



# Synthesis and biological activity of fibrate-based acyl- and alkyl-phenoxyacetic methyl esters and 1,2-dihydroquinolines

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## Abstract

A series of highly potent antihyperlipidemic agents constituted by a fibrate-based structure was recently reported by our group, whose synthesis started from isovanillin derivatives. In the interest of evaluating the bioisosteric effect of the vanillin-based isomers on their antihyperlipidemic activity, the present study focuses on the synthesis of 5-acyl-1-phenoxyacetic methyl esters **5a–c** and their saturated side-chain 5-alkyl-1-phenoxyacetates **6a–c**. Their strong *in vivo* effect was associated with the inhibition of HMG-CoA reductase. Since 1,2-dihydroquinolines inhibit this enzyme, a series of such heterocycles (**9a–d**) was prepared by our efficient regioselective, one-step, solvent-free method. Apart from showing hypolipidemic activity *in vivo*, some of the compounds displayed antifungal, antioxidant and cytotoxic activity *in vitro*. The binding mode of four compounds at the active site of HMGRh was examined with docking simulations, observing an interaction with most of the amino acids targeted by simvastatin.

**Keywords** Antihyperlipidemic activity · 2-Acyl-1-hydroxyphenoxyacetic esters · 2-Alkyl-1-hydroxyphenoxyacetic esters · 1,2-Dihydroquinolines · HMG-CoA reductase · Antioxidant activity

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## Introduction

Hyperlipidemia, which represent a significant cardiovascular risk factor, is a growing public health problem worldwide (Thagizadeh et al. 2019). It is mainly associated with diseases such as atherosclerosis (Yuan et al. 2007; Steinberg 2005) and ischemic heart disease (Mehta et al.

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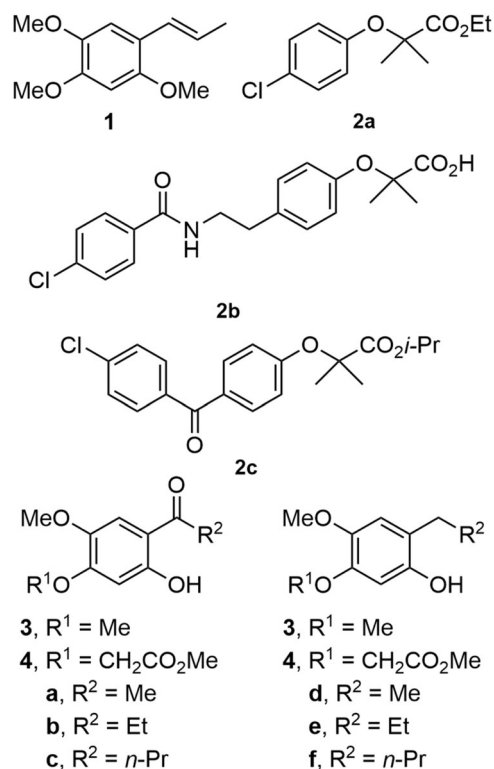
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2014). Obesity and diabetes may contribute substantially to the elevated prevalence of dyslipidemia (Van Gaal et al. 2006; Aguilar-Salinas et al. 2001). When lifestyle changes (e.g., diet, exercise, and weight loss) fail to control dyslipidemia, healthcare systems turn to the prescription of lipid-lowering medication (Gielen et al. 2009; Grundy et al. 2005). Although many synthetic drugs are available to treat hyperlipidemia, they are related with multiple adverse effects, such as myopathy, rhabdomyolysis, and liver toxicity (Chalasanani 2005; Graham et al. 2004; Elisaf et al. 2008; Hodel 2002; Joy and Hegele 2009; Williams and Feely 2002; Davidson et al. 2007).

Statins are among the most commonly used drugs for the clinical management of hypercholesterolemia (Maron et al. 2000; Wierzbicki 2001; Miyazaki et al. 2004; Vivancos et al. 1999). Their adverse effects include rhabdomyolysis, cognitive decline, neuropathy, pancreatic, and hepatic dysfunction, and an increased risk of cancer (Golomb and Evans 2008; Rotta-Bonfin et al. 2015). The mechanism of their activity has been established as the inhibition of the 3-hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) reductase (HMGR), an enzyme involved in the biosynthesis of cholesterol (Singh et al. 2009; Haines et al. 2013).

Among other agents inhibiting HMGR (Menéndez et al. 2001; Liu and Yeh 2002; Parker et al. 1993; Sung et al. 2004; Bradfute and Simoni 1994; Harada-Shiba et al. 1995; Panini et al. 1986) are cholestin (Man et al. 2002), diosgenin (Raju and Bird 2007), lanosterol analogs (Trzaskos et al. 1993),  $\beta$ -sitosterol (Field et al. 1997), and the natural product  $\alpha$ -asarone (**1**) (active metabolite of the Yucatan peninsula (Mexico) tree called Elemuy (*Mosannona depressa* (Baill.) Chatrou (Campos-Ríos and Chiang Cabrera 2006)) resulting in an antihyperlipidemic effect (Chamorro et al. 1993).  $\alpha$ -Asarone (**1**) has also been reported to inhibit HMGR (Singh et al. 2009; Rodríguez-Páez et al. 2003) and exhibit antifungal (Momin and Nair 2002; Lee et al. 2004) and antithrombotic (Poplawsky et al. 2000) activity.

Fibrates such as clofibrate (**2a**) (Oliver 2012; Mohamadzadeh et al. 2013), bezafibrate (**2b**) (BIP study group 2000), and fenofibrate (**2c**) (Uchida et al. 2011) are phenoxyacetic-based compounds with a potent hypotriglyceridemic activity (Lalloyer and Staels 2010; Jover-Fernández and Hernández-Mijares 2012). Unlike statins, their mechanism of action consists of binding to and activating peroxisome proliferator activated receptor  $\alpha$  (PPAR $\alpha$ ), a transcription factor (Schoonjans et al. 1996; Willson et al. 2000; Fazio and Linton 2004; Shaikh and Ali 2018). Fibrates enhance HDL cholesterol but can cause myopathy, cholelithiasis, nausea, syndrome of inappropriate secretion of antidiuretic hormone ADH (SIADH), and liver injury (The Field Study 2005; Okopień et al. 2018). These adverse side effects were among other motives that

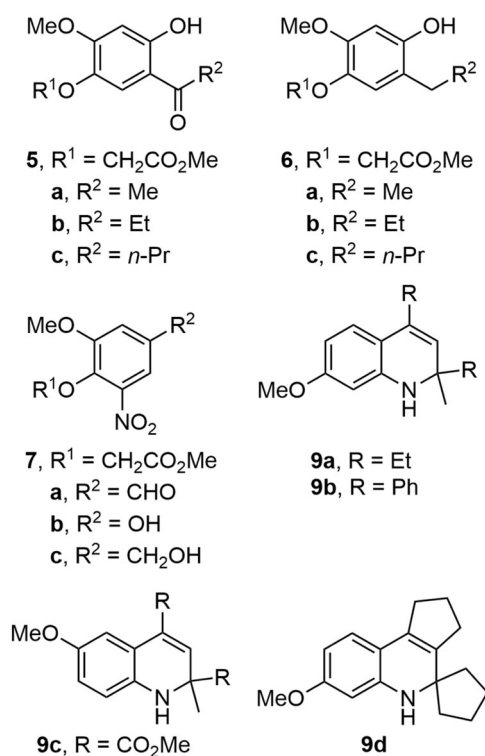


**Fig. 1** Structures of  $\alpha$ -asarone (**1**), clofibrate (**2a**), bezafibrate (**2b**), fenofibrate (**2c**),  $\alpha$ -asarone analogs **3a–f**, and fibrate analogs **4a–f**

clofibrate was withdrawn from the market in some countries. The biological activity of phenoxy acids and their derivatives is exceptionally wide-ranging, being antihyperlipidemic, hypoglycemic, antimicrobial, antiviral, antitubercular, anti-inflammatory, analgesic, antioxidant, anticancer, and anti-hypertensive (Shaheen et al. 2016).

In the search of new synthetic hypolipidemic agents with fewer or no side effects, our group previously synthesized a series of  $\alpha$ -asarone (**1**) analogs that showed the desired activity in potent form (Argüelles et al. 2010; Cruz et al. 2003). Docking studies revealed the binding mode of **1** with the active site of the human HMGR (HMGRh) enzyme (Medina-Franco et al. 2005). Moreover, the oxyacetic group at the C-2 position of **1** (Argüelles et al. 2010), mimicking the fibrate structure, proved to have a significant lipid-lowering activity (Labarrios et al. 1999; Zúñiga et al. 2005). Particularly, the series of  $\alpha$ -asarone- **3a–f** and phenoxyacetic-based derivatives **4a–f** (Fig. 1) (all prepared from isovanillin) displayed a sharp reduction in cholesterol, LDL-cholesterol, and triglycerides, indicating HMGRh inhibition (Mendieta et al. 2014).

On the other hand, there is evidence of diverse types of biological activity of 1,2-dihydroquinolines, found to act as antidiabetic, anti-inflammatory (Hegedüs et al. 2007), antibacterial (Johnson et al. 1989), antioxidant and cytotoxic (Dorey et al. 2000; Błaszczuk and Skolimowski 2007; Błaszczuk et al. 2013; de Koning 2002) (Błaszczuk et al.



**Fig. 2** Structures of the synthesized fibrate-based analogs **5a–c**, **6a–c**, and **7a–c**, and the 1,2-dihydroquinolines **9a–d**

2006; Błaszczuk and Skolimowski 2005; Błaszczuk and Skolimowski 2006) agents. Likewise, analogous 1,2,3,4-tetrahydroquinolines have shown antifungal (Chander et al. 2016a), antimicrobial (Chander et al. 2016b), and antioxidant (Dorey et al. 2000) activity, which has been associated with their capacity of lipid peroxidation and the inhibition of HMGR (Hegedüs et al. 2007). Consequently, these heterocycles may also produce a general hypolipidemic, antioxidant and antifungal effect.

The first aim of the present study was to examine the possible impact on the antihyperlipidemic activity caused by the isomeric position of the substituents in the benzene ring of vanillin-based isomers. Thus, three series of fibrate-based compounds were synthesized starting from vanillin (**8**) and their structure elucidated. The series consisted of 5-acyl-1-phenoxyacetic methyl esters **5a–c**, their saturated side-chain 5-alkyl-1-phenoxyacetates **6a–c** and nitrophenoxyacetates **7a–c** (Fig. 2). Due to the HMGR-related activity of hydroquinolines, a series of the latter was synthesized by using our regioselective one-step method (Gutiérrez et al. 2013). These compounds, **9a–d**, were evaluated *in vivo* as hypolipidemic agents, and *in vitro* for their antifungal, antioxidant, and anticancer effect. The mechanism of action of the test compounds on the HMGRh structure was explored with docking simulations.

## Materials and methods

### Chemistry

Melting points were determined with an Electrothermal capillary melting point apparatus. IR spectra were recorded on an FT-IR Perkin Elmer 2000 spectrophotometer. <sup>1</sup>H (300 or 500 MHz) and <sup>13</sup>C (75.4 or 125 MHz) NMR spectra were recorded on Varian Mercury-300 or Varian VNMR System instruments, with TMS as internal standard. Mass spectra (MS) were taken in the electron impact (EI) mode on Hewlett-Packard 5971A and Thermo-Finnigan Polaris Q spectrometers. High-resolution mass spectra (HRMS) were obtained (EI) on a Jeol JSM-GCMateII spectrometer. Commercial reagents were used as received from Aldrich and anhydrous solvents were obtained by a distillation process. Thin layer chromatography was performed on precoated silica gel plates (Merck 60F<sub>254</sub>). Silica gel (230–400 mesh) was used for column chromatography. All air moisture sensitive reactions were carried out with oven-dried glassware under nitrogen atmosphere. Prior to use, acetone and DMF were distilled from 4Å; molecular sieves and methylene chloride from calcium hydride. K<sub>2</sub>CO<sub>3</sub> was dried overnight at 120 °C before use. All other reagents were used without further purification. The preparation of compounds **12** (Grenier et al. 2000; Kiss et al. 2010; Rashid et al. 2018) and **9a–d** has been previously described (Gutiérrez et al. 2013).

