

## Synthesis, optimization, and evaluation of glycosylated naphthalimide derivatives as efficient and selective insect #-N-acetylhexosaminidase OfHex1 inhibitors

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1     **Synthesis, optimization, and evaluation of glycosylated naphthalimide**  
2     **derivatives as efficient and selective insect  $\beta$ -N-acetylhexosaminidase**  
3             **OfHex1 inhibitors**

4  
5     Shengqiang Shen,<sup>†,#</sup> Lili Dong,<sup>†,#</sup> Wei Chen,<sup>‡</sup> Renjie Wu,<sup>†</sup> Huizhe Lu,<sup>†</sup> Qing Yang,<sup>\*,‡</sup> and Jianjun Zhang<sup>\*,†</sup>

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8  
9     **ABSTRACT:** Insect chitinolytic  $\beta$ -N-acetylhexosaminidase OfHex1, from the agricultural pest *Ostrinia*  
10     *furnacalis* (Guenée), is considered as a potential target for green pesticide design. In this study, rational  
11     molecular design and optimization led to the synthesis of compounds **15r** ( $K_i=5.3 \mu\text{M}$ ) and **15y** ( $K_i=2.7 \mu\text{M}$ )  
12     that had superior activity against OfHex1 than previously reported lead compounds. Both **15r** and **15y** had  
13     high selectivity towards OfHex1 over HsHexB (human  $\beta$ -N-acetylhexosaminidase B) and hOGA (human  
14     O-GlcNAcase). In addition, to investigate the basis for the potency of glycosylated naphthalimides against  
15     OfHex1, molecular docking and MD simulations were performed to study possible binding modes.  
16     Furthermore, the *in vivo* biological activity of target compounds with efficient OfHex1 inhibitory potency were  
17     assayed against *Myzus persicae*, *Plutella xylostella*, and *Ostrinia furnacalis*. This present work indicates that  
18     glycosylated naphthalimides can be further developed as potential pest control and management agents  
19     targeting OfHex1.

20     **KEYWORDS:** Glycosylated naphthalimids,  $\beta$ -N-acetylhexosaminidase, OfHex1, inhibitors, molecular

21 docking, MD simulations

## 22 INTRODUCTION

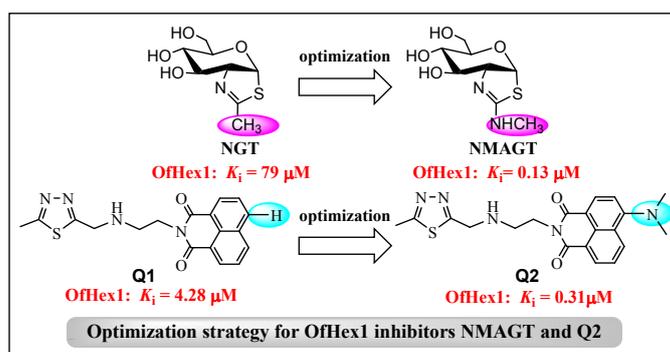
23 Chitin is a homopolymer of  $\beta$ -1,4-linked *N*-acetyl- $\beta$ -D-glucosamine (GlcNAc), and is the second most  
24 abundant polysaccharide in nature.<sup>1</sup> Chitin is also a key structural component of the fungal cell wall, nematode  
25 eggshell, and arthropod exoskeleton.<sup>2</sup> Moreover, the cuticles of the integument and peritrophic membranes of  
26 the midgut have found to contain chitin.<sup>3</sup> Chitin degradation is catalyzed by two members of the glycoside  
27 hydrolase family, GH18 chitinases (EC 3.2.1.14) and GH20  $\beta$ -*N*-acetylhexosaminidases (EC 3.2.1.52).<sup>4,5</sup>  
28 GH18 chitinases catalyze the cleavage of chitin into shorter chito-oligosaccharides, then GH20  
29  $\beta$ -*N*-acetylhexosaminidases hydrolyze these oligosaccharides into *N*-acetyl-D-glucosamine (GlcNAc) from  
30 terminal non-reducing.<sup>4,5</sup> Chitin degradation is essential to the growth and maturation of insects.<sup>6,7</sup> Interference  
31 with insect chitin degradation disrupts molting, pupation and eclosion processes, eventually leading to insect  
32 death.<sup>6,8</sup> Chitin is completely absent from higher plants and mammals,<sup>5</sup> and thus the inhibition of chitin  
33 degradation is a promising strategy for the development of green pesticides.<sup>5-10</sup>

34 GH20  $\beta$ -*N*-acetylhexosaminidase OfHex1 from the destructive agricultural pest, *Ostrinia furnacalis*  
35 (Guenée), is an important enzyme during insect chitin degradation.<sup>11</sup> The physiological function of OfHex1  
36 shown via RNA interference is to degrade of old cuticles of *O. furnacalis*.<sup>12</sup> Furthermore, the crystal structure  
37 of OfHex1 (PDB: 3NSM) revealed the unique structural feature, with long substrate-binding pocket containing  
38 two subsites (-1 and +1).<sup>11</sup> The -1 subsite is responsible for catalysis substrates, whilst the +1 subsite is  
39 responsible for binding related moieties of substrates to enhance affinity and specificity.<sup>11,12</sup> The  
40 crystallography of OfHex1 has attracted intense research attention and provides a solid foundation for the  
41 design of specific inhibitors.<sup>10,13-15</sup>

42 To date, a number of OfHex1 inhibitors have been reported, including TMG-chitotriomycin<sup>12</sup>, PUGNAc<sup>9</sup>,  
43 NAG-thiazoline (**NGT**)<sup>13</sup>, naphthalimides<sup>14-15</sup>, phlegmacin B1<sup>7</sup>, berberine<sup>16</sup>, thiazolyldrazones<sup>10</sup>. Among  
44 these inhibitors, NAG-thiazoline is the only molecule designed as an analog of the oxazolinium reaction  
45 intermediate, but shows poor potency against OfHex1 with  $K_i$  values of 79  $\mu\text{M}$ .<sup>13</sup> Subsequently, **NMAGT**  
46 bearing a methylamino substituent on the thiazoline ring was designed and synthesized by Yang, and exhibited  
47 higher inhibitory activity ( $K_i$  value of 0.13  $\mu\text{M}$ ) against OfHex1.<sup>13</sup> This suggested that the enlargement of the  
48 functional group on the thiazoline could improve inhibitory activity against OfHex1. Naphthalimide  
49 derivatives, including **Q1** and **Q2**, are an important class of non-carbohydrate inhibitors of OfHex1.<sup>14</sup> In  
50 particular, **Q1** has been shown to exhibit inhibitory activity towards OfHex1 with a  $K_i$  value of 4.28  $\mu\text{M}$ .<sup>14</sup> **Q2**  
51 containing a dimethylamino group at the naphthalimide exhibited a  $K_i$  value of 0.31  $\mu\text{M}$  against OfHex1.<sup>14</sup> The  
52 complex crystal structures of **Q1**-OfHex1 (PDB: 3WMB) and **Q2**-OfHex1 (PDB: 3WMC) show that the  
53 thiadiazole moiety bound the -1 subsite of the active pocket, and that the naphthalimide group was sandwiched  
54 by residues Val327 and Trp490 at the +1 subsite.<sup>14</sup> In addition, related to the naphthalimide moiety of **Q1** in  
55 OfHex1, the 4-dimethylaminonaphthalimide of **Q2** rotates approximately 180° to Trp490.<sup>14</sup> This  
56 conformational change ultimately results in tight binding of **Q2** to OfHex1, leading to a 13-fold increase in  
57 potency compared with **Q1**.<sup>14</sup> Thus, the 4-substituent at the naphthalimide group exerts a critical effect on the  
58 potency against OfHex1 (Figure 1).

