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1 Synthesis of new halogenated flavonoid-based isoxazoles: in vitro and in silico

2 evaluation of *a*-amylase inhibitory potential, a SAR analysis and DFT studies

3

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19	
20	
21	Abstract
22	Carbohydrates are the main source of calories in most diets, and α -amylase is considered one of the
23	main enzymes that initiate their digestion. The inhibition of α -amylase is therefore considered to be
24	a therapeutic strategy for the treatment of disorders of glucose absorption, such as dental caries,

25 periodontal disease, overweight, obesity and diabetes. On the other hand, flavonoids, isoxazoles and

26 halogenated derivatives are of great interest in medicinal chemistry due to their considerable 27 bioactivities. Therefore, this work describes the design, the synthesis and the α -amylase inhibitory potential of new halogenated flavonoid-based isoxazoles, and their biological properties. In fact, the 28 29 condensation of a previously synthesized halogenated flavonol with different aryl nitrile oxides 30 afford thirteen new hybrid cycloadducts (2a-m). Their structures were characterized by ¹H NMR, 31 ¹³C NMR and HRMS analysis. The physicochemical properties of these compounds (2a-m) were 32 also evaluated. The newly synthesized cycloadducts were evaluated in vitro for their α -amylase 33 inhibitory activity and excellent results were noted. The compound **2b** (IC₅₀= 16.2 \pm 0.3 μ M) exhibited the highest anti- α -amylase activity comparable to that of the standard substance 34 (Acarbose, IC₅₀= 15.7 \pm 0.2 μ M). The study of the Structure-Activity Relationship (SAR) was 35 36 sufficiently discussed based on the Molecular Docking analysis and the Density Functional Theory 37 (DFT) studies.

38

39 Keywords: Flavonoid-based isoxazoles, Cycloaddition, DFT study, α-Amylase inhibition,
40 Molecular docking, SAR analysis.

41

42 **1. Introduction**

43 Diabetes mellitus and overweight, although being non-communicable diseases, constitute a major 44 public health problem around the world [1-2]. On the other hand, carbohydrates are known as the main source of calories in most diets. Therefore, inhibiting the digestion of foods rich in 45 46 carbohydrates and blocking the absorption of monosaccharides can decrease the intake of calories, 47 which promotes the weight loss and the fight against obesity [3]. In addition, the succession of 48 carbohydrate consumption leads to postprandial hyperglycemia in diabetic subjects [4-5]. Diabetes 49 mellitus is primarily typified by uncontrolled blood sugar levels [3,6]. Thus, inhibiting carbohydrate 50 breakdown implements a fundamental strategy to control diabetes.

51 Furthermore, α -amylase (α -1,4-glucan-4-glucanohydrolase, EC.3.2.1.1) is an enzyme which has 52 the most important contribution to start the digestion of carbohydrates [7-8]. Indeed, α -amylase is a 53 digestive enzyme which hydrolyzes α -1,4-glycosidic bonds of carbohydrates, such as glycogen, 54 starches and other polysaccharides, to generate oligosaccharides of variable length capable of being 55 hydrolyzed by other digestive enzymes into monosaccharides which can be absorbed by cells [7-8]. 56 α -Amylase inhibitors reduce the glycemic index by delaying the hydrolysis of carbohydrates. The 57 inhibition of α -amylase enzyme, leading to a drop in blood glucose levels, is considered a 58 therapeutic strategy for the treatment of carbohydrate absorption disorders, such as dental caries, 59 periodontal disease, overweight, obesity and diabetes [8]. However, many commercial α -amylase 60 inhibitors have complained of intestinal side effects [4]. In order to avoid the unwanted side effects 61 of current drugs, many efforts are devoted to the discovery of new inhibitors of carbohydrate 62 digestion.

63 Flavonoids are secondary metabolites constituting the largest group of natural polyphenols. They 64 are commonly found in plants and responsible for the attractive colors of flowers and fruits [9]. The 65 flavonoid structure has a C₁₅ carbon skeleton consisting of two aromatic rings (A) and (B) linked 66 together by a C_3 chain forming the heterocycle (C) [10]. Moreover, 2-phenylbenzopyrane 67 flavonoids were classified into many categories, distinguished by cyclization, degree of 68 unsaturation and oxidation and by the pyranic cycle (C) substitution [10-11]. On the other hand, 69 numerous flavonoid compounds, such as rutin, quercetagetin, scutellarein, 8-prenylapigenin, fisetin, 70 luteolin and quercetin (Figure 1), were reported as antidiabetics [12-17] and α -amylase inhibitors 71 [5,18-19], together with different other biological properties [20-27]. However, the broad biological 72 profile of flavonoids has encouraged chemical researchers to synthesize these remarkable 73 compounds with special chemical substituents and targeted bioactivities [28-32].

74

Fig. 1. Example of flavonoids previously reported as antidiabetics.

76 Moreover, in a previous study, it was shown that the presence of bromine atom was found to be 77 essential in the structure of the most inhibitory derivatives of the α -amylase enzyme [33], thus 78 dictating their presence as one of the main parts of the structure of future α -amylase inhibitors. On 79 the other hand, the metabolic stability, the strong electron withdrawing character and the lipophilic 80 nature of the trifluoromethyl moiety allow an increase in their demand for the development of 81 bioactive compounds [34]. Otherwise, lipophilicity, an important physicochemical property of 82 bioactive compounds, has a molecule penetration role through the cell membrane [35], thus 83 increasing the binding to receptor pockets, which supposes a powerful activity. Hence, many 84 important drugs are trifluoromethylated compounds, viz. fluphenazine (antipsychotic, anticancer) 85 [36], leflunomide (antirheumatic) [37], celecoxib (COX-2 selective inhibitor) [38], fluoxetine 86 (antidepressant) [39], and fluazinam (fungicide) [40] (Figure 2). In addition, a trifluoromethylated 5-amino-nicotinic acid derivative has been reported to exhibit promising α -amylase and α -87 88 glucosidase inhibitors [41].

89

90

Fig. 2. Structures of trifluoromethylated drugs.

91

92 Furthermore, isoxazole (1,2-oxazole) is a five-membered aromatic monocyclic heteroarene. It is of 93 a great importance in medicinal chemistry. In addition, natural products containing isoxazole ring, 94 such as ibotenic acid (neurotoxin and hallucinogenic) and muscimol (hallucinogenic and GABAA receptor antagonist) (Figure 3a) [42-45], have aroused the interest from pharmacologists and 95 96 chemists because of their potent biological and pharmacological properties. Therefore, the isoxazole 97 heterocycle has been extensively used as a key building block for drugs such as valdecoxib, 98 parecoxib (COX-2 inhibitors) [46], zonisamide (anti-convulsant) [47] and leflunomide 99 (antirheumatic) [37]. Furthermore, 1,2-oxazole derivatives are reported to possess a wide spectrum

- of biological properties such as antidiabetics (Figure 3b) [48], along with other bioactivities viz.
 analgesic, anti-inflammatory, anti-HIV, and anti-cancer [49-53].
- 102

Fig. 3. (a) Natural compounds containing isoxazole ring. (b) Previously reported isoxazole
 derivatives as antidiabetic agents. (c) Reported flavonoid-based isoxazoles with antidiabetic
 properties.

106

107 To the best of our knowledge, there is only one previous study reported that flavonoid-based 108 isoxazoles exhibit antidiabetic properties, which have improved the glucose consumption in insulin 109 resistant HepG2 cells. Also, it turned out to be important to indicate that the studied structures were 110 limited to the 3-(4-cyanophenyl)isoxazole derivatives (Figure 3c) [54].

111 According to the potential α -amylase inhibition and antidiabetic activities of flavonoids and 112 isoxazole derivatives and the inspiration from the findings cited above, the present work was 113 performed to discover new hybrid molecules, where a halogenated flavonoid was joined to a 3-114 arylisoxazole moiety via a methylene linker in order to increase the α -amylase inhibitory potency. Hence, the structure-activity relationship was in silico approved by the molecular docking analysis. 115 116 In addition, the characterization of the structural and the vibrational spectroscopic properties of the synthesized molecules were performed using DFT methods at the B3LYP/6-311++G (d,p) level of 117 118 theory. The physicochemical properties of these compounds were evaluated using the software 119 HyperChem 8.03, and from which various physicochemical and quantum descriptor parameters 120 were deduced.

121

122 **2. Material and methods**

123 2.1. General experimental procedures

¹H NMR (300 MHz), DEPT 135 and ¹³C NMR (75 MHz) spectra were recorded in deuterated 124 chloroform and dimethyl sulfoxide- d_6 with a Bruker AC-300 spectrometer. All chemical shifts (δ) 125 126 were reported in parts per million (ppm), coupling constants (J) were given in Hertz (Hz) and 127 residual non-deuterated solvent resonance was used as internal reference. DCI-HRMS spectra were 128 run in a GCT 1er Waters in the positive ion mode. ESI-HRMS were recorded with ESI-TOF (LCT, 129 Waters) in the positive ion mode using the reflectron mode. All chemical reactions were monitored by commercial TLC plates (Silica gel 60, F₂₅₄). All used solvents were freshly distilled prior to use. 130 131 Büchi 510 apparatus was used to determine melting points.

132 *2.2. Chemistry*

133 2.2.1. General procedure for the synthesis of the flavonol (1)

134 The preparation of the halogenated flavonol (1), previously synthesized in our laboratory was 135 reported according to the procedure adopted by Znati et al. (2019) [32]. A mixture of 2 mL of 5-136 bromo-2-hydroxyacetophenone (16.61 mmol), 2.89 g of 4-(trifluoromethyl)benzaldehyde (16.61 137 mmol) and 2 g of NaOH (50 mmol) were added to 100 mL of methanol in a round-bottom flask 138 (500 mL) equipped with a magnetic stirrer. The obtained solution was heated at reflux for 3 hours. 139 The reaction mixture was cooled to room temperature for 15 min. Afterwards, 100 mL of NaOH 140 (0.5M), then 6.84 mL of H₂O₂ (35%) were added and the solution was stirred for 2 hours at room 141 temperature. The reaction mixture was poured to give a precipitate into ice-water which is then 142 recovered by filtration to obtain the flavonol (1). The halogenated flavonol (1) was obtained with a 143 yield of 87%.

