## **RSC Advances**



View Article Online

View Journal | View Issue

### PAPER

Check for updates

Cite this: RSC Adv., 2018, 8, 39893

Received 10th October 2018 Accepted 22nd November 2018

DOI: 10.1039/c8ra08391f

rsc.li/rsc-advances

# 2'-Deoxy-5-(hydroxymethyl)cytidine: estimation in human cancer cells with a simple chemosensor<sup>†</sup>

Himadri Sekhar Sarkar, 📴 a Shampa Kundu, a Sujoy Das, 回 a Pulak Kumar Maiti, <sup>b</sup> Sukhendu Mandal 🔟 <sup>b</sup> and Prithidipa Sahoo 🛈 \*a

A pyrrole-based rhodamine conjugate (CS-1) has been developed and characterized for the selective detection and quantification of 2'-deoxy-5-(hydroxymethyl)cytidine (5hmC) in human cancer cells with a simple chemosensing method.

2'-Deoxy-5-(hydroxymethyl)cytidine (5hmC) is found in both neuronal cells and embryonic stem cells. It is a modified pyrimidine and used to quantify DNA hydroxymethylation levels in biological samples1-3 as it is capable of producing interstrand cross-links in double-stranded DNA. It is produced through an enzymatic pathway carried out by the Ten-Eleven Translocation (TET1, TET2, TET3) enzymes, iron and 2-oxoglutarate dependent dioxygenase.<sup>4-7</sup> In the DNA demethylation process, methylcytosine is converted to cytosine and generates 5hmC as an intermediate in the first step of this process which is then further oxidized to 5-formylcytosine (fC) and 5-carboxycytosine (caC) of very low levels compared to the cytosine level.8 Though the biological function of 5hmC in the mammalian genome is still not revealed, the presence of a hydroxymethyl group can regulate gene expression (switch ON & OFF). Reports say that in artificial DNA 5hmC is converted to unmodified cytosine when introduced into mammalian cells.9,10

Levels of 5hmC substantially vary in different tissues and cells. It is found to be highest in the brain, particularly in nervous system and in moderate percentage in liver, colon, rectum and kidney tissues, whereas it is relatively low in lung and very low in breast and placenta.11,12 The percentage of 5hmC content is much less in cancer and tumor tissues compared to the healthy ones. The reason behind this loss is the absence of TET1, TET2, TET3, IDH1, or IDH2 mutations in most of the human cancer cells which means decrease of methylcytosine oxidation.13-15 This loss of 5hmC in cancer cells is being used as a diagnostic tool for the detection of early-stage of malignant disease. Few analytical methods<sup>16-19</sup> such as glucosyltransferase assays, tungsten-based oxidation systems, and TET-assisted bisulfite sequencing (TAB-Seq) or oxidative bisulfite sequencing (oxBS-Seq) protocols are now developed to

differentiate 5hmC from other nucleotide which are naturally occurred. There are also few methods such as liquid chromatography/tandem mass spectroscopy (LC/MS-MS), which determine the level of 5hmC in mammalian cancer cell.<sup>20-22</sup> However, these procedures are highly toxic and expensive due to requirement of catalyzation through enzymes or heavy metal ion and these techniques require expertise, facilities, much time and costs even beyond standard DNA sequencing. As a result, these detection techniques are currently inappropriate for the high-throughput screening of genome-wide 5hmC levels (performance comparison is shown in Table S1, ESI†).

Among all reputed methods fluorescence detection method using chemosensors is significantly important due to its indispensable role in medicinal and biological applications.<sup>23–27</sup> Chemosensors have been effectively explored to monitor biochemical processes and assays through *in situ* analysis in living systems and abiotic samples with much less time and cost.

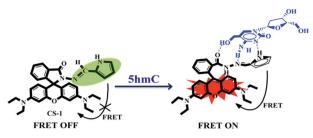
In this contribution we prepared and characterize (Scheme S1 and Fig. S1–S3, ESI†) a pyrrole–rhodamine based chemosensor (**CS-1**) which shows efficient and selective fluorescence signal for 5hmC in aqueous medium (Scheme 1). A transparent single crystal of **CS-1** (Fig. 1) was obtained by slow evaporation of the solvent from a solution of **CS-1** in CH<sub>3</sub>CN. It crystallizes as monoclinic with space group  $P2_1/n$  (Fig. S4 and Table S2, ESI†).

