

Article

# Spectroscopic Evidence, Evaluation of Biological Activity and Prediction of the Safety Profile of Fatty Hydroxamic Acids Derived from Olive Oil Triacylglycerides

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**Abstract:** A fatty hydroxamic acid (FHA) mixture synthesized from olive oil triacylglycerides by hydroxylaminolysis and composed predominantly of oleyl and linoleyl hydroxamic acid (OHA and LHA, respectively) was characterized by means of IR, Raman, MS and 1D and 2D <sup>1</sup>H- and <sup>13</sup>C-NMR spectroscopy. The ratio of OHA and LHA (4:1) was confirmed by MALDI TOF/TOF mass spectrometry. The radical scavenging and the Fe<sup>2+</sup>-chelating activity, as well as antioxidant activity in β-carotene-linoleic acid and the Fe<sup>3+</sup>-reducing power assays of the FHAs yielded positive results. The results of FHA cell toxicity on normal fibroblast (BJ) and a tumour cell line (HeLa) revealed that the normal cell line was sensitive to FHAs within the entire range of applied concentrations (5 × 10<sup>-4</sup> mg/mL to 5 × 10<sup>-1</sup> mg/mL), while the HeLa tumour cell line was sensitive only at the highest FHA concentration (5 × 10<sup>-1</sup> mg/mL). *In silico* target prediction indicated cannabinoid receptors 1 and 2, the fatty-acid amide hydrolase 1 and histone deacetylases as the most probable targets of OHA and LHA. According to ADMET predictor analysis, the safety profiles of OHA and LHA are comparable to that of SAHA (vorinostat) the histone deacetylase inhibitor in use as an antineoplastic and immunomodulating agent.

**Keywords:** fatty hydroxamic acids; synthesis; spectroscopy; antioxidant activity; cytotoxic activity; biological targets; ADMET; safety profile

## 1. Introduction

Over the last decades there has been a rapid increase in literature dealing with the development of hydroxamic acid-based derivatives and their potential therapeutic applications as antibacterial [1-2], antifungal [3-5], antitumor [6-11], antihypertensive and antiinflammatory drugs [12]. Several hydroxamic acids (HAs) are important therapeutic agents in diverse pharmacological groups such as nonsteroidal antiinflammatory and antirheumatic drugs (*ibuproxam*, *oxametacib* and *bufexamac*), psychoanaleptics (*adrafinil*), antineoplastic and immunomodulating agents (*hydroxycarbamide* or *hydroxyurea*, *vorinostat*, *panobinostat* and *belinostat*), and iron chelating agents which are used to remove excess iron from the body in anemia and thalassemia (*deferroxamine*) [13, 14]. Recently, HAs have attracted a lot of attention as their inhibitory effect on histone deacetylases (HDACs) and the associated applicability in tumour treatment were observed [8, 10, 15-17]. Although many hydroxamic acid-based HDAC inhibitors [18] are currently in discovery and undergoing preclinical phases, the number of HDACIs that have been approved for the market still remains low.

Among the abundant available data on HAs, there is considerably less evidence about biological activity and the potential pharmacotherapeutic usefulness of monounsaturated and polyunsaturated fatty hydroxamic acids (MUFHAs and PUFHAs, respectively) that can also be of pharmaceutical and biological interest [19, 20]. FHAs can be prepared by the usual methods used to prepare HAs, *i.e.*, from fatty carboxylic acid ester or carboxylic acid chloride with either hydroxylamine or its *O*- and *N*-substituted derivatives [14, 21]. For instance, LHA was synthesized from linoleic acid (LA) *via* its reactive intermediates and hydroxylamine. The biological activity evaluation of LHA revealed that it possesses an inhibitory activity against several enzymes involved in the arachidonic acid metabolism relevant for lipid signalling in mammals, *e.g.*, various lipoxygenases and iron-containing enzymes involved in the oxygenation of polyunsaturated fatty acids (PUFAs) [22-24]. The other synthetic possibilities included convenient hydroxylaminolysis of different vegetable oils, *e.g.*, *Jatropha Curcas* seed oil [25], canola, palm, and palm kernel oil [26--28] as well as palm olein, palm stearin, or corn oil [20]. However, owing to the diversity of fatty acids present in different vegetable oil triacylglycerides, a mixture of FHAs has been the resulting product of this type of reaction, which is difficult to separate due to the physicochemical similarities of its components. Therefore, FHAs prepared from different vegetable oils were tested as a mixture for their antimicrobial activity against *E. coli* and *S. aureus* [29-31], antifungal activity [5], and antioxidant activity [32].

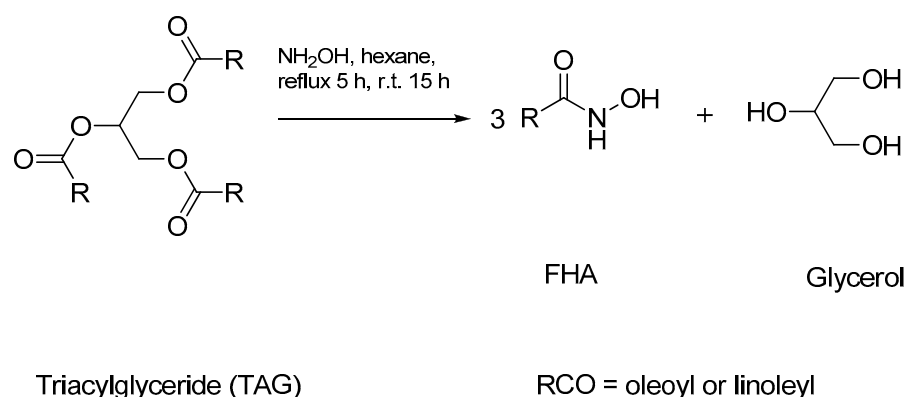
The biological activity of hydroxamic acids, as well as their toxic potential is based on hydroxamic acid moiety (R-CONHOH). Metabolites such as acyl nitroso, HNO, nitric oxide, and metal complexes of the parent drug are designated as the main actors in the physiological effects [33]. The carcinogenicity and mutagenicity of HAs following toxification either *via* Lossen rearrangement to the corresponding reactive isocyanates or *via* *O*-sulfonation or *O*-acetylation of hydroxyl moiety to the corresponding esters of great reactivity in aqueous media have also been reported [34, 35].

Within the framework of our research on the synthesis and biological activity testing of diverse derivatives bearing the HA moiety [7, 36, 37] as well as the content determination of olive oil from autochthonous Croatian olive cultivars [38], the aim of this study was to evaluate the antioxidant and cytotoxic activity of the FHA mixture and explore the safety profile of its main components, *i.e.*, oleyl and linoleyl hydroxamic acid (OHA and LHA, respectively) *in silico*.

## 2. Results and Discussion

### 2.1. Synthesis and spectroscopic evidence

The fatty hydroxamic acid mixture (FHAs) used in this study was synthesized from olive oil triacylglycerides (TAG) by hydroxylaminolysis according to a previously published procedure (Scheme 1) [39].



**Scheme 1.** The synthesis of fatty hydroxamic acids (FHAs).

The structural analysis of the obtained FHA mixture was carried out by FTIR, Raman, one- and two-dimensional, homo- and hetero-nuclear  $^1\text{H}$ - and  $^{13}\text{C}$ -NMR spectroscopy, and by MALDI-TOF-TOF mass spectrometry. Here we report the main findings.

Since oleic acid (OA, 18:1n-9) is the most abundant monounsaturated fatty acid (MUFA) and linoleic acid (LA, 18:2n-6) the most abundant polyunsaturated fatty acid (PUFA) in olive oil triacylglycerides (TAG), where their concentrations in the total content of fatty acids (FAs) range from 55% to 83% for OA and from 3.5% to 21.0% for LA [40], their corresponding hydroxamic acids were expected to be present in the reaction product, *i.e.*, FHAs. And the question that has arisen is whether there are certain discernible amounts of LHA and/or other PUFHAs in addition to OHA in the synthesized FHAs mixture that may be of relevance for biological effects because small amounts of other PUFAs may also be present in olive oil TAGs, *e.g.*, linolenic acid (LNA, 18:3n-6) in 0.4% to 1.9%, while eicosanoic acids are present in trace amounts. As it is the case with the separation of MUFAa and PUFAs due to similarities in their physicochemical properties [41], we encountered such difficulty with our FHA mixture and therefore continued to study FHAs with different spectroscopic methods in order to elucidate its main components.

#### 2.1.1. FTIR and Raman spectroscopy

The FTIR spectra of precursor olive oil and FHAs are presented in Figure S1 and the most significant bands of FHAs in Table S1 in the supplementary material. The FTIR results provided clear and convincing evidence regarding the chemical structure of FHAs, mainly from amide type carbonyl and OH/NH stretching bands. The broad band from  $\nu$  3650  $\text{cm}^{-1}$  to 2500  $\text{cm}^{-1}$  corresponds

to -CONHOH moiety, and the peak at  $\nu$  3285  $\text{cm}^{-1}$  was assigned to N-H stretching vibrations. The O-H moiety in -CONHOH group was most likely associated *via* inter- and/or intra-molecular hydrogen bonds and due to that produced a very broad vibration that overlapped with the stretching of the methyl and methylene groups. In addition to these findings, in the FTIR spectrum of the FHAs there were two stretching vibrations for the C=O group at  $\nu$  1664  $\text{cm}^{-1}$  and 1624  $\text{cm}^{-1}$  that belonged to amide I and amide II vibrations, respectively. The weak band at  $\nu$  3001  $\text{cm}^{-1}$  originated from =C-H stretching, thus indicating the presence of one or more unsaturated moieties in the alkyl chain. Vibration bands of the methyl group appeared at 2952  $\text{cm}^{-1}$  and 2872  $\text{cm}^{-1}$ , while stretching bands of methylene groups appeared at 2918  $\text{cm}^{-1}$  and 2849  $\text{cm}^{-1}$ . Strong and medium bands at 1464  $\text{cm}^{-1}$  and 1441  $\text{cm}^{-1}$ , respectively, were attributed to the deformation of C-H groups, while the medium strong band at 1096  $\text{cm}^{-1}$ , may have been due to skeletal vibrations, *e.g.*, the stretching of -C-C- bonds.