144 6-bromo-3-hydroxy-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one (1)

145 Red solid, yield: 87%, mp: 270°C. DCI-HRMS $[M + H]^+$ calcd. for $(C_{16}H_9BrF_3O_3)^+$: 384.9687,

146 found: 384.9680. ¹H NMR (DMSO- d_6 , 300 MHz) δ_H 8.11 (1H, d, J = 2.4 Hz, H-5), 7.71 (1H, dd, J_1

147 = 9.0, J_2 = 2.4 Hz, H-7), 7.64 (1H, d, J = 9.0 Hz, H-8), 7.71 (2H, d, J = 8.4 Hz, H-2',6'), 8.87 (2H,

148 d, J = 8.4 Hz, H-3',5'). ¹³C NMR (DMSO- d_6 , 75 MHz) δ_C 154.9 (C-2), 141.8 (C-3), 179.5 (C-4),

149 127.0 (C-5), 114.5 (C-6), 133.9 (C-7), 121.1 (C-8), 152.7 (C-9), 122.7 (C-10), 139.4 (C-1'), 124.5

150 (C-2',6'), 124.7 (q, ${}^{3}J_{CF}$ = 3.6 Hz, C-3',5'), 125.1 (q, ${}^{2}J_{CF}$ = 31.0 Hz, C-4'), 124.9 (q, ${}^{1}J_{CF}$ = 270.1 Hz, 151 C-7').

152 2.2.2. General procedure for the synthesis of the dipolarophile (2)

To a mixture of 3 g of the synthesized flavonol (1, 7.81 mmol) in anhydrous DMF, 2 equivalents of K₂CO₃ were added. The solution is kept under stirring for 15 min in an ice bath. Subsequently, 2 equivalents of propargyl bromide are then added. The reaction mixture was stirred for 2 hours at room temperature and the reaction was monitored by TLC analysis. Once the reaction is complete, the residue is diluted with 200 mL of distilled water and through sintered glass, the formed precipitate was then filtered. The dipolarophile **2** was obtained with an excellent yield, 96%.

159 6-bromo-3-(prop-2-yn-1-yloxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one (2)

160 Yellowish solid, yield: 96%, mp: 125°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.41 (1H, d, J = 2.4 Hz,

161 H-5), 7.81 (1H, dd, $J_1 = 8.9$, $J_2 = 2.4$ Hz, H-7), 7.48 (1H, d, J = 8.9 Hz, H-8), 7.81 (2H, d, J = 8.2

162 Hz, H-2',6'), 8.27 (2H, d, J = 8.2 Hz, H-3',5'), 5.06 (2H, d, J = 2.4 Hz, H-1"), 2.35 (1H, t, J = 2.4

163 Hz, H-3"). ¹³C NMR (CDCl₃, 75 MHz) δ_C 155.4 (C-2), 139.4 (C-3), 173.8 (C-4), 128.7 (C-5), 118.8

164 (C-6), 137.1 (C-7), 120.2 (C-8), 154.3 (C-9), 125.5 (C-10), 134.3 (C-1'), 129.5 (C-2',6'), 125.6 (q,

165 ${}^{3}J_{CF} = 3.6 \text{ Hz}, \text{ C-3}', 5'), 132.7 \text{ (q, } {}^{2}J_{CF} = 32.8 \text{ Hz}, \text{ C-4}'), 123.9 \text{ (q, } {}^{1}J_{CF} = 270.7 \text{ Hz}, \text{ C-7}'), 59.6 \text{ (C-1}''),$

166 78.4 (C-2"), 76.9 (C-3").

167 2.2.3. General procedure for the synthesis of the halogenated flavonoid-based isoxazoles 2a-m

A mixture of 50 mg of the dipolarophile (**2**), 1.5 equivalents of the appropriate hydroximyl chloride, equivalents of the triethylamine and 0.1 equivalent of CuI were dissolved at room temperature in 2 mL of DMF. The solution was subjected for 5 min to a microwave irradiation at 250 W. Then, the solution was diluted with distilled water and extracted three times with ethyl acetate (30 mL). After *in vacuo* solvent removing, the resulting residue was purified by silica gel column chromatography,

- eluted with a mixture of petroleum ether : ethyl acetate (8:2). The desired products 2a-m were
 obtained with good yields ranging from 72 to 94%.
- 175 2.2.3.1. 6-bromo-3-((3-phenylisoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-
- 176 *4-one* (2*a*)
- 177 Yellowish solid, yield: 90%, mp: 164°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.3 Hz,
- 178 H-5), 7.81 (1H, dd, $J_1 = 8.9$, $J_2 = 2.3$ Hz, H-7), 7.45 (1H, m, H-8), 7.73 (2H, d, J = 8.3 Hz, H-2',6'),
- 179 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.44 (1H, s, H-4"), 5.42 (2H, s, H-6"), 7.67 (2H, m, H-2"',6"'), 7.45
- 180 (3H, m, H-3^{'''},4^{'''},5^{'''}). ¹³C NMR (CDCl₃, 75 MHz) δ_{C} 155.5 (C-2), 139.7 (C-3), 173.7 (C-4), 128.6
- 181 (C-5), 118.9 (C-6), 137.3 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-
- 182 2',6'), 125.6 (q, ${}^{3}J_{CF}$ = 4.0 Hz, C-3',5'), 132.8 (q, ${}^{2}J_{CF}$ = 32.8 Hz, C-4'), 123.8 (q, ${}^{1}J_{CF}$ = 270.8 Hz, C-
- 183 7'), 167.8 (C-3"), 103.0 (C-4"), 162.5 (C-5"), 63.8 (C-6"), 128.7 (C-1""), 126.9 (C-2"",6""), 129.1 (C-
- 184 3‴,5‴), 130.4 (C-4‴).
- 185 2.2.3.2. 6-bromo-3-((3-(4-fluorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H186 chromen-4-one (2b)

187 Yellowish solid, yield: 94%, mp: 176°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.3 Hz, 188 H-5), 7.82 (1H, dd, $J_1 = 8.9$, $J_2 = 2.3$ Hz, H-7), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3189 Hz, H-2',6'), 8.11 (2H, d, J = 8.3 Hz, H-3',5'), 6.41 (1H, s, H-4"), 5.41 (2H, s, H-6"), 7.66 (2H, dd, $J_1 = 8.6, J_2 = 5.3$ Hz, H-2",6"), 7.13 (2H, t, J = 8.6 Hz, H-3",5"). ¹³C NMR (CDCl₃, 75 MHz) δ_C 190 191 155.5 (C-2), 139.8 (C-3), 173.6 (C-4), 128.6 (C-5), 119.0 (C-6), 137.3 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.7$ Hz, C-3',5'), 132.8 (q, ${}^{2}J_{CF} = 32.8$ 192 193 Hz, C-4'), 123.8 (q, ${}^{1}J_{CF}$ = 271.0 Hz, C-7'), 168.0 (C-3"), 102.8 (C-4"), 161.6 (C-5"), 63.8 (C-6"), 124.9 (d, ${}^{4}J_{CF}$ = 3.5 Hz, C-1""), 128.9 (d, ${}^{3}J_{CF}$ = 8.5 Hz, C-2"",6""), 116.3 (d, ${}^{2}J_{CF}$ = 21.8 Hz, C-3"",5""), 194 195 164.1 (d, ${}^{1}J_{CF} = 248.7$ Hz, C-4"').

196 2.2.3.3. 6-bromo-3-((3-(4-chlorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H197 chromen-4-one (2c)

Yellowish solid, yield: 93%, mp: 191°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz, H-5), 7.82 (1H, dd, $J_1 = 8.9$, $J_2 = 2.4$ Hz, H-7), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3Hz, H-2',6'), 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.42 (1H, s, H-4"), 5.41 (2H, s, H-6"), 7.61 (2H, d, J= 8.6 Hz, H-2",6"), 7.42 (2H, d, J = 8.6 Hz, H-3",5"). ¹³C NMR (CDCl₃, 75 MHz) $\delta_{\rm C}$ 155.5 (C-2), 139.8 (C-3), 173.6 (C-4), 128.6 (C-5), 119.0 (C-6), 137.3 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-203 10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ³ $J_{\rm CF} = 3.6$ Hz, C-3',5'), 132.8 (q, ² $J_{\rm CF} = 32.6$ Hz, C-4'), 123.8 (q, ¹ $J_{\rm CF} = 271.2$ Hz, C-7'), 168.2 (C-3"), 102.8 (C-4"), 161.6 (C-5"), 63.8 (C-6"), 127.2 (C-

- 205 1""), 129.4 (C-2"",6""), 128.2 (C-3"",5""), 136.5 (C-4"").
- 206 2.2.3.4. 6-bromo-3-((3-(4-bromophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H207 chromen-4-one (2d)

208 Yellowish solid, yield: 91%, mp: 194°C. ESI-HRMS $[M + H]^+$ calcd. for $(C_{26}H_{15}Br_2F_3NO_4)^+$: 209 619.9320, found: 619.9296. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz, H-5), 7.82 210 $(1H, dd, J_1 = 8.9, J_2 = 2.4 Hz, H-7), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3 Hz, H-2', 6'),$ 211 8.11 (2H, d, J = 8.3 Hz, H-3',5'), 6.43 (1H, s, H-4"), 5.41 (2H, s, H-6"), 7.58 (2H, d, J = 8.8 Hz, H-2",6"), 7.54 (2H, d, J = 8.8 Hz, H-3",5"). ¹³C NMR (CDCl₃, 75 MHz) δ_{C} 155.4 (C-2), 139.8 (C-3), 212 213 173.6 (C-4), 128.6 (C-5), 119.0 (C-6), 137.3 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.8 214 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.4$ Hz, C-3',5'), 132.9 (q, ${}^{2}J_{CF} = 32.6$ Hz, C-4'), 123.8 (q, 215 ${}^{1}J_{CF} = 271.5 \text{ Hz}, \text{ C-7'}, 168.2 \text{ (C-3'')}, 102.7 \text{ (C-4'')}, 161.7 \text{ (C-5'')}, 63.8 \text{ (C-6'')}, 127.7 \text{ (C-1''')}, 128.4 \text{ (C-6'')}, 127.7 \text{ (C-1''')}, 128.7 \text{ (C-6'')}, 127.7 \text{ (C-6''')}, 127.7 \text{ (C-6''')}, 127.7 \text{ (C-6$ 216 (C-2^{'''},6^{'''}), 132.4 (C-3^{'''},5^{'''}), 124.7 (C-4^{'''}).

