



Selective binding and controlled release of anticancer drugs by polyanionic cyclodextrins

Jian-Guang Cheng^a, Hua-Jiang Yu^a, Yong Chen^a, Yu Liu^{a,b,*}

^a College of Chemistry, State Key Laboratory of Elemento-Organic Chemistry, Nankai University, PR China

^b Collaborative Innovation Center of Chemical Science and Engineering (Tianjin), Tianjin 300071, PR China

ARTICLE INFO

Article history:

Received 6 February 2018

Revised 8 March 2018

Accepted 8 March 2018

Available online 10 March 2018

Keywords:

Polyanionic cyclodextrin

Selective binding

Controlled release

Anticancer drugs

Supramolecular chemistry

ABSTRACT

The binding stoichiometry, binding constants, and inclusion mode of some water-soluble negatively charged cyclodextrin derivatives, i.e. heptakis-[6-deoxy-6-(3-sulfanylpropanoic acid)]- β -cyclodextrin (**H1**), heptakis-[6-deoxy-6-(2-sulfanylacetic acid)]- β -cyclodextrin (**H2**), mono-[6-deoxy-6-(3-sulfanylpropanoic acid)]- β -cyclodextrin (**H3**) and mono-[6-deoxy-6-(2-sulfanylacetic acid)]- β -cyclodextrin (**H4**), with three anticancer drugs, i.e. irinotecan hydrochloride; topotecan hydrochloride; doxorubicin hydrochloride, were investigated by means of ¹H NMR, UV-Vis spectroscopy, mass spectra and 2D NMR. Polyanionic cyclodextrins **H1-H2** showed the significantly high binding abilities of up to 2.6×10^4 – 2.0×10^5 M⁻¹ towards the selected anticancer drugs, which were nearly 50–1000 times higher than the corresponding Ks values of native β -cyclodextrin. In addition, these polyanionic cyclodextrins also showed the pH-controlled release behaviors. That is, the anticancer drugs could be efficiently encapsulated in the cyclodextrin cavity at a pH value similar to that of serum but sufficiently released at an endosomal pH value of a cancer cell, which would make these cyclodextrin derivatives the potential carriers for anticancer drugs.

© 2018 Elsevier Ltd. All rights reserved.

1. Introduction

Recently, numerous effective anticancer drugs have been used for the treatment of various human and animal cancers. Among them, irinotecan hydrochloride (**CPT-11**), topotecan hydrochloride (**TPT**) and doxorubicin hydrochloride (**DOX**) are three prominent leader compounds. **CPT-11** and **TPT** are both water-soluble semi-synthetic derivatives of the alkaloid camptothecin.^{1,2} **CPT-11** exhibits remarkable antitumor activity in clinical trials against a variety of human tumors,^{3–5} including colorectal cancer, lung cancer and malignant lymphoma.^{6–8} **TPT** is used clinically in the treatment of relapsed ovarian, lung cancer, and cervical cancer.^{9–12} **DOX** is a chemotherapeutic agent used for the treatment of a wide variety of human malignancies with an anthracycline structure, which consists of an aglycon, adriamycinone, combined with an amino sugar, daunosamine.^{13–16} On the other hand, cyclodextrins (CDs), a class of cyclic oligosaccharides linked by 1,4-glucose bonds, are water-soluble, nontoxic, compounds commercially available at low price,^{17–23} and their torus-shaped cavity can bind various inorganic/organic/biological molecules. This excellent