### Methyl 2-(4-formyl-2-methoxyphenoxy)acetate (**10**)

Under an N<sub>2</sub> atmosphere, a mixture of vanillin (**8**) (10.00 g, 65.72 mmol), methyl bromoacetate (16.463 g, 98.58 mmol), and dry K<sub>2</sub>CO<sub>3</sub> (18.165 g, 131.44 mmol) in anhydrous acetone (100 mL) was heated to 60 °C for 4 h. The reaction mixture was filtered and the solvent removed under vacuum. The residue was purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 8:2) to give **10** (13.41 g, 91%) as a white solid. *R*<sub>f</sub> 0.48 (hexane/EtOAc, 6:4); m.p. 92–93 °C. IR (KBr):  $\bar{\nu}$  2994, 1740, 1677, 1591, 1508, 1452, 1404, 1285, 1261, 1224, 1136, 1028, 861, 807, 735 cm<sup>-1</sup>. <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>):  $\delta$  3.81 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.96 (s, 3H, OCH<sub>3</sub>), 4.80 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 6.89 (d, *J* = 8.5 Hz, 1H, H-6'), 7.42 (dd, *J* = 8.5, 1.8 Hz, 1H, H-5'), 7.44 (d, *J* = 1.8 Hz, 1H, H-3'), 9.87 (s, 1H, CHO). <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>):  $\delta$  52.4 (CO<sub>2</sub>CH<sub>3</sub>), 56.0 (OCH<sub>3</sub>), 65.8 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 109.8 (C-3'), 112.3 (C-6'), 126.1 (C-5'), 131.2 (C-4'), 149.9 (C-2'), 152.4 (C-1'), 168.5 (CO<sub>2</sub>CH<sub>3</sub>), 190.7 (CHO). MS (70 eV): *m/z* 224 (M<sup>+</sup>, 100), 165 (25), 150 (30), 137 (35), 119 (36), 105 (20), 95 (30), 77 (31), 59 (32). HRMS (EI): calcd. for C<sub>11</sub>H<sub>12</sub>O<sub>5</sub> [M]<sup>+</sup> 224.0685; found: 224.0685.

### Methyl 2-(4-hydroxy-2-methoxyphenoxy)acetate (11)

A mixture of **10** (10,000 g, 44.64 mmol) and *m*CPBA (77%) (15,000 g, 66.96 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (100 mL) was stirred at room temperature (rt) for 12 h. The solvent was removed under vacuum, and MeOH (100 mL) and HCl (6N) (5 mL) were added. The reaction mixture was stirred at rt for 1 h before removing the solvent under vacuum. A saturated aqueous solution of NaHCO<sub>3</sub> was added until neutral, then extracted with EtOAc (4 × 50 mL). The organic layer was dried (Na<sub>2</sub>SO<sub>4</sub>) and the solvent removed under vacuum. The residue was purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 7:3) to afford **11** (6.814 g, 72%) as a white solid. *R*<sub>f</sub> 0.43 (hexane/EtOAc, 6:4); m.p. 89–90 °C. IR (KBr):  $\bar{\nu}$  3431, 2958, 1732, 1612, 1512, 1479, 1433, 1309, 1208, 1136, 1081, 1032, 953, 837, 793, 766, 712 cm<sup>-1</sup>. <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>):  $\delta$  3.79 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.80 (s, 3H, OCH<sub>3</sub>), 4.62 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 4.96 (br, 1H, OH), 6.30 (dd, *J* = 8.5, 2.7 Hz, 1H, H-5'), 6.46 (d, *J* = 2.7 Hz, 1H, H-3'), 6.75 (d, *J* = 8.5 Hz, 1H, H-6'). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>):  $\delta$  52.2 (CO<sub>2</sub>CH<sub>3</sub>), 55.7 (OCH<sub>3</sub>), 67.6 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 100.7 (C-3'), 106.0 (C-5'), 116.7 (C-6'), 140.9 (C-1'), 150.7 (C-2'), 151.8 (C-4'), 170.2 (CO<sub>2</sub>CH<sub>3</sub>). HRMS (EI): calcd. for C<sub>10</sub>H<sub>12</sub>O<sub>5</sub> [M]<sup>+</sup> 212.0685; found: 212.0681.

### General procedure for preparing 5-acylphenoxyacetic methyl esters 5a–c

Under an N<sub>2</sub> atmosphere and at rt, a mixture of **11** (1.0 mol equiv.) and of the respective acyl chloride (1.5 mol equiv.) was stirred for 10 min. Subsequently, BF<sub>3</sub>·OEt<sub>2</sub> (1.0 mol equiv.) was added, stirred at rt for 20 min, and then heated to 80 °C for 4 h. Ice was poured into the reaction mixture, followed by adding a saturated aqueous solution of NaHCO<sub>3</sub> dropwise until neutral, then washed with EtOAc (3 × 50 mL). The organic extract was dried (Na<sub>2</sub>SO<sub>4</sub>) and the solvent removed under vacuum. The residue was purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 85:15).

### Methyl 2-(5-acetyl-4-hydroxy-2-methoxyphenoxy)acetate (5a)

Following the general procedure, **11** (1.000 g, 4.72 mmol) was mixed with acetyl chloride (0.555 g, 7.07 mmol) and BF<sub>3</sub>·OEt<sub>2</sub> (0.670 g, 4.72 mmol) to provide **5a** (0.91 g, 76%) as a white solid. *R*<sub>f</sub> 0.31 (hexane/EtOAc, 7:3); m.p. 125–126 °C. IR (KBr):  $\bar{\nu}$  3472, 2974, 2372, 1744, 1640, 1614, 1503, 1440, 1365, 1334, 1265, 1202, 1163, 1058, 1007, 975, 814, 710 cm<sup>-1</sup>. <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>):  $\delta$  2.53 (s, 3H, CH<sub>3</sub>CO), 3.80 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.90 (s, 3H,

OCH<sub>3</sub>), 4.62 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 6.46 (s, 1H, H-3'), 7.29 (s, 1H, H-6'), 12.61 (s, 1H, OH). <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>):  $\delta$  26.3 (CH<sub>3</sub>CO), 52.2 (CO<sub>2</sub>CH<sub>3</sub>), 56.1 (OCH<sub>3</sub>), 68.1 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 100.9 (C-3'), 111.9 (C-5'), 118.3 (C-6'), 139.6 (C-1'), 157.8 (C-2'), 161.2 (C-4'), 169.5 (CO<sub>2</sub>CH<sub>3</sub>), 202.2 (CH<sub>3</sub>CO). MS (70 eV): *m/z* 254 (M<sup>+</sup>, 51), 239 (6), 195 (7), 181 (100), 153 (14), 135 (15), 107 (9). HRMS (EI): calcd for C<sub>12</sub>H<sub>14</sub>O<sub>6</sub> [M]<sup>+</sup>: 254.0790; found: 254.0787.

### Methyl 2-(4-hydroxy-2-methoxy-5-propionylphenoxy)acetate (5b)

Following the general procedure, **11** (1.000 g, 4.72 mmol) was mixed with propanoyl chloride (0.654 g, 7.07 mmol) and BF<sub>3</sub>·OEt<sub>2</sub> (0.670 g, 4.72 mmol) to obtain **5b** (0.91 g, 76%) as a white solid. *R*<sub>f</sub> 0.40 (hexane/EtOAc, 7:3); m.p. 105–106 °C. IR (KBr):  $\bar{\nu}$  3482, 2979, 2942, 1748, 1638, 1616, 1565, 1504, 1439, 1375, 1319, 1249, 1193, 1158, 1069, 1007, 975, 945, 820, 707 cm<sup>-1</sup>. <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>):  $\delta$  1.22 (t, *J* = 7.2 Hz, 3H, CH<sub>3</sub>CH<sub>2</sub>CO), 2.90 (q, *J* = 7.2 Hz, 2H, CH<sub>3</sub>CH<sub>2</sub>CO), 3.80 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.90 (s, 3H, OCH<sub>3</sub>), 4.62 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 6.46 (s, 1H, H-3'), 7.33 (s, 1H, H-6'), 12.70 (s, 1H, OH). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>):  $\delta$  8.3 (CH<sub>3</sub>CH<sub>2</sub>CO), 31.2 (CH<sub>3</sub>CH<sub>2</sub>CO), 52.1 (CO<sub>2</sub>CH<sub>3</sub>), 56.0 (OCH<sub>3</sub>), 68.0 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 100.9 (C-3'), 111.3 (C-5'), 117.5 (C-6'), 139.5 (C-1'), 157.4 (C-2'), 161.1 (C-4'), 169.5 (CO<sub>2</sub>CH<sub>3</sub>), 204.9 (CH<sub>3</sub>CH<sub>2</sub>CO). MS (70 eV): *m/z* 268 (M<sup>+</sup>, 66), 239 (66), 195 (100), 138 (22), 121 (14). HRMS (EI): *m/z* calcd for C<sub>13</sub>H<sub>16</sub>O<sub>6</sub> [M]<sup>+</sup>: 268.0947; found: 268.0946.

### Methyl 2-(5-butyryl-4-hydroxy-2-methoxyphenoxy)acetate (5c)

Following the general procedure, **11** (1.000 g, 4.72 mmol) was mixed with butanoyl chloride (0.753 g, 7.07 mmol) and BF<sub>3</sub>·OEt<sub>2</sub> (0.670 g, 4.72 mmol) to produce **5c** (0.892 g, 67%) as a white solid. *R*<sub>f</sub> 0.46 (hexane/EtOAc, 7:3); m.p. 82–83 °C. IR (KBr):  $\bar{\nu}$  3482, 2954, 1748, 1637, 1511, 1442, 1384, 1336, 1283, 1253, 1197, 1162, 1075, 1019, 986, 900, 848, 793, 703 cm<sup>-1</sup>. <sup>1</sup>H NMR (300 MHz, CDCl<sub>3</sub>):  $\delta$  1.01 (t, *J* = 7.5 Hz, 3H, CH<sub>3</sub>(CH<sub>2</sub>)<sub>2</sub>CO), 1.75 (sext, *J* = 7.5 Hz, 2H, CH<sub>3</sub>CH<sub>2</sub>CH<sub>2</sub>CO), 2.84 (t, *J* = 7.5 Hz, 2H, CH<sub>3</sub>CH<sub>2</sub>CH<sub>2</sub>CO), 3.81 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.91 (s, 3H, OCH<sub>3</sub>), 4.63 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 6.46 (s, 1H, H-3'), 7.32 (s, 1H, H-6'), 12.78 (s, 1H, OH). <sup>13</sup>C NMR (75 MHz, CDCl<sub>3</sub>):  $\delta$  13.8 (CH<sub>3</sub>(CH<sub>2</sub>)<sub>2</sub>CO), 18.1 (CH<sub>3</sub>CH<sub>2</sub>CH<sub>2</sub>CO), 39.9 (CH<sub>3</sub>CH<sub>2</sub>CH<sub>2</sub>CO), 52.2 (CO<sub>2</sub>CH<sub>3</sub>), 56.1 (OCH<sub>3</sub>), 68.0 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 100.87 (C-3'), 111.4 (C-5'), 117.5 (C-6'), 139.5 (C-1'), 157.4 (C-2'), 161.2 (C-4'), 169.5 (CO<sub>2</sub>CH<sub>3</sub>), 204.6 (CH<sub>3</sub>(CH<sub>2</sub>)<sub>2</sub>CO). MS (70 eV): *m/z* 282 (M<sup>+</sup>, 66), 239 (68), 209 (100), 167 (14), 138 (20), 107 (10), 69 (16).