The most intense bands in the Raman spectrum of the FHAs (Table S1 in supplementary material) have to do with the stretching of the CH<sub>2</sub> groups,  $\nu_{\text{as}}$  2932  $\text{cm}^{-1}$  and  $\nu_{\text{sym}}$  2847  $\text{cm}^{-1}$ . The C=O stretching produced two bands that arose due to the stretching of the C=O group of amide moiety in the Raman spectrum of the FHAs, the medium one at 1659  $\text{cm}^{-1}$  and one very weak band at 1622  $\text{cm}^{-1}$ . The bending of the C-H bond in the methylene group appeared at 1441  $\text{cm}^{-1}$  as a very strong and broad band. The strong band at 1295  $\text{cm}^{-1}$  most likely arose due to C-N stretching. The stretching of the -C-C- group of medium intensity appeared at 1095  $\text{cm}^{-1}$  and 1062  $\text{cm}^{-1}$ . Bands at lower wavenumbers were either weak or very weak. FTIR and Raman spectra of the FHAs may be useful for the assessment of unsaturated moieties from which the FHAs mixture was composed. Band positions and/or band shapes attributed to the stretching and/or bending modes that include an H-atom imply the presence of hydrogen bonds.

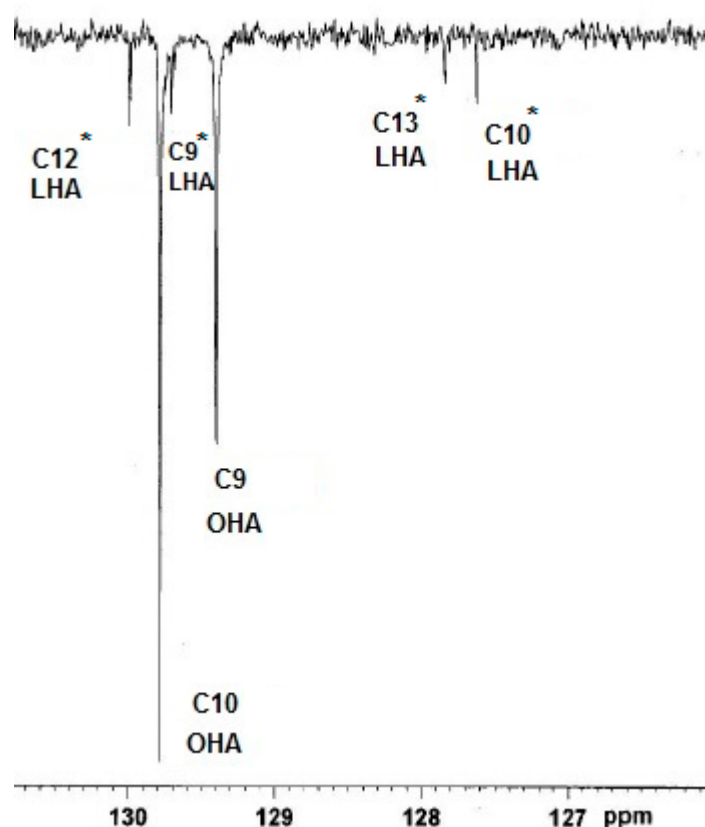
### 2.1.2. NMR spectroscopy

The <sup>1</sup>H-NMR spectrum of the FHA mixture (in CDCl<sub>3</sub>) and chemical structures of its main components, *i.e.*, OHA and LHA, with labelled proton atoms are displayed in Figure S2 and Table S2 in the supplementary material. Analogous protons in LHA are labelled with an asterisk (\*) (Figure S2, Table S2). The proton assignments were made on the basis of observed chemical shifts ( $\delta$  ppm) in the 1D <sup>1</sup>H-NMR spectrum and 2D HMQC (Heteronuclear Multiple Quantum Coherence), 2D HMBC (Heteronuclear Multiple Bond Coherence) and <sup>1</sup>H-<sup>1</sup>H COSY (Correlated spectroscopy) experiments (Figure S2, Figure S4, Figure S5 and Figure S6, respectively), as well. The observed chemical shifts were in good agreement with the available literature data [42, 25]. The <sup>1</sup>H-NMR spectrum of the FHAs was characterized by eight signals that belong to eight chemically and magnetically nonequivalent protons. Protons of the -CO-NHOH group were observed as a broad and very weak signal at  $\delta$  8.674 ppm, while the multiplet signal at  $\delta$  5.339 ppm was assigned to olefinic protons (-HC=CH-) in OHA (H9, H10) and LHA (H9\*, H10\*, H12\* and H13\*). The signal at  $\delta$  0.880 ppm (t) was assigned to protons in the methyl groups, *i.e.*, H-18 and H-18\*. Methylene protons (-CH<sub>2</sub>-) in the OHA (H2, H3, H4 to H7, H8, H11, H12 to H17) and in LHA (H2\*, H3\*, H4\* to H7\*, H8\*, H11\*, H15\* to H17) were observed from  $\delta$  1.254 to 2.777 ppm depending on the position of each methylene group in the alkyl chains of the FHAs. Thus methylene protons H4 to H7, H12 to H17, H4\* to H7\*, H15\* to H17\* were observed at  $\delta$  1.250 to 1.290 ppm (m) while methylene protons

H8, H11, H8\* and H14\*, adjacent to the olefinic bond, *i.e.*,  $-\underline{\text{CH}_2}-\text{CH}=\text{CH}-$  were observed as multiplet (m) at  $\delta$  1.988 to 2.040 ppm. Methylene protons H2 and H2\*, in the immediate vicinity to the  $-\text{CO}-\text{NHOH}$  group were observed downfield as a triplet (t) at  $\delta$  2.135 ppm and methylene protons H3 and H3\* at 1.621 ppm (m). However, a triplet (t) signal of very low intensity at  $\delta$  2.766 ppm indicated the presence of LHA and this signal corresponds to methylene protons H11\*. Although this signal may come from LHA ( $-\text{CH}=\text{CH}-\underline{\text{CH}_2}-\text{CH}=\text{CH}-$ ) and/or LNHA ( $-\text{CH}=\text{CH}-\underline{\text{CH}_2}-\text{CH}=\text{CH}-\underline{\text{CH}_2}-\text{CH}=\text{CH}-$ ) and these signals may be overlap to some extent, the assignment was made in favour of LHA methylene protons H11\* because of the lack of an additional characteristic signal of methyl protons ( $-\underline{\text{CH}_3}$ ) of LNA (n-3) which is usually separated from  $-\underline{\text{CH}_3}$  of n-9 and n-6 FAs and was found to be at  $\delta$  0.970 ppm compared to oleoyl and linoleyl methyl protons observed at  $\delta$  0.880 ppm [43, 44].

On the basis of these findings, it was concluded that the obtained FHAs mixture was either free of LNHA or its minor presence was below the discernible amount relevant to the  $^1\text{H}$ NMR spectroscopy.

The experimental  $^{13}\text{C}$ -NMR data of FHAs recorded in deuterated chloroform ( $\text{CDCl}_3$ ) (Figure S3 in supplementary material) and theoretical data computed for OHA and LHA using nmrshiftdb (<http://nmrshiftdb.nmr.uni-koeln.de>) and ChemDraw Ultra v. 11.0  $^{13}\text{C}$  NMR software packages, together with available literature data for  $^{13}\text{C}$ -NMR of LHA are presented in Table S2 in the supplementary material. Analysis of the  $^{13}\text{C}$ -NMR spectrum of FHAs confirmed the presence of signals characteristic for 18:1 and 18:2 FHAs (Figure 1). The distinction and assignments of carbon atoms that belong to OHA from those of LHA were made on the basis of characteristic chemical shifts ( $\delta$  ppm), signal intensities, and by comparing experimental data with the computed theoretical spectra of OHA and LHA, as well as with the available literature data for  $^{13}\text{C}$ -NMR of LHA [24]. The carbon atoms involved in the carbonyl ( $\text{C}=\text{O}$ ) and methyl ( $-\text{CH}_3$ ) groups were observed as singlets at  $\delta$  171.73 and 13.99 ppm, respectively, for both, OHA and LHA, while all other carbon atoms were separated with intensities that were in favour of the OHA as a predominant component in the FHA mixture. The observed six signals of olefinic carbon atoms in the region of  $\delta$  127.78 to 130.14 ppm (Table S2, Figure 1) in the  $^{13}\text{C}$ -NMR spectrum of FHAs indicated the presence of six nonequivalent carbon atoms involved in the  $\text{C}=\text{C}$  bonds. Although  $^{13}\text{C}$ -NMR cannot be considered a quantitative technique compared to  $^1\text{H}$ -NMR, the observed signals at distinct chemical shifts with obviously different signal intensities were assigned so that the OHA (C9,  $\delta$  129.56 ppm and C10, 129.94 ppm) was determined as the prevailing component of FHAs with lower proportions of LHA (C9\*, C10\*, C12\* and C13\*,  $\delta$  129.86, 127.78, 130.14 and 127.99, respectively). No signs of inadvertent  $Z \rightarrow E$  isomerization of OHA and LHA during the synthesis were observed.



