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Abstract: Background: Dipeptidyl peptidase-4 (DPP-4) inhibitors have emerged as anti-hyperglycemic agents that improve glycemic control in type 2 diabetic patients, either as monotherapy or in combination with other antidiabetic drugs. Methods: A novel series of dihydropyrimidine phthalimide hybrids was synthesized and evaluated for their in vitro and in vivo DPP-4 inhibition activity and selectivity using alogliptin as reference. Oral glucose tolerance test was assessed in type 2 diabetic rats after chronic treatment with the synthesized hybrids ± metformin. Cytotoxicity and antioxidant assays were performed. Additionally, molecular docking study with DPP-4 and structure activity relationship of the novel hybrids were also studied. Results: Among the synthesized hybrids, 10g, 10i, 10e, 10d and 10b had stronger in vitro DPP-4 inhibitory activity than alogliptin. Moreover, an in vivo DPP-4 inhibition assay revealed that 10g and 10i have the strongest and the most extended blood DPP-4 inhibitory activity compared to alogliptin. In type 2 diabetic rats, hybrids 10g, 10i and 10e exhibited better glycemic control than alogliptin, an effect that further supported by metformin combination. Finally, 10j, 10e, 10h and 10d had the highest radical scavenging activity in DPPH assay. Conclusions: Hybrids 10g, 10i and 10e are potent DPP-4 inhibitors which may be beneficial for T2DM treatment.

Keywords: DPP-4 inhibitors; T2DM; OGTT; DPPH; dihydropyrimidine; phthalimide

1. Introduction

Diabetes mellitus is a rapidly growing chronic disease that threatens human health worldwide. T2DM is characterized by insulin resistance and impaired glucose homeostasis, resulting in hyperglycemia [1]. Currently, there are a number of antidiabetic drugs including sulfonylureas, meglitinides, thiazolidinediones, biguanides, α-glucosidase inhibitors aimed at suppressing hepatic glucose output, stimulating insulin release, mitigating glucose absorption, and increasing peripheral glucose utilization [2,3]. Unfortunately, the use of these drugs is associated with various side effects including weight gain and hypoglycaemia which deteriorate the optimum glycemic control over a longer time [4,5]. Therefore, many newer approaches with novel mechanism of action have been emerged for better T2DM management. Among these newer approaches, dipeptidyl peptidase-4 (DPP-4) inhibitors that gained extensive interest in T2DM treatment with proved long term efficacy and better glycemic control. Additionally, DPP-4 inhibitors play an important role in regeneration and differentiation of pancreatic β-cells and also proved to be well tolerated as well as decreasing the risk of hypoglycemia and cardiovascular side effects [6].
DPP-4 is widely expressed in most tissues and body fluids including the biliary tract, kidney, gastrointestinal tract, uterus and liver [7]. It is the key regulatory enzyme and a signaling factor of incretin hormones (insulin-stimulating hormones, glucagon-like peptide (GLP-1), and glucose-dependent insulinotropic polypeptide (GIP)) [8]. These hormones play a crucial role in promoting β-cell proliferation, increasing insulin biosynthesis, reducing β-cell apoptosis [9,10]. As a consequence of rapid degradation by DPP-4, both GLP-1 and GIP have very short half-life which is 1–2 minutes for the former and 7 minutes for the latter [11,12]. Currently, there are several structurally diverse DPP-4 inhibitors some of them have been approved on the market such as sitagliptin, alogliptin, linagliptin, gemigliptin, anagliptin and teneligliptin [13,14].

On the other hand, pyrimidines represented one of the most important versatile scaffolds in medicinal chemistry [15] that possess many activities including antidiabetic [16], anti-inflammatory [17], anticancer [18], antibacterial [19], antihypertensive [20], antifungal [21] and antitubercular [22]. Moreover, some of the synthesized N-methylated and N-benzylated pyrimidinediones have shown in vitro DPP-4 inhibitory activity in significant range [23]. In addition, excellent IC₅₀ values were recorded for novel pyrimidinedione derivatives when evaluated for their DPP-4 inhibitory activity and in vivo anti-hyperglycemic efficacy [24].

On the other hand, phthalimide-containing compounds are precious synthetic scaffolds that showed different biological activities such as: antihyperglycemic [25], anti-inflammatory [26], antimicrobial [27], antihyperlipidimic [28], anticonvulsant [29], antiviral [30] and anticancer effects [31]. Additionally, a novel series of potent DPP-4 inhibitors bearing an N-phenylphthalimide skeleton has been synthesized and displayed higher potency compared to a well-known DPP-4 specific inhibitor [32,33]. Recently, some non-competitive DPP-4 inhibitors have been synthesized from a series of benzo[4,5]thieno[2,3-d]pyrimidine phthalimide derivatives and have shown better activity compared to DPP-4 inhibitors synthesized from amine precursors [34].

Based on the aforementioned information, we aimed at combining pyrimidine and phthalimide moieties into a single candidate template for the purpose of synergistic activity (Figure 1). To better elucidate their DPP-4 inhibitory activity, our synthesized hybrids were subjected to in vitro DPP inhibition assay utilizing human recombinant DPP-4, DPP-8 and DPP-9; alogliptin used as a reference standard. Moreover, the effect of our compounds on in vivo DPP-4 activity was assessed in SD rats, where DPP-4 activity was evaluated in the blood over 2 days using alogliptin as a reference standard. Additionally, to gain better understanding of the role of our compounds on T2DM, we studied the effect of chronic treatment of the synthesized hybrids 10a–j in presence and absence of metformin (MET) on HFD-induced T2DM in rats. Also, the target hybrids were tested for any possible antioxidant activity using DPPH method. Finally, the effect of synthesized hybrids 10a–j on cell viability was examined on normal hepatic LO2 cells using MTT assay, to check for any cytotoxic effect. Furthermore, structure activity relationship and molecular docking analysis of the titled hybrids were performed to get a distinct insight about the binding mode and interactions of the compounds with DPP-4 active site.
Figure 1. Diagram represents design of novel DPP-4 inhibitors.

2. Result and Discussion

2.1. Chemistry

The synthetic protocols applied to synthesize the target compounds 10a–j are illustrated in Scheme 1. Substituted acetoephene 1 reacted with various aromatic aldehydes 2 in 10% sodium hydroxide solution following Claisen-Schmidt condensation conditions to afford exclusively the E-chalcone isomers 3a–j in high yields [35]. These chalcones 3a–j were further reacted with thiourea (4) in the presence of sodium hydroxide in abs. ethanol to give 4,6-diaryl-3,4-dihydropyrimidine-2(1H)-thiones 5a–j. The structures of compounds 5e, g, i and j were confirmed on the basis of their spectral data (FT-IR, $^1$H-NMR, $^{13}$C-NMR, MS) and elemental analyses (see Section 4).

Reduction of phthalimide gave phthalide (6) which on refluxing with potassium phthalimido-$o$-toluic acid (8). Reaction of 8 with thionyl chloride brought about $\alpha$-phthalimido-$o$-toluyl chloride (9) [36]. Treatment of 4,6-diaryl-3,4-dihydropyrimidine-2(1H)-thiones 5a–j with 9 in dichloromethane, and in the presence of triethyl amine (TEA) as a base brought about the target compounds (4,6-diaryl-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoindolin-2-yl)methyl) benzothioates 10a–j. Notably, the S-substituted product was isolated as nucleophilicity of sulphur is greater than nitrogen and no N-substituted product was isolated. The structure of 10a–j hybrids were elucidated on the bases of both spectral data (IR, $^1$H-NMR, $^{13}$C-NMR, MS) and elemental analyses.

