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A fluorogenic 2D glycosheet for the simultaneous identification of human- and avian-receptor specificity in influenza viruses†

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The switch in glycan-receptor specificity of influenza viruses may cause their interspecies transmission. However, tools that can simultaneously unveil the different receptor specificities of a single virus strain, in a homogeneous solution, have been elusive. Here we show a simple yet effective “2D glycosheet” that is capable of simultaneously identifying the human- and avian-glycan-receptor specificity of influenza viruses. Two fluorophores with different emission colors are coupled with avian- or human-glycan receptors. Then, the glycan-fluorophore conjugates with different receptors and emission colors are co-assembled on a 2D molybdenum disulfide platform, producing a duplexed, fluorogenic 2D glycosheet. While the system shows a single emission color with viruses containing a single receptor specificity, in the presence of a virus (H7N9) containing dual (human- and avian-) receptor specificity the system produces both emission colors. Interestingly, the use of graphene oxide and carbon nanotubes as the material substrate also produces dual emission with H7N9, thus enabling a new generation of low-dimensional glyco-materials for effectively probing the switch in receptor-specificity of influenza viruses.

Introduction

Influenza A viruses are segmented, negative-sense and single-stranded RNA viruses. They are classified into various subtypes according to the serological reactivity of their surface glycoproteins, hemagglutinin (HA) and neuraminidase (NA).¹ The HA protein is the key mediator of virus infection to the host

Conceptual insights

The possibility of an unpredictable reoccurrence of an influenza pandemic produces a tangible social anxiety and fear worldwide. While the influenza viruses mainly circulate in birds, mutations can cause the viruses to infect, and thus circulate in, humans. Hence, an effective surveillance technique for influenza is in urgent demand. Conventional techniques for the detection of influenza are both expensive and technically demanding requiring complicated manipulations. It is clear that a simple method to simultaneously identify both the bird- and human-infection risk of a single influenza virus strain is urgently required. We have developed a simple method using a 2D glycosheet material composite that is capable of simultaneously identifying both the human- and bird-infecting ability of an influenza virus that can circulate in humans (H7N9), using two different colored emissions. We believe, this research will lead to much needed early warning systems for the surveillance and evaluation of the influenza virus and therefore prevent future global pandemics.

cells with a receptor binding site (RBS) on its globular head, which determines the specificity of binding to α 2,3-sialic acid (SA) or α -2,6-SA receptors on host cells.² Generally, avian influenza viruses have a preference for α 2,3-SA and attach to cells of the lower respiratory tract, type II pneumocytes and non-ciliated cells. In contrast, influenza viruses infecting mammals have a preference for α 2,6-SA and attach to cells of the upper respiratory tract, primarily to ciliated cells.^{3,4} However, mutations and re-assortments within the RBS can switch the receptor-binding specificity of the viruses and therefore result in the interspecies transmission of influenza viruses.^{5–7} The traits of HA-receptor binding preference can be harnessed for rapid identification of influenza A virus.⁸ Although many elegant methods have been developed for the detection of influenza viruses,^{9–11} a tool that can concomitantly probe the avian- and human-receptor specificity of a single virus strain in a homogeneous solution has been elusive. Here we develop a 2D glycosheet by the co-assembly of two different fluorophore-glycan conjugates to a 2D material for the simultaneous identification of the dual-receptor specificity of a single influenza virus stain.

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The development of 2D materials has become a very topical research area in many industrial fields since these thin-layer sheets show a number of superior properties to their bulk counterparts.^{12–15} Of particular interest is the incorporation of 2D materials into the development of biomedical material composites.^{16–19} These materials have shown promise in sensing biomacromolecules, cell imaging and disease theranostics.^{20–29} Here we describe the use of a 2D molybdenum disulfide (2D MoS₂, a graphene analogue widely used in the design of biomaterials) for the simple construction of a 2D glycosheet that can simultaneously identify the human- and avian-receptor specificity in influenza viruses.

