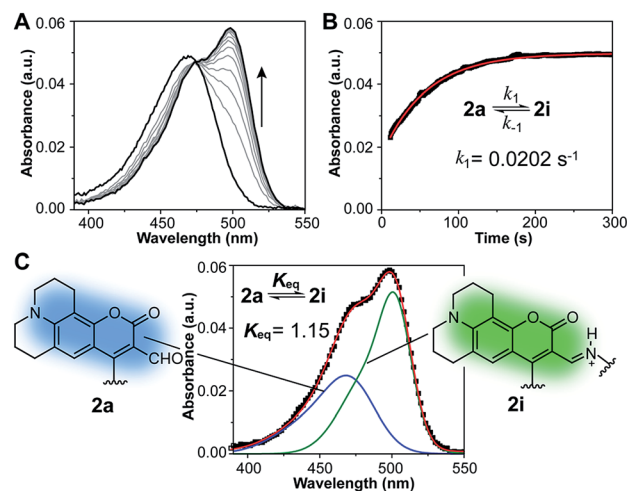


Fig. 2 Spectroscopic changes observed upon intramolecular imine formation from **2a** in aqueous buffer at pH 8.0 and 37 °C. (A) Absorption spectra during equilibration of a ~1.5 μM solution of amine **2a**. (B) Absorbance at 499 nm over time (black squares) fitted to an integrated rate law for a reversible first order reaction (approach to equilibrium, red line). (C) Deconvolution of the absorption spectrum (black squares) of equilibrated sample showing individual contributions of the aldehyde (blue line) and imine (green line) to the overall absorption.



Table 1 Photophysical properties of HDAC activity probe **1** and deacetylated product^{ab}

Compound	$\epsilon_{471} (\times 10^3 \text{ M}^{-1} \text{ cm}^{-1})$	$\epsilon_{499} (\times 10^3 \text{ M}^{-1} \text{ cm}^{-1})$	Φ_{499}^c
Probe 1	23.4 (7)	6.0 (2)	0.53 (4)
Equilibrated mixture of 2a and 2i	22.6 (9)	28 (1)	0.21 (3)

^a Values determined from measurements in 50 mM HEPES, 100 mM KCl, pH 8.0 at 37 °C. Errors in the last significant figure indicated in parenthesis. ^b Deacetylated product obtained from acid deprotection of compound **3a**. ^c Fluorescein in 0.1 N NaOH ($\Phi_{492} = 0.95$) used as standard.

products **2a** and **2i** is overall less emissive than the parent acetylated probe (Table 1). This difference could be explained by a possible quenching effect of the free amine to the coumarin fluorophore in **2a**, leading to its lower contribution to the emission.

The absorption spectra were employed to estimate the equilibrium constant for the intramolecular imine formation leading to macrocycle **2i**. The spectrum of an equilibrated mixture of **2a** and **2i** was deconvoluted as a sum of Gaussian functions (Fig. 2C and S3†) representing the individual contributions of the aldehyde and imine to the overall absorption. In this process, the spectrum of aldehyde **2a** was assumed to be identical to that of the acetylated probe **1** measured independently. Derivatives with linkers of various lengths between the fluorophore and the lysine mimic, *i.e.* compounds **3b** and **3c**, were prepared and evaluated in order to assess the optimal length of the tether for intramolecular imine formation (Fig. S1†). The equilibrium mixture of the deacetylated product **2a**, with a four-carbon alkyl linker, showed the largest fraction of imine in solution ($K_{\text{eq}} = 1.15 \pm 0.04$) and the largest optical change compared to analogues with shorter and longer linkers ($K_{\text{eq}} = 0.77 \pm 0.07$ and 0.51 ± 0.02 for the equilibrium mixture obtained after deprotection of **3b** and **3c**, respectively).

Kinetics of the reversible imine formation step were studied *via* pH jump experiments, starting from pure amine samples obtained by quantitative cleavage of the Boc-protected compounds **3a–c**. Forward and reverse rate constants (Table S2†) were determined from a nonlinear fit of the absorption at 499 nm as a function of time (Fig. 2B and S4†). In all cases, equilibrium was reached within a few minutes, with forward rate constants, k_1 , ranging between 0.0071 and 0.0202 s^{-1} for the three derivatives. Among these, the species with a four-carbon alkyl linker between fluorophore and lysine mimic displayed the fastest equilibration. The advantageous kinetics combined with favorable thermodynamics for imine formation in this model compound led us to choose acetylated probe **1** as our best candidate for the development of a deacetylation assay.