property enables the wide application of CDs in fields of molecular recognition and molecular assembly. Among the CD family, the most used one is β -CD that contains 7 glucose units.^{24–28} Nevertheless, the complex stability constants (Ks) between native β -CD and anticancer drugs (**CPT-11**, **TPT** and **DOX**) are very limited,²⁹ i.e. 2.6×10^2 M⁻¹ for β -CD/**CPT-11** pair, 8.8×10^3 M⁻¹ for β -CD/**TPT** pair, and 2.1×10^2 M⁻¹ for β -CD/**DOX** pair respectively, which greatly restricts the application of β -CD as carriers of anticancer drugs. Recently, the negatively charged CD derivatives have attracted more attention because of their potential applications in drug delivery. For example, Zhang et al. reported a negatively charged CD named ORG25969 as a good acceptor to give an extraordinarily high binding affinity towards rocuronium bromide (Ks up to 10^7 M⁻¹), and thus can be clinically used as a reversal agent in the post-operative recovery.³⁰ Wenz and Apostolakis et al. synthesized a series of negatively charged CDs and researched their binding behaviors with camptothecin. The result showed that the stabilities of camptothecin complexes obtained from solubility measurements of negatively charged CD derivatives were significantly higher than those of other reported CD derivatives.^{31,32} Herein, we selected four negatively charged CD derivatives, i.e. heptakis-[6-deoxy-6-(3-sulfanylpropanoic acid)]- β -CD (**H1**), heptakis-[6-deoxy-6-(2-sulfanylacetic acid)]- β -CD (**H2**), mono-[6-deoxy-6-(3-sulfanylpropanoic acid)]- β -CD (**H3**) and mono-[6-

* Corresponding author at: College of Chemistry, State Key Laboratory of Elemento-Organic Chemistry, Nankai University, PR China.

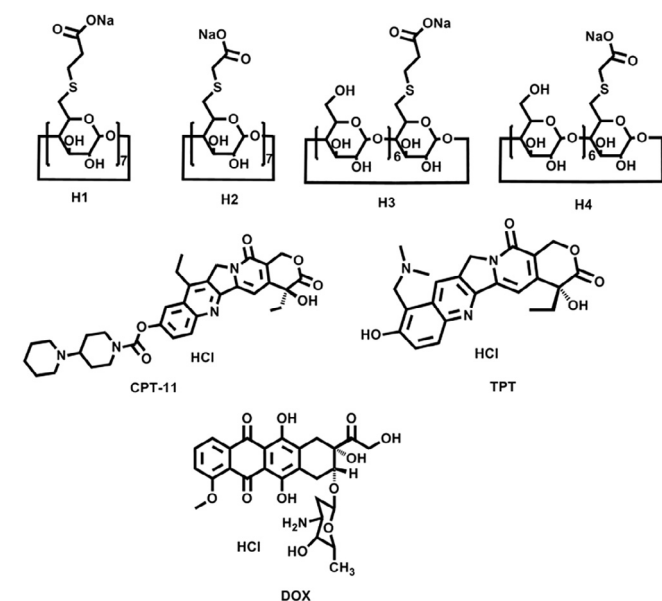
E-mail address: yuliu@nankai.edu.cn (Y. Liu).

deoxy-6-(2-sulfanylacetic acid)]- β -CD (**H4**),³³ and investigated their selective binding and controlled release behaviors towards anticancer drugs **CPT-11**, **TPT** and **DOX** (Scheme 1). Significantly, with binding abilities much stronger than those of most previously reported CD derivatives, these polyanionic CDs exhibited the pH-responsive release of drug in a cancer cell environment. That is, the polyanionic CD/anticancer drug complex was stable in a biological environment such as serum (pH 7.2), but efficiently released the encapsulated anticancer drug at pH 5.7 (endosomal pH values of a cancer cell).

2. Results and discussion

2.1. Job plots and binding constants of **H1-H4** and anticancer drugs

UV-vis spectroscopy was employed to determine the host-guest binding stoichiometry. As shown in Fig. 1, the Job plot of **H1/CPT-11** in water gave a maximum at molar fraction of 0.5, indicating that **H1** formed stoichiometric 1:1 inclusion complex with **CPT-11**. Moreover, the mass spectrum measurements (Figs. S31–S33) also demonstrated the formation of 1:1 inclusion complexes between cyclodextrin hosts and anticancer drugs. The quantitative investigation on the molecular binding behavior of **H1** with **CPT-11** was examined by means of UV-vis spectral titration, wherein the UV-vis spectra of a series of solutions containing the same amounts of **CPT-11** and different amounts of **H1** were measured to determine the binding constant between **CPT-11** and **H1**. As can be seen from Fig. S20, with the addition of **H1**, the absorbance maximum of **CPT-11** slightly decreased, accompanied by the appreciable red shift of maximum wavelength. By using the nonlinear least-squares method,³⁴ the stability constants (K_s) values could be calculated as $(1.7 \pm 0.2) \times 10^5 \text{ M}^{-1}$ according to the sequential changes of absorbance intensity of **CPT-11** with the different concentrations of **H1**. Similar 1:1 binding stoichiometry was also found in the association of hosts **H1-H4** with anticancer drugs **CPT-11**, **TPT** and **DOX**, and the corresponding stability constants (K_s) were determined (Fig. 2) and listed in Table 1. Moreover, we also tried to use isothermal titration calorimetry to determine the binding constants. However, the isothermal titration calorimetry experiment required the higher concentrations, and the inclusion complex formed precipitate under such a concentration.