Results and discussion

It has been reported that the human- and avian-adapted HA of influenza viruses preferentially binds to the α 2,6- and α 2,3-sialyl glycan receptors, respectively.² The switch in the receptor specificity is a key factor for virus interspecies transmission.⁷ Therefore, we set out to conjugate two different fluorescence reporters to the regioisomeric glycans. A cyan-emitting coumarin (CMN) derivative and a red-emitting dicyanomethylene 4*H*-pyran (DCM) derivative were used to couple with an avian-receptor (Neu5Ac α 2,3Gal- β 1,4Glc) and a human-receptor (Neu5Ac α 2,6Gal- β 1,4Glc) (Fig. 1 and Scheme S1, ESI†), producing diverse glycan–fluorophore conjugates (CMN23, CMN26, DCM23 and DCM26) with different emission colors. We then took pairs of these probes (CMN23/DCM26 or CMN26/DCM23) and co-assembled them using van der Waals interactions³⁰ with a 2D MoS₂ platform to construct the fluorogenic 2D glycosheets (Fig. 2a). We predict that a single emission color would be produced for influenza viruses with a single glycan receptor specificity, whereas both emission colors should be produced in the presence of a virus strain with dual receptor specificity (Fig. 2b).

The four desired glycan–fluorophore conjugates were synthesized by coupling the sialyl glycans to amino-CMN and amino-DCM intermediates (Scheme S1, ESI†). 2D MoS₂ was prepared using a liquid exfoliation method, with the sheet-like morphologies confirmed by high-resolution transmission electron microscopy (Fig. S1, ESI†).³¹ With these materials in hand, we first tested the interaction between a single probe and 2D MoS₂ using fluorescence spectroscopy (Fig. S2, ESI†). We observed concentration-dependent fluorescence quenching with all four probes containing different

Fig. 1 Structures of the sialyl-glycan–fluorophore conjugates.

Fig. 2 Schematic illustration of (a) the co-assembly of sialyl-glycan–fluorophore conjugates with different emission colors to 2D MoS₂, producing the 2D glycosheet and (b) the use of the 2D glycosheet for the identification of the single or dual receptor specificity in influenza viruses.

sialyl glycans and fluorophores with increasing 2D MoS₂ (Fig. S2a, ESI†). This is in agreement with the known quenching ability of this 2D material for fluorescence species closely attached to its surface.²⁴ Interestingly, the subsequent addition of an influenza virus strain with 2,3- (*i.e.* H10N8: A/Hunan/3-9/2007) or 2,6-receptor (*i.e.* H1N1pdm09: A/California/08/2009) specificity to the probe-material composite resulted in a concentration dependent fluorescence recovery of the probes (Fig. S2b, ESI†). This fluorescence recovery was determined to be selective in terms of receptor-specificity (*i.e.* H10N8 for CMN23 and DCM23 and H1N1pdm09 for CMN26 and DCM26) (Fig. S3, ESI†). These preliminary results suggest that (1) 2D MoS₂ is a suitable quenching platform for assembling fluorophore-conjugated sialyl glycans and that (2) the fluorescence enhancement of the material composite formed is dependent on the receptor-specificity of the influenza viruses, irrespective of the coupled fluorophore.

Having obtained these promising results, we then set out to construct 2D glycosheets using the co-assembly of a pair of CNM/DCM probes with different glycan receptors with the 2D MoS₂.²⁰ We first determined that broad-band fluorescence was recorded for the aqueous mixture of both CMN26/DCM23 and CMN23/DCM26 at a single excitation wavelength (430 nm) (Fig. 3a). The use of a single excitation wavelength facilitates the detection procedure since the majority of previous multiplexed sensing systems require the interchange of multiple excitation wavelengths to produce the different emissions.³² Then, we determined that the entire broad-band fluorescence emission could be quenched, in a concentration-dependent manner, with increasing 2D MoS₂ (Fig. 3a), suggesting the powerful quenching ability of this 2D material for fluorophore mixtures.

To further corroborate the glycosheet assembly, a variety of other techniques including dynamic light scattering (DLS), Raman spectroscopy and zeta potential measurements were performed.

Fig. 3 (a) Fluorescence titration of the **CMN26/DCM23** (1/2 μ M) or **CMN23/DCM26** (1/2 μ M) mixture with increasing 2D MoS₂ (0–300 μ g mL⁻¹) in phosphate buffered saline (0.01 M, pH 7.4) (excitation wavelength: 430 nm). (b) Dynamic light scattering of 2D MoS₂, glycosheet **1** (**CMN26/DCM23**) and **2** (**CMN23/DCM26**). (c) Stacked Raman spectrum of 2D MoS₂, glycosheet **1** (**CMN26/DCM23**) and **2** (**CMN23/DCM26**). (d) Zeta potential of 2D MoS₂, glycosheet **1** (**CMN26/DCM23**) and **2** (**CMN23/DCM26**).