**Figure 1.** A part of  $^{13}\text{C}$  APT (Attached-Proton-Test) spectrum of FHAs mixture recorded in  $\text{CDCl}_3$ , displaying olefinic carbons of OHA (a pair of larger signals, *i.e.*, C9 129.56 ppm and C10 129.94 ppm) and LHA (two pairs of smaller signals, *i.e.*, C9\* 129.86, C10\* 127.78, C12\* 130.14 and C13\* 127.99 ppm).

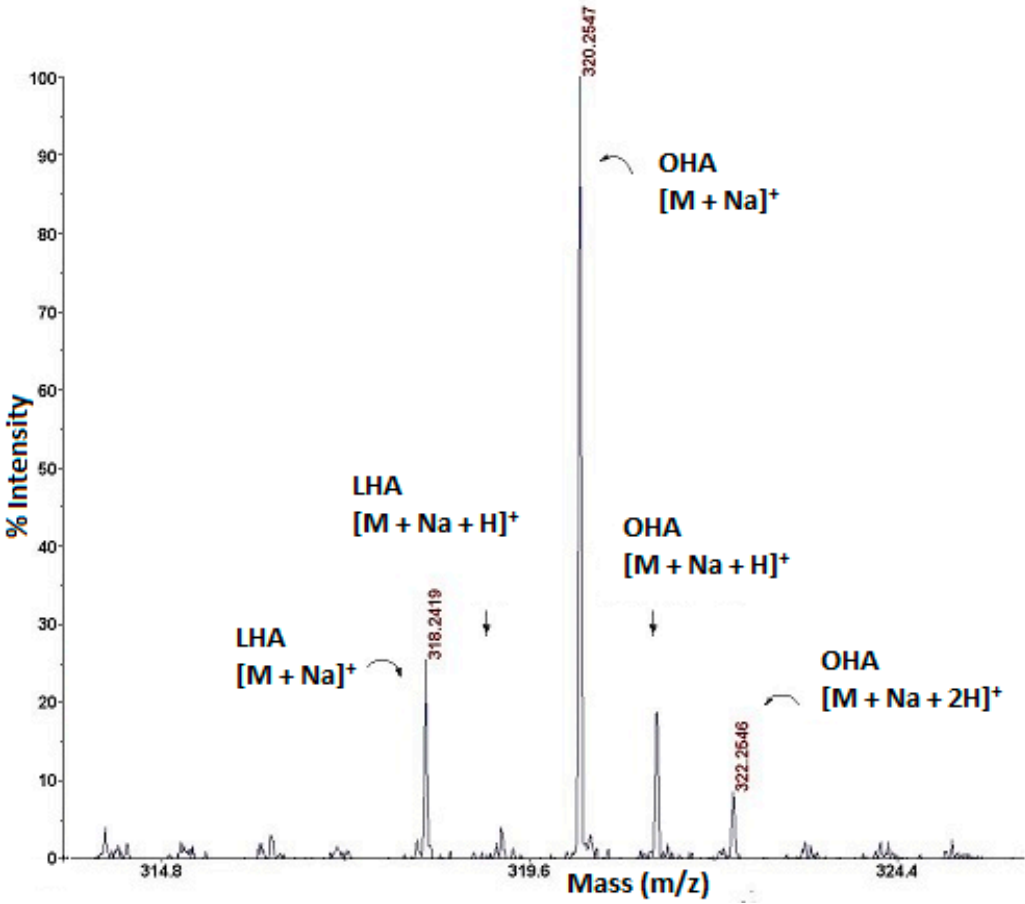
Obviously, NMR techniques alone were not able to provide detailed information about the molar proportion of different fatty hydroxamic acids in the FHA mixture which was also due to the fact that NMR is a less sensitive technique compared to traditional chromatographic methods and mass spectrometry. Therefore, we continued to study the FHA mixture by matrix-assisted laser desorption/ionization (MALDI) time-of-flight/time-of-flight (TOF-TOF) high resolution tandem mass spectrometry (MS).

### 2.1.3. MALDI-TOF/TOF MS spectrometry

In order to obtain a more precise ratio of components in the FHA mixture, MALDI-TOF-TOF MS was applied. A part of the MALDI-TOF/TOF tandem mass spectrum of synthesized FHAs in the region of mono- and polyunsaturated FHAs, *i.e.*,  $m/z$  310 – 325, is presented in Figure 2. A broader MS spectrum is presented in Figure S7 in the supplementary material, as well. Components of the FHA mixture, *i.e.*, OHA and LHA were observed as singly charged sodium alkali adducts, therefore their molecular ions were shifted by the mass of sodium, *i.e.*, 22.99. Thus the molecular ions of the predominantly present OHA were observed as their sodium adducts  $[\text{M} + \text{Na}]^+$  at  $m/z$  320.25 and  $[\text{M} + \text{Na} + \text{H}]^+$  at  $m/z$  321.25 while molecular ions of LHA, *i.e.*,  $[\text{LHA} + \text{Na}]^+$  at 318.24 and  $[\text{LHA} + \text{Na} + \text{H}]^+$  at 319.24. The ratio of the two main  $m/z$  intensities, *i.e.*,  $I_{m/z} [\text{OHA} + \text{Na}]^+ / I_{m/z} [\text{LHA} + \text{Na}]^+$  was found to be 4:1. On the basis of the results of the MALDI TOF/TOF MS analysis it was



concluded that the amount of LHA in the FHA mixture was 20% and of OHA 80%. The signal at  $m/z$  322.25 was attributed to the  $[OHA + Na + 2H]^+$  however, the possible presence of saturated stearic hydroxamic acid (SHA) due to its presence in olive oil TAG (0.5% to 5.0%) cannot be completely excluded through this finding since its molecular ion sodium adduct  $m/z$   $[SHA + Na]^+$  has the same  $m/z$  as  $[OHA + Na + 2H]^+$ .



**Figure 2.** A part of the MALDI-TOF/TOF tandem mass spectrum of the FHA mixture in the region of  $m/z$  314.0 – 325.0 with molecular ions of OHA and LHA represented as their sodium alkali adducts  $[M + Na]^+$ ,  $[M + Na + H]^+$  and  $[M + Na + 2H]^+$ .

2.2. Biological evaluation in vitro

So far, FHA products prepared from different plant oils have been evaluated for their biological activity mostly without specifying the exact composition of the mixture, and so have asserted that the fatty hydroxamic acid (FHA) from the seed oil of *Cyperus esculentus* showed antioxidant activity tested by DPPH radical scavenging assay as the concentration reduced below 0.05 mg/mL while the antioxidant capacity reduced once the concentration became higher and FHA tended toward being a pro-oxidant. *C. esculentus* has been also presented as a potential source of feed stock for the synthesis of a relatively cheap and non-toxic FHA [32]. Here we report the antioxidant and cytotoxic activity of a FHA mixture obtained from olive oil triacylglycerides with its main components OHA and LHA (80 : 20, ww).

2.2.1. Antioxidant activity

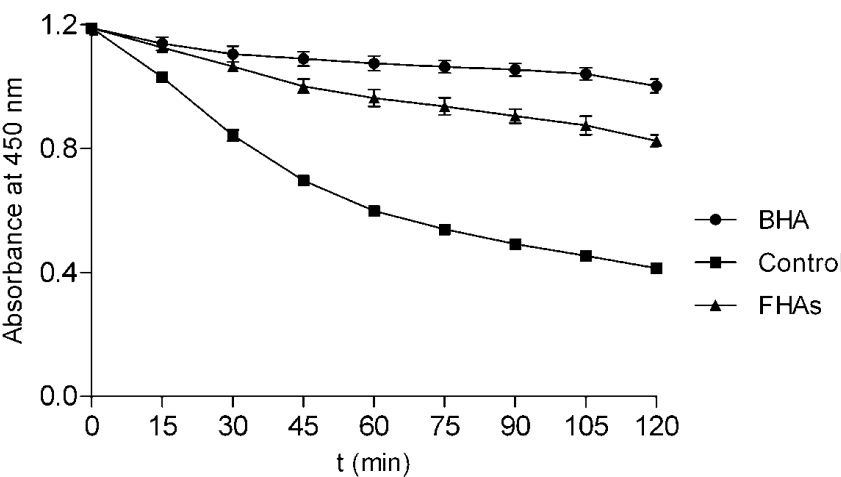
Four different methods for the assessment of FHA antioxidant activity were used in this study: radical scavenging activity (RSA), chelating activity of Fe<sup>2+</sup> ions (ChA), antioxidant activity in β-carotene bleaching assay (AOA), and the reducing power assay (RP).

The capability of FHAs to scavenge free radicals was assessed in reaction with the relatively stable 1,1-diphenyl-2-picrylhydrazyl (DPPH) free radical. FHAs demonstrated notable antiradical activity, albeit lower than butylated hydroxyanisol (BHA), a widely used food antioxidant. The obtained result for FHAs, RSA EC<sub>50</sub> = 235.66 ± 54.01 µg/mL, corresponds to the concentration that scavenges 50% of the DPPH free radicals present in the solution (Table 1).