In the IR spectra the following peaks were observed: 3476–3384 cm$^{-1}$ (NH), 3071–2918 cm$^{-1}$ (C=C). Beside the δ values of OCH$_3$, CH$_3$, Nme$_2$, NH$_2$ side chains in the $^1$H-NMR spectra of the hybrids 10b, 10d, 10e, 10h and 10j which appear at 3.76, 2.34, 2.52, 6.19 and 6.17, respectively, the δ values of the other functional groups fall in the following ranges: NH: 11.52–11.19 ppm; Ar-H: 8.51–6.75 ppm; HC=CH-Ar: 6.68–6.35 ppm; CH-Ar: 5.29–5.27 ppm; CH$_2$: 5.08–5.01 ppm. Also, the $^{13}$C-NMR spectra of some selected compounds were resolved in appropriate regions confirming the assigned structures. Both mass spectra and elemental analyses of 10a–j hybrids are a further support of their structures by giving correct molecular formula and molecular ion peaks.

Reported non-competitive selective DPP-4 inhibitor
Scheme 1. Synthesis of the target compounds 10a–j. Reagents and conditions: (a) NaOH, dil. EtOH, rt; (b) NaOH, abs. EtOH; (c) DMF, reflux; (d) SOCl₂, reflux, dry benzene, distillation; (e) abs. dichloromethane, TEA, rt.

2.2. Biological Evaluation

2.2.1. In Vitro DPP Inhibition Assay

All the target hybrids listed in Table 1 were evaluated for the inhibition of human recombinant DPP-4 and alogliptin was chosen as a reference compound. Additionally, all
the synthesized hybrids were tested for the inhibition of DPP-8 and DPP-9 which is closely related to DPP-4 and whose inhibition is associated with great toxicity.

| Compound No. | DPP-4 IC$_{50}$ (nM) * | DPP-8 IC$_{50}$ (µM) * | DPP-9 IC$_{50}$ (µM) * |
|--------------|------------------------|------------------------|------------------------|
| 10a          | 19.74 ± 1.27           | >100                   | >100                   |
| 10b          | 3.78 ± 0.23            | >100                   | >100                   |
| 10c          | 14.71 ± 0.82           | >100                   | >100                   |
| 10d          | 2.19 ± 0.14            | >100                   | >100                   |
| 10e          | 1.42 ± 0.11            | >100                   | >100                   |
| 10f          | 31.52 ± 2.41           | >100                   | >100                   |
| 10g          | 0.51 ± 0.03            | >100                   | >100                   |
| 10h          | 10.37 ± 0.54           | >100                   | >100                   |
| 10i          | 0.66 ± 0.04            | >100                   | >100                   |
| 10j          | 7.35 ± 0.36            | >100                   | >100                   |
| Alogliptin   | 5.71 ± 0.21            | >100                   | >100                   |

* Data are means of four replicates with standard error (Mean ± SE).

Interestingly, hybrids 10g and 10i identified as the most potent DPP-4 inhibitors with picomolar potency, exhibiting IC$_{50}$ values of 0.51 and 0.66 nM, respectively; which represented 11.2, 8.6 fold superior activity than alogliptin. The potent DPP-4 inhibitory activity of these compounds may be attributed to electronic behavior of the nitro group in ring B of hybrid 10g and the small size of the furanyl group in ring B of hybrid 10i, which facilitated the formation of hydrogen bond with DPP-4 active site.

Of considerable interest, the mesomeric effect raised from 4-Me$_2$N group in ring B as well as 4-bromo substitutions in ring A of hybrid 10e markedly boosted DPP-4 inhibition resulted in low nanomolar potency activity (1.42 nM, 4 fold increased activity than that of alogliptin). Moreover, introducing an electron donating groups (methyl group in ring A and methoxy group in ring B) retained the potent DPP-4 inhibitory activity of hybrids 10b and 10d, whose activities were 1.5 fold and 2.6 fold the activity of alogliptin with IC$_{50}$ values of 3.78, 2.19 nM, respectively. According to the basis of electron donating properties, hybrid 10j expected to be the most potent DPP-4 inhibitor but on the contrary, the DPP-4 inhibitory potency is eroded (IC$_{50}$ = 7.35 nM). The probable reason could be the worse fit of the naphthyl group in ring B with more steric inhibition into the receptor pocket compared to the phenyl ring in the other synthesized compounds. On the other hand, the enzyme inhibition activity was dramatically decreased in the unsubstituted phenyl group and upon introducing the electron withdrawing chlorine atom in 10h, 10c and hybrids 10f which was confirmed by their high IC$_{50}$ values compared to alogliptin. Out of these, it might be concluded that the substituent electronic character and the steric volume of the phenyl ring considerably influenced the DPP-4 inhibition activity.

On the other hand, all the synthesized hybrids targeted DPP-8 and DPP-9 at IC$_{50}$ exceeding 100 µM which indicates that our compounds have superior selectivity for DPP-4 and neglected affinity toward DPP-8 and DPP-9.

### 2.2.2. Effects of Synthesized Hybrids (10a–j) on Cell Viability

The cytotoxic effect of the synthesized hybrids 10a–j was examined on normal hepatic LO2 cells using MTT assay. The test was done using 3 different concentrations (10 µM, 30 µM and 50 µM) for each compound. As shown in (Figure 2a), hybrid 10c had the most cytotoxic effect with 72.57% and 72.50% cell viability at 30 µM and 50 µM concentration, respectively. However, hybrid 10i came in the second place with cell viability percent of 74.60 at 10 µM concentration, followed by hybrid 10h with cell viability percent of 77.47 at 50 µM concentration, as presented in (Figure 2b). While, the cell viability percent for other synthesized hybrids ranged from 80.13 to 101.2. In conclusion, our compounds showed weak cytotoxic effect against hepatic LO2 cells.
for other synthesized hybrids ranged from 80.13 to 101.2. In conclusion, our compounds showed weak cytotoxic effect against hepatic LO2 cells.

**Figure 2.**

(a) Effect of hybrids 10a–e on viability of normal hepatic LO2 cells. (n = 5). (b) Effect of hybrids 10f–j on viability of normal hepatic LO2 cells (n = 5). *** Significant from control group at p < 0.001, ** Significant from control group at p < 0.01, * Significant from control group at p < 0.05.

### 2.2.3. Effect of Synthesized Hybrids (10a–j) on In Vivo DPP-4 Activity

The effect of the synthesized hybrids 10a–j on blood DPP-4 activity was investigated in SD rats, as shown in (Figure 3a,b). The hybrids were administrated in a single oral dose of 10 mg/kg and in vivo DPP-4 activity was evaluated over 2 days, using alogliptin as reference compound. Notably, among all tested hybrids and aloglipin, hybrid 10g has both the strongest and longest DPP-4 inhibitory action, with 18.45% and 47% DPP-4 blood activity at 12 h and 24 h, respectively, followed by hybrid 10i with DPP-4 blood activity of 18.8% and 49.9% at 12 h and 24 h, respectively. While, alogliptin achieved DPP-4 blood activity of 20.95% and 56.1% at 12 h and 24 h, respectively. Importantly, hybrids 10g and 10i also showed extended DPP-4 inhibitory activity at 48 h with blood DPP-4 activity of
73.3% and 76%, respectively, while, alogliptin DPP-4 blood activity was 97.05%. Worthily, hybrids 10g, 10i and 10e had the strongest in vitro and in vivo DPP-4 inhibiting activity.

Figure 3. (a). The in vivo DPP-4 activity of 10a–e hybrids and alogliptin within 48 h. (n = 3). (b) The in vivo DPP-4 activity of 10f–j hybrids and alogliptin within 48 h. (n = 3).