59 In our previous studies, we presented glycosylated naphthalimide derivatives as promising lead compounds  
60 for  $\beta$ -*N*-acetyl-D-hexosaminidase inhibitors.<sup>15,17,18</sup> In particular, inhibitors **15k**<sup>17</sup> (**1**,  $K_i$  =16.3  $\mu\text{M}$ , unpublished  
61 data), **6e**<sup>15</sup> (**2**,  $K_i$  =22.4  $\mu\text{M}$ , unpublished data), and **6f**<sup>15</sup> (**3**,  $K_i$  =21.8  $\mu\text{M}$ ) exhibited higher inhibitory potency  
62 towards OfHex1. Considering the crucial roles of OfHex1 to agricultural pest control, we further modified

63 these structures to improve the inhibitory efficiency. Prompted by these observations, we retained the frame  
 64 structure of glycosylated naphthalimides and focused on two aspects of optimization, namely the 2-acetamido  
 65 group at the glycosyl moiety (to increase affinity at the -1 the subsite) and the 4-substituent at the  
 66 naphthalimide (to increase affinity at the +1 subsite) (Figure 1). Accordingly, several classes of glycosylated  
 67 naphthalimide derivatives were synthesized and the inhibitory activity against OfHex1 was evaluated. To  
 68 investigate the selectivity of these naphthalimides, their inhibitory capabilities towards human  
 69  $\beta$ -*N*-acetylhexosaminidase B (HsHexB) and human O-GlcNAcase (hOGA) were assessed. This work may  
 70 provide useful information for future design of eco-friendly pesticides.



↓ Inspiration and molecular design

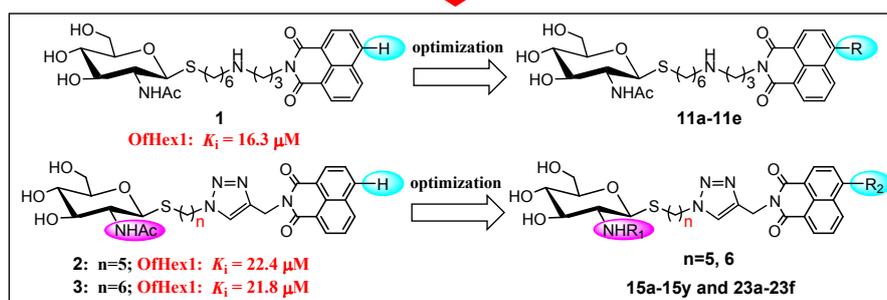


Figure 1. Design of novel glycosylated naphthalimide derivatives for OfHex1.

## MATERIALS AND METHODS

**Materials.** All commercial materials were commercially available and treated with standard methods before use. With TMS as an internal reference, a Bruker AVANCE600 spectrometer was used to record  $^1\text{H}$  NMR-300

76 MHz and  $^{13}\text{C}$  NMR-75 MHz in  $\text{CDCl}_3$  or  $\text{DMSO}-d_6$  at  $25^\circ\text{C}$ . The Bruker Daltonics Bio-TOF-Q III mass  
77 spectrometer (Bruker Co., Karlsruhe, Germany) was used to give high-resolution mass spectra (HRMS).  
78 Reaction progress was monitored using thin layer chromatography (TLC) on silica gel GF254 plates with  
79 detection by charring with 15% (v/v)  $\text{H}_2\text{SO}_4$  in MeOH or by UV light (254 nm).

80 **Synthetic Chemistry.** Detailed synthetic procedures and characterization data for all of the synthesized  
81 compounds are given in the Supporting Information.

82 **Enzyme Preparation.** OfHex1 was overexpressed in *Pichia pastoris* and then purified according to  
83 previous methods.<sup>12</sup> HsHexB was obtained as described in the literature.<sup>17</sup> Human O-GlcNAcase (hOGA) was  
84 overexpressed in *Escherichia coli* BL21(DE3) and purified by IMAC as described previously.<sup>19</sup>

85 **Enzyme Inhibitory Activity Assays.** The inhibitory activities of OfHex1, HsHexB, and hOGA were  
86 assayed in end-point experiments using 4-methylumbelliferyl N-acetyl- $\beta$ -D- glucosaminide (4-MU-GlcNAc)  
87 as the substrate. OfHex1 and hOGA were assayed in 20 mM sodium phosphate buffer (pH 6.5), HsHexB was  
88 assayed in 20 mM sodium citrate buffer (pH 4.5). In a final assay volume of 100  $\mu\text{L}$ , the enzyme was  
89 pre-incubated with inhibitors in buffer for 10 min at  $30^\circ\text{C}$ , then 40  $\mu\text{M}$  substrate (4-MU-GlcNAc) was added.  
90 After incubation for a further 20 min at  $30^\circ\text{C}$ , the reaction was terminated by the addition of 100  $\mu\text{l}$  0.5 M  
91 sodium carbonate solution. The fluorescence of the liberated MU was quantitated using a Varioskan Flash  
92 microplate reader (Thermo Fisher Scientific, Waltham, MA, USA) at an excitation of 366 nm and emission of  
93 445 nm. When determining the  $\text{IC}_{50}$  values, various inhibitor concentrations were used to detect the  
94 corresponding inhibition rates. The inhibition constant ( $K_i$ ) was obtained using Dixon plots by changing the  
95 concentration of the 4-MU-GlcNAc at a constant concentration (40, 20 and 10  $\mu\text{M}$ ).

96 **Molecular Docking.** The crystal structure of OfHex1 in complex with PUGNAc (PDB ID:3OZP)<sup>9</sup> was

97 retrieved from the Protein Data Bank and used as the starting model for molecular docking employing the  
98 Sybyl Software (Version 7.3).<sup>20</sup> Prior to docking calculations, the structures were optimized using MMFF94  
99 force field. For the protein, all water molecules were removed and missing hydrogen atoms were added.  
100 Subsequently, the ligand protomol, appropriate putative ligand pose, was generated by ligand mode based on  
101 the Hammerhead scoring function with the molecular similarity algorithm in the active domain of the  
102 receptor.<sup>21-23</sup> Finally, molecular dockings were performed using the Surflex–Dock algorithm.

103 **Molecular Dynamics (MD) Simulations.** To obtain the convincing conformations, the molecular dynamics  
104 (MD) simulations were carried out after docking. MD simulations of three systems (OfHex1 in complex with  
105 ligands **15r**, **15y**, and **23f**) were performed using the Amber14 package<sup>24</sup>. AMBER03 force field<sup>25</sup> was selected  
106 for the protein, and GAFF force field<sup>26</sup> was selected for the ligand. Each system was immersed in a truncated  
107 octahedral box with TIP3P water molecules and electrostatic neutralized by adding appropriate number of  
108 counterions (Cl<sup>-</sup> or Na<sup>+</sup>). Initially, Sander module was used to realize the energy minimization of the system.  
109 The hydrogen atoms and water molecules were minimized with the 2500 cycles of steepest descent followed  
110 by the conjugated-gradient methods. Then, 2500 cycles of the steepest-descent and 2500 cycles of the  
111 conjugated gradient algorithm were carried out to minimize all atoms of the systems. After that, the system  
112 was heated gradually from 0 to 300 K in the NVT ensemble and equilibrated in 300 K. Finally, MD  
113 Simulations of 30 ns were performed at a constant temperature of 300 K and pressure of 1 atm employing the  
114 PMEMD module. The SHAKE method<sup>27</sup> was applied to constrain hydrogen atoms, and the particle mesh  
115 Ewald (PME) method<sup>28</sup> was used to treat the long-range electrostatic interactions under periodic boundary  
116 conditions.

