



Article Metabolomic Profile and Cytotoxic Activity of Cissus incisa Leaves Extracts

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Copyright: © 2021 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Abstract: Cissus incisa leaves have been traditionally used in Mexican traditional medicine to treat certain cancerous illness. This study explored the metabolomic profile of this species using untargeted technique. Likewise, it determined the cytotoxic activity and interpreted all data by computational tools. The metabolomic profile was developed through UHPLC-QTOF-MS/MS for dereplication purposes. MetaboAnalyst database was used in metabolic pathway analysis and the network topological analysis. Hexane, chloroform/methanol, and aqueous extracts were evaluated on HepG2, Hep3B, HeLa, PC3, A549, and MCF7 cancer cell lines and IHH immortalized hepatic cells, using Cell Titer proliferation assay kit. Hexane extract was the most active against Hep3B (IC₅₀ = $27 \pm 3 \mu g/mL$), while $CHCl_3/MeOH$ extract was the most selective (SI = 2.77) on the same cell line. A Principal Component Analysis (PCA) showed similar profiles between the extracts, while a Venn diagram revealed 80 coincident metabolites between the bioactive extracts. The sesquiterpenoid and triterpenoid biosynthesis pathway was the most significant identified. The Network Pharmacology (NP) approach revealed several targets for presqualene diphosphate, phytol, stearic acid, δ -tocopherol, ursolic acid and γ -linolenic acid, involved in cellular processes such as apoptosis. This work highlights the integration of untargeted metabolomic profile and cytotoxic activity to explore plant extracts, and the NP approach to interpreting the experimental results.

Keywords: Cissus incisa; Vitaceae; extracts; untargeted metabolomics; cytotoxicity; network pharmacology

1. Introduction

Cancer is the second leading cause of death globally, responsible for an estimated 9.6 million deaths in 2018. Although important medical and technological advances have been made, conventional therapies directed against cancer have severe side effects and complications such as serious toxicities and development of resistance. In this point, the exploration and discovery of anticancer drugs from medicinal plants is playing an important role [1]. From ancient times, several medicinal plants have been consumed by patients in order to prevent and treat cancer, as an alternative therapy. These plants have been used because of their wealth in anticarcinogenic and chemoprotective potentials. Natural extracts from medicinal plants are a key source of antitumor agents with applicability in anticancer modern therapy [2]. It is known that the synergistic effects of plant extracts of a group of metabolites on a biological activity can play a role together, rather than as a single compound.

Recently, untargeted metabolomics have become a useful tool for the simultaneous analysis of many compounds in vegetal extracts. In contrast to targeted analyses, this

technique allows the uncovering of as many groups of metabolites as possible without necessarily identifying or quantifying a particular compound [3]. Mass Spectrometry (MS) in combination with high-performance chromatographic separation is considered the most universal approach for metabolome purposes by its sensitivity, specificity, and demonstrated efficiency in the analysis of plant metabolomes. Moreover, it is known that multivariate statistical techniques are frequently used in these studies, and for exploratory data analysis the PCA can be successfully applied [3,4].

On the other hand, a novel paradigm called NP has gained appreciation as method for omics data integration and multitarget drug development, which combines network biology and polypharmacology approaches. NP attempts to understand metabolites actions and interactions with multiple targets. Currently, this approach is getting attention in cancer research from natural products, since these products aim multiple protein targets and thus, are linked to many types of cancers [5].

Mexico stands out for its broad culture into traditional medicine. Despite the rich experience regarding the use of plants to treat diseases, very few have been studied regarding their phytochemical and pharmacological content. One under-explored species is *Cissus incisa* (Nutt.) Des Moul. Ex S. Watson (syn. *C. trifoliata*), which belongs to Vitaceae family. This plant is native to southern United States and northern of Mexico. It is fast growing and blooms in the summer. Leaves of this plant are used into traditional Mexican medicine to treat skin infections and tumors [6,7].

Because of our interest in giving scientific authentication and explanation of the traditional use of *C. incisa*, the antibacterial potential of some phytocompounds and extracts have been previously determined [8]. Further investigations on CHCl₃/MeOH extract led to isolation of several compounds such as: ceramides, cerebrosides, β -sitosterol, β -sitosterol-D-glucopyranoside, α -amyrin-3-*O*- β -D-glucopyranoside, and 2,3-dihydroxypropyl tetracosanoate [9,10]. Another study reported the chemical and biological profile of the stems of this plant [11]. In spite of the above, and to the best of our knowledge, there are no previous investigations about the cytotoxic activities related to the leaves of this plant.

Accordingly, in this work, the untargeted metabolomic technique was used to explore three extracts from *C. incisa* leaves, by UHPLC-QTOF-MS/MS. Metabolomic fingerprints were obtained by accurate mass measurements, and multivariate analyzes were applied to determine the phytochemical content of the extracts. In addition, the cytotoxic activity of extracts was evaluated on six human cancer cells lines. The integration of the metabolomic study and the cytotoxic activity revealed the cytotoxic metabolites from the bioactive extracts. Finally, a network pharmacology approach was applied to interpret the experimental results.

2. Results and Discussion

2.1. Metabolomic Profile Analysis of the Extracts

Metabolomic profiling of the extracts from *C. incisa* leaves by UHPLC-QTOF-MS/MS for dereplication purposes, led to the identification of 171, 260, and 114 metabolites in the hexane, CHCl₃/MeOH and aqueous extracts, respectively (Tables S1–S3 in Supplementary Material). Putative identification of compounds detected were made consulting several databases, such as: MEDLINE_Metabolites, Dictionary of Natural Products, KNApSAcK, PubChem, LIPID MAPS, and Human Metabolome Database (HMDB).

Based on the normalized areas data, three common primary metabolites were detected as the most abundant among the three extracts: two glycerophospholipids and a fatty acyl glycoside (Tables S1–S3). The percentages of abundance of each phytocompound in the hexane, CHCl₃/MeOH and aqueous extracts were as follows: (0.7491, 0.4864, 1.1213); (0.7480, 0.4856, 1.1222) and (0.7430, 0.4792, 1.0945), respectively. These results are fully comprehensible because glycerophospholipids are the most plentiful phospholipids localized in large amounts in plant cell membranes. In plants, approximately one-third of the organic phosphorus compounds are found in phosphoglycerolipids. In addition, glycerophospholipids participate in cell signaling and as an anchor for proteins in cell membranes [12]. Fatty acyl glucosides, meanwhile, are amphipathic compounds mainly produced by bacteria, yeast, fungi, marine invertebrates, and plants. Recent studies have demonstrated that they play an important role in plant-insect and plant-fungus interactions [13].

Regarding secondary metabolites, α -amyrin acetate and α -tocopherolquinone were the most abundant compounds in the hexane extract. In the CHCl₃/MeOH extract, the most abundant secondary metabolites were kazinol A and ursolic acid 3-*O*- α -L-arabinopyranoside. Meanwhile in the aqueous, armillane and chabrosterol were found to be the most plentiful compounds (Tables S1–S3).

As far as we know, this is the first time that a metabolomic fingerprint of *C. incisa* leaves is reported, thus contributing to the scientific knowledge of this species. A PCA scores plot was obtained (Figure 1) from a multivariate statistical analysis. The PCA showed close metabolomic profiles for the three analyzed extracts. In Figure 1, a similar composition is observed regarding the presence of fatty acyls, sphingolipids, sterols, glycerolipids, prenol lipids, and terpenoids; although their ratio within the extracts is variable. Thirty-three common compounds between these extracts were found (Figure 2, Table S4). Additionally, 80 common compounds were detected only in the hexane and CHCl₃/MeOH, which are included in Table 1.

The findings presented here agree with those reported by Kumar et al. [14] and Chipiti et al. [15] for the leaf extracts of *C. quadrangularis* and *C. cornifolia*, respectively.



Figure 1. PCA plot of *C. incisa* leaves extracts. PC1 (64.99%), PC2 (35.01%). Within the PCA graph the hexane extract is represented with brown color, CHCl₃/MeOH in red and the aqueous one in blue. Different classes of metabolites identified in each extract are also represented.



Figure 2. Venn diagram showing the common compounds among the three extracts (n = 33), and the common compounds among the most active extracts (hexane and CHCl₃/MeOH, n = 80).