Scheme 1. Chemical structures of host and guest.

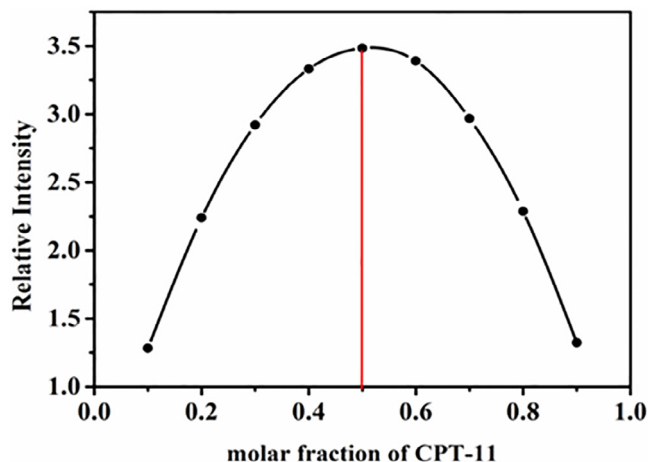


Fig. 1. Job plot for the binding of **H1** with **CPT-11** in water at 25 °C, indicating a 1:1 stoichiometry. The changes of absorbance were measured at 369 nm, and the total concentration was maintained at 0.1 mM.

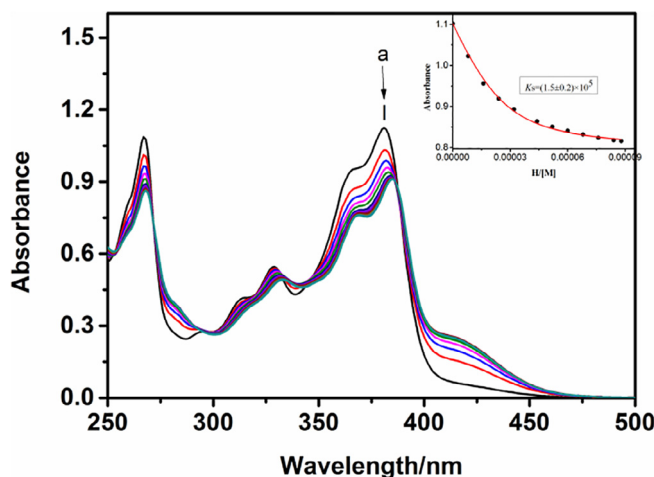


Fig. 2. UV-vis spectral titration of **TPT** upon addition of **H1** in H_2O at 25 °C. The nonlinear least-squares analysis (inset) of the differential absorbance to calculate the complex stability constant. The changes of absorbance were measured at 381 nm. ($[\text{TPT}] = 0.05 \text{ mM}$, $[\text{H1}] = 0.00, 0.01, 0.02, 0.04, 0.06, 0.08, 0.10, 0.12, 0.14, 0.16, 0.18, 0.20, 0.22 \text{ mM}$ from a to l).