2.2.4. Effect of Chronic Treatment of Compounds 10a–j with or without MET on HFD-Induced Type 2 Diabetic rats

HFD significantly induced insulin resistance in SD rats as evident by extremely significant increase in the AUC of OGTT in non-treated diabetic group compared to control group, as shown in (Figure 4a,b). We studied the chronic effect of oral administration of hybrids 10a–j at a dose of 10 mg/kg/day in absence and presence of MET, on insulin resistance in type 2 diabetic rats. In absence of MET, hybrids 10g, 10i and 10e significantly improved glucose tolerance above alogliptin, as evident by the reduction in the AUC of OGTT when compared to non-treated diabetic group, (Figure 5a,b). Moreover, oral administration of MET (150 mg/kg/day) together with hybrids 10a–j, further enhanced insulin sensitivity with a profound reductions in AUC of OGTT almost in all treated groups.
Interestingly, MET addition to 10g, 10i and 10e-treated groups reduced AUC of OGTT by 22.93%, 21.7% and 22.82%, respectively. Accordingly, glucose tolerance in 10g/MET treated group reached a normal level with AUC equals 13787 ± 201 mg.min/dL compared to 14305 ± 318 mg.min/dL for normal control group. Similarly, addition of MET to alogliptin treated group reduced AUC of OGTT from 20835 ± 146 mg.min/dL in alogliptin treated group to 15451 ± 110 mg.min/dL in MET/alogliptin treated group.

Figure 4. (a) Chronic effect of hybrids 10a–j and alogliptin administration on blood glucose levels during an OGTT in type 2 diabetic rats. Data are presented as mean ± SEM (n = 7). (b) Chronic effect of combined administration of 10a–j/MET, alogliptin/MET on blood glucose levels during an OGTT in type 2 diabetic rats. Data are presented as mean ± SEM (n = 7).
2.2.5. Antioxidant Activity

The in vitro antioxidant activity of the synthesized hybrids 10a–j was tested depending on DPPH method using ascorbic acid as standard. The results in Table 2 revealed that all the synthesized compounds showed good antioxidant activity, wherein hybrids 10j, 10e, 10h and 10d have the highest radical scavenging activity in DPPH assay at 100 µM concentration.
Table 2. Antioxidant activity of hybrids 10a–j using the DPPH method.

| Compound No. (100 µM) | % DPPH Scavenging * |
|-----------------------|----------------------|
| 10a                   | 73.85 ± 2.46         |
| 10b                   | 74.28 ± 1.69         |
| 10c                   | 52.39 ± 0.93         |
| 10d                   | 80.16 ± 1.38         |
| 10e                   | 87.21 ± 2.58         |
| 10f                   | 80.16 ± 1.38         |
| 10g                   | 84.74 ± 2.76         |
| 10h                   | 69.11 ± 1.44         |
| 10i                   | 93.52 ± 1.65         |

* Data are means of four replicates with standard error (Mean ± SE).

2.2.6. Molecular Docking Study: Docking at the DPP-4 active site

To verify the protein-ligand binding properties and the key structural features required for the inhibitory activity of the novel synthesized hybrids 10a–j with the protein DPP-4, a molecular docking study was performed using the molecular docking program MOE. The results of energy binding scores and binding interactions were depicted in Table 3 and Figure 6. The 3D crystal structure of DPP-4 was obtained from the Protein Data Bank [PDB code 3G0B, resolution 2.25 Å (www.rcsb.org, accessed on 10 February 2021)] and alogliptin was used as a reference DPP-4 inhibitor. DPP-4 active site is formed of two main binding pockets Figures S1 and S2 in addition to Figure S2 extensive subsite pocket. Figure S1 pocket is formed of highly hydrophobic residues Ser 630, Tyr 631, Tyr 547, Val 656, Trp 659, Tyr 662, Tyr 666, Asn 710, Val 711 and His 740), Figure S2 pocket (formed by Glu 205 and Glu 206 dyad and Arg 125) [37]. The Figure S2 pocket is surrounded by residues of Val 207, Ser 209, Arg 358, Phe 357, Lys 544, Trp 627, Trp 629 and Asp 708 which constitute the Figure S2 extensive subsite [38].

Table 3. Types of binding interactions and energy scores (kcal/mol) for hybrids 10a–j and alogliptin at the DPP-4 active site.

| Compound No. | Types of Interactions                                                                 | Energy Scores |
|--------------|---------------------------------------------------------------------------------------|---------------|
| 10a          | - One hydrogen bonding interaction with Arg 125 (3.14 Å).                              | −8.68         |
| 10b          | - One π-cation interaction with Lys 554 (4.06 Å)                                      | −9.12         |
|              | - One π-hydrogen bonding interaction with Trp 629 (4.23 Å).                           |               |
| 10c          | - Two hydrogen bonding interaction with Arg 125 and Tyr 547 (3.18 Å, 2.77 Å).          | −7.74         |
|              | - One π-hydrogen bonding interaction with His 126 (4.03 Å).                           |               |
| 10d          | - Three hydrogen bonding interaction with Arg 125 (2.85 Å, 3.01 Å, 3.22 Å).            | −8.14         |
|              | - One hydrogen bonding interaction with Tyr 457 (2.71 Å).                              |               |
|              | - One π-hydrogen bonding interaction with Tyr 547 (4.20 Å).                            |               |
| 10e          | - Three hydrogen bonding interaction with Arg 125 (2.83 Å, 3.01 Å, 3.24 Å).            | −8.15         |
|              | - One hydrogen bonding interaction with Tyr 547 (2.73 Å).                              |               |
| 10f          | - One hydrogen bonding interaction with Asp 545 (3.14 Å).                              | −7.74         |
|              | - One π-cation interaction with Lys 554 (4.87 Å).                                     |               |
| 10g          | - Two hydrogen bonding interaction with Lys 554 and Glu 206 (2.97 Å and 3.09 Å).       | −9.45         |
| 10h          | - Two hydrogen bonding interaction with ASP 545 and VAL 546 (3.07 Å and 2.83 Å).       | −9.32         |
| 10i          | - Two hydrogen bonding interaction with Tyr 547 and Glu 206 (2.89 Å and 3.63 Å).       | −8.50         |
|              | - Two hydrogen bonding interaction with Arg 125 (3.34 Å and 2.91 Å).                   |               |
Table 3. Cont.

| Compound No. | Types of Interactions                                                                 | Energy Scores |
|--------------|----------------------------------------------------------------------------------------|---------------|
| 10j          | - Three hydrogen bonding interaction with Glu 205, Glu 206 and Tyr 631 (3.16 Å, 3.12 Å and 3.53 Å).<br>- One H-π interaction with Trp 629 (4.33 Å).<br>- One π-H interaction with Tyr 456 (4.21 Å). | −7.72         |
| Alogliptin   | - Five hydrogen bonding interaction with Glu 205, Glu 206, Tyr 631, Arg 125 (2.78 Å and 2.96 Å, 2.90 Å, 3.35 Å and 2.97 Å).<br>- Three ionic bond with Glu 206 and Glu 205 (2.96 Å, 2.90 Å and 3.95 Å). | −7.92         |

Figure 6. Cont.
Figure 6. Docking and binding pattern of compounds 10g (a,b), 10i (e,f) and alogliptin (g,h) showing interactions with different amino acid residues found in the active site of DPP-4 (PDB code: 3G0B).

The SiteMap analysis highlighted the importance of the Tyr 547, Lys 554, and Trp 629 residues as a top-ranked binding pocket of DPP-4 in the formation of the enzyme–inhibitor complex [39]. Moreover, the hydroxyl group of Tyr 547 plays an oxygen-stabilizing role in the DPP-4 catalytic mechanism as well as Trp 629 residue forms catalytic Figure S2 pocket in this protease [40]. The DPP-4 inhibitory activity is markedly increased when the inhibitor can interact to Figures S1 and S2 pockets. Interestingly, studies suggested that only few inhibitors can bind into Figure S2 extensive subsite and enhancing the bioactivity of DPP-4 inhibitors through the interactions with Lys 554 so it is the target of DPP-4 inhibitor selectivity [41].