117 **In Vivo Activity.** Target compounds that efficiently inhibited OfHex1 were further tested for their

118 insecticidal activity against *Myzus persicae* and *Plutella xylostella* according to previous studies.<sup>29-30</sup> Test  
119 compounds were dissolved in DMSO and diluted to a final concentration of 600 µg/mL and 200 µg/mL.  
120 Hexaflumuron was used as a positive control and DMSO was used as a negative control. *Myzus persicae* and  
121 *Plutella xylostella* were treated for 48h and 72h, respectively. Each bioassay was repeated in triplicate.

122 *Ostrinia furnacalis* larvae were raised using an artificial diet with a relative humidity of 70% at 26–28 °C.  
123 Day 1 third-instar larvae were selected for feeding experiments. Compounds **15r** and **15y** were dissolved in  
124 DMSO and diluted with an artificial diet to a final concentration of 600 µg/mL. In the control group, the  
125 artificial diet contained an equal volume of DMSO. Each group contained 30 individual larvae that were  
126 continuously fed for 5 days.<sup>7,16</sup>

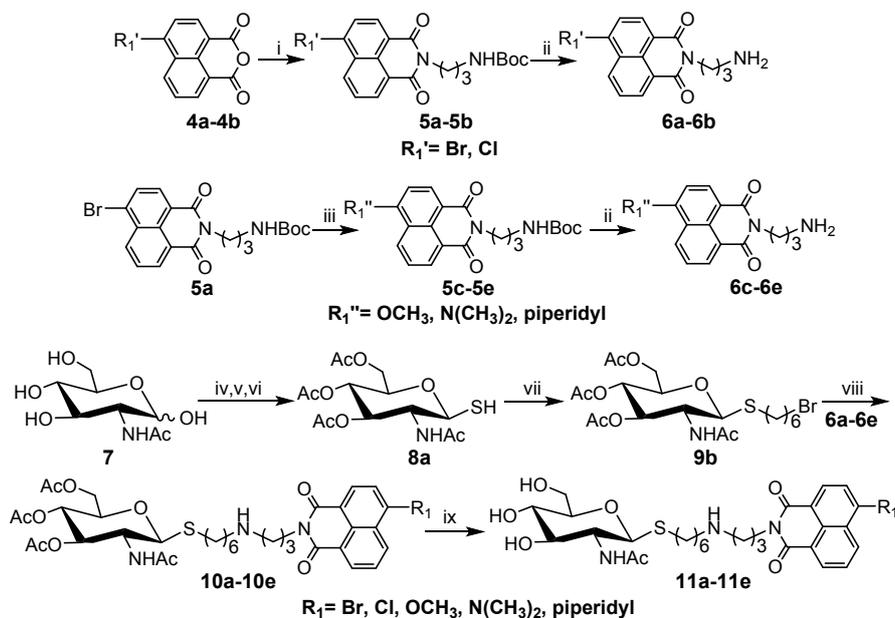
## 127 RESULTS AND DISCUSSION

128 **Synthesis of Target Compounds 11a-11e and 15a-15t.** The target compounds **11a-11e** are outlined in  
129 Scheme 1. Briefly, the key intermediates **6a-6e**<sup>18,31</sup> and **9b**<sup>17</sup> were obtained according to published methods  
130 (Schemes S1-S3). Treatment of **9b** with excess naphthalimides **6a-6e** in the presence of K<sub>2</sub>CO<sub>3</sub> and CH<sub>3</sub>CN  
131 yielded the acetyl-protected compounds **10a-10e**. The deacetylation of **10a-10e** via methanol-ammonia  
132 catalysis resulted in **11a-11e** (Scheme S3).

133 To investigate the influence of the 2-substituent group (at glycosyl moiety) and the 4-substituent group (at  
134 naphthalimide) on the inhibitory activity and selectivity against OfHex1, triazole group-bearing  
135 thioglycosyl-naphthalimides **15a-15t** were synthesized (Scheme 2). Bromides **9a-9d** were prepared according  
136 to literature procedures<sup>15</sup>, which on further reaction with NaN<sub>3</sub> yielded key intermediates **13a-13d** (Scheme  
137 S6). Meanwhile, naphthalimide derivatives **12a-12e** were prepared as previously reported<sup>15,32</sup> (Schemes  
138 S4-S5). Subsequently, azides **13a-13d** were reacted with **12a-12e** in 2:1 THF/water mixture via Cu-catalyzed

139 cycloaddition to form precursors **14a-14t**. Finally, deprotection of **14a-14t** was performed using  
 140 methanol-ammonia to produce the target compounds **15a-15t** (Scheme S6).

141

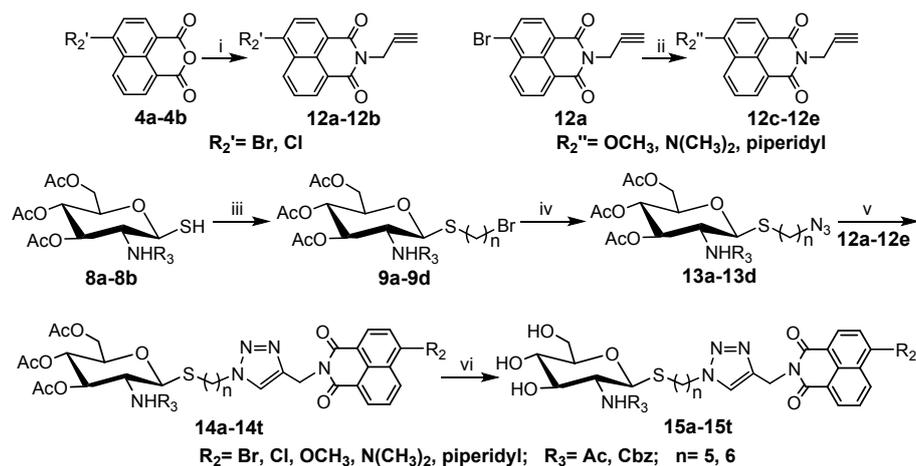


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143

144 **Scheme 1.** Synthesis of glycosylated naphthalimide derivatives **11a-11e**. (i) *tert*-butyl (3-aminopropyl) carbamate, EtOH; (ii)  
 145 DCM, CF<sub>3</sub>COOH; (iii) MeOH, KOH for **5c**; dimethylamine, 2-methoxyethanol for **5d**; piperidine, 2-methoxyethanol for **5e**; (iv)  
 146 AcCl; (v) thiourea, acetone; (vi) Na<sub>2</sub>S<sub>2</sub>O<sub>5</sub>, DCM, H<sub>2</sub>O; (vii) 1,6-dibromohexane, K<sub>2</sub>CO<sub>3</sub>, acetone, H<sub>2</sub>O; (viii) **6a-6e**, K<sub>2</sub>CO<sub>3</sub>,  
 147 CH<sub>3</sub>CN; (ix) NH<sub>3</sub>, MeOH.

148



149

150 **Scheme 2.** Synthesis of glycosylated naphthalimide derivatives **15a-15t**. (i) 2-propynylamine, EtOH; (ii) MeOH, KOH for **12c**;  
 151 dimethylamine, 2-methoxyethanol for **12d**; piperidine, 2-methoxyethanol for **12e**; (iii) a,  $\omega$ - dibromoalkane,  $\text{K}_2\text{CO}_3$ , acetone,  
 152  $\text{H}_2\text{O}$ ; (iv)  $\text{NaN}_3$ ,  $\text{K}_2\text{CO}_3$ , acetone,  $\text{H}_2\text{O}$ ; (v) **12a-12e**,  $\text{CuSO}_4$ , sodium ascorbate, THF,  $\text{H}_2\text{O}$ ; (vi)  $\text{NH}_3$ , MeOH.