- 217 2.2.3.5. 6-bromo-3-((3-(p-tolyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen218 4-one (2e)
- 219 Yellowish solid, yield: 90%, mp: 172°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz,
- 220 H-5), 7.81 (1H, dd, $J_1 = 8.9$, $J_2 = 2.4$ Hz, H-7), 7.45 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3
- 221 Hz, H-2',6'), 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.41 (1H, s, H-4"), 5.41 (2H, s, H-6"), 7.56 (2H, d, J
- 222 = 8.0 Hz, H-2^{'''},6^{'''}), 7.24 (2H, d, J = 8.0 Hz, H-3^{'''},5^{'''}), 2.40 (3H, s, H-7^{'''}). ¹³C NMR (CDCl₃, 75

223 MHz) δ_C 155.5 (C-2), 139.7 (C-3), 173.7 (C-4), 128.6 (C-5), 118.9 (C-6), 137.2 (C-7), 120.3 (C-8),

224 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.7$ Hz, C-3',5'), 132.8 (q,

225 ${}^{2}J_{CF} = 32.7 \text{ Hz}, \text{ C-4'}$, 123.8 (q, ${}^{1}J_{CF} = 270.5 \text{ Hz}, \text{ C-7'}$), 167.6 (C-3"), 102.9 (C-4"), 162.5 (C-5"),

- 226 63.8 (C-6"), 125.9 (C-1""), 126.8 (C-2"",6""), 129.8 (C-3"",5""), 140.5 (C-4""), 21.6 (C-7"").
- 227 2.2.3.6. 6-bromo-3-((3-(4-(tert-butyl)phenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)228 4H-chromen-4-one (2f)
- Yellowish solid, yield: 90%, mp: 118°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.44 (1H, d, J = 2.4 Hz, 229 230 H-5), 7.82 (1H, dd, *J*₁ = 8.9, *J*₂ = 2.4 Hz, H-7), 7.46 (1H, d, *J* = 8.9 Hz, H-8), 7.73 (2H, d, *J* = 8.3 231 Hz, H-2',6'), 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.42 (1H, s, H-4"), 5.43 (2H, s, H-6"), 7.61 (2H, d, J = 8.5 Hz, H-2''', 6'''), 7.46 (2H, d, J = 8.5 Hz, H-3''', 5'''), 1.36 (9H, s, H-8''').¹³C NMR (CDCl₃, 75 232 233 MHz) δ_C 155.5 (C-2), 139.7 (C-3), 173.7 (C-4), 128.6 (C-5), 118.9 (C-6), 137.2 (C-7), 120.3 (C-8), 234 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.8$ Hz, C-3',5'), 132.8 (q, 235 ${}^{2}J_{CF} = 32.7$ Hz, C-4'), 123.8 (q, ${}^{1}J_{CF} = 270.8$ Hz, C-7'), 167.5 (C-3"), 103.0 (C-4"), 162.4 (C-5"), 236 63.8 (C-6"), 125.8 (C-1""), 126.1 (C-2"",6""), 126.7 (C-3"",5""), 153.7 (C-4""), 35.0 (C-7""), 31.4 (C-8""). 237 2.2.3.7. 6-bromo-3-((3-(4-nitrophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-
- 238 *chromen-4-one* (**2***g*)

239 Yellowish solid, yield: 72%, mp: 177°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.44 (1H, d, J = 2.4 Hz, 240 H-5), 7.84 (1H, dd, $J_1 = 8.9$, $J_2 = 2.4$ Hz, H-7), 7.48 (1H, d, J = 8.9 Hz, H-8), 7.74 (2H, d, J = 8.3241 Hz, H-2',6'), 8.12 (2H, d, J = 8.3 Hz, H-3',5'), 6.57 (1H, s, H-4"), 5.43 (2H, s, H-6"), 7.87 (2H, d, J = 8.9 Hz, H-2",6"), 8.31 (2H, d, J = 8.9 Hz, H-3",5"). ¹³C NMR (CDCl₃, 75 MHz) δ_{C} 155.5 (C-2), 242 243 139.8 (C-3), 173.6 (C-4), 128.7 (C-5), 119.1 (C-6), 137.4 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.7 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF}$ = 3.9 Hz, C-3',5'), 132.9 (q, ${}^{2}J_{CF}$ = 32.7 Hz, C-4'), 244 245 124.0 (q, ${}^{1}J_{CF} = 271.0 \text{ Hz}, \text{ C-7'}$), 169.0 (C-3"), 102.9 (C-4"), 160.8 (C-5"), 63.8 (C-6"), 134.8 (C-246 1""), 124.4 (C-2"",6""), 127.8 (C-3"",5""), 149.1 (C-4"").

247 2.2.3.8. 6-bromo-3-((3-(4-methoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-

248 *4H-chromen-4-one* (2*h*)

- 249 Yellowish solid, yield: 94%, mp: 152°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.42 (1H, d, J = 2.3 Hz,
- 250 H-5), 7.81 (1H, dd, $J_1 = 8.9$, $J_2 = 2.3$ Hz, H-7), 7.45 (1H, d, J = 8.9 Hz, H-8), 7.72 (2H, d, J = 8.3
- 251 Hz, H-2',6'), 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.38 (1H, s, H-4"), 5.40 (2H, s, H-6"), 7.60 (2H, d, J
- 252 = 8.7 Hz, H-2"',6"'), 6.95 (2H, d, J = 8.7 Hz, H-3"',5"'), 3.86 (3H, s, H-7"'). ¹³C NMR (CDCl₃, 75
- 253 MHz) δ_C 155.4 (C-2), 139.7 (C-3), 173.7 (C-4), 128.6 (C-5), 118.9 (C-6), 137.2 (C-7), 120.3 (C-8),
- 254 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.7$ Hz, C-3',5'), 132.8 (q,
- 255 ${}^{2}J_{CF} = 32.7 \text{ Hz}, \text{ C-4'}, 123.8 \text{ (q, } {}^{1}J_{CF} = 270.6 \text{ Hz}, \text{ C-7'}, 167.5 \text{ (C-3'')}, 102.8 \text{ (C-4'')}, 162.1 \text{ (C-5'')},$
- 256 63.8 (C-6"), 121.2 (C-1""), 128.3 (C-2"",6""), 114.5 (C-3"",5""), 161.3 (C-4""), 55.5 (C-7"").
- 257 2.2.3.9. 6-bromo-3-((3-(4-butoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H258 chromen-4-one (2i)
- 259 Yellowish solid, yield: 93%, mp: 126°C. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz, 260 H-5), 7.81 (1H, dd, *J*₁ = 8.9, *J*₂ = 2.4 Hz, H-7), 7.45 (1H, d, *J* = 8.9 Hz, H-8), 7.73 (2H, d, *J* = 8.3 261 Hz, H-2',6'), 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.37 (1H, s, H-4"), 5.40 (2H, s, H-6"), 7.58 (2H, d, J 262 = 8.8 Hz, H-2^{'''},6^{'''}), 6.94 (2H, d, J = 8.8 Hz, H-3^{'''},5^{'''}), 4.01 (2H, t, J = 6.5 Hz, H-7^{'''}), 1.80 (2H, m, 263 H-8"''), 1.52 (2H, m, H-9"'), 1.00 (3H, t, J = 7.3 Hz, H-10"'). ¹³C NMR (CDCl₃, 75 MHz) δ_{C} 155.5 (C-2), 139.7 (C-3), 173.7 (C-4), 128.6 (C-5), 118.9 (C-6), 137.2 (C-7), 120.3 (C-8), 154.2 (C-9), 264 265 125.5 (C-10), 133.7 (C-1'), 129.4 (C-2',6'), 125.5 (q, ${}^{3}J_{CF} = 3.6$ Hz, C-3',5'), 132.7 (q, ${}^{2}J_{CF} = 32.6$ 266 Hz, C-4'), 123.8 (q, ${}^{1}J_{CF} = 270.9$ Hz, C-7'), 167.3 (C-3"), 102.8 (C-4"), 162.1 (C-5"), 63.7 (C-6"), 267 120.8 (C-1"'), 128.2 (C-2",6"'), 115.0 (C-3",5"'), 160.9 (C-4"'), 68.0 (C-7"'), 31.4 (C-8"'), 19.4 (C-9"'), 14.0 (C-10""). 268
- 269 2.2.3.10. 6-bromo-3-((3-(4-butoxy-3-chlorophenyl)isoxazol-5-yl)methoxy)-2-(4-
- 270 (trifluoromethyl)phenyl)-4H-chromen-4-one (2j)

271 Yellowish solid, yield: 91%, mp: 140°C. ESI-HRMS $[M + H]^+$ calcd. for $(C_{30}H_{23}BrClF_3NO_5)^+$: 648.0400, found: 648.0397. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.42 (1H, d, J = 2.4 Hz, H-5), 7.81 272 273 $(1H, dd, J_1 = 8.9, J_2 = 2.4 Hz, H-7), 7.45 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3 Hz, H-2', 6'),$ 274 8.10 (2H, d, J = 8.3 Hz, H-3',5'), 6.38 (1H, s, H-4"), 5.39 (2H, s, H-6"), 7.69 (1H, d, J = 2.1 Hz, H-2"'), 6.95 (1H, d, J = 8.6 Hz, H-5"'), 7.50 (1H, dd, $J_1 = 8.6$, $J_2 = 2.1$ Hz, H-6"'), 4.08 (2H, t, J = 6.5275 276 Hz, H-7"'), 1.86 (2H, m, H-8"'), 1.56 (2H, m, H-9"'), 1.01 (3H, t, J = 7.4 Hz, H-10"'). ¹³C NMR 277 (CDCl₃, 75 MHz) δ_C 155.4 (C-2), 139.7 (C-3), 173.6 (C-4), 128.6 (C-5), 118.9 (C-6), 137.3 (C-7), 278 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} = 3.6$ Hz, C-279 3',5'), 132.8 (q, ${}^{2}J_{CF}$ = 32.6 Hz, C-4'), 123.8 (q, ${}^{1}J_{CF}$ = 269.9 Hz, C-7'), 167.8 (C-3"), 102.6 (C-4"), 280 161.3 (C-5"), 63.8 (C-6"), 121.7 (C-1""), 128.7 (C-2""), 123.8 (C-3""), 156.3 (C-4""), 113.4 (C-5""), 281 126.4 (C-6"'), 69.2 (C-7"'), 31.3 (C-8"'), 19.4 (C-9"'), 14.0 (C-10"').