Table 1.	Common	compounds	identified i	n the	hexane	and	CHCl ₃ /MeC	OH extracts	(UHPLC-	QTOF-MS	/MS)-	and
metabolo	mic pathwa	ays.										

Identified Metabolites	Molecular Formula	Accurate Mass	Metabolite Class	Related Pathway
<i>α</i> -Tocopherolquinone *	C ₂₉ H ₅₀ O ₃	446.3760	Diterpenoid	
Alpinumisoflavone dimethyl ether *	$C_{22}H_{20}O_5$	364.1311	Flavonoid	
7,9,13,17-tetramethyl-75,14S-dihydroxy- 2E,4E,8E,10E,12E,16-octadecahexaenoic acid *	$C_{22}H_{32}O_4$	360.2301	Fatty acid derivative	
7-oxo-β-Sitosterol	$C_{29}H_{48}O_2$	428.3654	Sterol	
Heneicosan-2-one *	$C_{21}H_{42}O$	310.3236	Fatty Acyl	
Gancaonin R *	$C_{24}H_{30}O_4$	382.2144	Stilbene	
1-Monoacylglycerol	$C_{21}H_{36}O_4$	352.2614	Acyl glycerol	
14-O-(α-L-rhamnopyranosyl)-7 <i>S</i> ,14 <i>R</i> -dihydroxy- 7,9,13,17-tetramethyl-2 <i>E</i> ,4 <i>E</i> ,8 <i>E</i> ,10 <i>E</i> ,12 <i>E</i> ,16 <i>E</i> - octadecahexaenoic acid	$C_{28}H_{42}O_8$	506.2880	Fatty acid glycoside	
Calycanthidine *	$C_{23}H_{28}N_4$	360.2314	Alkaloid	
1,2,6α,6β,9,9,12α-Heptamethyl-10-[(3,4,5-trihydroxyoxan- 2-yl)oxy]-1,2,3,4,4α,5,6,6α,6β,7,8,8α,9,10,11,12,12α,12β,13, 14β-icosahydropicene-4α-carboxylate	$C_{35}H_{56}O_7$	588.4026	Terpenoid	
2-Heptadecylfuran *	C ₂₁ H ₃₈ O	306.2923	Heteroaromatic compound	
5-Methoxy-3-(2R-acetoxy-pentadecyl)-1,4-benzoquinone	$C_{24}H_{38}O_5$	406.2719	Quinone	
Phytol *	$C_{20}H_{40}O$	296.3079	Diterpenoid	
Oxyacanthine *	$C_{37}H_{40}N_2O_6\\$	608.2886	Lignan	
Yucalexin B16 *	$C_{20}H_{28}O_2$	300.2089	Diterpenoid	
Campesteryl <i>p</i> -coumarate	$C_{37}H_{54}O_3$	546.4072	Steroid ester	
1-dodecanoyl-glycero-3-phospho-(1'-sn-glycerol)	$C_{18}H_{37}O_9P$	428.2175	Glycerophospholipid	
10,13-Epoxy-11-methyloctadeca-10,12-dienoic acid *	C ₁₉ H ₃₂ O ₃	308.2351	Fatty Acyl derivative	
Spheroidenone	$C_{41}H_{58}O_2$	582.4437	Carotene derivative	

Identified Metabolites	Molecular Formula	Accurate Mass	Metabolite Class	Related Pathway
2-Monopalmitoylglycerol	C ₁₉ H ₃₈ O ₄	330.2771	Monoglyceride	
Doristerol	C ₂₇ H ₄₆ O	386.3549	Sterol	
(12 <i>S</i> ,15 <i>S</i>)-15- <i>O</i> -demethyl-10,29-dideoxy-11,12- dihydro-striatin C	$C_{25}H_{38}O_{6}$	434.2668	Terpene	
δ -Tocopherol *	$C_{27}H_{46}O_2$	402.3498	Prenol lipid	Ubiquinone and other terpenoid-quinone biosynthesis
16β-16-Hydroxy-3-oxo-1,12-oleanadien-28-oic acid	$C_{30}H_{44}O_4$	468.3240	Triterpene	
(3 <i>E</i>)-4-(2,3-dihydroxy-2,5,5,8α-tetramethyl- decahydronaphthalen-1-yl)but-3-en-2- one *	$C_{18}H_{30}O_3$	294.2195	Sesquiterpenoid	
3β ,18 β -3-Methoxy-11-oxo-12-oleanen-30-oic acid *	$C_{31}H_{48}O_5$	484.7104	Triterpenoid	
(1-cyano-2-methylprop-2-en-1-yl) 9Z,12Z-octadecadienoate	C ₂₃ H ₃₇ NO ₂	359.2824	Fatty Acyl	
Fragarin	$C_{21}H_{21}O_{10}$	434.1207	Flavonoid	
Flavoxate	C ₂₄ H ₂₅ NO ₄	391.1784	Flavonoid	
β -Citraurinene	C ₃₀ H ₄₂ O	418.3236	Triterpenoid	
N-(3-hydroxy-dodecanoyl)-homoserine lactone *	$C_{16}H_{29}NO_4$	299.2097	Fatty Acyl	
1-(11Z,14Z-eicosadienoyl)-glycero-3-phosphate	C ₂₃ H ₄₃ O ₇ P	462.2746	Glycerophospholipid	Glycerophospholipid metabolism
Cavipetin D	$C_{25}H_{38}O_5$	418.2719	Diterpenoid	
10-Methoxyheptadec-1-en-4,6-diyne-3,9-diol *	$C_{18}H_{28}O_3$	292.2038	Fatty Acyl	
1-pentadecanoyl-2-arachidonoyl-sn-glycero-3-phosphate	C ₃₈ H ₆₇ O ₈ P	682.4574	Glycerophospholipid	
Diisobutyl phthalate	$C_{16}H_{22}O_4$	278.1516	Pollutant	
Lucidone A *	$C_{24}H_{34}O_5$	402.2406	Sesquiterpenoid	
1-(9Z,12Z-octadecadienoyl)-rac-glycerol	$C_{21}H_{38}O_4$	354.2770	Glycerolipid	
(3S,5R,6S,7E,9x)-7-Megastigmene-3,6,9-triol9-glucoside	$C_{19}H_{34}O_8$	390.2253	Fatty acyl glycosides	
5,7,4'-Trimethoxyflavan	$C_{18}H_{20}O_4$	300.1362	Flavonoid	
all-trans-Heptaprenyl diphosphate *	$C_{35}H_{60}O_7P_2$	654.3814	Prenol lipid	
1-(1Z-octadecenyl)-2-(5Z,8Z,11Z,14Z,17Z- eicosapentaenoyl)-glycero-3-phospho-(1'-sn-glycerol)	C44H77O9P	780.5305	Glycerophospholipid	
Heliotrine *	C ₁₆ H ₂₇ NO ₅	313.1889	Member of pyrrolizines	
(all-E)-6'-Apo-y-caroten-6'-al	C ₃₂ H ₄₂ O	442.3272	Prenol lipid	
1-dodecanoyl-sn-glycero-3-phosphocholine	$C_{20}H_{42}NO_7P$	439.2699	Glycerophospholipid	
1-(9Z-hexadecenoyl)-2-(11Z-eicosenoyl)-glycero-3- phosphoserine	C ₄₂ H ₇₈ NO ₁₀ P	787.5363	Glycerophospholipid	
Stylisterol B	$C_{28}H_{46}O_4$	446.3396	Sterol Lipid	
Stylisterol A *	C ₂₈ H ₄₆ O ₃	430.3447	Sterol Lipid	
Cucurbitacin E *	$C_{32}H_{44}O_8$	556.3036	Triterpenoid	
(22E,24R)-Stigmasta-4,22-diene-3,6-dione	$C_{29}H_{44}O_2$	424.3341	Lipid	
Junceic acid *	$C_{21}H_{30}O_3$	330.2195	Prenol lipid	
2-Hydroxy-6-tridecylbenzoic acid *	$C_{20}H_{32}O_3$	320.2351	phenolic compound	
1-docosanoyl-glycero-3-phospho-(1'-sn-glycerol)	C ₂₈ H ₅₇ O ₉ P	568.3740	Glycerophospholipid	
Gibberellin A12 aldehyde *	C ₂₀ H ₂₈ O ₃	316.2038	Prenol lipid	Diterpenoid biosynthesis
Matricin *	$C_{17}H_{22}O_5$	306.1467	Prenol lipid	
19-α-19-hydroxy-3,11-dioxo-12-ursen-28-oic acid	C ₃₀ H ₄₄ O ₅	484.3188	Triterpenoid	
<i>α</i> -Amyrin acetate *	$C_{32}H_{52}O_2$	468.3967	Triterpenoid	

Table 1. Cont.

