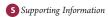
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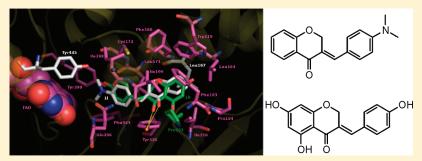
Homoisoflavonoids: Natural Scaffolds with Potent and Selective Monoamine Oxidase-B Inhibition Properties

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ABSTRACT:



A series of homoisoflavonoids [(E)-3-benzylidenechroman-4-ones 1a-w, 3-benzyl-4H-chromen-4-ones 2a-g, and 3-benzylchroman-4-ones 3a-e] have been synthesized and tested in vitro as inhibitors of human monoamine oxidase isoforms A and B (hMAO-A and hMAO-B). Most of the compounds were found to be potent and selective MAO-B inhibitors. In general, the (E)-3-benzylidenechroman-4-ones 1a-w showed activities in the nano- or micromolar range coupled with high selectivity against hMAO-B. The reduction of the exocyclic double bond results in compounds 3a-e selective against isoform B and active in the micromolar range. In contrast, the endocyclic migration of the double bond (compounds 2a-g) generally produces the loss of the inhibitory activity or a marked reduction in potency. (E)-3-(4-(Dimethylamino)benzylidene)chroman-4-one (11) and (E)-5,7-dihydroxy-3-(4-hydroxybenzylidene)chroman-4-one (1h) were the most interesting compounds of the entire series of inhibitors, showing hMAO-B affinity better than the selective inhibitor selegiline. Molecular modeling studies have been carried out to explain the selectivity of the most active homoisoflavonoids 1h and 11.

■ INTRODUCTION

Monoamine oxidase (MAO) is an ubiquitous membrane-bound, flavin-containing enzyme, which is particularly abundant in the liver and brain. MAO is located intracellularly in the mitochondrial outer membranes of neuronal, glial, and other cells and catalyzes the oxidative deamination of monoamine neurotransmitters and xenobiotic amines to the corresponding aldehydes with consumption of oxygen and production of hydrogen peroxide, thereby affecting the concentrations of neurotransmitter amines and many xenobiotic ones. 2—4

Currently, two MAO isoforms, encoded by separate genes sharing a common intron/exon organization,^{4–6} have been identified based on their differential substrate specificity and inhibitor sensitivity,^{7–9} tissue and cell distribution,¹⁰ and gene expression characteristics.^{11,12}

MAO-A preferentially deaminates serotonin, epinephrine, and norepinephrine and is selectively inhibited by low concentrations of clorgyline, whereas MAO-B preferentially metabolizes β -phenylethylamine and benzylamine and is inhibited by selegiline. Both isoenzymes deaminate dopamine, tyramine, and tryptamine. 1

Because of the important function played by MAO in the metabolism of neurotransmitters, MAO inhibitors can be useful in the treatment of many psychiatric and neurological diseases. In fact, MAO-A inhibitors act as antidepressant and antianxiety agents, whereas MAO-B inhibitors are used alone or in combination to treat Alzheimer's and Parkinson's diseases. ¹³

Recognition of the importance of MAO as a drug target for treatment of neurological disorders has produced an enormous interest in the development of inhibitors of these enzymes. Various structural classes of potent MAO inhibitors, including

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Figure 1. Some representative structures of nonselective and selective MAO inhibitors.

hydrazide, amide, thiazole, imidazole, oxazolidinone, oxadiazolone, diacylurea derivatives, etc., have been identified. ^{14–27} In spite of considerable progress in understanding the interactions of the two enzyme forms with their corresponding substrates, no general rules are yet available for the rational design of potent and selective MAO inhibitors. There are many different structures of MAO inhibitors partly because the active sites of the MAO enzymes are ambiguous, which limits the design of potent selective MAO inhibitors. Figure 1 shows the chemical structures of some representative MAO inhibitors used in research or clinical practice.

The aim of the present study was the identification of novel potent MAO inhibitors that could serve as potential lead molecules for drug discovery.

Homoisoflavonoids constitute a small class of natural products prevalently isolated from the bulbs, rhizomes, or roots of several genera of *Hyacinthaceae* and *Caesalpinioideae*. These compounds are structurally related to the more widespread flavonoids and, according to their structural features, can be classified into three types: 3-benzylidenechroman-4-ones, 3-benzyl-4*H*-chromen-4-ones, and 3-benzylchroman-4-ones. Several natural and synthetic homoisoflavonoids, like the related flavonoids, were found to possess various biological properties such as antifungal, antiviral, 29-31 antimutagenic, 32,33 antiproliferative, 34 antioxidant, 35,36 antiallergic and antihistaminic, 37 anti-inflammatory, 38 and protein tyrosine kinase (PTK) inhibitor activities. 39 However, no data are available on the inhibitory activity of homoisoflavonoids on MAOs

Recently, several substituted chalcones and flavanones were reported as potent and selective inhibitors of the B-isoform of human MAO (hMAO). 40,41 The structural relationship of homoisoflavonoids to chalcones and flavanones prompted us to investigate their inhibitory activity against the A and B isoforms of hMAO. Initially, we tested the synthetic (*E*)-3-benzylidenechroman-4-ones 1a-g, 3-benzyl-4-chromones 2a-g, and 3-benzylchroman-4-ones 3a-e, previously studied as antipicornavirals. $^{29-31}$ Because all the (*E*)-3-benzylidenechroman-4-ones 1a-g, which may be regarded as rigid analogues of chalcones, were potent and selective inhibitors of the B isoform of hMAO, we planned the synthesis of a larger panel

of (E)-3-benzylidenechroman-4-ones to further investigate the structure—activity relationships. The MAO recognition of the most active compounds, **1h** and **1l**, was investigated by docking experiments performed using available Protein Data Bank (PDB) structures as receptor models.

■ CHEMISTRY

The (E)-3-benzylidenechroman-4-ones 1a-w were synthesized by acid-catalyzed condensation of the appropriate chroman-4-one with substituted benzaldehydes (Scheme 1).

In general, the target compounds were prepared in good yield using 85% phosphoric acid as the acid catalyst and heating the mixture at 80 °C for 6 h. Under these conditions, the reaction of chroman-4-one with N-(4-formylphenyl)acetamide gave a mixture of amide 1k and the corresponding amine 1j, which were easily separated by column chromatography. Starting from substituted chroman-4-ones or 2-substituted benzaldehydes, treatment with 85% phosphoric acid provided the corresponding chromanones (1d, 1e, 1g, 1i, and 1s-v) in very low yields. To obtain these compounds more efficiently, the condensations were carried out in dry ethyl alcohol saturated with dry hydrogen chloride. A single stereoisomer was obtained with both procedures. The 1 H NMR spectra allow the E configuration of the double bound to be assigned on the basis of the chemical shift of the olefinic proton, ranging from 7.6 to 7.9 ppm.