The chelation of metal ions (ChA) such as copper and especially iron ions by certain compounds decreases their pro-oxidant activity [45]. This aspect of antioxidant activity is particularly important because in the Fenton reaction hydroxyl radical production is directly related to the concentration of the iron ion or other transition ions [46]. The results are expressed as ChA EC<sub>50</sub>, the concentration that chelates 50% of the Fe<sup>2+</sup> ions present in the solution. In this study the chelating activity of FHAs was compared with two chelating standards, ethylenediaminetetraacetic acid (EDTA) and quercetin (Qn) (Table 1). FHAs showed lower activity (ChA EC<sub>50</sub> = 1226.53 ± 58.33 µg/mL) than the strong ion chelator, EDTA (ChA EC<sub>50</sub> = 12.64 ± 2.81 µg/mL). However, the FHA mixture was as active as Qn (EC<sub>50</sub> = 1177.96 ± 175.73 µg/mL), a natural ion chelator, present in many types of fruits and vegetables.

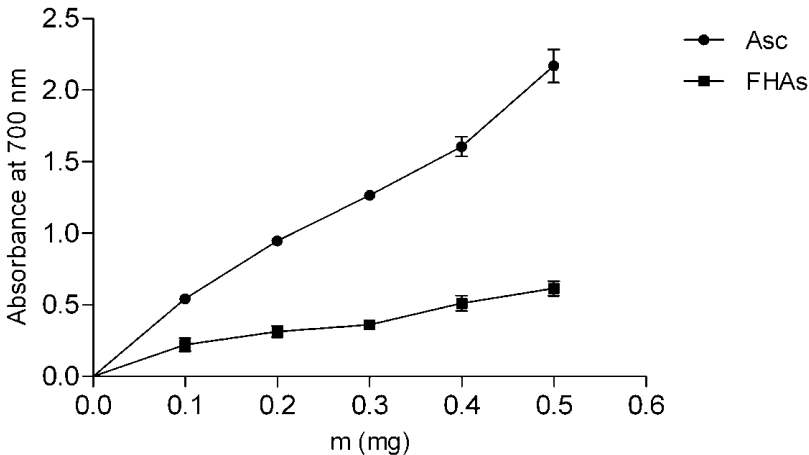
Antioxidant activity in β-carotene linoleic acid assay is frequently employed as a test for measuring total antioxidant activity (AOA) of plant extracts by the oxidation of the aqueous emulsion of β-carotene and linoleic acid [47, 48]. This assay measures the capacity of antioxidants to inhibit the formation of conjugated diene hydroperoxide arising from linoleic acid oxidation. Thus, the assay gives insight into the inhibitory effect of substances on lipid peroxidation [49]. Antioxidant activity is expressed as percentage (%) of inhibition relative to the control and AOA EC<sub>50</sub> (µg/mL) is defined as the concentration of an antioxidant required to inhibit the 50% degradation of beta-carotene relative to the control under the given experimental conditions. The FHA concentration (AOA EC<sub>50</sub> = 55.71 ± 1.29 µg/mL) (Table 1) showed that the FHAs were able to significantly reduce the rate of degradation of β-carotene in comparison with water as control. However, FHAs activity is comparable but somewhat lower than the activity of BHA (AOA EC<sub>50</sub> = 45.61 ± 0.11 µg/mL) (Figure 3, Table 1).





**Figure 3.** Inhibition of  $\beta$ -carotene-linoleic acid emulsion bleaching by the FHAs and BHA. Values are presented as mean  $\pm$  SD (n = 3).

Literature reports suggest that the antioxidant activity of some antioxidants and plant extracts is associated with their reducing power (RP  $EC_{0.5}$ ) which terminates free radical chain reactions [50]. The reducing power of the investigated FHAs mixture increased linearly with the concentration (Figure 4). Coefficients of determination ( $r^2$ ) for ascorbic acid (Asc) and FHAs were 0.9807 and 0.9166, respectively. In this assay, ascorbic acid demonstrated significantly better reducing power (RP  $EC_{0.5}$  =  $97.028 \pm 10.87 \mu\text{g/mL}$ ) than FHAs (RP  $EC_{0.5}$  =  $395.71 \pm 47.37 \mu\text{g/mL}$ ) (Table 1). The RP  $EC_{0.5}$  as a sample concentration providing 0.5 of absorbance ( $EC_{50}$ ) was calculated by plotting absorbance at 700 nm against the corresponding sample concentration.



**Figure 4.** Reducing power activities of FHAs and ascorbic acid (Asc). Values are presented as mean  $\pm$  SD (n = 3).

**Table 1.** The results of DPPH radical scavenging activity (RSA EC<sub>50</sub>), Fe<sup>2+</sup> chelating activity (Ch EC<sub>50</sub>), antioxidant activity in β-carotene-linoleic acid assay (AOA EC<sub>50</sub>) and Fe<sup>3+</sup>-reducing power (RP EC<sub>0.5</sub>) of FHAs composed of OHA and LHA (4 : 1). Values (μg/ml) are presented as means ± SD (n = 3).

Sample	RSA EC <sub>50</sub> (μg/mL)	ChA EC <sub>50</sub> (μg/mL)	AOA EC <sub>50</sub> (μg/mL)	RP EC <sub>0.5</sub> (μg/mL)
FHAs	235.66 ± 54.01 <sup>1*</sup>	1226.53 ± 58.33 <sup>2*</sup>	55.71 ± 1.29 <sup>1*</sup>	395.71 ± 47.37 <sup>3*</sup>
Standard	24.46 ± 2.54 <sup>1</sup>	12.64 ± 2.81 <sup>2</sup> 1177.96 ± 175.73 <sup>4</sup>	45.61 ± 0.11 <sup>1</sup>	97.028 ± 10.87 <sup>3</sup>

<sup>1</sup> Butylated hydroxyanisol

<sup>2</sup> EDTA

<sup>3</sup> Ascorbic acid

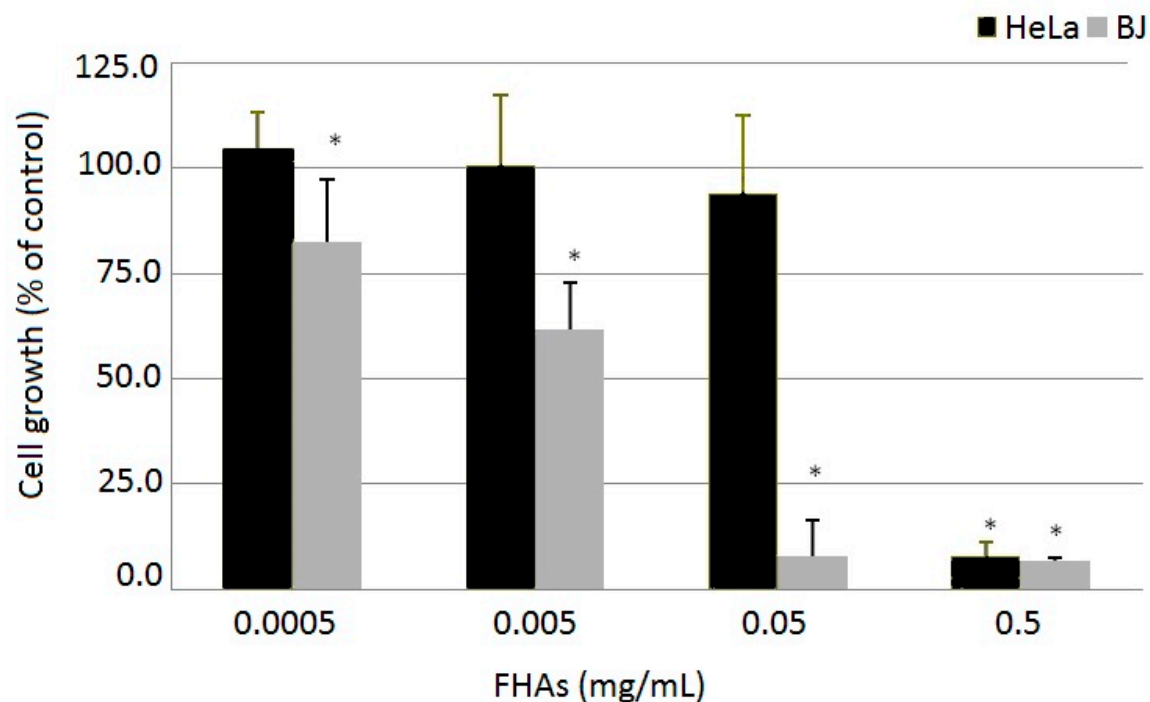
<sup>4</sup> Quercetin

\*Statistical differences with the corresponding standards (*P* < 0.05) within the column.

The results of these studies demonstrated that the FHA mixture possessed moderate antioxidant activity.

2.2.2. Cytotoxic activity

The results of the MTT colorimetric assay revealed that the addition of the FHAs to normal (BJ) and tumour (HeLa) cell lines at a final concentration of 0.5 mg/mL suppressed the cell growth of both of the tested lines by more than 90% (Figure 5). The antiproliferative effect on the normal cell line at 0.05 mg/mL was strong (92% of cell died), while moderate growth suppression (8% of dead cells) was observed in the tumour cell line. The BJ and Hela cell line have different modes of biological response to the FHAs applied at the tested concentration, *i.e.*, the growth of normal cells was strongly suppressed while the tested tumour cell line was not as sensitive. The established IC<sub>50</sub> value for HeLa (80.5 μg/mL) and BJ (7.92 μg/mL) confirmed a ten times higher FHA cytotoxicity for normal cells compared with malignant HeLa cells.