$\begin{array}{c c c c c c c c } \hline (5\alpha,25R)\mbox{-}Spirostan-3,6\mbox{-}dione & C_{27}H_{40}O_4 & 428.2926 & Sterol \\ \hline \hline Ent-9\mbox{-}L1\mbox{-}phytoP * & C_{18}H_{28}O_4 & 308.1988 & Fatty Acyl \\ \hline & & & & & \\ \gamma\mbox{-}Linolenic Acid * & & & & \\ \hline & & & & \\ \gamma\mbox{-}Linolenic Acid * & & & & \\ \hline & & & & \\ C_{18}H_{30}O_2 & 278.2246 & Fatty acid & & \\ \hline & & & & \\ Fatty acid & & & \\ \hline & & & \\ respane & & & \\ \hline & & & \\ Austroinulin * & & & \\ \hline & & & \\ Presqualene diphosphate * & & \\ \hline & & & \\ \hline$	Identified Metabolites	Molecular Formula	Accurate Mass	Metabolite Class	Related Pathway
Ent-9-L1-phytoP* $C_{18}H_{28}O_4$ 308.1988Fatty Acyl γ -Linolenic Acid* $C_{18}H_{30}O_2$ 278.2246Fatty acidBiosynthesis of unsaturated fatty acidsCrispane $C_{20}H_{32}O_3$ 320.2351TerpeneAustroinulin* $C_{20}H_{34}O_3$ 322.2508DiterpenoidPresqualene diphosphate* $C_{30}H_{52}O_7P_2$ 586.3188Terpenoid	(5α,25R)-Spirostan-3,6-dione	C ₂₇ H ₄₀ O ₄	428.2926	Sterol	
γ -Linolenic Acid * $C_{18}H_{30}O_2$ 278.2246 Fatty acidBiosynthesis of unsaturated fatty acidsCrispane $C_{20}H_{32}O_3$ 320.2351 TerpeneAustroinulin * $C_{20}H_{34}O_3$ 322.2508 DiterpenoidPresqualene diphosphate * $C_{30}H_{52}O_7P_2$ 586.3188 Terpenoid	Ent-9-L1-phytoP *	C ₁₈ H ₂₈ O ₄	308.1988	Fatty Acyl	
$\begin{tabular}{ c c c c c } \hline Crispane & C_{20}H_{32}O_3 & 320.2351 & Terpene \\ \hline Austroinulin* & C_{20}H_{34}O_3 & 322.2508 & Diterpenoid \\ \hline Presqualene diphosphate* & C_{30}H_{52}O_7P_2 & 586.3188 & Terpenoid \\ \hline \end{array} \end{tabular}$	γ -Linolenic Acid *	C ₁₈ H ₃₀ O ₂	278.2246	Fatty acid	Biosynthesis of unsaturated fatty acids
Austroinulin *C20H34O3322.2508DiterpenoidPresqualene diphosphate *C30H52O7P2586.3188TerpenoidSesquiterpenoid and triterpenoid biosynthesis; Steroid	Crispane	$C_{20}H_{32}O_3$	320.2351	Terpene	
Presqualene diphosphate *C_{30}H_{52}O_7P_2586.3188TerpenoidSesquiterpenoid and triterpenoid biosynthesis; Steroid	Austroinulin *	$C_{20}H_{34}O_3$	322.2508	Diterpenoid	
biosynthesis	Presqualene diphosphate *	$C_{30}H_{52}O_7P_2$	586.3188	Terpenoid	Sesquiterpenoid and triterpenoid biosynthesis; Steroid biosynthesis
(-)-Folicanthine * $C_{24}H_{30}N_4$ 374.2470 Indoles derivative	(–)-Folicanthine *	$C_{24}H_{30}N_4$	374.2470	Indoles derivative	
1-Octadecanoyl-2-docosanoyl-sn-glycero-3-phosphate C43H85O8P 760.5982 Glycerophospholipid	1-Octadecanoyl-2-docosanoyl-sn-glycero-3-phosphate	$C_{43}H_{85}O_8P$	760.5982	Glycerophospholipid	
Ursolic acid C ₃₀ H ₄₈ O ₃ 456.3603 Triterpenoid	Ursolic acid	$C_{30}H_{48}O_3$	456.3603	Triterpenoid	
Amabiline *C15H25NO4283.1784Carboxylic ester	Amabiline *	$C_{15}H_{25}NO_4$	283.1784	Carboxylic ester	
(–)-Epicatechin 3'-O-sulfate C ₁₅ H ₁₄ O ₉ S 370.0358 Flavonoid	(–)-Epicatechin 3'-O-sulfate	$C_{15}H_{14}O_9S$	370.0358	Flavonoid	
Stearic acid * $C_{18}H_{36}O_2$ 284.2715Fatty acidBiosynthesis of unsaturated fatty acids	Stearic acid *	C ₁₈ H ₃₆ O ₂	284.2715	Fatty acid	Biosynthesis of unsaturated fatty acids
Grandifloric acid * C ₂₀ H ₃₀ O ₃ 318.2194 Terpene	Grandifloric acid *	$C_{20}H_{30}O_3$	318.2194	Terpene	
1,2-Dihexadecanoylphosphatidylglycerol phosphate C ₃₈ H ₇₆ O ₁₃ P ₂ 802.4761 Glycerophospholipid	1,2-Dihexadecanoylphosphatidylglycerol phosphate	$C_{38}H_{76}O_{13}P_2$	802.4761	Glycerophospholipid	
Yucalexin B5 * C ₂₀ H ₂₆ O ₃ 314.1881 Terpene	Yucalexin B5 *	$C_{20}H_{26}O_3$	314.1881	Terpene	
6-Deoxohomodolichosterone * C ₂₉ H ₅₀ O ₄ 462.3709 Sterol Lipid	6-Deoxohomodolichosterone *	$C_{29}H_{50}O_4$	462.3709	Sterol Lipid	
7',8'-Dihydro-8'-hydroxycitraniaxanthin * C ₃₃ H ₄₄ O ₃ 488.3290 Triterpenoid	7',8'-Dihydro-8'-hydroxycitraniaxanthin *	C ₃₃ H ₄₄ O ₃	488.3290	Triterpenoid	
5,6-Epoxy-5,6-dihydro-10'-apo-β,γ-carotene-3,10'-diol * C ₂₇ H ₃₈ O ₃ 410.2820 Carotenoid	5,6-Epoxy-5,6-dihydro-10'-apo-β,γ-carotene-3,10'-diol *	C ₂₇ H ₃₈ O ₃	410.2820	Carotenoid	
<i>N</i> -tetradecanoyl glutamine * $C_{19}H_{36}N_2O_4$ 356.2675 Fatty Acyl	N-tetradecanoyl glutamine *	$C_{19}H_{36}N_2O_4$	356.2675	Fatty Acyl	
Stigmastane-3,6-dione *C29H48O2428.3654Sterol Lipid	Stigmastane-3,6-dione *	C ₂₉ H ₄₈ O ₂	428.3654	Sterol Lipid	
2-Stearyl citric acid * C ₂₄ H ₄₄ O ₇ 444.3087 Tricarboxylic acid	2-Stearyl citric acid *	C ₂₄ H ₄₄ O ₇	444.3087	Tricarboxylic acid	
1-(4Z,7Z,10Z,13Z,16Z,19Z-Docosahexaenoyl)-2-(13Z- docosenoyl)- <i>sn</i> -glycero-3- phosphocholine * C ₅₂ H ₉₀ NO ₈ P 887.6404 Glycerophospholipid	1-(4Z,7Z,10Z,13Z,16Z,19Z-Docosahexaenoyl)-2-(13Z- docosenoyl)- <i>sn</i> -glycero-3- phosphocholine *	C ₅₂ H ₉₀ NO ₈ P	887.6404	Glycerophospholipid	
17-Phenyl heptadecanoic acid * C ₂₃ H ₃₈ O ₂ 346.2872 Fatty Acyl	17-Phenyl heptadecanoic acid *	C ₂₃ H ₃₈ O ₂	346.2872	Fatty Acyl	

Table 1. Cont.