While the size distribution of the 2D MoS₂ was in the low sub-micrometer range,³¹ the glycosheets (**1**: **CMN23/DCM26** and **2**: **CMN26/DCM23**) seemed to be larger (Fig. 3b). This suggests the attachment of the glycan–fluorophore conjugates to the 2D MoS₂. We also observed typical Raman shifts for the 2D MoS₂ at 404 and 378 cm⁻¹, which could be ascribed to the A_{1g} (out-of-plane vibration of sulfur) and E_{12g} (in-plane relative motion between sulfur and molybdenum) modes of hexagonal MoS₂ (Fig. 3c).³³ However, the intensity ratio of E_{12g} to A_{1g} of the glycosheets (**1** and **2**) increased with respect to 2D MoS₂ alone, suggesting that the presence of the fluorophore–glycan conjugates perturbed the in-plane relative motion between the S and Mo atoms of the material.³⁴ In addition, the zeta potentials of the glycosheets were more negative than that of the 2D material alone (Fig. 3d). This is probably a result of the coating of the negatively charged sialyl glycans on the surface of the material.

Next, we tested the sensing performance of the duplexed 2D glycosheets for influenza viruses with different or mixed receptor specificities. The avian-specific H10N8, human specific H1N1pdm09 and an H7N9 “6+2” reassortant virus (A/Anhui/1/2013 derived HA and NA in the background of A/Puerto Rico/8/1934), with human-infecting properties (*i.e.* the HA can bind to 2,6-sialyl

Fig. 4 Normalized fluorescence change of (a) the **CMN26/DCM23** (1/2 μ M) 2D glycosheet (2D MoS₂: 70 μ g mL⁻¹) and (b) the **CMN23/DCM26** (1/2 μ M) 2D glycosheet (2D MoS₂: 70 μ g mL⁻¹) with increasing H10N8 (A/Hunan/3-9/2007), H1N1pdm09 (A/California/08/2009) and H7N9 (A/Anhui/1/2013) in phosphate buffered saline (0.01 M, pH 7.4) (excitation: 430 nm). For the original fluorescence spectra, see Fig. S4, ESI†.

glycans in addition to the avian specific 2,3-sialyl glycans)^{35–37} were used. A fluorescence titration assay indicated that the presence of the viruses with single-receptor-specificity only enhanced the fluorescence of the corresponding sialyl glycan (*i.e.* H10N8 for **CMN23** and **DCM23** and H1N1pdm09 for **CMN26** and **DCM26**) (Fig. 4 and Fig. S4, ESI†). This phenomenon was consistently observed for both **CMN23/DCM26** (Fig. 4a) and **CMN26/DCM23** 2D glycosheets (Fig. 4b). In contrast, the presence of H7N9 caused both the 2,3- and 2,6-glycans to enhance (Fig. 4). These results are in accordance with the previously reported receptor specificity of the virus strains.³⁶

In addition, we observed that the quenched cyan and red fluorescence of **CMN26** and **DCM23** could be recovered by H1N1pdm09 and H10N8, respectively, whereas the presence of H7N9 produced a mixed color of both glycan–fluorophore conjugates (Fig. 5a). In addition, confocal laser-scanning microscopy was carried out to corroborate the selective glycan–virus binding. Incubation of the 2D glycosheet with H1N1pdm09 and H10N8 resulted in the emergence of **CMN26** and **DCM23** fluorescence, respectively. In contrast, both fluorescence emissions were seen upon incubation of the material with H7N9 (Fig. 5b). These results together suggest the amenability of our 2D glycosheet for the simultaneous identification of human- and avian-glycan receptors in a single influenza virus strain.