**Figure 5.** The cytotoxic effects of the FHAs. Cell line growth is displayed as percentage (%) after 72 h of incubation at the final concentration range (0.0005 - 0.5 mg/mL) evaluated by MTT colorimetric technique. Data represent the mean value of three independent experiments done in triplicate. Statistically significant change ( $p < 0.05$ ) is presented by an asterisk (\*).

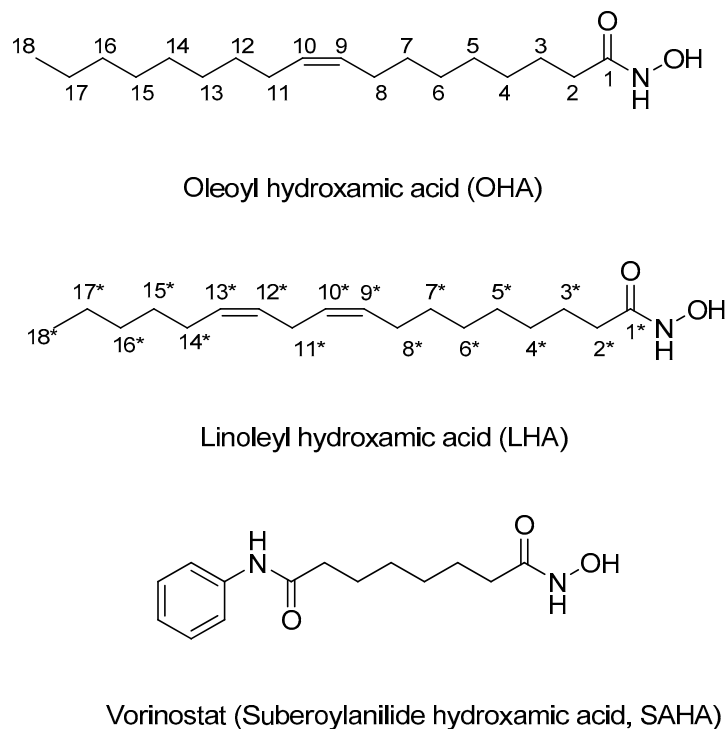
2.3. *In silico* prediction of biological targets and safety profile

2.3.1. Target prediction

Potential biological on- and off-targets of OHA and LHA molecules were evaluated by the SwissTargetPrediction web server that provides a prediction for small molecules using a combination of 2D and 3D similarity measures (<http://www.swisstargetprediction.ch>). It compares the query molecule against a library of 280000 compounds active on more than 2000 targets of 5 different organisms [51, 52]. The outcome of this analysis yielded fifteen of the most probable targets in *Homo sapiens* for each query molecule. Three main target classes were revealed: the 'enzyme other' with an overall frequency of 80% for OHA and 87% for LHA, the 'membrane receptor' class with a frequency of 13% for both OHA and LHA, and the 'ion channel' class with a 7% frequency for OHA. The revealed specific targets and their probabilities are displayed in Table 2.

In addition to histone deacetylases 1 to 9 (HDAC1 – HDAC9) with probabilities (P) from 0.4 to 0.5 for both OHA and LHA, the most probable targets that include cannabinoid receptor 1 (CB1) (P = 0.6 for OHA and 0.8 for LHA) and cannabinoid receptor 2 (CB2) (P = 0.4 for OHA and 0.3 for LHA) as well as fatty-acid amide hydrolase 1 (FAAH) (P = 0.6 for OHA and 0.5 for LHA) were also predicted (Table 2). The prediction for SAHA (suberoylanilide hydroxamic acid, vorinostat) the first

HDAC inhibitor registered by the FDA, revealed that histone deacetylases 1 to 11 (HDAC1 - HDAC11) are the main targets of this molecule with the highest probability ( $P = 1.0$ ). Although not as high, the probabilities of interactions of OHA and FHA with histone deacetylases could be a good starting point for further research regarding the development of more prominent HDAC inhibitor properties for tumour treatment. Chemical structures of oleoyl hydroxamic acid (OHA), linoleyl hydroxamic acid (LHA) and suberoylanilide hydroxamic acid (SAHA) are displayed in Figure 6.



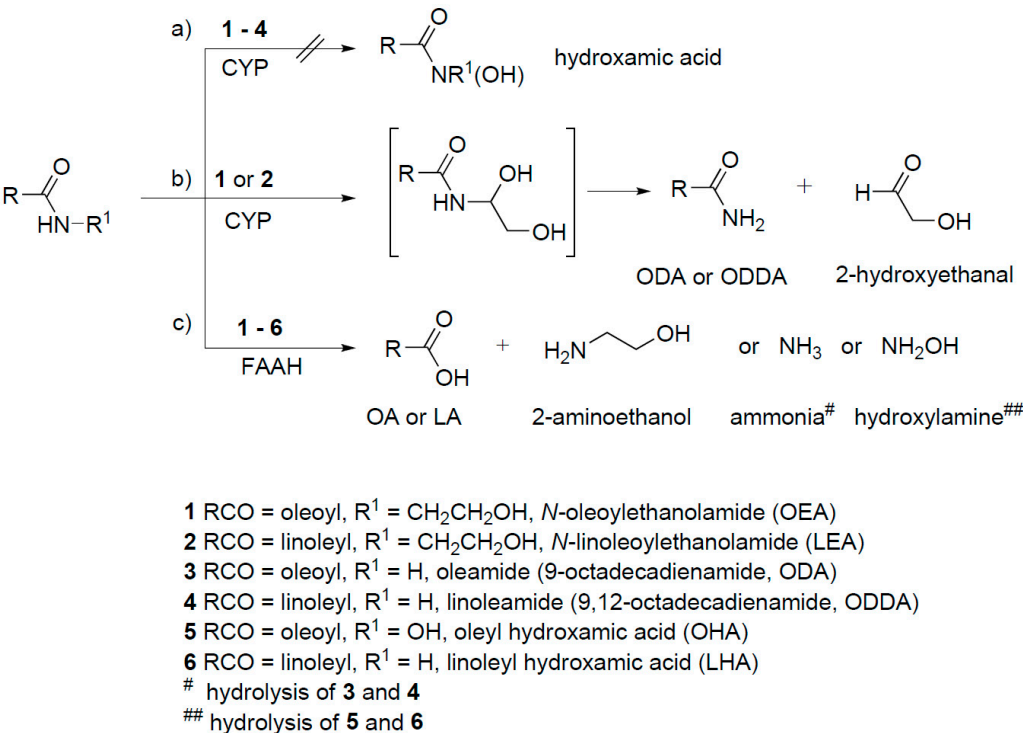
**Figure 6.** Chemical structures of oleoyl hydroxamic acid (OHA), linoleyl hydroxamic acid (LHA) and suberoylanilide hydroxamic acid (SAHA)

The probability of interactions with FAAH in humans assumes the inhibition of this enzyme, which is involved in hydrolytic degradation of endogenous fatty acid amides (FAAs) and among them the signalling endocannabinoid molecule anandamide (*N*-arachidonylethanolamine, AEA) to arachidonic acid and ethanolamine [53, 54], as well as oleamide (*cis*-9,10-octadecanoamide, ODA), the primary amide of oleic acid which is an endogenous sleep-inducing substance, to oleic acid and ammonia [55].

Anandamide was also found in chocolate together with two substances that might mimic the effects of anandamide, *N*-oleylethanolamide (OEA) and *N*-linoleylethanolamide (LEA) [56]. More recently, many new molecules have been investigated as possible FAAH inhibitors [57]. From the predicted interactions of OHA and LHA with CB1 and CB2, as well as with FAAH, it could be assumed that these molecules may also mimic the effects of *N*-oleylethanolamide and *N*-linoleylethanolamide on anandamide, as well as oleamide in the body based on their similarities with these endogenous messengers.

By increasing the anandamide level, its physiological effects either in the central or peripheral nervous system, *via* cannabinoid receptors CB1 or CB2, respectively, will consequently be increased

as will the influence on its many physiological functions to *e.g.*, eating, sleep patterns or pain *via* CB1 as well as immunity, mainly *via* CB2 receptors. The potential metabolic pathways of FAAs normally present in the body (**1** – OEA, **2** – LEA, **3** – OA and **4** – LA) as well as xenobiotic OHA (**5**) and LHA (**6**) are displayed in Scheme 2.



**Scheme 2.** The possible metabolic pathway of endogenous fatty acid amides (FAAs) and FHAs to corresponding metabolites *via*: a) *N*-hydroxylation, b) *N*-dealkylation, and c) hydrolysis.

While *N*-hydroxylation of aromatic amides by CYP enzymes to corresponding hydroxamic acids is common reaction [34], the *N*-hydroxylation of aliphatic amides is not usual therefore the potential formation of OHA and LHA from endogenous amides (**1** – **4**) will not be possible (Scheme 2, pathway a). On the other hand, OA and LA can be formed by CYP enzymes from OEA and LEA, respectively *via* *N*-dealkylation of amides **1** and **2** (Scheme 2, pathway b). All these endogenous and xenobiotic substances (**1** – **6**) bearing the  $-C(=O)NH-$  moiety are subject to hydrolysis by FAAH (Scheme 2, pathway c).