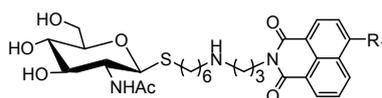
153 **Evaluation of Enzyme Inhibitory Activity.** The target compounds **11a-11e** and **15a-15t** were evaluated for  
 154 their inhibitory activity against OfHex1, HsHexB, and hOGA (Tables 1-3).

155 As shown in Table 1, the substituents at the 4-position of naphthalimide moiety (**11a-11e**) improved the  
 156 inhibitory potency against OfHex1 compared to naphthalimide bearing no 4-substituent group (compound **1**).  
 157 In particular, naphthalimides **11a-11c** bearing smaller functional groups (Br, Cl, OMe) exhibited a minor  
 158 increase in activity. Naphthalimides **11d-11e** bearing a basic substituent ( $\text{N(CH}_3)_2$  or piperidyl) resulted in a  
 159 more effective inhibitory potency. Among these inhibitors, compound **11d** bearing the dimethylamino group  
 160 showed the highest inhibitory activity against OfHex1 with an  $\text{IC}_{50}$  value of  $13.4 \mu\text{M}$  (Table 3). Additionally,  
 161 **11d** showed no inhibitory activities against HsHexB and hOGA, which exhibited improved selectivity  
 162 compared to lead compound **1**.

163 As shown in Table 2, bioassay results of **15a-15t** revealed that 2-substituent of the glycosyl moiety and  
 164 4-substituent of naphthalimide significantly influenced the potency and selectivity of these compounds

165 towards OfHex1. Specifically, upon increasing the size from acetyl (Ac) to benzyloxycarbonyl (Cbz), a loss in  
 166 inhibitory activity against OfHex1 was observed. These results suggest that a large substituent at the 2-position  
 167 of the sugar ring leads to steric hindrance, ultimately decreasing the binding affinity of the inhibitors and  
 168 OfHex1. The 4-substituted group on naphthalimide moiety increased the inhibitory potency against OfHex1  
 169 compared to lead compounds **2** and **3**. In detail, inhibitors bearing 4-bromo (**15a-15b**) and 4-piperidyl  
 170 (**15q-15r**) groups possessed improved potency towards OfHex1 compared to inhibitors bearing Cl (**15e-15f**),  
 171 OCH<sub>3</sub> (**15i-15j**), and N(CH<sub>3</sub>)<sub>2</sub> (**15m-15n**) substituents. The length of the linker between glycosyl and triazole  
 172 also affected the activity against OfHex1, and increasing the length of the carbon atoms from five to six  
 173 slightly increased their activity (e.g. **15b**>**15a** or **15r**>**15q**). Further IC<sub>50</sub> determination showed that **15r** had  
 174 the highest inhibitory potency against OfHex1 (IC<sub>50</sub> = 6.4 μM) in **15a-15t** and more selective than lead  
 175 compounds **2** and **3** (Table 3).

176

177 **Table 1. Inhibition rates of 11a-11e against OfHex1, HsHexB, and hOGA at 10 μM.**

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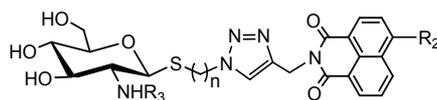
Compd	R <sub>1</sub>	Inhibition rate at 10 μM (%)		
		OfHex1	HsHexB	hOGA
<b>11a</b>	Br	39.6 ± 3.1	13.9 ± 0.6	21.0 ± 0.5
<b>11b</b>	Cl	46.0 ± 1.8	9.2 ± 0.2	18.3 ± 2.2
<b>11c</b>	OMe	42.6 ± 1.6	12.5 ± 1.1	7.7 ± 1.6
<b>11d</b>	N(CH <sub>3</sub> ) <sub>2</sub>	<b>54.6 ± 0.8</b>	<b>13.9 ± 1.7</b>	<b>12.7 ± 0.3</b>

<b>11e</b>	piperidyl	48.3 ± 0.7	13.6 ± 0.5	14.2 ± 0.8
<b>1</b>	H	35.2 ± 1.7	20.1 ± 0.1	9.9 ± 0.3

179

180 **Table 2. Inhibition rates of 15a-15t against OfHex1, HsHexB, and hOGA at 10 μM.**

181



Compd	substituent group			Inhibition rate at 10 μM (%)		
	R <sub>2</sub>	R <sub>3</sub>	n	OfHex1	HsHexB	hOGA
<b>15a</b>	Br	Ac	5	66.0 ± 1.3	11.2 ± 1.4	14.1 ± 0.5
<b>15b</b>	Br	Ac	6	68.1 ± 2.5	6.7 ± 0.8	17.6 ± 0.3
<b>15c</b>	Br	Cbz	5	8.7 ± 1.2	7.9 ± 0.1	4.0 ± 0.4
<b>15d</b>	Br	Cbz	6	15.3 ± 3.6	7.4 ± 0.3	13.0 ± 0.5
<b>15e</b>	Cl	Ac	5	37.6 ± 1.5	10.1 ± 1.8	11.4 ± 0.7
<b>15f</b>	Cl	Ac	6	49.2 ± 2.8	10.3 ± 2.0	17.8 ± 0.3
<b>15g</b>	Cl	Cbz	5	19.6 ± 2.6	4.8 ± 0.7	7.3 ± 1.2
<b>15h</b>	Cl	Cbz	6	11.2 ± 1.3	3.7 ± 0.8	10.9 ± 0.4
<b>15i</b>	OMe	Ac	5	42.2 ± 2.9	10.5 ± 0.4	11.5 ± 0.6
<b>15j</b>	OMe	Ac	6	60.1 ± 0.6	8.0 ± 1.1	13.8 ± 0.5
<b>15k</b>	OMe	Cbz	5	19.6 ± 1.4	2.1 ± 1.5	7.2 ± 0.9
<b>15l</b>	OMe	Cbz	6	11.2 ± 3.0	12.4 ± 0.2	10.9 ± 0.8
<b>15m</b>	N(CH <sub>3</sub> ) <sub>2</sub>	Ac	5	56.9 ± 0.5	8.1 ± 0.3	13.1 ± 0.8

<b>15n</b>	N(CH <sub>3</sub> ) <sub>2</sub>	Ac	6	57.9 ± 2.1	9.7 ± 0.9	13.3 ± 0.2
<b>15o</b>	N(CH <sub>3</sub> ) <sub>2</sub>	Cbz	5	24.2 ± 2.4	9.3 ± 0.7	8.2 ± 0.5
<b>15p</b>	N(CH <sub>3</sub> ) <sub>2</sub>	Cbz	6	19.7 ± 1.7	6.1 ± 1.9	10.9 ± 0.7
<b>15q</b>	piperidyl	Ac	5	67.5 ± 1.8	5.5 ± 0.9	14.0 ± 1.1
<b>15r</b>	piperidyl	Ac	6	<b>75.1 ± 0.4</b>	<b>7.1 ± 1.7</b>	<b>12.6 ± 0.9</b>
<b>15s</b>	piperidyl	Cbz	5	16.3 ± 2.9	4.3 ± 0.6	4.9 ± 0.5
<b>15t</b>	piperidyl	Cbz	6	6.5 ± 2.3	7.9 ± 0.4	6.2 ± 0.7
<b>2</b>	H	Ac	5	29.2 ± 0.3	16.4 ± 3.2	10.3 ± 1.8
<b>3</b>	H	Ac	6	30.6 ± 1.1	16.1 ± 0.7	14.0 ± 1.1