* Not detected in aqueous extract.

2.2. Cytotoxic Activity

Cytotoxic activity of *C. incisa* leaves extracts is also reported here for the first time, which was determined on six human cancer cells. The experimental results are shown in Table 2. According to the National Cancer Institute of the United States of America, an extract is considered active if it achieves an $IC_{50} \leq 30 \ \mu\text{g/mL}$ on tumor cells [16]. In this sense, the hexane extract exhibited cytotoxic activity on Hep3B ($IC_{50} = 27 \pm 3 \ \mu\text{g/mL}$) and HepG2 ($IC_{50} = 30 \pm 6 \ \mu\text{g/mL}$), being the most active extract. In the case of CHCl₃/MeOH extract, it was less active on hepatocellular cancer cells, reaching $IC_{50} = 39 \pm 3 \ \mu\text{g/mL}$ and $31 \pm 2 \ \mu\text{g/mL}$, respectively. Previously, Opoku et al. [17] reported the antiproliferative activity of MeOH extract of *C. quadrangularis* against the HepG2 cell line with 36.58% of inhibition of proliferation. On the other hand, the hexane extract exhibited certain cytotoxicity on Hela and A549 cancer lines ($IC_{50} = 40 \pm 2 \ \text{and} 52 \pm 2 \ \mu\text{g/mL}$) and PC3 ($57 \pm 4 \ \mu\text{g/mL}$).

	Hexane Extract		CHCl ₃ /MeC	OH Extract	Aqueous	Extract	Paclitaxel	
Cell Lines	IC ₅₀ (μg/mL)	SI	IC ₅₀ (μg/mL)	SI	IC ₅₀ (μg/mL)	SI	IC ₅₀ (μg/mL)	SI
HepG2	30 ± 6	1.5	39 ± 3	2.21	>100	ND	$64 imes 10^{-3}$	1.24
Hep3B	27 ± 3	1.66	31 ± 2	2.77	>100	ND	$33 imes 10^{-3}$	2.41
HeLa	40 ± 2	ND	61 ± 4	ND	>100	ND	$4.78 imes 10^{-3}$	ND
A549	52 ± 2	ND	77 ± 6	ND	>100	ND	$5.12 imes 10^{-3}$	ND
PC3	76 ± 5	ND	57 ± 4	ND	>100	ND	10.2×10^{-3}	ND
MCF7	74 ± 6	ND	50.7 ± 6	ND	>100	ND	$4.27 imes 10^{-3}$	ND
IHH	45 ± 3		86 ± 5		>100		$79.4 imes 10^{-3}$	

Table 2. Cytotoxic activity of *C. incisa* leaves extracts.

Values expressed are \pm SD of three independent experiments (*n* = 3); ND = not determined.

The Selectivity Index (SI) was determined only for hepatocellular carcinoma cell lines, since they were the most sensitive of all tested (Table 2). It has been reported that SI values less than 2 can indicate toxicity for an extract or a pure compound towards mammal cells [18]. CHCl₃/MeOH extract gave a SI = 2.77 on Hep3B, and SI = 2.21 on HepG2, surpassing the values of the control (Paclitaxel) on the same cell lines (2.41 and 1.24, respectively). As consequence, the CHCl₃/MeOH extract from *C. incisa* leaves was the most selective.

The aqueous extract obtained by successive extractions did not show cytotoxic activity in any cancer cell line tested. Different results were obtained by Sáenz et al. [19] evaluating the aqueous extract of *C. sicyoides* leaves (direct extraction) on HEp-2 cells finding a $IC_{50} = 43.2 \pm 2.4 \mu g/mL$. In addition, our aqueous extract did not show cytotoxicity in immortalized cells ($IC_{50} > 100 \mu g/mL$), which is a good first step for further safety studies of the total extract of *C. incisa* aerial parts.

2.3. Metabolomics Pathway Analysis

Based on the biological properties displayed by the hexane and $CHCl_3/MeOH$ extracts, we focused on exploring the 80 common metabolites among these extracts, using the Metabolomics Pathway Analysis (MetPA). As a consequence, the most relevant pathways involving these metabolites were identified, in this case, nine networks were revealed (see in Table 1). The threshold of impact was set to 0.10. The pathway is considered to be closely related if its impact value is higher than this value.

The results obtained from MetPA shows four important routes in plants operation, belonging to their primary metabolism: Linoleic acid metabolism, alpha-Linolenic acid metabolism, Glycerophospholipid metabolism, and Fatty acid biosynthesis. However, the most significant pathway identified via MetPA are those related to the biosynthesis of secondary metabolites, specially terpenes and sterols. This is consistent, since terpenoids and sterols from leaves exhibite a multifunctionality role in plants: more specialized chemical interactions and protection in the abiotic and biotic environment [20]. The results from pathway analysis are presented in detail in Table 3, and only the pathway with higher impact is presented graphically (Figure 3).

No.	Pathway Name	Total *	Expected	Hits *	Raw p *	Holm p *	FDR p *	Impact *
1	Biosynthesis of unsaturated fatty acids	22	0.21	2	1.72×10^{-2}	1.00	1.00	0.00
2	Linoleic acid metabolism	4	0.04	1	3.73×10^{-2}	1.00	1.00	0.00
3	Sesquiterpenoid and triterpenoid biosynthesis	24	0.23	1	$2.05 imes 10^{-1}$	1.00	1.00	0.20374
4	alpha-Linolenic acid metabolism	28	0.26	1	2.35×10^{-1}	1.00	1.00	0.00
5	Diterpenoid biosynthesis	28	0.26	1	$2.35 imes 10^{-1}$	1.00	1.00	0.07625
6	Glycerophospholipid metabolism	37	0.35	1	$2.99 imes 10^{-1}$	1.00	1.00	0.07614
7	Ubiquinone and other terpenoid-quinone biosynthesis	38	0.36	1	3.06×10^{-1}	1.00	1.00	0.02227
8	Steroid biosynthesis	45	0.42	1	$3.52 imes 10^{-1}$	1.00	1.00	0.02644
9	Fatty acid biosynthesis	56	0.56	1	$4.18 imes10^{-1}$	1.00	1.00	0.00

Table 3. Results from Pathway Analysis with MetaboAnalyst.

* Total is the total number of compounds in the pathway; the Hits is the actually matched number from the user uploaded data; the Raw *p* is the original *p* value calculated from the enrichment analysis; the Holm *p* is the *p* value adjusted by Holm-Bonferroni method; the FDR *p* is the *p* value adjusted using False Discovery Rate; the Impact is the pathway impact value calculated from pathway topology analysis.



Figure 3. Pathway analysis (A) metabolome view, (B) pathway with higher impact.

2.4. Correspondence between Metabolomic Profiling and Cytotoxic Activity

The distribution of the 80 coincident compounds is presented in a heat map (Figure 4), which contains the normalized relative areas of these metabolites, identified in the hexane, $CHCl_3/MeOH$ and the aqueous extracts. The heat map also shows the distribution between the three extracts of the cytotoxic metabolites reported against the same cell lines included in this study (or some related ones). It can be seen that most of the cytotoxic compounds are found in a higher proportion within the hexane extract.