282 2.2.3.11. 6-bromo-3-((3-(3-chloro-4-methoxyphenyl)isoxazol-5-yl)methoxy)-2-(4283 (trifluoromethyl)phenyl)-4H-chromen-4-one (2k)

284 Yellowish solid, yield: 90%, mp: 141°C. ESI-HRMS $[M + H]^+$ calcd. for $(C_{27}H_{17}BrClF_3NO_5)^+$: 605.9931, found: 605.9924. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz, H-5), 7.82 285 286 $(1H, dd, J_1 = 8.9, J_2 = 2.4 Hz, H-7), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, m, H-2', 6'), 8.11 (2H, H-1), 8.16 (2H,$ 287 d, J = 8.2 Hz, H-3',5'), 6.39 (1H, s, H-4"), 5.40 (2H, s, H-6"), 7.71 (1H, m, H-2"), 6.97 (1H, d, J = 288 8.5 Hz, H-5"'), 7.54 (1H, dd, $J_1 = 8.5$, $J_2 = 1.9$ Hz, H-6"'), 3.96 (3H, s, H-7"'). ¹³C NMR (CDCl₃, 75) 289 MHz) δ_C 155.4 (C-2), 139.7 (C-3), 173.6 (C-4), 128.5 (C-5), 118.9 (C-6), 137.3 (C-7), 120.3 (C-8), 290 154.2 (C-9), 125.5 (C-10), 133.7 (C-1'), 129.4 (C-2',6'), 125.5 (q, ${}^{3}J_{CF} = 3.2$ Hz, C-3',5'), 132.7 (q, 291 $^{2}J_{CF} = 32.7$ Hz, C-4'), 123.8 (q, $^{1}J_{CF} = 269.7$ Hz, C-7'), 167.8 (C-3"), 102.6 (C-4"), 161.1 (C-5"), 292 63.7 (C-6"), 122.0 (C-1""), 128.7 (C-2""), 123.3 (C-3""), 156.6 (C-4""), 112.2 (C-5""), 126.5 (C-6""), 293 56.4 (C-7"").

- 294 2.2.3.12. 6-bromo-3-((3-(3-bromo-4-methoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-
- 295 (trifluoromethyl)phenyl)-4H-chromen-4-one (2l)

296 Yellowish solid, yield: 89%, mp: 164°C. ESI-HRMS $[M + H]^+$ calcd. for $(C_{27}H_{17}Br_2F_3NO_5)^+$: 649.9426, found: 649.9409. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.43 (1H, d, J = 2.4 Hz, H-5), 7.82 297 298 $(1H, dd, J_1 = 8.9, J_2 = 2.4 Hz, H-7), 7.46 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3 Hz, H-2', 6'),$ 299 8.11 (2H, d, J = 8.3 Hz, H-3',5'), 6.40 (1H, s, H-4"), 5.40 (2H, s, H-6"), 7.88 (1H, d, J = 2.0 Hz, H-300 2"'), 6.94 (1H, d, J = 8.5 Hz, H-5"'), 7.59 (1H, dd, $J_1 = 8.5$, $J_2 = 2.0$ Hz, H-6"'), 3.95 (3H, s, H-7"'). 301 ¹³C NMR (CDCl₃, 75 MHz) δ_C 155.5 (C-2), 139.7 (C-3), 173.6 (C-4), 128.6 (C-5), 119.0 (C-6), 302 137.3 (C-7), 120.3 (C-8), 154.3 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} =$ 3.7 Hz, C-3',5'), 132.8 (q, ${}^{2}J_{CF}$ = 33.0 Hz, C-4'), 123.8 (q, ${}^{1}J_{CF}$ = 270.8 Hz, C-7'), 167.9 (C-3"), 303 304 102.7 (C-4"), 161.1 (C-5"), 63.8 (C-6"), 122.5 (C-1""), 131.9 (C-2""), 112.5 (C-3""), 157.6 (C-4""), 305 112.2 (C-5"'), 127.3 (C-6"'), 56.6 (C-7"').

306 2.2.3.13.6-bromo-3-((3-(3,4-dimethoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-

308 Yellowish solid, yield: 90%, mp: 148°C. ESI-HRMS $[M + H]^+$ calcd. for $(C_{28}H_{20}BrF_3NO_6)^+$: 309 602.0426, found: 602.0423. ¹H NMR (CDCl₃, 300 MHz) $\delta_{\rm H}$ 8.41 (1H, d, J = 2.4 Hz, H-5), 7.80 310 $(1H, dd, J_1 = 8.9, J_2 = 2.4 Hz, H-7), 7.45 (1H, d, J = 8.9 Hz, H-8), 7.73 (2H, d, J = 8.3 Hz, H-2', 6'),$ 311 8.11 (2H, d, J = 8.3 Hz, H-3',5'), 6.41 (1H, s, H-4"), 5.40 (2H, s, H-6"), 7.31 (1H, d, J = 1.9 Hz, H-312 2"), 6.89 (1H, d, J = 8.3 Hz, H-5"), 7.13 (1H, dd, $J_1 = 8.3$, $J_2 = 1.9$ Hz, H-6"), 3.93 (6H, s, H-7", 8"). 313 ¹³C NMR (CDCl₃, 75 MHz) δ_C 155.4 (C-2), 139.7 (C-3), 173.6 (C-4), 128.6 (C-5), 118.9 (C-6), 314 137.2 (C-7), 120.3 (C-8), 154.2 (C-9), 125.6 (C-10), 133.8 (C-1'), 129.4 (C-2',6'), 125.6 (q, ${}^{3}J_{CF} =$ 315 3.8 Hz, C-3',5'), 132.7 (q, ${}^{2}J_{CF} = 32.7$ Hz, C-4'), 123.8 (q, ${}^{1}J_{CF} = 270.8$ Hz, C-7'), 167.5 (C-3"), 316 102.8 (C-4"), 162.2 (C-5"), 63.8 (C-6"), 121.4 (C-1""), 109.4 (C-2""), 151.0 (C-3""), 149.6 (C-4""), 317 111.3 (C-5"'), 120.1 (C-6"'), 56.2 (C-7"'), 56.2 (C-8"').

318 2.3. α -Amylase inhibitory assay

319 The inhibition of α -amylase enzyme assays was performed according to the protocol described by 320 Chortani et al. (2021) [55]. Each compound was dissolved in the minimum of DMSO at different

321 concentrations and then diluted in sodium phosphate buffer (0.02 M, pH 6.9 with 0.006 M NaCl). 322 The DMSO level does not exceed 1% in the mixture. After that, 50 µL of each sample solution was 323 added to 50 μ L of α -amylase solution (0.5 mg/mL dissolved in the same buffer medium. After pre-324 incubation for 10 min at 25°C, 50 µL of the starch solution (1% in buffer solution) was added to the 325 mixture. After incubation at 25°C for 10 min, the reaction was mixed with 100 µL of 3,5-326 dinitrosalicylic acid (DNS) solution. At this point, the test tubes were placed in a boiling water bath 327 for 5 min, followed by cooling to room temperature. Next, each solution was diluted with 1 mL of 328 distilled water. Over time, the absorbance was measured at 540 nm by Thermo Scientific™ 329 Multiskan[™] GO Microplate Spectrophotometer using a 96-well microplate.

330 The α -amylase inhibitory activity was determined as follows:

Inhibition (%) =
$$100[(Abs_{Control} - Abs_{Sample})/(Abs_{Control})]$$

Where Abs_{Control} is the absorbance of the solution without sample (buffer instead of the sample solution) and Abs_{Sample} is the absorbance of the mixture with the solution of the sample to be tested. The concentration of the compound providing 50% inhibition (IC₅₀) was calculated by plotting inhibition percentages against sample concentrations. Acarbose was used as a positive control. Each assay was conducted in triplicate.

337 2.4. Molecular Docking procedure

338 The chemical structures of acarbose and the synthesized compounds (1, 2 and 2a-m) were 339 generated, optimized and their energies were minimized using ACD (3D viewer) software [56]. The 340 crystal structure of Aspergillus oryzae α -amylase protein (PDB: 7TAA) was downloaded from the 341 RCSB data bank [57]. The protein was prepared by removing water molecules and the complexed 342 inhibitor ligand. Then, the polar hydrogens were added followed by appending Kollman charges. Hence, the grid box, with dimensions of 65 x 52 x 52 points, spacing of 0.375 Å and centered with 343 344 coordinates x: 38.725, y: 44.127, and z: 25.633, was generated based on acarbose binding position 345 in the target protein binding site. The Molecular Docking analyzes of acarbose and the synthesized 346 compounds (1, 2 and 2a-m) were performed using AutoDock Vina software [58]. Molecule-enzyme
347 interactions were drawn and construed by employing the Biovia Discovery Studio Visualizer
348 (BIOVIA, D. S. (2017)).

349 *2.5. DFT studies*

350 The optimization of all geometries was performed by Density Functional Theory (DFT) using the 351 Gaussian 09 set of programs [59]. Geometry optimization of all molecules was carried out using 352 DFT methods at the B3LYP/6-311++G (d,p) level of theory [60] with Grimme D3 corrections, updated to include dispersion correlation [61] parameterized by Becke-Johnson (D3BJ). For these 353 354 systems, full geometry optimization in the C1 point group was carried out, followed by harmonic 355 frequencies simulations to confirm that they correspond to true minima (all positive frequencies). 356 All the calculations were done in DMF, using conductor like polarizable continuum (CPCM). The conceptual DFT descriptors [62-64] have been computed using the so-called frozen core 357 358 approximation and the Koopmans theorem [65]. In addition, the stability of flavonoid-based isoxazoles was investigated with natural bond orbital analysis (NBO) [66] using NBO6 to 359 360 determine the contribution of different interactions in the stability of these reactive compounds, 361 precisely compound 2b, the most bioactive one. This evaluation was done at B3LYP/6-311++G(d,p) level of theory. Compound **2b**, the most α -amylase inhibitor, was taken as a reference 362 363 in this study.

On the other hand, Quantitative Structure-Activity Relationship (QSAR) modeling was used as implemented in HyperChem 8.03 software [67] to identify the physicochemical properties of halogenated flavonoid-based isoxazole derivatives. Moreover, different types of descriptors viz. topological descriptors (polarizability (Pol), molar refractivity (MR), partition coefficient octanol/water (log P), molar volume (MV), surface area grid (SAG), molar weight (MW), electronic descriptors (LUMO (ELUMO) and HOMO (EHOMO) energies), and geometric descriptors (solvent accessible hardness and potential) were used [68-71].

371 2.6. Statistical analysis

372 Statistical analysis was carried out using Graph Pad Prism 7.0 (Graph Pad Software Inc., CA, and 373 USA). All the data are presented as the mean ± standard error of the mean (SEM). The difference 374 between two groups was evaluated using Student's t-test. Significant difference among three or 375 more groups was determined by one-way ANOVA with a post hoc analysis.