HRMS (EI):  $m/z$  calcd for  $C_{14}H_{18}O_6$   $[M]^+$ : 282.1103; found: 282.1100.

### General procedure for preparing 5-alkylphenoxyacetic methyl esters 6a–c

Under an  $N_2$  atmosphere and at rt, a mixture of Zn/Hg and HCl (36%) (1.0 g/1.0 mL, 10.0 mol equiv.) was stirred for 5 min before being poured into a flask. Addition was made of the corresponding acyl derivative **5a–c** (1.0 mol equiv.), followed by a mixture of MeOH/HCl (1.0/0.2 mL). After heating to 60 °C for 4 h, ice was poured in and a saturated aqueous solution of  $NaHCO_3$  added dropwise until neutral, and then washed with EtOAc (3 × 30 mL). The organic extract was dried ( $Na_2SO_4$ ) and the solvent was removed under vacuum. The residue was purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 95:5).

#### Methyl 2-(5-ethyl-4-hydroxy-2-methoxyphenoxy)acetate (6a)

Following the general procedure, **5a** (1.000 g, 3.94 mmol) was mixed with Zn/Hg (10.47 g, 39.4 mmol) to furnish **6a** (0.68 g, 72%) as a white solid.  $R_f$  0.26 (hexane/EtOAc, 7:3); m.p. 66–67 °C. IR (KBr):  $\bar{\nu}$  3402, 2961, 2925, 2866, 1750, 1717, 1616, 1525, 1447, 1423, 1367, 1293, 1237, 1204, 1122, 1069, 1007, 857, 747, 706  $cm^{-1}$ .  $^1H$  NMR (500 MHz,  $CDCl_3$ ):  $\delta$  1.17 (t,  $J = 7.5$  Hz, 3H, H-2''), 2.52 (q,  $J = 7.5$  Hz, 2H, H-1''), 3.78 (s, 6H,  $CO_2CH_3$ , OCH<sub>3</sub>), 4.62 (s, 2H,  $CH_2CO_2CH_3$ ), 4.77–4.84 (m, 1H, OH), 6.42 (s, 1H, H-3'), 6.73 (s, 1H, H-6').  $^{13}C$  NMR (125 MHz,  $CDCl_3$ ):  $\delta$  14.2 (C-2''), 22.3 (C-1''), 52.1 ( $CO_2CH_3$ ), 55.9 (OCH<sub>3</sub>), 68.0 ( $CH_2CO_2CH_3$ ), 101.1 (C-3'), 117.7 (C-6'), 120.9 (C-5'), 141.0 (C-1'), 148.8 (C-2', C-4'), 170.1 ( $CO_2CH_3$ ). HRMS (EI):  $m/z$  calcd for  $C_{12}H_{16}O_5$   $[M]^+$ : 240.0998; found: 240.1000.

#### Methyl 2-(4-hydroxy-2-methoxy-5-propylphenoxy)acetate (6b)

Following the general procedure, **5b** (1.000 g, 3.73 mmol) was mixed with Zn/Hg (9.93 g, 37.3 mmol) to form **6b** (0.663 g, 70%) as a white solid.  $R_f$  0.30 (hexane/EtOAc, 7:3); m.p. 54–55 °C. IR (KBr):  $\bar{\nu}$  3390, 3019, 2955, 2867, 1871, 1708, 1619, 1528, 1450, 1419, 1377, 1302, 1198, 1120, 1078, 1008, 884, 866, 843, 768, 739, 703  $cm^{-1}$ .  $^1H$  NMR (300 MHz,  $CDCl_3$ ):  $\delta$  0.93 (t,  $J = 7.4$  Hz, 3H, H-3''), 1.57 (sext,  $J = 7.4$  Hz, 2H, H-2''), 2.46 (t,  $J = 7.4$  Hz, 2H, H-1''), 3.73 (s, 3H, OCH<sub>3</sub>), 3.78 (s, 3H,  $CO_2CH_3$ ), 4.61 (s, 2H,  $CH_2CO_2CH_3$ ), 5.45 (br s, 1H, OH), 6.42 (s, 1H, H-3'), 6.69 (s, 1H, H-6').  $^{13}C$  NMR (75 MHz,  $CDCl_3$ ):  $\delta$  13.8 (C-3''), 23.0 (C-2''), 31.3 (C-1''), 52.1 ( $CO_2CH_3$ ), 55.7 (OCH<sub>3</sub>),

67.8 ( $CH_2CO_2CH_3$ ), 100.9 (C-3'), 118.1 (C-6'), 119.5 (C-5'), 140.5 (C-1'), 148.5 (C-2'), 149.1 (C-4'), 170.3 ( $CO_2CH_3$ ). HRMS (EI):  $m/z$  calcd for  $C_{13}H_{18}O_5$   $[M]^+$ : 254.1154; found: 254.1151.

#### Methyl 2-(5-butyl-4-hydroxy-2-methoxyphenoxy)acetate (6c)

Following the general procedure, **5c** (1.000 g, 3.55 mmol) was mixed with Zn/Hg (9.43 g, 35.5 mmol), resulting in **6c** (0.65 g, 68%) as a white solid.  $R_f$  0.31 (hexane/EtOAc, 7:3); m.p. 49–50 °C. IR (KBr):  $\bar{\nu}$  3448, 3019, 2952, 2860, 1719, 1617, 1529, 1449, 1417, 1367, 1306, 1206, 1123, 1078, 1013, 939, 894, 856, 829, 759, 714  $cm^{-1}$ .  $^1H$  NMR (500 MHz,  $CDCl_3$ ):  $\delta$  0.94 (t,  $J = 7.5$  Hz, 3H, H-4''), 1.35 (sext,  $J = 7.5$  Hz, 2H, H-3''), 1.53 (qu,  $J = 7.5$  Hz, 2H, H-2''), 2.45 (t,  $J = 7.5$  Hz, 2H, H-1''), 3.79 (s, 3H,  $CO_2CH_3$ ), 3.80 (s, OCH<sub>3</sub>), 4.61 (s, 2H,  $CH_2CO_2CH_3$ ), 6.42 (s, 1H, H-3'), 6.70 (s, 1H, H-6').  $^{13}C$  NMR (125 MHz,  $CDCl_3$ ):  $\delta$  13.9 (C-4''), 22.4 (C-3''), 29.0 (C-1''), 32.1 (C-2''), 52.1 ( $CO_2CH_3$ ), 56.0 (OCH<sub>3</sub>), 68.0 ( $CH_2CO_2CH_3$ ), 101.1 (C-3'), 118.4 (C-6'), 119.4 (C-5'), 141.0 (C-1'), 148.9 (C-2', C-4'), 170.0 ( $CO_2CH_3$ ). HRMS (EI):  $m/z$  calcd for  $C_{14}H_{20}O_5$   $[M]^+$ : 268.1311; found: 268.1303.

#### Methyl 2-(4-Formyl-2-methoxy-6-nitrophenoxy)acetate (7a)

Under  $N_2$  atmosphere, a mixture of **12** (10.000 g, 50.76 mmol), dry  $K_2CO_3$  (14.010 g, 101.52 mmol) and methyl bromoacetate (15.10 g, 98.7 mmol) in anhydrous DMF (50 mL) was stirred at 60 °C for 4 h (Brown et al. 2010). The reaction mixture was filtered, the solvent removed under vacuum, and the residue purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 8:2) to give **7a** (9150 g, 67%) as a yellow solid.  $R_f$  0.66 (hexane/EtOAc, 1:1); m.p. 96–97 °C. IR (KBr):  $\bar{\nu}$  3385, 3086, 3014, 2956, 2851, 1752, 1701, 1604, 1541, 1466, 1427, 1357, 1289, 1218, 1202, 1141, 1046, 918, 872, 772, 680  $cm^{-1}$ .  $^1H$  NMR (300 MHz,  $CDCl_3$ ):  $\delta$  3.78 (s, 3H,  $CO_2CH_3$ ), 3.99 (s, 3H, OCH<sub>3</sub>), 4.90 (s, 2H,  $CH_2CO_2CH_3$ ), 7.64 (d,  $J = 1.8$  Hz, 1H, H-3'), 7.88 (d,  $J = 1.8$  Hz, 1H, H-5'), 9.93 (s, 1H, CHO).  $^{13}C$  NMR (75 MHz,  $CDCl_3$ ):  $\delta$  52.3 ( $CO_2CH_3$ ), 56.8 (OCH<sub>3</sub>), 69.4 ( $CH_2CO_2CH_3$ ), 113.5 (C-3'), 119.7 (C-5'), 131.7 (C-4'), 145.2 (C-1'), 153.6 (C-2'), 168.5 ( $CO_2CH_3$ ), 188.9 (CHO). HRMS (EI):  $m/z$  calcd for  $C_{11}H_{11}NO_7$   $[M]^+$ : 269.0535; found: 269.0541.

#### Methyl 2-(4-hydroxy-2-methoxy-6-nitrophenoxy)acetate (7b)

A mixture of **7a** (1.000 g, 3.72 mmol) and *m*CPBA (77%) (1.240 g, 5.57 mmol) in  $CH_2Cl_2$  (10 mL) was stirred at rt for 4 h before removing the solvent under vacuum. MeOH

(10 mL) and HCl (37%) (1 mL) were added, the mixture was stirred at rt for 2 h, and the solvent was removed under vacuum. An aqueous saturated solution of NaHCO<sub>3</sub> was added until neutral, then extracted with EtOAc (4 × 50 mL). The organic layer was dried (Na<sub>2</sub>SO<sub>4</sub>), the solvent removed under vacuum, and the residue purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 1:1) to afford **7b** (0.580 g, 61%) as a pale yellow solid. *R<sub>f</sub>* 0.33 (EtOAc); m.p. 110–111 °C. IR (KBr):  $\bar{\nu}$  3070, 2952, 2643, 1732, 1609, 1542, 1490, 1466, 1421, 1354, 1289, 1222, 1202, 1114, 1066, 1017, 918, 897, 865, 854, 814, 775, 755, 736, 700 cm<sup>-1</sup>. <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>):  $\delta$  3.79 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.99 (s, 3H, OCH<sub>3</sub>), 4.89 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 7.80 (d, *J* = 2.0 Hz, 1H, H-3'), 8.12 (d, *J* = 2.0 Hz, 1H, H-5'). <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>):  $\delta$  52.3 (CO<sub>2</sub>CH<sub>3</sub>), 56.8 (OCH<sub>3</sub>), 69.5 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 116.8 (C-3'), 118.7 (C-5'), 124.6 (C-6'), 144.6 (C-1'), 153.0 (C-2'), 168.6 (CO<sub>2</sub>CH<sub>3</sub>), 169.1 (C-4'). HRMS (EI): *m/z* calcd for C<sub>10</sub>H<sub>11</sub>NO<sub>7</sub> [M]<sup>+</sup>: 257.0536; found: 257.0531.