Figure 4. Heat map with the distribution according to the normalized areas of the 80 shared metabolites, identified in the active extracts. Metabolites with previous reports of cytotoxic activity are highlighted in red. Aque *: Aqueous extract; Ch/Me *: CHCl₃/MeOH extract; Hex *: Hexane extract; Comp *: Compounds: 1: 1-(1Z-octadecenvl)-2-(5Z,8Z,11Z,14Z,17Z-eicosapentaenoyl)-glycero-3-phospho-(1'-sn-glycerol) **2**: 1,2-dihexadecanoylphosphatidylglycerol phosphate; 3: (3S,5R,6S,7E,9x)-7-megastigmene-3,6,9-triol 9-glucoside; 4: α -amyrin acetate; 5: α -tocopherolquinone; 6: 1-(9Z-hexadecenoyl)-2-(11Z-eicosenoyl)-glycero-3-phosphoserine; 7:gibberellin A12 aldehyde; 8: 16β-16-hydroxy-3-oxo-1,12-oleanadien-28-oic acid; 9: (3E)-4-(2,3-dihydroxy-2,5,5,8a-tetramethyl-decahydronaphthalen-1-yl)but-3-en-2one; 10: N-(3-hydroxy-dodecanoyl)-homoserine lactone; 11: (all-E)-6'-apo-y-caroten-6'-al; 12: campesteryl p-coumarate; 13: 1-docosanoyl-glycero-3-phospho-(1'-sn-glycerol); 14: 1,2,6a,6b,9,9,12a-heptamethyl-10-[(3,4,5-trihydroxyoxan-2yl)oxy]-1,2,3,4,4a,5,6,6a,6b,7,8,8a,9,10,11,12,12a,12b,13,14b-icosahydropicene-4a-carboxylate; 15: 7,9,13,17-tetramethyl-75,14S-dihydroxy-2E,4E,8E,10E,12E,16-octadecahexaenoic acid; 16: calycanthidine; 17: not identified; 18:5,7,4'trimethoxyflavan; 19:alpinumisoflavone dimethyl ether; 20: gancaonin R; 21: β -citraurinene; 22: spheroidenone; 23: 6-deoxohomodolichosterone; 24: 3β,18β-3-methoxy-11-oxo-12-oleanen-30-oic acid; 25: (5α,25R)-spirostan-3,6-dione; 26: 5,6-epoxy-5,6-dihydro-10'-apo- β_{γ} -carotene-3,10'-diol; 27: stylisterol B; 28: stylisterol A; 29: grandifloric acid; 30:phytol; 31: 7',8'-Dihydro-8'-hydroxycitraniaxanthin; 32:γ-linolenic acid; 33: Amabiline; 34: cucurbitacin E; 35: 2-stearyl citric acid; 36: (1cyano-2-methylprop-2-en-1-yl) 9Z,12Z-octadecadienoate; 37:ursolic acid; 38: ent-9-L1-phytoP; 39: 17-phenyl heptadecanoic acid; **40**: 1-dodecanoyl-glycero-3-phospho-(1'-sn-glycerol); **41**: 14-O-(α -L-rhamnopyranosyl)-7*S*,14*R*-dihydroxy-7,9,13,17tetramethyl-2E,4E,8E,10E,12E,16E-octadecahexaenoic acid; 42: 1-octadecanoyl-2-docosanoyl-sn-glycero-3-phosphate; 43: 10-methoxyheptadec-1-en-4,6-diyne-3,9-diol; 44: (125,155)-15-O-demethyl-10,29-dideoxy-11,12-dihydro-striatin C; 45: fragarin; 46: 5-methoxy-3-(2R-acetoxy-pentadecyl)-1,4-benzoquinone; 47: Yucalexin B16; 48: Junceic acid; 49: Yucalexin B5; 50: N-tetradecanoyl glutamine; 51: 10,13-Epoxy-11-methyloctadeca-10,12-dienoic acid; 52: 19α-19-hydroxy-3,11-dioxo-12-ursen-28-oic acid; 53: 1-monoacylglycerol; 54: (–)-folicanthine; 55: 7-oxo- β -sitosterol; 56: 1-dodecanoyl-sn-glycero-3-phosphocholine; 57: 1-(9Z,12Z-octadecadienoyl)-rac-glycerol; 58: 1-(11Z,14Z-eicosadienoyl)-glycero-3-phosphate; 59: Flavoxate; 60: (-)-Epicatechin 3'-O-sulfate; 61: 2-Heptadecylfuran; 62: matricin; 63: heneicosan-2-one; 64: austroinulin; 65: heliotrine; 66: doristerol; 67: crispane; 68: stearic acid; 69: 2-hydroxy-6-tridecylbenzoic acid; 70: stigmastane-3,6dione; 71: 1-pentadecanoyl-2-arachidonoyl-sn-glycero-3-phosphate; 72: oxyacanthine; 73: all-trans-Heptaprenyl diphosphate; 74: cavipetin D; 75: presqualene diphosphate; 76: lucidone A; 77: δ-tocopherol; 78: 1-(4Z,7Z,10Z,13Z,16Z,19Zdocosahexaenoyl)-2-(13Z-docosenoyl)-sn-glycero-3-phosphocholine; 79: (22E,24R)-stigmasta-4,22-diene-3,6-dione; 80: 2monopalmitoylglycerol. * tentative assignment based on accurate mass.

> As it presented in Table 2, hexane and $CHCl_3/MeOH$ extracts had similar cytotoxic results on the hepatocellular cells (even if the hexane extract was more active on Hep3B). These similarities can be explaining by the chemical content, these extracts include 80 common metabolites (Venn diagram Figure 2), showing a correspondence between the metabolomics profiles of the active extracts and the cytotoxic activity on Hep3B and HepG2 cell lines. These cells share common characteristics (for instance Wnt/ β -catenin activation [21], providing a unique platform for parallel comparisons, but also HepG2 and Hep3B are from different ethnic origins. Some differential gene expression (for instance; HepG2 cells are known to contain wild-type p53 whereas Hep3B cells are p53 deficient), provide a broad spectrum of mechanisms, particularly for apoptosis induction. Several studies suggested that phytosterols and terpenes disturb the cell cycle and induce apoptosis by activating caspases 3 and 9 in cancer cells. Particularly triterpenes and its derivates glycosides have shown effect against cancer cells and induction of apoptosis mechanism [22]. These phytocompounds are present in both extracts (hexane and CHCl₃/MeOH) (Figure 4).

It is necessary to point out that some of these 80 shared metabolites have been previously reported with cytotoxic activity against hepatocellular cancer cells: (5) α -tocopherolquinone, (30) phytol, (29) grandifloric acid, (34) cucurbitacin E, (4) α -amyrin acetate, (37) ursolic acid, (32) δ -linolenic acid, (72) oxyacanthine, (68) stearic acid, and (62) matricin of which, the first six are terpenoids, including three triterpenes. The presence of these cytotoxic metabolites may explain the cytotoxicity of the extracts (numbering is according to the heat map, Figure 4).

 α -Tocopherolquinone was dereplicated with molecular formula (C₂₉H₅₀O₃) and accurate mass 446.3760. This diterpene has reported good cytotoxic activity on HepG2 cells $(IC_{50} = 6.97 \pm 0.5 \,\mu g/mL)$ [23]. Another terpene dereplicated, phytol ($C_{20}H_{40}O$; accurate mass 296.3079) selectively inhibited the growth of the HepG2 cells with an IC_{50} value of $78 \pm 3.45 \,\mu$ M [24]. Another study showed that phytol exerted antitumor effect in hepatocellular carcinoma cells by activation of caspases 9/3 [25]. The triterpene cucurbitacin E (formula suggested $C_{32}H_{44}O_8$; accurate mass 556.3036) exhibited antiproliferative action on Hep3B cancer cells through inhibition of Wnt/ β -catenin activation [26]. Meanwhile, $C_{32}H_{52}O_2$ (468.3967) identified as α -amyrin acetate, and showed moderate activity on HepG2 = 148.9 \pm 1.80 μ M [27]. Other triterpene, ursolic acid (C₃₀H₄₈O₃; accurate mass 456.3603) is distributed among the three extracts; it has been widely studied in relation with anticancer properties. In Hep3B cell lines, ursolic acid has reduced the tumorigenesis in vivo, enhancing apoptosis in tumor tissues [28], and exerting antiangiogenic action [29]. A different work showed that ursolic acid displayed effects on cell viability, DNA fragmentation, mitochondrial membrane potential on human liver cancer HepG2 ($IC_{50} = 4$ μ M) and Hep3B (IC₅₀ = 8 μ M) cells [30]. A study showed in vivo that γ -linolenic acid $(C_{18}H_{30}O_2; 278.2246)$ reduced the proliferative and angiogenic effect of carcinoma hepatocellular induced in Wistar rats, by activation of a mitochondrial mediated apoptosis pathway [31]. Likewise, oxyacanthine (C₃₇H₄₀N₂O₆, 608.28863) attenuated cell proliferation ability and promoted cell apoptosis in mammary, prostatic, liver cancers cells [32], while stearic acid [33] ($C_{18}H_{36}O_2$; 284.2715) and grandifloric acid [34] (terpene; $C_{20}H_{30}O_3$; 318.2194) had the same action on HepG2 cells. A recent study determined the antiproliferative activity of extracts from Australian plants leaves that contained matricin ($C_{17}H_{22}O_5$; 306.1467), a prenol lipid, on HepG2 cells [35].