376

377 **3. Results and discussion**

378 *3.1. Chemistry*

The 5-bromo-2-hydroxyacetophenone was condensed with 4-(trifluoromethyl)benzaldehyde in the presence of NaOH for 3 hours at the reflux of methanol, yielding the 5-bromo-4'-(trifluoromethyl)-2-hydroxychalcone. Afterwards, the cyclization of the ring C and the oxidation of carbon C-3 are performed by adding H_2O_2 and NaOH. The reaction mixture was stirred for 2 hours at room temperature (Scheme 1). The compound **1** was obtained in 87% yield after precipitation in icewater.

385

386 Scheme 1. Synthetic pathway of the halogenated flavonol (1).

387

388 The structure of flavonol (1) was confirmed by spectroscopic analysis and by comparison with 389 literature data [32].

390 On the other hand, 1,3-dipolar cycloaddition is considered the most important synthetic way of a 391 wide range of isoxazoles. Furthermore, the regiospecific synthesis of isoxazoles catalyzed by Cu(I) 392 under microwave irradiation is based on the [3 + 2] cycloaddition between terminal alkynes 393 (dipolarophiles) and aryl nitrile oxides (dipoles) providing only 3,5-disubstituted regioisomers 394 [53,72]. The approach to target the halogenated flavonoid-based isoxazoles (**2a-m**), firstly started by the preparation of the dipolarophile **2** *via* the propargylation of the hydroxyl group in position C-3 of flavonol **1**, in anhydrous DMF for 2 hours at room temperature, in the presence of K_2CO_3 , as illustrated in Scheme 2. The dipolarophile **2** was prepared with an excellent yield (96%).

- 399
- 400

Scheme 2. Synthetic pathway of dipolarophile (2).

401

402 The dipolarophile structure (2) was established according to its spectral data. Indeed, in addition to 403 the signals corresponding to the protons and carbons introduced by the flavonol (1), new signals of the propargyl moiety were observed in the ¹H and ¹³C NMR spectra. The ¹H NMR spectrum of 2 404 405 recorded in the CDCl₃ at 300 MHz, showed a doublet at $\delta_{\rm H}$ 5.06 (2H, d, J = 2.4 Hz) attributable to 406 the methylene protons H-1" and a triplet at $\delta_{\rm H}$ 2.35 (1H, t, J = 2.4 Hz) attributable to the proton of the ethynyl group (H-3"). The ¹³C NMR spectrum confirmed the introduction of the propargyl 407 408 group by the observation of new signals at δ_C 78.4, 76.9 and 59.6 attributable to C-2", C-3" and C-409 1", respectively.

The 1,3-dipolar cycloaddition reaction was applied in a regiospecific approach using the terminal alkyne (2) and the different hydroximyl chlorides **a-m** variously substituted. Hydroximyl chlorides **a-m** are the main precursors for the *in situ* generation of aryl nitrile oxides. All the reactions were conducted under microwave irradiation (250 W) in DMF in the presence of triethylamine and CuI for 5 minutes (Scheme 3).

415

416 **Scheme 3.** Synthetic pathway of halogenated flavonoid-based isoxazoles **2a-m**.

417

The newly prepared 3,5-disubstituted isoxazoles (2a-m) were obtained in good yields ranging from
72 to 94%. Their structures are given in Figure 4.

420

421 422 Fig. 4. Structures of the halogenated flavonoid-based isoxazoles (2a-m).

423 The structures of the synthesized halogenated flavonoid-based isoxazoles **2a-m** (Figure 4) were determined by means of ¹H, ¹³C NMR and DEPT 135 spectra. The ¹H NMR spectra of these 424 425 compounds showed a singlet resonating at $\delta_{\rm H}$ 6.30-6.60 attributable to the methine proton H-4" of the isoxazole ring, a singlet at $\delta_{\rm H}$ 5.35-5.45 attributable to the methylene protons H-6" and other 426 427 signals in the aromatic proton zone relating to the protons introduced by the aryl group. Moreover, these structures were also confirmed by ¹³C NMR and DEPT 135 spectra, showing all the expected 428 429 signals from the carbons, essentially the aromatic carbons introduced by the used hydroximyl 430 chlorides, the carbon of the methine C-4" in the isoxazole ring resonating at $\delta_{\rm C}$ 102.5-103.0, and the 431 carbon whose signal is inverted in the DEPT 135 spectrum resonating at $\delta_{\rm C}$ 63.5-63.9 attributable to 432 the unique methylene C-6".

433 *3.2. Evaluation of α-Amylase inhibition*

Inhibition is one of successful therapeutic means for controlling the activity of enzymes. Therefore, inhibiting the enzymatic activity of the α -amylase is a good way to manage diabetes, obesity and to control overweight, by hindering carbohydrate digestion. In this article, fifteen synthesized compounds (1, 2 and 2a-m) were evaluated for their α -amylase inhibitory activities. The results were expressed in IC₅₀ (µM) and given in Table 1.

According to the results, the synthesized molecules (1, 2 and 2a-m) possess significant anti- α amylase activity with IC₅₀ values ranging from 16.2 ± 0.3 to 33.3 ± 0.4 µM. The cycloadduct 2b (R₁= F, R₂= H; IC₅₀= 16.2 ± 0.3 µM) exhibited the highest α -amylase inhibitory activity, this significant value was found to be comparable to that of the standard substance used (Acarbose, IC₅₀= 15.7 ± 0.2 µM). The compounds 2h, 2i, 2k and 2m displayed considerable α -amylase inhibitory effects with IC₅₀ values ranging from 17.3 ± 0.1 to 19.3 ± 0.4 µM. The rest of compounds were found to be slightly less active against the α -amylase enzyme (IC₅₀ = 20.8 ± 0.2 - 33.3 ± 0.4 μ M). The inhibitory α -amylase activity carried out by the newly synthesized compounds is consistent with previous reports. Nie et al. (2020) [54] reported that flavonoid isoxazole compounds could display valuable scaffolds for the discovery of antidiabetic drugs.

449 Furthermore, the halogenated compounds in *para* position of the phenyl group attached to the isoxazole ring (2b, 2c and 2d) performed significant inhibitions of α -amylase enzyme, with IC₅₀ 450 451 values ranging from 16.2 ± 0.3 to $21.2 \pm 0.2 \mu$ M. This finding is in good agreement with the 452 literature, showing that the presence of halogen atoms (F, Cl and Br) in the structure was essential 453 for a potent α -amylase inhibition [33,41]. In addition, the structure-activity relationship study 454 allowed to conclude that the more the inductive attractor (-I) and mesomeric donor (+ M) effects 455 increase, the greater the inhibitory power of α -amylase (activity of fluorinated derivative **2b** (R₁= F, 456 $R_2 = H$; $IC_{50} = 16.2 \pm 0.3 \mu M$) is higher than that of the chlorinated one 2c ($R_1 = Cl, R_2 = H$; $IC_{50} =$ 457 20.8 \pm 0.2 μ M), followed by the brominated one 2d (R₁= Br, R₂= H; IC₅₀= 21.2 \pm 0.2 μ M)). These 458 findings were found to be consistent with literature data indicating that the α -amylase inhibition 459 increases with inductive attractor (-I) and mesomeric donor (+M) effects of halogen atoms [6,41].

460 On the other hand, compound **2h** (R_1 = OMe, R_2 = H; IC₅₀= 19.3 ± 0.4 µM), bearing a *para*-methoxy 461 group, exhibited an interesting α -amylase inhibitory activity. This finding is in agreement with 462 previous results [41]. Moreover, compound 2i ($R_1 = O$ -*n*-Bu, $R_2 = H$; IC₅₀= 17.3 ± 0.1 μ M), bearing 463 a *para*-substituted butoxy group, perceived higher activity than that of *para*-methoxylated one (2h). 464 The electronic effects of alkoxyl groups (-I and +M) comparable to those of halogens, may explain 465 their significant activities. Besides, the methoxylation or the halogenation in the *meta* position of the *para*-alkoxylated compounds (2h and 2i), to give the compounds 2j-m (IC₅₀= $17.6 \pm 0.3 - 30.9$ 466 467 $\pm 0.5 \,\mu$ M), did not show a well-defined effect, they reduced the enzyme activity of the α -amylase in 468 some cases and increased it in others, but 2i remains the most active alkoxylated derivative.

469 Likewise, the compound 2g ($R_1 = NO_2$, $R_2 = H$; IC₅₀= 24.7 ± 0.3 μ M), bearing a *para*-nitro group, 470 revealed a moderate activity, in agreement with the report of Nawaz et al. (2020) [41]. On the other 471 hand, unsubstituted (2a: $R_1 = H$, $R_2 = H$; $IC_{50} = 33.2 \pm 0.8 \mu M$), *para*-methylated (2e: $R_1 = CH_3$, $R_2 =$ 472 H; IC₅₀= 32.8 ± 1.0 μ M) and *para-tert*-butylated (**2f**: R₁= *tert*-Bu, R₂= H; IC₅₀= 33.3 ± 0.4 μ M) 473 compounds did not offer remarkable activities compared to their aforementioned analogues. This 474 finding shows that the increase in the donor inductive effect (+I) does not have practically an effect 475 on the α -amylase activity. This result is consistent with the reports of Nawaz et al. (2020) and 476 Kanwal et al. (2021) [6,41].

477

478 **Table 1**

479 α -Amylase inhibition (IC₅₀ μ M), binding energy (kcal/mol) and interaction detail of compounds **1**, **2** and 480 **2a-m** docked in the ABC pocket (PDB: 7TAA).

481

482 *3.3. Molecular Docking Studies*

The *in silico* experiments can show the ability of molecules to fit well into the active pocket of the target enzymes. The binding to several residues, especially with the key residues, can explain the experimental activities of the tested compounds [73-77].

486 α -Amylase enzyme (EC 3.2.1.1), a glycosylhydrolase, hydrolyzes α -1,4-glycosidic bonds in starches, such as amylose. The crystal structure of α -amylase enzyme (PDB: 7TAA) is composed of 487 488 a chain A with a sequence length of 478 amino acids [78]. α -Amylase is one of the most important 489 key enzymes responsible for carbohydrate digestion [7]. Thus, blocking α -amylase enzyme offers a 490 strategy to controlling diabetes and helping in the management of obesity [4,5]. Extensive 491 molecular docking analysis was executed to explore the binding mode, determine the likely 492 interactions of synthesized molecules (1, 2 and 2a-m) within the hydrophobic binding pocket of 493 ABC (Modified acarbose hexasaccharide) of the protein crystal structure of Aspergillus oryzae α - 494 amylase enzyme (PDB: 7TAA) [78] using AutoDock Vina software and rationalize the observed *in* 495 *vitro* α -amylase inhibitory activities of the newly synthesized compounds.