(*E*)-5,7-Dihydroxy-3-(4-hydroxybenzylidene)chroman-4-one (1h) was synthesized by saponification of the corresponding tribenzoate following published procedures.⁴²

The reduction of (E)-3-(4-hydroxy-3-nitrobenzylidene)chroman-4-one (1q) with tin chloride in ethyl alcohol and concentrated hydrogen chloride provided the corresponding amino derivative (1r).

The (E)-3-benzylidenechroman-4-ones $\mathbf{1a} - \mathbf{g}$ were converted into the corresponding 3-benzyl-4-chromones $\mathbf{2a} - \mathbf{g}$ or 3-benzylchroman-4-ones $\mathbf{3a} - \mathbf{e}$ according to procedures described previously.²⁹

■ BIOCHEMISTRY

The potential effects of the tested compounds on hMAO activity were investigated by measuring their effects on the production of hydrogen peroxide (H₂O₂) from p-tyramine, using the Amplex Red MAO assay kit (Molecular Probes, Inc., Eugene, Oregon, U.S.) and microsomal MAO isoforms prepared from insect cells (BTI-TN-5B1-4) infected with recombinant baculovirus containing cDNA inserts for hMAO-A or hMAO-B (Sigma-Aldrich Química S.A., Alcobendas, Spain). In this study hMAO activity was evaluated using the above method following the general procedure described previously by us.⁴³ The tested drugs (new compounds and reference inhibitors) themselves were unable to react directly with the Amplex Red reagent, which indicates that these compounds do not interfere with the measurements. On the other hand, in our experiments and under our experimental conditions, the control activity of hMAO-A and hMAO-B (using p-tyramine as a common substrate for both isoforms) was 165 \pm 2 pmol of p-tyramine oxidized to p-hydroxyphenylacetaldehyde/min (n = 20).

The results of hMAO-A and hMAO-B inhibition studies are expressed as IC_{50} and reported in Tables 1 and 2 together with the hMAO-B selectivity indexes (SI = IC_{50} MAO-A/ IC_{50} MAO-B).

Scheme 1^a

$$\begin{array}{c} R \\ R \\ R \\ R \\ \end{array}$$

$$\begin{array}{c} R \\ R \\ \end{array}$$

1a R = R¹ = R² = R³ = R⁴ = R⁵ = H 1b R = R¹ = R² = R³ = R⁵ = H, R⁴ = OH 1c R = R¹ = R² = R³ = R⁵ = H, R⁴ = CI 1d R = R² = R³ = R⁵ = H, R¹ = R⁴ = CI 1e R¹ = R² = R³ = R⁵ = H, R = R⁴ = CI 1f R = R¹ = R² = R³ = R⁵ = H, R⁴ = OCH₃ 1g R¹ = R² = R³ = R⁵ = H, R = R⁴ = OCH₃ 1i R = R² = R³ = R⁴ = R⁵ = H, R¹ = CI 1j R = R¹ = R² = R³ = R⁵ = H, R⁴ = NH₂ 1k R = R¹ = R² = R³ = R⁵ = H, R⁴ = NHCOCH₃ 1l R = R¹ = R² = R³ = R⁵ = H, R⁴ = NHCOCH₃ $\begin{array}{l} \text{1m R} = \text{R}^1 = \text{R}^2 = \text{R}^5 = \text{H}, \, \text{R}^3 = \text{OCH}_3, \, \text{R}^4 = \text{OH} \\ \text{1n R} = \text{R}^1 = \text{R}^2 = \text{R}^5 = \text{H}, \, \text{R}^3 = \text{OH}, \, \text{R}^4 = \text{OCH}_3 \\ \text{1o R} = \text{R}^1 = \text{R}^2 = \text{R}^5 = \text{H}, \, \text{R}^3 = \text{R}^4 = \text{OH} \\ \text{1p R} = \text{R}^1 = \text{R}^2 = \text{R}^5 = \text{H}, \, \text{R}^3 = \text{COOH}, \, \text{R}^4 = \text{OH} \\ \text{1q R} = \text{R}^1 = \text{R}^2 = \text{R}^5 = \text{H}, \, \text{R}^3 = \text{NO}_2, \, \text{R}^4 = \text{OH} \\ \text{1s R} = \text{R}^1 = \text{R}^3 = \text{R}^5 = \text{H}, \, \text{R}^2 = \text{OH}, \, \text{R}^4 = \text{OCH}_3 \\ \text{1t R} = \text{R}^1 = \text{R}^3 = \text{R}^5 = \text{H}, \, \text{R}^2 = \text{R}^4 = \text{OCH}_3 \\ \text{1u R} = \text{R}^1 = \text{R}^4 = \text{R}^5 = \text{H}, \, \text{R}^2 = \text{R}^3 = \text{OCH}_3 \\ \text{1v R} = \text{R}^1 = \text{R}^5 = \text{H}, \, \text{R}^2 = \text{R}^3 = \text{R}^4 = \text{OCH}_3 \\ \text{1w R} = \text{R}^1 = \text{R}^2 = \text{H}, \, \text{R}^3 = \text{R}^4 = \text{R}^5 = \text{OCH}_3 \\ \end{array}$

^a Reagents and conditions: (i) 85% H₃PO₄, 80° C, 6 h; (ii) dry EtOH sat. HCl, room temp, 48 h.

In Table 1 are presented the data obtained for the (E)-3benzylidenechroman-4-ones 1a-w. Most of these homoisoflavonoids selectively inhibited the enzymatic activity of hMAO-B in the nanomolar or micromolar range. Derivatives 1h, 1k, 1o, and 1r were also able to inhibit hMAO-A in the micromolar range. Among these analogues, 1k and 1r were essentially nonselective (SI = 2 and 3, respectively) whereas 1h and 1o exhibited hMAOB selectivity to different extents (SI = 1331 and 34, respectively). Only the acid 1p was found to be completely inactive toward both isoforms up to the highest concentration tested (100 μM). (E)-3-[4-(Dimethylamino)benzylidene]chroman-4one (11) was the most potent and selective hMAO-B inhibitor identified in this study, exhibiting an IC50 of 8.51 nM and SI of >11751. In comparison with the reference inhibitor selegiline, compound 11 showed higher hMAO-B affinity and selectivity. The replacement of the dimethylamino group with the amino group resulted in compound 1j, about 100-fold less potent and selective as a hMAO-B inhibitor (IC₅₀ = 879.18 nM, SI > 114) than 11. Interestingly, acetylation of the amino group (compound 1k) produced a further decrease of hMAO-B inhibitory activity and a dramatic reduction of selectivity (IC₅₀ = 1.06 μ M, SI = 2). (E)-5,7-Dihydroxy-3-(4-hydroxybenzylidene)chroman-4-one (1h) showed an affinity for hMAO-B ($IC_{50} = 8.61 \text{ nM}$) comparable to that for 1l. Although 1h was able to inhibit both hMAO isoforms, it still exhibited an excellent hMAO-B selectivity (SI = 1331). Although less active than 1h and 1l, (E)-5,7dichloro-3-(4-chlorobenzylidene)chroman-4-one (1e) showed hMAO-B affinity and selectivity (IC₅₀ = 13.03 nM, SI > 7675) better than those of the reference compound, selegiline.

