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Isoindoline-based fluorogenic probes bearing a self-immolative linker for the sensitive and selective detection of O-GlcNAcase activity†

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O-GlcNAcase (OGA) is implicated in several important biological and disease-relevant processes. Here, we synthesized fluorogenic probes for OGA by grafting GlcNAc directly or using a self-immolative linker to the hydroxyl position of 4-hydroxyisoindoline (BHID), a typical excited-state intramolecular proton transfer (ESIPT) probe. The probe was used for a fluorogenic assay to determine the half maximal inhibitory concentration of a known OGA inhibitor and differentiate between OGA and hexosaminidase when GlcNAc is replaced by GlcNPr, where a propionyl group is used instead of an acetyl group.

O-GlcNAcylation is a typical post-translational modification implicated in a diverse range of biological processes by introducing *N*-acetylglucosamine (GlcNAc) to the serine or threonine residue of proteins. Two key enzymes are involved in this modification process O-GlcNAc transferase (OGT) that transfers GlcNAc to biomolecules, and O-GlcNAcase (OGA) that deglycosylates GlcNAc reversibly. Both enzymes have been shown to be associated with a variety of biological processes, and abnormal OGA expression was found to be linked with a number of diseases including diabetes, cancer, Alzheimer's disease and

cardiovascular disease.^{1,2} This highlights the importance of developing effective sensing tools for OGA.

In addition to traditional biochemical techniques such as quantitative real-time polymerase chain reaction and western blot, activity-based assays are commonly used for the analysis of enzymatic activities. While colorimetric assays are generally used for enzymatic assays, fluorogenic probes have been extensively developed to visualize the activity of an enzyme of interest in live cells.^{3,4} A general strategy to design such probes relies on the introduction of the substrate molecule of a specific enzyme to a fluorescent dye, inhibiting the electron or charge transfer process of the dye system, thus quenching the fluorescence.⁵ Then after selective removal of the substrate by the enzyme, the fluorescence is restored. With this rationale, fluorogenic probes for many diverse glycosidases have been developed for the intracellular and *in vivo* imaging of their activities.^{6,7} Glycosidase-activatable prodrugs have also been developed for therapy of cancer and senescence.^{8,9}

Several fluorogenic probes based on the sensing mechanisms described above have been reported for OGA detection.^{10–15} However, it is still necessary to expand the range of designs that can be incorporated into molecular probes for

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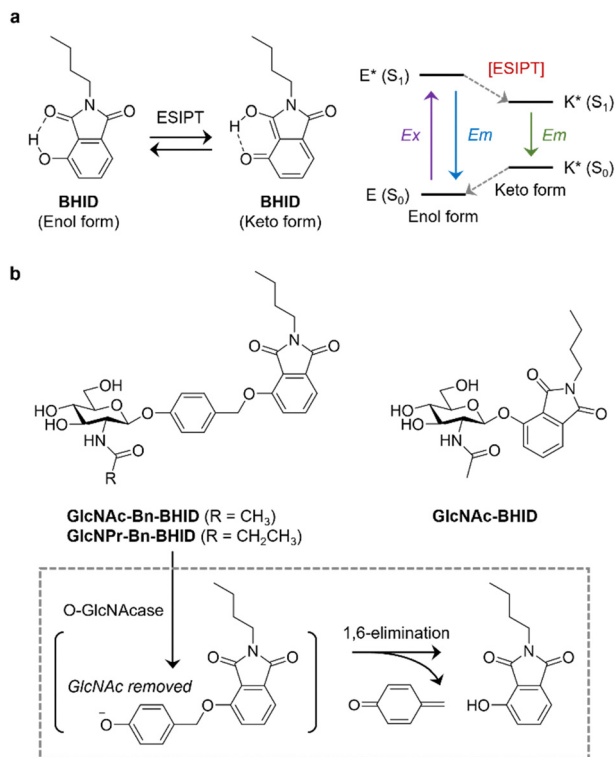


Fig. 1 (a) Switching between Enol to Keto form of **BHID** via ES IPT and a simplified scheme highlighting the ES IPT process. (b) Structure of the fluorogenic probes developed in this study for OGA detection, and a scheme illustrating the sensing mechanism.