All the synthesized molecules (**1**, **2** and **2a-m**) and acarbose, used as a standard substance, were *in silico* studied. Binding energies and interaction details (Number of interactions, number of interacting amino acids, interacting amino acids and hydrogen bonds) of ligands with target enzyme are presented in Table 1.

500 From the *in silico* docking results (Table 1), it was observed that compounds 1 and 2 exhibited good 501 binding energies (-7.5 and -7.6 kcal/mol, respectively), within the binding site of the α -amylase 502 enzyme (PDB: 7TAA). These values are comparable to that of the standard substance (Acarbose, -503 7.9 kcal/mol), which will allow them to fit favorably into the active site of the target enzyme. 504 Interestingly, the newly synthesized halogenated flavonoid-based isoxazoles (2a-m) gave the 505 expected results. They are properly in the bonding pose and exhibited magnificent binding energies 506 ranging from -9.6 to -8.0 kcal/mol (Table 1). This would allow everyone to fit favorably into the 507 active site of the α -amylase enzyme, better than acarbose.

508 The binding poses of acarbose and the four potent α -amylase inhibitor compounds (2b, 2i, 2k and 509 2m) in the target enzyme are shown in Figure 5.

- 510
- 511

Fig. 5. The binding pose of acarbose, 2b, 2i, 2k and 2m in α -amylase enzyme.

512

513 Furthermore, the binding mode of acarbose and the four most active compounds (2b, 2i, 2k and 514 2m) with α -amylase enzyme are discussed in the following.

515 Acarbose, the compound used as a standard inhibitor substance of α -amylase enzyme, exhibited a 516 good binding energy of -7.9 kcal / mol. Its mode of binding shows that it is involved in twelve non-517 covalent interactions with nine amino acids (Table 1). Perfectly, it formed nine conventional 518 hydrogen bonds with the residues GLN-35, HIS-80, TRP-83, ASP-206, LYS-209, GLU-230, LEU- 519 232, ASP-297 and ASP-340. In addition, three carbon hydrogen bonds have been implicated with
520 LYS-209 and ASP-340 (Figure 6).

521

Fig. 6. Acarbose and compound 2b, 2i, 2k et 2m fits into the hydrophobic binding pocket of ABC
in PDB: 7TAA.

524

525 Compound 2b, the most potent *in vitro* inhibitor of α -amylase enzyme in the tested series of 526 halogenated flavonoid-based isoxazoles, exhibited an excellent binding energy of -9.6 kcal/mol. Its 527 binding mode shows that it is implicated in fifteen non-covalent interactions with nine amino acids 528 (Table 1). Perfectly, the 3-(3-(4-fluorophenyl)isoxazol-5-yl)methoxyl moiety forms three halogen 529 (Fluorine) bonds with GLU-230 and ASP-206 residues, a Pi-Pi T-shaped bond with TYR-82, a Pi-530 cation and conventional hydrogen bonds with ARG-344, and Pi-anion and carbon hydrogen bonds 531 with ASP-340 residue. Additionally, the trifluoromethyl group linked to the aromatic ring B of the 532 flavonoid skeleton is involved in two alkyl interactions with LEU-166 and LEU-173. Moreover, 533 flavonoid skeleton is involved in a conventional hydrogen binding with GLN-35, a Pi-anion bond 534 with ASP-340, and two Pi-Pi Stacked and a Pi-alkyl bonds with TYR-75 (Figure 6).

535 The compound 2i carried out an attractive binding energy of -9.3 kcal/mol. The binding mode of 536 this derivative shows that it is involved in seventeen non-covalent interactions with eleven amino 537 acids (Table 1). Consequently, the 3-(3-(4-butoxyphenyl)isoxazol-5-yl)methoxy group interacts in 538 the active binding site by a carbon hydrogen bond with GLU-230, an alkyl bond with LEU-232, a 539 Pi-alkyl binding with HIS-210, a Pi-Pi T-shaped bond with TYR-82, a Pi-cation and two 540 conventional hydrogen bonds with ARG-344, a carbon hydrogen and a Pi-anion interactions with 541 ASP-340 residue. Furthermore, the trifluoromethyl moiety is involved in a conventional hydrogen 542 bond with GLY-167, and two alkyl interactions with LEU-166 and LEU-173. In addition, flavonoid 543 skeleton is involved in a conventional hydrogen bond with GLN-35, a Pi-anion and a carbon 544 hydrogen bonds with ASP-340, and two Pi-Pi Stacked and a Pi-alkyl bonds with TYR-75 (Figure545 6).

546 On the other hand, the compound $2\mathbf{k}$ exhibited a significant binding energy of -8.8 kcal/mol. Its 547 binding mode shows that it is implicated in fourteen non-covalent interactions with eleven amino 548 acids (Table 1). Thoroughly, the 3-(3-(3-chloro-4-methoxyphenyl)isoxazol-5-yl)methoxyl moiety 549 forms two conventional hydrogen bonds with TYR-83 and HIS-80, two carbon hydrogen bonds 550 with ASP-206 and ASP-297, two Pi-alkyl interactions with HIS-122 and TYR-82, a Pi-Pi Stacked 551 binding with TYR-82, and a Pi-cation bond with the ARG-344. Furthermore, the trifluoromethyl group attached to C-4' of the flavonoid skeleton is involved in two conventional hydrogen bonds 552 553 with GLY-167 and TRP-83, and an alkyl bond with the LEU-173. On the other hand, flavonoid 554 skeleton interacts by a conventional hydrogen binding with GLN-35 and two Pi-Pi Stacked bonds 555 with TYR-75 (Figure 6).

556 Furthermore, the compound **2m** exhibited an interesting binding energy of -8.7 kcal/mol. Its binding 557 mode shows that it is involved in fourteen non-covalent interactions with seven amino acids (Table 558 1). Accordingly, molecular insight of docking analysis suggests that the 3-(3-(3,4-559 dimethoxyphenyl)isoxazol-5-yl)methoxyl moiety forms three carbon hydrogen bonds with ASP-560 206 and GLU-230 residues, two Pi-sigma and a Pi-alkyl bonds with the LEU-166. Besides, the 561 trifluoromethyl group is involved in a conventional hydrogen binding with HIS-122, a halogen 562 (Fluorine) bond with TYR-82, two Pi-alkyl bonds with TYR-82 and HIS-122 and an alkyl 563 interaction with the LEU-173. In addition, the flavonoid skeleton is involved in a Pi-alkyl and two 564 Pi-Pi Stacked bonds with the TYR-75 residue (Figure 6).

565 In the Aspergillus oryzae α -amylase enzyme (PDB: 7TAA), the nucleophile and the catalytic 566 acid/base residues, which are responsible for the cleavage of glycosidic bonds, are ASP-206 and 567 GLU-230 amino acids, respectively [7]. Therefore, interactions between protein and ligand, 568 primarily *via* the GLU-230 and ASP-206 residues, are the best way to block the cleavage of

569 glycosidic bonds. In fact, the low binding energy, the correct binding pose, the ability to interact 570 with key residues of glycosidic cleavage (GLU-230 and ASP-206) and the large number of 571 interactions in the active pocket of the target enzyme can explain the strong inhibitory α -amylase 572 effects establishing by almost compounds.

573 *3.4. DFT studies*

574 3.4.1. Vibrational analysis

575 The anharmonic frequencies, certain characteristic sites of the most active product: 6-bromo-3-((3-576 (4-fluorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one (2b,577 IC₅₀= 16.2 \pm 0.3 μ M), are given in Table 2. These data are computed at B3LYP/6311++G (d,p) 578 levels of theory and using second order perturbation theory (VPT2) approach as implemented in 579 GAUSSIAN. This flavonoid-based isoxazole derivative is a planar molecule. It belongs to the Cs 580 point group. It possesses 94 a' and 50 a" vibrational modes (Table S1). Since there are no 581 experimental spectra available for this compound, our data can be used for the assignment of these 582 spectra whenever measured.

583

Table 2

585 Anharmonic vibrational frequencies (in cm⁻¹) of 6-bromo-3-((3-(4-fluorophenyl)isoxazol-5-586 yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one as computed using B3LYP in 587 conjunction with the 6-311++ G (d,p) basis set. Also their assignment was given.

588

589 3.4.2. Characterization of the halogenated flavonoid-based isoxazoles

590 Structures of the thirteen halogenated flavonoid-based isoxazoles 2a-m are optimized at the 591 B3LYP/ 6311++G (d,p) level of theory. The optimized structures of newly synthesized molecules 592 are given in Table S2 of the supplementary material. Close examination of these structures of the 593 halogenated flavonoid-based isoxasoles shows that the 3-phenylisoxazole moiety within these 594 compounds has a very close structure. Indeed, we compute C=O, C=C, C=N, N-O and O-C 595 distances of 1.24, 1.43, 1.32, 1.39, 1.37 (all values are given in Å), respectively.

596 Thus, the functionalization of the isoxasole ring slightly disturb its structure. This is confirmed by 597 the vibrational analysis performed for this series of analogs. Anharmonic frequencies can be 598 obtained after scaling the corresponding harmonic frequencies by the appropriate scaling factor (by 0.9662 for $\omega > 1000$ cm⁻¹ and by 0.9833 for $\omega < 1000$ cm⁻¹) as discussed in previous report [79]. 599 Table 3 presents the respective data for C=O, N-O, C=N, and C-O elongations. This table shows 600 that similar C=N (of~1443.7 cm⁻¹), C=O (of ~1604.9 cm⁻¹), O-N (of ~1271.8 cm⁻¹), C-O (of 601 ~1362.3 cm⁻¹), elongations values are computed for all compounds. Indeed, they differ by less than 602 20 cm⁻¹ within the series. Again, this is due to slight perturbation of the heterocycle upon 603 604 functionalization.

605

606 **Table 3**

607 Scaled anharmonic vibrational frequencies (in cm⁻¹) (scaling factor 0.9662 [79]) of some elongation 608 modes of flavonoid-based isoxazole derivatives as computed using B3LYP/6-311++G (d,p).

