




Article

# Synthesis of Chromen-4-One-Oxadiazole Substituted Analogs as Potent $\beta$ -Glucuronidase Inhibitors

Muhammad Taha <sup>1,\*</sup>, Fazal Rahim <sup>2</sup>, Muhammad Ali <sup>3</sup>, Muhammad Naseem Khan <sup>4</sup>, Mohammed A. Alqahtani <sup>5</sup>, Yasser A. Bamarouf <sup>5</sup>, Mohammed Gollapalli <sup>5</sup>, Rai Khalid Farooq <sup>6</sup>, Syed Adnan Ali Shah <sup>7,8</sup> , Qamar Uddin Ahmed <sup>9</sup>  and Zainul Amiruddin Zakaria <sup>10,11,\*</sup> 

- <sup>1</sup> Department of Clinical Pharmacy, Institute for Research and Medical Consultations (IRMC), Imam Abdulrahman Bin Faisal University, P.O. Box 1982, Dammam 31441, Saudi Arabia
- <sup>2</sup> Department of Chemistry, Hazara University, Mansehra 21300, Khyber Pakhtunkhwa, Pakistan; fazalstar@gmail.com
- <sup>3</sup> Natural and Medical Sciences Research Center, University of Nizwa, P.O. Box 33, Birkat Al Mauz, Nizwa 616, Sultanate of Oman; m\_alisaad@yahoo.com
- <sup>4</sup> Department of Chemistry, COMSATS Institute of Information Technology, University Road, Abbottabad 22060, KPK, Pakistan; naseemkhan.tareen@gmail.com
- <sup>5</sup> Department of Computer Information Systems, College of Computer Science & Information Technology, Imam Abdulrahman Bin Faisal University, P.O. Box 1982, Dammam 31441, Saudi Arabia; maqhtani@iau.edu.sa (M.A.A.); yabamarouf@iau.edu.sa (Y.A.B.); magollapalli@iau.edu.sa (M.G.)
- <sup>6</sup> Department of Neuroscience Research, Institute for Research and Medical Consultations (IRMC), Imam Abdulrahman Bin Faisal University, P.O. Box 1982, Dammam 3144, Saudi Arabia; rkfarooq@iau.edu.sa
- <sup>7</sup> Faculty of Pharmacy, Universiti Teknologi MARA Puncak Alam Campus, 42300 Bandar Puncak Alam, Selangor D.E., Malaysia; benzene301@yahoo.com
- <sup>8</sup> Atta-ur-Rahman Institute for Natural Products Discovery (AuRIns), Universiti Teknologi MARA Puncak Alam Campus, 42300 Bandar Puncak Alam, Selangor D.E., Malaysia
- <sup>9</sup> Department of Pharmaceutical Chemistry, Kulliyah of Pharmacy, International Islamic University Malaysia, 25200 Kuantan, Pahang DM, Malaysia; quahmed@iiu.edu.my
- <sup>10</sup> Department of Biomedical Science, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia
- <sup>11</sup> Halal Institute Research Institute, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia
- \* Correspondence: mtaha@iau.edu.sa (M.T.); zaz@upm.edu.my (Z.A.Z.); Tel.: +00966502057370 (M.T.); +603-8947-2111 (Z.A.Z.); Fax: +603-8943-6178 (Z.A.Z.)

Received: 26 December 2018; Accepted: 2 February 2019; Published: 18 April 2019



**Abstract:** Chromen-4-one substituted oxadiazole analogs **1–19** have been synthesized, characterized and evaluated for  $\beta$ -glucuronidase inhibition. All analogs exhibited a variable degree of  $\beta$ -glucuronidase inhibitory activity with  $IC_{50}$  values ranging in between  $0.8 \pm 0.1$ – $42.3 \pm 0.8$   $\mu$ M when compared with the standard D-saccharic acid 1,4 lactone ( $IC_{50} = 48.1 \pm 1.2$   $\mu$ M). Structure activity relationship has been established for all compounds. Molecular docking studies were performed to predict the binding interaction of the compounds with the active site of enzyme.

**Keywords:** chromen-4-one; oxadiazole; synthesis;  $\beta$ -glucuronidase inhibition; molecular docking; SAR

## 1. Introduction

$\beta$ -Glucuronidase (E.C.3.2.1.31) is one of the most extensively studied enzymes in the metabolic hydrolysis of conjugated compounds, and eliminates large number of toxic compounds from body as glucuronides [1]. It acts as a glycoside hydrolase enzyme which induces to breaking of glucuronosyl-O-bonds [2]. In human body  $\beta$ -glucuronidase is present in many body fluids and

organs [3]. Increased activity of  $\beta$ -glucuronidase has resulted in various health issues that include renal disorders [4], urinary tract infections [5], epilepsy [6], renal transplant rejection [7], neoplasm of bladder [8] and breast cancers [9]. Bacterial  $\beta$ -glucuronidase and  $\beta$ -glucosidase in the human colon are involved in the metabolism and activation of xenobiotics derived from dietary compounds [10]. The enzymes are identified to be mediators of colon cancer (CRC). Many carcinogenic dietary compounds are metabolized in the liver and then conjugated to glucuronic acid before being excreted with bile into the small intestine. In the colon, bacterial  $\beta$ -glucuronidase hydrolyzes the conjugates, thereby releasing the parent compounds or their activated hepatic metabolites [11]. Epidemiological studies have shown that populations at high risk for colorectal cancer have high levels of fecal  $\beta$ -glucuronidase activity and CRC patients have significantly higher levels of the enzyme than healthy controls [12].

Chromone (1,4-benzopyrone) is a derivative of benzopyran with a substituted keto group on the pyran ring. Chromones (4H-chromen-4-ones) constitute a well-known class of naturally occurring oxygen-containing heterocycles. These compounds exhibit important biological properties such as anticancer [13], cytotoxic [14], antioxidant [15], anti-inflammatory [16], antifungal activities [17].

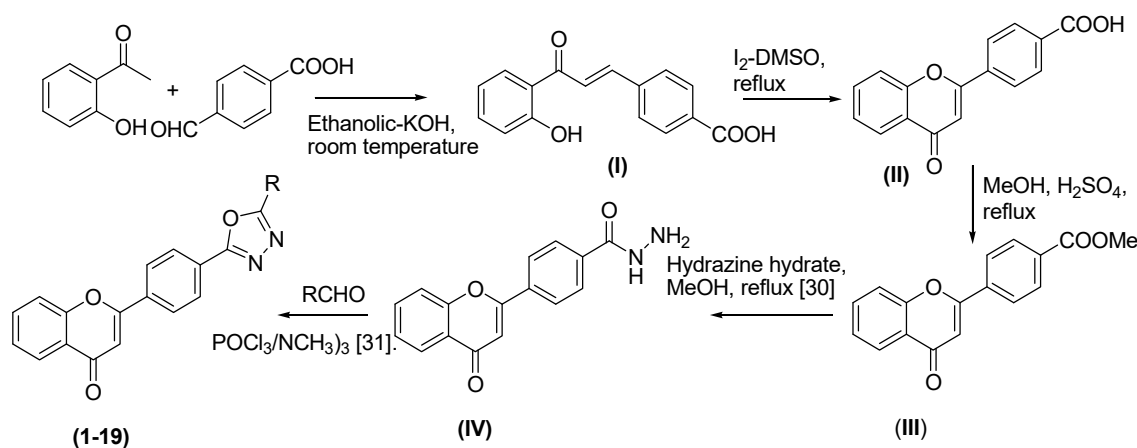
1,3,4-Oxadiazole containing compounds have remarkable biological and pharmacological properties. Recently, differently substituted 1,3,4-oxadiazoles have been reported for their biological activities such as antibacterial [18], antifungal [19], insecticidal [20], antiviral [21], inflammatory [22], antitubercular [23], antitumor [24] and analgesic activity [25].

Our research group has been working on synthesis of variety of heterocycles and explored a huge biological potential of these compounds [26–32]. Present work belongs to a simple and reliable synthetic protocol that has been developed for the synthesis of a series of chromone substituted oxadiazole pharmacophores in another effort towards developing more novel agents against  $\beta$ -glucuronidase enzyme. This is beyond any doubt that combination of two or more biologically active compounds grants newer molecules with complementary pharmacophoric potentials or with different mechanisms of actions. The unity of pharmacophoric approach, as a result, could possibly address the active site of different targets more effectively and could offer the possibility to overcome the issue of drug resistance and unwanted effects [33–36]. The outcomes of this merger have already been the subject of attention in finding new antimicrobials [37–41].