Spectrophotometric and spectrofluorimetric titrations were carried out to understand the **CS-1**–5hmC interaction with 1 : 1 binding stoichiometry (Fig. S5, ESI†) upon adding varying concentrations of 5hmC to a fixed concentration of **CS-1** (1  $\mu$ M) in aqueous medium at neutral pH. Upon the addition of increasing concentrations of the 5hmC, a clear absorption band ( $K_a = 4.47 \times 10^5 \text{ M}^{-1}$ , Fig. S6, ESI†) appeared to be centered at 556 nm with increasing intensity (Fig. 2a). On the other hand, for the fluorescence emission spectra of **CS-1** (Fig. 2b), upon irradiation at 325 nm, an emission maxima at 390 nm was observed, which was attributed to the fluorescence emission from the donor unit *i.e.* the pyrrole moiety of **CS-1**. When 5hmC were added, due to rhodamine moiety **CS-1** showed a 95-fold

<sup>&</sup>lt;sup>a</sup>Department of Chemistry, Visva-Bharati University, Santiniketan-731235, India. E-mail: prithidipa@hotmail.com

<sup>&</sup>lt;sup>b</sup>Department of Microbiology, University of Calcutta, Kolkata-700073, India

<sup>&</sup>lt;sup>†</sup> Electronic supplementary information (ESI) available. CCDC 1855229. For ESI and crystallographic data in CIF or other electronic format see DOI: 10.1039/c8ra08391f



Scheme 1 5hmC-induced FRET OFF-ON mechanism of the chemosensor CS-1.

increase in fluorescence at 565 nm ( $K_a = 4.61 \times 10^5 \text{ M}^{-1}$ , Fig. S7, ESI<sup>†</sup>) with the detection limit of 8 nM (Fig. S8, ESI<sup>†</sup>). The binding of 5hmC induces opening of the spirolactam ring in **CS**-**1**, inducing a shift of the emission spectrum. Subsequently, increased overlap between the emission of the energy-donor (pyrrole) and the absorption of the energy-acceptor (rhodamine) greatly enhances the intramolecular FRET process,<sup>28,29</sup> producing an emission from the energy acceptor unit in **CS-1**.

In order to establish the sensing selectivity of the chemosensor **CS-1**, parallel experimentations were carried out with other pyrimidine/purine derivatives such as 5-methylcytosine, cytosine, cytidine, thymine, uracil, 5-hydroxymethyluracil, adenine and guanine. Comparing with other pyrimidine/purine derivatives the abrupt fluorescence enhancement was found upon addition of 5hmC to **CS-1** while others do not make any fluorescence changes under UV lamp (Fig. 3, lower panel). Furthermore, the prominent color change from colorless to deep pink allows 5hmC to be detected by naked eye (Fig. 3, upper panel). The above observation shows consistency with the fluorescence titration experiments where no such binding of **CS-1** with other pyrimidine/purine derivatives was found (Fig. S9, ESI<sup>†</sup>).

pH titration reveals that **CS-1** becomes fluorescent below pH 5 due to the spirolactam ring opening of rhodamine. However, it is non-fluorescent at pH range of 5–13. Upon addition of 5hmC to **CS-1** shows deep red fluorescence in the pH range of 5–8 (Fig. S10, ESI†). Considering the biological application and the practical applicability of the chemosensor pH 7.4 has been preferred to accomplish all experiments successfully.

In <sup>1</sup>H NMR titration (Fig. S11, ESI<sup>†</sup>), the most interesting feature is the continuous downfield shift of aromatic protons on

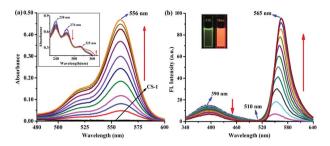


Fig. 2 (a) UV-vis absorption spectra of CS-1 (1  $\mu$ M) upon gradual addition of 5hmC up to 1.2 equiv. in H<sub>2</sub>O-CH<sub>3</sub>CN (15 : 1, v/v) at neutral pH. (b) Fluorescence emission spectra of CS-1 (1  $\mu$ M) upon addition of 1.2 equiv. of 5hmC in H<sub>2</sub>O-CH<sub>3</sub>CN (15 : 1, v/v) at neutral pH ( $\lambda_{ex} = 325$  nm).