OGA. Excited-state intramolecular proton transfer (ES IPT) is a photochemical process that describes the switch of a molecule from the enol to keto form upon light excitation, causing its fluorescence emission features to change. Fluorescent probes based on the ES IPT mechanism are advantageous in terms of large Stokes shift and flexibility in fine-tuning the ES IPT states by simply blocking the proton donor site.¹⁶ A survey of the literature indicates the importance and popularity of using fluorescent ES IPT dyes for biosensing and bioimaging.¹⁷

Here, we develop ES IPT-based OGA probes, and demonstrate the importance of incorporating a self-immolative linker to improve sensitivity (Fig. 1). We chose 2-butyl-4-hydroxyisoindoline-1,3-dione (**BHID**), which has been used to develop fluorogenic probes for detecting hydrazine and peroxyxynitrite,^{18,19} as the fluorophore because of its accessibility and simplicity for structural modification. As such, to the hydroxyl position of **BHID** GlcNAc was added with or without a “self-immolative” spacer.²⁰ It has been reported that the presence of the spacer enhances the sensitivity of fluorogenic probes for glycosidases probably because of a reduced steric hindrance.²¹ GlcNAc was introduced to the hydroxyl group of *p*-hydroxybenzaldehyde by stirring in acetonitrile with silver oxide and potassium iodide, followed by reduction and bromination to produce intermediate **GlcNAc-Bn-Br**. Then, this intermediate underwent Williamson ether reaction with **BHID**, followed by deacetylation to afford **GlcNAc-Bn-BHID** in moderate yield (Scheme S1, ES I⁺). A control probe **GlcNAc-BHID** without the spacer was also prepared directly

introducing GlcNAc to the phenolic OH of **BHID** followed by deprotection (Scheme S1, ES I⁺).

It is known that in addition to OGA, other endogenous enzymes such as *N*-acetyl- β -D-hexosaminidases A (HexA) can also unselectively remove GlcNAc from proteins. A previous report indicated that the replacement of the *N*-acetyl group with an *N*-propionyl group in the GlcNAc structure achieved selective binding to OGA over HexA due to better ligand-binding.^{22–24} Therefore, a GlcNPr (in which a propionyl was used to modify the 2-amino group of the sugar instead of acetyl group) donor was used to glycosylate the spacer, followed by coupling to **BHID** by the same protocol as described above. The resulting **GlcNPr-Bn-BHID** probe was used to achieve selectivity sensing of OGA over a commercial Hex.

With the probes in hand, we determined their sensing properties. The ES IPT properties of **BHID** were determined in different solvents by fluorescence spectroscopy (Fig. 2a). We found that it exhibited two emission peaks with different Stokes shift. In aprotic polar solvent (acetonitrile) a short emission was observed, while a longer emission was seen in protic solvent (deionized water and PBS). This dual-emission reflected the enol and keto form in the ES IPT process, confirming the ES IPT mechanism of our fluorophore.

Then, we expressed OGA by BL21(DE3) Competent *Escherichia coli* for enzymatic analysis. To establish the activity-based assay, a fluorescence microplate reader was used. **GlcNAc-BHID** and **GlcNAc-Bn-BHID** were incubated with OGA in an optimized buffer system of NaCl–NaH₂PO₄–BSA

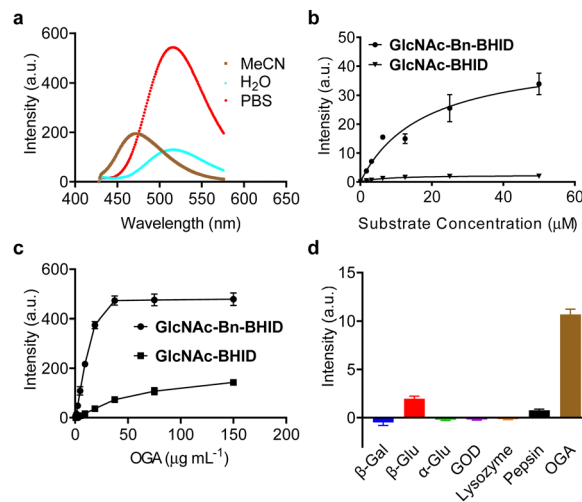


Fig. 2 (a) Stacked emission spectra of **BHID** (50 μ M) in different indicated solvents (λ_{ex} = 405 nm). (b) Fluorescence enhancement of **GlcNAc-Bn-BHID** (50 μ M) and **GlcNAc-BHID** (50 μ M) at different substrate concentrations in the presence of OGA (1 μ g mL⁻¹) measured in buffer (100 mM NaCl, 0.1% bovine serum albumin (BSA) and 50 mM NaH₂PO₄, pH 7.0) with an excitation of 405 nm. (c) Fluorescence enhancement of **GlcNAc-Bn-BHID** (50 μ M) and **GlcNAc-BHID** (50 μ M) with increasing concentrations of OGA measured in buffer (100 mM NaCl, 0.1% bovine serum albumin (BSA) and 50 mM NaH₂PO₄, pH 7.0) with an excitation of 405 nm. (d) Fluorescence change of **GlcNAc-Bn-BHID** (50 μ M) in the presence of different enzymes measured in phosphate buffered saline (PBS, 0.01 M, pH 7.4) with an excitation of 405 nm.