609

610 3.4.3. Natural bond orbital (NBO) and molecular electrostatic potential (MESP) surface of
611 flavonoid-based isoxazole derivatives (2a-m)

The information contained in molecular electrostatic potentials (MESP) has been harnessed for determination of structure, reactivity, electron diffraction and scattering and also the energetics of a wide variety of organic and inorganic, as well as biological molecular systems [80]. In fact, MESP displays simultaneously the molecular size and shape as well as positive, negative and neutral electrostatic potential regions in terms of the electrostatic surface [81]. Figure 7 shows the MESP surface map of the most active compound (**2b**: 6-bromo-3-((3-(4-fluorophenyl))isoxazol-5vl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4*H*-chromen-4-one; IC₅₀= 16.2 ± 0.3 μ M). The red color 619 refers to negative potential regions as around the oxygen atom (carbonyl/ C=O) of the pyran ring. 620 For nitrogen, we compute mostly negative potentials with different magnitudes. These regions are 621 susceptible for electrophilic attacks. Figure 7 shows also that the regions around the hydrogens 622 attached to the phenyl ring are blue (positive potentials). These regions are susceptible for 623 nucleophilic attacks. This analysis agrees with the net charges calculated using natural bond orbital 624 (NBO) model (Figure 8). The present analyses allow identifying the hydrogen bond donor (HBD) 625 and hydrogen bond acceptor (HBA) sites (Figure 7). These characteristics may be helpful for a 626 qualitative understanding of the electrostatic interactions that may take place between reagents or 627 enzyme active sites and the flavonoid-based isoxazole derivatives (2a-m) under study.

628

Fig. 7. 3D MESP surface map (left) for 6-bromo-3-((3-(4-fluorophenyl)isoxazol-5-yl)methoxy)-2(4-(trifluoromethyl)phenyl)-4*H*-chromen-4-one (2b) and its hydrogen bond acceptor (HBA) sites
(right). The red regions correspond to negative potentials and the blue regions to positive potentials.

Fig. 8. B3LYP/6-311++G (d,p) net charges calculated using natural bond orbital (NBO) in
 compound 2b.

635

636 3.4.4. Structural and physicochemical property relationships

The structures of the flavonoid-based isoxazole derivatives are shown in Figure 4. QSAR properties of compounds **2a-m**, such as octanol-water partition coefficient (logP), molar refractivity (MR), polarizability (Pol), surface bounded molecular volume, molecular weight (MW), Hardness (η), potential (μ), HOMO energy (E_{HOMO}, energy of the highest occupied molecular orbital) and LUMO energy (E_{LUMO}, energy of the lowest unoccupied molecular orbital) [82], were listed in Table 4. The polarizability and molar refractivity are connected to the size and the molecular weight of the halogenated flavonoid-based isoxazoles (**2a-m**). This result is in agreement with the formula of 644 Lorentz-Lorenz [83-84] which provides a relation between polarizability, molar refractivity and 645 volume. These findings can be proven by the compound 2a (6-bromo-3-((3-phenylisoxazol-5-646 yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one, $R_1 = H$, $R_2 = H$), the smallest molecule of the newly synthesized series, whose molecular weight, polarizability and molar 647 refractivity are equal to 542.31 amu, 47.79 Å³ and 140.54Å³, respectively. On the other hand, the 648 649 (6-bromo-3-((3-(4-butoxy-3-chlorophenyl)isoxazol-5-yl)methoxy)-2-(4compound 2i 650 (trifluoromethyl)phenyl)-4H-chromen-4-one, $R_1 = O$ -*n*-Bu, $R_2 = Cl$) has the highest molecular 651 weight (648.86 amu), therefore we note a growth in the polarizability and molar refractivity, 57.69 $Å^3$ and 165.50 $Å^3$, respectively. 652

Several physicochemical descriptors, derived from conceptual DFT, are given in Table 4. They describe the reactivity of the studied molecules and can explain the effect of various substituents on our core [85], such as chemical potential μ ($\mu = \left(\frac{\partial \mathbf{E}}{\partial N}\right)_{\nu(\mathbf{r})}$) and hardness η ($\eta = \left(\frac{\partial^2 \mathbf{E}}{\partial N^2}\right)_{\nu(\mathbf{r})}$) with $E_{\nu(\mathbf{r})}$ is the total energy of the system, N: number of electrons and v(r) indicates that the derivatives are taken at constant external potential [86]. These indexes are often approximated through frontier orbital eigenvalues, leading to $\mu \cong (\mathbf{E}_{HOMO} + \mathbf{E}_{LUMO})/2$ and $\eta \cong (\mathbf{E}_{LUMO} - \mathbf{E}_{HOMO})$ [87].

659 HOMO and LUMO energies are related to the reactivity of the molecule: molecules with electrons at accessible (near-zero) HOMO levels tend to be good nucleophiles because it does not cost much 660 661 to donate these electrons toward making a new bond. Similarly, molecules with lower LUMO energies tend to be good electrophiles because it does not cost much to place an electron into such 662 663 an orbital. The energy gap which expresses the energy difference between two important FMOs 664 (HOMO-LUMO) [88], signifies the chemical hardness-softness, the kinetic stability and the reactivity of molecules. Hard molecules with large gap are more stable and less reactive. In the 665 666 opposite case, soft molecules having a gap are more reactive and less stable [89-90]. As can be 667 seen, compound **2**c (6-bromo-3-((3-(4-chlorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4*H*-chromen-4-one, $R_1 = Cl$, $R_2 = H$) has the highest value of LUMO (E = -668

669	2.651eV) with energy gap (ΔE = 4.147 eV), hardness (η = 2.072 eV) and potential (μ = -4.725 eV).
670	Thus, this compound is the most stable one (Figure 9). Compound 2i (6-bromo-3-((3-(4-
671	butoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4 <i>H</i> -chromen-4-one, $R_1 = O_1$
672	butyl, R ₂ = H) has the highest HOMO (-6.224 eV) and the smallest ΔE value (ΔE = 3.578 eV). Thus,
673	this compound is the most reactive one (Figure 9). Indeed, a group with a negative inductive effect
674	(-I) like (Cl, F, OCH ₃) substituted in para position of the 3-phenylisoxazole core have an
675	influence on the α -amylase inhibition activity for instance 2c (IC ₅₀ = 20.8 ± 0.2 μ M), on the other
676	hand 2i (IC ₅₀ = 17.3 ± 0.1 μ M). This order is proportion to the noted biological activity. In overall,
677	this result supports the experimental trend.

678

679 **Fig. 9.** HOMO, LUMO orbitals and their energy gap (ΔE gap) of the compounds **2c** and **2i** at 680 B3LYP/6–311++G (d, p) level of DFT.

681

For good absorption and permeability, the molecules must satisfy the Lipinski rules [91]. The rule describes the important molecular properties for a drug pharmacokinetics in the human body, including their absorption, distribution, metabolism and excretion. Lipinski's rule states that, LogP ≤ 5 , molecular weight (MW) ≤ 500 Da, H-bond donors (HBD) ≤ 5 and H-bond acceptors (HBA) \leq 10.

The lipophilicity is proportional to the hydrophobic character of the substituent group [92]. The LogP is the most useful parameter for the characterization of hydrophobicity (and polarity) of compounds. It is an important variable in QSAR studies because the distribution of chemicals between fatty and aqueous phases of a biological system could totally account for the variation in activities. A positive value for logP denotes a higher concentration in the lipid phase. A drug targeting the central nervous system (CNS) should ideally have a logP value around 2 for oral and intestinal absorption the idea value is 1.35–1.8, while a drug intended for sub-lingual absorption should have a logP value >5. All our compounds respect this rule [93].

Table 4 reveals logP (3.53-0.23), compound **2f** (6-bromo-3-((3-(4-(tert-butyl)phenyl)isoxazol-5yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one, R_1 = tert-butyle, R_2 = H) has high value of logP (3.53). This compound generally has a good intestinal absorption owing to a good balance between solubility and passive diffusion permeability. In fact, the metabolism is minimized because of the lower binding with metabolic enzymes.

The most active compounds (**2b**, **2i**, **2k** and **2m**) have logP value (1.62, 2.43, 1 and 0.23, respectively). They are very lipophilic. Therefore, they have a good permeability through biological membrane, a better binding to plasma proteins, elimination by metabolism but a poor solubility.

To a Lipophilic ligand efficiency (LipE) is defined as: LipE = pIC50 - logP. As a rough guide, medicinal

compounds in drug-like space have LipE values in the range 2-7. Compound **2m** (6-bromo-3-((3-

705 (3,4-dimethoxyphenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one,

 R_1 = OCH₃, R_2 = OCH₃) has the highest LipE value (4.50) and it is deemed to be the most optimal compound (Table 4).

Lipinski's rule states that, $LogP \le 5$, molecular weight (MW) ≤ 500 Da, H-bond donors (HBD) ≤ 5 and H-bond acceptors (HBA) ≤ 10 . If molecules violating more than one of these parameters, they can have problems with bioavailability and high probability of failure to display drug-likeness.

Therefore, the obtained results reveal that the thirteen halogenated flavonoid-based isoxazole compounds respect the Lipinski rules except for the rule relating to molecular mass molecular weight (MW) \leq 500 Da. Indeed, when the molecular weight is less than 500 Da it can be easily transported, diffused and absorbed through the cell membrane compared to heavy molecules. All selected compounds have (MW) > 500 Da. However, in 2014, Doak et al. [94] shown that strict reliance on the rule of 5 (Ro5) may have resulted in lost opportunities, particularly for difficult targets. To identify opportunities for oral drug discovery beyond the Ro5 (bRo5), they have comprehensively analyzed drugs and clinical candidates with molecular weight (MW) > 500 Da. They conclude that oral drugs are found far bRo5 and properties such as intramolecular hydrogen bonding, macrocyclization, dosage and formulations can be used to improve bioavailability. Therefore, it suggested that these compounds do not have any problems with oral bioavailability.

722

723 **Table 4**

Values of chemical descriptors used in the regression analysis. The energies of the HOMO (E_{HOMO} , eV) and LUMO (E_{LUMO} , eV), energy gap (ΔE , eV), hardness (η , eV), softness, (S, eV), potential (μ , eV), electrophilicity (μ , eV) octanol-water partition coefficient (log P), polarizability (Pol, Å³), molar refractivity (MR, Å³), volume (V, Å³), surface area (SAG, Å²), molar weight (MW, amu). HBA and HBD are the numbers of the hydrogen bond acceptor and donor sites.