The results for inhibitory effects and selectivity of the 3-benzyl-4H-chromen-4-ones 2a-g and 3-benzylchroman-4-ones 3a-e on hMAO isoforms are reported in Table 2.

Several of these compounds (2e, 2f, 3a, and 3c-e) showed selective hMAO-B inhibitory activities, whereas the 3-benzyl-4chromones 2a-d and 2g and the 3-benzylchroman-4-one 3b were essentially inactive toward both isoforms up to the highest concentration tested (100 μ M). Disappointly, the endocyclic migration of the double bond produces a loss of the inhibitory activity (compounds 2a-d, 2g) or a marked reduction in potency (compounds 2e and 2f) with respect to the corresponding (E)-3-benzylidenechroman-4-ones 1a-g. The reduction of the double bond results in compounds 3a-e, generally more potent and selective than the corresponding 3-benzyl-4H-chromen-4-ones 2a-g. The only exception was 5,7-dichloro-3-(4chlorobenzyl)-4H-chromen-4-one (2e) that exhibited submicromolar potency toward hMAO-B, whereas the corresponding 3-benzylchroman-4-one (3e) was about 10-fold less active and less selective. However, all the 3-benzylchroman-4-ones 3a-e were less potent hMAO-B inhibitors than the corresponding (*E*)-3-benzylidenechroman-4-ones 1a-e.

Table 3 shows the results of the reversibility and irreversibility tests for the most effective compounds 1h and 1l. hMAO-A and hMAO-B inhibition was irreversible in the presence of 1h as shown by the lack of enzyme activity restoration after repeated washing. Similar results were obtained for compound 1l and selegiline against hMAO-B. On the contrary, significant recovery of hMAO-A activity was observed after repeated washing of moclobemide, indicating that this drug is a reversible inhibitor of hMAO-A.

■ DOCKING STUDIES

Docking studies were carried out to evaluate the binding modes of this class of homoisoflavonoids with respect to both

Table 1. IC₅₀ and SI for the Inhibitory Effects of (E)-3-Benzylidenechroman-4-ones 1a—w and Reference Inhibitors on the Enzymatic Activity of Human Recombinant MAO Isoforms Expressed in Baculovirus Infected BTI Insect Cells^a

							IC ₅₀		
compd	R	\mathbb{R}^1	\mathbb{R}^2	\mathbb{R}^3	\mathbb{R}^4	R^5	MAO-A	MAO-B	SI^d
1a	Н	Н	Н	Н	Н	Н	f	$479.70 \pm 19.37 \text{ nM}$	>208 ^h
1b	Н	Н	Н	Н	ОН	Н	f	$55.37\pm2.18~\text{nM}$	>1806 ^h
1c	Н	Н	Н	Н	Cl	Н	f	$154.23 \pm 6.93 \text{ nM}$	>648 ^h
1d	Н	Cl	Н	Н	Cl	Н	f	$31.82\pm1.19~\text{nM}$	>3143 ^h
1e	Cl	Н	Н	Н	Cl	Н	f	$13.03\pm0.79~\text{nM}$	>7675 ^h
1f	Н	Н	Н	Н	OCH ₃	Н	f	$58.90\pm2.09~\text{nM}$	>1698 ^h
1g	OCH_3	Н	Н	Н	OCH ₃	Н	f	$1.57\pm0.03\mu\mathrm{M}$	>64 ^h
1h	OH	Н	Н	Н	ОН	Н	$11.46 \pm 0.43 \mu\text{M}^b$	$8.61\pm0.12~\text{nM}$	1331
1i	Н	Cl	Н	Н	Н	Н	f	$331.11 \pm 9.63 \text{ nM}$	>302 ^h
1j	Н	Н	Н	Н	NH_2	Н	f	$879.18 \pm 45.31 \text{ nM}$	>114 ^h
1k	Н	Н	Н	Н	NHCOCH ₃	Н	$2.10\pm0.11\mu\mathrm{M}^{c}$	$1.06\pm0.04\mu\mathrm{M}$	2.0
11	Н	Н	Н	Н	$N(CH_3)_2$	Н	f	$8.51\pm0.32~\text{nM}$	$> 11751^h$
1m	Н	Н	Н	OCH_3	ОН	Н	g	$104.34\pm2.61~\text{nM}$	>958 ^h
1n	Н	Н	Н	ОН	OCH ₃	Н	f	$476.51 \pm 21.60 \text{ nM}$	>210 ^h
10	Н	Н	Н	ОН	ОН	Н	$4.74 \pm 0.21 \mu\text{M}^b$	$140.52 \pm 4.71 \text{ nM}$	34
1p	Н	Н	Н	СООН	ОН	Н	f	f	
1q	Н	Н	Н	NO_2	ОН	Н	f	$490.84 \pm 12.73 \; nM$	>204 ^h
1r	Н	Н	Н	NH_2	ОН	Н	$1.44 \pm 0.51 \mu\text{M}^b$	$483.88 \pm 16.21 \text{ nM}$	3.0
1s	Н	Н	OH	Н	OCH ₃	Н	g	$1.33 \pm 0.07 \mu\mathrm{M}$	>75 ^h
1t	Н	Н	OCH_3	Н	OCH ₃	Н	f	$247.51 \pm 6.31 \text{ nM}$	>404 ^h
1u	Н	Н	OCH_3	OCH_3	Н	Н	f	$1.92\pm0.08\mu\mathrm{M}$	>52 ^h
1v	Н	Н	OCH_3	OCH_3	OCH ₃	Н	f	$1.31\pm0.04\mu\mathrm{M}$	>76 ^h
1w	Н	Н	Н	OCH_3	OCH ₃	OCH_3	f	$20.69\pm0.37~\mu\mathrm{M}$	>4.8 ^h
clorgyline							$4.46 \pm 0.32 \text{ nM}^b$	$61.35\pm1.13~\mu\mathrm{M}$	0.000073
selegiline							$67.25 \pm 1.02 \mu\text{M}^b$	$19.60\pm0.86~\text{nM}$	3431
iproniazid							$6.56\pm0.76\mu\mathrm{M}$	$7.54\pm0.36\mu\mathrm{M}$	0.87
moclobemide							$361\pm19.37~\mu\mathrm{M}$	e	< 0.36