#### Methyl 2-(4-(hydroxymethyl)-2-methoxy-6-nitrophenoxy) acetate (**7c**)

Under N<sub>2</sub> atmosphere and at 0 °C, a mixture of **7a** (1.000 g, 3.71 mmol) and NaBH<sub>4</sub> (0.070 g, 1.85 mmol) in MeOH (10 mL) was stirred for 2 h before removing the solvent under vacuum. The residue was purified by column chromatography over silica gel (20 g/g of crude, hexane/EtOAc, 1:1) to provide **7c** (0.86 g, 86%) as a white solid. *R<sub>f</sub>* 0.16 (hexane/EtOAc, 1:1); m.p. 97–98 °C. IR (KBr):  $\bar{\nu}$  3481, 2921, 1747, 1535, 1452, 1390, 1355, 1276, 1241, 1208, 1135, 1062, 972, 919, 850, 810, 780 cm<sup>-1</sup>. <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>):  $\delta$  2.47 (br, 1H, OH), 3.79 (s, 3H, CO<sub>2</sub>CH<sub>3</sub>), 3.90 (s, 3H, OCH<sub>3</sub>), 4.68 (s, 2H, CH<sub>2</sub>OH), 4.73 (s, 2H, CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 7.14 (d, *J* = 1.5 Hz, 1H, H-3'), 7.29 (d, *J* = 1.5 Hz, 1H, H-5'). <sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>):  $\delta$  52.2 (CO<sub>2</sub>CH<sub>3</sub>), 56.4 (OCH<sub>3</sub>), 63.7 (CH<sub>2</sub>OH), 69.6 (CH<sub>2</sub>CO<sub>2</sub>CH<sub>3</sub>), 113.7 (C-5'), 114.2 (C-3'), 138.1 (C-4'), 139.2 (C-1'), 144.5 (C-6'), 153.3 (C-2'), 169.1 (CO<sub>2</sub>CH<sub>3</sub>). HRMS (EI): *m/z* calcd for C<sub>11</sub>H<sub>13</sub>NO<sub>7</sub>: 271.0692; found: 271.0687.

#### Hypolipidemic activity

The hypolipidemic effect of the compounds was studied on male ICR mice weighing 25–30 g (Birmex, Mexico City), which were housed in hanging metal cages and maintained at 24 ± 2 °C and 50 ± 10% relative humidity on a 12 h light/dark cycle (lights on at 8:00 a.m.). Food (a standard pellet diet, Rodent Diet 5001, PMI Nutrition International, Brentwood, MO) and water were freely available. All animals appeared healthy throughout the dosing period, maintaining normal food intake and weight gain. Post-sacrifice analysis

showed no gross abnormalities in any treated mice. All animals were handled and maintained in accordance with ethical principles and regulations specified by the Animal Care and Use Committee of our institution and the standards of the National Institutes of Health of Mexico.

An aqueous solution of Triton WR 1339 was administered intraperitoneally (ip) to mice (400 mg/kg) and after 1 h the test compounds (25, 50, or 100 mg/kg, dissolved in saline, or saline-Tween) were injected ip. Blood was taken from a retro-orbital puncture 24 h later, and the plasma levels of total cholesterol (TC) and triglycerides (TG) were determined in duplicate with commercially available kits. Values are expressed as the mean from 6 animals per compound.

#### Antifungal activity

The in vitro antimicrobial effect of compounds **5a–c**, **6a–b**, **7a**, **7c**, and **9a–d** was evaluated with susceptibility assays based on the microdilution techniques of the Antimicrobial Susceptibility Testing Protocols standardized by the Clinical and Laboratory Standards Institute (CLSI). The M38-A2 method was used for the four strains of filamentous fungi (CLSI 2002; Espinel-Ingroff and Canton 2007a): *Aspergillus fumigatus* ATCC-16907, *Trichosporon cutaneum* ATCC-28592, *Rhizopus oryzae* ATCC-10329 and *Mucor hiemalis* ATCC-8690. The M27-A3 method was applied for the three strains of ATCC *Candida* yeasts (CLSI 2008; Espinel-Ingroff and Cantón 2007b; Pfaller and Diekema 2012): *C. albicans* ATCC-10231, *C. utilis* ATCC-9226 and *C. tropicalis* ATCC-13803. The test compounds were soluble in dimethyl sulfoxide (DMSO), in accordance with the Antimicrobial Susceptibility Testing Protocols. Hence, the antifungal assays were performed with this solvent.

The minimum inhibitory concentration (MIC) of all compounds and the reference drug (itraconazole), expressed in µg/mL, was determined in 96-well plates, utilizing RPMI-1640 culture medium buffered with 3-(*N*-morpholino)propanesulfonic acid (MOPS, Sigma) and dilutions of 16 to 0.03 µg/mL (according to the procedure outlined in the instructions for each type of fungus).

The inoculum for the yeasts was prepared with culture colonies of 24-h growth strain in SDA medium and resuspending them in a tube of saline solution (0.85% NaCl) to obtain a concentration of ~1–5 × 10<sup>6</sup> CFU/mL. The necessary amount of saline solution was added to adjust the optical density to 0.5 McFarland. Subsequently, a 1:1000 dilution was made with RPMI medium. For filamentous fungi, the inoculum was elaborated from a culture having undergone 7 days of growth at 35 °C on PDA agar, a medium that induces the formation of conidia or sporangiospores. To remove the conidia, the culture was

introduced and handled in Tween 20, then resuspended in saline solution. The particles were allowed to settle for 5 min, the supernatant transferred to another tube and shaken vigorously for 15 s. Because the size of the conidia is different for each species, the optical density required to reach a concentration of  $1\text{--}5 \times 10^6$  CFU/mL varied. Adjustment to the desired value was accomplished with the McFarland Turbidity Standard No. 0.15, based on the methods of the CLSI.

Compounds **5–7** and **9** were prepared (at 10 mM) and tested for their inhibition of *Fusarium oxysporum* with the PDA dilution method (Lalitha 2004). Briefly, PDA medium was sterilized in an autoclave at 121 °C for 20 min, followed by the addition of one of the compounds diluted in DMSO (2%). The plates were inoculated with *F. oxysporum* and then incubated at 26–28 °C for 72 h. All experiments were carried out in triplicates ( $n = 12$ ), and the data are expressed as the mean. The growth inhibition efficiency of each compound was compared with captan (10 mM), the reference drug, and validated by ANOVA and the Tukey test ( $p < 0.05$ ) on the SAS V9.0 program (SAS 2014).

## Cytotoxic activity

### Cell culture

Cell lines of human cervical cancer (HeLa), prostate cancer (DU-145), breast cancer (MDA-231), and no tumoral cell line (HaCaT) were cultivated in Dulbecco's Minimum Essential Medium (DMEM) containing 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin (100/100 U/mL). Cells were cultured as adherent monolayers and maintained at 37 °C and 5% humidity (Freshney 2010).

### Cell viability assay

Cell viability was determined by an MTT (3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide) assay. Cells were harvested, counted, and transferred to 96-well plates at 3000 cells per well to be incubated for 24 h. Upon completion of this time, cells were treated with compounds **5a–b**, **6a–b**, and **9a–c** at concentrations of 10, 20, 40, 60, 80, and 100  $\mu\text{g/mL}$  with 1% DMSO in cell culture medium, then again incubated for 24 h. Subsequently, microscope images were obtained for each cell line exposed to each compound. Cells without treatment were used as the viability control, 1% DMSO as the vehicle control, and methotrexate as the control for comparison of the respective median lethal concentrations ( $\text{LC}_{50}$ ).

In each well, 50  $\mu\text{L}$  of a solution of MTT in culture medium (100  $\mu\text{g}/\mu\text{L}$ ) were added and cells were incubated at 37 °C for 4 h. The culture medium was removed and 100  $\mu\text{L}$  of isopropanol was added to each well. Optical density in

each well was measured on an ELISA microplate reader at 595 nm. The result was expressed as the percentage of the viability of control cells and the  $\text{LC}_{50}$  was calculated (van Meerloo et al. 2011).

## Data analysis

Data analysis was carried out by nonparametric one-way ANOVA and a post hoc with Dunn's test, run on Sigma Plot software version 12 (Motulsky 1999). All data are expressed as the mean  $\pm$  SD, with significance set at  $p < 0.05$ . Each experiment was performed in triplicate.

## Antioxidant activity

### DPPH radical scavenging assay

The scavenging of free radicals by the chromones was assessed by using the 2,2-diphenyl-1-picrylhydrazyl radical (DPPH<sup>\*</sup>) assay, as previously described (Cevallos-Casals and Cisneros-Zevallos 2003) with slight modifications. A concentration of 10 mM of each compound was prepared, to which a solution of DPPH<sup>\*</sup> (133.33  $\mu\text{M}$ ) was added at a ratio of 1:3 (v/v). The mixture was incubated at 37 °C for 30 min and read at 517 nm. Scavenging capacity (SC) was expressed as the percentage decrease in DPPH at 10 mM:

$$\text{SC}\% = [(A_{\text{control}} - A_{\text{test}})/A_{\text{control}}] \times 100$$

where  $A_{\text{control}}$  is the absorbance of the DPPH solution (control) and  $A_{\text{test}}$  is the absorbance of the DPPH solution plus a compound.

### ABTS radical activity

The free radical scavenging capacity was quantified based on a slightly modified version of the previously reported 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) radical cation (ABTS<sup>•+</sup>) assay (Gallardo et al. 2016; Zhang et al. 2013). The radical cation was prepared by dissolving the stock solution of ABTS (7 mM in distilled water) with  $\text{K}_2\text{S}_2\text{O}_8$  solution (2.45 mM) at ratio of 1:1 (v/v) and then leaving the mixture to stand in the dark at rt for 16 h. For the evaluation of antioxidant activity, the ABTS<sup>•+</sup> solution was diluted with absolute ethanol until reaching the absorbency of  $0.700 \pm 0.02$  at 734 nm. Taking 10  $\mu\text{L}$  of the resulting solution at different concentrations (10, 1.0 and 0.1 mM), 1 mL of ABTS solution was added and after 6 min absorbance was read at 734 nm. All tests were performed in triplicate and the mean was centered. Finally, the percentage inhibition of ABTS absorbance was calculated by the aforementioned formula for the DPPH<sup>\*</sup> assay. The data are expressed as Trolox equivalent antioxidant capacity (TEAC), developing a standard curve in a range of 0.5 to 3.5 mM.

## Molecular docking study

### Protein and ligand structures

The crystallographic structures of human HMGRh in complex with a selective inhibitor (simvastatin) were retrieved from the protein data bank (PDB) (<http://www.rcsb.org/>) with the code 1HW9 (Istvan and Deisenhofer 2001). Before docking simulations were run, the coordinates of the protein were set, water molecules removed, hydrogen atoms added to the polar atoms (considering pH at 7.4), and Kollman charges assigned. The 3D structures of simvastatin were downloaded from the Zinc database (Irwin and Shoichet 2005). The structures of the ligands were sketched in 2D with ChemSketch (<https://www.acdlabs.com/resources/freeware/chemsketch/>) and optimized with AM1 on Gaussian 98 software (Frisch et al. 2004) to obtain the minimum energy conformation for docking studies.

### Molecular docking

The protein-ligand interaction was observed on Autodock version 4.0 and AutoDockTools (Morris et al. 2009). All the possible rotatable bonds, torsion angles, atomic partial charges and non-polar hydrogens were determined for each ligand. For HMGRh, the grid dimensions in AutoDockTools were  $62 \times 82 \times 106 \text{ \AA}^3$  with points separated by  $0.375 \text{ \AA}$ , centered at  $X = 5.564$ ,  $Y = -6.681$ , and  $Z = 4.637$ . The hybrid Lamarckian Genetic Algorithm was applied for minimization, utilizing default parameters. A total of one hundred docking runs were conducted, adopting the conformation with the lowest binding energy (kcal/mol) for all further simulations. The script and files were prepared and the docking results visualized on AutoDockTools, then edited in Discovery 4.0 Client (Dassault Systèmes 2016).