Accordingly, the encapsulation and loading efficiency of anticancer drugs by hosts were calculated and listed in Table 1. As seen in Table 1, the native β -CD only showed very poor binding ability towards the selected anticancer drugs. Possessing an anionic side arm on the β -CD rim, host **H3** or **H4** showed the moderate binding ability (1.02×10^3 – $1.7 \times 10^4 \text{ M}^{-1}$) towards anticancer drugs owing to the electrostatic interactions between the anionic side arm of host and the cationic guest. However, host **H1** or **H2** showed a significantly increased binding ability towards anticancer drugs up to 2.6×10^4 – $2.0 \times 10^5 \text{ M}^{-1}$, which was nearly 50–1000 times higher than the corresponding K_s values of native β -CD. A possible reason may be that the seven anionic side arms on **H1** or **H2** (either of **H1** or **H2** possesses 7 negative charges) gave the greatly strengthened electrostatic interactions with the cationic guest. Moreover, the extended cavity formed by seven side arms may also provide the additional van der Waals and hydrophobic interactions towards the accommodated drug. As a result, host **H1** exhibited the fairly high encapsulation efficiency (>75%) and loading efficiency (>18%) towards the selected anticancer drugs when the concentrations of anticancer drugs and hosts were fixed at 0.1 mM, which enables it as a good candidate of anticancer drug carriers. The anti-

Table 1
Stability constants (K_s), encapsulation efficiency and loading efficiency of anticancer drugs by negatively charged cyclodextrins in water at 25 °C.

Host	Guest	K_s/M^{-1}	Encapsulation efficiency/%	Loading efficiency/%
β -CD	CPT-11	2.6×10^2 [36]	2.29	1.26
	TPT	8.8×10^3 [37]	36.0	14.5
	DOX	2.1×10^2 [38]	2.02	1.03
H1	CPT-11	$(1.7 \pm 0.2) \times 10^5$	78.5	25.7
	TPT	$(1.5 \pm 0.2) \times 10^5$	77.3	18.6
	DOX	$(2.0 \pm 0.31) \times 10^5$	80.0	24.4
H2	CPT-11	$(2.6 \pm 0.2) \times 10^4$	54.3	18.7
	TPT	$(4.4 \pm 0.4) \times 10^4$	62.4	15.8
	DOX	$(2.6 \pm 0.2) \times 10^4$	54.3	17.4
H3	CPT-11	$(5.1 \pm 0.5) \times 10^3$	27.1	13.6
	TPT	$(1.7 \pm 0.2) \times 10^4$	47.3	17.4
	DOX	$(2.3 \pm 0.1) \times 10^3$	16.2	7.53
H4	CPT-11	$(4.1 \pm 0.1) \times 10^3$	23.8	12.0
	TPT	$(9.7 \pm 0.8) \times 10^3$	37.7	14.0
	DOX	$(1.02 \pm 0.07) \times 10^3$	8.53	4.02

cancer drugs encapsulation efficiency and loading efficiency was calculated by the following formulas:³⁵

$$\text{encapsulation efficiency}(\%) = (m_{\text{loaded}}/m_D) \times 100$$

$$\text{loading efficiency}(\%) = (m_{\text{loaded}}/m_{cd}) \times 100$$

$$m_{\text{loaded}} = \frac{1}{2} M_D v \left[([H]_0 + [G]_0 + \frac{1}{K_s}) - \sqrt{([H]_0 + [G]_0 + \frac{1}{K_s})^2 - 4[H]_0[G]_0} \right]$$

where m_{loaded} is the mass of anticancer drugs that formed inclusion complex with hosts, m_D is the mass of anticancer drugs added, m_{cd} is the mass of hosts, $[G]_0$ is the initial concentration of anticancer drugs added, $[H]_0$ is the initial concentration of hosts, and M_D , v and K_s are molecular weight of anticancer drugs, volume of the solution, and the stability constants of the inclusion complex, respectively. In addition, the solubility and stability of inclusion complexes in 10% serum solution were also investigated. The results showed that the solubility could reach 0.5 mmol/mL, and the complex could keep stable for at least 24 h.