 $[^]a$ All IC $_{50}$ values shown in this table are the mean \pm SEM from five experiments. b Level of statistical significance: P < 0.01 versus the corresponding IC $_{50}$ values obtained against MAO-B, as determined by ANOVA/Dunnett's. c Level of statistical significance: P < 0.05 versus the corresponding IC $_{50}$ values obtained against MAO-B, as determined by ANOVA/Dunnett's. d SI: hMAO-B selectivity index = IC $_{50}$ (hMAO-A)/IC $_{50}$ (hMAO-B). c Inactive at 1 mM (highest concentration tested). f Inactive at 100 μ M (highest concentration tested). g 100 μ M inhibits the corresponding MAO activity by approximately 40–50%. At higher concentration the compounds precipitate. h Values obtained under the assumption that the corresponding IC $_{50}$ against MAO-B is the highest concentration tested (100 μ M).

isoforms of human MAO. Crystallographic structures were selected from the PDB to build our theoretical receptors models (Experimental Section). Taking into account the experimental hMAO inhibition data, compounds 1h and 1l were chosen for our molecular modeling investigation. Although the irreversible inhibitors usually form a covalent bond with a residue on the enzyme, they may also act by other mechanisms. 44 The so-called tight-binding inhibitors, owing to the very low dissociation constant, may show kinetics similar to the kinetics of covalent irreversible inhibitors. In some cases, the inhibitors may rapidly bind to the enzyme in a low-affinity enzyme-inhibitor (EI) complex and then undergoes a slower rearrangement to a very tightly bound EI^* complex. For instance, R-(-)-deprenyl, an irreversible MAO inhibitor, forms a noncovalent complex with MAO as an initial and reversible step. Afterward, the interaction of R-(-)-deprenyl with MAO leads to a reduction of the enzyme-bound FAD and concomitant oxidation of the inhibitor. This oxidized inhibitor is able to form a covalent bond at the N-5 position of FAD. 45 The initial noncovalent binding to MAO has

been also described for other MAO inhibitors (e.g., clorgyline derivatives). ⁴⁶ Moreover, we have already docked noncovalent ligands ^{40,41,47} showing an irreversible profile similar to **1h** and **1l**.

Both homoisoflavonoids were submitted to a conformational search revealing, not surprisingly, only two conformers within 10 kcal/mol from the global minimum energy. Using the AutoDock Vina method, 48 the most populated structures of 1h and 1l were submitted to flexible docking simulation with respect to both hMAO-A and hMAO-B receptor models (Experimental Section). The theoretical complexes were evaluated taking into account two interaction energy descriptors. Best interaction energy (IE) is the AutoDock Vina lowest ligand-target interaction energy computed with the entire complex ensemble generated by the docking program. In both cases, good qualitative agreements with experimental inhibition data have been obtained (Table 4).

The most stable binding modes of both homoisoflavonoids in the hMAO-A and -B active sites were graphically inspected (Figure 2).

Table 2. IC₅₀ and SI for the Inhibitory Effects of 3-Benzyl-4*H*-chromen-4-ones 2a-g and 3-Benzylchroman-4-ones 3a-e and Reference Inhibitors on the Enzymatic Activity of Human Recombinant MAO Isoforms Expressed in Baculovirus Infected BTI Insect Cells^a

							IC_{50}		
compd	R	\mathbb{R}^1	\mathbb{R}^2	\mathbb{R}^3	R^4	\mathbb{R}^5	MAO-A	MAO-B	SI^c
2a	Н	Н	Н	Н	Н	Н	e	e	
2b	Н	Н	Н	Н	OH	Н	e	f	
2c	Н	Н	Н	Н	Cl	Н	e	e	
2d	Н	Cl	Н	Н	Cl	Н	e	f	
2e	Cl	Н	Н	Н	Cl	Н	f	$438.60 \pm 27.48 \; \text{nM}$	>228 ^g
2f	Н	Н	Н	Н	OCH_3	Н	e	$21.59\pm1.06\mu\mathrm{M}$	>4.6 ^g
2g	OCH_3	Н	Н	Н	OCH_3	Н	e	f	
3a	Н	Н	Н	Н	Н	Н	e	$58.53 \pm 2.24 \mu{ m M}$	>1.7g
3b	Н	Н	Н	Н	OH	Н	e	f	
3c	Н	Н	Н	Н	Cl	Н	e	$7.37 \pm 0.31 \mu\mathrm{M}$	>14 ^g
3d	Н	Cl	Н	Н	Cl	Н	e	$3.07 \pm 0.11 \mu\mathrm{M}$	>33 ^g
3e	Cl	Н	Н	Н	Cl	Н	e	$4.59\pm0.19\mu\mathrm{M}$	>22 ^g
clorgyline							$4.46\pm0.32~\mathrm{nM}^b$	$61.35 \pm 1.13 \mu{ m M}$	0.000073
selegiline							$67.25 \pm 1.02 \mu\text{M}^b$	$19.60\pm0.86~\text{nM}$	3431
iproniazid							$6.56\pm0.76\mu\mathrm{M}$	$7.54\pm0.36\mu\mathrm{M}$	0.87
moclobemide						,	$361 \pm 19.37 \mu\mathrm{M}$	d	< 0.36

 $[^]a$ All IC $_{50}$ values shown in this table are the mean \pm SEM from five experiments. b Level of statistical significance: P < 0.01 versus the corresponding IC $_{50}$ values obtained against MAO-B, as determined by ANOVA/Dunnett's. c SI: hMAO-B selectivity index = IC $_{50}$ (hMAO-A)/IC $_{50}$ (hMAO-B). d Inactive at 1 mM (highest concentration tested). c Inactive at 100 μ M (highest concentration tested). f 100 μ M inhibits the corresponding MAO activity by approximately 40–50%. At higher concentration the compounds precipitate. g Values obtained under the assumption that the corresponding IC $_{50}$ against MAO-B is the highest concentration tested (100 μ M).

Table 3. Reversibility and Irreversibility of hMAO-B Inhibition of Derivatives 1h and 11^a

	% hMAO-A inhibition			
compd	before washing	after repeated washing		
1h (50 μ M) moclobemide (500 μ M)	68.26 ± 4.34 85.98 ± 4.03	61.76 ± 4.98 11.45 ± 0.58^{b}		

	% hMAO-B inhibition			
compd	before washing	after repeated washing		
1h (10 nM)	48.66 ± 6.39	36.22 ± 4.26		
11 (10 nM)	56.34 ± 5.37	58.67 ± 6.60		
selegiline (20 nM)	53.28 ± 2.59	56.34 ± 2.01		

^a Each value is the mean \pm SEM from five experiments (n = 5). ^b Level of statistical significance: P < 0.01 versus the corresponding % MAO-A or MAO-B inhibition before washing, as determined by ANOVA/Dunnett's.