**Table 2.** The biological targets for OHA, LHA, as well as for histone deacetylase inhibitor SAHA (suberoylanilide hydroxamic acid, vorinostat) predicted by SwissTargetPrediction.

Target class	Target	Probability (P)		
		OHA	LHA	SAHA
Membrane receptor	Cannabinoid receptor 1	0.6	0.8	
Membrane receptor	Cannabinoid receptor 2	0.4	0.3	
Enzyme	Fatty-acid amide hydrolase 1	0.6	0.5	
Enzyme	Histone deacetylase 1 to 11	0.4 -0.5 <sup>1</sup>	0.4 - 0.5 <sup>1</sup>	1.0

Enzyme	Corticosteroid 11-beta-dehydrogenase isozyme 1 and 2	0.3	0.3
Enzyme	Hydroxysteroid 11-beta-dehydrogenase 1-like protein		0.3
Ion channel	Transient receptor potential cation channel subfamily V member 1	0.3	
Enzyme/Unclassified	Complex		1.0
Serine protease	Lipid-phosphate phosphatase		0.6
Enzyme	FAD-linked sulfhydryl oxidase ALR		0.6
Metallo protease	22 kDA interstitial collagenase		0.5

<sup>1</sup> Histone deacetylase 1 to 9 (HDAC1 to HDAC9)

<sup>2</sup> Fatty acid-binding protein liver/Unclassified

2.3.2. ADMET properties prediction and safety profile of OHA and LHA

Although many studies have been published on the preparation and biological evaluation of FHAs obtained from plant oils, there is insufficient evidence about their safety profile. Therefore, the results of an oral administration of FHA from *Cyperus esculentus* seed oil in rats receiving a dose of 5 mg/kg/day showed no adverse effects [32]. However, in rats receiving a dose of 15 mg/kg/day (the least observed adverse effect level) a significant increase in alkaline phosphatase activity and triglycerides and creatinine levels were observed together with moderate hyper-albuminemia and hypoalbuminemia that resulted in an increased albumin/globulin ratio. In addition, the study on the aquatic toxicity of the golden algae *Prymnesium parvum*, implicated in fish and aquatic animal deaths globally revealed six fatty acid amides (FAAs) and LHAs as the toxins responsible for the observed aquatotoxicity [58]. The poor pharmacokinetics and toxicity are important causes of costly late-stage failures in drug development and these areas should be considered as early as possible in the drug discovery process. Therefore, early data on absorption, distribution, metabolism, excretion and toxicity (ADMET) are needed. Regarding the aforementioned issues, *in silico* approaches are valuable tools as they increase our ability to predict and model the most relevant pharmacokinetic, metabolic, and toxicity endpoints that accelerate the drug discovery process [59].

In this study, FHAs were screened for undesirable ADMET properties *in silico*, using the liability scoring system known as ADMET Risk™ (ADMET Predictor™ 8.1, Simulations Plus, Inc., USA). Here we report the results of this *in silico* study in which the ADMET properties relevant to the toxic profile of OHA and LHA, the main components of the FHA mixture, were predicted by ADMET Predictor™ [60]. The obtained results were compared with the predicted parameters obtained for their precursor fatty acids (OA and LA). Since HDACs were predicted as potential biological targets of OHA and LHA (with probabilities P = 0.4 – 0.5), SAHA (P = 1.0) was also included *in silico* study. The results are presented in Table 3. In addition to these predicted data, the ADMET Predictor Metabolism Module (Human Cytochrome P450) which contains two models (the classification of compounds as general inhibitors of 5 major CYP isoforms and the qualitative and quantitative inhibition models for CYP3A4 with either midazolam or testosterone as the substrate) predicted the extensive LHA CYP metabolism in comparison to other investigated molecules. Thus



sites of the CYP 1A2 metabolism in LHA were predicted at the following C-atoms with corresponding hydroxylated metabolites: C18\* ( $\omega$ ), C17\* ( $\omega$ -1), C16\* ( $\omega$ -2), C14\*, and C11\* and additionally, a site of CYP3A4 at C8\*. The predicted CYP metabolism of OA and LA by CYP 2E1 and CYP 1A1, respectively, involved the hydroxylation of corresponding C-atoms in each molecule: C18 and C18\* ( $\omega$ ), C17 and C17\* ( $\omega$ -1), as well as C16 and C16\* ( $\omega$ -2). Interestingly, for OHA no atoms were predicted to be sites of the CYP metabolism. The main sites (C-atoms) of the CYP metabolism of SAHA were aromatic C-atoms at positions *ortho* and *para* (C2' and C4', respectively) towards the alkyl chain substitution in the SAHA molecule. The ADMET analysis predicted the inhibition of the testosterone (ti) metabolism by CYP 3A4 for LHA and LA as well as SAHA.

Both pairs, *i.e.*, OHA and LHA as well as OA and LA were characterized by high lipophilicity (MlogP = 4.141 to 4.261) compared to the lipophilicity of SAHA (MlogP = 2.057) (Table 3). The low absorption scores expressed as S-Absn Risk (Table 3) were predicted for all of the investigated molecules and the highest score was obtained for OA (S-Absn Risk = 3). The common predicted parameters of overall ADMET risk for all of the investigated molecules (except for SAHA) according to these predictions corresponded to the big number of rotational bonds (RB), lipophylicity ( $\omega$ , Mlog) and water solubility (Sw).

TOX Risk<sup>TM</sup>, the toxicity liability score composed of 7 rules (hERG, acute rat toxicity, rat and mouse carcinogenicity, hepatotoxicity, liver enzyme elevation, and Ames mutagenicity) predicted the highest score of 3 that corresponded to hepatotoxicity (SG and Hp) and mutagenicity (Mu) for LHA, the score of 2 for SAHA (Hp and Mu) and the lowest score of 1 for OHA, OA and LA (Mu, Hp and Hp, respectively) (Table 3).

On the basis of an *in silico* Amest test of the ADMET Predictor<sup>TM</sup> modules, mutagenicity in *S. typhimurium* was predicted for hydroxamic acids (OHA, LHA and SAHA) and not for carboxylic acids (OA and LA) which implies that CONHOH moiety contributes to the potential mutagenic effect of the FHAs.

Regarding the environmental toxicity observed with LHA [58], in this study biodegradable toxicity was predicted for all of the investigated molecules except for SAHA. The environmental toxicity as a consequence of a high bioconcentration factor (BCF) defined as the ratio of the chemical concentration in biota to that in water at steady state, as a result of absorption *via* the respiratory surface, was predicted to be as follows: TOX BCF<sup>TM</sup> of LA (48.945) > OA (31.734) > LHA (6.280) > OHA (4.453) > SAHA (1.275) (Table 3).

On the basis of the predicted TOX Risk<sup>TM</sup> scores, as well as overall ADMET Risk<sup>TM</sup> scores displayed in this order: LHA (6.846) > LA (5.987) > OA (5.000) > OHA (4.130) > SAHA (3.197), it can be concluded that LHA is the most toxic molecule in this study with the worse safety profile (Table 3).

Table 3. Computed molecular descriptors and the predicted ADMET properties of FHAs *i.e.*, OHA and LHA, for their corresponding precursor fatty acids, oleic and linoleic acid (OA and LA, respectively) and SAHA (suberoylanilide hydroxamic acid, vorinostat) by ADMET Predictor™ [60]

Acid	$M_r$ <sup>1</sup>	MlogP <sup>2</sup>	ADMET Risk <sup>3</sup>	ADMET Code	CYP Risk <sup>4</sup>	CYP Code	TOX MUT Risk <sup>5</sup>	TOX MUT Code	TOX Risk <sup>6</sup>	TOX Code	S-Absn Risk <sup>7</sup>	S-Absn Risk Code	TOX Rat <sup>17</sup>	TOX BRM Rat <sup>18</sup>	TOX BRM Mouse <sup>19</sup>	BCF <sup>20</sup>
OHA	297.484	4.238	4.130	RB <sup>8</sup> , ow <sup>9</sup> , Sw <sup>10</sup> , fu <sup>11</sup> , Mu <sup>12</sup>	0	None	2	m1 <sup>20</sup> , S2 <sup>21</sup>	1	Mu	2.966	RB, ow, Sw	4015.411	111.18 1	1034.993	4.453
LHA	295.468	4.141	6.846	RB, ow, Sw, SG <sup>13</sup> , Hp <sup>14</sup> Mu, 1A <sup>15</sup> , ti <sup>16</sup>	1.225	1A, ti	2	m1, S2	3	SG, Hp, Mu	2.621	RB, ow, Sw	2722.121	79.991	1056.419	6.28
OA	282.470	4.261	5.000	RB, ow, Sw, fu, Hp	0	None	0	None	1	Hp	3.000	RB, ow, Sw	7019.417	536.65 2	743.482	31.734
LA	280.454	4.165	5.987	RB, ow, Sw, fu, Hp, ti	1.000	ti	0	None	1	Hp	1.000	RB, ow, Sw	5050.166	411.21 4	761.805	48.945
SAHA	297.484	2.057	3.197	RB, Hp, Mu, 1A	0.697	1A	2	m1, S2	2	Hp, Mu	0.500	RB	1460.041	10.165	1522.536	1.275