The marked optical changes and rapid equilibration of the probe after amine deprotection suggested that our design would enable continuous, real-time detection of enzymatic deacetylation by spectroscopic methods without the need for chemical extraction of the product or the addition of developing reagents typically used in commercial assays. To test this notion, compound **1** was used as a substrate for human recombinant HDAC3 in a 1 : 1 stoichiometry complex with the deacetylase activation domain of NCOR1.²³ An aliquot of

enzyme complex was diluted in HEPES buffer at pH 8.0 and 37 °C, and treated with 12 μM of the probe. Product formation was observed over time (Fig. 3A) as revealed by the appearance of the absorption band centered at 499 nm corresponding to the coumarin imine. A negative control was conducted by treating the enzyme with 1.2 μM of a reversible HDAC inhibitor, trichostatin A (TSA), prior to addition of the probe. For this fully inhibited sample, no significant changes were observed in the absorption spectrum over time. A similar sample treated with chemically deacetylated probe **2a** was employed as positive control. Absorption values at 499 nm obtained from negative and positive control experiments were employed to estimate the signal corresponding to zero and 100% conversion levels, respectively, and to calibrate the optical response of the probe over time (Fig. 3B). A 74% probe conversion was achieved over 100 min,²⁴ with an initial velocity of $22 \text{ U } \mu\text{g}^{-1}$ of protein.

The applicability of probe **1** as a deacetylase activity probe was further demonstrated using fluorescence emission measurements. Over the course of the reaction catalyzed by HDAC3/NCOR1, deacetylated product formation was detected by the change in emission intensity at 523 nm upon excitation at 499 nm (Fig. 3C). Positive and negative control experiments

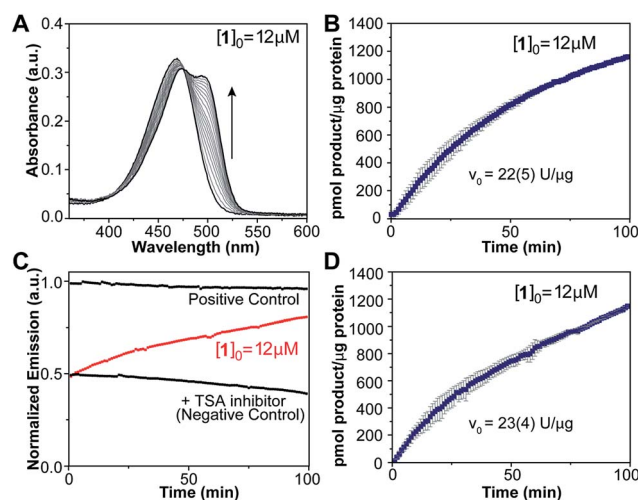


Fig. 3 Deacetylation of probe **1** in a sample of HDAC3/NCOR1 complex (100 nM enzyme) at pH 8.0 and 37 °C. (A) Absorption spectrum over time. (B) Product formation over time, calculated from changes in absorption at 499 nm. (C) Change in fluorescence emission ($\lambda_{\text{Ex}} = 499 \text{ nm}$, $\lambda_{\text{Em}} = 523 \text{ nm}$) for probe **1** (red) and corresponding negative and positive controls (black). (D) Product formation over time calculated from changes in emission intensity. Reported activity values were estimated from the slope of the curve in the first 10 minutes.



were conducted in the same fashion as for the absorption-based assays, and product formation over time was calibrated based on the signal obtained from these experiments. A slow decline in emission intensity was observed for both controls over time, presumably as a result of some photobleaching caused by prolonged exposure to the excitation light source. The reaction progress curve obtained from the fluorescence experiment (Fig. 3D) resembles those obtained from the absorption spectroscopy experiments, revealing 73% conversion in 100 min and similar initial velocity. Overall, these results indicate that changes in optical properties of our probe enable the real-time detection of enzyme activity, with good Z' values of 0.88 for absorption and 0.82 for fluorescence detection methods (Table S3†).²⁵

To explore the isoform specificity of the probe, we conducted spectrophotometric measurements in the presence of representative examples of other classes of deacetylases, namely HDAC6 (Class IIa) and Sirt1 (Class III) in addition to HDAC3 (Class I). The probe was rapidly converted in the presence of HDAC6, but no significant conversion was detected with up to 100 nM Sirt1 and 0.3 mM NAD⁺ under the typical conditions of our experiments (Fig. S5 and S6†).