the pyrrole moiety of **CS-1** upon gradual addition of 5hmC. This may be explained as the decrease in electron density of the pyrrole moiety upon binding with 5hmC through hydrogen bonding. Xanthene protons to be shifted downfield upon spirolactam ring opening indicates the probe to coordinate with 5hmC and electrons are accumulated around 5hmC. In <sup>13</sup>C NMR titration the spiro cycle carbon peak at 65 ppm was shifted to 138 ppm along with a little downfield shift of the aromatic region of **CS-1** (Fig. S12, ESI†). This coordination led to the spiro cycle opening and changes to the absorption and emission spectra, further evident by mass spectrometry (Fig. S13, ESI†), which corroborates the stronger interaction of **CS-1** with 5hmC.

The experimental findings were validated by density functional theory (DFT) calculations using the 6-31G+(d,p) method basis set implemented at Gaussian 09 program. Energy optimization calculations presented the conformational changes at the spirolactam position of **CS-1** while 5hmC takes part to accommodate a probe molecule. After **CS-1**–5hmC complexation the energy is minimized by 19.45 kcal from the chemosensor **CS-1**, indicating a stable complex structure (Fig. 4 and Table S3, ESI†). This theoretical study strongly correlates the experimental findings.

The desirable features of **CS-1** such as high sensitivity and high selectivity at physiological pH encouraged us to further evaluate the potential of the chemosensor for imaging 5hmC in live cells (Fig. 5). A549 cells (Human cancer cell A549, ATCC no. CCL-185) treated with **CS-1** (1  $\mu$ M) exhibited weak fluorescence, whereas a deep red fluorescence signal was observed in the cells stained with **CS-1** (1  $\mu$ M) and 5hmC (10  $\mu$ M), which is in good



Fig. 1 ORTEP diagram of CS-1 (ellipsoids are drawn at 40% probability level).

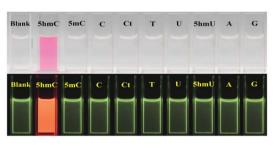


Fig. 3 Visible color (top) and fluorescence changes (bottom) of CS-1 (1  $\mu$ M) in aqueous medium upon addition of 1.2 equiv. of various pyrimidine/purine derivatives ( $\lambda_{ex} = 325$  nm) in H<sub>2</sub>O-CH<sub>3</sub>CN (15 : 1, v/ v) at neutral pH.

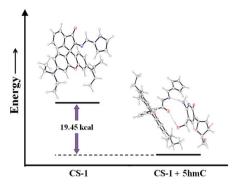


Fig. 4 Energy diagram showing the energy differences between CS-1 and CS-1–5hmC complex.

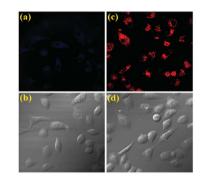
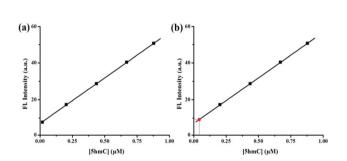


Fig. 5 Confocal microscopic images of A549 cells treated with CS-1 and 5hmC. (a) Cells treated with only CS-1 at 1  $\mu$ M concentration. (b) Bright field image of (a). (c) Cells treated with CS-1 and 5hmC at concentration 10  $\mu$ M. (d) Bright field image of (c). All images were acquired with a 60× objective lens with the applied wavelengths: For (a) and (b),  $E_{ex} = 341$  nm,  $E_{em} = 414$  nm, filter used: DIDS; for (c) and (d)  $E_{ex} = 550$  nm,  $E_{em} = 571$  nm, filter used: Rhod-2.



**Fig. 6** (a) Calibration curve obtained for the estimation of 5hmC. (b) Estimation of the concentration of 5hmC (red point) from the calibration curve.

agreement with the FRET OFF–ON profile of the chemosensor **CS-1** in presence of 5hmC, thus corroborating the in-solution observation (Fig. S14, ESI<sup>†</sup>). Cytotoxicity assay measurement shows that the chemosensor **CS-1** does not have any toxicity on the tested cells and **CS-1**–5hmC complex does not exert any significant adverse effect on cell viability at tested concentrations (Fig. S15, ESI<sup>†</sup>). As far as we are aware, this is the first report where we are executing the possible use of the pyrrole–rhodamine based chemosensor for selective recognition of 5hmC in living cells. These findings open an avenue for future biomedical applications of the chemosensor to recognize 5hmC.