183 **Table 3. IC<sub>50</sub> values of representative compounds for OfHex1, HsHexB, and hOGA.**

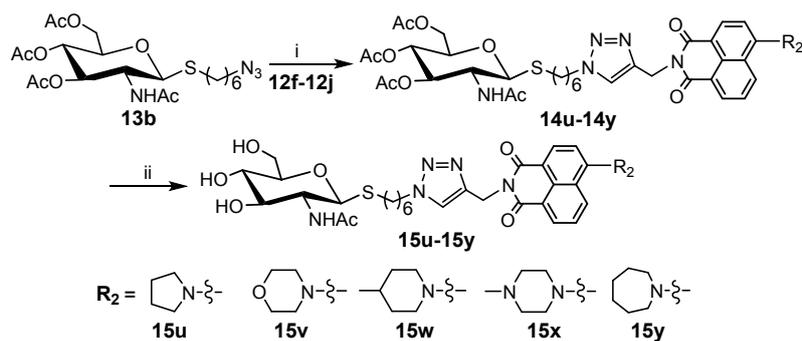
Compd	IC <sub>50</sub> (μM)		
	OfHex1	HsHexB	hOGA
<b>11d</b>	13.4 ± 1.2	> 100	> 100
<b>15a</b>	8.5 ± 0.8	> 100	> 100
<b>15b</b>	8.3 ± 1.0	> 100	66.9 ± 2.1
<b>15j</b>	9.7 ± 0.3	> 100	> 100
<b>15m</b>	11.4 ± 1.1	> 100	> 100
<b>15n</b>	10.9 ± 0.1	> 100	> 100
<b>15q</b>	7.6 ± 0.9	> 100	> 100
<b>15r</b>	<b>6.4 ± 0.3</b>	<b>&gt; 100</b>	<b>&gt; 100</b>
<b>1</b>	18.0 ± 1.4	55.1 ± 0.5	> 100

2	23.9 ± 0.9	97.5 ± 1.6	> 100
3	22.5 ± 0.2	94.2 ± 1.9	> 100

---

184 **Optimization of Inhibitor 15r.** The highest performing compound from the first stage, **15r**, was selected  
185 for further structural optimization. Based on structure–activity relationships, we fixed the 2-acetyl on the  
186 glycosyl moiety and selected a linker with six carbon atoms (n=6, Scheme 2). Then we replaced the  
187 4-piperidyl group with a similar sized nitrogen-containing cycloalkane. Thus, five glycosyl-naphthalimides  
188 **15u-15y** were synthesized (Scheme 3). The synthetic route of **15u-15y** were identical to **15a-15t** and outlined  
189 in Schemes S7 and S8. The analysis of **15u-15y** against OfHex1 (Table 4) showed that the size of the  
190 nitrogen-containing cycloalkane could affect inhibitory efficiency. Shrinking the 4-piperidyl group of **15r** to  
191 4-pyrrolyl (**15u**) enhanced the activity (IC<sub>50</sub> value of 6.4 μM to 5.1 μM). Enlargement of the piperidyl (**15r**) to  
192 azepanyl (**15y**) resulted in a 1-fold increase in inhibitory efficiency. Moreover, the addition of a methyl group  
193 at the 4-position of piperidyl (**15w**) led to slightly increased potency. However, changing the piperidyl group  
194 to morpholino (**15v**) or methylpiperazinyl (**15x**) decreased the potency against OfHex1, particularly for **15x**,  
195 which resulted in a significantly weakened inhibitory effect with IC<sub>50</sub> value of 87.1 μM. Among these  
196 inhibitors, **15y** (IC<sub>50</sub> = 3.1 μM against OfHex1, IC<sub>50</sub> >100 μM against HsHexB, IC<sub>50</sub> = 90.5 μM against  
197 hOGA) exhibited excellent potency and selectivity against OfHex1, confirming the correctness of our  
198 optimization strategy.

199



200

201 **Scheme 3.** Synthesis of glycosylated naphthalimide derivatives **15u-15y**. (i) **12f-12j**, CuSO<sub>4</sub>, sodium ascorbate, THF, H<sub>2</sub>O; (iv)202 NH<sub>3</sub>, MeOH.203 **Table 4. Inhibitory activity of the optimized compounds 15u-15y against OfHex1, HsHexB, and hOGA**

Compd	R <sub>2</sub>	IC <sub>50</sub> (μM)		
		OfHex1	HsHexB	hOGA
<b>15u</b>		5.1 ± 0.3	> 100	68.8 ± 2.4
<b>15v</b>		9.4 ± 0.9	> 100	> 100
<b>15w</b>		6.0 ± 0.2	> 100	71.7 ± 1.9
<b>15x</b>		87.1 ± 3.5	> 100	7.8 ± 0.6
<b>15y</b>		<b>3.1 ± 0.4</b>	> 100	<b>90.5 ± 3.2</b>
<b>15r</b>		<b>6.4 ± 0.3</b>	> 100	> 100

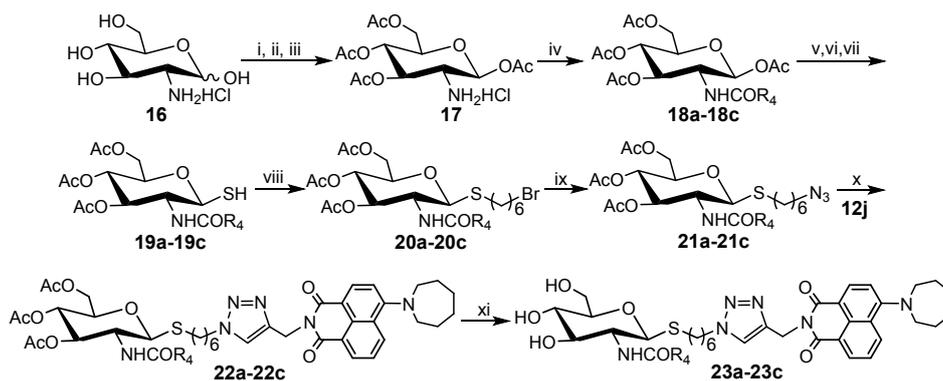
204

205 **Further Optimization of Inhibitor 15y.** Considering that NMAGT bearing a methylamino on thiazoline

206 was approximately 600-fold more potent than **NGT** (bearing a methyl on thiazoline) against OfHex1<sup>13</sup>, we  
207 further attempted to modify the 2-position on glycosyl moiety of **15y** to improve its potency and selectivity.  
208 Our previous experience highlighted that the large substituents at 2-position (i.e. Cbz, Table 2) would result in  
209 activity loss, and so the methyl group of 2-acetamido was replaced with smaller substituents (i.e. NCH<sub>3</sub>, CF<sub>3</sub>,  
210 Et, Pr). Accordingly, compounds **23a-23c** (Scheme 4) bearing Et, NCH<sub>3</sub>, CF<sub>3</sub> and compounds **23e-23f**  
211 (Scheme 5) bearing *n*-Pr and *i*-Pr were synthesized. The specific synthesis methods are shown in Schemes S9  
212 and S10.