Interestingly, we also demonstrated that the use of two other widely employed low-dimensional quenching platforms,³⁸ 2D graphene oxide (GO) and 1D carbon nanotubes (CNT), instead of the 2D MoS₂, as the substrate also produced a similar result (Fig. 6 and Fig. S5, ESI†). We first observed a concentration-dependent fluorescence quenching for the mixture of **CMN23** and **DCM26** in the presence of increasing GO (Fig. S5a) and CNT (Fig. S5e) (ESI†). Then, we observed that while a single fluorescence was selectively enhanced (Fig. S5b and c for GO and

Fig. 5 (a) Fluorescence emission color change of the **CMN26/DCM23** 2D glycosheet with H10N8, H1N1pdm09 and H7N9 in phosphate buffered saline (0.01 M, pH 7.4), as excited by a portable UV lamp (365 nm). The emission colors of **CMN26**, **DCM23** and **CMN26/DCM23** were used as a reference. (b) Confocal laser scanning microscopy of the **CMN26/DCM23** 2D glycosheet in the presence of different influenza virus strains (excitation: 425–475 nm; CMN emission channel: 425–475 nm; DCM emission channel: 570–620 nm).

Fig. S5f and g for CNT) (ESI[†]) in the presence of H10N8 and H1N1pdm09, both fluorescence emissions were observed in the presence of H7N9 (Fig. 6a and Fig. S5d (ESI[†]) for GO and Fig. 6b and Fig. S5h (ESI[†]) for CNT). This suggests that the system established tolerates a wide range of low-dimensional material substrates.

We performed a series of experiments in order to determine the stability of the 2D glycosheet. It was found that the **CMN23/DCM26** glycosheet was stable to increasing pH (Fig. S6a, ESI[†]) and increasing salt strength (Fig. S6b, ESI[†]). Also, the material did not show any fluorescence increase with a series of proteins including bovine serum albumin, IgG, lysozyme and pepsin (Fig. S6c, ESI[†]), suggesting that the detection system would not be adversely influenced by environmental factors. Furthermore, we used two additional viruses (the simian adenovirus AdC68 and the human adenovirus AdHu7) with the 2D glycosheet.³⁹ Our results indicate that the material was not responsive to these two adenoviruses (Fig. S7, ESI[†]).

The limit of detection (LOD) for the **CMN23/DCM26** glycosheet towards H1N1pdm09 and H10N8 was determined to be 0.04 and 0.02 HAU 50 μL^{-1} , respectively, and that for H7N9 with dual receptor specificity was determined to be 0.07 (**CMN23**) and 0.06 (**DCM26**) HAU 50 $\mu\text{g mL}^{-1}$ (Fig. S8, ESI[†]). We performed a fluorescence kinetic assay to determine how quickly the 2D glycosheet functions. The results indicated that a detection equilibrium was quickly established for both fluorescence emissions and within 5 min (Fig. S9, ESI[†]).

Fig. 6 Normalized fluorescence change of (a) the **CMN23/DCM26** (1/2 μM) 2D glycosheet (based on 2D graphene oxide: 40 $\mu\text{g mL}^{-1}$) and (b) the **CMN23/DCM26** (1/2 μM) 2D glycosheet (based on 1D carbon nanotube: 4.8 $\mu\text{g mL}^{-1}$) with increasing H10N8, H1N1pdm09 and H7N9 in phosphate buffered saline (0.01 M, pH 7.4) (excitation: 430 nm). For the original fluorescence spectra, see Fig. S5, ESI[†].

The most commonly used technique for influenza A virus detection is the hemagglutination assay, which is quasi-quantitative. This assay must be followed by an additional hemagglutinin inhibition assay including complicated manipulation procedures and long detection time. ELISA (enzyme-linked immunosorbent assay) is another frequently used method with good sensitivity. However, it requires the use of antibodies, which are expensive and sluggish to produce. Whereas the sensitivity of our 2D glycosheet is comparable to, or lower than, most of the competing methods developed including surface plasmon resonance,^{40a} HPLC,^{40b} gold nanoparticle based probes^{40c} and quartz crystal microbalance,^{40d} and the speed of detection appears to be the fastest (Table S1, ESI[†]). More importantly, our duplexed 2D glycosheet is a unique system to simultaneously determine the dual receptor specificity of a single influenza A virus in a homogeneous solution.