The concentration of 5hmC was also quantified from A549 human cancer cells. Lysate of  $10^7$  A549 cells was added to 1  $\mu$ M of **CS-1** and the fluorescence signal was recorded. Presence of 5hmC in these cancer cells was detected with the help of **CS-1**–5hmC standard fluorescence curve (Fig. 6) using the selective detection ability of the chemosensor **CS-1**.

From the standard curve it was found that the concentration of 5hmC in the tested sample was  $0.034 \,\mu$ M present in 16.7 mm<sup>3</sup> A549 cell volume (Table 1). The above result was authenticated by spiking a real sample with a known concentration of 5hmC and different fluorescence signals were observed by adding known volumes from each sample to the **CS-1**. The recovery of the spiked samples was estimated to be over 99% (Table S4, ESI†). Assay of 5hmC was further validated from multiple samples of A549 human cancer cells using **CS-1**. Increasing fold of fluorescence signals was also statistically validated after calculating the Z' value (Table S5, ESI†). All tested samples shows the Z' score value more than 0.9, indicating an optimized and validated assay of 5hmC.

#### Conclusions

In conclusion, a chemosensor **CS-1** has been developed for the rapid detection of 5hmC with low cost. The selective detection and quantification of 5hmC was successfully demonstrated in human cancer cells at neutral pH with very low concentration. With this potentiality of **CS-1** one can successfully apply this method to estimate 5hmC in disease cancer tissues and other biological samples of patients with metabolic dysfunction or various carcinomas.

#### Conflicts of interest

There is no conflicts to declare.

Sample	<b>CS-1</b> used (μM)	Initial 5hmC used	Addition of exogenous 5hmC (µM)	Amount of 5hmC derived from fluorescence signal (µM)	Fluorescence signal recovery (%)
1	1	5hmC present in 16.7 mm <sup>3</sup> A549 cell	0	0.034	_
2	1	volume	1	1.028	99.4
3	1		3	4.019	99.6
4	1		5	5.012	99.5

 Table 1
 Quantification of 5hmC in human cancer cell A549

#### Acknowledgements

P. S. acknowledges SERB-DST, Govt. of India for awarding her the young scientist grant [Project file no. SB/FT/CS-021/2014]. H. S. S. is sincerely thankful to UGC, Delhi for research fellowship. Authors are also grateful to DBT-IPLS facility, CU.

#### Notes and references

- 1 T. Le, K. P. Kim, G. Fan and K. F. Faull, *Anal. Biochem.*, 2011, **412**, 203–209.
- 2 A. M. Krais, Y. J. Park, C. Plass and H. H. Schmeiser, *Epigenetics*, 2011, **6**, 560–565.
- 3 T. A. Clark, X. Lu, K. Luong, Q. Dai, M. Boitano, S. W. Turner, C. He and J. Korlach, *BMC Biol.*, 2013, **11**, 1–10.
- 4 L. M. Iyer, M. Tahiliani, A. Rao and L. Aravind, *Cell Cycle*, 2009, **8**, 1698–1710.
- 5 C. Loenarz and C. J. Schofield, Chem. Biol., 2009, 16, 580-583.
- 6 S. Ito, A. C. D'Alessio, O. V. Taranova, K. Hong, L. C. Sowers and Y. Zhang, *Nature*, 2010, **466**, 1129–1133.
- 7 M. Ko, Y. Huang, A. M. Jankowska, U. J. Pape, M. Tahiliani,
  H. S. Bandukwala, J. An, E. D. Lamperti, K. P. Koh,
  R. Ganetzky, X. S. Liu, L. Aravind, S. Agarwal,
  J. P. Maciejewski and A. Rao, *Nature*, 2010, 468, 839–843.
- M. Tahiliani, K. P. Koh, Y. Shen, W. A. Pastor, H. Bandukwala, Y. Brudno, S. Agarwal, L. M. Iyer, D. R. Liu, L. Aravind and A. Rao, *Science*, 2009, **324**, 930–935.
- 9 M. A. Hahn, P. E. Szabó and G. P. Pfeifer, *Genomics*, 2014, 104, 314–323.
- 10 D.-Q. Shi, I. Ali, J. Tang and W.-C. Yang, *Front. Genet.*, 2017, 8, 100.
- M. Münzel, D. Globisch, T. Bruckl, M. Wagner, V. Welzmiller, S. Michalakis, M. Muller, M. Biel and T. Carell, *Angew. Chem., Int. Ed. Engl.*, 2010, 49, 5375–5377.
- 12 S. G. Jin, X. Wu, A. X. Li and G. P. Pfeifer, *Nucleic Acids Res.*, 2011, **39**, 5015–5024.