213 The inhibitory activity data of compounds **23a-23f** are exhibited in Table 5. Unfortunately, when slightly  
214 increasing the size of the CH<sub>3</sub> group (at the 2-acetamido) to Et, NHCH<sub>3</sub>, CF<sub>3</sub> substituents, the reduced activity  
215 against OfHex1 was observed. Moreover, the larger substituents led to the lower inhibitory potency.  
216 Specifically, replacement of the CH<sub>3</sub> to CF<sub>3</sub> or Et led to a 2-fold reduction in potency (from **15y** to **23c** or **23a**).  
217 The loss of activity from **23a** to **23b** suggested that NH of the the NHCH<sub>3</sub> group was detrimental to the affinity  
218 compared to CH<sub>2</sub> of the Et group at this position. When the substituent was modified from CH<sub>3</sub> to *n*-Pr (**23e**),  
219 the inhibitory activity was reduced 10-fold. This reduction continued with *i*-Pr (**23f**) and OBn (**23d**) groups  
220 leading to a loss of OfHex1 potency (IC<sub>50</sub> >100 μM). Additionally, enlargement of the functional group at the  
221 2-position of glycosyl moiety promoted selectivity towards OfHex1 over hOGA (compared with the IC<sub>50</sub>  
222 values of **15y**, **23a**, **23b**, **23c** towards hOGA), and compounds **23a-23f** showed no inhibitory potency towards  
223 HsHexB.

224



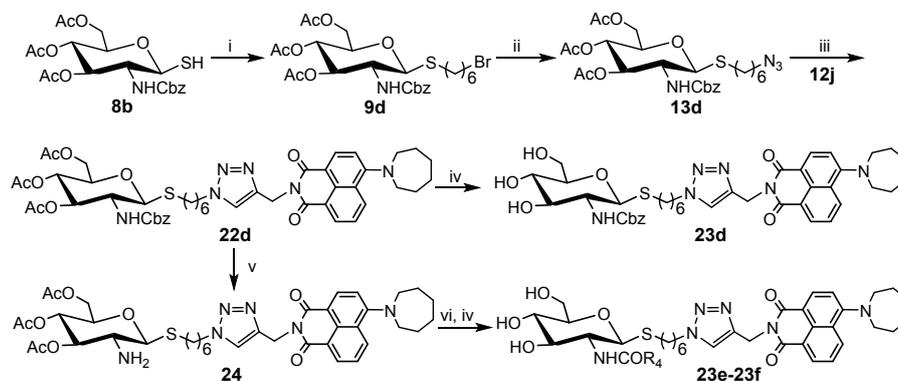
227 **Scheme 4.** Synthesis of glycosylated naphthalimide derivatives **23a-23e**. (i) *p*-anisaldehyde, NaOH, H<sub>2</sub>O; (ii) Py, Ac<sub>2</sub>O; (iii)

228 acetone, HCl, H<sub>2</sub>O; (iv) Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, RCOCl for **18a-18b**; Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, TFAA for **18c**; (v) HBr, CH<sub>3</sub>COOH, CH<sub>2</sub>Cl<sub>2</sub>; (vi)

229 thiourea, acetone; (vii) Na<sub>2</sub>S<sub>2</sub>O<sub>5</sub>, CH<sub>2</sub>Cl<sub>2</sub>, H<sub>2</sub>O; (viii) 1,6-dibromoalkane, K<sub>2</sub>CO<sub>3</sub>, acetone, H<sub>2</sub>O; (ix) NaN<sub>3</sub>, acetone, H<sub>2</sub>O; (x) **12j**,

230 CuSO<sub>4</sub>, sodium ascorbate, THF, H<sub>2</sub>O; (xi) NH<sub>3</sub>, MeOH.

231



234 **Scheme 5.** Synthesis of glycosylated naphthalimide derivatives **23d-23f**. (i) 1,6-dibromoalkane, K<sub>2</sub>CO<sub>3</sub>, acetone, H<sub>2</sub>O; (ii) NaN<sub>3</sub>,

235 acetone, H<sub>2</sub>O; (iii) **12j**, CuSO<sub>4</sub>, sodium ascorbate, THF, H<sub>2</sub>O; (iv) NH<sub>3</sub>, MeOH; (v) Pd/C, H<sub>2</sub>, THF, MeOH, HCl; (vi) R<sub>4</sub>COCl,

236 Et<sub>3</sub>N, CH<sub>2</sub>Cl<sub>2</sub>, DMAP.

237

238 **Table 5. Inhibitory activity of the optimized compounds 23a-23f against OfHex1, HsHexB, and hOGA**

Compd	R <sub>4</sub>	IC <sub>50</sub> (μM)		
		OfHex1	HsHexB	hOGA
<b>23a</b>	Et	<b>6.8 ± 0.5</b>	> 100	<b>94.8 ± 0.9</b>
<b>23b</b>	NHCH <sub>3</sub>	7.2 ± 0.7	> 100	97.1 ± 3.4
<b>23c</b>	CF <sub>3</sub>	<b>6.4 ± 0.4</b>	> 100	<b>98.9 ± 1.4</b>
<b>23e</b>	<i>n</i> -Pr	30.2 ± 1.8	> 100	> 100
<b>23f</b>	<i>i</i> -Pr	> 100	> 100	> 100
<b>23d</b>	OBn	> 100	> 100	> 100
<b>15y</b>	CH <sub>3</sub>	<b>3.1 ± 0.4</b>	> 100	<b>90.5 ± 3.2</b>

239

240 **Inhibitory Mechanism of Glycosylated Naphthalimides for OfHex1.** Two representative inhibitors,  
 241 namely **15r** and **15y**, were selected to investigate the inhibitory mechanism of these glycosylated  
 242 naphthalimides. Dixon plots of **15r** and **15y** against OfHex1 were performed. As shown in Figure 2, **15r** and  
 243 **15y** were competitive inhibitors, with  $K_i$  values of  $5.3 \pm 0.2 \mu\text{M}$  and  $2.7 \pm 0.1 \mu\text{M}$  against OfHex1,  
 244 respectively.

245 To investigate the basis for the potency of these glycosylated naphthalimides towards OfHex1, the binding  
 246 modes of **15r**, **15y** and **23f** with OfHex1 were carried out via molecular docking. As shown in Figure 3a, the  
 247 glycosyl moiety from all three inhibitors was found to bind to -1 subsite of OfHex1 and the naphthalimide  
 248 group extended out from the active pocket. Moreover, the conformation of inhibitor **15r** was similar to that of

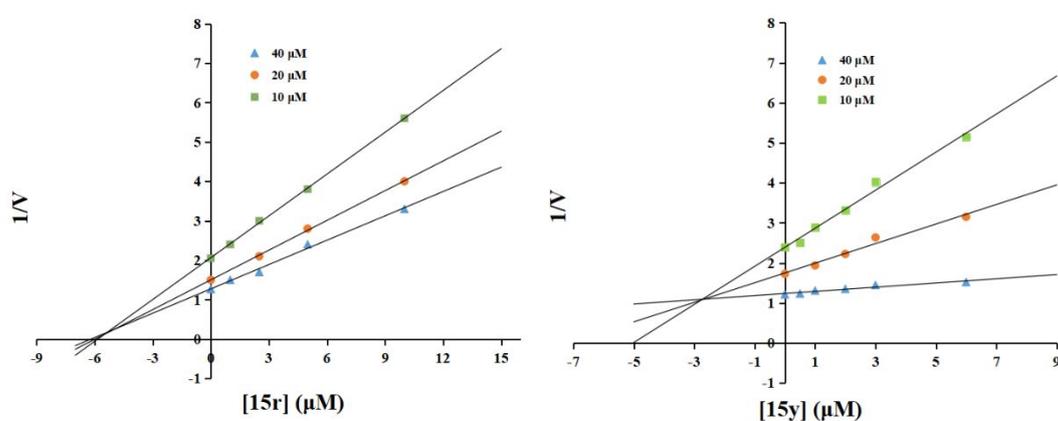
249 **15y** but buried shallower. For the binding mode of **23f**, the naphthalimide group had a great degree of rotation  
250 compared to **15r**, which moved away from residues Gly485 and Ala486.