An important application of activity assays is the evaluation of inhibitors as potential drug candidates for HDAC-targeting therapeutics. To evaluate the performance of our probe, we constructed dose response curves for the inhibition of HDAC3/NCOR1 by valproic acid, a drug used in the treatment of seizures and bipolar disorders, which has been shown to target histone deacetylases²⁶ (Fig. 4). Samples of enzyme equilibrated with the desired concentrations of the drug were incubated with probe 1 and deactivated by heat shock at 95 °C after 3.5 hours. An EC₅₀ value in the sub-millimolar range was estimated from either absorption or fluorescence measurements, which is within the range of IC₅₀ values reported for HDAC3.^{27,28}

Finally, we sought to gauge the performance of our probe for the detection of deacetylase activity in cell lysates, in which the HDAC complexes are largely intact and should display closer to native activity. An aliquot of HeLa nuclear extract was diluted in HEPES buffer at pH 8.0 and 37 °C, and treated with the probe for a total concentration of 4 μM. Product formation was monitored over the course of 4 hours by measuring the increase in absorption at 510 nm (Fig. 5A),²⁹ or fluorescence emission at

523 nm upon excitation at 499 nm (Fig. 5C). Positive and negative controls were conducted in the same fashion as those described for the experiments with purified proteins.

For the fully HDAC-inhibited sample treated with TSA, a rapid increase in absorption was observed over the first 5 min of the experiment, followed by a slow, small increase over the course of hours. We attributed these changes to nonspecific binding of the probe to nucleophiles present in the cell extract, as confirmed by the absence of deacetylated product in the HPLC analysis of the sample. For comparison, an association constant $K < 4 \text{ M}^{-1}$ was determined for the intermolecular reaction of the probe with 6-aminohexanoic acid in aqueous buffer at pH 8.0 and 37 °C. In this regard, Hooker and coworkers have exploited the intermolecular reaction of an enzymatically-released aldehyde moiety with endogenous cellular nucleophiles as a means to increase the intracellular retention of a fluorescent HDAC probe.³⁰

Subtraction of the signal caused by background reactions, as determined by the negative control, revealed the changes in absorption associated with the formation of the deacetylated product in our assay (Fig. 5B). Using the positive and negative controls to calibrate the optical signal, we estimated 5–10%

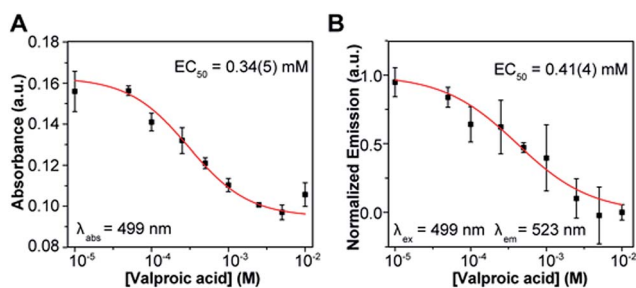


Fig. 4 Dose response curve for inhibition of 37 nM HDAC3/NCOR1 activity by valproic acid. (A) Curve constructed from absorption readings at 499 nm. (B) Data obtained from fluorescence readings ($\lambda_{\text{ex}} = 499 \text{ nm}$, $\lambda_{\text{em}} = 523 \text{ nm}$).