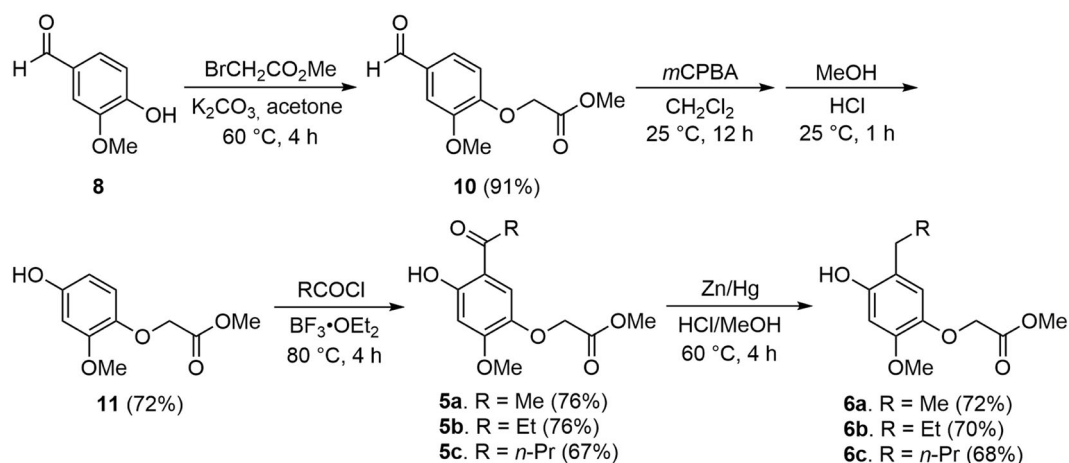
## Results and discussion

### Chemistry

The synthesis of the series of methyl phenoxyacetates **5a–c** and **6a–c** was based on the functionalization of vanillin (**8**) as the starting material (Scheme 1). Firstly, the hydroxy group was protected by the acetate moiety through a Williamson alkylation with methyl bromoacetate under basic conditions (Argüelles et al. 2010) to give compound **10** in high yield. The conversion of the formyl group to the hydroxy group was carried out by a Baeyer–Villiger rearrangement with *m*CPBA, followed by hydrolysis of the formyl ester to furnish the corresponding acylphenoxyacetic esters **5a–c** in fairly good yields (Scheme 1). Conversion of this series into the alkylphenoxyacetic esters **6a–c** took place through a Clemmensen reduction to provide the expected products in good yields.

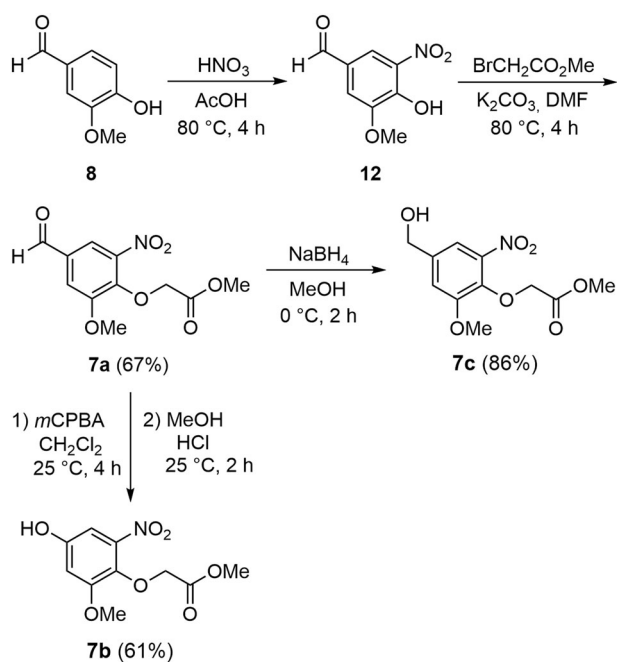
The structures of all compounds were established by  $^1\text{H}$  NMR,  $^{13}\text{C}$  NMR, and HRMS. The full assignment of proton and carbon signals was achieved by 2D NMR experiments (HMQC and HMBC). The selective formation of the tetrasubstituted benzene ring was ascertained by the two characteristic singlets for the *para* aromatic protons. The presence of the carbonyl group of the series of acyl derivatives **5a–c** was verified by the signal appearing at around 204 ppm in the  $^{13}\text{C}$  NMR spectra.

The series of nitro compounds **7a–c** was readily prepared by direct nitration of vanillin (**8**) to provide the known 5-nitrovanillin (**12**) (Grenier et al. 2000; Kiss et al. 2010;



**Scheme 1** Synthesis of phenol **10**, acyl phenols **5a–c**, and alkyl phenols **6a–c**





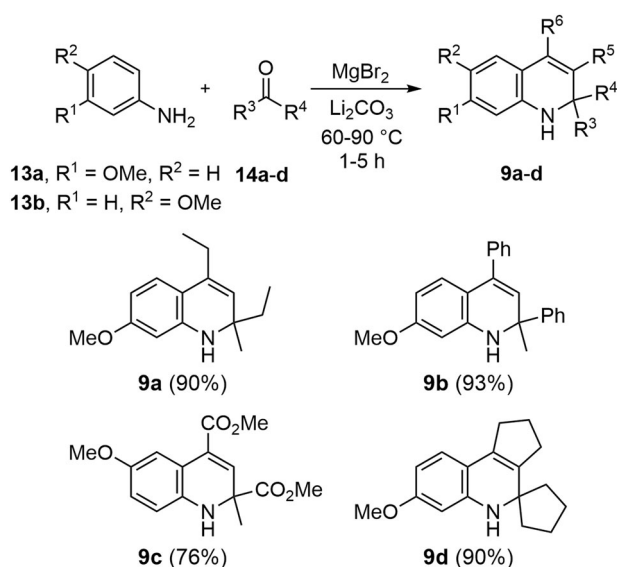
**Scheme 2** Synthesis of methyl nitrophenoxyacetates **7a-c** from vanillin (**8**)

Rashid et al. 2018) (Scheme 2). The latter was treated with methyl bromoacetate under basic conditions to obtain the nitrophenoxyacetate **7a** in a 67% yield for the two steps. Baeyer–Villiger rearrangement produced the respective formate, which was hydrolyzed under acid conditions to deliver **7b** in 61% yield. Moreover, reduction of the formyl group of **7a** with sodium borohydride resulted in the hydroxymethyl derivative **7c** in 86% yield.

The synthesis of 1,2-dihydroquinolines **9a-d** was achieved through a single-step reaction, according to the published procedure (Gutiérrez et al. 2013) (Scheme 3). Thus, analogs **9a-b** were prepared in high yields by reacting *m*-anisidine (**13a**) with ketones **14a-b**, under solvent-free conditions and in the presence of magnesium bromide as the catalyst and lithium carbonate as the base. Meanwhile, 1,2-dihydroquinoline **9c** was formed by using *p*-anisidine (**13b**) and methyl pyruvate (**14c**) under similar reaction conditions. Finally, the spiro-polycyclic 1,2-dihydroquinoline **9d** was furnished in high yield via the reaction of **13a** with cyclopentanone (**14d**). The physical data and NMR spectra of the resulting 1,2-dihydroquinolines were in agreement with the previous report (Gutiérrez et al. 2013).

### Hypolipidemic activity

The *in vivo* hypolipidemic screening of the compounds **5a-c**, **6a-c**, **7a**, and **9a-d** was performed on male ICR mice subjected to tyloxapol-induced hyperlipidemia (Silva et al. 2001; Kourounakis et al. 2002). The possible mechanism of lipid-lowering agents is commonly evaluated with Triton WR 1339



**Scheme 3** Synthesis of 1,2-dihydroquinolines **9a-d**

(tyloxapol), a non-ionic surfactant (Levine and Saltzman 2007; Zeniya and Reuben 1988; Edelstein et al. 1985). It causes a drastic rise in serum triglycerides and cholesterol levels by increasing HMGR activity (Kuroda et al. 1977; Goldfard 1978). To diminish the concentration of blood serum lipids, it is necessary to inhibit the synthesis of endogenous cholesterol (elevated after treatment with tyloxapol), limit the absorption of lipoprotein lipids, and stimulate the excretion of the latter (Korolenko et al. 2010). Many clinically used drugs accomplish some of these functions, such as ezetimibe (inhibitor of cholesterol absorption), fibrates (PPAR- $\alpha$  agonists) and statins (HMGR inhibitors).

The series of analogs **5a-c**, **6a-c**, **7a**, and **9a-d** were administered *ip* at doses of 25, 50, and 100 mg/kg to hyperlipidemic mice, and their effect on the plasma levels (mg/dL) of total cholesterol, and triglycerides was examined with a procedure similar to that employed in previous studies with related compounds (Argüelles et al. 2010). Since preliminary assays showed a very modest decrease in cholesterol by **7a-c**, and considering that the nitro group can be cytotoxic (Chung et al. 2011; Olender et al. 2018), only **7a** was subjected to further assessment of hypolipidemic activity.

All compounds evaluated, including the reference (simvastatin), sharply depleted plasma levels of cholesterol and triglycerides, in the range of 44–75% and 51–93%, respectively, vs. the hyperlipidemic control animals. A significant difference existed between the values of the control and most treatments (Table 1). Moreover, it is worth noticing that almost all compounds significantly decrease triglyceride levels, since it is critical to diminish the concentration of the latter because in excess they induce the formation of atheroma, leading to an imminent cardiovascular disease (Veseli et al. 2017).

**Table 1** Effect of the test compounds on the serum lipid profile of male ICR mice<sup>a</sup>

Compound	Dose mg/ Kg/day	Cholesterol	Triglycerides
<b>Normal diet</b>		-78.15 ± 0.07*	-80.02 ± 0.08*
<b>Simvastatin</b>	17	-76.38 ± 0.19*	-79.10 ± 0.17*
<b>Tyloxapol</b>	400	100 ± 1.03 <sup>b</sup>	100 ± 0.33 <sup>c</sup>
<b>5a + Tyloxapol</b>	25	-67.29 ± 6.78*	-83.74 ± 2.44*
	50	-75.44 ± 6.27*	-93.40 ± 1.25*
	100	-58.5 ± 7.35*	-86.46 ± 2.31*
<b>5b + Tyloxapol</b>	25	-75.20 ± 3.52*	-81.42 ± 2.51*
	50	-57.49 ± 7.74*	-87.49 ± 1.25*
	100	-52.53 ± 11.61*	-88.95 ± 1.31*
<b>5c + Tyloxapol</b>	25	-73.43 ± 4.70*	-92.04 ± 2.51*
	50	-64.93 ± 11.06*	-91.4 ± 1.79*
	100	-74.02 ± 5.14*	-85.70 ± 1.84*
<b>6a + Tyloxapol</b>	25	-72.84 ± 0.25*	-20.77 ± 0.79
	50	-74.73 ± 0.09*	-80.59 ± 0.11*
	100	-65.40 ± 0.69*	-73.95 ± 0.38*
<b>6b + Tyloxapol</b>	25	-69.89 ± 0.49*	4.11 ± 1.14
	50	-61.03 ± 0.60*	4.11 ± 2.18
	100	-70.89 ± 0.58*	-9.55 ± 0.83
<b>6c + Tyloxapol</b>	25	-71.07 ± 0.70*	-76.25 ± 0.34
	50	-69.30 ± 0.30*	-75.72 ± 0.19*
	100	-72.72 ± 0.41*	-78.22 ± 0.29*
<b>7a + tyloxapol</b>	25	-58.43 ± 0.53	-67.13 ± 0.40*
	50	-59.03 ± 0.78	-66.98 ± 0.60*
	100	-17.46 ± 2.68	-43.01 ± 0.37
<b>9a + tyloxapol</b>	25	-72.27 ± 0.11*	-76.22 ± 0.24*
	50	-63.16 ± 0.62*	-63.26 ± 0.62*
	100	-72.72 ± 0.24*	-70.47 ± 0.44*
<b>9b + tyloxapol</b>	25	-68.82 ± 0.29*	-67.37 ± 0.55*
	50	-44.98 ± 1.45*	-34.11 ± 0.54
	100	-73.78 ± 0.33*	-75.25 ± 0.22*
<b>9c + tyloxapol</b>	25	-51.70 ± 1.01	-60.13 ± 0.46*
	50	28.22 ± 7.82	-66.94 ± 0.57*
	100	-65.99 ± 0.4	-82.79 ± 0.20*
<b>9d + tyloxapol</b>	25	-72.72 ± 0.42*	-73.92 ± 0.21*
	50	-63.99 ± 0.72*	-61.10 ± 0.75*
	100	-62.01 ± 0.39*	-51.36 ± 0.39*