729

730 Conclusion

731 In summary, this work reported a design and synthesis of new molecules described as halogenated flavonoid-based isoxazoles. Chemical structures were established by ¹H NMR, ¹³C NMR and 732 733 HRMS analysis. The obtained compounds were evaluated for their α -amylase inhibitory activities 734 and excellent results were noted. In addition, analysis of the structure-activity relationship revealed 735 that alkoxylated and halogenated derivatives were found to be the most active compounds. Based 736 on the *in silico* docking analysis, the halogenated flavonoid-based isoxazoles showed low binding 737 energies, a large number of interactions and correct binding poses in the active pocket of the target 738 enzyme. In addition, all the synthesized cycloadducts were modeled by DFT calculations using 6-739 311++G (d,p) basis set. The chemical stability, electrical transport properties, the geometrical 740 parameters and frontier molecular orbitals of the halogenated flavonoid-based isoxazoles 2a-m 741 were determined and analyzed. Quantitative effects of the molecular structure of the cycloadducts

742 **2a-m** on their α -amylase inhibitory activity were found. The obtained results made it possible to 743 conclude that the newly synthesized compounds have no problem of oral bioavailability. These 744 hybrid molecules can be considered as promising candidates for investigating new anti-obesity 745 agents and antidiabetics.

746

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750

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

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764

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Fig. 1. Example of flavonoids previously reported as antidiabetics.



Fig. 2. Structures of trifluoromethylated drugs.



Fig. 3. (a) Natural compounds containing isoxazole ring (b) Previously reported isoxazole derivatives as antidiabetic agents (c) Reported flavonoid-based isoxazoles with antidiabetic properties.



Fig. 4. Structures of halogenated flavonoid-based isoxazoles (2a-m).



Fig. 5. The binding pose of acarbose, 2b, 2i, 2k and 2m in α -amylase enzyme.



Fig. 6. Acarbose and compound 2b, 2i, 2k et 2m fits into the hydrophobic binding pocket of ABC in PDB: 7TAA.



Fig. 7. 3D MESP surface map (left) for 6-bromo-3-((3-(4-fluorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4*H*-chromen-4-one (**2b**) and **its** hydrogen bond acceptor (HBA) sites (right). The red regions correspond to negative potentials and the blue regions to positive potentials.



Fig. 8. B3LYP/6-311++G (d,p) net charges calculated using natural bond orbital (NBO) in

compound 2b.



Fig. 9. HOMO, LUMO orbitals and their energy gap (ΔE gap) of the compounds 2c and 2i at B3LYP/6–311++G (d, p) level of DFT.

 α -Amylase inhibition (IC₅₀ μ M), binding energy (kcal/mol) and interaction detail of compounds 1, 2 and

Comp.	α-Amylase	Binding	Interaction detail:
	inhibition	energy	NI/NIAA: IAA
	$(IC_{50} \mu M)$	(kcal/mol)	
1	21.47 ± 0.62	-7.5	10/6: GLN-35*, TYR-75, TYR-82, HIS-122*, ASP-206, ASP-
			340
2	24.32 ± 0.46	-7.6	14/9: TYR-75, HIS-80, TYR-82, TRP-83, HIS-210*, GLU-230,
			HIS-296, ASP-340, ARG-344
2a	33.15 ± 0.77	-9.3	9/7: GLN-35*, TYR-75, HIS-80, TYR-82, LEU-166, ASP-340,
			ARG-344*
2b	16.20 ± 0.31	-9.6	15/9 : GLN-35*, TYR-75, TYR-82, LEU-166, LEU-173, ASP-
			206 , GLU-230 , ASP-340, ARG-344*
2c	20.76 ± 0.19	-8.7	12/7 : HIS-80, TYR-82, TRP-83, TYR-155, LEU-166, GLY-
			167*, LEU-173, ASP-206, LEU-232, ASP-340
2d	21.17 ± 0.16	-8.3	14/8: TYR-75, TYR-82, HIS-122*, LEU-166, ASP-206, LYS-
			209, HIS-210, LEU-232
2e	32.80 ± 1.05	-9.5	12/8 : GLN-35*, TYR-75, TYR-82, LEU-166, LEU-173, ASP-
			297, ASP-340, ARG-344*
2f	33.26 ± 0.41	-8.5	12/9: GLN-35*, TYR-75, HIS-80, TYR-82, TRP-83, GLY-167*,
			LEU-173, ASP-340, ARG-344*
2g	24.71 ± 0.33	-8.6	11/11: GLN-35*, TYR-75, HIS-80, TYR-82, LEU-166, GLY-
			167*, LEU-173, ARG-204*, HIS-296*, ASP-340, PRO-341
2h	19.25 ± 0.39	-8.7	12/9: HIS-80*, TYR-82, LEU-166, GLY-167*, LEU-173, TYR-
			155, ASP-206 , LEU-232, ASP-340
2i	17.33 ± 0.13	-9.3	17/11 : GLN-35*, TYR-75, TYR-82, LEU-166, GLY-167*, LEU-
			173, HIS-210, GLU-230, LEU-232, ASP-340, ARG-344**
2j	23.99 ± 0.29	-8.0	12/10: GLN-35*, TYR-75, TYR-82, TRP-83*, HIS-122, GLY-
			167*, LEU-173, ASP-297, PRO-341, ARG-344
2k	17.58 ± 0.26	-8.8	14/11: GLN-35*, TYR-75, HIS-80*, TYR-82, TRP-83**, HIS-
			122, GLY-167*, LEU-173, GLU-230 , ASP-297, ARG-344
21	30.86 ± 0.47	-9.4	13/8: GLN-35*, TYR-75, HIS-80, TYR-82, LEU-166, ASP-297,
			ASP-340, ARG-344**
2m	18.36 ± 0.14	-8.7	14/7 : TYR-75, TYR-82, HIS-122*, LEU-166, LEU-173, ASP-
			206, GLU-230
Acarbose	15.74 ± 0.21	-7.9	11/9: GLN-35*, HIS-80*, TRP-83*, ASP-206*, LYS-209*,
			GLU-230*, LEU-232*, ASP-297*, ASP-340*

2a-m docked in the ABC pocket (PDB: 7TAA).

NI: Number of interactions, NIAA: Number of interacting amino acids, IAA: Interacting amino acids, *= One hydrogen bond, **= Two hydrogen bonds.

Anharmonic vibrational frequencies (in cm⁻¹) of 6-bromo-3-((3-(4-fluorophenyl)isoxazol-5-yl)methoxy)-2-(4-(trifluoromethyl)phenyl)-4H-chromen-4-one as computed using B3LYP in conjunction with the 6-311++ G (d,p) basis set. Also their assignment was given.

Sym.	B3LYP	Assignment
a'	3179.1	C–H stretching
a'	3155.1	C–H stretching
a'	1606.0	C=C stretching
a'	1604.9	C=O stretching
a'	1443.9	C=N stretching
a'	1362.3	C–O stretching
a'	1271.3	N–O in-plane bending
a'	677.8	C-Br stretching
a'	967.6	N–O–C in-plane-bending
a'	661.5	Wagging C=O
a"	622.4	Molecule deformation in-plane-bending
a"	575.6	Molecule deformation in-plane-bending
a"	268.5	Rings deformation out-of-plane-bending
a"	227.4	Molecule deformation out-of-plane-bending
a"	201.1	Molecule deformation out-of-plane-bending

Scaled anharmonic vibrational frequencies (in cm⁻¹) (scaling factor 0.9662 [74]) of some elongation modes of flavonoid-based isoxazole derivatives as computed using B3LYP/6-311++G (d,p).



Values of chemical descriptors used in the regression analysis. The energies of the HOMO (E_{HOMO} , eV) and LUMO (E_{LUMO} , eV), energy gap (ΔE , eV), hardness (η , eV), softness, (S , eV), potential (μ , eV), electrophilicity (μ , eV) octanol-water partition coefficient (log P), polarizability (Pol, Å³), molar refractivity (MR, Å³), volume (V, Å³), surface area (SAG, Å²), molar weight (MW, amu). HBA and HBD are the numbers of the hydrogen bond acceptor and donor sites.

Comp.	PIC50*	LUMO	HOMO	ΔE	S	μ	η	щ	LipE	LogP	MW	Pol	R	SAG	Vol	HBD	HBA
2a	4.479	-2.654	-6.785	4.131	0.242	-4.720	2.065	-5.393	2.259	2.22	542.31	47.79	140.54	715.55	1234.17	0	5
2b	4.790	-2.655	-6.798	4.142	0.241	-4.726	2.071	-5.393	3.170	1.62	560.30	47.70	140.67	720.80	1243.37	0	5
2c	4.682	-2.651	-6.799	4.147	0.241	-4.725	2.070	-5.383	2.682	2.00	576.75	49.71	145.26	750.2	1286.74	0	5
2d	4.674	-2.651	-6.786	4.135	0.241	-4.719	2.067	-5.385	2.404	2.27	621.20	50.41	148.07	754.11	1305.49	0	5
2e	4.484	-2.652	-6.686	4.033	0.247	-4.669	2.016	-5.405	2.114	2.37	556.34	49.62	144.82	751.15	1292.68	0	5
2f	4.478	-2.647	-6.686	4.039	0.247	-4.667	2.019	-5.392	0.948	3.53	598.42	55.13	158.45	819.07	1431.13	0	5
2g	4.607	-3.273	-6.807	3.533	0.283	-5.040	1.766	-7.189	3.207	1.40	587.31	49.50	146.76	752.84	1300.06	0	8
2h	4.715	-2.645	-6.293	3.648	0.274	-4.469	1.824	-5.475	3.485	1.23	572.33	50.26	146.92	765.39	1318.05	0	6
2i	4.761	-2.645	-6.224	3.578	0.279	-4.435	1.789	-5.496	2.331	2.43	614.42	55.76	160.79	849.52	1477.87	0	6
2j	4.619	-2.650	-6.791	4.140	0.241	-4.720	2.070	-5.383	2.409	2.21	648.86	57.69	165.50	869.93	1518.66	0	6
2k	4.754	-2.649	-6.795	4.145	0.241	-4.722	2.072	-5.379	3.754	1.00	606.78	52.19	151.63	783.73	1360.29	0	6
21	4.510	-2.652	-6.789	4.137	0.241	-4.721	2.068	-5.386	2.230	1.28	542.31	47.79	140.54	715.55	1234.17	0	6
2m	4.736	-2.652	-6.565	3.913	0.255	-4.609	1.956	-5.428	4.506	0.23	560.30	47.70	140.67	720.80	1243.37	0	7
*PIC50= -logIC50																	



- ✓ Synthesis
- ✓ In vitro Inhibition of *a*-Amylase Activity
- ✓ Structure-Activity Relationship Analysis
- ✓ In silico Docking and DFT Studies