It was obvious from the results depicted in Table 3 that nearly all of the novel hybrids achieved energy binding scores higher than that of alogliptin. Additionally, hybrids 10b, 10d, 10e, 10g, 10i bind efficiently with the key amino acids located in the receptor pockets which explained the potent DPP-4 inhibitory activity of these hybrids. Of considerable interest, hybrid 10g recorded the highest energy binding score and the lowest in vitro DPP-4 IC₅₀ inhibitory activity among the other synthesized hybrids via binding mainly to the Figure S2 pocket and Figure S2 extensive subsite through formation of two strong hydrogen bonds between both oxygen of the free nitro group and phthalimido carbonyl group with Glu 206 and Lys 554 which have a vital role in the enhancement of DPP-4 affinity. However, the binding mode showed that the phthalimido carbonyl oxygen and S atom of hybrid 10i participated in a hydrogen-bonding network with Arg 125 and Glu 206 residues in Figure S2 pocket together with another strong hydrogen bond formed with Tyr 547 in Figure S1 pocket. Similarly, compound 10d was buried in Figures S1 and S2 pockets via formation of four hydrogen bonds as well as π-H bond with Arg 125 and Tyr 547.

Notably, one hydrogen bond is formed between one of the phthalimido carbonyl oxygen of compound 10e and free OH group of Tyr 547 residue in Figure S1 pocket together with another three hydrogen bonds between the other carbonyl oxygen and the free NH₂ of Arg 125 in Figure S2 pocket. However, hybrid 10b stacked with the side chain of Lys 554 and Trp 629 in Figure S2 extensive subsite via weak π-cation and π-hydrogen interactions which explained its higher IC₅₀ value comparable to the previously mentioned hybrids but its DPP-4 inhibitory activity still higher than that of alogliptin. On the other hand, in Figure S2 pocket, the nitrogen of the free amino group of hybrid 10j is responsible for binding with Glu 205 and Glu 206 residues through two hydrogen bonds together with another hydrogen bond formed with Tyr 631 in Figure S1 pocket. Moreover, the phenyl ring of that compound provided additional binding affinity with Figure S2 extensive subsite via H-π interaction with Trp 629 residue while the naphthyl ring forms another π-H stacking with Tyr 456 side chain.

However, the electron withdrawing effect of chlorine atom in hybrid 10c negatively affects its binding with the receptor site and hence its DPP-4 inhibitory activity. In addition
hybrid 10a binds weakly with the receptor site via only one hydrogen bonding with Arg 125. Meanwhile, 10f and 10h exhibited weak DPP-4 inhibitory activity as they could not be able to bind to the crucial amino acids responsible for the activity. Finally, it can be concluded that most of the titled synthesized hybrids interact with DPP-4 protein in distinct manner and thus are able to inhibit its activity efficiently.

3. Structure Activity Relationship

Based on the aforementioned results, it becomes clear that the electronic nature of the substituents and the steric factor greatly affect the biological activity of the synthesized hybrids, so the present SAR study focused on the effect of substituents on position 4 in the aryl groups that located in positions 2 and 4 of dihydropyrimidine as in hybrids 10a–h, as well as the furanyl and 1-naphthyl groups in hybrids 10i and 10j. It is worth noting that positions 2 and 4 in dihydropyrimidine are equivalent due to the migration of hydrogens and electron transition which is cleared in (Figure 7).

Moreover, the structures of most of the title compounds 10a–j contain a 4-chlorophenyl group. Consequently, the compounds under investigation are related structurally to the substituents attached to the other phenyl group. On the basis of the electron donating effect of both the 1-naphthyl and 4-aminophenyl groups and the lipophilic character of the latter, hybrid 10j should be the top ranked electron donor and consequently expected to be the most potent DPP-4 inhibitor via formation of more strong H-bonds with the receptor binding sites but on the contrary, the steric factor of the 1-naphthyl and 4-aminophenyl groups hinder the efficient binding with the receptor binding site, indicating that the steric factor is more effective, but it is still potent enough for inhibiting the enzyme activity as its IC₅₀ was slightly less than that of alogliptin. Interestingly, hybrid 10g achieved the most potent DPP-4 inhibitory activity owing to the electron donating properties and the unique electronic ionic behavior of the nitro group leading to higher ability to form strong H-bond with Figure S2 extensive subsite as evidenced by its higher energy binding score.

Additionally, the DPP-4 inhibitory activity is markedly increased thanks to the lipophilic character and the small size of the furanyl group in hybrid 10i which enhance its fit to the receptor site, and accordingly its ability to form strong H-bond with the receptor site as evidenced by its molecular docking results. On the other hand, the lipophilic electron donating character of the methyl group, via hyperconjugative effect, in hybrid 10d resulted in an increase of its ability to form more H-bonds which, in turn, reflected on the marked difference in DPP-4 inhibitory IC₅₀ values compared to that of the corresponding electron-withdrawing chlorine-atom in hybrid 10c. Moreover, the mesomeric effect raised from 4-Me₂N group as well as 4-bromo atom in 10e hybrid seemed to be the most effective factor.
than hyperconjugative effect of 4-methyl and mesomeric effect of 4-chloro in hybrid 10d, that led to stabilization of H-bonding and consequently better DPP-4 inhibiting activity. Meanwhile, the lipophilic electron donating methoxy group of hybrid 10b experienced much higher DPP-4 inhibiting activity than the hydrophilic electron-withdrawing chlorine-atom hybrid 10f which exhibited much weaker binding interaction with the key amino acid residue in the receptor site as evidenced by its high IC\textsubscript{50} value. Finally, it is concluded from the aforementioned SAR results that the electronic nature, the steric factor and lipophilicity of the substituents greatly affected the biological activity of the system incorporated.

4. Materials and Methods

4.1. Animals

Male Sprague Dawley (SD) rats (6–7 weeks old and 100–120g body weight) were purchased from Abu-rawash Animal House (Giza, Egypt). Rats were allowed to acclimatize for one week with free access to food and water and kept under controlled laboratory conditions of humidity and 12 h light-dark cycle. All experimental protocols were approved by the Research Ethics Committee of the Faculty of Pharmacy, Port-said University (R 20.5.1527).

4.2. Experimental Design

After acclimatization period, rats were randomly divided into 25 groups, each of seven rats. The first group was normal control group which maintained on normal rat chow for the entire experiment period (16 weeks). The second group served as diabetic control group in which T2DM was induced by HFD that consisted of casein (30%), raw beef fat (Suet, 40%), wheat flour (7%), glucose (10%), salt mixture (6%), vitamin mixture (3%), and bran (4%), for 16 weeks. The remaining groups consumed HFD for 16 weeks, however, in the last four weeks, HFD was administrated together with each compound from our synthesized hybrids 10a–j at a dose of 10 mg/kg/day, orally, or with alogliptin (Takeda Pharmaceuticals, Deerfield, IL, USA) at a dose of 20 mg/kg/day, orally, with and without MET (Alfa Aesar, ThermoFisher Inc., Haverhill, MA, USA) at a dose of 150 mg/kg/day, orally.

4.3. In Vitro DPP-4, DPP-8 and DPP-9 Inhibition Assay

DPP-4 inhibition was assessed, as previously described \cite{42,43}, based on a continuous fluorometric assay method in which the substrate Gly-Pro-AMC (Gly-Pro-7-amido-4-methylcoumarin hydrobromide) was split by DPP-4 to release fluorescent aminomethyl coumarin (AMC). The liberated AMC was continuously monitored using an excitation wavelength of 360 nm and an emission wavelength of 460 nm using an EnVision microplate reader (PerkinElmer, Waltham, MA, USA). The reaction mixture consisted of different concentrations of the synthesized hybrids 10a–j, 20 µU/µL of recombinant human DPP-4 (Sigma Aldrich, St. Louis, MO, USA), 10 µM Gly-Pro-AMC (Sigma Aldrich, St. Louis, MO, USA) and assay buffer (150 mM NaCl, 25 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES), 0.12 mg/ml bovine serum albumin (BSA)) and pH of 7.5. The assay was carried out in quadruplicate. IC\textsubscript{50} was calculated using GraphPad Prism 8 software (San Diego, CA, USA). Similar to the DPP-4 assay, DPP-8 and DPP-9 inhibition assay was performed and the pH of the assay buffer was 8.