Docking simulations generated several low energy binding modes of 1h and 1l within both hMAO isoforms. These poses revealed the ability of these ligands to alternatively recognize the FAD with both benzyl and chromone rings (see Supporting Information). Interestingly, the hMAO-A best poses of the 1h and 1l chromone rings were located toward the enzymatic cofactor (Figure 2a). Most of the ligand—enzyme interactions appeared to be identical. The main difference between 1h and 1l hMAO-A recognition was due to the presence of a phenol OH in the former compound that established a hydrogen bond with FAD N5 atom. Moreover, 1h, attracted by the FAD, resulted in

Table 4. Comparison between Experimental IC_{50} Data and Theoretical Binding Affinities of 1h and 1l with Respect to hMAO-A and -B

	hM	MAO-A	hM	hMAO-B		
parameter	1h	11	1h	11		
IC ₅₀ a	11460.00	>100000.00	8.61	8.51		
$\Delta G_{ m bind}^{\;\;b}$	-6.30	-1.88	-9.10	-9.35		
best IE c	-7.00	-3.00	-9.50	-10.40		

^a Experimental inhibition data (nM). ^b Free energy of binding (kcal/mol). ^c Best interaction energy (kcal/mol).

exclusive contact with Tyr69; conversely 1l was involved in attractive hydrophobic contacts with Asn181 and Tyr444. The effects of these different interactions on binding energy are highlighted in Table 4.

The best poses of these compounds in the hMAO-B binding site were different from the A isoform. These configurations displayed the chromone ring far from the FAD cofactor. In both 1h and 11 hMAO-B complexes the interaction pattern was similar: one hydrogen bond between the carbonyl oxygen of the ligands and Tyr326-OH and several almost identical hydrophobic contacts. The slightly better interaction of 1l can be attributed to its more hydrophobic nature compared with 1h. Actually the chromone ring was surrounded by a large lipophilic cage (Phe103, Pro104, Trp119, Leu164, Leu167, Phe168, Leu171, Ile199, and Ile316) where 1h, because of its hydrophilic substituents, was disfavored. Finally, the hMAO-B selectivity of both compounds can be attributed to (a) the larger set of residues interacting with hMAO-B and (b) the different hydrogen bond

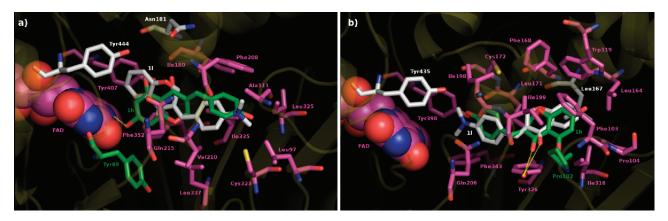


Figure 2. Most representative binding modes of 1h (green) and 1l (white) into (a) hMAO-A and (b) hMAO-B catalytic sites. Common 1h and 1l interacting residues are reported in magenta carbon atoms sticks. Exclusive 1h or 1l interacting residues are displayed according to the corresponding ligands. The FAD cofactor is depicted using the spacefill representation. The rest of the enzyme is shown in yellow, transparent cartoon. Yellow dotted lines indicate hydrogen bonds.

contributions, stronger in hMAO-B (Tyr326-OH···O=1h/1l) than in hMAO-A (FAD-N5···HO-1h).

■ CONCLUSION

In the current study we have synthesized and evaluated a series of (*E*)-3-benzylidenechroman-4-ones 1a-w, 3-benzyl-4*H*-chromen-4-ones 2a-g, and 3-benzylchroman-4-ones 3a-e as inhibitors of h-MAO isoforms A and B. In general, the active compounds showed potent and selective activity toward the B isoform. In particular, several (*E*)-3-benzylidenechroman-4-ones exhibited potencies in the nanomolar range. Two derivatives, 1h and 1l, were the most potent and selective hMAO-B inhibitors with higher potency than the reference inhibitor selegiline and, in the latter case, even higher selectivity. Molecular docking studies suggest that stronger enzyme—inhibitor hydrogen bonds and hydrophobic contacts in the hMAO-B active site can explain the selectivity of both inhibitors for this isoform.

■ EXPERIMENTAL SECTION

Chemistry. Chemicals were purchased from Sigma-Aldrich and used without further purification. Melting points were determined on a Stenford Research Systems OptiMelt (MPA-100) apparatus and are uncorrected. ¹H NMR and ¹³C NMR spectra were recorded in DMSO d_{6} , unless indicated otherwise, on a Bruker AM-400 spectrometer, using TMS as internal standard. IR spectra were recorded in KBr disks on an FT-IR PerkinElmer Spectrum 1000. All compounds were routinely checked by thin-layer chromatography (TLC), ¹H NMR, and ¹³C NMR. TLC was performed on silica gel or aluminum oxide fluorescent coated plates (Merck, Kieselgel, or aluminum oxide 60 F254). Components were visualized by UV light. Compound purity was determined by elemental analysis and was confirmed to be \geq 95% for all compounds. (E)-3-Benzylidenechroman-4-ones (1a-g), 3-benzyl-4-chromones (2a-g), and 3-benzylchroman-4-ones (3a-e) were prepared as previously described by us.²⁹ (E)-5,7-Dihydroxy-3-(4-hydroxybenzylidene)chroman-4-one (1h) was obtained by saponification of the corresponding tribenzoate as previously reported. 42

General Procedure for the Synthesis of (*E*)-3-Benzylidenechroman-4-ones 1l—q and 1w. A mixture of the appropriate chroman-4-one (0.01 mol) and substituted benzaldehyde (0.01 mol) in 85% phosphoric acid (20 mL) was heated at 80 °C for 6 h. After cooling, the mixture was diluted with ice—water (and made alkaline for the preparation of compound 11). The solid was filtered off, washed with water, and crystallized.

(*E*)-3-(4-(Dimethylamino)benzylidene)chroman-4-one (1l). Yield: 77%. Mp = 151-153 °C (lit. 150-153 °C)³⁴ from EtOH. The compound exhibited spectroscopic data identical to those previously reported.³⁴

(*F*)-3-(4-Hydroxy-3-methoxybenzylidene)chroman-4-one (1m). Yield: 71%. Mp = 130–131 °C (lit. 126–129 °C)⁴⁹ from ethyl alcohol. IR (KBr): 3200, 1655 cm⁻¹. ¹H NMR: δ (ppm) 9.75 (bs, 1H, OH), 7.88 (dd, 1H, H5, J_{5-6} = 8.0 Hz, J_{5-7} = 1.6 Hz), 7.71 (t, 1H, =CH, J_{all} = 1.8 Hz), 7.58 (ddd, 1H, H7, J_{6-7} = 7.2 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.6 Hz), 7.12 (ddd, 1H, H6, J_{6-7} = 7.2 Hz, J_{5-6} = 8.0 Hz, J_{6-8} = 1.0 Hz), 7.07 (d, 1H, H2', $J_{2'-6'}$ = 1.7 Hz), 7.05 (dd, 1H, H8, J_{7-8} = 8.4 Hz, J_{6-8} = 1.0 Hz), 6.93 (dd, 1H, H6', $J_{2'-6'}$ = 1.7 Hz), 6.90 (d, 1H, H5', $J_{5'-6'}$ = 8.4 Hz), 5.47 (d, 2H, H2, J_{all} = 1.7 Hz), 3.84 (s, 3H, OCH₃). ¹³C NMR: δ (ppm) 180.9, 160.4, 148.8, 147.6, 137.2, 135.8, 127.7, 127.1, 125.2, 124.4, 121.8, 121.6, 117.7, 115.6, 114.8, 67.5, 55.6.