<sup>1</sup> Relative molecular mass; <sup>2</sup> Lipophilicity (according to Moriguchi); <sup>3</sup> Overall ADMET toxicity; <sup>4</sup> Risk of the toxicity due to biotransformation by CYP enzymes; <sup>5</sup> The risk of mutagenicity in *S. Typhimurium*; <sup>6</sup> Risk of overall toxicity, including mutagenicity; <sup>7</sup> Low absorption risk; <sup>8</sup> RB – number of rotatable bonds (too flexible); <sup>9</sup> ow S+logP, S+logD at pH 7,4 and MlogP (too lipophilic); <sup>10</sup> Sw (low water solubility); <sup>11</sup> fu –low fraction unbound in plasma; <sup>12</sup> Mu – mutagenicity in a panel of *in silico* Ames tests with and without metabolic activation; <sup>13</sup> SG – hepatotoxicity based on another kind of serum enzyme profile; <sup>14</sup> Hp – hepatotoxicity based on one kind of serum enzyme profile; <sup>15</sup> excessive clearance by CYP 1A2; <sup>16</sup> ti – inhibition of testosterone metabolism by CYP 3A4. <sup>17</sup> Acute lethal toxicity in rat, the TD<sub>50</sub> value of a particular compound in units of mg/kg/day; <sup>18</sup> TD<sub>50</sub> value of a particular compound in units of mg/kg/day, TD<sub>50</sub> – dose of a substance administered orally to rats over the course of their lifetimes that results in the appearance of tumours in 50% of their population); <sup>19</sup> TD<sub>50</sub> value in mice mg/kg/day; <sup>20</sup> BCF – environmental bio-concentration factor;

484

485 **4. Materials and Methods**

486

487 The olive oil from autochthonous Croatian olive cultivars was donated from a producer in the  
488 town of Brela (Dalmatia, Croatia) and it served as the main source material for the preparation of  
489 the FHAs. Reagents, starting materials and solvents were purchased from common commercial  
490 suppliers: sodium hydroxide p.a. from Kemika (Croatia), hydroxylamine hydrochloride p.a., ferric  
491 (III) chloride and hydrochloric acid 37% from Merck (Germany), methanol for HPLC, hexane p.p.a.  
492 and dichloromethane p.p.a. from Sigma-Aldrich (Germany). Butylated hydroxyanisole (BHA),  
493 2,2-diphenyl-1-picrylhydrazyl (DPPH),  $\beta$ -carotene, linoleic acid, Tween-40 (polyoxyethylene  
494 sorbitan monopalmitate) and quercetin were purchased from Sigma-Aldrich Chemical Co. (USA).  
495 The other chemicals and solvents used in this study were of analytical grade. Precoated Merck silica  
496 gel 60 F<sub>254</sub> plates were used for thin-layer chromatography. The HeLa (ATCC® CCL-2™) tumor cell  
497 line (human, cervix, epithelial adenocarcinoma) and BJ (ATCC® CRL-2522™) normal cell line  
498 (human, skin, foreskin, fibroblast) were purchased from American Type Culture Collection,  
499 Manassas, VA, USA.

500 The melting point was determined at Stuart SMP3 melting point (Bibby Sterilin Ltd, Stone,  
501 Strafkodshire, UK) and is uncorrected. Mid-IR (MIR) spectra were recorded in KBr pellets using  
502 ABB Bomem MB102 Fourier-transform infrared spectrometer (Quebec, Canada). The FT Raman  
503 spectrum was recorded on a Bruker Equinox 55 IR spectrometer (Germany) equipped with a FRA  
504 106/S module and Nd-YAG laser (1064 nm the wavelength of excitation). 1D and 2D homo- and  
505 hetero-nuclear <sup>1</sup>H- and <sup>13</sup>C-NMR spectra were recorded on a Bruker-Avance 600 MHz NMR  
506 spectrometer (Germany) operating at 600.133 MHz for the <sup>1</sup>H nucleus and 150.917 MHz for the <sup>13</sup>C  
507 nucleus. Samples were measured from CDCl<sub>3</sub> solution at 25 °C (298 K). Chemical shifts ( $\delta$ ) are in ppm,  
508 and are referred to the tetramethylsilane (TMS). The following 1D and 2D measurement techniques  
509 were used: standard <sup>1</sup>H and <sup>13</sup>C gated proton decoupling, APT, COSY, HMQC and HMBC. The 2D  
510 NMR spectra were measured in pulsed field gradient mode (z-gradient). Theoretical NMR data were  
511 computed for OHA and LHA using nmrshiftdb (<http://nmrshiftdb.nmr.uni-koeln.de>) and  
512 ChemDraw Ultra v. 11.0 <sup>13</sup>C NMR software packages. The analyses of the FHAs were achieved  
513 using matrix-assisted laser desorption/ionization time-of-flight/time-of flight mass spectrometry  
514 (MALDI TOF/TOF MS). A model 4800 Plus MALDI TOF/TOF analyser (Applied Biosystems Inc.,  
515 Foster City, CA, USA) equipped with the Nd:YAG laser (355 nm wavelength frequency 200 Hz and  
516 the pulse of 3-7 ns) was used. Colorimetric measurements were made by an Elisa microplate reader  
517 (iMark, BIO RAD, Hercules, CA, USA). Biological targets were predicted using the  
518 SwissTargetPrediction web server at <http://www.swisstargetprediction.ch/> and ADMET properties  
519 were predicted using ADMET Predictor™ (Simulations Plus, USA).

520

521

522 **4.1. Synthesis of fatty hydroxamic acids (FHAs)**

523

524 To a solution of olive oil (1.00 g) in hexane (15 mL), 5 mL of 2 M hydroxylamine in methanol  
525 solution prepared according to a previously published procedure [39] was added (Scheme 1). The  
526 reaction mixture was stirred at 130 rpm and 72 °C for 5 h and then at room temperature for 15 h.

The residue was filtered off and the hexane layer was separated and washed with 2 M HCl and water, dried over sodium sulphate, and evaporated in a vacuum. The crude product was purified by column chromatography (silica gel, dichloromethane/methanol, 9:1) to obtain the FHA product as a white solid residue (100 mg/g, expressed in mg per 1 g of olive oil). Following isolation and purification, the qualitative analysis test was performed by addition of iron (III) ions dissolved in hydrochloric acid to the solution of FHAs in methanol and the resulting purple coloured complex indicated the presence of hydroxamic acid groups.

#### 4.2. Characterization of fatty hydroxamic acids by spectroscopic methods

*Fatty hydroxamic acid mixture*: white solid; m.p. 64-69 °C;

FT-IR (KBr)  $\nu_{\max}$ : 3285 (-NH-OH), 3001 (-CH=CH-), 2952 (-CH<sub>3</sub>), 2918 (-CH<sub>2</sub>-), 1664 (-CO-, amide I), 1624 (-CO-, amide II) cm<sup>-1</sup>; data see in Table S-1 in the supplementary material;

Raman ( $\nu_{\max}$ /cm<sup>-1</sup>): data see in Table S-1 in the supplementary material.

<sup>1</sup>H-NMR (CDCl<sub>3</sub>, 600 MHz) data see in Table S-2 and Figure S-2 in the supplementary material

<sup>13</sup>C-NMR (CDCl<sub>3</sub>, 150 MHz) data see in Table S-2 and Figure S-3 in the supplementary material

MALDI TOF/TOF MS: m/z 320 (100) [M<sub>OHA</sub> + Na]<sup>+</sup>, m/z 318 (25) [M<sub>LHA</sub> + Na]<sup>+</sup>; all data see in Figure 2 and Figure S-7 in the supplementary material).

#### 4.3. Biological evaluation in vitro

##### 4.3.1. Antioxidant activity

*Free radical scavenging activity (RSA)* was evaluated by the scavenging of 1,1-diphenyl-2-picrylhydrazyl (DPPH) radicals according to a previously described method [61] with some modifications. A methanolic solution of DPPH (20 µL, 0.735 mg/mL) was added to 130 µL of either methanol (negative control) or FHA methanolic solutions of various concentrations. The mixture was vortexed for 1 min and then left to stand at room temperature in the dark. After 30 min absorbance was read at 545 nm. A standard antioxidant, the butylated hydroxyanisole (BHA), was used as positive control. RSA was expressed as the concentration that scavenges 50% of DPPH free radicals (EC<sub>50</sub>). RSA was calculated using the following equation:

$$X(\%) = \frac{A_{\text{control}} - A_{\text{sample}}}{A_{\text{control}}} \cdot 100 \quad (\text{Equation 1}), \text{ where } X \text{ represents the RSA in percentage } (\%)$$

while  $A_{\text{control}}$  is methanol and  $A_{\text{sample}}$  is FHAs solution in methanol. The concentration of FHAs sample that provides 50 % inhibition (EC<sub>50</sub>) was calculated by plotting the inhibition percentages against the concentrations of the sample.

*The chelating activity (ChA)* of FHAs toward ferrous ions (Fe<sup>2+</sup>) was studied according to modified literature procedures [62]. In brief, to an aliquot of the methanolic solution of FHAs (150 µL), 0.25 mM FeCl<sub>2</sub> solution (50 µL) was added. After 5 min, the reaction was initiated by adding 1.0 mM ferrozine solution (100 µL). Absorbance at 545 nm was recorded after 10 min of incubation at room temperature. A reaction mixture containing methanol (150 µL) instead of FHA solution

served as control. Quercetin (Qn) and EDTA were used as the chelating standards. The chelating activity (ChA) was expressed as the concentration that chelates 50% of  $\text{Fe}^{2+}$  ions ( $\text{EC}_{50}$ ). ChA was calculated using Equation 1, where X represents ChA while  $A_{\text{control}}$  is the absorbance of the negative control, i.e., blank solution without test compound and  $A_{\text{sample}}$  is the absorbance of the FHAs solution in methanol.