4.4. Cytotoxicity Test

We assessed any possible cytotoxic effect of our synthesized hybrids 10a–j on hepatic LO2 cells using MTT assay. The cells, in DMEM (Gibco, Invitrogen), were seeded in 96 well plates at density of 1 × 104/well. The synthesized compounds were added to the cells and incubated for 48 h at 37 °C and 5% CO\textsubscript{2}. Then, MTT reagent was added and the cells were re-incubated for further 4 h. After discarding the supernatants, the formed formazan crystals were dissolved in DMSO. The intensity of color was assessed using microplate reader (Bio-Rad Laboratories, Hercules, CA, USA) at 490 nm wavelength.
4.5. In Vivo DPP-4 Inhibition Assay

The assay was performed using male SD rats \( n = 3 \) per group, as previously described [44]. The synthesized hybrids 10a–j were suspended in 0.5 carboxymethyl cellulose (CMC) and administrated orally at a single dose of 10 mg/kg.

Blood samples were collected at time points 0, 6, 12, 24, 36 and 48 h. Plasma samples were extracted and kept at \(-20^\circ C\) until analysis time. Similar to the in vitro DPP-4 assay principle, plasma samples were mixed with equal volumes of Gly-Pro-AMC in assay buffer (100 mM HEPES, 0.1 mg/ml BSA) and pH of 7.5. The liberated AMC was continuously monitored using an excitation wavelength of 360 nm and an emission wavelength of 460 nm using EnVision microplate reader (PerkinElmer, Waltham, MA, USA). The data were expressed as% blood DPP-4 activity.

4.6. In Vivo Oral Glucose Tolerance Test (OGTT) in SD Rats

At the end of treatment regimen that lasted for four weeks; using our synthesized hybrids 10a–j with and without MET, rats were fasted overnight (12 h) with free access to water. Oral glucose tolerance test (OGTT) was performed as previously described [45]. Briefly, blood samples were collected from retro-orbital sinuses veins using glass capillaries at 0 time, 30, 60, 90 and 120 min after oral glucose loading (2 g/kg) and glucose concentration was determined with an automatic blood glucose meter (Accu-Chek; Roche Diagnostics, Rotkreuz, Switzerland).

4.7. Antioxidant Activity (DPPH Method)

DPPH (1,1-diphenyl-2-picrylhydrazyl) free radical is stable at rt and has a purple colour with a maximum absorption at 517 nm. This free radical is reduced by antioxidant molecules into 1,1-diphenyl-2-picrylhydrazine with resultant discoloration of free radical; this discoloration is stoichiometric with regard to the captured electrons. Accordingly, the fluctuation in the absorbance is an indicator for the antioxidant activity. Our synthesized hybrids 10a–j (100 µM) were mixed with DPPH (100 µM) and left for 20 min at rt, the absorbance was measured at 517 nm.

4.8. Statistical Analysis

Results were expressed as means ± standard error of the mean (SEM) and were analyzed for statistically significant differences using one-way analysis of variance (ANOVA). P values less than 0.05, 0.01 and 0.001 were considered significant. GraphPad Prism® 8 was used for statistical calculations (GraphPad Software, St. Louis, MO, USA).

4.9. Molecular Docking

Molecular docking studies of the synthesized hybrids 10a–j and alogliptin were performed using MOE software. Ligands were built into the builder interface of the MOE program and their energies were minimized until a root mean square deviation (RMSD) gradient of 0.01 kcal/mol and root mean square (RMS) distance of 0.1 Å with MMFF94× (Merck molecular force field 94×) force-field and the partial charges were automatically calculated. The X-ray crystallographic structure of DPP-4 (PDB code: 3G0B) was downloaded from protein data bank (www.rcsb.org, accessed on 10 February 2021). The enzymes were prepared, the hydrogens were added then the atoms connection and type were checked with automatic correction. The obtained poses were studied and the poses showed best ligand-enzyme interactions were selected and stored for energy calculations.

4.10. Chemistry

Melting points were determined by open tube capillary on a Mel-Temp hot stage apparatus (Stuart, Staffordshire, UK) on the Celsius scale and are uncorrected. Reactions were monitored by thin-layer chromatography on silica gel PF254 plates (Merck, Darmstadt, Germany) using dichloromethane/ethyl acetate (9:1) as mobile phase, and spots were detected with UV analyses lamp at 254 and 366 nm. IR spectra were recorded as KBr disks.
on a Fourier Transform (FT-IR) spectrophotometer (Bruker, Billerica, MA, USA). \textsuperscript{1}H-NMR and \textsuperscript{13}C-NMR spectra were recorded on a Bruker NMR spectrometer operating at 400 MHz and 100 MHz, respectively, using TMS as internal reference. Chemical shifts (δ values) are given in parts per million (ppm) using CDCl\textsubscript{3} (7.26) or DMSO-\textsubscript{d6} (2.49) as solvents and coupling constants (J) in Hz. Splitting patterns are designated as follows: s, singlet; bs, broad singlet; d, doublet; t, triplet; m, multiplet. GCMS: Mass spectra was recorded on a LCMS-QQQ instrument (Agilent, Santa Clara, CA, USA). Elemental analyses were performed on a 2400 CHN elemental analyzer (Perkin–Elmer, Waltham, MA, USA) and were within ± 0.4% of theoretical values unless otherwise specified.

4.11. Materials

Chemicals and solvents are of commercial grade, and purchased from Alfa Aesar (Haverhill, MA, USA), Cambrian Chemicals (Ontario, CA), Sigma-Aldrich (St. Louis, MO, USA), Acros Organics (Haverhill, MA, USA), Fluka (Gillingham, Dorset, UK), Merck (Ontario, CA), and El-Nasr Pharmaceutical Chemical Company (Cairo, Egypt). Chemicals were used without purification. Solvents were purified following the reported methods.

4.12. Synthesis

4.12.1. General Procedure for Synthesis of 4,6-diaryl-3,4-dihydropyrimidine-2(1H)-thiones 5d–g, i, j [53]

A mixture of 1,3-diaryl-2-propene-1-ones 3 (2.404 mmol), prepared by Claisen-Schmidt condensation between substituted acetophenones 1, benzaldehydes 2, and thiourea 4 (2.404 mmol) and NaOH (2.404 mmol) in absolute ethanol (25 mL) was heated under reflux for 4–6 h. The reaction progress was monitored by TLC using dichloromethane/ethyl acetate (9:1). The reaction mixture was cooled to rt and poured onto crushed ice. The solid product was filtered off, washed with water and crystallized from ethanol to yield the following pure titled compounds.

6-(4-Bromophenyl)-4-(4-(dimethylamino)phenyl)-3,4-dihydropyrimidine-2(1H)-thione (5e)

Yield: 0.633 g, 68% (orange solid); m.p. 112–3 °C; IR (KBr, cm\textsuperscript{-1}) v: 3195 (NH), 3084, 2925 (Ar-CH), 1679 (C=N), 1523, 1482 (C=C), 1174 (NH—C=C), 99.3 (NH—C=C), 55.5 (CHAr), 21.2 (CH\textsubscript{3}). GCMS m/z (rel. abundance): 388 (M.+ 100), 372 (21), 355 (60), 332 (17), 301 (16), 268 (65), 232 (31), 186 (21), 157 (23), 120 (17). Anal. Calcd. for \textsuperscript{13}C\textsubscript{18}H\textsubscript{18}BrN\textsubscript{2}S (388.32); C, 55.67; H, 4.67; N, 10.32; S, 8.26. Found: C, 55.93; H, 4.72; N, 10.11; S, 8.25.
6-(4-Chlorophenyl)-4-phenyl-3,4-dihydropyrimidine-2(1H)-thione (5f)
Yield: 0.384 g, 66% (white solid); m.p. 162–3 °C; IR (KBr, cm⁻¹) ν: 3188 (NH), 3028, 2979 (Ar-CH), 1671 (C=N), 7.70–7.27 (m, 17H, ArH), 6.72 (d, 1H, J = 4.8 Hz, H5), 5.24 (d, 1H, J = 4.8 Hz, H4). GCMS m/z (rel. abundance): 300 (M.+, 67), 285 (40), 254 (48), 225 (100), 219 (55), 189 (32), 161 (49), 137 (31), 103 (14). Anal. Calcd. for C₁₆H₁₃ClN₂S (300.81); C, 63.88; H, 4.36; N, 9.31; S, 10.66. Found: C, 64.06; H, 4.22; N, 10.41; S, 8.14.