(*E*)-3-(3-Hydroxy-4-methoxybenzylidene)chroman-4-one (1n). Yield: 78%. Mp = 192–193 °C from EtOH. IR (KBr): 3150, 1650 cm⁻¹. ¹H NMR: δ (ppm) 9.34 (bs, 1H, OH), 7.88 (dd, 1H, H5, J_{5-6} = 7.9 Hz, J_{5-7} = 1.8 Hz), 7.64 (t, 1H, =CH, J_{all} = 1.8 Hz), 7.58 (ddd, 1H, H7, J_{6-7} = 7.2 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.8 Hz), 7.13 (ddd, 1H, H6, J_{6-7} = 7.2 Hz, J_{5-6} = 7.9 Hz, J_{6-8} = 1.0 Hz), 7.07–7.03 (m, 2H, H8, H5'), 6.94–6.91 (m, 2H, H2', H6'), 5.43 (d, 2H, H2, J_{all} = 1.8 Hz), 3.85 (s, 3H, OCH₃). ¹³C NMR: δ (ppm) 180.9, 160.4, 149.4, 146.5, 136.8, 135.9, 128.3, 127.1, 126.5, 122.9, 121.8, 121.5, 117.7, 117.2, 112.0, 67.4, 55.6.

(*E*)-3-(3,4-Dihydroxybenzylidene)chroman-4-one (10). Yield: 54%. Mp = 227–230 °C (lit. 224–225 °C)⁴⁹ from EtOH. IR (KBr): 3400, 1640 cm⁻¹. ¹H NMR: δ (ppm) 9.50 (bs, 2H, 2OH), 7.87 (dd, 1H, H5, $J_{5-6} = 7.8$ Hz, $J_{5-7} = 1.7$ Hz), 7.61 (t, 1H, =CH, $J_{all} = 1.8$ Hz), 7.58 (ddd, 1H, H7, $J_{6-7} = 7.4$ Hz, $J_{7-8} = 8.4$ Hz, $J_{5-7} = 1.7$ Hz), 7.12 (ddd, 1H, H6, $J_{6-7} = 7.4$ Hz, $J_{5-6} = 7.8$ Hz, $J_{6-8} = 1.0$ Hz), 7.05 (dd, 1H, H8, $J_{7-8} = 8.4$ Hz, $J_{6-8} = 1.0$ Hz), 6.89–6.80 (m, 3H, H2', H5', H6'), 5.43 (d, 2H, H2, $J_{all} = 1.8$ Hz). ¹³C NMR: δ (ppm) 180.9, 160.3, 147.8, 145.3, 137.2, 135.8, 127.3, 127.1, 125.2, 123.4, 121.7, 121.6, 117.7, 115.8, 67.5.

(*E*)-2-Hydroxy-5-[(4-oxochroman-3-ylidene)methyl]benzoic Acid (1p). Yield: 74%. Mp = 234–238 °C from EtOH. IR (KBr): 3500–2300, 1650, 1600 cm⁻¹. ¹H NMR: δ (ppm) 11.87 (bs, 1H, COOH), 7.87–7.80 (m, 2H, H5, H2'), 7.69 (bs, 1H, =CH), 7.63–7.54 (m, 2H, H6', H7), 7.12–7.01 (m, 3H, H5', H6, H8), 5.40 (d, 2H, H2, $J_{\rm all}$ = 1.4 Hz). ¹³C NMR: δ (ppm) 180.9, 171.2, 162.0, 160.5, 137.4, 136.0, 135.6, 132.7, 129.4, 127.1, 125.0, 121.8, 121.4, 117.8, 117.7, 113.5, 67.2.

(*E*)-3-(4-Hydroxy-3-nitrobenzylidene)chroman-4-one (1q). Yield: 77%. Mp = 203–205 °C from EtOAc. IR (KBr): 3240, 1655, 1530, 1300 cm⁻¹. ¹H NMR: δ (ppm) 11.65 (bs, 1H, OH), 7.94 (d, 1H, H2', $J_{2'-6'}$ = 2.2 Hz), 7.88 (dd, 1H, H5, J_{5-6} = 7.9 Hz, J_{5-7} = 1.8 Hz), 7.70 (bs, 1H, =CH), 7.64 (dd, 1H, H6', $J_{2'-6'}$ = 2.2 Hz $J_{5'-6'}$ = 8.7 Hz), 7.60 (ddd, 1H, H7, J_{6-7} = 7.3 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.8 Hz), 7.23 (d, 1H, H5', $J_{5'-6'}$ = 8.7 Hz), 7.13 (ddd, 1H, H6, J_{6-7} = 7.3 Hz, J_{5-6} = 7.9 Hz, J_{6-8} = 0.9 Hz), 7.05 (dd, 1H, H8, J_{7-8} = 8.4 Hz, J_{6-8} = 0.9 Hz), 5.44 (d, 2H, H2, J_{all} = 1.8 Hz). ¹³C NMR: δ (ppm) 180.5, 160.5, 152.8, 137.2, 136.5, 135.1, 134.4, 130.2, 127.2, 127.1, 124.9, 121.9, 121.3, 119.3, 117.8, 67.2.

(*E*)-3-(3,4,5-Trimethoxybenzylidene)chroman-4-one (1w). Yield: 63%. Mp = 105-106 °C (lit. 108-109 °C)⁵⁰ from EtOAc/petroleum ether. IR (KBr): 1660 cm⁻¹. ¹H NMR: δ (ppm) 7.90 (dd, 1H, HS, $J_{5-6} = 7.8$ Hz, $J_{5-7} = 1.8$ Hz), 7.73 (bs, 1H, =CH), 7.60 (ddd, 1H, H7, $J_{7-8} = 8.4$ Hz, $J_{6-7} = 7.2$ Hz, $J_{5-7} = 1.8$ Hz), 7.14 (ddd,1H, H6, $J_{6-7} = 7.2$ Hz, $J_{5-6} = 7.8$ Hz, $J_{6-8} = 1.0$ Hz), 7.06 (dd, 1H, H8, $J_{7-8} = 8.4$ Hz, $J_{6-8} = 1.0$ Hz), 6.77 (s, 2H, H2′, H6′), 5.50 (d, 2H, H2 $J_{all} = 1.8$ Hz), 3.84 (s, 6H, 2OCH₃), 3.74 (s, 3H, OCH₃). ¹³C NMR: δ (ppm) 181.0, 160.6, 152.8, 138.9, 136.9, 136.2, 130.0, 129.2, 127.2, 121.9, 121.5, 117.8, 108.0, 67.4, 60.1, 56.0.