## 2. Result and Discussions

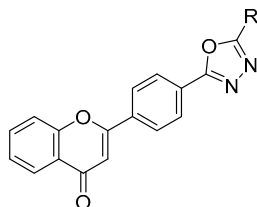
### 2.1. Chemistry

The synthesis of the novel series of chromene-based oxadiazole  $\beta$ -glucuronidase inhibitors 1–19 commenced with the reaction of 2-hydroxyacetophenone and *p*-formylbenzoic acid in ethanolic potassium hydroxide at room temperature (Scheme 1).



**Scheme 1.** Synthesis of chromene based oxadiazole derivatives 1–19.

The adduct **I** underwent oxidative cyclization to form 4-substituted chromenone analog **II**. This acid analog **II** was esterified to give **III** and reacted with hydrazine hydrate to form the corresponding hydrazide **IV** which upon reaction with the corresponding aldehydes under the described conditions formed the desired oxadiazoles 1–19 Figure 1. All synthesized compounds were characterized using various spectroscopic methods.



**Figure 1.** Structure of the basic skeleton of the chromen-4-one based oxadiazole derivatives 1–19.

## 2.2. $\beta$ -Glucuronidase Activity

4-Substituted chromenone-based oxadiazole derivatives 1–19 were synthesized and evaluated for  $\beta$ -glucuronidase inhibitory potential. All the synthesized analogs showed potent inhibitory potential, ranging between  $0.8 \pm 0.1 \mu\text{M}$  to  $42.3 \pm 0.8 \mu\text{M}$  when compared with the standard D-saccharic acid 1,4 lactone ( $\text{IC}_{50}$  value =  $48.1 \pm 1.2 \mu\text{M}$ ) (Table 1). The structure activity relationship has been established on the basis of substitution pattern on the phenyl ring.

**Table 1.** Chromen-4-one based oxadiazole derivatives 1–19.

S. No	R	$\text{IC}_{50} \pm \text{SEM}^a$	S. No	R	$\text{IC}_{50} \pm \text{SEM}^a$
1		$2.6 \pm 0.1$	11		$26.1 \pm 0.5$
2		$5.1 \pm 0.2$	12		$17.2 \pm 0.4$
3		$2.1 \pm 0.1$	13		$32.20 \pm 0.6$
4		$1.1 \pm 0.05$	14		$39.5 \pm 0.7$
5		$3.8 \pm 0.2$	15		$20.1 \pm 0.5$
6		$0.8 \pm 0.1$	16		$14.9 \pm 0.4$
7		$9.4 \pm 0.3$	17		$34.7 \pm 0.6$
8		$13.0 \pm 0.3$	18		$24.5 \pm 0.5$
9		$8.6 \pm 0.3$	19		$42.3 \pm 0.8$
10		$21.3 \pm 0.4$		D-saccharic acid 1,4 lactone <sup>c</sup>	$48.1 \pm 1.2$

<sup>a</sup> standard error mean; <sup>c</sup> standard drug.

The compound **6** ( $IC_{50} = 0.8 \pm 0.1 \mu\text{M}$ ), an *ortho*-fluoro analog was found to be the most potent among the series. The greater potential shown by this analog might be due to the electron withdrawing fluoro group. If we compare it with other fluoro analogs like compound **4** ( $IC_{50} = 1.1 \pm 0.05 \mu\text{M}$ ) a *para*-fluoro analog and compound **5** ( $IC_{50} = 3.8 \pm 0.2 \mu\text{M}$ ), a *meta*-fluoro analog, the analog **6** was found superior than the analogs **4** and **5**. The little bit difference in their potential is appears to be due to the difference in the position of fluoro groups on phenyl ring Figure 2.

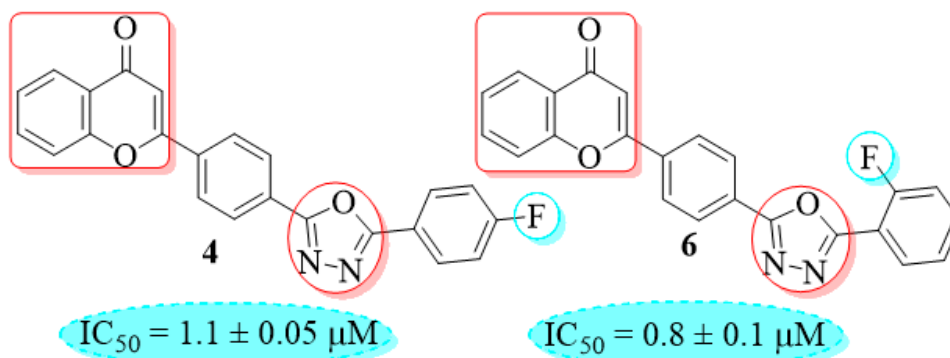


Figure 2. *ortho*-Fluoro and *para*-fluoro analogs (most potent ones).

The compound **3**, an *ortho*-chloro analog ( $IC_{50} = 2.1 \pm 0.1 \mu\text{M}$ ) was found to be the second most potent analog among the series if we compare it with other chloro analogs like compound **1** ( $IC_{50} = 2.6 \pm 0.1 \mu\text{M}$ ) a *para*-chloro analog and compound **2** ( $IC_{50} = 5.1 \pm 0.2 \mu\text{M}$ ) a *meta*-fluoro analog. The greater potential is again seemingly due to position of substituent(s) on phenyl ring.

Similarly, compound **9**, an *ortho* methyl analog ( $IC_{50} = 8.6 \pm 0.3 \mu\text{M}$ ) showed outstanding inhibitory potential. If we compare it with other methyl analogs like analog **7** ( $IC_{50} = 9.4 \pm 0.3 \mu\text{M}$ ) a *para*-methyl analog and compound **8** ( $IC_{50} = 13.0 \pm 0.3 \mu\text{M}$ ) a *meta*-methyl analog, compound **9** is superior.

Similarly compound **12** ( $IC_{50} = 17.2 \pm 0.4 \mu\text{M}$ ), an *ortho*-nitro analog showed greater potential if we compare it with other nitro analogs like compound **11** ( $IC_{50} = 26.1 \pm 0.5 \mu\text{M}$ ) a *meta*-nitro analog and compound **10** ( $IC_{50} = 21.3 \pm 0.4 \mu\text{M}$ ) a *para*-nitro analog. The same pattern was observed in other analogs with different substituents.

A closer look upon the structures of compounds and their corresponding  $IC_{50}$  values revealed that the presence of different substitutions i.e., fluoro, chloro, methyl, nitro, methoxy and pyridinyl on the aromatic side chain is of great importance. It was also observed that either an EWG or EDG on the phenyl part showed great potential but the slight difference in potential was mainly affected by the position of the substituent.

Further to above, it is evident from the results that the presence of halogens or alkyl groups on benzene ring influenced the inhibitory potentials significantly. Similarly, nitro substitutions reduced the potential of inhibition. Apart from direct interactions of these attachments, it can be easily argued that these substitutions significantly affected the electronic distributions and intensities at the active sites. In many cases, oxadiazole part also played a significant role in forming hydrogen bond interactions with the active site of enzyme. Most important of these positions are Asp 207, Glu 451 and Glu 540. To better understand the binding interactions of the most active analogs, molecular docking analysis were performed.

### 2.3. Molecular Docking Studies

Experimental results were further validated by carrying out molecular docking studies and more information was obtained concerning binding modes of synthesized compounds within the active site of  $\beta$ -Glucuronidase. The X-ray structure of  $\beta$ -glucuronidase was retrieved from the Protein Data Bank with PDB ID: 1BHG. All synthesized compounds were docked into the active site of protein and

conformations selected for each docked compound were of least energy and were pictured to further understand the interactions between ligands and receptor.