(bovine serum albumin) at pH 7.0.^{22,25} We first evaluated the fluorescence response of the two probes with increasing substrate concentrations at an OGA concentration of $1 \mu\text{g mL}^{-1}$ (Fig. 2b). A substrate concentration of $50 \mu\text{M}$ was found to be required for the fluorescence to reach equilibrium. Over the entire substrate concentration range, the fluorescence of **GlcNAc-Bn-BHID** was constantly more intense than that of **GlcNAc-BHID**, suggesting the former being more sensitive. Indeed, a ~ 3 -fold larger K_m (Michaelis constant) was determined for **GlcNAc-Bn-BHID** than **GlcNAc-BHID** by using a nonlinear curve fitting method in GraphPad Prism 9.5.0 (Table. S1, ESI[†]), indicating a larger catalytic efficiency of **GlcNAc-Bn-BHID**. Then, with a fixed substrate concentration ($50 \mu\text{M}$), increasing concentrations of OGA was added to the system. We determined that equilibrium was reached for **GlcNAc-Bn-BHID** with $37.5 \mu\text{g mL}^{-1}$ OGA, and a larger OGA concentration was required to fully activate the fluorescence of **GlcNAc-BHID** (Fig. 2c). These results corroborate the importance of the spacer for improving OGA sensitivity. Measurement using fluorescence spectroscopy also confirmed the fluorescence “turn-on” response of **GlcNPr-Bn-BHID** in the presence of OGA (Fig. S1, ESI[†]).

Next, the selectivity of **GlcNAc-Bn-BHID** for OGA was determined. A range of other glycosidases including β -galactosidase (β -Gal), α -glucosidase (α -Glu) and β -glucosidase (β -Glu), and other physiological enzymes including lysozyme, pepsin and glucose oxidase (GOD) were used. All enzymes were separately incubated with the probe in microplate wells for 90 min, followed by fluorescence detection in the microplate reader. As shown in Fig. 2d, the results indicated that the unselective enzymes hardly enhanced the fluorescence of **GlcNAc-Bn-BHID**. A slightly enhanced fluorescence was observed for β -glucosidase, which might be the result of structural similarity between glucose and GlcNAc.

To prove its real-world applicability, we sought to exploit the probe for evaluating the inhibitory activity of OGA inhibitors. A commercial OGA probe, **4-MUF-NAG**, was used as reference. In addition, a known OGA inhibitor, **Thiamet-G**, was used for the experiments.²⁶ Lineweaver–Burk plots were obtained for **GlcNAc-Bn-BHID** (Fig. 3a) and **4-MUF-NAG** (Fig. 3b) with different inhibitor concentrations (0.17 nM to $10 \mu\text{M}$) in the presence of OGA (Table. S2, ESI[†]). The fluorescence of both probes decreased gradually when the concentration of inhibitor increased. The half maximal inhibitory concentration (IC_{50}) of **Thiamet-G** was determined to be 0.8 nM and 1.7 nM for the **GlcNAc-Bn-BHID** and **4-MUF-NAG**-based assay, respectively. The IC_{50} determined from the **GlcNAc-Bn-BHID**-based assay agrees with that reported previously,²⁶ suggesting the reliability of our ESIP-based probe for screening OGA inhibitors.

Finally, the probes were used to differentiate OGA and HexA. It is important to achieve selective OGA detection over other enzymes to better elaborate the biological functions. In particular, HexA unselectively hydrolyzes GlcNAc and *N*-acetylgalactosamine (GalNAc) from biomacromolecules in the lysosome.²⁷ A previous report indicated that *N*-propionyl substitution significantly enhances OGA binding of *N*-glucosamines over HexA.