Other dereplicated compounds with promising anticancer activities reported are: (19) alpinumisoflavone dimethyl ether ($C_{22}H_{20}O_5$; 364.1311) that suppress the proliferation, migration/invasion, tumor angiogenesis and metastasis, and the promotion of apoptosis in various cancers: human oral epidermoid carcinoma KB cells ($IC_{50} = 4.13 \ \mu g/mL$), and murine leukemia P-388 ($IC_{50} = 4.31 \ \mu g/mL$) cells [36] and (10) N-(3-hydroxy-dodecanoyl)-homoserine lactone ($C_{16}H_{29}NO_4$; 299.2097) with pro-apoptotic activities [37] (28) Stylisterol A ($C_{28}H_{46}O_3$; 430.3447), (27) stylisterol B ($C_{28}H_{46}O_4$; 446.3396), [38] and (7) gibberellin A12 aldehyde [39] ($C_{20}H_{28}O_3$; 316.2038) have been found to have antiproliferative action in cancer cells, even with apoptosis induction.

In the heat map (Figure 4), it can be observed that the aforementioned metabolites occur in the hexane extract, justifying why this extract is the most active of the three tested. While, only (**55**) 7-oxo- β -sitosterol [40] (C₂₉H₄₈O₂; 428.3654), (**46**) 5-methoxy-3-(2*R*-acetoxy-pentadecyl)-1,4-benzoquinone [41] (C₂₄H₃₈O₅; 406.2719), (**69**) 2-hydroxy-6-tridecylbenzoic acid [42] (C₂₀H₃₂O₃; 320.2351), (**70**) stigmastane-3,6-dione [43] (C₂₉H₄₈O₂; 428.3654), and (**18**) 5,7,4'-trimethoxyflavan [44] (C₁₈H₂₀O₄; 300.1362) appear preferably in the CHCl₃/MeOH extract, and have also displayed anticancer effects. Therefore, all the metabolites presented so far are involved in the cytotoxic activity of the active extracts.

On the other hand, there are other experimental results from cytotoxic assays that are worth discussing. As we presented earlier, two cell lines (HeLa and A549) were more susceptible to hexane extract than CHCl₃/MeOH extract (Table 2). In this regard, the fold change analysis detected 30 up-regulated phytocompounds in hexane extract (Table 4). In contrast, PC3 and MCF7 cell lines were more sensitive to the CHCl₃/MeOH extract than the hexane one according to Table 2. Thirty-eight up-regulated compounds were

identified in the $CHCl_3/MeOH$ extract by the fold change analysis. Table 4 also contains these compounds, along with the previous studies against Hela, A549, MCF7 and/or related cell lines.

Table 4. Fold change analysis results in the hexane and CHCl₃/MeOH extracts.

Metabolites	Molecular Formula	Accurate Mass	Up-Regulation [Hexane Extract]	Up-Regulation [CHCl ₃ /MeOH Extract]	Biological Activity/ References			
	He	eLa, A549 and	/or related cell lines					
α-Tocopherolquinone	C ₂₉ H ₅₀ O ₃	446.3760	Yes	-	[23]			
Alpinumisoflavone dimethyl ether	$C_{22}H_{20}O_5$	364.1311	Yes	-	H2108 (IC ₅₀ = 33.5 μM); H1299 (IC ₅₀ = 38.8 μM) [36]			
7,9,13,17-tetramethyl-7 <i>S</i> ,14 <i>S</i> - dihydroxy-2 <i>E</i> ,4 <i>E</i> ,8 <i>E</i> ,10 <i>E</i> ,12 <i>E</i> ,16- octadecahexaenoic acid	C ₂₂ H ₃₂ O ₄	360.2301	Yes	-				
Heneicosan-2-one	$C_{21}H_{42}O$	310.3236	Yes	-				
Gancaonin R	$C_{24}H_{30}O_4$	382.2144	Yes	-				
Calycanthidine	$C_{23}H_{28}N_4$	360.2314	Yes	-				
2-Heptadecylfuran	$C_{21}H_{38}O$	306.2923	Yes	-				
Phytol	$C_{20}H_{40}O$	296.3079	Yes	-	Hela (IC $_{50}$ = 15.51 ± 0.76 µM); A549 (IC $_{50}$ = 56.98 ± 2.68 µM) [45]			
Yucalexin B16	$C_{20}H_{28}O_2$	300.2089	Yes	-				
δ -Tocopherol	C ₂₇ H ₄₆ O ₂	402.3498	Yes	-	[46]			
(3E)-4-(2,3-dihydroxy-2,5,5,8α- tetramethyl-decahydronaphthalen-1- yl)but-3-en-2-one	C ₁₈ H ₃₀ O ₃	294.2195	Yes	-				
3β,18β-3-Methoxy-11-oxo-12-oleanen- 30-oic acid	C ₃₁ H ₄₈ O ₅	484.7104	Yes	-				
N-(3-hydroxy-dodecanoyl)- homoserine lactone	C ₁₆ H ₂₉ NO ₄	299.2097	Yes	-	[37]			
10-methoxyheptadec-1-en-4,6-diyne- 3,9-diol	C ₁₈ H ₂₈ O ₃	292.2038	Yes	-				
Lucidone A	$C_{24}H_{34}O_5$	402.2406	Yes	-				
Stylisterol A	$C_{28}H_{46}O_3$	430.3447	Yes	-	HeLa (IC ₅₀ = 14.1 μM) [38]			
Cucurbitacin E	$C_{32}H_{44}O_8$	556.3036	Yes	-	[26]			
gibberellin A12 aldehyde	$C_{20}H_{28}O_3$	316.2038	Yes	-	[39]			
α-Amyrin acetate	$C_{32}H_{52}O_2$	468.3967	Yes	-	[27]			
Ent-9-L1-phytoP	$C_{18}H_{28}O_4$	308.1988	Yes	-	[47]			
γ -Linolenic Acid	C ₁₈ H ₃₀ O ₂	278.2246	Yes	-	[31]			
(–)-Folicanthine	C24H30N4	374.2470	Yes	-	A549 (IC ₅₀ = 7.76 μM) [48]			
Amabiline	C ₁₅ H ₂₅ NO ₄	283.1784	Yes	-				
Grandifloric acid	C ₂₀ H ₃₀ O ₃	318.2194	Yes	-	[34]			
Yucalexin B5	$C_{20}H_{26}O_3$	314.1881	Yes	-				
7′,8′-Dihydro-8′- hydroxycitraniaxanthin	C ₃₃ H ₄₄ O ₃	488.3290	Yes	-				
5,6-Epoxy-5,6-dihydro-10'-apo-β,γ- carotene-3,10'-diol	C ₂₇ H ₃₈ O ₃	410.2820	Yes	-				
N-tetradecanoyl glutamine	$C_{19}H_{36}N_2O_4$	356.2675	Yes	-				
2-Stearyl citric acid	C ₂₄ H ₄₄ O ₇	444.3087	Yes	-				
17-phenyl heptadecanoic acid	C ₂₃ H ₃₈ O ₂	346.2872	Yes	-				
PC3, MCF7 and/or related cell lines								