Binding mode of compound **6**, the most active member ( $IC_{50} = 0.80 \pm 0.10 \mu\text{M}$ ) among the series was analyzed. Three conventional hydrogen bond interactions were observed between the fluorine atom at *ortho* position and side chain amino acids including Tyr 508 and Lys 606 with a internuclear distance of 2.63 Å, 2.66 Å and 1.83 Å respectively. Meanwhile, C-H and N-H groups of Asn 484 were involved in forming carbon hydrogen bond with ether oxygen of chromene-4-one ring (2.88 Å) and  $\pi$ -donor hydrogen bond with the central arene ring of compound **6** (3.14 Å). Significant residues like Asp 207, Glu 451 and Glu 540 were not involved in hydrogen bonding due to absence of hydrogen bond donor groups on compound **6** instead, these residues were involved in  $\pi$ -anion interactions as shown in Figure 3a. Furthermore, the ligand /receptor complex was further stabilized by  $\pi$ - $\pi$  T-shaped and  $\pi$ - $\pi$  Stacked interactions between indole ring of Trp 587 and phenol ring of Tyr 508 respectively.

Compound **4**, the second most active member with  $IC_{50} = 1.10 \pm 0.05 \mu\text{M}$  was evaluated and a conventional hydrogen bond between fluorine atom at *para* position and N-H group of Tyr 487 with a internuclear distance of 2.22 Å and carbon hydrogen bond between nitrogen atom of pyrazole ring of compound **4** and C-H (at imidazole ring) of His 509 (2.58 Å) were detected. Meanwhile, the catalytic residue Glu 451, as in case of compound **6** was also involved in forming two  $\pi$ -Anion interactions with the chromene-4-one ring of compound **4**. Side chain residues involved in  $\pi$ - $\pi$  T-Shaped interactions with compound **4** were Tyr 504 and Tyr 508 as shown in Figure 3b.

Analysis of interactions between third most active compound **3** ( $IC_{50} = 2.10 \pm 0.10 \mu\text{M}$ ) and active site of  $\beta$ -Glucuronidase revealed that it was not involved in conventional hydrogen bonding. It was found that the ether oxygen atom of chromene-4-one ring and central arene ring of compound **3** were involved in forming carbon hydrogen bond (2.77 Å) and  $\pi$ -Donor hydrogen bond (3.26 Å) with the C-H and N-H groups of Asn 484 respectively. The chlorine atom presents at *ortho* position formed so-called halogen bond of distance 2.94 Å and bond angle  $\sim 153^\circ$  with the carboxyl oxygen of Glu 451. Other interactions shown by compound **3** were almost the same as those shown by compound **6** (Figure 3c).

The least active compound **19** ( $IC_{50} = 42.30 \pm 0.80 \mu\text{M}$ ) among the series failed to form many interactions as those made by active compounds including conventional hydrogen bonding and halogen bonding. It was involved in forming carbon hydrogen and  $\pi$ -Donor hydrogen bond with Asn 484 as formed by compound **3** but with one longer bond distance i.e., 2.89 Å and 3.2 Å. Other interactions resulted between compound **19** and Asp 207, Glu 451, Tyr 508, Glu 540, Trp 587 of the side chain were  $\pi$ -Anion (3.87 Å),  $\pi$ -Anion (4.11 Å),  $\pi$ - $\pi$  Stacked (4.27 Å),  $\pi$ -Anion (3.51) and  $\pi$ - $\pi$  T-shaped (5.19 Å and 5.29 Å) respectively (Figure 3d).

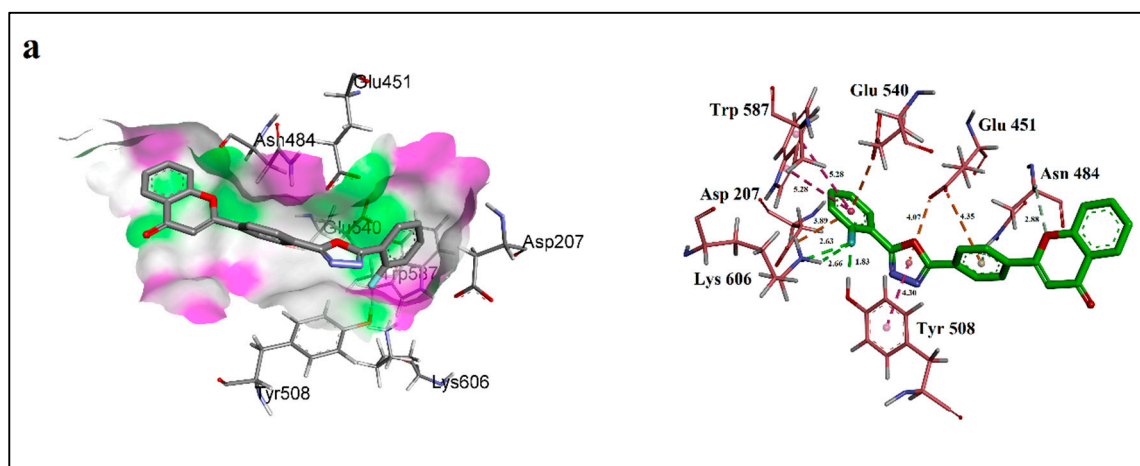
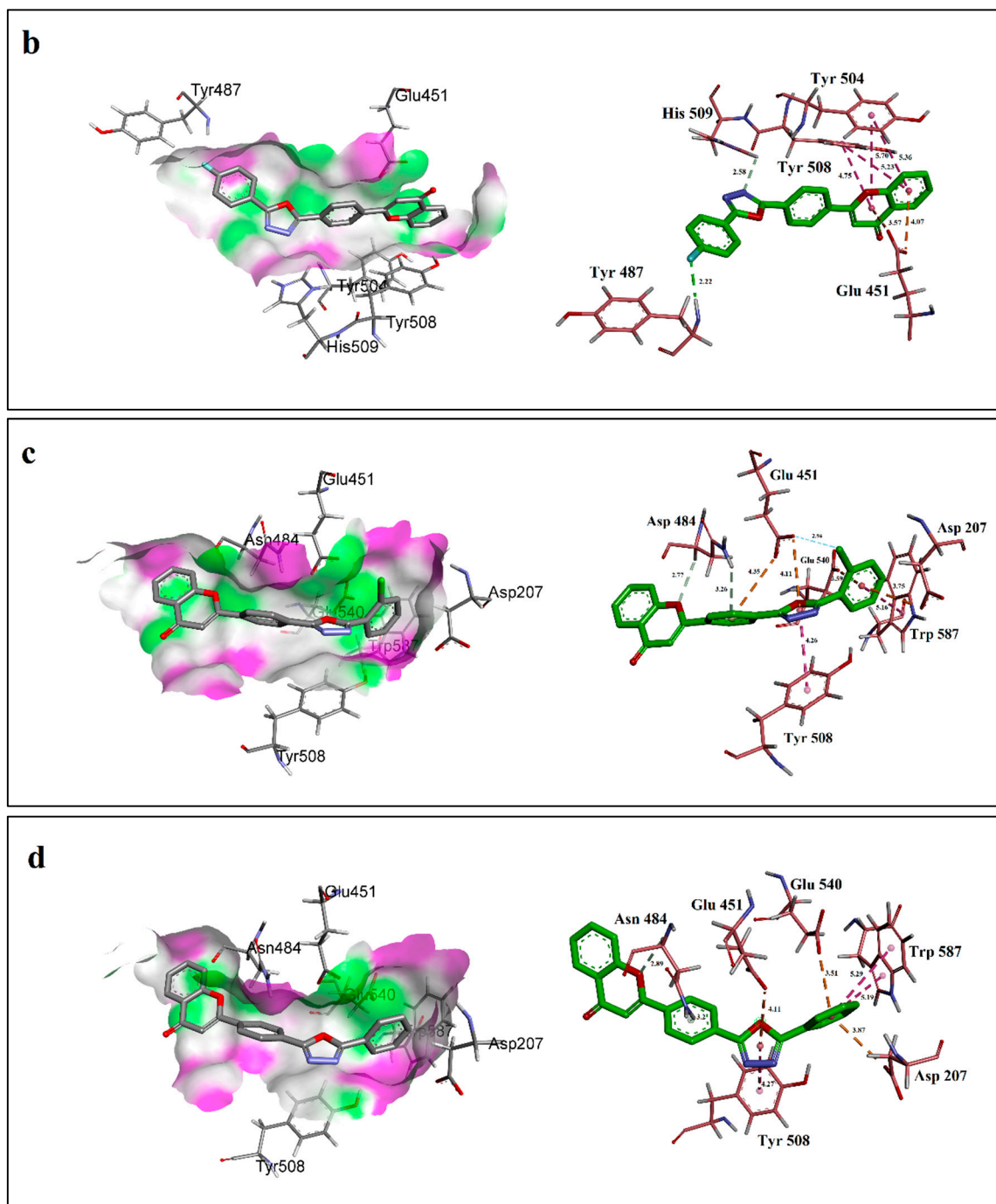


Figure 3. Cont.