Conclusions

To conclude, we have developed a simple yet effective 2D glycosheet using supramolecular co-assembly of two fluorophore–glycan conjugates to a low-dimensional material platform. The fluoro-genic material composite formed has proven to be effective for the simultaneous identification of the dual-receptor specificity of a single influenza virus strain. This research allows for the effective interrogation of the receptor specificity switch for seasonal influenza viruses.

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Notes and references

- 1 N. J. Cox, G. Neumann, R. O. Donis and Y. Kawaoka, *Topley & Wilson's Microbiology and Microbial Infections*, John Wiley & Sons, Ltd, 2010.
- 2 Y. Suzuki, T. Ito, T. Suzuki, R. E. Holland, T. M. Chambers, M. Kiso, H. Ishida and Y. Kawaoka, *J. Virol.*, 2000, **74**, 11825–11831.
- 3 K. Shinya, M. Ebina, S. Yamada, M. Ono, N. Kasai and Y. Kawaoka, *Nature*, 2006, **440**, 435–436.
- 4 M. N. Matrosovich, T. Y. Matrosovich, T. Gray, N. A. Roberts and H. D. Klenk, *Proc. Natl. Acad. Sci. U. S. A.*, 2004, **101**, 4620–4624.
- 5 C. R. Parrish and Y. Kawaoka, *Annu. Rev. Microbiol.*, 2005, **59**, 553–586.
- 6 J. Stevens, O. Blixt, T. M. Tumpey, J. K. Taubenberger, J. C. Paulson and I. A. Wilson, *Science*, 2006, **312**, 404–410.
- 7 Y. Wang, *N. Engl. J. Med.*, 2013, **368**, 2348–2349.
- 8 A. García-Sastre and M. Schmolke, *Lancet*, 2014, **383**, 676–677.
- 9 S. Fukui, T. Feizi, C. Galustian, A. M. Lawson and W. Chai, *Nat. Biotechnol.*, 2002, **20**, 1011–1017.
- 10 T. Feizi and W. Chai, *Nat. Rev. Mol. Cell Biol.*, 2004, **5**, 582–588.
- 11 C. Lee, M. A. Gaston, A. A. Weiss and P. Zhang, *Biosens. Bioelectron.*, 2013, **42**, 236–241.
- 12 Q. H. Wang, K. Kalantar-Zadeh, A. Kis, J. N. Coleman and M. S. Strano, *Nat. Nanotechnol.*, 2012, **7**, 699–712.
- 13 M.-R. Gao, Y.-F. Xu, J. Jiang and S.-H. Yu, *Chem. Soc. Rev.*, 2013, **42**, 2986–3017.
- 14 H. Wang, H. Feng and J. Li, *Small*, 2014, **10**, 2165–2181.
- 15 Y. Sun, S. Gao, F. Lei, C. Xiao and Y. Xie, *Acc. Chem. Res.*, 2015, **48**, 3–12.
- 16 M. Pumera and A. H. Loo, *Trends Anal. Chem.*, 2014, **61**, 49–53.
- 17 Y. Chen, C. Tan, H. Zhang and L. Wang, *Chem. Soc. Rev.*, 2015, **44**, 2681–2701.
- 18 D. Chimene, D. L. Alge and A. K. Gaharwar, *Adv. Mater.*, 2015, **27**, 7261–7284.
- 19 (a) X.-P. He and H. Tian, *Small*, 2016, **12**, 144–160; (b) H.-L. Zhang, X.-L. Wei, Y. Zang, J.-Y. Cao, S. Liu, X.-P. He, Q. Chen, Y.-T. Long, J. Li, G.-R. Chen and K. Chen, *Adv. Mater.*, 2013, **25**, 4097–4101; (c) X.-P. He, Q. Deng, L. Cai, C.-Z. Wang, Y. Zang, J. Li, G.-R. Chen and H. Tian, *ACS Appl. Mater. Interfaces*, 2014, **6**, 5379–5382; (d) X. Sun, B. Zhu, D.-K. Ji, Q. Chen, X.-P. He, G.-R. Chen and T. D. James, *ACS Appl. Mater. Interfaces*, 2014, **6**, 10078–10082; (e) X.-P. He, B.-W. Zhu, Y. Zang, J. Li, G.-R. Chen, H. Tian and Y.-T. Long, *Chem. Sci.*, 2015, **6**, 1996–2001; (f) D.-K. Ji, Y. Zhang, X.-P. He and G.