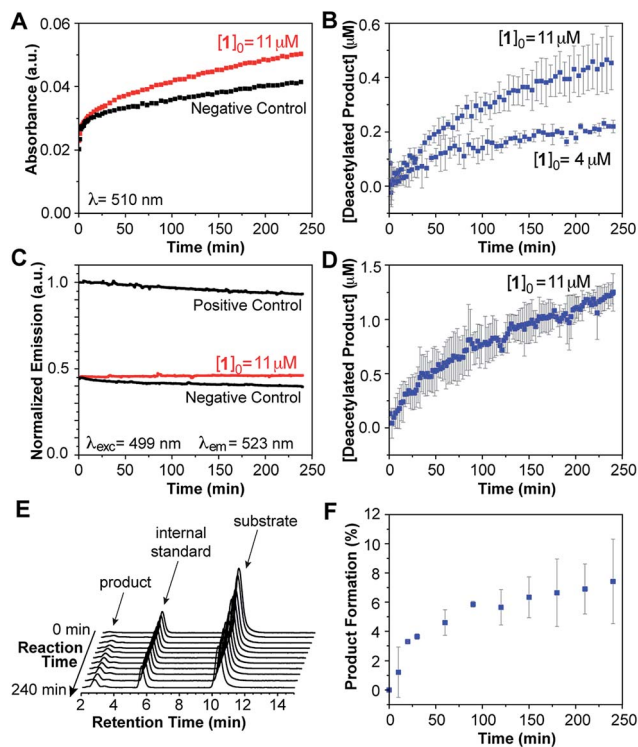


Fig. 5 Deacetylation of probe 1 in diluted HeLa nuclear extract at pH 8.0 and 37 °C. (A) Change in absorption at 510 nm over time for probe 1 (red) and negative control (black). (B) Concentration of deacetylated product over time calculated from changes in absorption. (C) Change in fluorescence emission ($\lambda_{\text{ex}} = 499 \text{ nm}$, $\lambda_{\text{em}} = 523 \text{ nm}$) for probe 1 (red) and corresponding negative and positive controls (black). (D) Concentration of deacetylated product over time calculated from changes in emission intensity. (E) HPLC chromatograms of a reaction mixture showing the appearance of deacetylated product (standard = coumarin 6H). (F) Product formation over time estimated by HPLC.



product formation over 4 hours.³¹ Higher probe concentrations led to proportionally faster deacetylation, and similar conversion levels at the end of the run. However, limits in the solubility of the probe beyond ~14 μM precluded studies at much higher concentrations. Future adjustments in the molecular design will be aimed at fine-tuning the properties of the probe to provide a wider working range of concentrations for kinetic studies.

Lastly, HPLC was employed to monitor the progress of the deacetylation of probe **1** in a sample of diluted HeLa extract (pH 8.0 and 37 °C), treated with coumarin 6H as an internal standard (Fig. 5E). Aliquots were taken at various time points over the course of 4 hours and mixed with equal volumes of acetonitrile to precipitate the protein component. Product concentration estimated from chromatographic analysis of the supernatant at various time points (Fig. 5F) matched the values calculated from the spectroscopic studies, thus indicating that the latter indeed reveal, in real time, the progress of the reaction through a faster and much simpler analysis.

Conclusions

In summary, we have developed a small-molecule probe that enables single-step detection of HDAC activity from simple spectrophotometric and fluorescence measurements, based on a new application of an intramolecular imine formation process with an amine-reactive coumarin fluorophore. We have illustrated the utility of the probe with the first examples of continuous, real-time measurement of deacetylation activity in samples of purified enzymes and in cell lysates. The convenience of the optical readouts and good Z' values make our probe a valuable tool for evaluation of HDAC inhibitors with potential therapeutic application. The wavelength shift observed in the absorption spectrum upon imine formation is an attractive feature, as it may enable ratiometric determination of substrate conversion. Finally, with a rapid and continuous response, and no need for additional reagents or extraction steps, we envision that our small-molecule probe may serve as a starting point in the development of new methods for the real time measurement of deacetylase activity in cells, a valuable tool in the epigenetics research arena.

Acknowledgements

This work was supported by an NYU Whitehead Fellowship for Junior Faculty in Biomedical and Biological Sciences to D.B. and a Margaret Strauss Kramer Fellowship to D.R. The Bruker Avance 400 and 500 NMR spectrometers were acquired with support of NSF under award number CHE-01162222. The authors thank Harriet Lloyd and Boris Ryabtsev for assistance during initial studies.