2.2. Binding mode of **H1** with anticancer drugs

2D NMR spectroscopy is an essential method to investigate binding mode between host and guest. As shown in Fig. 3, we could

see NOE correlations between protons of **CPT-11** and interior protons of **H1** from the ROESY spectrum of an equimolar mixture of **H1** with **CPT-11**. The cross peak A was assigned to NOE correlations between H18/H23 protons of **CPT-11** and H3 protons of **H1**, and the cross peak B was assigned to NOE correlations between H28 protons of **CPT-11** and H3/H5 protons of **H1**, and H5 protons gave stronger NOE correlations than H3 protons. Moreover, the cross peak C was assigned to NOE correlations between H11/H12/H13/H14 protons of **CPT-11** and H3 protons of **H1**. Therefore, we deduced that the **CPT-11** guest entered the **H1** cavity from the wide side. Based on the ROESY and molecular simulation experiments, the geometry of inclusion complex of H3 with CPT-11 was proposed where the **CPT-11** guest entered the **H1** cavity from the wide side.

2.3. Controlled release of anticancer drugs

In addition to the UV–vis spectral changes, the association with polyanionic CDs also led to the obvious fluorescence and color changes of anticancer drugs, which could be readily monitored by fluorescence spectra or naked eyes. For example, the fluorescence intensity of **DOX** showed the obvious decrease, and its color turned dark, after the association with **H1** (Fig. 4). In the control experiment, the pH dependence of DOX fluorescence changes in the absence of β -CD hosts was relatively limited. Therefore, the flu-

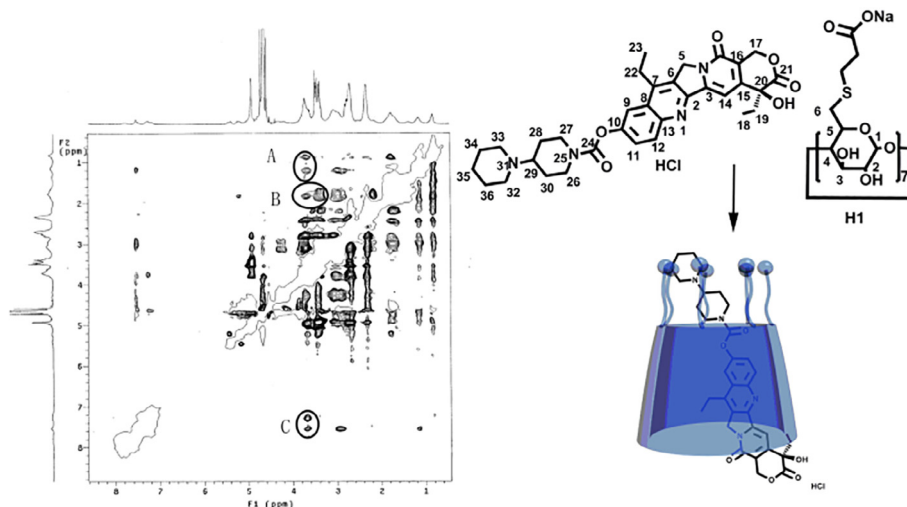


Fig. 3. 2D ROESY spectra of **H1**•**CPT-11** complex in D_2O . The cross peaks indicate intermolecular interactions between **CPT-11** and **H1**.

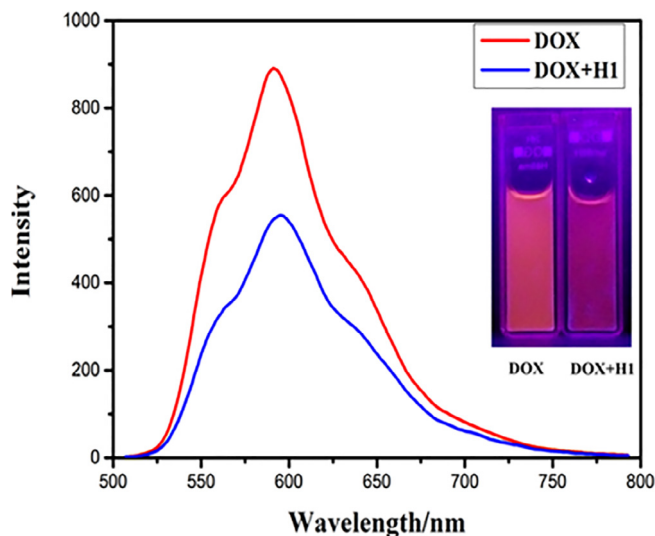


Fig. 4. Fluorescence spectral changes of **DOX** upon addition of **H1**. inset: photographs of fluorescence changes of **DOX** upon addition of **H1** at 25 °C. ([**DOX**] = 0.05 mM, [**H1**] = 0.05 mM.)