251 In an effort to further illustrate the appropriate binding modes of compounds **15r**, **15y** and **23f** in complex  
252 with OfHex1, 30 ns MD simulations were performed (Figures 3b-3c). The root-mean-square deviations  
253 (RMSD) values were maintained at around 1.5-3.1 Å for these three systems, suggesting the complexes  
254 underwent reasonable conformational changes (Figure 3b). The conformations of **15r**, **15y** and **23f** combined  
255 with OfHex1 at 30 ns of the MD simulations were superimposed (Figure 3c).

256 In comparison to the docking conformations (Figure 3a), the alkyl chain of **15r** and **15y** showed a high  
257 degree of folding after MD simulations (Figure 3c). Thus, naphthalimide moiety of **15r** and **15y** could bind to  
258 +1 subsite of the OfHex1 pocket with high affinity, forming strong  $\pi$ - $\pi$  stacking interactions with Trp490 and  
259 van der Waals interactions with Val327 (Figures 3c-3e). The glycosyl moiety of **15r** was found to bind to the  
260 -1 subsite via H-bonding interactions with residues His246, Glu297, Asp299, Glu367, and the triazole ring  
261 bound with residue Tyr475 at a distance of 2.1 Å (Figure 3d). These interactions in -1 subsite are much less  
262 than that of PUGNAc ( $K_i = 0.1 \mu\text{M}$  against OfHex1)<sup>9</sup>, which may suggest that the efficiency of these  
263 glycosylated naphthalimids is in micromolar range. As a comparison, the glycosyl moiety of **15y** buried  
264 shallower than **15r**, which might be due to the increased size of cycloalkane (from piperidyl to azepanyl)  
265 leading to the steric hindrance in -1 subsite (Figure 3c). Accordingly, the glycosyl moiety of **15y** formed four  
266 fewer hydrogen bonds than **15r** (five hydrogen bonds) with residues Arg220, Glu297, Asp299, Tyr475  
267 (Figures 3d-3e). However, the triazole ring of **15y** could form three hydrogen bonds (more than **15r**) with  
268 residues Trp483, Val484, and Trp490 (Figures 3d-3e). On the basis of these observations, **15y** ( $K_i = 2.7 \mu\text{M}$ )  
269 ultimately displayed a higher inhibitory potency against OfHex1 than **15r** ( $K_i = 5.3 \mu\text{M}$ ).

270 It was noteworthy that the conformation of **23f** significantly differed to **15r** and **15y** (Figures 3c and 3f). In  
271 detail, the glycosyl moiety of **23f** was buried at the entrance of the OfHex1 pocket and only interacted with  
272 Arg 220, whilst the triazole bearing naphthalimide moiety extended out from the active pocket (Figure 3f).  
273 These results revealed that the branched and large *i*-Pr group led to the glycosyl moiety of **23f** being unable to  
274 move deeply into the -1 subsite of OfHex1, explaining the basis for the loss of activity.

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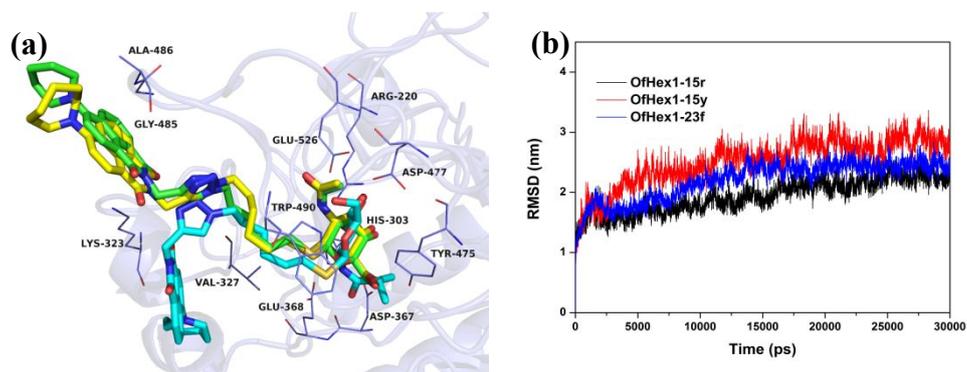


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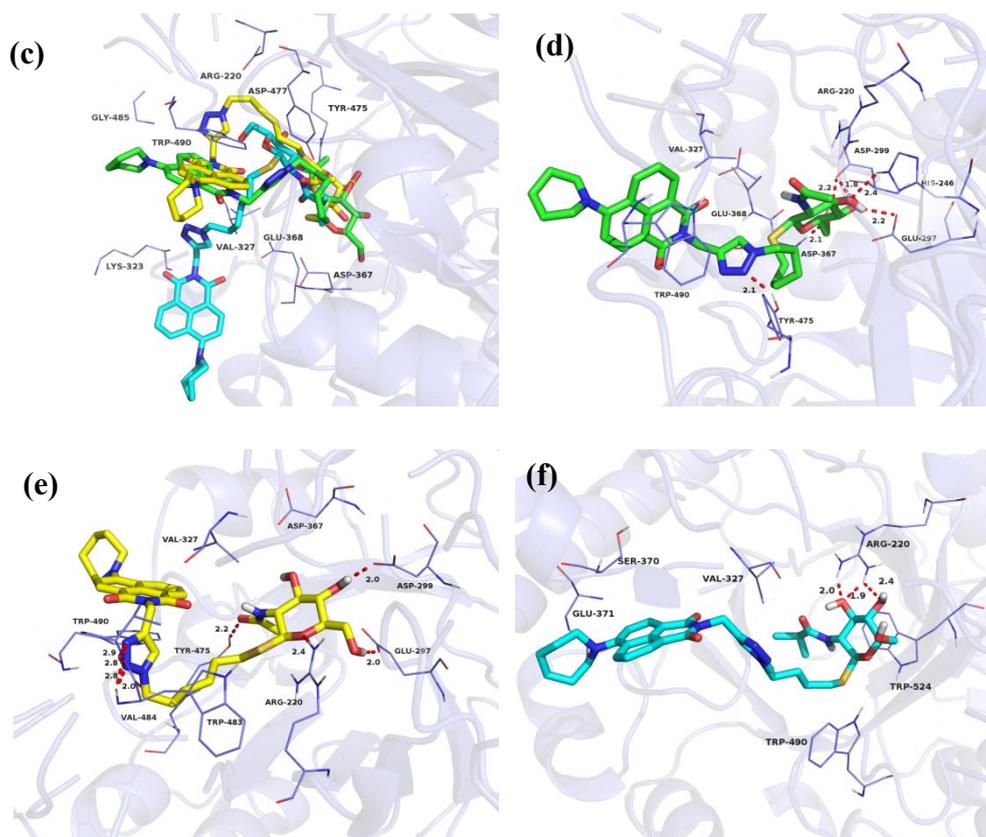
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Figure 2. Dixon plots for inhibitors **15r** and **15y** against OfHex1

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280

281

282 Figure 3. Binding mechanism of **15r**, **15y** and **23f** with OfHex1 revealed by molecular docking and MD simulations. (a)  
 283 Molecular docking and (c) 30 ns MD simulations of OfHex1 in complex with **15r**, **15y** and **23f**. (b) RMSD changes of **15r**, **15y**  
 284 and **23f** in the active pocket of OfHex1. Specific binding modes of (d) OfHex1-**15r**, (e) OfHex1-**15y**, and (f) OfHex1-**23f**  
 285 systems at 30 ns after MD simulations. The protein is shown as a cartoon style. Compound **15r** is shown in green, **15y** is shown  
 286 in yellow, **23f** is shown in cyan (colored according to the element).