6-(4-Chlorophenyl)-4-(4-nitrophenyl)-3,4-dihydropyrimidine-2(1H)-thione (5g)
Yield: 0.589 g, 71% (grey solid); m.p. 210–12 °C; IR (KBr, cm⁻¹) ν: 3155 (NH), 3070 (Ar-CH), 1678 (C=N), 1552, 1493 (C=C), 1177 (C=S). ¹H-NMR (CDCl₃, 400 MHz): δ 10.28 (s, 1H, 1NH), 9.72 (s, 1H, 1NH), 8.30 (d, 2H, J = 8.1 Hz, ArH), 8.10 (d, 2H, J = 8.1 Hz, ArH), 7.59 (d, 2H, J = 8.1 Hz, ArH), 7.42 (d, 2H, J = 8.1 Hz, ArH), 6.43 (d, 1H, J = 4.8 Hz, H5), 5.26 (d, 1H, J = 4.8 Hz, H4). ¹³C-NMR (CDCl₃, 100 MHz): δ 176.6 (CS), 144.5, 140.0, 135.0, 132.4, 131.2, 130.0, 128.4, 126.2, 123.8 (ArC), 101.9 (NH-C=C), 52.0 (CHAr). GCMS m/z (rel. abundance): 345 (M.+, 100), 312 (50), 310 (6), 270 (26), 255 (9), 223 (16), 190 (9), 176 (16), 134 (16), 122 (28). Anal. Calcd. for C₁₆H₁₂ClN₃O₂S (345.80); C, 55.57; H, 3.50; N, 12.15; S, 9.27. Found: C, 55.33; H, 3.62; N, 12.23; S, 9.15.

6-(4-Chlorophenyl)-4-(furan-2-yl)-3,4-dihydropyrimidine-2(1H)-thione (5i)
Yield: 0.509 g, 73% (yellow solid); m.p. 165–7 °C; IR (KBr, cm⁻¹) ν: 3187 (NH), 3104, 2984 (Ar-CH), 1676 (C=N), 1552, 1493 (C=C), 1177 (C=S). ¹H-NMR (CDCl₃, 400 MHz): δ 10.08 (s, 1H, 1NH), 9.86 (s, 1H, 1NH), 7.69–7.39 (m, 7H, 4 ArH & 3 furyl-H), 6.23 (d, 1H, J = 4.8 Hz, H5), 5.35 (d, 1H, J = 4.8 Hz, H4). GCMS m/z (rel. abundance): 290 (M.+, 100), 257 (32), 235 (15), 215 (32), 190 (16), 179 (42), 159 (23), 131 (13), 115 (5). Anal. Calcd. for C₁₄H₁₁ClN₂OS (290.77); C, 57.83; H, 3.81; N, 9.63; S, 11.03. Found: C, 58.14; H, 3.69; N, 9.48; S, 10.83.

6-(4-Aminophenyl)-4-(naphthalen-1-yl)-3,4-dihydropyrimidine-2(1H)-thione (5j)
Yield: 0.555 g, 73% (pale yellow solid); m.p. 155–6 °C; IR (KBr, cm⁻¹) ν: 3217 (NH), 3051, 2924 (Ar-CH), 1669 (C=N), 1559, 1464 (C=C), 1177 (C=S). ¹H-NMR (CDCl₃, 400 MHz): δ 9.28 (s, 1H, 1NH), 9.23 (s, 1H, 1NH), 8.16 (d, 1H, J = 8.1 Hz, ArH), 7.88–7.53 (m, 6H, 4 ArH & 3 furyl-H), 6.68 (bs, 2H, NH₂), 6.30 (d, 1H, J = 4.8 Hz, H5), 6.68 (bs, 2H, NH₂), 6.30 (d, 1H, J = 4.8 Hz, H4). GCMS m/z (rel. abundance): 331 (M.+, 100), 298 (50), 274 (43), 256 (23), 239 (15), 204 (23), 160 (17), 136 (16), 119 (8), 112 (10). Anal. Calcd. for C₂₀H₁₇N₃S (331.43); C, 72.48; H, 5.17; N, 12.68; S, 9.67. Found: C, 72.22; H, 4.93; N, 12.53; S, 9.98.2.

4.12.2. General Procedure for Synthesis of (4,6-diaryl-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisindolin-2-yl)methyl)benzothioates 10a–j

A solution of equimolar amounts of α-phthlimido-o-tolyl chloride (9, 0.668 mmol) and an appropriate 4,6-diaryl-3,4-dihydropyrimidine-2(1H)-thione 5a–j (0.668 mmol) in 10 mL absolute dichloromethane was stirred at rt for 4–6 h. The progress of the reaction was monitored periodically by TLC using dichloromethane/ethyl acetate (9:1) as mobile phase. The reaction mixture was then treated with brine, extracted with ethyl acetate, treated with dil. hydrochloric acid, sod. bicarbonate and finally washed with water. The organic solvent was dried with anhydrous sodium sulfate and evaporated under vacuum. The solid product was purified by crystallization from the appropriate solvent to give the title hybrids 10a–j.
S-(4,6-Diphenyl-1,6-dihydropyrimidin-2-yl)2-((1,3-dioxoisoindolin-2-yl)methyl)benzothioate (10a)

Yield: 0.233 g, 66% (white solid, ethanol); m.p. 147–8 °C; IR (KBr, cm⁻¹) ν: 3399 (NH), 3071,3011 (Ar-CH), 1772,1716 (CO), 1603 (C=N), 1547,1491 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.46 (s, 1H, NH), 7.90–721 (m, 18 H, ArH), 6.42 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.12 (d, 1H, J = 4.8 Hz, CHAr), 5.06 (s, 2H, CH₂). ¹³C-NMR (DMSO-d₆, 100 MHz): δ 189.7 (CO), 168.3 (2CO), 156.4, 144.0, 140.5, 136.0, 136.7, 135.5, 135.0, 132.0, 131.4, 129.3, 128.4, 127.6, 127.0, 126.7, 126.3, 126.4, 124.0 (ArC), 107.8 (C=CH-Ar), 55.1 (CHAr). GCMS m/z (rel. abundance): 529 (M+, 36), 452 (29), 392 (43), 338 (20), 280 (100), 265 (60), 236 (20), 223 (10), 173 (35), 146 (28). Anal. Calcd for C₃₂H₂₅N₃O₇S (529.61); C, 72.75; H, 4.38; N, 7.93; S, 6.05. Found: C, 72.34; H, 4.30; N, 8.07; S, 6.23.