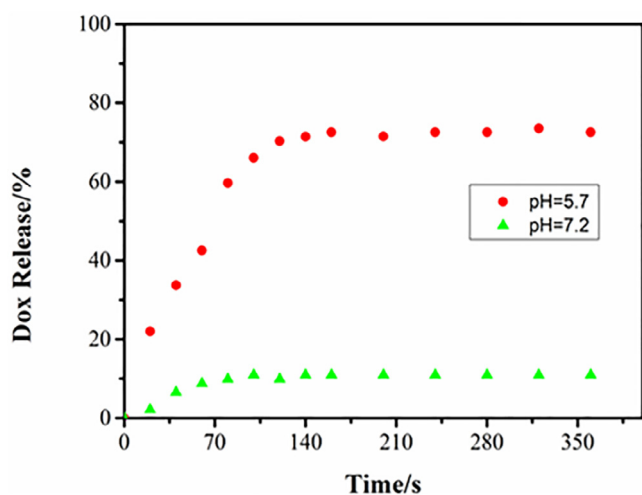


Fig. 5. Release of **H1/DOX** at different pH in 0.01 M sodium phosphate buffer (pH 7.2 and pH 5.7) at 25 °C. ([**DOX**] = 0.05 mM, [**H1**] = 0.05 mM.)

orescence spectra were used to investigate the controlled release behaviors of polyanionic CD/anticancer drug systems at different physiological pH values. As shown in Fig. 5, a very slow release of **DOX** was observed at pH 7.2, indicating that **H1/DOX** was stable in a biological environment such as serum. However, the release rate was significantly enhanced when the polyanionic CD/anticancer drug system was placed in an acidic solution. That is, > 70% was released at pH 5.7 (endosomal pH values of a cancer cell) within 140 s. The binding/release of **DOX** at different pH was mainly owing to the different binding abilities of **H1** towards **DOX** at different pH. That is, the binding ability of **H1** towards **DOX** was much stronger at pH 7.2, which led to the lower release level at pH 7.2. In addition, the UV–vis spectroscopy experiment demonstrated that the drug binding/release could be reversible for several times. Other polyanionic CD/anticancer drug systems also showed the similar controlled release behaviors of their drugs. This pH-responsive release of drug in a cancer cell environment will not only improve its cytotoxic efficacy against tumor cells but also reduce the toxicity of drug to normal tissues.

3. Conclusion

In conclusion, the selective binding and controlled release behaviors of negatively charged CDs with some anticancer drugs were investigated by means of ^1H NMR, UV–Vis spectroscopy, mass spectra and 2D NMR, and their binding constants (K_s) can reach 10^5 M^{-1} level, which is higher than most previously reported CD derivatives. Therefore, we deduce that the present polyanionic CD/anticancer drug systems may find their application potential in cancer therapy.

Acknowledgment

We thank NNSFC (21432004, 21672113, 21772099 and 91527301) for financial support.

A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <https://doi.org/10.1016/j.bmc.2018.03.013>.