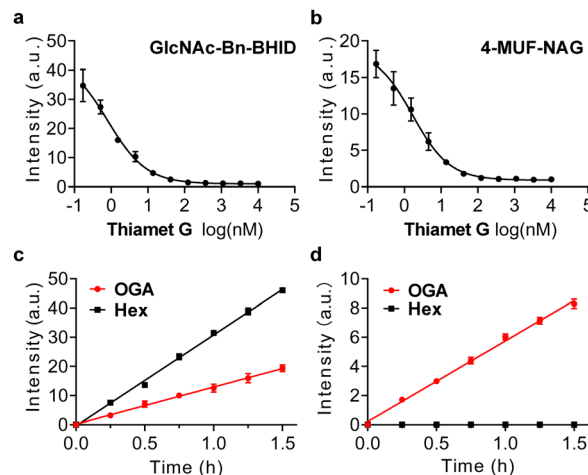


Fig. 3 Plotting the fluorescence intensity change of (a) **4-MUF-NAG** ($50 \mu\text{M}$) and (b) **GlcNAc-Bn-BHID** ($50 \mu\text{M}$) in the presence of increasing concentrations of **Thiamet G**, a known OGA inhibitor. Time-dependent fluorescence intensity change of (c) **GlcNAc-Bn-BHID** ($50 \mu\text{M}$) and (d) **GlcNPr-Bn-BHID** ($50 \mu\text{M}$) in the presence of commercial hexosaminidase ($0.03 \text{ U } \mu\text{g}^{-1}$) and OGA ($1 \mu\text{g mL}^{-1}$). All measurements were done in buffer with an excitation wavelength of 405 nm .

Thus, we determined whether our **GlcNAc-Bn-BHID** and **GlcNPr-Bn-BHID** could be used to differentiate the enzymes in a fluorogenic assay.

We found that in the presence of both OGA and a commercial hexosaminidase (from *Canavalia ensiformis*) used as a homogenous surrogate for human HexA,²⁸ the fluorescence of **GlcNAc-Bn-BHID** enhanced in a concentration-dependent manner (Fig. 3c). However, when **GlcNPr-Bn-BHID** was used as the substrate, a time-dependent fluorescence enhancement was only observed for OGA, and the presence of commercial hexosaminidase did not cause the probe to fluoresce with time (Fig. 3d). The better sensitivity of **GlcNAc-Bn-BHID** than **GlcNPr-Bn-BHID** was probably a result of the steric effects of the propionyl group that partially compromises glucosamine binding. However, the observation that **GlcNPr-Bn-BHID** does not respond to commercial hexosaminidase suggests its potential to be used for OGA-selective biosensing and bioimaging in complicated biological samples where both enzymes co-exist. As such, our ongoing research will focus on developing fluorogenic OGA probes suitable for cell imaging.^{4,29–31}

In summary, we have developed the first ESIP-based fluorogenic probe for OGA, which is an important glycosidase implicated in a variety of disease-relevant biological processes. The presence of a self-immolative linker was shown to significantly improve the sensitivity of the probe. We also established an enzymatic assay that exhibited time- and concentration-dependent fluorescence recovery in the presence of recombinant OGA. We also determined the IC_{50} of a known OGA inhibitor and achieved the differentiation between OGA and commercial hexosaminidase. We believe that the chemical probe developed as part of this research can provide useful strategies for the study of OGA-based biological events and OGA-based inhibitor screening.



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Data availability

All data supporting this research are included in the main article and/or ESI.†

Conflicts of interest

The authors declare no conflict of interest.