S-(4-(4-Chlorophenyl)-6-(4-methoxyphenyl)-1,6-dihydropyrimidin-2-yl)2-((1,3-dioxoisoindolin-2-yl)methyl)benzothioate (10b)

Yield: 0.278 g, 70% (pale yellow solid, ethanol); m.p. 142–3 °C; IR (KBr, cm⁻¹) ν: 3450 (NH), 3064,2918 (Ar-CH), 1772, 1725 (CO), 1676 (C=N), 1594,1489 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.49 (s, 1H, NH), 7.99–6.92 (m, 16 H, ArH), 6.28 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.20 (d, 1H, J = 4.8 Hz, CHAr), 5.08 (s, 2H, CH₂), 3.76 (s, 3H, OCH₃). GCMS m/z (rel. abundance): 593 (M+, 100), 562 (50), 462 (34), 382 (43), 355 (61), 329 (23), 264 (32), 186 (26), 160 (35), 132 (20). Anal. Calcd for C₃₅H₂₄ClN₃O₇S (593.08); C, 66.72; H, 4.07; N, 7.07; S, 5.40. Found: C, 66.39; H, 4.22; N, 8.11; S, 5.91.

S-(6-Bis-(4-chlorophenyl)-1,6-dihydropyrimidin-2-yl)2-((1,3-dioxoisoindolin-2-yl)methyl)benzothioate (10c)

Yield: 0.258 g, 65% (yellow solid, ethanol/chloroform); m.p. 118–19 °C; IR (KBr, cm⁻¹) ν: 3432 (NH), 3064, 2964 (Ar-CH), 1772,1616 (CO), 1672 (C=N), 1592, 1489 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.43 (s, 1H, NH), 7.92–7.19 (m, 16H, ArH), 6.42 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.19 (d, 1H, J = 4.8 Hz, CHAr), 5.06 (s, 2H, CH₂). GCMS m/z (rel. abundance): 598 (M+, 100), 487 (86), 466 (20), 568 (37)521 (17), 398 (28), 358 (18), 302 (42), 266 (59), 275 (27), 190 (19), 160 (30), 147 (46), 132 (23). Anal. Calcd for C₃₅H₂₆Cl₂N₃O₇S (598.50); C, 64.22; H, 3.54; N, 7.02; S, 5.36. Found: C, 64.60; H, 3.37; N, 7.21; S, 5.30.

S-(6-(4-Chlorophenyl)-4-(p-tolyl)-1,6-dihydropyrimidin-2-yl)2-((1,3-dioxoisoindolin-2-yl)methyl)benzothioate (10d)

Yield: 0.282 g, 73% (yellow solid, ethanol); m.p. 144–6 °C; IR(KBr, cm⁻¹) ν: 3413 (NH), 3061,3041, 2922 (Ar-CH), 1771,1715 (CO), 1677 (C=N), 1545, 1490 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.45 (s, 1H, NH), 7.89–7.18 (m, 16H, ArH), 6.34 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.20 (d, 1H, J = 4.8 Hz, CHAr), 5.08 (s, 2H, CH₂), 2.34 (s, 3H, CH₃). GCMS m/z (rel. abundance): 577.5 (M+, 31), 487 (70), 466 (100), 457 (20), 385 (30), 375 (21), 314 (39), 297 (81), 264 (57), 217 (10), 160 (26), 146 (22), 132 (16). Anal. Calcd for C₃₅H₂₄ClN₃O₇S (578.08); C, 68.56; H, 4.18; N, 7.27; S, 5.55. Found: C, 68.84; H, 4.01; N, 7.37; S, 5.23.

S-(4-(4-Bromophenyl)-6-(4-(dimethylamino)phenyl)-1,6-dihydropyrimidin-2-yl)2-((1,3-dioxoisoindolin-2-yl)methyl)benzothioate (10e)

Yield: 0.304 g, 70% (brown solid, ethanol/chloroform); m.p. 135–6 °C; IR (KBr, cm⁻¹) ν: 3476 (NH), 2920 (Ar-CH), 1770,1715 (CO), 1674 (C=N), 1578,1485(C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.45 (s, 1H, NH), 8.08–6.75 (m, 16H, ArH), 6.34 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.18 (d, 1H, J = 4.8 Hz, CHAr), 5.05 (s,2H, CH₂), 2.52 (s, 6H, NMe₂). GCMS m/z (rel. abundance): 651 (M+, 41), 569 (19), 546 (100), 531 (13), 519 (18), 415 (21), 362 (49), 302
S-(4-(4-Chlorophenyl)-6-phenyl-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoxindolin-2-yl)methyl)benzothioate (10f)

Yield: 0.282 g, 75% (pale yellow solid, ethanol); m.p. 143–4 °C; IR (KBr, cm⁻¹) ν: 3464 (NH), 3061, 2970 (Ar-CH), 1772, 1716 (CO), 1677 (C=N), 1557, 1490 (C=C). GCMS m/z (rel. abundance): 563.5 (M+, 100), 543 (48), 443 (43), 414 (38), 389 (18), 375 (36), 300 (28), 265 (31), 227 (24), 161 (36), 146 (27), 132 (17). Anal. Calcd. for C₃₂H₂₂ClN₃O₃S (564.05); C, 68.14; H, 3.93; N, 7.45; S, 5.68. Found: C, 67.83; H, 4.17; N, 7.27; S, 5.55.

S-(4-(4-Chlorophenyl)-6-(4-nitrophenyl)-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoxindolin-2-yl)methyl)benzothioate (10g)

Yield: 0.280 g, 69% (brown solid, ethanol/chloroform); m.p. 161–3 °C; IR (KBr, cm⁻¹) ν: 3418 (NH), 3030, 2926 (Ar-CH), 1774, 1715 (CO), 1623 (C=N), 1521 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.52 (s, 1H, NH), 8.51–7.15 (m, 16H, ArH), 6.39 (d, 1H, J = 4.8 Hz, CH=C-Ar), 5.21 (d, 1H, J = 4.8 Hz, CHAr), 5.02 (s, 2H, CH₂). ¹³C-NMR (DMSO-d₆, 100 MHz): δ 188.4 (CO), 168.3 (2CO), 157.5, 148.0, 146.1, 142.0, 137.9, 137.7, 135.3, 135.0, 133.7, 132.5, 131.8, 129.8, 129.0, 128.1, 127.4, 126.1, 125.1, 124.6, 124.5 (ArC), 112.0 (C=CH—CHAr), 45.2 (CHAr). GCMS m/z (rel. abundance): 608.5 (M+, 50), 551 (17), 532 (25), 504 (55), 446 (31), 373 (100), 346 (22), 311 (25), 306 (9), 264 (39), 232 (21), 188 (31), 163 (16). Anal. Calcd. for C₃₂H₂₁ClN₄O₅S (609.05); C, 63.11; H, 3.48; N, 9.20; S, 5.26. Found: C, 62.88; H, 3.59; N, 9.18; S, 5.17.

S-(4-(4-Aminophenyl)-6-(4-chlorophenyl)-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoxindolin-2-yl)methyl)benzothioate (10h)

Yield: 0.285 g, 74% (orange solid, ethanol/chloroform); m.p. 185–7 °C; IR (KBr, cm⁻¹) ν: 3430 (NH), 3060, 2942 (Ar-CH), 1772, 1716 (CO), 1677 (C=N), 1545, 1490 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.48 (s, 1H, NH), 8.10–6.98 (m, 16H, ArH), 6.68 (d, 1H, J = 4.8 Hz, CH=C-Ar), 6.19 (bs, 2H, NH₂), 5.23 (d, 1H, J = 4.8 Hz, CHAr), 5.03 (s, 2H, CH₂). GCMS m/z (rel. abundance): 578.5 (M+, 47), 514 (9), 432 (22), 394 (11), 342 (100), 315 (49), 282 (46), 264 (34), 238 (27), 196 (17), 140 (14), 132 (11). Anal. Calcd. for C₃₂H₂₁ClN₄O₄S (579.07); C, 66.37; H, 4.00; N, 9.68; S, 5.54. Found: C, 66.20; H, 3.83; N, 9.83; S, 5.50.