*The antioxidant activity (AOA) in  $\beta$ -carotene-linoleic acid assay* of FHAs was evaluated as described previously [62]. Tween 40 (200 mg) and  $\beta$ -carotene solution in chloroform (1.0 mL,  $\gamma = 0.2 \text{ mgL}^{-1}$ ) were mixed. After removing chloroform in a rotary evaporator, linoleic acid (20 mg) and aerated distilled water (30 mL) were added to the oily residue with vigorous stirring. Aliquots (200  $\mu\text{L}$ ) of the thus obtained emulsion were added to sample solutions in methanol (50  $\mu\text{L}$ ). After adding the emulsion to the FHA solution in methanol (sample solution), the reaction mixture was incubated at 50 °C for 1 h. During that period, the absorbance was measured at 450 nm at 15-minute intervals, starting immediately after sample preparation ( $t = 0 \text{ min}$ ) until the end of the experiment ( $t = 120 \text{ min}$ ). A reaction mixture containing methanol (50  $\mu\text{L}$ ) instead of the sample solution served as control. Butylated hydroxyanisol (BHA) was used as an antioxidant standard. The antioxidant activity was calculated using Equation 1 where X represents AOA, while  $R_{\text{control}}$  and  $R_{\text{sample}}$  are the average bleaching rates of the water control and antioxidant (test compound or BHA standard), respectively.

*The reducing power (RP) assay* of FHAs was conducted as previously described [63] with some modifications. In brief, each FHA solution of different concentrations (0.1 – 0.5 mg/mL) in distilled water (0.5 mL) was mixed with 1.25 mL of 0.2 M sodium phosphate buffer (pH 6.6) and 1.25 mL of 1% (mv) potassium ferricyanide. The mixture was incubated for 20 min at 50°C after which 1.25 mL of 10% trichloroacetic acid (m/v) was added and the mixture was centrifuged at 2795 rpm. The upper layer (1.25 mL) was mixed with 1.25 mL of deionized water and 0.25 mL of 0.1% (m/v) ferric chloride. Absorbance was measured at 700 nm against water as a blank. Ascorbic acid was used for comparison.

*Statistical analysis.* All assays regarding *antioxidant activity* were performed in triplicate. The results are expressed as mean  $\pm$  SD. Statistical comparisons were made using Student's t-test or one-way ANOVA, followed by Dunnett's *post-hoc* test for multiple comparison with the control. Statistical analyses were performed using the JMP V6 from SAS software (SAS Institute, Cary, NC, USA).

#### 4.3.2. Cytotoxic activity

Cytotoxic effects were determined by MTT [(3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetra sodium bromide)] assay *in vitro* [64]. MTT is a pale yellow substrate that cleaved by living cells to yield a dark blue formazan product in a process that requires active mitochondria. Thus the amount of cleaved MMT is directly proportional to the number of viable cells present, which is quantified by colorimetric methods. The experiments were carried out on one normal human cell line (ATCC® CRL-2522™) of foreskin fibroblast (BJ) between 28-29 passages and one epithelial tumour cell line (cervix adenocarcinoma - HeLa, ATCC® CCL-2™HeLa). Cells were cultured in Dulbecco's modified Eagle medium – DMEM (Gibco) supplemented with 10% heat-inactivated foetal bovine serum (FBS, Gibco, EU), 2 mM glutamine, and 100 U/0.1mg penicillin/streptomycin. All cells were grown as

monolayer in tissue culture flasks (BD Falcon, Germany) in humidified atmosphere under the conditions of 37 °C/5% of CO<sub>2</sub> gas in a CO<sub>2</sub> incubator (Shell Lab, Sheldon Manufacturing, USA). The trypan blue dye exclusion method was used to assess the cell viability.

The FHA sample was prepared as a stock solution (5 mg/mL) in DMSO. Working solutions (0.005 up to 5 mg/mL) were made in purified water prior to the experiment. Cells were seeded in 96 micro well flat bottom plates (Greiner, Frickenhausen, Austria) at concentration 2x10<sup>4</sup> cells/mL and left overnight in the CO<sub>2</sub> incubator allowing them to attach to the plate surface. After 72 h the compound addition growth medium was discarded and 5 mg/mL of MTT was added. After 4 h incubation at 37 °C the water insoluble MTT-formazan crystals were dissolved in DMSO. Absorbance was measured at 595 nm on an Elisa microplate reader (iMark, BIO RAD, Hercules, CA, USA). Controlled, the cells were grown under the same conditions and treated with appropriate concentration of solvent used for the preparation of the tested compound. All experiments were performed at least three times in triplicate. The percentage of cell growth (PG) was calculated using the following equation

$$PG = \frac{A - A_{blank}}{A_{control} - A_{blank}} \cdot 100 \quad (\text{Equation 2}), \text{ where } A \text{ means absorbance of FHAs solution, the blank}$$

medium is medium without cells containing MTT and DMSO while control is cell containing DMSO.

The GI<sub>50</sub> value, defined as compound concentration (mg/mL) leading to cellular viability reduction by 50%, was calculated and used as a parameter to compare cytotoxicity among the tested substances. STATISTICA 11 software and Kolmogorov-Smirnov two-sample tests were used to statistically evaluate the data obtained from the MTT test.

#### 4.3.3. In silico prediction of biological targets and ADMET properties

##### 4.3.3.1. In silico prediction of biological targets

Biological on- and off-targets of OH, LHA, OA, LA and SAHA were predicted using SwissTargetPrediction, a web server accessible free of charge and without login requirement at <http://www.swisstargetprediction.ch/>.

##### 4.3.3.1. In silico prediction of ADMET properties

ADMET properties of OHA, LHA, OA, LA and SAHA were evaluated using ADMET Predictor™ Version 8.1 (Simulations Plus Inc., USA).



## 5. Conclusions

FHAs were synthesized by hydroxylaminolysis from olive oil triacylglycerides as starting material. The obtained spectroscopic data revealed that the isolated product corresponds to the FHAs mixture of OHA and LHA (ratio 4:1). The results of preliminary *in vitro* assays and antitumor activity (HeLa cell line) revealed that the FHA mixture possessed moderate antioxidant and antiradical properties, as well as antitumor activity. The results of biological activity testing and computed data for FHAs highlighted OHA and LHA as promising lead-compounds for further research.

Although many hydroxamic acid-based HDAC inhibitors are currently in discovery and preclinical phases, the number of HDACIs that have been approved for the market, still remains low. Hence, there is permanent interest in the synthesis of new hydroxamic acid derivatives and their biological evaluation, so any new chemical entity in this class can be considered for investigation as a potentially new active drug.

**Supplementary Materials:** **Figure S1:** A parallel display of FTIR spectra of olive oil recorded as liquid on potassium bromide (KBr) pellet (**A**) and fatty hydroxamic acids (FHAs) recorded in potassium bromide (KBr) pellet (**B**), **Figure S2:** The  $^1\text{H}$ -NMR spectrum of FHAs mixture recorded in  $\text{CDCl}_3$  and chemical structures of its main components, *i.e.*, OHA and LHA with indicated signals for each proton groups, **Figure S3:**  $^{13}\text{C}$ -NMR spectrum of fatty hydroxamic acid (FHAs) mixture and chemical structures of its main components, *i.e.*, oleoyl and linoleyl hydroxamic acid (OHA and LHA, respectively) with labeled C-atoms, **Figure S4:** The 2D NMR HMQC (Heteronuclear Multiple Quantum Coherence) spectrum of FHAs in the region of aliphatic and olefinic protons displaying correlations through one bond. (The signal at 77.0 ppm ( $^{13}\text{C}$ -NMR) and 7.24 ppm ( $^1\text{H}$ -NMR) belong to  $\text{CDCl}_3$  solvent), **Figure S5:** A part of 2D NMR HMBC (Heteronuclear Multiple Bond Coherence) spectrum of FHAs displaying correlations through multiple bonds in the region of aliphatic and olefinic protons, **Figure S6:** A part of  $^1\text{H}$ - $^1\text{H}$ -COSY spectrum of FHAs in the region of aliphatic and olefinic protons, **Figure S7:** A MALDI-TOF/TOF tandem mass spectrum of FHAs mixture representing the region of unsaturated fatty hydroxamic acids, *i.e.*,  $m/z$  310 – 323 with molecular ions of OHA and LHA as their sodium alkali adducts  $[\text{M} + \text{Na}]^+$ ,  $[\text{M} + \text{Na} + \text{H}]^+$  and  $[\text{M} + \text{Na} + 2\text{H}]^+$ , **Table S1.** The list of vibrational frequencies and their assignments in FTIR (KBr) and Raman spectra of recorded FHAs, **Table S2.**  $^{13}\text{C}$ - and  $^1\text{H}$ -NMR spectroscopic data (recorded at 175 and 700 MHz, respectively;  $\text{CDCl}_3$ ) of fatty hydroxamic acids mixture (FHAs) consisted of oleoyl (OHA) and linoleyl hydroxamic acid (LHA\*) and theoretical<sup>2,3</sup> spectral data of OHA and LHA\* and literature spectral data of LHA.

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