-R. Chen, *J. Mater. Chem. B*, 2015, **3**, 6656–6661; (g) D.-K. Ji, Y. Zhang, Y. Zang, W. Liu, X. Zhang, J. Li, G.-R. Chen, T. D. James and X.-P. He, *J. Mater. Chem. B*, 2015, **3**, 9182–9185; (h) D.-K. Ji, Y. Zhang, Y. Zang, J. Li, G.-R. Chen, X.-P. He and H. Tian, *Adv. Mater.*, 2016, **28**, 9356–9363; (i) D. Xie, D.-K. Ji, Y. Zhang, J. Cao, H. Zheng, L. Liu, Y. Zang, J. Li, G.-R. Chen, T. D. James and X.-P. He, *Chem. Commun.*, 2016, **52**, 9418–9421; (j) M. Wahiba, X.-Q. Feng, Y. Zang, T. D. James, J. Li, G.-R. Chen and X.-P. He, *Chem. Commun.*, 2016, **52**, 11689–11692; (k) S. Guo, J. Chen, B.-Y. Cai, W.-W. Chen, Y.-F. Li, X. Sun, G.-R. Chen, X.-P. He and T. D. James, *Mater. Chem. Front.*, 2017, **1**, 61–64; (l) Y.-H. Ma, W.-T. Dou, Y.-F. Pan, L.-W. Dong, Y.-X. Tan, X.-P. He, H. Tian and H.-Y. Wang, *Adv. Mater.*, 2017, **29**, 1604253.
- 20 Y. Zhang, B. Zheng, C. Zhu, X. Zhang, C. Tan, H. Li, B. Chen, J. Yang, J. Chen, Y. Huang, L. Wang and H. Zhang, *Adv. Mater.*, 2015, **27**, 935–939.
- 21 L. Cheng, J. Liu, X. Gu, H. Gong, X. Shi, T. Liu, C. Wang, X. Wang, G. Liu, H. Xing, W. Bu, B. Sun and Z. Liu, *Adv. Mater.*, 2014, **26**, 1886–1893.
- 22 T. Liu, C. Wang, X. Gu, H. Gong, L. Cheng, X. Shi, L. Feng, B. Sun and Z. Liu, *Adv. Mater.*, 2014, **26**, 3433–3440.
- 23 W. Yin, L. Yan, J. Yu, G. Tian, L. Zhou, X. Zheng, X. Zhang, Y. Yong, J. Li, Z. Gu and Y. Zhao, *ACS Nano*, 2014, **8**, 6922–6933.
- 24 C. Zhu, Z. Zeng, H. Li, F. Li, C. Fan and H. Zhang, *J. Am. Chem. Soc.*, 2013, **135**, 5998–6001.
- 25 B. L. Li, H. L. Zou, L. Lu, Y. Yang, J. L. Lei, H. Q. Luo and N. B. Li, *Adv. Funct. Mater.*, 2015, **25**, 3541–3550.
- 26 H. Fan, Z. Zhao, G. Yan, X. Zhang, C. Yang, H. Meng, Z. Chen, H. Liu and W. Tan, *Angew. Chem., Int. Ed.*, 2015, **54**, 4801–4805.
- 27 S. S. Chou, B. Kaehr, J. Kim, B. M. Foley, M. De, P. E. Hopkins, J. Huang, C. J. Brinker and V. P. Dravid, *Angew. Chem., Int. Ed.*, 2013, **52**, 4160–4164.
- 28 L. Cheng, C. Yuan, S. Shen, X. Yi, H. Gong, K. Yang and Z. Liu, *ACS Nano*, 2015, **9**, 11090–11101.
- 29 J. Nam, W.-G. La, S. Hwang, Y. S. Ha, N. Park, N. Won, S. Jung, S. H. Bhang, Y.-J. Ma, Y.-M. Cho, M. Jin, J. Han, J.-Y. Shin, E. K. Wang, S. G. Kim, S.-H. Cho, J. Yoo, B.-S. Kim and S. Kim, *ACS Nano*, 2013, **7**, 3388–3402.
- 30 Y. Ling, Z. Gu, S.-G. Kang, J. Luo and R. Zhou, *J. Phys. Chem. C*, 2016, **120**, 6796–6803.
- 31 J. N. Coleman, M. Lotya, A. O'Neill, S. D. Bergin, P. J. King, K. Young, A. Gaucher, S. De, R. J. Smith, I. V. Shvets, S. K. Arora, G. Stanton, H.-Y. Kim, K. Lee, G. T. Kim, G. S. Duesberg, T. Hallam, J. J. Boland, J. J. Wang, J. F. Donegan, J. C. Grunlan, G. Moriarty, A. Shmeliov, R. J. Nicholls, J. M. Perkins, E. M. Grievson, K. Theuwissen, D. W. McComb, P. D. Nellist and V. Nicolosi, *Science*, 2011, **311**, 568–571.
- 32 D.-K. Ji, G.-R. Chen, X.-P. He and H. Tian, *Adv. Funct. Mater.*, 2015, **25**, 3483–3487.
- 33 C. Lee, H. Yan, L. E. Brus, T. F. Heinz, J. Hone and S. Ryu, *ACS Nano*, 2010, **4**, 2695–2700.