				TI D 1 d	
Metabolites	Molecular Formula	Accurate Mass	Up-Regulation [Hexane Extract]	Up-Regulation [CHCl ₃ /MeOH Extract]	Biological Activity/ References
4'-O-Geranylisoliquiritigenin	C ₂₅ H ₂₈ O ₄	392.1988	-	Yes	MDB-MB-231 (IC ₅₀ = 125.5 μM) [49]
1-Monoacylglycerol	$C_{20}H_{34}NO_{4}$	352.2619	-	Yes	
Sanguisorbin B	C ₃₅ H ₅₆ O ₇	588.4026	-	Yes	
Oxyacanthine	$C_{37}H_{40}N_2O_6$	608.2886	-	Yes	[32]
3-Methyl-5-pentyl-2- furannonanoic acid	$C_{19}H_{32}O_3$	308.2351	-	Yes	
2-Monopalmitoylglycerol	$C_{19}H_{38}O_4$	330.2771	-	Yes	
15-hydroxy-5,9-dimethyl-14- methylidenetetracyclo[11.2.1.01,10.04,9] hexadecane-5-carboxylic acid	$C_{20}H_{30}O_3$	318.2194	-	Yes	
4-hydroxy-8-cis-sphingenine	$C_{18}H_{37}NO_3$	315.2773	-	Yes	
N-(5-aminopentyl)-N'-(5-{[4-({5- [butylidene(oxido)-lambda(5)- azanyl]pentyl}amino)-4- oxobutanoyl](hydroxy)amino}pentyl)- N-hydroxybutanediamide	C ₂₇ H ₅₂ N ₆ O ₇	572.3897	-	Yes	
1,2-di-(9Z-pentadecenoyl)-sn-glycerol	C ₃₃ H ₆₀ O ₅	536.4441	-	Yes	
4,2',4'-Trihydroxy-3',5'- diprenylchalcone	$C_{25}H_{28}O_4$	392.1988	-	Yes	[50]
5,7,4'-Trimethoxyflavan	$C_{18}H_{20}O_4$	300.1362	-	Yes	
all-trans-Heptaprenyl diphosphate	$C_{35}H_{60}O_7P_2\\$	654.3814	-	Yes	
Heliotrine	$C_{16}H_{27}NO_5$	313.1889	-	Yes	
(22E,24R)-Stigmasta-4,22-diene-3,6- dione	$C_{29}H_{44}O_2$	424.3341	-	Yes	
Methyl 9R-hydroxy-10E,12E- octadecadienoate	$C_{19}H_{34}O_3$	310.2508	-	Yes	[51]
1,2-di-(9Z,12Z-octadecadienoyl)-sn- glycerol	$C_{39}H_{68}O_5$	616.5067	-	Yes	
N-(1,3-dihydroxypropan-2- yl)hexadecanamide	$C_{21}H_{30}O_3$	330.2195	-	Yes	
2-Hydroxy-6-tridecylbenzoic acid	$C_{20}H_{32}O_3$	320.2351	-	Yes	$\begin{array}{l} MDA\text{-}MB\text{-}231\\ (IC_{50}=117.25\ \mu\text{M});\\ 4\text{T-1}\ (IC_{50}=102.39\ \mu\text{M})\ [42] \end{array}$
β-isorenieratane	$C_{40}H_{72}$	552.5634	-	Yes	
Matricin	$C_{17}H_{22}O_5$	306.1467	-	Yes	[35]
9,10,13-trihydroxy-octadecanoic acid	$C_{18}H_{36}O_5$	332.2563	-	Yes	
Austroinulin	$C_{20}H_{34}O_3$	322.2508	-	Yes	
Butyl 3- <i>Ο-β-</i> D-glucopyranosyl-butanoate	$C_{14}H_{26}O_8$	322.1628	-	Yes	
Presqualene diphosphate	$C_{30}H_{52}O_7P_2$	586.3188	-	Yes	
2-O-(β-D-galactopyranosyl-(1->6)-β- D-galactopyranosyl) 23,3R-dihydroxynonanoic acid	$C_{21}H_{38}O_{14}$	514.2262	-	Yes	
Catechin 3-O-rutinoside	$C_{27}H_{34}O_{15}$	598.1898	-	Yes	[52]
Stearic acid	$C_{18}H_{36}O_2$	284.2715	-	Yes	[33]
1-O-(2R-methoxy-4Z-heneicosenyl)- sn-glycerol	$C_{25}H_{50}O_4$	414.3709	-	Yes	
Ammothamnidin	$C_{25}H_{28}O_5$	408.1937	-	Yes	
6-Deoxohomodolichosterone	C ₂₉ H ₅₀ O ₄	462.3709	-	Yes	

Table 4. Cont.

Metabolites	Molecular Formula	Accurate Mass	Up-Regulation [Hexane Extract]	Up-Regulation [CHCl ₃ /MeOH Extract]	Biological Activity/ References
1-hexadecyl-glycero-3-phospho- (1' <i>-sn-</i> glycerol)	$C_{22}H_{47}O_8P$	470.3009	-	Yes	
28-Glucopyranosyl-3- methyloleanolic acid	C37H60O8	632.4285	-	Yes	
Stigmastane-3,6-dione	$C_{29}H_{48}O_2$	428.3654	-	Yes	
N-(dodecanoyl)-sphing-4-enine	C ₃₀ H ₅₉ NO ₃	481.4503	-	Yes	
Phyllospadine	$C_{21}H_{21}NO_6$	383.1369	-	Yes	
Tricosylic acid	$C_{23}H_{46}O_2$	354.3498	-	Yes	
1-Docosahexaenoyl-2-erucoyl-sn- glycero-3-phosphocholine	C ₅₂ H ₉₀ NO ₈ P	887.6404	-	Yes	

Table 4. Cont.

2.5. Network Pharmacology (NP)

NP approach was used to explore metabolite/gen/disease interaction in the cancer context. The results displayed the synergist activity of some metabolites to achieve anticancer effect. Some compounds such as presqualene diphosphate, phytol, stearic acid, δ -tocopherol, ursolic acid and γ -linolenic acid are directly involved in the five sub-networks identified. Figure 5A–C shows three interaction networks selected. Figure 5A is about the most noteworthy network by the largest number of concerned nodes. Some key genes identified in this network are recognized for the National Center for Biotechnology Information (NCBI) for their significant role in drug discovery [53]: CASP3 (caspase 3), the protein encoded by this gene plays a central role in the execution-phase of cell apoptosis. In addition, two nuclear receptors (PARP1; NR3C1) involved in the regulation of several important cellular processes such as differentiation, proliferation, and in the recovery of cell from DNA damage. Likewise, three different genes (BAX, BCL2, STAT3) whose encoded proteins were implicated in cell growth and apoptosis. Two well-known signaling molecules (PTPN6 and PTPN3) were also identified, which regulate a variety of cellular processes including cell growth, differentiation, mitotic cycle, and oncogenic transformation. Last, DNA topoisomerase (TOP2A) was also recognized. The gene encoding this enzyme functions as the target for several anticancer agents and a variety of mutations in this gene have been associated with the development of drug resistance [53].

The analysis of the pharmacological network revealed, as targets, several genes involved in the inflammatory response, which occurs in various pathological conditions, such as cancer (CXCL8, ALOX5 and ALOX15) [53]. Phytol (network Figure 5B), targets PPAR α . This gene is implied in cell proliferation, cell differentiation, and immune and inflammation responses. Along with presqualene diphosphate (Figure 5C), phytol targets genes encoding proteins involved in drug metabolism and synthesis of cholesterol, steroids, and other lipids (CYP46A1 and FDFT1, respectively) [53].

Summarizing, the current work presents for the first time the metabolomic fingerprint of *C. incisa* leaves, and the cytotoxic properties of their extracts. Untargeted metabolomics profiles through UHPLC-QTOF-MS and multivariate analyzes allowed to determine the phytochemical similarities and differences between the three extracts and to understand their cytotoxic effects by the presence of bioactive metabolites. The hexane extract achieved remarkable cytotoxicity on hepatocellular cancer cells, hence, coupling the metabolome data with its biological activity could support a targeted isolation focused on the predicted active metabolites. The NP approach used was successful for the interpretation of the experimental results, because the metabolites that contribute to the cytotoxic activity and the molecular pathways involved were revealed.



Figure 5. (A–C). Graphs of the selected interaction networks.

3. Materials and Methods

3.1. Vegetal Material and Extracts Preparation

Cissus incisa (Nutt.) Des Moul. Ex S. Watson was collected in Rayones, Nuevo Leon, Mexico (Latitude: 25.0167°, Longitude: –100.05°, Altitude: 900 m) on 10 October 2016. The identification was made by the biologist Ph.D. Mauricio Gonzalez Ferrara (Autonomous University of Nuevo Leon, San Nicolás de los Garza, Mexico). The collected species were deposited in the herbarium of Biological Sciences Faculty of the Autonomous University of Nuevo Leon with Voucher 027499. The plant name has been checked with http://www.theplantlist.org, accessed on 3 February 2020.