\*Significantly different from the tyloxapol group ( $p < 0.05$ )

<sup>a</sup>Expressed as a percentage of the group treated with tyloxapol only (mean ± standard error,  $n = 6$ )

<sup>b</sup>141.16 mmol/L

<sup>c</sup>1.26 mmol/L

The best hypolipidemic agents were the acylphenoxycetic esters **5a–c**, depleting cholesterol and triglycerides to almost the same extent as the positive control (simvastatin). A similar result was observed for the analogous series

derived from isovanillin **4a–f** (Mendieta et al. 2014), as well as for the analogs to  $\alpha$ -asarone **3a–f** (Mendieta et al. 2014). However, an inverse hypolipidemic effect was found for other homologs structurally closer to  $\alpha$ -asarone (Cruz et al. 2001), among which the most potent activity was displayed by those containing an alkyl vs. acyl side chain.

Actually, the alkyl side chain would be expected to act as an important hypocholesterolemic pharmacophore, because of its strong lipophilic binding interaction with the HMGR active site (Argüelles et al. 2010). Nevertheless, there was no relation between the side chain length and the hypocholesterolemic effect, evidenced by the comparable reduction in serum cholesterol values of the homologs in the **5a–c** and **6a–c** series. Regardless of the possible synergetic effect of the acyl and alkyl side chains on the test compounds, the phenoxyacetic ester frame appears to play the main role as the pharmacophore moiety (Table 1), as has been reported (Argüelles et al. 2010; Cruz et al. 2003; Labarrios et al. 1999; Zúñiga et al. 2005; Mendieta et al. 2014).

Interestingly, among the four tested 1,2-dihydroquinolines, **9a**, **9b**, and **9d** displayed high hypocholesterolemic and hypotriglyceridemic activity, which is relevant and almost unique for this kind of heterocycles (Lagu et al. 2007; Matsuda et al. 2007; Guo et al. 2017). The change of substituents in the A ring of the heterocyclic frame does not seem to have a significant influence on the hypolipidemic effect of the compounds, suggesting that the 1,2-dihydroquinoline scaffold could be the active pharmacophore of these potential hypolipidemic agents. Despite the non-significant hypocholesterolemic activity of **9c**, it was able to diminish the level of triglycerides significantly.

Although a dose-response relationship cannot be established for each member of the series, in many cases the 25 mg/kg dose (the closest to that of the control group) produced the greatest effect (Table 1).

### Antifungal activity

Statins have been reported to produce in vitro activity against several human pathogenic fungi, including *Candida* spp. and *Aspergillus* spp. (Qiao et al. 2007; Cabral et al. 2013). Hence, the test compounds (except **6c** and **7b**) and itraconazole (the reference) were tested in vitro against four filamentous fungi (*Aspergillus fumigatus* ATCC-16907, *Trichosporon cutaneum* ATCC-28592, *Rhizopus oryzae* ATCC-10329, and *Mucor hiemalis* ATCC-8690) and three yeast specimens (*Candida albicans* ATCC-10231, *C. utilis* ATCC-9226, and *C. tropicalis* ATCC-13803) (Table 2). The MIC values (expressed in  $\mu\text{g/mL}$ ) of all compounds were determined in 96-well plates with 3-(*N*-morpholino)propanesulfonic acid (MOPS) as the buffer. Standardized microbiological methods developed by the CLSI (Fothergill 2012;

**Table 2** In vitro antifungal activities of most synthesized compounds (MIC,  $\mu\text{g/mL}$ )

Compound	Yeast fungi			Filamentous fungi			
	<i>C. alb.</i>	<i>C. uti.</i>	<i>C. trop.</i>	<i>M. hie.</i>	<i>A. fum.</i>	<i>R. ory.</i>	<i>T. cut.</i>
<b>5a</b>	8	8	4	16	16	16	8
<b>5b</b>	8	8	8	16	16	8	16
<b>5c</b>	2	16	16	16	16	8	16
<b>6a</b>	0.5	8	16	16	16	16	16
<b>6b</b>	0.5	16	16	16	16	1	16
<b>7a</b>	8	16	4	16	16	16	2
<b>7c</b>	4	8	16	16	16	16	16
<b>9a</b>	4	16	16	8	16	8	16
<b>9b</b>	16	16	8	16	16	1	16
<b>9c</b>	4	16	8	16	16	16	16
<b>9d</b>	1	8	16	16	16	16	16
Standard <sup>a</sup>	0.03	0.25	0.06	4	1	1	8

*C. alb.* *Candida albicans*, *C. uti.* *Candida utilis*, *C. trop.* *Candida tropicalis*, *M. hie* *Mucor hiemalis*, *A. fum* *Aspergillus fumigatus*, *R. ory* *Rhizopus oryzae*, *T. cut* *Trichosporon cutaneum*

<sup>a</sup>Itraconazole

La Regina et al. 2008; Ramírez-Villalva et al. 2017; García-Vanegas et al. 2019) were employed. The sensitivity of the filamentous microorganisms was determined by the microdilution M38-A method (CLSI 2002; Espinel-Ingroff and Canton 2007a), and that of the yeast fungi with the M27-A3 method (CLSI 2008; Espinel-Ingroff and Cantón 2007b; Pfaller and Diekema 2012).

A good effect was produced against *C. albicans* by **6a** and **6b**, though not as good as the inhibition elicited by itraconazole. Contrarily, a lower antifungal activity was found for **5c** and **9d**. None of the present derivatives were active against the yeast fungi *C. utilis* and *C. tropicalis*. Compared with the effect of itraconazole, only **6b** and **9b** demonstrated good growth inhibition of the filamentous fungus *R. oryzae*, and **5a** and **7a** of *T. cutaneum*.

No structure-activity correlation was detected among the active compounds in regard to the growth inhibition of *C. albicans*. However, homologs **6** (containing C-5 alkyl groups) gave a greater antifungal effect than homologs **5** (with the C-5 acyl moiety). Overall, these results indicate that the presence of the methyl phenoxyacetate and 1,2-dihydroquinoline scaffolds seems to impact antifungal capacity.

Fusariosis is one of the most common infections in humans (Garnica and Nucci 2013; Guarro 2013). *Fusarium oxysporum* is an invasive pathogen known to cause infections like keratitis and onychomycosis in immunocompetent individuals, but in immunocompromised patients the infections are frequently fatal (Garnica and Nucci 2013; Guarro and Gené 1995; Cordoba-Guijarro et al. 2002; Olivares et al. 2005). Although the optimal treatment against fusariosis is unclear, the modestly effective drugs consist of antifungal azoles and amphotericin B. In vitro

data demonstrates general resistance of *Fusarium* to the available antifungal drugs with poor MIC values (Guarro 2013; Cordoba-Guijarro et al. 2002).

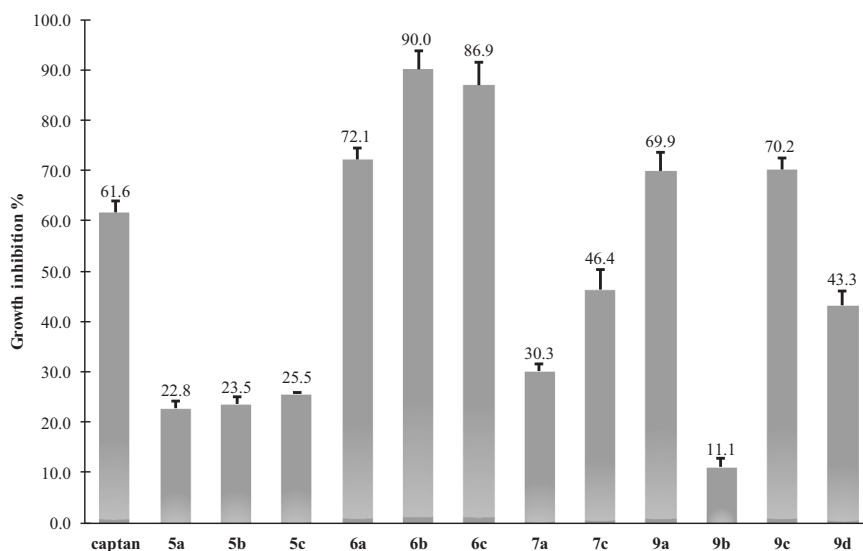
Consequently, the series **5–7** and **9** were tested as inhibitors of the mycelial growth of the pathogen *F. oxysporum*, using the agar dilution method and captan as the positive control. Captan, a fungicide utilized in agriculture, cosmetics and pharmaceuticals (U.S. Environmental Protection Agency 1984) inhibits the mycelial growth of *Fusarium* spp. (Türkkan and Erper 2015). According to previous studies, it also increases cell vulnerability to oxidative stress (Moreno-Aliaga et al. 1999; Inoue et al. 2018).

The compounds evaluated induced significant cell growth inhibition, from 72 to 90% for **6a–c**, and 70% for **9a** and **9c** (Fig. 3). Interestingly, the presence of the alkyl side chain of derivatives **6** significantly improved growth inhibition, with **6b** displaying the best result (90%). 1,2-Dihydroquinolines **9a** and **9c** contain either an alkyl side chain or alkyl carboxylate.

### Cytotoxic activity

The cytotoxic effect of **5a–b**, **6a–b**, and **9a–c** was determined through the MTT (3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide) cell viability assay with cell lines of human cervical cancer (HeLa), prostate cancer (DU-145), breast cancer (MDA-231) and normal skin keratinocytes (HaCaT) (van Meerloo et al. 2011). Death was observed in up to 90% of the cells exposed to these compounds in concentrations higher than 40  $\mu\text{g/mL}$ . In the MDA-231 and DU-145 cell lines, there was up to 50% cell death at 80–100  $\mu\text{g/mL}$ . HeLa and HaCaT cells showed 100% viability following exposure to the non-active

**Fig. 3** Growth inhibition of *F. oxysporum* produced by **5a–c**, **6a–c**, **7a**, **7c**, and **9a–d** (at 10 mM) after 72 h of incubation. Data are expressed as the mean  $\pm$  standard deviation ( $n = 12$ ) and are significantly different from the control ( $p < 0.05$ )



**Table 3** Median lethal concentration (LC<sub>50</sub>) (μg/mL) calculated for derivatives **6a–b** in the four cell lines evaluated

Compound	Cell line/values	MDA-231	DU-145	HeLa	HaCaT <sup>a</sup>
<b>6a</b>	LC <sub>50</sub> (IC 95%)	24 (8–40)	27 (19–35)	44 (25–62)	19 (11–27)
<b>6b</b>	LC <sub>50</sub> (IC 95%)	–	–	73 (60–85)	69 (53–85)
Methotrexate <sup>b</sup>	LC <sub>50</sub> (IC 95%)	>100	>100	>100	>100

A 95% confidential interval (CI 95%) was calculated for each LC<sub>50</sub>

<sup>a</sup>No tumoral cell line

<sup>b</sup>The reference drug

compounds. In cells incubated exclusively with the vehicle (DMSO), no death occurred, and the decrease in cell growth relative to the control was non-significant.