Synthesis of (*E*)-3-(4-Aminobenzylidene)chroman-4-one (1j) and (*E*)-*N*-(4-[(4-Oxochroman-3-ylidene)methyl]phenyl) acetamide (1k). A mixture of chroman-4-one (0.01 mol) and 4-aminobenzaldehyde (0.01 mol) in 85% $\rm H_3PO_4$ (20 mL) was heated at 80 °C for 6 h. After cooling, the mixture was diluted with ice—water and alkalinized with 2 N NaOH. The solid was filtered off and washed with water. The obtained mixture of (*E*)-3-(4-aminobenzylidene)chroman-4-one (1j) and (*E*)-N-(4-[(4-oxochroman-3-ylidene)methyl]phenyl)-acetamide (1k) was separated by column chromatography on silica gel, eluting with AcOEt/petroleum ether (1:1).

(*E*)-3-(4-Aminobenzylidene)chroman-4-one (1j). Yield: 30%. Mp = 164–66 °C from EtOAc/petroleum ether. IR (KBr): 3430, 3340, 1650 cm⁻¹. ¹H NMR: δ (ppm) 7.85 (dd, 1H, H5, J_{5-6} = 7.8 Hz, J_{5-7} = 1.8 Hz), 7.62 (bs, 1H, =CH), 7.55 (ddd, 1H, H7, J_{6-7} = 7.2 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.8 Hz), 7.21 (d, 2H, H2', H6', $J_{2'-3'}$ = 8.6 Hz), 7.10 (ddd, 1H, H6, J_{6-7} = 7.2 Hz, J_{5-6} = 8.2 Hz, J_{6-8} = 1.0 Hz), 7.03 (dd, 1H, H8, J_{7-8} = 8.4 Hz, J_{6-8} = 1.0 Hz), 6.66 (d, 2H, H3', H5', $J_{2'-3'}$ = 8.6 Hz), 5.95 (bs, 2H, NH₂), 5.43 (d, 2H, H2, J_{all} = 1.8 Hz). ¹³C NMR: δ (ppm) 180.6, 160.2, 151.2, 137.8, 135.4, 133.0, 127.0, 124.6, 121.8, 121.6, 121.0, 117.6, 113.5, 67.8.

(*E*)-*N*-(4-[(4-Oxochroman-3-ylidene)methyl]phenyl)acetamide (1k). Yield: 25%. Mp = 175–176 °C from EtOAc/petroleum ether. IR (KBr): 3300, 1664, 1660 cm $^{-1}$. ¹H NMR: δ (ppm) 10.23 (s, 1H, NH), 7.88 (dd, 1H, H5, J_{5-6} = 7.8 Hz, J_{5-7} = 1.6 Hz), 7.74–7.70 (m, 3H, H2′, H6′, =CH), 7.59 (ddd, 1H, H7, J_{6-7} = 7.1 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.6 Hz), 7.44 (d, 2H, H3′, H5′, $J_{2'.3'}$ = 8.6 Hz), 7.13 (ddd, 1H, H6, J_{6-7} = 7.1 Hz, J_{5-6} = 7.8 Hz, J_{6-8} = 0.7 Hz), 7.06 (dd, 1H, H8, J_{7-8} = 8.4 Hz, J_{6-8} = 0.7 Hz), 5.45 (d, 2H, H2, J_{all} = 1.6 Hz), 2.09 (s, 3H, CH₃). ¹³C NMR: δ (ppm) 181.0, 168.7, 160.5, 140.8, 136.3, 136.0, 131.5, 129.1, 128.3, 127.2, 121.9, 121.5, 118.7, 117.8, 67.5, 24.1.

General Procedure for the Synthesis of (*E*)-3-Benzylidenechroman-4-ones 1i and 1s–v. A mixture of the appropriate chroman-4-one (0.01 mol) with substituted benzaldehyde (0.01 mol) in dry EtOH saturated with HCl (40 mL) was stirred at room temperature for 48 h. After this time, the mixture was diluted with ice—water and the solid filtered off and washed with water. The product was purified by column chromatography on silica gel or by crystallization.

(*E*)-3-Benzylidene-6-chlorochroman-4-one (1i). Yield: 86%. Mp = 158–159 °C (lit. 148–150 °C)⁵¹ from EtOAc. IR (KBr): 1660 cm⁻¹. ¹H NMR (CDCl₃): δ (ppm) 7.98 (d, 1H, H5, J_{5-7} = 2.6 Hz), 7.90 (bs, 1H, =CH), 7.49–7.37 (m, 4H, H7, H3'-H5'), 7.33–7.30 (m, 2H, H2', H6'), 6.93 (d, 1H, H8, J_{7-8} = 8.8 Hz), 5.35 (d, 2H, H2, J_{all} = 1.8 Hz). ¹³C NMR (DMSO d₆): δ (ppm) 180.2, 159.2, 137.3, 135.7, 133.5, 130.3, 129.8, 128.8, 126.0, 125.9, 124.8, 122.4, 120.3, 67.5.

(*E*)-3-(2-Hydroxy-4-methoxybenzylidene)chroman-4-one (1s). Purified by column chromatography, eluting with AcOEt/petroleum ether (1:5). Yield: 56%. Mp = 158–160 °C from EtOH. IR (KBr): 3124, 1655 cm⁻¹. ¹H NMR: δ (ppm) 10.34 (bs, 1H, OH), 7.89–7.86 (m, 2H, =CH, H5), 7.57 (ddd, 1H, H7, J_{6-7} = 7.2 Hz, J_{7-8} = 8.4 Hz, J_{5-7} = 1.8 Hz), 7.12 (ddd, 1H, H6, J_{6-7} = 7.2 Hz, J_{5-6} = 7.9 Hz, J_{6-8} = 1.0 Hz), 7.08 (d, 1H, H6', $J_{5'-6'}$ = 8.2 Hz), 7.04 (dd, 1H, H8, J_{7-8} = 8.4 Hz, J_{6-8} = 1.0 Hz), 6.53–6.49 (m, 2H, H3', H5'), 5.33 (d, 2H, H2, J_{all} = 1.8 Hz), 3.77 (s, 3H, OCH₃). ¹³C NMR: δ (ppm) 181.2, 162.2, 160.5, 158.6, 135.8, 132.8, 131.7, 127.3, 127.1, 121.8, 121.7, 117.8, 114.0, 105.4, 101.1, 67.9, 55.2.