**Figure 3.** Ligand–protein interactions of compound (a) 6, (b) 4, (c) 3 and (d) 19 with the active site of  $\beta$ -glucuronidase (1BHG) created by using Discovery Studio 17.2. The left side displays 3D maps of the hydrogen bonds of the compounds in the binding site. The right side shows the 2D interaction patterns. Dashed lines show the interactions among the ligand and the amino acids of the protein. Distances are in Angstrom.

### 3. Conclusions

Chromen-4-one based oxadiazole derivatives 1–19 have been synthesized and evaluated for  $\beta$ -glucuronidase inhibitory potential. All analogs displayed potent  $\beta$ -glucuronidase inhibitory potential ranging between  $0.8 \pm 0.1$  to  $42.3 \pm 0.8 \mu\text{M}$  as a compare to standard D-saccharic acid 1,4 lactone ( $\text{IC}_{50} = 48.1 \pm 1.2 \mu\text{M}$ ). The fluoro derivatives were found to be the more potent among the series. A molecular docking study indicated that the top ranked conformations of almost all compounds

were well accommodated inside the active site of  $\beta$ -glucuronidase enzyme and they were involved in various types of interactions with the active site residues of the  $\beta$ -glucuronidase enzyme.

## 4. Experimental

### 4.1. General Information

The intermediates 4-(4-oxo-4H-chromen-2-yl)benzoic acid (**II**), methyl 4-(4-oxo-4H-chromen-2-yl)benzoate (**III**) and 4-(4-oxo-4H-chromen-2-yl)benzohydrazide (**IV**) were synthesized as reported in our previous paper [42].

### 4.2. General Procedure for Synthesis of Flavone-Based Oxadiazoles

Compound **IV** (0.5 mmol) and a substituted arylcarboxylic acid (0.5 mmol) were mixed in a 50 mL round bottom flask then  $\text{POCl}_3$  (5 mL) was added dropwise [33]. The mixture was refluxed for 4–5 h while the reaction progress was monitored using TLC until completion, when the reaction mixture cooled to room temperature and poured onto crushed ice.  $\text{NaHCO}_3$  solution was added and the resulting solid mass that precipitated out was filtered, dried, and recrystallized from methanol in good to excellent yields.

### 4.3. 2-(4-(5-(4-Chlorophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (**1**)

Solid; Yield 75%; m.p. 288–289 °C;  $^1\text{H-NMR}$  ( $\text{MeOD-}d_4$ , 500 MHz):  $\delta$  8.05 (d,  $J = 8.5$  Hz, 2H), 7.96 (d,  $J = 8.5$  Hz, 2H), 7.84 (d,  $J = 8.0$  Hz, 2H), 7.75 (dd,  $J = 8.0, 2.0$  Hz, 1H), 7.44 (d,  $J = 8.0$  Hz, 2H), 7.20 (s, 1H), 7.17 (td,  $J = 8.5$  Hz, 1.0 Hz, 1H), 6.94–6.91 (m, 2H);  $^{13}\text{C-NMR}$  (125 MHz,  $\text{DMSO-}d_6$ ):  $\delta$  177.4, 164.5, 164.4, 163.6, 156.2, 135.2, 132.6, 132.2, 132.2, 131.6, 130.1, 128.5, 128.5, 127.8, 127.8, 127.1, 127.1, 126.0, 125.1, 124.0, 123.2, 116.0, 104.6; HR-EI-MS:  $m/z$  Calcd for  $\text{C}_{23}\text{H}_{13}\text{ClN}_2\text{O}_3$ ,  $[\text{M}]^+$  414.0770; found 414.0752.

### 4.4. 2-(4-(5-(3-Chlorophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (**2**)

Solid; Yield 79%; m.p. 282–283 °C;  $^1\text{H-NMR}$  ( $\text{DMSO-}d_6$ , 500 MHz):  $\delta$  8.04 (s, 4H), 7.83 (s, 1H), 7.72–7.70 (m, 2H), 7.47 (d,  $J = 4.5$  Hz, 2H), 7.31 (s, 1H), 7.16 (t,  $J = 8.5$  Hz, 1H), 6.94 (d,  $J = 8.5$  Hz, 1H), 6.85 (t,  $J = 8.0$  Hz, 1H);  $^{13}\text{C-NMR}$  (125 MHz,  $\text{DMSO-}d_6$ ):  $\delta$  177.3, 166.2, 164.4, 156.1, 137.8, 135.1, 134.1, 130.1, 129.9, 129.8, 127.6, 127.6, 127.2, 127.2, 127.0, 126.1, 125.5, 125.5, 125.2, 123.6, 124.2, 116.3, 104.6; HR-EI-MS:  $m/z$  Calcd for  $\text{C}_{23}\text{H}_{13}\text{ClN}_2\text{O}_3$ ,  $[\text{M}]^+$  414.0771; found 414.0751.

### 4.5. 2-(4-(5-(2-Chlorobenzyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (**3**)

Solid; Yield 81%; m.p. 276–277 °C;  $^1\text{H-NMR}$  ( $\text{DMSO-}d_6$ , 500 MHz):  $\delta$   $^1\text{H-NMR}$  ( $\text{MeOD-}d_4$ , 500 MHz):  $\delta$  8.24 (dd,  $J = 8.0, 2.0$  Hz, 1H), 8.02 (d,  $J = 8.5$  Hz, 2H), 7.96 (d,  $J = 8.5$  Hz, 2H), 7.70 (dd,  $J = 8.0, 2.0$  Hz, 1H), 7.42 (d,  $J = 8.5$  Hz, 1H), 7.43–7.40 (m, 2H), 7.22–7.18 (m, 2H), 6.94–6.91 (m, 2H);  $^{13}\text{C-NMR}$  (125 MHz,  $\text{DMSO-}d_6$ ):  $\delta$  177.4, 166.3, 164.1, 156.0, 138.0, 135.3, 134.5, 130.8, 130.6, 129.1, 127.5, 127.5, 127.1, 127.1, 127.0, 126.9, 126.0, 125.2, 124.2, 123.7, 123.2, 116.2, 104.7; HR-EI-MS:  $m/z$  Calcd for  $\text{C}_{23}\text{H}_{13}\text{ClN}_2\text{O}_3$ ,  $[\text{M}]^+$  414.0772; found 414.0754.

### 4.6. 2-(4-(5-(4-Fluorophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (**4**)

Solid; Yield 82%; m.p. 299–300 °C;  $^1\text{H-NMR}$  ( $\text{DMSO-}d_6$ , 500 MHz):  $\delta$  8.00–7.96 (m, 4H), 7.78–7.75 (m, 3H), 7.33–7.30 (m, 3H), 7.18 (t,  $J = 7.0$  Hz, 1H), 6.94 (d,  $J = 8.0$  Hz, 1H), 6.88 (t,  $J = 7.0$  Hz, 1H);  $^{13}\text{C-NMR}$  (125 MHz,  $\text{DMSO-}d_6$ ):  $\delta$  177.8, 164.6, 164.6, 163.8, 163.0, 156.1, 135.0, 130.5, 129.3, 129.3, 127.6, 127.6, 126.8, 126.8, 125.6, 125.5, 124.1, 123.6, 121.5, 116.3, 115.8, 115.8, 104.7; HR-EI-MS:  $m/z$  Calcd for  $\text{C}_{23}\text{H}_{13}\text{FN}_2\text{O}_3$ ,  $[\text{M}]^+$  384.0910; found 384.0894.

#### 4.7. 2-(4-(5-(3-Fluorophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (5)

Solid; Yield 80%; m.p. 294–295 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.02 (d, *J* = 7.5 Hz, 2H), 87.96 (d, *J* = 7.5 Hz, 2H), 7.74–7.71 (m, 2H), 7.60 (d, *J* = 7.0 Hz, 1H), 7.42 (d, *J* = 8.5 Hz, 1H), 7.20–7.17 (m, 3H), 6.96–6.982 (m, 2H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.6, 164.4, 164.4, 163.5, 162.1, 135.1, 130.4, 128.0, 127.6, 127.5, 127.5, 127.3, 127.3, 127.1, 127.1, 125.9, 124.1, 123.6, 123.0, 116.0, 115.8, 115.3, 104.3; HR-EI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>13</sub>FN<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 384.0911; found 384.0892.