Only compounds **6a**, **b** exhibited cytotoxicity (Table 3). Derivative **6a** produced an exponential decline in cell viability in all cell lines, including non-tumor cells (HaCaT). Derivative **6b** (at 100 μg/mL) caused greater cytotoxicity in the cervical carcinoma cell line (HeLa) and HaCaT (control) than the other cell lines. Hence the activity of these compounds was non-selective, a characteristic that may be improved by future modifications in their structure. Their selectivity should be thoroughly studied in various cell lines (Xiao et al. 2018) to seek specificity for tumor cells.

### Antioxidant activity

Antioxidants are agents capable of protecting a biological target against oxidative stress, DNA mutations and cell damage, helping to prevent many human diseases. For example, oxidative stress plays a pivotal role in the development of diabetes, cancer, and cardiovascular disease (Pisoschi and Pop 2015). The antioxidant potential was herein evaluated for the synthesized compounds for two main reasons that suggest a free radical scavenging capacity of these derivatives. Firstly, alkylphenols **6a–c** are analogs

of  $\alpha$ -asarone (**1**), which has antioxidant properties (Pages et al. 2010). Secondly, the 1,2-dihydroquinoline ethoxyquin is a known antioxidant feed additive (Dorey et al. 2000; Błaszczuk and Skolimowski 2007; Błaszczuk et al. 2013; de Koning 2002; Błaszczuk et al. 2006; Błaszczuk and Skolimowski 2005; Błaszczuk and Skolimowski 2006).

The antioxidant activity of the series of analogs **5a–c**, **6a–c**, **7a**, **7c**, and **9a–d** was assessed by using the radical scavenging method with the 2,2-diphenyl-1-picrylhydrazyl radical (DPPH<sup>•</sup>) and 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) radical cation (ABTS<sup>•+</sup>). Antioxidant activity was expressed as the percentage of decrease in DPPH and ABTS (Table 4). Butylated hydroxytoluene (BHT) was the positive control for the DPPH assay, and the TEAC (mM) was the unit for expressing the data of the ABTS assay (Huang et al. 2005).

The **6a–c** and **9a–d** series showed good radical scavenging capacity in the DPPH assay. The IC<sub>50</sub> values for **6a–c** were in a narrow range (0.24–0.25 mM) and for **9a–d** in a much wider range (0.12–1.52 mM). Moreover, a stronger antioxidant effect was detected for **6a–c**, **9a**, and **9c** than BHT (based on the IC<sub>50</sub> determination). For **6a–c**, the presence of an alkyl side chain apparently promoted antioxidant activity (Gallardo et al. 2016), while the existence of an acyl group on the side chain notably diminished it.



**Table 4** DPPH<sup>•</sup> and ABTS<sup>•+</sup> cation radical scavenging activity of **5a–c**, **6a–c**, **7a**, **7c**, and **9a–d**<sup>a</sup>

Compound	DPPH Scavenging activity		ABTS Scavenging activity	
	% (10 mM)	(IC <sub>50</sub> mM)	% (10 mM)	(mM Trolox/g)
<b>5a</b>	40.6 ± 0.60		13.40 ± 0.20	0.244 ± 0.002
<b>5b</b>	15.9 ± 0.61		11.68 ± 1.38	0.220 ± 0.014
<b>5c</b>	19.1 ± 0.60		13.23 ± 0.71	0.216 ± 0.007
<b>6a</b>	95.3 ± 1.82	0.241 ± 0.009	98.80 ± 0.01	1.509 ± 0.001
<b>6b</b>	94.2 ± 1.50	0.255 ± 0.016	98.61 ± 0.01	1.424 ± 0.001
<b>6c</b>	94.2 ± 1.10	0.244 ± 0.012	98.62 ± 2.39	1.349 ± 0.001
<b>7a</b>	0.2 ± 0.32		0.91 ± 0.37	0.050 ± 0.004
<b>7c</b>	0.9 ± 0.24		0.25 ± 0.48	0.058 ± 0.005
<b>9a</b>	93.5 ± 1.94	0.127 ± 0.030	97.81 ± 3.94	1.552 ± 0.039
<b>9b</b>	93.6 ± 2.49	1.519 ± 0.023	97.53 ± 0.01	1.095 ± 0.001
<b>9c</b>	94.3 ± 1.16	0.238 ± 0.022	98.78 ± 0.01	1.244 ± 0.001
<b>9d</b>	91.0 ± 1.93	1.52 ± 0.015	68.07 ± 0.044	1.00 ± 0.021
<b>BHT</b>	85.02 ± 3.33	0.84 ± 0.08	–	–

<sup>a</sup>The DPPH scavenging activity is calculated as the IC<sub>50</sub>, and the ABTS scavenging activity as the mM Trolox equivalent (TEAC, mM). Data are expressed as the mean ± standard deviation (*n* = 6), with significant differences considered at *p* < 0.05

This characteristic was not affected by the side chain length. Regarding the 1,2-dihydroquinolines **9a–d**, the presence of a hindered group, such as the phenyl (**9b**) or the cycloalkyl (**9d**) groups, played a negative role, slightly reducing the DPPH<sup>•</sup> scavenging activity (IC<sub>50</sub> for both **9b** and **9d** was 1.52 mM). Contrarily, the alkyl (**9a**) and methoxycarbonyl (**9c**) groups improved the antioxidant effect, being almost sevenfold greater with **9a** (IC<sub>50</sub> = 0.12 mM) than BHT (IC<sub>50</sub> = 0.84 mM). The significant activity of **9a** can be attributed to the lipophilicity of the 2-ethyl and 2-methyl groups (Dorey et al. 2000). Interestingly, the methoxy group attached at either the C-6 or C-7 position of the quinoline ring did not substantially affect the DPPH<sup>•</sup> radical scavenging capacity, although its presence seems to have contributed considerably to the antioxidant effect (Dorey et al. 2000; Błaszczuk and Skolimowski 2007; Błaszczuk et al. 2013; de Koning 2002).

Series **6a–c** and **9a–d** also displayed good ABTS<sup>•+</sup> radical inhibition (Table 4) with **6a** and **9a** giving the best ABTS<sup>•+</sup> scavenging capacity (1.51 and 1.55 mM of TEAC). There was a similar pattern for the DPPH<sup>•</sup> radical scavenging activity, finding the best results for **9a** and no effect for compounds **5a–c**, **7a** and **7c**. In summary, the phenoxyacetic ester and 1,2-dihydroquinoline scaffolds demonstrated a significant antioxidant activity, which was improved by the presence of an alkyl side chain or a polar carboxylate group.

### Docking of analogs **5**, **6**, and **9** on human HMGR

The in vitro inhibition of HMGRh has been demonstrated for **3** and **4**, showing a significant structure-binding contact similarities with respect to simvastatin (Mendieta et al.

2014). Hence, a docking study was carried out for the most active hypolipidemic derivatives (**5b**, **6a**, **9a**, and **9d**) of the four series herein evaluated.

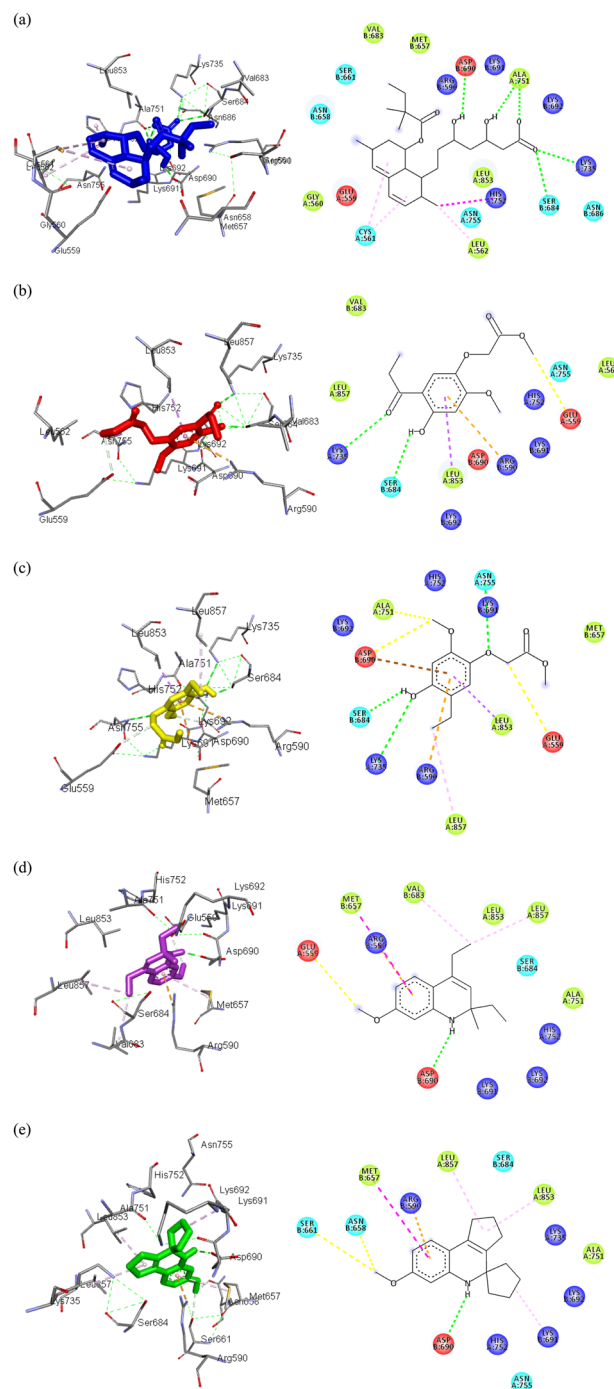
Docking studies were conducted on Autodock version 4.0 and AutoDockTools (Morris et al. 2009) to explore the binding mode of the test compounds and the reference drug (simvastatin) at the active site of HMGRh (retrieved from the PDB; code: 1HW9) (Istvan and Deisenhofer 2001). The binding energy was calculated and the interaction residues identified in each case (Table 5).

The binding modes of **5b**, **6a**, **9a**, **9d**, and simvastatin (Fig. 4) involve the amino acid side chains of the active site of the enzyme, including Glu559, Arg590, Asp690, Lys691, and Asn755. In a previous study by our group (Mendieta et al. 2014), these residues were identified and Glu559 and Lys691 were found to be the key residues (Andrade-Pavón et al. 2017; Andrade-Pavón et al. 2019). The acylphenoxyacetic ester **5b** had a better binding energy (−7.47 kcal/mol) than the alkylphenoxyacetic ester **6a** (−5.93 kcal/mol). The docking data correlate with the results of the in vivo assessment of their hypolipidemic activity, since **5a–c** decreased the levels of cholesterol and triglycerides to a greater extent than **6a–c**.