287

288 **Insecticidal Activity.** The glycosylated naphthalimides that showed higher inhibitory activity against  
 289 OfHex1 were further selected to evaluate their insecticidal activity. As shown in Table 6, the activity of  
 290 compounds **15b** and **15y** were higher than the control drug hexaflumuron towards *Myzus persicae* with the  
 291 mortalities of 41.5% and 47.2% at a concentration of 200  $\mu\text{g}/\text{mL}$ . Moreover, Compounds **15m**, **15n** and **23a**  
 292 displayed > 50% mortality against *Myzus persicae* at 600  $\mu\text{g}/\text{mL}$ , and the remaining compounds exhibited

20

293 relatively weak activity. For *Plutella xylostella*, compounds **15m**, **15n**, **15r**, **15v**, **15w** and **15y** exhibited high  
 294 activity with mortality rates over 50% at 600  $\mu\text{g/mL}$ , which were lower than those of hexaflumuron (93.3%).  
 295 Although many of the compounds displayed moderate or a lack of insecticidal activity against *Plutella*  
 296 *xylostella*, a smaller body size of the treated insects was noted after 72 hours feeding, compared to the negative  
 297 control groups. The structure–activity relationships of these glycosylated naphthalimides against *Myzus*  
 298 *persicae* and *Plutella xylostella* showed that the  $\text{N}(\text{CH}_3)_2$  substituent at the 4-position of the naphthalimides  
 299 was perhaps the active functional group for insecticidal activity.

300

301 **Table 6. Insecticidal activity of glycosylated naphthalimides (mortality (%)  $\pm$  SD).**

Compd	Myzus persicae		Plutella
			xylostella
	600 $\mu\text{g/mL}$	200 $\mu\text{g/mL}$	600 $\mu\text{g/mL}$
<b>11d</b>	46.2 $\pm$ 2.8		16.7 $\pm$ 0.9
<b>15a</b>	7.9 $\pm$ 0.5		33.3 $\pm$ 1.2
<b>15b</b>	76.7 $\pm$ 3.7	41.5 $\pm$ 0.9	23.3 $\pm$ 0.8
<b>15j</b>	19.6 $\pm$ 3.5		10.0 $\pm$ 0.8
<b>15m</b>	74.1 $\pm$ 1.1	38.8 $\pm$ 2.5	53.3 $\pm$ 1.6
<b>15n</b>	68.0 $\pm$ 1.7	29.4 $\pm$ 1.2	66.7 $\pm$ 2.4
<b>15q</b>	31.9 $\pm$ 2.5		33.3 $\pm$ 2.0
<b>15r</b>	48.9 $\pm$ 3.6		56.7 $\pm$ 1.2
<b>15u</b>	39.2 $\pm$ 0.5		13.3 $\pm$ 1.2

<b>15v</b>	9.5 ± 0.9		50.0 ± 0.0
<b>15w</b>	43.1 ± 1.3		53.3 ± 0.9
<b>15y</b>	82.7 ± 2.4	47.2 ± 1.9	63.3 ± 3.1
<b>23a</b>	55.1 ± 4.1	17.1 ± 1.7	43.3 ± 1.2
<b>23b</b>	39.2 ± 3.0		16.7 ± 0.4
<b>23c</b>	26.2 ± 0.8		20.0 ± 1.6
<b>Hexaflumuron</b>	75.6 ± 2.2	40.7 ± 3.1	93.3 ± 0.5

302

303 **Effects of 15r and 15y on *Ostrinia furnacalis* Larvae.** Compounds **15r** and **15y** were the most effective  
 304 OfHex1 inhibitors observed in this study. Both compounds thus fed to 3th-instar day 1 *Ostrinia furnacalis*  
 305 larvae for 5 days. As shown in Figure 4, the larvae of inhibitor-containing feeding groups (Figures 4c and  
 306 4d) exhibited slower growth than the control group (Figure 4b). Especially inhibitor **15y** led to the death of >  
 307 76 % of larvae after 5 days of feeding. The *Ostrinia furnacalis* larvae that survived had significantly reduced  
 308 body sizes (Figure 4d). These results suggested that OfHex1 inhibitors are promising pest control reagents.



309

310 Figure 4. Bioactivity of OfHex1 inhibitors **15r** and **15y** for *Ostrinia furnacalis* larvae. (a) 3th-instar day 1 larvae prior to  
 311 inhibitor treatment; larvae of (b) DMSO, (c) **15r**, (d) **15y** containing diet-fed group after 5 days of feeding.

312

313 In conclusion, we present the molecular design, synthesis, and inhibitory activity against OfHex1, HsHexB,  
314 and hOGA of glycosylated naphthalimides **11a-11e** and **15a-15t**. As a result, compound **15r** exhibited the  
315 higher efficiency against OfHex1 with a  $K_i$  value of 5.3  $\mu\text{M}$  and excellent selectivity ( $\text{IC}_{50} > 100 \mu\text{M}$  against  
316 HsHexB and hOGA). After in-depth SAR studies, an azepanyl naphthalimide derivative, compound **15y** ( $K_i$   
317 = 2.7  $\mu\text{M}$  against OfHex1,  $\text{IC}_{50} > 100 \mu\text{M}$  against HsHexB,  $\text{IC}_{50} = 90.5 \mu\text{M}$  against hOGA), was identified as a  
318 promising OfHex1 inhibitor. Moreover, molecular docking and MD simulations studies of **15r**, **15y** and **23f**  
319 were performed to allow us to rationalize the basis for the potency of these glycosylated naphthalimides  
320 towards OfHex1. In addition, the efficient OfHex1 inhibitors were further selected to evaluate their insecticidal  
321 activity against *Myzus persicae* and *Plutella xylostella*. The results showed that compounds **15b** and **15y** had  
322 higher insecticidal activity against *Myzus persicae* than the commercial pesticide hexaflumuron. Furthermore,  
323 feeding experiments demonstrated that the OfHex1 inhibitors **15r** and **15y** could effectively inhibit the growth  
324 of *Ostrinia furnacalis* larvae. Thus, the structure–activity relationships combined with docking and MD  
325 simulations studies provide direction for the further structural optimization for OfHex1, and the development  
326 of green pest control and management agents.

327

328 ASSOCIATED CONTENT

329 ■ **Supporting Information**330 Experimental procedures, Molecular docking and MD simulations results,  $^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectrum.331 ■ **AUTHOR INFORMATION**

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335 **Author Contributions**

336 #These authors contributed equally to this work. All authors have given approval to the final version of the  
337 manuscript.

338 **Notes**

339 The authors declare no competing financial interest.

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343 **■ ABBREVIATIONS**

344 GH18, glycoside hydrolase family 18; GH20, glycoside hydrolase family 20; MD, molecular dynamics.

345

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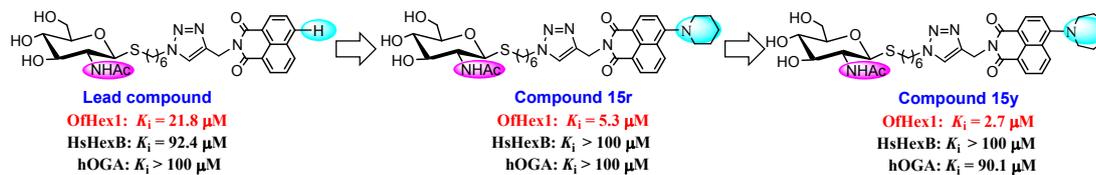
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