Notes and references

- 1 T. Kouzarides, *Cell*, 2007, **128**, 693–705.
- 2 C. B. Yoo and P. A. Jones, *Nat. Rev. Drug Discovery*, 2006, **5**, 37–50.
- 3 M. A. Gluzak and E. Seto, *Oncogene*, 2007, **26**, 5420–5432.
- 4 D. Kölle, G. Brosch, T. Lechner, A. Lusser and P. Loidl, *Methods*, 1998, **15**, 323–331.
- 5 Y. Zhang, G. LeRoy, H.-P. Seelig, W. S. Lane and D. Reinberg, *Cell*, 1998, **95**, 279–289.
- 6 K. Hoffmann, G. Brosch, P. Loidl and M. Jung, *Nucleic Acids Res.*, 1999, **27**, 2057–2058.
- 7 D. Wegener, F. Wirsching, D. Riester and A. Schwienhorst, *Chem. Biol.*, 2003, **10**, 61–68.
- 8 K. Dhara, Y. Hori, R. Baba and K. Kikuchi, *Chem. Commun.*, 2012, **48**, 11534–11536.
- 9 R. Baba, Y. Hori, S. Mizukami and K. Kikuchi, *J. Am. Chem. Soc.*, 2012, **134**, 14310–14313.
- 10 M. Minoshima, T. Matsumoto and K. Kikuchi, *Anal. Chem.*, 2014, **86**, 7925–7930.
- 11 K. Hoffmann, B. Heltweg and M. Jung, *Arch. Pharm.*, 2001, **334**, 248–252.
- 12 D. Wegener, C. Hildmann, D. Riester and A. Schwienhorst, *Anal. Biochem.*, 2003, **321**, 202–208.
- 13 Bioluminogenic variants of this method have been reported as well. See: F. Halley, J. Reinshagen, B. Ellinger, M. Wolf, A. L. Niles, N. J. Evans, T. A. Kirkland, J. M. Wagner, M. Jung, P. Gribbon and S. Gul, *J. Biomol. Screening*, 2011, **16**, 1227–1235.
- 14 R. Baba, Y. Hori and K. Kikuchi, *Chem.-Eur. J.*, 2015, **21**, 4695–4702.
- 15 A. Vannini, C. Volpari, G. Filocamo, E. C. Casavola, M. Brunetti, D. Renzoni, P. Chakravarty, C. Paolini, R. De Francesco, P. Gallinari, C. Steinkühler and S. Di Marco, *Proc. Natl. Acad. Sci. U. S. A.*, 2004, **101**, 15064–15069.
- 16 M. S. Finnin, J. R. Donigian, A. Cohen, V. M. Richon, R. A. Rifkind, P. A. Marks, R. Breslow and N. P. Pavletich, *Nature*, 1999, **401**, 188–193.
- 17 A. V. Bieliauskas and M. K. H. Pflum, *Chem. Soc. Rev.*, 2008, **37**, 1402–1413.
- 18 K. E. Secor and T. E. Glass, *Org. Lett.*, 2004, **6**, 3727–3730.
- 19 R.-Y. Tang, P. Zhong and Q.-L. Lin, *Synthesis*, 2007, **2007**, 85–91.
- 20 E. K. Feuster and T. E. Glass, *J. Am. Chem. Soc.*, 2003, **125**, 16174–16175.
- 21 C. M. Cooley, K. S. Hettie, J. L. Klockow, S. Garrison and T. E. Glass, *Org. Biomol. Chem.*, 2013, **11**, 7387–7392.
- 22 K. S. Hettie, X. Liu, K. D. Gillis and T. E. Glass, *ACS Chem. Neurosci.*, 2013, **4**, 918–923.
- 23 V. Perissi, K. Jepsen, C. K. Glass and M. G. Rosenfeld, *Nat. Rev. Genet.*, 2010, **11**, 109–123.
- 24 HPLC analysis of the mixture at the end of each experiment was used to verify the amount of product obtained, based on calibration conducted with pure samples prepared independently.
- 25 J.-H. Zhang, T. D. Y. Chung and K. R. Oldenburg, *J. Biomol. Screening*, 1999, **4**, 67–73.
- 26 M. Göttlicher, S. Minucci, P. Zhu, O. H. Krämer, A. Schimpf, S. Giavara, J. P. Sleeman, F. Lo Coco, C. Nervi, P. G. Pelicci and T. Heinzel, *EMBO J.*, 2001, **20**, 6969–6978.
- 27 D. M. Fass, R. Shah, B. Ghosh, K. Hennig, S. Norton, W.-N. Zhao, S. A. Reis, P. S. Klein, R. Mazitschek,



- R. L. Maglathlin, T. A. Lewis and S. J. Haggarty, *ACS Med. Chem. Lett.*, 2011, **2**, 39–42.
- 28 N. Gurvich, O. M. Tsygankova, J. L. Meinkoth and P. S. Klein, *Cancer Res.*, 2004, **64**, 1079–1086.
- 29 Absorption at 510 nm shows a smaller contribution from the aldehyde than absorption at 499 nm.
- 30 H. V. K. Diyabalanage, G. C. van de Bittner, E. L. Ricq and J. M. Hooker, *ACS Chem. Biol.*, 2014, **9**, 1257–1262.
- 31 Some variability in the conversion rate was observed between experiments conducted with different batches of the nuclear extract. In every case, however, the amount of product formed estimated by the spectroscopic measurements were in accordance with the values calculated by chromatographic analysis of the mixture at the end of the run.