- 34 P. T. K. Loan, W. Zhang, C.-T. Lin, K.-H. Wei, L.-J. Li and C.-H. Chen, *Adv. Mater.*, 2014, **26**, 4838–4844.
- 35 Y. Shi, W. Zhang, F. Wang, J. Qi, Y. Wu, H. Song, F. Gao, Y. Bi, Y. Zhang, Z. Fan, C. Qin, H. Sun, J. Liu, J. Haywood, W. Liu, W. Gong, D. Wang, Y. Shu, Y. Wang, J. Yan and G. F. Gao, *Science*, 2013, **342**, 243–247.
- 36 (a) X. Xiong, S. R. Martin, L. F. Haire, S. A. Wharton, R. S. Daniels, M. S. Bennett, J. W. McCauley, P. J. Collins, P. A. Walker, J. J. Skehel and S. J. Gamblin, *Nature*, 2013, **499**, 496–499; (b) M. Wang, W. Zhang, J. Qi, F. Wang, J. Zhou, Y. Bi, Y. Wu, H. Sun, J. Liu, C. Huang, X. Li, J. Yan, Y. Shu, Y. Shi and G. F. Gao, *Nat. Commun.*, 2015, **6**, 5600; (c) T. R. Maines, A. Jayaraman, J. A. Belser, D. A. Wadford, C. Pappas, H. Zeng, K. M. Gustin, M. B. Pearce, K. Viswanathan, Z. H. Shriver, R. Raman, N. J. Cox, R. Sasisekharan, J. M. Katz and T. M. Tumpey, *Science*, 2009, **325**, 484–487.
- 37 S. G. Vachieri, X. Xiong, P. J. Collins, P. A. Walker, S. R. Martin, L. F. Haire, Y. Zhang, J. W. McCauley, S. J. Gamblin and J. J. Skehel, *Nature*, 2014, **511**, 475–477.
- 38 (a) X.-P. He, Y. Zang, T. D. James, J. Li and G.-R. Chen, *Chem. Soc. Rev.*, 2015, **44**, 4239–4248; (b) X.-P. He, Y. Zang, T. D. James, J. Li, G.-R. Chen and J. Xie, *Chem. Commun.*, 2017, **53**, 82–90.
- 39 D. Zhou, X. Zhou, A. Bian, H. Li, H. Chen, J. C. Small, Y. Li, W. Giles-Davis, Z. Xiang and H. C. J. Ertl, *Nat. Protoc.*, 2010, **5**, 1775–1785.
- 40 (a) H. Bai, R. Wang, B. Hargis, H. Lu and Y. Li, *Sensors*, 2012, **12**, 12506–12518; (b) L. Urbas, B. Košir, M. Peterka, B. Pihlar, A. Štrancar and M. Barut, *J. Chromatogr. A*, 2011, **1218**, 2432–2437; (c) Y. Liu, L. Zhang, W. Wei, H. Zhao, Z. Zhou, Y. Zhang and S. Liu, *Analyst*, 2015, **140**, 3989–3995; (d) R. Wang and Y. Li, *Biosens. Bioelectron.*, 2013, **42**, 148–155.