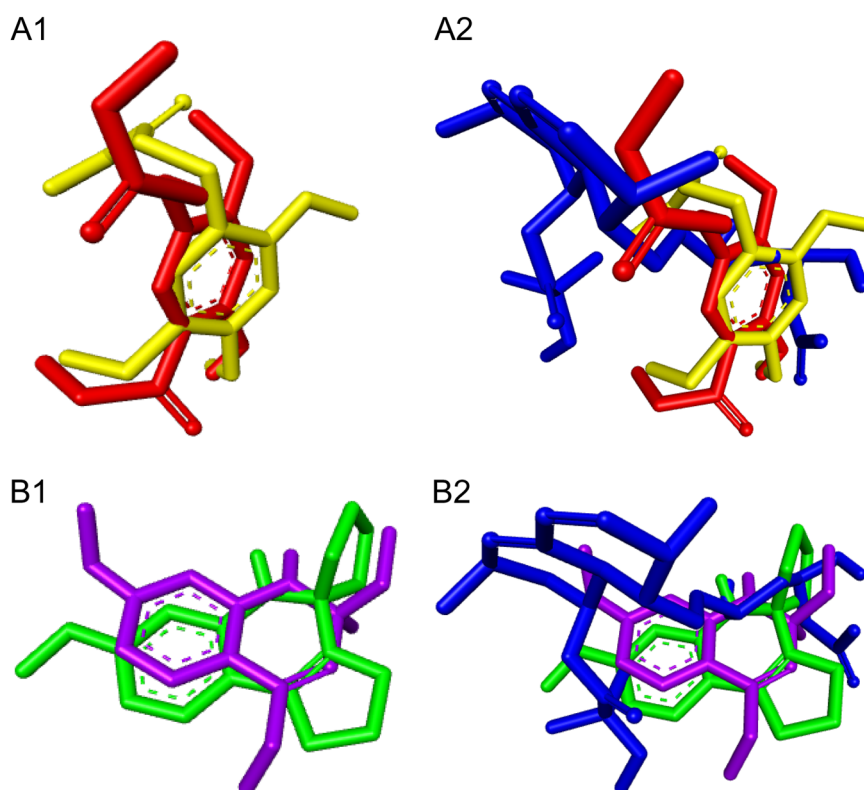
Compounds **5b**, **6a**, and simvastatin share key hydrophilic interactions, such as conventional hydrogen and carbon-hydrogen bonding to the active site of the HMGRh enzyme, mainly with amino acid side chains Glu559, Ser684, Asp690, and Lys735. There are some key similarities in the interactions. For instance, the hydroxyl group at C-4 of the aromatic ring of both **5b** and **6a** interact with the same residue (Ser684) as the carboxylate group of simvastatin. Moreover, there is an interaction with Lys735 by

**Table 5** Docking results for **5b**, **6a**, **9a**, **9d**, and simvastatin at the active site of HMGRh

Compound	Binding energy $\Delta G$ (kcal/mol)	Interacting residues	Polar interactions	Hydrophobic interactions
Simvastatin	-8.45	Glu559, Gly560, Cys561, Leu562, Arg590, Met657, Asn658, Ser661, Ser684, Asn686, Asp690, Lys691, Lys692, Lys735, Ala751, His752, Asn755, Leu853, Leu857	O-H $\cdots$ O (Ser684) O-H $\cdots$ O (Asp690) O-H $\cdots$ O (Ala751) O $\cdots$ H-N (Lys735) O-H $\cdots$ O (Ser684) O $\cdots$ H-N (Lys735) C-H $\cdots$ O (Glu559) O-H $\cdots$ O (Ser684) O $\cdots$ H-N (Lys735) O $\cdots$ H-N (Asn755) C-H $\cdots$ O (Glu559) C-H $\cdots$ O (Asp690) C-H $\cdots$ O (Ala751) N-H $\cdots$ O (Asp690) C-H $\cdots$ O (Glu559)	Alkyl (Cys561, Leu562) $\pi$ -alkyl (His752) $\pi$ -cation (Arg590) $\pi$ -sigma (Leu853) Alkyl (Leu857) $\pi$ -cation (Arg590) $\pi$ -sigma (Leu853) $\pi$ -anion (Asp690) Alkyl (Val683, Leu857) $\pi$ -alkyl (Met657) Alkyl (Lys691, Leu853, Leu857) $\pi$ -cation (Arg590) $\pi$ -alkyl (Met657)
<b>5b</b>	-7.47	Glu559, Leu562, Arg590, Val683, Ser684, Asp690, Lys691, Lys692, Lys735, His752, Asn755, Leu853, Leu857	O-H $\cdots$ O (Ser684) O $\cdots$ H-N (Lys735) C-H $\cdots$ O (Glu559)	Alkyl (Leu857) $\pi$ -cation (Arg590) $\pi$ -sigma (Leu853) $\pi$ -anion (Asp690)
<b>6a</b>	-5.93	Glu559, Arg590, Met657, Ser684, Asp690, Lys691, Lys692, Lys735, Ala751, His752, Asn755, Leu853, Leu857	O-H $\cdots$ O (Ser684) O $\cdots$ H-N (Lys735) C-H $\cdots$ O (Glu559)	Alkyl (Leu857) $\pi$ -cation (Arg590) $\pi$ -sigma (Leu853) $\pi$ -anion (Asp690)
<b>9a</b>	-8.22	Glu559, Arg590, Met657, Val683, Ser684, Asp690, Lys691, Lys692, Lys735, Ala751, His752, Leu853, Leu857	O-H $\cdots$ O (Ser684) O $\cdots$ H-N (Lys735) C-H $\cdots$ O (Glu559)	Alkyl (Val683, Leu857) $\pi$ -cation (Arg590) $\pi$ -alkyl (Met657)
<b>9d</b>	-8.21	Arg590, Met657, Asn658, Ser661, Ser684, Asp690, Lys691, Lys692, Lys735, Ala751, His752, Asn755, Leu853, Leu857	N-H $\cdots$ O (Asp690) C-H $\cdots$ O (Asp658) C-H $\cdots$ O (Asp661)	Alkyl (Lys691, Leu853, Leu857) $\pi$ -cation (Arg590) $\pi$ -alkyl (Met657)

**Fig. 4** Predicted binding mode of simvastatin (**a**), **5b** (**b**), **6a** (**c**), **9a** (**d**), and **9d** (**e**) at the active site of HMGRh (1HW9) using Autodock version 4.0 and AutoDockTools. The 3D model portrays select amino acid residues bound by the ligands at the active site of HMGRh. Only the hydrophilic bonds are shown for better clarity. In the 2D model, the following interactions are denoted with dotted lines: conventional hydrogen bonds (green), carbon-hydrogen (yellow), alkyl (light pink),  $\pi$ -anion (brown),  $\pi$ -cation (orange),  $\pi$ -sigma (purple), and  $\pi$ -alkyl (dark pink). The amino acid residues are illustrated as: hydrophobic (green), polar (cyan), positively charged (blue), and negatively charged (red)

**Fig. 5** A1 Overlay of the docking poses of **5b** (red) and **6a** (yellow). B1 Overlay of **9a** (purple) and **9d** (green). A2–B2 The binding mode is compared between the two pairs of compounds and simvastatin (blue)



the hydroxyl group at the C-4 position of **6a**, the carbonyl groups of **5b** and simvastatin. The three compounds also have comparable hydrophobic interactions, such as  $\pi$ -sigma with the amino acid Leu853. Due to their strongly activated aromatic ring, compounds **5b** and **6a** share electrostatic interactions of the  $\pi$ -cation type with the Arg590 residue. The carbonyl oxygen atom of **5b** forms an additional hydrogen bond contact with the side chain of Lys735. This residue may affect the enzymatic activity of **6a** by interacting with the alkyl side chain. The latter observations are in agreement with the *in vivo* assessment of hypolipidemic activity, which was improved by the polar-induced effect of the carbonyl group in the chain.

The results support our hypothesis that by replacing the C-4 methoxy group of **1** by a hydroxyl group, and maintaining a hydrogen bond network comparable to the one existing in **3** and **4**, the hypolipidemic effect of **5b** and **6a** should be conserved.

For 1,2-dihydroquinolines **9a** and **9d**, on the other hand, a very similar binding energy was found (−8.22 and 8.21 kcal/mol, respectively) (Table 5). Furthermore, **9a** had the binding energy closest to the value for simvastatin. It is likely that the hydrocarbon portion present in both **9a**, **9d** and the reference drug enhances the number of the interactions within the active site as well as the overall stability (Figs 4 and 5). In addition, **9a** and **9d** share hydrophilic interactions between their polar NH and the Asp690 residue

of the enzyme, involving conventional hydrogen bonds and carbon-hydrogen bonds. These interactions are comparable to those observed for the acylphenoxyacetic esters **5**, the alkylphenoxyacetic esters **6**, and the hydroxy group of simvastatin. The presence of a high electron density aromatic ring in **9a** and **9d** led to  $\pi$ -cation interactions with Arg590, and to  $\pi$ -alkyl and alkyl interactions with the side chains of Met657 and Leu857, respectively.

Compounds **5b** and **6a** adopt similar orientation at the active site of the enzyme, especially for the activated aromatic ring (Fig. 5). The orientation of 1,2-dihydroquinolines **9a** and **9d** is also alike, mainly due to the benzoheterocyclic frame. They adopt a conformation in which the polar and the hydrophobic functional groups occupy a position similar to some of the groups in simvastatin with comparable polarity, although the structure of the latter is different. Despite this difference, compounds **5b**, **6a**, **9a**, and **9d** interact with most amino acids in the active site of the enzyme that are targeted by the reference drug. Based on the docking data, **5a–c**, **6a–c**, and **9a–d** are likely to have the same mechanism of action as simvastatin.

To the best of our knowledge, there are no reports of molecular docking of 1,2-dihydroquinolines like **9a** and **9d** in the active site of the HMGRh enzyme. The interaction energy calculated from docking was correlated with the *in vivo* hypolipidemic effect.

## Conclusions

Most members of the three series of acyl phenols (**5a–c**) and alkyl phenols (**6a–c**) and 1,2-dihydroquinolines (**9a–d**) showed a potent hypolipidemic effect, resulting in an over 67% decrease in serum cholesterol and triglycerides at the lowest doses (25 g/kg). Using the most effective derivatives (**5b**, **6a**, **9a**, and **9d**), docking studies were carried out on human HMGR finding that the different polar and non-polar functional groups of the ligands exhibited strong and multiple interactions with the active site of this enzyme. Hence, the mechanism of action of these compounds probably involves the inhibition of HMGR. However, due to the structural similarity of analogs **5** and **6** with fibrates, the activation of PPAR $\alpha$  cannot be ruled out as a competitive mechanism participating in a reduction of the level of triglycerides.

Regarding antifungal potential, the same three series caused a moderate to robust growth inhibition of *C. albicans*, *R. oryzae*, *T. cutaneum* and *F. oxysporum*, which are highly pathogenic fungi. For the latter fungus, derivatives **6a–c** and **9a** and **9c** proved to be more active than the positive control (captan). On the other hand, derivative **6a** produced cytotoxicity not only in all the cancer cell lines (MDA-231, DU-145 and HeLa) but also in normal cells (HaCaT), indicating a lack of selectivity. Finally, both the **6a–c** and **9a–d** series demonstrated good antioxidant capacity, evidenced by the strong DPPH and ABTS scavenging activity. Indeed, **6a–c** and **9a** and **9c** were more active than the positive control (BHT).

Overall, the phenoxyacetic acid esters derived from vanillin and 1,2-dihydroquinolines showed promise as frames for novel hypolipidemic, antifungal, anticancer and antioxidant agents. Thus, the corresponding structures may be advantageous as templates for the design of new compounds with more potent pharmacological activity.

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## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest.

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