#### 4.8. 2-(4-(5-(2-Fluorophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (6)

Solid; Yield 78%; m.p. 291–292 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.20 (t, *J* = 7.0 Hz, 1H), 8.03 (d, *J* = 8.0 Hz, 2H), 7.97 (d, *J* = 8.0 Hz, 2H), 7.72 (dd, *J* = 7.0, 20 Hz, 1H), 7.44 (d, *J* = 8.5 Hz, 1H), 7.25 (t, *J* = 8.0 Hz, 1H), 7.22–7.18 (m, 3H), 6.97–6.95 (m, 2H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.5, 164.4, 164.4, 163.5, 158.2, 156.1, 135.0, 130.1, 129.0, 127.2, 127.2, 126.8, 126.8, 125.6, 125.2, 125.0, 124.9, 123.8, 123.4, 123.3, 116.0, 114.5, 104.4; HR-EI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>13</sub>FN<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 384.0912; found 384.0895.

#### 4.9. 2-(4-(5-(*p*-Tolyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (7)

Solid; Yield 83%; m.p. 270–271 °C, <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.03 (d, *J* = 8.0 Hz, 2H), 7.96 (d, *J* = 8.0 Hz, 2H), 7.74–7.71 (m, 3H), 7.25 (d, *J* = 8.0 Hz, 2H), 7.23–7.20 (m, 2H), 6.97–6.95 (m, 2H), 2.43 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.4, 164.6, 164.6, 163.5, 156.0, 142.1, 135.1, 131.5, 130.2, 127.3, 127.3, 127.2, 127.2, 126.8, 126.8, 126.2, 126.2, 125.6, 125.2, 123.7, 123.5, 116.0, 104.3, 21.2; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 380.1161; found 380.1145.

#### 4.10. 2-(4-(5-(*m*-Tolyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (8)

Solid; Yield 85%; m.p. 264–265 °C, <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.03 (m, 4H), 7.72(s, 1H), 7.56 (s, 1H), 7.51 (d, *J* = 7.0 Hz, 1H), 7.42 (s, 1H), 7.32 (t, *J* = 7.0 Hz, 1H), 7.24 (d, *J* = 7.0 Hz, 1H), 7.16 (t, *J* = 8.5 Hz, 1H), 6.95(d, *J* = 7.5 Hz, 1H), 6.89 (t, *J* = 7.5 Hz, 1H), 2.39 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.5, 164.5, 164.5, 163.6, 156.2, 139.0, 135.2, 130.3, 129.1, 129.0, 128.8, 127.3, 127.3, 126.7, 126.7, 126.0, 125.9, 125.2, 124.5, 123.9, 123.2, 116.0, 104.3, 21.5; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 380.1160; found 380.1143.

#### 4.11. 2-(4-(5-(*o*-Tolyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (9)

Solid; Yield 80%; m.p. 260–261 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.02–7.97 (m, 5H), 7.31–7.28 (m, 4H), 7.85 (d, *J* = 7.0 Hz, 1H), 7.16 (t, *J* = 7.0 Hz, 1H), 6.94 (s, 1H), 6.88 (t, *J* = 7.0 Hz, 1H), 2.44 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.3, 164.3, 164.3, 163.5, 156.1, 137.1, 136.8, 135.1, 130.2, 129.4, 128.5, 127.3, 127.3, 127.2, 126.7, 126.7, 126.1, 125.8, 125.2, 123.6, 123.2, 116.0, 104.4, 18.5; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 380.1161; found 380.1141.

#### 4.12. 2-(4-(5-(4-Nitrophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (10)

Solid; Yield 85%; m.p. 304–305 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.29–8.25 (m, 2H), 8.03–7.99 (m, 6H), 7.72 (d, *J* = 7.5 Hz, 1H), 7.34 (s, 1H), 7.17 (t, *J* = 7.0 Hz, 1H), 6.94 (d, *J* = 8.0 Hz, 1H), 6.89 (t, *J* = 7.0 Hz, 1H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.4, 164.4, 164.4, 163.4, 156.0, 147.8, 135.1, 132.0, 131.0, 131.0, 130.2, 128.6, 128.6, 127.2, 127.2, 127.0, 127.0, 125.6, 125.2, 123.7, 123.3, 116.0, 104.5; HR-EI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>13</sub>N<sub>3</sub>O<sub>5</sub>, [M]<sup>+</sup> 411.0855; found 411.0839.

#### 4.13. 2-(4-(5-(3-Nitrophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (11)

Solid; Yield 82%; m.p. 296–297 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.56–8.52 (m, 2H), 8.27(d, *J* = 7.0 Hz, 1H), 8.14 (d, *J* = 7.0 Hz, 1H), 8.03–7.98 (m, 3H), 7.74–7.71 (m, 2H), 7.35 (s, 1H), 7.15 (t, *J* = 7.5 Hz, 1H), 6.94 (d, *J* = 7.0 Hz, 1H), 6.88 (t, *J* = 7.0 Hz, 1H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.5, 164.4, 164.4, 163.4, 156.0, 148.3, 135.1, 133.8, 130.2, 130.1, 127.4, 127.4, 127.0, 126.8, 126.8, 125.8, 125.2, 123.8, 123.8, 123.2, 122.6, 116.0, 104.4; HR-EI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>13</sub>N<sub>3</sub>O<sub>5</sub>, [M]<sup>+</sup> 411.0855; found 411.0837.



**4.14. 2-(4-(5-(2-Nitrophenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (12)**

Solid; Yield 84%; m.p. 287–288 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.15–8.06 (m, 6H), 7.90–7.70 (m, 3H), 7.32 (s, 1H), 7.15 (t, *J* = 7.0 Hz, 1H), 6.94 (d, *J* = 7.0 Hz, 1H), 6.86 (t, *J* = 7.0 Hz, 1H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.6, 164.5, 164.5, 163.5, 156.1, 146.8, 135.2, 135.1, 131.5, 130.2, 129.8, 128.2, 127.3, 127.3, 126.8, 126.8, 125.6, 125.2, 124.2, 123.6, 123.1, 116.0, 104.3; HR-EI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>13</sub>N<sub>3</sub>O<sub>5</sub>, [M]<sup>+</sup> 411.0855; found 411.0833.

**4.15. 2-(4-(5-(4-Methoxyphenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (13)**

Solid; Yield 88%; m.p. 307–308 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.08 (br. s, 4H), 7.75–7.70 (m, 3H), 7.33 (s, 1H), 7.18 (t, *J* = 7.0 Hz, 1H), 7.01 (d, *J* = 8.5 Hz, 2H), 6.94 (d, *J* = 7.5 Hz, 1H), 6.88 (t, *J* = 7.5 Hz, 1H), 3.80 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.4, 164.4, 164.4, 163.5, 160.5, 156.1, 135.1, 130.3, 129.0, 127.2, 127.2, 126.8, 126.8, 125.6, 125.2, 123.8, 123.2, 116.0, 115.8, 115.8, 114.6, 114.6, 104.4, 56.1; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>4</sub>, [M]<sup>+</sup> 396.1110; found 396.1095.

**4.16. 2-(4-(5-(3-Methoxyphenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (14)**

Solid; Yield 84%; m.p. 299–300 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.11 (d, *J* = 7.0 Hz, 2H), 8.03 (d, *J* = 7.0 Hz, 2H), 7.72 (d, *J* = 6.0 Hz, 1H), 7.60 (s, 1H), 7.36–7.32 (m, 2H), 7.24–7.20 (m, 2H), 6.98 (d, *J* = 7.0 Hz, 1H), 6.95–6.93 (m, 2H), 3.92 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.5, 164.3, 164.3, 163.5, 161.0, 156.0, 135.0, 130.2, 130.1, 130.0, 127.2, 127.2, 126.6, 126.6, 125.6, 125.2, 123.7, 123.2, 119.6, 116.0, 114.2, 111.2, 104.3, 55.4; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>4</sub>, [M]<sup>+</sup> 396.1110; found 396.1093.