(*E*)-3-(2,4-Dimethoxybenzylidene)chroman-4-one (1t). Yield: 64%. Mp = 123-125 °C (lit. 133-135 °C)³⁴ from MeOH. The compound exhibited spectroscopic data identical to those previously reported.³⁴

(*F*)-3-(2,3-Dimethoxybenzylidene)chroman-4-one (1u). Yield: 53%. Mp = 108-109 °C EtOAc/petroleum ether. IR (KBr): 1673 cm⁻¹. ¹H NMR: δ (ppm) 7.90 (dd, 1H, H5, $J_{5-6} = 7.8$ Hz, $J_{5-7} = 1.7$ Hz), 7.82 (bs, 1H, =CH), 7.61 (ddd, 1H, H7, $J_{7-8} = 8.9$ Hz, $J_{6-7} = 7.2$ Hz, $J_{5-7} = 1.7$ Hz), 7.22-7.12 (m, 3H, H6, H5′, H6′), 7.05 (d, 1H, H8, $J_{7-8} = 8.3$ Hz), 6.83 (dd, 1H, H4′, $J_{4'.5'} = 6.9$ Hz, $J_{4'.6'} = 2.2$ Hz), 5.29 (d, 2H, H2 $J_{\rm all} = 1.8$ Hz), 3.86 (s, 3H, OCH₃), 3.74 (s, 3H, OCH₃). ¹³C NMR: δ (ppm) 181.2, 160.8, 152.5, 147.6, 136.3, 132.2, 131.3, 127.6, 127.3, 124.1, 122.0, 121.8, 121.5, 118.0, 114.8, 67.6, 60.6, 55.8.

(*F*)-3-(2,3,4-Trimethoxybenzylidene)chroman-4-one (1v). Yield: 43%. Mp = 116–118 °C (lit. viscous oil)⁵⁰ from acetone. IR (KBr): 1668 cm⁻¹. ¹H NMR: δ (ppm) 7.89 (dd, 1H, H5, J_{5-6} = 7.8 Hz, J_{5-7} = 1.8 Hz), 7.79 (bs, 1H, =CH), 7.60 (ddd, 1H, H7, J_{7-8} = 8.9 Hz, J_{6-7} = 7.2 Hz, J_{5-7} = 1.8 Hz), 7.13 (ddd, 1H, H6, J_{6-7} = 7.2 Hz, J_{5-6} = 7.8 Hz, J_{6-8} = 1.0 Hz), 7.05 (dd, 1H, H8, J_{7-8} = 8.9 Hz, J_{6-8} = 1.0 Hz), 6.98 (s, 1H, H6', $J_{2'-6'}$ = 8.7 Hz), 6.92 (s, 1H, H5', $J_{2'-6'}$ = 8.7 Hz), 5.33 (d, 2H, H2, J_{all} = 1.8 Hz), 3.88 (s, 3H, OCH3), 3.83 (s, 3H, OCH3), 3.80 (s, 3H, OCH3). ¹³C NMR: δ (ppm) 181.1, 160.6, 155.3, 152.7, 141.7, 136.0, 132.0, 129.5, 127.2, 125.6, 121.9, 121.6, 120.3, 117.8, 107.7, 67.7, 61.3, 60.5, 56.0.

Synthesis of (*E***)-3-(3-Amino-4-hydroxybenzylidene)chroman-4-one (1r).** Tin(II) chloride (0.03 mol) in EtOH (15 mL) was added to a suspension of (*E*)-3-(4-hydroxy-3-nitrobenzylidene)chroman-4-one (1q) (0.005 mol) in EtOH(35 mL) and concentrated HCl (75 mL). The mixture was refluxed for 2 h under stirring. After cooling, the mixture was neutralized with 2 N NaOH and the solid was filtered off. The product was purified by column chromatography on silica gel (AcOEt/petroleum ether 1:1). Yield: 46%. Mp = 192–195 °C from EtOAc/petroleum ether. IR (KBr): 3460, 3380, 1655 cm⁻¹. ¹H NMR: δ (ppm) 9.80 (bs, 1H, OH), 7.86 (d, 1H, H5, J_{5-6} = 7.6 Hz), 7.59–7.55 (m, 2H, =CH, H7), 7.12 (t, 1H, H6, J_{5-6} = J_{6-7} = 7.6 Hz), 7.04 (d, 1H, H8, J_{7-8} = 8.3 Hz), 6.78–6.75 (m, 2H, H2', H6'), 6.61 (d, 1H, H5', $J_{5'-6'}$ = 7.9 Hz), 5.43 (d, 2H, H2, J_{all} = 1.5 Hz), 4.76 (bs, 2H, NH₂). ¹³C NMR: δ (ppm) 181.0, 160.4, 146.5, 137.9, 137.0, 135.7, 127.1, 126.8, 125.3, 121.8, 121.7, 120.9, 117.7, 115.9, 114.3, 67.7.

Pharmacological Studies. Determination of hMAO Isoform Activity. The effects of the test compounds on hMAO isoform enzymatic activity were evaluated by a fluorimetric method following the experimental protocol previously described. ⁵²

Reversibility and Irreversibility Experiments. To evaluate whether some of the tested compounds (1h and 11) are reversible or irreversible hMAO-B inhibitors, an effective centrifugation—ultrafiltration method (so-called repeated washing) was used. 40

Molecular Modeling. Compounds 1h and 1l were built by means of the Maestro GUI.⁵³ Conformational properties of both molecules have been investigated by means of 1000 steps of Monte Carlo (MC) search as implemented in Macromodel, version 7.2.⁵⁴ Each conformer was energy-minimized using the AMBER*⁵⁵ force field in united atoms

notation. Water solvent effects have been taken into account by means of the implicit model GB/SA. The global minimum energy structures of **1h** and **1l** were considered for the next docking simulations. Crystallographic structures, deposited into the Protein Data Bank (PDB) Twith codes 2Z5X and 2V60, were considered, after graphical manipulation (fixing missing atoms and/or bond order, adding nonaliphatic H atoms), as receptor models of hMAO-A and hMAO-B, respectively. The cocrystallized ligands harmine and 7-(3-chlorobenzyloxy)-4-carboxaldehyde-coumarin, respectively, for 2Z5X and 2V60, were removed, FAD double bonds were fixed, and hydrogen atoms were added onto both proteins and cofactors.

According to the AutoDock Vina⁴⁸ docking methodology, a regular box of 15 625 ų, centered on the FAD N5, was considered as the active site in both hMAO-A and -B receptor models. The global minimum energy conformers of 1h and 1l were submitted to flexible docking simulations. The estimation of the $\Delta G_{\rm bind}$ was carried out according to the following equation:

$$\Delta G_{\text{bind}} = \sum_{c=1}^{n} \frac{(\text{IE}_c)(P_c)}{100}$$

where ΔG_{bind} is the Boltzmann averaged binding free energy, c is the configuration of the ligand—target complex, n is the maximum number of configurations, IE $_c$ is the AutoDock Vina interaction energy of the configuration c, and P_c is the Boltzman population at 300 K for configuration c.

ASSOCIATED CONTENT

Supporting Information. Elemental analysis results (C, H, N, Cl) of tested compounds and molecular modeling details. This material is available free of charge via the Internet at http://pubs.acs.org.

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■ ABBREVIATIONS USED

MAO, monoamine oxidase; MAO-A, monoamine oxidase A; MAO-B, monoamine oxidase B; PTK, protein tyrosine kinase; hMAO, human MAO; PDB, Protein Data Bank

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