Notes and references

- 1 Y. Li, M. Xie, L. Men and J. Du, *Int. J. Mol. Med.*, 2019, **44**, 363–374.
- 2 J. Ma, C. Hou and C. Wu, *Chem. Rev.*, 2022, **122**, 15822–15864.
- 3 J. J. Zhang, X. Z. Chai, X. P. He, H. J. Kim, J. Y. Yoon and H. Tian, *Chem. Soc. Rev.*, 2019, **48**, 683–722.
- 4 W. T. Dou, H. H. Han, A. C. Sedgwick, G. B. Zhu, Y. Zang, X. R. Yang, J. Yoon, T. D. James, J. Li and X. P. He, *Sci. Bull.*, 2022, **67**, 853–878.
- 5 Y. Kim, H. Li, J. Choi, J. Boo, H. Jo, J. Y. Hyun and I. Shin, *Chem. Soc. Rev.*, 2023, **52**, 7036–7070.
- 6 X. Chai, H. H. Han, A. C. Sedgwick, N. Li, Y. Zang, T. D. James, J. Zhang, X. L. Hu, Y. Yu, Y. Li, Y. Wang, J. Li, X. P. He and H. Tian, *J. Am. Chem. Soc.*, 2020, **142**, 18005–18013.
- 7 H. Tian, W. Lin, X. L. Hu, J. B. Wang, M. Y. Zhang, Y. Zang, X. Y. Wu, J. Li, T. D. James and X. P. He, *Org. Chem. Front.*, 2023, **10**, 2913–2917.
- 8 D. L. Shi, W. W. Liu, Y. Gao, X. M. Li, Y. Y. Huang, X. K. Li, T. D. James, Y. Guo and J. Li, *Nat. Aging.*, 2023, **3**, 297–312.
- 9 M. Chang, Y. Dong, H. Xu, A. B. Cruickshank-Taylor, J. S. Kozora, B. Behpour and W. Wang, *Angew. Chem., Int. Ed.*, 2024, **63**, e202315425.
- 10 J. Y. Hyun, S. H. Park, C. W. Park, H. B. Kim, J. W. Cho and I. Shin, *Org. Lett.*, 2019, **21**, 4439–4442.
- 11 H. Jung, S. H. Park, W. H. Yang, J. W. Cho and I. Shin, *Sens. Actuators, B*, 2022, **367**, 132093.
- 12 J. Boo, J. Lee, Y. H. Kim, C. H. Lee, B. Ku and I. Shin, *Front. Chem.*, 2023, **11**, 1133018.
- 13 S. Cecioni and D. J. Vocadlo, *J. Am. Chem. Soc.*, 2017, **139**, 8392–8395.
- 14 E. J. Kim, D. O. Kang, D. C. Love and J. A. Hanover, *Carbohydr. Res.*, 2006, **341**, 971–982.
- 15 V. S. Borodkin, K. Raffie, N. Selvan, T. Aristotelous, I. Navratilova, A. T. Ferenbach and D. M. F. Van Aalten, *ACS Chem. Biol.*, 2018, **13**, 1353–1360.
- 16 A. C. Sedgwick, L. Wu, H. H. Han, S. D. Bull, X. P. He, T. D. James, J. L. Sessler, B. Z. Tang, H. Tian and J. Yoon, *Chem. Soc. Rev.*, 2018, **47**, 8842–8880.
- 17 H. Gu, W. J. Wang, W. Y. Wu, M. L. Wang, Y. R. Liu, Y. J. Jiao, F. Wang, F. Wang and X. Q. Chen, *Chem. Commun.*, 2023, **59**, 2056–2071.
- 18 S. Hu, J. Wang, M. Luo, Z. Wu, Y. Hou and X. Chen, *Anal. Bioanal. Chem.*, 2021, **413**, 5463–5468.
- 19 L. Wu, X. Tian, D. J. Lee, J. Yoon, C. S. Lim, H. M. Kim and T. D. James, *Chem. Commun.*, 2021, **57**, 11084–11087.
- 20 S. Gnaïm and D. Shabat, *Acc. Chem. Res.*, 2014, **47**, 2970–2984.
- 21 Q. Wu, Q. H. Zhou, W. Li, T. B. Ren, X. B. Zhang and L. Yuan, *ACS Sens.*, 2022, **7**, 3829–3837.
- 22 M. S. Macauley, G. E. Whitworth, A. W. Debowski, D. Chin and D. J. Vocadlo, *J. Biol. Chem.*, 2005, **280**, 25313–25322.
- 23 M. González-Cuesta, P. Sidhu, R. A. Ashmus, A. Males, C. Proceviat, Z. Madden, J. C. Rogalski, J. A. Busmann, L. J. Foster, J. M. Garcia Fernández, G. J. Davies, C. Ortiz Mellet and D. J. Vocadlo, *J. Am. Chem. Soc.*, 2022, **144**, 832–844.
- 24 B. Li, H. Li, C. W. Hu and J. Jiang, *Nat. Commun.*, 2017, **8**, 666.
- 25 Y. Gao, L. Wells, F. I. Comer, G. J. Parker and G. W. Hart, *J. Biol. Chem.*, 2001, **276**, 9838–9845.
- 26 S. A. Yuzwa, M. S. Macauley, J. E. Heinonen, X. Shan, R. J. Dennis, Y. He, G. E. Whitworth, K. A. Stubbs, E. J. McEachern, G. J. Davies and D. J. Vocadlo, *Nat. Chem. Biol.*, 2008, **4**, 483–490.
- 27 J. J. Morsby and B. D. Smith, *Bioconjugate Chem.*, 2022, **33**, 544–554.
- 28 Q. Chen, P. Guo, L. Liu, T. Liu, X. H. Qian and Q. Yang, *Biochimie*, 2014, **97**, 152–162.
- 29 X.-P. He and H. Tian, *Chem*, 2018, **4**, 246–268.
- 30 L. Dong, M.-Y. Zhang, H.-H. Han, Y. Zang, G.-R. Chen, J. Li, X.-P. He and S. Vidal, *Chem. Sci.*, 2022, **13**, 247–256.
- 31 J.-B. Jiao, G.-Z. Wang, X.-L. Hu, Y. Zang, S. Maisonneuve, A. C. Sedgwick, J. L. Sessler, J. Xie, J. Li, X.-P. He and H. Tian, *J. Am. Chem. Soc.*, 2020, **142**, 1925–1932.