**4.17. 2-(4-(5-(2-Methoxyphenyl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (15)**

Solid; Yield 84%; m.p. 286–287 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.64 (s, 1H), 8.05–8.00 (m, 4H), 7.72 (d, *J* = 7.5 Hz, 1H), 7.53 (d, *J* = 7.0 Hz, 1H), 7.34 (s, 1H), 7.30 (t, *J* = 7.0 Hz, 1H), 7.15 (t, *J* = 7.0 Hz, 1H), 6.95–6.90 (m, 4H) 3.94 (s, 3H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.3, 164.3, 163.5, 162.5, 157.2, 156.0, 135.1, 130.2, 130.0, 129.6, 127.2, 127.2, 126.4, 126.4, 125.1, 125.6, 123.5, 123.1, 121.3, 116.5, 116.0, 112.5, 104.2, 56.0; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>4</sub>, [M]<sup>+</sup> 396.1109; found 396.1090.

**4.18. 2-(4-(5-(Pyridin-2-yl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (16)**

Solid; Yield 80%; m.p. 270–271 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.30 (d, *J* = 7.0 Hz, 1H), 8.10–8.02 (m, 3H), 7.99–7.96 (m, 2H), 7.90 (t, *J* = 7.0 Hz, 1H), 7.70 (d, *J* = 7.0 Hz, 1H), 7.42 (d, *J* = 7.0 Hz, 1H), 7.24–7.20 (m, 2H), 6.96–6.92 (m, 2H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.2, 164.3, 164.3, 163.3, 157.2, 156.0, 149.1, 137.1, 135.0, 130.1, 127.2, 127.2, 126.4, 126.4, 125.6, 125.1, 124.4, 124.0, 123.5, 123.2, 116.0, 104.2; HR-EI-MS: *m/z* Calcd for C<sub>22</sub>H<sub>13</sub>N<sub>3</sub>O<sub>3</sub>, [M]<sup>+</sup> 367.0957; found 367.0941.

**4.19. 2-(4-(5-(Pyridin-3-yl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (17)**

Solid; Yield 82%; m.p. 280–281 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.58 (d, *J* = 5.0 Hz, 1H), 8.44–8.40 (m, 2H), 8.05–8.00 (m, 4H), 7.75 (d, *J* = 6.0 Hz, 1H), 7.52 (d, *J* = 6.0 Hz, 1H), 7.23–7.20 (m, 2H), 6.94–6.90 (m, 2H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.3, 164.3, 164.3, 163.3, 156.0, 152.6, 147.8, 135.0, 134.0, 130.1, 127.2, 127.2, 126.4, 126.4, 125.6, 125.1, 124.3, 124.0, 123.5, 123.2, 116.0, 104.2; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>4</sub>, [M]<sup>+</sup> 367.0957; found 367.0942.

**4.20. 2-(4-(5-(Pyridin-4-yl)-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (18)**

Solid; Yield 84%; m.p. 288–289 °C; <sup>1</sup>H-NMR (MeOD-*d*<sub>4</sub>, 500 MHz): δ 8.60 (d, *J* = 6.0 Hz, 2H), 8.04 (d, *J* = 6.0 Hz, 2H), 7.98 (d, *J* = 8.0 Hz, 2H), 7.84 (d, *J* = 6.0 Hz, 2H), 7.70 (dd, *J* = 7.0 Hz, 1.5 Hz, 1H), 7.22 (s, 1H), 7.18 (t, *J* = 7.0 Hz, 1H), 6.96–6.94 (m, 2H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.4, 164.2, 164.2, 163.2, 156.1, 149.7, 149.7, 143.5, 135.1, 130.2, 127.2, 127.2, 126.4, 126.4, 125.4, 125.1, 123.6, 123.2, 121.1, 121.1, 116.0, 104.2; HR-EI-MS: *m/z* Calcd for C<sub>24</sub>H<sub>16</sub>N<sub>2</sub>O<sub>4</sub>, [M]<sup>+</sup> 367.0957; found 367.0931.

#### 4.21. 2-(4-(5-Phenyl-1,3,4-oxadiazol-2-yl)phenyl)-4H-chromen-4-one (19)

Solid; Yield 80%; m.p. 280–281 °C; <sup>1</sup>H-NMR (DMSO-*d*<sub>6</sub>, 500 MHz): δ 8.39 (s, 1H), 8.8–8.03 (m, 6H), 7.72 (s, 1H), 7.23 (t, *J* = 7.0 Hz, 1H), 7.22–7.18 (m, 2H), 7.09 (d, *J* = 7.0 Hz, 1H), 6.94 (d, *J* = 7.0 Hz, 1H), 6.90 (t, *J* = 7.0 Hz, 1H), 6.80 (d, *J* = 7.0 Hz, 1H); <sup>13</sup>C-NMR (125 MHz, DMSO-*d*<sub>6</sub>): δ 177.2, 164.1, 164.1, 163.2, 156.0, 135.0, 130.1, 129.0, 129.0, 128.5, 127.3, 127.3, 127.1, 127.1, 126.3, 126.3, 125.4, 123.4, 123.1, 121.3, 116.0, 112.5, 104.2; HR-ESI-MS: *m/z* Calcd for C<sub>23</sub>H<sub>14</sub>N<sub>2</sub>O<sub>3</sub>, [M]<sup>+</sup> 366.1004; found 366.0990.

#### 4.22. β-Glucuronidase Assay

β-Glucuronidase activity was determined in accordance to method used by Taha et al. [43,44] by measuring absorbance at 405 nm of the *p*-nitrophenol formed substrate by a spectrophotometric method. The total reaction volume was 250 μLs. Reaction mixtures containing 5 μL of test compound solution, 185 μL of 0.1M acetate buffer and 10 μL of enzyme solution were incubated for 30 min at 37 °C. At 405 nm the plates were recorded on a multiplate reader (SpectraMax, Molecular Devices, Sunnyvale, CA, USA) after the addition of 50 μL of 0.4 mM *p*-nitrophenyl-β-D-glucuronide. Experiments were performed in triplicate [45]. To avoid precipitation, compound concentrations were decreased, and the volume of reaction was increased (200 μL). Precipitation probability was less thus addition of detergents was not needed.

#### 4.23. Molecular Docking Studies Assay

In silico binding interactions of synthesized derivatives against human-β-glucuronidase were inspected using the Autodock Vina software [46]. The X-ray crystal structure of human-β-glucuronidase (PDB ID: 1bhg) was retrieved from the Protein Data Bank [<http://www.rcsb.org/>]. Water molecules, ligands and Chain-B were removed from the downloaded PDB file by means of Discovery Studio Visualizer [47]. Two-dimensional structures of synthesized compounds were sketched and their energies were optimized using ChemBio3D Ultra software (Version; 14.0.0.117) (Cambridge Soft, PerkinElmer Inc., Boston, MA, USA). Both receptor and ligand files were changed into pdbqt format by adding charges “q” and autodock type “t” using Autodock Tools (ADT) [48]. Other parameters contained within configurational file were defined as follow; Grid Box of size 19 × × 19 Å centered at X = 81.25, Y = 85.15 and Z = 87. 35 so that the active site [49] is entirely enclosed, maximum number of modes were set 50, energy difference between finest and poorest binding mode was set 4 Kcal/mol and exhaustiveness = 32. Conformations with lowest free energy of binding were evaluated and visualized using Discovery Studio Visualizer.

**Author Contributions:** Conceptualization, M.T. and F.R.; methodology, M.T.; software, M.A.A., M.G. and Y.A.B. validation, M.G., M.A.A. and Y.A.B.; formal analysis, M.A.; investigation, M.N.K.; resources, R.K.F.; data curation, R.K.F. and S.A.A.S.; Writing—Original Draft preparation, M.T. and Z.A.Z.; Writing—Review and Editing, Q.U.A. and Z.A.Z.; visualization, S.A.A.S.; supervision, M.T.; project administration, Z.A.Z. and M.T.; funding acquisition, M.T. and Q.U.A.

**Funding:** This research was funded by Deanship Scientific Research Imam Abdulrahman Bin Faisal university for this study; grant number [2018-066-IRMC].

**Acknowledgments:** The authors would like to Knowledge Institute for Research and Medical Consultations (IRMC), Imam Abdulrahman Bin Faisal University, for providing excellent lab facilities for research.