Leaves were dried at room temperature for 2 weeks and then milled until obtaining 809 g of dried and grounded plant material. Sequential macerations were made using hexane (10 L), chloroform/methanol (1:1) (7 L), and water (10 L) yielding the organic extracts and the aqueous extract. The extractions were made at room temperature, following the same steps: filtration and vacuum distillation to dryness for the organic extracts. Whereas, a lyophilization was carried out to obtain the dry aqueous extract, yielding 11.6 g of hexane (1.43%), 84 g of CHCl₃/MeOH (10.38%), and 19.6 g of aqueous one (2.42%) of dry extracts.

3.2. UHPLC-QTOF-MS/MS Analysis

All solvents LCMS grade Baker (Thermo Fisher Scientific, Waltham, MA, USA) were filtered using membrane filter, NYLON 0.45-micron \times 47 mm (DS0215-4045, Thermo Fisher Scientific, Waltham, MA, USA). Three samples per extract were diluted independently (1 mg/mL) in MeOH (50%), sonicated 5 min \times 10,000 rpm and filtered using PTFE 0.20 μ m Syringe filter (721-1320 Thermo Scientific, Waltham, MA, USA), and transferred to a high-recovery MS Analyzed Type 1 borosilicate amber glass vial (5190-7041/5182-0717, Agilent Technologies, Santa Clara, CA, USA).

Reverse-phase liquid chromatography was performed at 20 °C, using an Agilent 1290 Infinity II Ultra High-Performance Liquid chromatography system (UHPLC Waters, Singapore, Singapore) and the column ZORBAX Eclipse Plus C18 HD 2.1 × 50 mm, 1.8 μ m (Agilent Technologies, Santa Clara, CA, USA). The mobile phase was delivered by a binary pump at a flow rate of 0.250 mL/min in a gradient elution using: LCMS grade water + 0.1% v/v formic acid (solvent A) and LCMS grade MeOH + 0.1% v/v formic acid (solvent

B) with the following gradient conditions: 0–6 min, from 30 to 100% solvent B; held at 100% B until 10 min; 10–11 min, from 100 to 30% B to return to original conditions. Injection volume was 5 μ L. Mass spectrometric analysis was performed using an Agilent 6545 Quadrupole Time of Flight (QTOF) LCMS with an electrospray ionization (ESI) source (Agilent Technologies, Waldbronn, Germany), in positive mode. Detection range of mass-to-charge ratio (*m*/*z*) was 100–3000. The nebulizer pressure was set at 35 psi, gas temperature of 320 °C, and a gas flow rate of 8 L/min.

3.3. Data Processing and Metabolic Pathway Analysis

The identification of metabolites was carried out using the METLIN Metabolites Database on Agilent MassHunter Qualitative Analysis B.08.00 software and the lists for data analysis were generated with compounds present in all the replicates of each extract. Putative assignments for each compound were made based on their accurate mass. Additionally the Dictionary of Natural Products, PubChem, (http://pubchem.ncbi.nlm.nih.gov/, accessed on 11 October 2020), LIPID MAPS (http://www.lipidmaps.org/tools, accessed on 11 October 2020), and Human Metabolome Database (HMDB) (http://www.hmdb.ca, accessed on 11 October 2020) were consulted. Principal component analysis (PCA), Venn diagram, and fold change analysis (cut off 2.0) were carried out for UHPLC-QTOF-MS/MS data on Mass Profiler Professional software. PCA presents the average of replicates by each extract. For all statistical tests performed, ANOVA with cut-off p < 0.05 was taken as significant. The metabolomics pathway analysis and the network topological analysis were performed with MetaboAnalyst (http://www.metaboanalyst.ca/, accessed on 3 February 2021) and STITCH (http://stitch.embl.de/, accessed on 3 February 2021). The metabolite-gene-disease interaction network was selected within the MetPA module, through the integration of network topological analysis, interactive network exploration, and functional enrichment analysis.

3.4. Cytotoxic Activity

3.4.1. Cell Lines

The extracts were evaluated for their cytotoxic activity in human cancer cells: PC3 (prostate ATCC[®] CRL-1435), Hep3B (hepatocellular ATCC[®] HB-8064), and HepG2 (hepatocellular ATCC[®] HB-8065), MCF7 (breast (ATCC[®] HTB-22), A549 (lung (ATCC[®] CCL-185), and HeLa (cervical ATCC[®] CCL-2), all were obtained from ATCC (American Type Culture Collection, Manassas, VA, USA). In addition, a cell line of immortalized human hepatocytes (IHH) was included as control of non-cancerous cells [54]. PC3 cells were cultured in RPMI-1640 medium (Sigma Aldrich, St. Louis, MO, USA), while Hep3B, HepG2, IHH, MCF7, A549 and HeLa in DMEM (Invitrogen, Thermo Fisher Scientific, Inc., Waltham, MA, USA) and supplemented with 10% fetal bovine serum (SFB, Invitrogen, Waltham, MA, USA) and with 2 mM glutamine, all cultures were incubated at 37 °C in a 5% CO₂ atmosphere.

3.4.2. IC₅₀ Determination

For the cytotoxic evaluation 4000 cells were cultured per well in 96-well plates. The concentrations used for the extracts and for positive control Paclitaxel were 100, 10, 1, 0.1, $0.01 \mu g/mL$ for a dose/response curve.

Prior to the assay, stock solutions of 20 mg/mL (20,000 μ g/mL) of each extract were prepared, (1 mg of extract dissolved in 50 μ L of DMSO) for organic extracts and sterile water for the aqueous one.

The solutions were prepared from this stock as follows. The concentration 100 μ g/mL was prepared from 2.5 μ L of a stock solution 20 mg/mL (20,000 μ g/mL) in 497.5 μ L of culture medium. The concentration 10 μ g /mL was prepared from 50 μ L of the 100 μ g/mL solution in 450 μ L of medium. The concentration 1 μ g/mL was prepared from 50 μ L of the 10 μ g/mL solution in 450 μ L of medium. The concentration 0.1 μ g/mL was prepared from 50 μ L of the 1 μ g/mL solution in 450 μ L of medium. The concentration 0.1 μ g/mL was prepared from 50 μ L of the 1 μ g/mL solution in 450 μ L of medium. The concentration 0.1 μ g/mL was prepared from 50 μ L of the 1 μ g/mL solution in 450 μ L of medium. The concentration 0.1 μ g/mL was prepared from 50 μ L of the 0.1 μ g/mL solution in 450 μ L of medium.

Subsequently, 100 μ L of each solution was added to its corresponding well. Treatment with extracts did not exceed 0.5% of DMSO. In addition, a solvent control was performed at this concentration, not observing cell growth inhibition, which guaranteed that the cytotoxic activity of each extract was associated with the chemical content present in each extract and not with the solvent.

Plates were incubated at 37 °C in 5% CO₂ atmosphere for 48 h. The number of viable cells in proliferation was then determined by the Cell Titer 96[®] aqueous solution cell proliferation assay kit (Promega, Madison, WI, USA) following the supplier's protocol. Cell viability was determined by absorbance at 450 nm using an automated ELISA reader. The experiments were performed in triplicate in three independent experiments. Data were expressed as means \pm SD and were analyzed in the Prism 5.0 statistical program, IC₅₀ values were determined by regression analysis [54].

3.4.3. Selectivity Index

The extracts were tested against IHH normal cell line [54] to determine the selectivity of the cytotoxic activity on hepatocellular lines. The Selectivity Index (SI) was calculated following previous reports [55]: SI = IC_{50} of extract in a normal cell line/ IC_{50} of the same extract in cancer cell line, where IC_{50} is the concentration required to kill 50% of the cell population.

Supplementary Materials: The following are available online at https://www.mdpi.com/article/ 10.3390/plants10071389/s1, Table S1: UHPLC-QTOF-MS/MS results for Hexane extract, Table S2: UHPLC-QTOF-MS/MS results for chloroform/methanol extract, Table S3: UHPLC-QTOF-MS/MS results for aqueous extract, Table S4: Common compounds in the Hexane and CHCl3-MeOH extracts.

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