S-(4-(4-Chlorophenyl)-6-(furan-2-yl)-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoxindolin-2-yl)methyl)benzothioate (10i)

Yield: 0.274 g, 74% (grey solid, ethanol); m.p. 146–8 °C; IR (KBr, cm⁻¹) ν: 3464 (NH), 3064, 2942 (Ar-CH), 1772, 1716 (CO), 1677 (C=N), 1545, 1490 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.19 (s, 1H, NH), 7.92–7.09 (m, 12 ArH and 3 furanyl-H), 6.25 (d, 1H, J = 4.8 Hz, CH=C-Ar), 6.19 (bs, 2H, NH₂), 5.23 (d, 1H, J = 4.8 Hz, CHAr), 5.03 (s, 2H, CH₂). GCMS m/z (rel. abundance): 553.5 (M+, 100), 511 (11), 449 (41), 393 (20), 315 (44), 287 (18), 254 (23), 238 (26), 264 (20), 160 (21), 132 (16). Anal. Calcd. for C₃₀H₂₀ClN₃O₄S (554.02); C, 65.04; H, 3.64; N, 7.58; S, 5.79. Found: C, 65.33; H, 3.78; N, 7.42; S, 5.51.

S-(4-(4-Aminophenyl)-6-(naphthalen-1-yl)-1,6-dihydropyrimidin-2-yl)-2-((1,3-dioxoisoxindolin-2-yl)methyl)benzothioate (10j)

Yield: 0.274 g, 74% (grey solid, ethanol); m.p. 146–8 °C; IR (KBr, cm⁻¹) ν: 3464 (NH), 3064, 2942 (Ar-CH), 1770, 1716 (CO), 1677 (C=N), 1502 (C=C). ¹H-NMR (DMSO-d₆, 400 MHz): δ 11.19 (s, 1H, NH), 7.92–7.09 (m, 12 ArH and 3 furanyl-H), 6.25 (d, 1H, J = 4.8 Hz, CH=C-Ar), 6.19 (bs, 2H, NH₂), 5.23 (d, 1H, J = 4.8 Hz, CHAr), 5.03 (s, 2H, CH₂). GCMS m/z (rel. abundance): 553.5 (M+, 100), 511 (11), 449 (41), 393 (20), 315 (44), 287 (18), 254 (23), 238 (26), 264 (20), 160 (21), 132 (16). Anal. Calcd. for C₃₀H₂₀ClN₃O₄S (554.02); C, 65.04; H, 3.64; N, 7.58; S, 5.79. Found: C, 65.33; H, 3.78; N, 7.42; S, 5.51.
5. Conclusions

A novel series of dihydropyrimidine phthalimide hybrids was synthesized, characterized and evaluated for their DPP-4 inhibitory activity, as well as, antioxidant activity. Hybrids 10g, 10i, 10e, 10d and 10b showed superior DPP-4 inhibitory activity compared to alogliptin, in the in vitro DPP-4 inhibition assay. Additionally, hybrids 10g and 10i have the most potent and longest DPP-4 inhibitory activity among the synthesized hybrids and alogliptin, in the in vivo study that carried on SD rats. Furthermore, investigation of antihyperglycemic effect of our synthesized hybrids in HFD-induced type 2 diabetic rats, revealed that hybrids 10g, 10i and 10e exhibited better glycemic control than alogliptin; this effect was further potentiated by addition of MET. Notably, glucose tolerance reached a normal level in 10g/MET treated group. Finally, hybrids 10j, 10e, 10h and 10d showed potent radical scavenging activity in DPPH assay.

Supplementary Materials: IR spectroscopy, $^1$H-NMR, $^{13}$C-NMR and mass spectroscopy of all the synthesized compounds are available online at [https://www.mdpi.com/1424-8247/14/2/144/s1](https://www.mdpi.com/1424-8247/14/2/144/s1), Figure S1: IR spectrum of compound 5a, Figure S2: IR Spectroscopy of compound 5b, Figure S3: IR Spectroscopy of compound 5e, Figure S4: IR Spectroscopy of compound 5d, Figure S5: IR Spectroscopy of compound 5f, Figure S6: IR Spectroscopy of compound 5i, Figure S7: IR Spectroscopy of compound 5g, Figure S8: IR Spectroscopy of compound 5b, Figure S9: IR Spectroscopy of compound 5i, Figure S10: IR Spectroscopy of compound 5j, Figure S11: IR Spectroscopy of compound 10a, Figure S12: IR Spectroscopy of compound 10b, Figure S13: IR Spectroscopy of compound 10c, Figure S14: IR Spectroscopy of compound 10d, Figure S15: IR Spectroscopy of compound 10e, Figure S16: IR Spectroscopy of compound 10f, Figure S17: IR Spectroscopy of compound 10g, Figure S18: IR Spectroscopy of compound 10h, Figure S19: IR Spectroscopy of compound 10i, Figure S20: IR Spectroscopy of compound 10j, Figure S21: $^1$H NMR Spectroscopy of Compound 5b, Figure S22: $^1$H-NMR spectrum of compound 5e, Figure S23: $^1$H NMR Spectroscopy of Compound 5d, Figure S24: $^1$H NMR Spectroscopy of Compound 5e, Figure S25: $^1$H NMR Spectroscopy of Compound 5f, Figure S26: $^1$H NMR Spectroscopy of Compound 5g, Figure S27: $^1$H NMR Spectroscopy of Compound 5h, Figure S28: $^1$H NMR Spectroscopy of Compound 5i, Figure S29: $^1$H NMR Spectroscopy of Compound 5j, Figure S30: $^1$H NMR Spectroscopy of Compound 10a, Figure S31: $^1$H NMR Spectroscopy of Compound 10b, Figure S32: $^1$H NMR Spectroscopy of Compound 10c, Figure S33: $^1$H NMR Spectroscopy of Compound 10d, Figure S34: $^1$H NMR Spectroscopy of Compound 10e, Figure S35: $^1$H NMR Spectroscopy of Compound 10f, Figure S36: $^1$H NMR Spectroscopy of Compound 10g, Figure S37: $^1$H NMR Spectroscopy of Compound 10h, Figure S38: $^1$H NMR Spectroscopy of Compound 10i, Figure S39: $^1$H NMR Spectroscopy of Compound 10j, Figure S40: $^{13}$C NMR Spectroscopy of Compound 5a, Figure S41: $^{13}$C NMR Spectroscopy of Compound 5d, Figure S42: $^{13}$C NMR Spectroscopy of Compound 5g, Figure S43: $^{13}$C NMR Spectroscopy of Compound 5b, Figure S44: $^{13}$C NMR Spectroscopy of Compound 10a, Figure S45: $^{13}$C NMR Spectroscopy of Compound 10b, Figure S46: $^{13}$C NMR Spectroscopy of Compound 10d, Figure S47: $^{13}$C NMR Spectroscopy of Compound 10g, Figure S48: $^{13}$C NMR Spectroscopy of Compound 10j, Figure S49: Mass Spectroscopy of Compound 5b, Figure S50: Mass Spectroscopy of Compound 5c, Figure S51: Mass Spectroscopy of Compound 5d, Figure S52: Mass Spectroscopy of Compound 5e, Figure S53: Mass Spectroscopy of Compound 5f, Figure S54: Mass Spectroscopy of Compound 5g, Figure S55: Mass Spectroscopy of Compound 5h, Figure S56: Mass Spectroscopy of Compound 5i, Figure S57: Mass Spectroscopy of Compound 5j, Figure S58: Mass Spectroscopy of Compound 10a, Figure S59: Mass Spectroscopy of Compound 10b, Figure S60: Mass Spectroscopy of Compound 10c, Figure S61: Mass Spectroscopy of Compound 10d, Figure S62: Mass Spectroscopy of Compound 10e, Figure S63: Mass Spectroscopy of Compound 10f, Figure S64: Mass Spectroscopy of Compound 10g, Figure S65:
Mass Spectroscopy of Compound 10h, Figure S66: Mass Spectroscopy of Compound 10i, Figure S67: Mass Spectroscopy of Compound 10j.

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