**Conflicts of Interest:** The authors declare no conflict of interest.

## References

1. Fishman, H.W. *Beta-Glucuronidase*; Academic Press: New York, NY, USA, 1983; p. 929.
2. Khan, K.M.; Ambreen, N.; Taha, M.; Halim, S.A. Structure-based design, synthesis and biological evaluation of β-glucuronidase inhibitors. *J. Comput. Aided Mol. Des.* **2014**, *28*, 577–585. [[CrossRef](#)] [[PubMed](#)]
3. Kunert-Keil, C.; Ritter, C.A.; Kroemer, H.K.; Sperker, B. Deconjugating enzymes: Sulphatases and glucuronidases. In *Enzyme Systems that Metabolise Drugs and Other Xenobiotics*; Ioannides, C., Ed.; John Wiley & Sons: Hoboken, NJ, USA, 2002; Volume 2, p. 521.

4. Gonick, H.C.; Kramer, H.J.; Schapiro, A.E. Urinary  $\beta$ -Glucuronidase Activity in Renal Disease. *Arch. Intern. Med.* **1973**, *132*, 63–69. [[CrossRef](#)]
5. Perez, J.L.; Berrocal, C.I.; Berrocal, L. Evaluation of a commercial beta-glucuronidase test for the rapid and economical identification of Escherichia coli. *J. Appl. Bacteriol.* **1986**, *61*, 541–545. [[CrossRef](#)] [[PubMed](#)]
6. Zeng, L.H.; Zhang, H.D.; Xu, C.J.; Bian, Y.J.; Xu, X.J.; Xie, Q.M.; Zhang, R.H. Neuroprotective effects of flavonoids extracted from licorice on kainate-induced seizure in mice through their antioxidant properties. *J. Zhejiang Univ. Sci. B* **2013**, *14*, 1004–1012. [[CrossRef](#)] [[PubMed](#)]
7. Pantscheva-Haschen, R.; Schulze, R.; Haerting, J.; Langkopf, B. An attempt to optimize the diagnosis of rejection after allogeneic kidney transplantation using urinary enzymes and multivariate statistics. *Med. Lab. Diagn.* **1988**, *29*, 179–187.
8. Paigen, K.; Peterson, J.; Paigen, B. Role of Urinary  $\beta$ -Glucuronidase in Human Bladder Cancer. *Cancer Res.* **1984**, *44*, 3620–3623. [[PubMed](#)]
9. Boyland, E.; Gasson, J.E.; Williams, D.C. Enzyme Activity in Relation to Cancer. *Br. J. Cancer* **1957**, *11*, 578–589. [[CrossRef](#)]
10. Gloux, K.; Barteau, O.; Eloumami, H.; Beguet, F.; Leclerc, M.; Dore, J. A metagenomic  $\beta$ -glucuronidase uncovers a coreadaptive function of the human intestinal microbiome. *Proc. Natl. Acad. Sci. USA* **2011**, *108*, 4539–4546. [[CrossRef](#)]
11. Zhu, Q.; Gao, R.; Wu, W.; Qin, H. The role of gut microbiota in the pathogenesis of colorectal cancer. *Tumor Biol.* **2013**, *34*, 1285–1300. [[CrossRef](#)]
12. Kim, D.H.; Jin, Y.H. Intestinal bacterial  $\beta$ -glucuronidase activity of patients with colon cancer. *Arch. Pharm. Res.* **2001**, *24*, 564–567. [[CrossRef](#)]
13. Huang, W.; Liu, M.-Z.; Li, Y.; Tan, Y.; Yang, G.-F. Design, syntheses, and antitumor activity of novel chromone and aurone derivatives. *Bioorg. Med. Chem.* **2007**, *15*, 5191–5197. [[CrossRef](#)]
14. Awadallah, F.M.; El-Waei, T.A.; Hanna, M.M.; Abbas, S.E.; Ceruso, M.; Oz, B.E.; Guler, O.O.; Supuran, C.T. Synthesis, carbonic anhydrase inhibition and cytotoxic activity of novel chromone-based sulfonamide derivatives. *Eur. J. Med. Chem.* **2015**, *96*, 425–435. [[CrossRef](#)]
15. Dias, M.M.; Machado, N.F.; Marques, M.P. Dietary chromones as antioxidants—The structural variable. *Food Funct.* **2011**, *2*, 595. [[CrossRef](#)]
16. Gomes, A.; Fernandes, E.; Silva, A.M.S.; Pinto, D.C.G.A.; Santos, C.M.M.; Cavaleiro, J.A.S.; Lima, J.L.F.C. Anti-inflammatory potential of 2-styrylchromones regarding their interference with arachidonic acid metabolic pathways. *Biochem. Pharmacol.* **2009**, *78*, 171–177. [[CrossRef](#)] [[PubMed](#)]
17. Prakash, O.; Kumar, R.; Parkash, V. Synthesis and antifungal activity of some new 3-hydroxy-2-(1-phenyl-3-aryl-4-pyrazolyl) chromones. *Eur. J. Med. Chem.* **2008**, *43*, 435–440. [[CrossRef](#)]
18. Farshori, N.N.; Banday, M.R.; Ahmad, A.; Khan, A.U.; Rauf, A. Synthesis, characterization, and in vitro antimicrobial activities of 5-alkenyl/hydroxyalkenyl-2-phenylamine-1,3,4-oxadiazoles and thiadiazoles. *Bioorg. Med. Chem. Lett.* **2010**, *20*, 1933–1938. [[CrossRef](#)] [[PubMed](#)]
19. Cui, Z.N.; Shi, Y.X.; Zhang, L.; Ling, Y.; Li, B.J.; Nishida, Y.; Yang, X.L. Synthesis and Fungicidal Activity of Novel 2,5-Disubstituted-1,3,4-oxadiazole Derivatives. *J. Agric. Food Chem.* **2012**, *60*, 11649–11656. [[CrossRef](#)]
20. Li, Y.H.; Zhu, H.J.; Chen, K.; Liu, R.; Khallaf, A.; Zhang, X.N.; Ni, J.P. Synthesis, insecticidal activity, and structure–activity relationship (SAR) of anthranilic diamides analogs containing oxadiazole rings. *Org. Biomol. Chem.* **2013**, *11*, 3979–3988. [[CrossRef](#)]
21. Akano, Y.; Shiga, F.; Asano, J.; Hori, W.; Anraku, T.; Uno, T. Synthesis and AMPA receptor antagonistic activity of a novel 7-imidazolyl-6-trifluoromethyl quinoxalinecarboxylic acid with a substituted phenyl group and improved its good physicochemical properties by introduced CF<sub>3</sub> group. *J. Bioorg. Med. Chem.* **2004**, *14*, 5107–5111.
22. Bansal, S.; Bala, M.; Suthar, S.S.; Choudhary, S.; Bhattacharya, S.; Bhardwaj, V.; Singla, S.; Joseph, A. Design and synthesis of novel 2-phenyl-5-(1,3-diphenyl-1H-pyrazz; fol-4-yl)-1,3,4-oxadiazoles as selective COX-2 inhibitors with potent anti-inflammatory activity. *Eur. J. Med. Chem.* **2014**, *80*, 167–174. [[CrossRef](#)] [[PubMed](#)]
23. Kucukguzel, S.G.; Oruc, E.E.; Rollas, S.; Sahin, F.; Ozbek, A. Synthesis, characterisation and biological activity of novel 4-thiazolidinones, 1,3,4-oxadiazoles and some related compounds. *Eur. J. Med. Chem.* **2002**, *37*, 197–206. [[CrossRef](#)]
24. Bondock, S.; Adel, S.; Etman, H.A.; Badria, F.A. Synthesis and antitumor evaluation of some new 1,3,4-oxadiazole-based heterocycles. *Eur. J. Med. Chem.* **2012**, *48*, 192–199. [[CrossRef](#)]