References

- Baudin E, Docao C, Gicquel C, et al. *Ann Oncol*. 2002;13:1806–1809.
- Joshi H, Sengupta A, Gavvala K, Hazra P. *RSC Adv*. 2014;4:1015–1024.
- Wierdl M, Morton CL, Nguyen NK, Redinbo MR, Potter PM. *Biochemistry*. 2004;43:1874–1882.
- Zhao H, Rubio B, Sapra P, et al. *Bioconjugate Chem*. 2008;19:849–859.
- Escoffre JM, Novell A, Serrière S, Lecomte T, Bouakaz A. *Mol Pharm*. 2013;10:2667–2675.
- Senter PD, Beam KS, Mixan B, Wahl AF. *Bioconjugate Chem*. 2001;12:1074–1080.
- Egreteau J, Boucher E, de Guibert S, et al. *Int J Gastrointest Cancer*. 2005;35:69–76.
- Raez LE, Rosado MF, Santos ES, Reis IM. *Lung Cancer*. 2004;45:131–132.
- Kim HR, Pereira CM, Han HY, Lee H. *J Anal Chem*. 2015;87:5356–5362.
- Mi Z, Malak H, Burke TG. *Biochemistry*. 1995;34:13722–13728.
- Rosca EV, Wright M, Gonitell R, Gedroyc W, Miller AD, Thanou M. *Mol Pharm*. 2015;12:1335–1346.
- Pan P, Li Y, Yu H, Sun H, Hou T. *J Chem Inf Model*. 2013;53:997–1006.
- Sultana R, Di Domenico F, Tseng M, et al. *J Proteome Res*. 2010;9:6232–6241.
- Cui H, Huan M-L, Ye W-L, et al. *Mol Pharm*. 2017;14:746–756.
- Aljabali AAA, Shukla S, Lomonossoff GP, Steinmetz NF, Evans DJ. *Mol Pharm*. 2013;10:3–10.
- Li H, Cui Y, Sui J, et al. *ACS Appl Mater Int*. 2015;7:15855–15865.
- Rekharsky MV, Inoue Y. *Chem Rev*. 1998;98:1875–1918.
- Villalonga R, Cao R, Fragoso A. *Chem Rev*. 2007;107:3088–3116.
- Takashima Y, Osaki M, Harada A. *J Am Chem Soc*. 2004;126:13588–13589.
- VanEtten RL, Sebastian JF, Clowes GA, Bender ML. *J Am Chem Soc*. 1967;89:3242–3253.
- Chen Y, Liu Y. *Adv Mater*. 2015;27:5403–5409.
- Ma X, Zhao Y. *Chem Rev*. 2015;115:7794–7839.
- Milović NM, Badžić JD, Kostić NM. *J Am Chem Soc*. 2004;126:696–697.
- Nelles G, Weisser M, Back R, Wohlfart P, Wenz G, Mittler-Neher S. *J Am Chem Soc*. 1996;118:5039–5046.
- Balabai N, Linton B, Napper A, Priyadarshy S, Sukharevsky AP, Waldeck DH. *J Phys Chem B*. 1998;102:9617–9624.
- Wang H, Liu K, Chen K-J, et al. *ACS Nano*. 2010;4:6235–6243.
- Baer AJ, Macartney DH. *Org Biomol Chem*. 2005;3:1448–1452.
- Schaschke N, Fiori S, Weyher E, et al. *J Am Chem Soc*. 1998;120:7030–7038.
- Liu Y, Chen G-S, Chen Y, Cao D-X, Ge Z-Q, Yuan Y-J. *Bioorg Med Chem*. 2004;12:5767–5775.
- Bom A, Bradley M, Cameron K, et al. *Angew Chem Int Ed*. 2002;41:265–270.
- Steffen A, Thiele C, Tietze S, et al. *Chem Eur J*. 2007;13:6801–6809.
- Wenz G, Strassnig C, Thiele C, Engelke A, Morgenstern B, Hegetschweiler K. *Chem Eur J*. 2008;14:7202–7211.
- Adam JM, Bennett DJ, Bom A, et al. *J Med Chem*. 2002;45:1806–1816.
- Zhang Y-M, Zhang X-J, Xu X, Fu X-N, Hou H-B, Liu Y. *J Phys Chem B*. 2016;120:3932–3940.
- Wang K, Guo D, Wang X, Liu Y. *ACS Nano*. 2011;5:2880–2894.
- Kang J, Kumar V, Yang D, Chowdhury PR, Hohl RJ. *Eur J Pharm Sci*. 2002;15:163–170.
- di Nunzio MR, Wang Y, Douhal A. *Photochem Photobiol A-Chem*. 2013;266:12–21.
- Swiech O, Mieczkowska A, Chmurski K, Bilewicz R. *J Phys Chem B*. 2012;116:1765–1771.