- 13 Y. Kudo, K. Tateishi, K. Yamamoto, S. Yamamoto, Y. Asaoka,
  H. Ijichi, G. Nagae, H. Yoshida, H. Aburatani and K. Koike, *Cancer Sci.*, 2012, **103**, 670–676.
- 14 H. Yang, Y. Liu, F. Bai, J. Y. Zhang, S. H. Ma, J. Liu, Z. D. Xu, H. G. Zhu, Z. Q. Ling, D. Ye, K. L. Guan and Y. Xiong, *Oncogene*, 2013, 32, 663–669.
- 15 L. I. Kroeze, B. A. van der Reijden and J. H. Jansen, *Biochim. Biophys. Acta*, 2015, **1855**, 144–154.
- 16 A. Szwagierczak, S. Bultmann, C. S. Schmidt, F. Spada and H. Leonhardt, *Nucleic Acids Res.*, 2010, **38**, e181.
- 17 C. X. Song, T. A. Clark, X. Y. Lu, A. Kislyuk, Q. Dai, S. W. Turner, C. He and J. Korlach, *Nat. Methods*, 2012, **9**, 75.
- 18 M. J. Booth, M. R. Branco, G. Ficz, D. Oxley, F. Krueger, W. Reik and S. Balasubramanian, *Science*, 2012, 336, 934.
- 19 M. J. Booth, T. W. B. Ost, D. Beraldi, N. M. Bell, M. R. Branco, W. Reik and S. Balasubramanian, *Nat. Protoc.*, 2013, 8, 1841.
- 20 L. T. Zhang, L. J. Zhang, J. J. Zhang, X. X. Ye, A. M. Xie, L. Y. Chen, J. X. Kang and C. Cai, *Bioanalysis*, 2013, 5, 839– 845.
- 21 O. K. Zahid, B. S. Zhao, C. He and A. R. Hall, *Sci. Rep.*, 2016, 6, 29565.
- 22 S. Yan, X. Xu, P. Guo, J. Hu, C. Wang, R. Huang, X. Weng, Y. Dua and X. Zhou, *RSC Adv.*, 2013, **3**, 12066.
- 23 P. Sahoo, H. S. Sarkar, S. Das, K. Maiti, M. R. Uddin and S. Mandal, *RSC Adv.*, 2016, **6**, 66774–66778.
- 24 H. S. Sarkar, S. Das, M. R. Uddin, S. Mandal and P. Sahoo, *Asian J. Org. Chem.*, 2017, **6**, 71–75.
- 25 H. S. Sarkar, S. Das, K. Rissanen and P. Sahoo, *Anal. Chem.*, 2017, **89**, 13054–13057.
- 26 H. S. Sarkar, S. Das, D. Mandal, M. R. Uddin, S. Mandal and P. Sahoo, *RSC Adv.*, 2017, 7, 54008–54012.
- 27 H. S. Sarkar, A. Ghosh, S. Das, P. Maiti, S. Maitra, S. Mandal and P. Sahoo, *Sci. Rep.*, 2018, **8**, 3402.
- 28 Y. Kurishita, T. Kohira, A. Ojida and I. Hamachi, *J. Am. Chem. Soc.*, 2010, **132**, 13290.
- 29 M. Tang, L. Wu, D. Wua, C. Huang, W. Zhu, Y. Xua and X. Qiana, *RSC Adv.*, 2016, **6**, 34996.