25. Husain, A.; Ajmal, M. Synthesis of novel 1,3,4-oxadiazole derivatives and their biological properties. *Acta Pharm.* **2009**, *59*, 223–233. [[CrossRef](#)] [[PubMed](#)]
26. Rahim, F.; Ullah, H.; Javid, M.T.; Wadood, A.; Taha, M.; Ashraf, M.; Shaukat, A.; Junaid, M.; Hussain, S.; Rehman, W.; et al. Synthesis, in vitro evaluation and molecular docking studies of thiazole derivatives as new inhibitors of  $\alpha$ -glucosidase. *Bioorg. Chem.* **2015**, *62*, 15–21. [[CrossRef](#)]
27. Ullah, H.; Rahim, F.; Taha, M.; Uddin, I.; Wadood, A.; Ali Shah, S.A.; Farooq, R.K.; Nawaz, M.; Wahab, Z.; Khan, K.M. Synthesis, molecular docking study and in vitro thymidine phosphorylase inhibitory potential of oxadiazole derivatives. *Bioorg. Chem.* **2018**, *78*, 58–67. [[CrossRef](#)]
28. Javid, M.T.; Rahim, F.; Taha, M.; Rehman, H.U.; Nawaz, M.; Wadood, A.; Imran, S.; Uddin, I.; Mosaddik, A.; Khan, K.M. Synthesis, in vitro  $\alpha$ -glucosidase inhibitory potential and molecular docking study of thiazole analogs. *Bioorg. Chem.* **2018**, *78*, 201–209. [[CrossRef](#)]
29. Taha, M.; Ismail, N.H.; Imran, S.; Selvaraj, M.; Rashwan, H.; Farhanah, F.U.; Rahim, F.; Kesavanarayanan, K.S.; Ali, M. Synthesis of benzimidazole derivatives as potent  $\beta$ -glucuronidase inhibitors. *Bioorg. Chem.* **2015**, *61*, 36–44. [[CrossRef](#)]
30. Taha, M.; Shah, S.A.A.; Afifi, M.; Imran, S.; Sultan, S.; Rahim, F.; Khan, K.M. Synthesis,  $\alpha$ -glucosidase inhibition and molecular docking study of coumarin based derivatives. *Bioorg. Chem.* **2018**, *77*, 586–592. [[CrossRef](#)]
31. Taha, M.; Shah, S.A.A.; Afifi, M.; Imran, S.; Sultan, S.; Rahim, F.; Ismail, N.H.; Khan, K.M. Synthesis, molecular docking study and thymidine phosphorylase inhibitory activity of 3-formylcoumarin derivatives. *Bioorg. Chem.* **2018**, *78*, 17–23. [[CrossRef](#)]
32. Taha, M.; Ismail, N.H.; Imran, S.; Wadood, A.; Rahim, F.; Khan, K.M.; Riaz, M. Hybrid benzothiazole analogs as antiurease agent: Synthesis and molecular docking studies. *Bioorg. Chem.* **2016**, *66*, 80–87. [[CrossRef](#)]
33. Hu, C.; Solomon, V.R.; Ulibarri, G.; Lee, H. The efficacy and selectivity of tumor cell killing by Akt inhibitors are substantially increased by chloroquine. *Bioorg. Med. Chem.* **2008**, *16*, 7888–7893. [[CrossRef](#)]
34. Hu, C.; Solomon, V.R.; Cano, P.; Lee, H. A 4-aminoquinoline derivative that markedly sensitizes tumor cell killing by Akt inhibitors with a minimum cytotoxicity to non-cancer cells. *Eur. J. Med. Chem.* **2010**, *45*, 705–709. [[CrossRef](#)]
35. Keith, C.T.; Borisy, A.A.; Stockwell, B.R. Multicomponent therapeutics for networked systems. *Nat. Rev. Drug Discov.* **2005**, *4*, 71–78. [[CrossRef](#)] [[PubMed](#)]
36. Mayur, Y.C.; Peters, G.J.; Prasad, V.V.S.R.; Lemos, C.; Sathish, N.K. Design of New Drug Molecules to be Used in Reversing Multidrug Resistance in Cancer Cells. *Curr. Cancer Drug Targets* **2009**, *9*, 298–306. [[CrossRef](#)]
37. Solomon, V.R.; Hu, C.; Lee, H. Hybrid pharmacophore design and synthesis of isatin–benzothiazole analogs for their anti-breast cancer activity. *Bioorg. Med. Chem.* **2009**, *17*, 7585–7592. [[CrossRef](#)]
38. Hubschwerlen, C.; Specklin, J.L.; Baeschlin, D.K.; Borer, Y.; Haefeli, S.; Sigwalt, C.; Schroeder, S.; Locher, H.H. Structure–activity relationship in the oxazolidinone–quinolone hybrid series: Influence of the central spacer on the antibacterial activity and the mode of action. *Bioorg. Med. Chem. Lett.* **2003**, *13*, 4229–4233. [[CrossRef](#)]
39. Bhanot, S.K.; Singh, M.; Chatterjee, N.R. The Chemical and Biological Aspects of Fluoroquinolones Reality and Dreams. *Curr. Pharm. Des.* **2001**, *7*, 311–335. [[CrossRef](#)]
40. Adamec, J.; Beckert, R.; Weiss, D.; Klimesova, V.; Waisser, K.; Mollmann, U.; Kaustova, J.; Buchta, V. Hybrid molecules of estrone: New compounds with potential antibacterial, antifungal, and antiproliferative activities. *Bioorg. Med. Chem.* **2007**, *15*, 2898–2906. [[CrossRef](#)] [[PubMed](#)]
41. Kouznetsov, V.V.; Gomez, A.-B. Recent developments in the design and synthesis of hybrid molecules based on aminoquinoline ring and their antiplasmodial evaluation. *Eur. J. Med. Chem.* **2009**, *44*, 3091–3113. [[CrossRef](#)] [[PubMed](#)]
42. Imran, S.; Taha, M.; Ismail, N.H.; Kashif, S.M.; Rahim, F.; Jamil, W.; Hariono, M.; Yusuf, M.; Wahab, H. Synthesis of novel flavone hydrazones: In-vitro evaluation of  $\alpha$ -glucosidase inhibition, QSAR analysis and docking studies. *Eur. J. Med. Chem.* **2015**, *105*, 156–170. [[CrossRef](#)]
43. Taha, M.; Ismail, N.H.; Imran, S.; Selvaraj, M.; Rahim, A.; Ali, M.; Siddiqui, S.; Rahim, F.; Khan, K.M. Synthesis of novel benzohydrazone-oxadiazole hybrids as  $\beta$ -glucuronidase inhibitors and molecular modeling studies. *Bioorg. Med. Chem.* **2015**, *23*, 7394–7404. [[CrossRef](#)] [[PubMed](#)]
44. Taha, M.; Ullah, H.; Muqarrabun, L.M.R.A.; Khan, M.N.; Rahim, F.; Ahmat, N.; Ali, M.; Perveen, S. Synthesis of bis-indolylmethanes as new potential inhibitors of  $\beta$ -glucuronidase and their molecular docking studies. *Eur. J. Med. Chem.* **2018**, *143*, 1757–1767. [[CrossRef](#)] [[PubMed](#)]

45. Jamil, W.; Perveen, S.; Shah, S.A.A.; Taha, M.; Ismail, N.H.; Perveen, S.; Ambreen, N.; Khan, K.M.; Choudhary, M.I. Phenoxyacetohydrazide Schiff Bases:  $\beta$ -Glucuronidase Inhibitors. *Molecules* **2014**, *19*, 8788–8802. [[CrossRef](#)] [[PubMed](#)]
46. Trott, O.; Olson, A.J. AutoDock Vina: Improving the speed and accuracy of docking with a new scoring function, efficient optimization, and multithreading. *J. Comput. Chem.* **2010**, *31*, 455–461. [[CrossRef](#)] [[PubMed](#)]
47. Dassault Systèmes BIOVIA, Discovery Studio Modeling Environment, Release 4.5, San Diego: Dassault Systèmes. 2015. Available online: <https://www.3dsbiovia.com/products/collaborative-science/biovia-discovery-studio/> (accessed on 20 December 2018).
48. Morris, G.M.; Huey, R.; Lindstrom, W.; Sanner, M.F.; Belew, R.K.; Goodsell, D.S. AutoDock4 and AutoDockTools4: Automated docking with selective receptor flexibility. *J. Comput. Chem.* **2009**, *30*, 2785–2791. [[CrossRef](#)]
49. Stierand, K.; Maass, P.C.; Rarey, M. Molecular complexes at a glance: Automated generation of two-dimensional complex diagrams. *Bioinformatics* **2006**, *22*, 1710–1716. [[CrossRef](#)]

**Sample Availability:** Samples of the compounds are available from the authors.



© 2019 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<http://creativecommons.org/licenses/by/4.0/>).