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[Luz A. Guerrero-Lagunes](#) , [Lucero M. Ruiz-Posadas](#) ^{*} , [Jorge Cadena-Iñiguez](#) ,
[Ramon Marcos Soto-Hernández](#) , [Carlos H. Avendano-Arrazate](#) , [Juan F. Aguirre-Medina](#) ,
[Celeste Soto-Mendoza](#) , Juan F. Aguirre-Cadena

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Review Paper

Bioprospecting of *Bixa Orellana* L. for the Selection of Characters with Biological Activity

Luz A. Guerrero-Lagunes ¹, Lucero M. Ruiz-Posadas ^{1,*}, Jorge Cadena-Iñiguez ²,
Ramón Marcos and Soto-Hernández ¹, Carlos H. Avendaño-Arrazate ³, Juan F. Aguirre-Medina ⁴,
Celeste Soto-Mendoza ¹ and Juan F. Aguirre-Cadena ⁴

¹ Colegio de Postgraduados, Campus Montecillo. Km. 36.5 Carretera Federal México-Texcoco. Montecillo. Texcoco, Edo. De México. C.P. 56264. México, C.P. 56264

² Colegio de Postgraduados, Campus San Luis Potosí, Innovación en Manejo de Recursos Naturales, Iturbide 73, Salinas de Hidalgo, San Luis Potosí, México, C.P. 78600

³ Instituto Nacional de Investigaciones Forestales, Agrícolas y Pecuarias, Centro Nacional de Recursos Genéticos, Blvd. de la Biodiversidad #400 Rancho las Cruces Tepatitlán de Morelos, Jalisco, México, C.P. 47600

⁴ Faculty of Agricultural Sciences—Campus IV, Autonomous University of Chiapas, Junction of the Coastal Highway and the Town of Huehuetán, Chiapas. C.P. 30660, México

* Correspondence: lucpo@colpos.mx

Abstract: A meta-analysis of 28 sources of information was conducted, considering different variables in *Bixa orellana*, with the aim of identifying bioprospective variables. Variables were approached, such as the organ of extraction, extraction method, 63 biochemical classes, and 20 for biological activity, and their states were codified. The statistical analysis was developed through a cladistics analysis using the WinClada version 1.00.08 84,85 software and the explicative accumulated variance was determined through a descriptive multivariate analysis and multiple correspondence analysis (MCA). The tree obtained showed the genotype Africa 1 as the one closest to the basal state. After Africa 1, 9 clades are derived and the genotypes Colombia3 and Colombia5 were the most evolved. The analyses demonstrated that in *B. orellana* L., the genotypes from India, Brazil and Yucatán present anticancer activity against the cell lines U251, MCF-7, HeLa, NCI-H460, PC-3, A549 and HT-29, as well as biological activity against *Staphylococcus aureus*, *Escherichia coli* and *Pseudomonas aeruginosa*, related primarily with biochemical compounds such as geranylgeraniol, ellagic acid, and carotenoids (bixin and norbixin), naringenin and alkaloids. The conditions of reproductive isolation of the genotypes mentioned before providing the ideal agroclimatic conditions to produce compounds with biological activity.

Keywords: achiote; biological activity; cancer

1. Introduction

Bixa orellana L. (Bixaceae), known as achiote or annatto, is a native shrub from Central and South America, grown in some tropical countries of the world, such as Peru, Mexico, Brazil, Colombia, Ecuador, Indonesia, India, Kenya and Eastern Africa [1]. Both the seeds and the leaves have been used in traditional medicine to treat constipation, fever, stomach acidity, asthma, scabies, ulcers and diarrhea [2].

The seeds, as coloring, condiment and thanks to compounds like bixin and norbixin, are of interest for the food, textile, chemical, pharmaceutical and cosmetic industries [3]. In addition, tannins, flavonoids, phenolic compounds, saponins, alkaloids, terpenoids, anthraquinones and glucosides have been identified [4–7]. The content of bixin, norbixin and geranylgeraniol confers biological activity with therapeutic purposes [8–10].

The carotenoids, apocarotenoids, terpenes, terpenoids, sterols, and aliphatic compounds are the main compounds that are found in every part of this plant, for which a wide range of pharmacological activities have been researched [11]. Their biological activity has been demonstrated for the control of bacteria and fungi [9,12]. The antioxidant activity has been demonstrated by various studies [7,9,13], and also the anticancer activity in cell lines of medical interest [8,10,14,15]; therefore, it has been included among nutraceutical foods. Because of its broad biological activity, *B. orellana* L. is a source for the development of new drugs with pharmacological activity, so there is the possibility of identifying morphological and phytochemical variables with a bioprospective approach in use, under the premise that the bioprospective meta-analysis eases the identification of the genotype, its character or outstanding phytochemical variable, and the state of the character, specifying the statistical validity and reducing possible contradictions in the literature.

2. Materials and Methods

An analysis of the studies published in the Scopus, Science Direct, Scifinder, Springer and Google Scholar databases was carried out, using the search terms achiote, *B. orellana* L., phytochemicals, pharmacology, cancer, biological activity, antibacterial activity, anticancer, cytotoxic and antioxidant activity. From this, n=56 results were identified, and when criteria of plant organ and biological activity identified in each publication were applied, the sample was reduced to n=28. All the studies included were studies that addressed the phytochemical characterization and biological activity of extracts from *B. orellana* L. (Table 1).

The studies included were conducted in Africa (n=1), United States (n=1), Philippines (n=1), Ecuador (n=1), Bangladesh (n=1), South Korea (n=1), Nigeria (n=2), Yucatán Mexico (n=2), Colombia (n=5), India (n=5), and Brazil (n= 6). The last three led the phytochemical and biological activity research of *B. orellana* L. A database was elaborated with the information, codifying the variables and their different states (Table 2), made up by the following: organ of the plant used, extraction methods, biochemical classes, groups of compounds, phenols and phenolic acids, flavonoids, tannins, monoterpenes, sesquiterpenes, diterpenes, triterpenes, tetraterpenes, alkaloids, cyanogenic glucosides, antimicrobial and anticancer activity.

Table 1. Publications included in the bioprospecting meta-analysis in *Bixa orellana* L.

Variable	Genotype	Research focus	References
Biological activity	África1; USA1; Colombia1; Philippines1; India1; Colombia2; Colombia3; India2; Nigeria1; South Korea1; India3; Colombia4; Philippines2; Yucatán1; Brazil1; Bangladesh1; Brazil2; Colombia5; Brazil3; India4; Indonesia1; Brazil5; India5; Yucatán2; Ecuador1	Antimicrobial, anticancer, antioxidant and hepatoprotective activity of leaves and/or seeds of <i>Bixa orellana</i>	[1,4,6–9,11–27]
	África1; USA1; Colombia1; Philippines1; India1; Colombia2; Colombia3; India2; Nigeria1; South Korea1; Nigeria2; India3; Colombia4; Philippines2; Yucatán1; India4; Colombia5; Brazil3; India5; Yucatán2; Ecuador1	Phytochemical characterization of extracts and essential oil of leaves and seeds of <i>Bixa orellana</i>	[1,4–28]
Biochemistry			

Table 2. Characters and character states of *Bixa orellana* L., for the bioprospecting analysis.

Number	Variable	Variable status
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1	Extraction organ	Leaves=1, Seeds=2
2	Extraction method	Absent=0, Aqueous and ethanolic extracts=1, Solvent system=2, Methanol=3, Ethanol=4, Petroleum ether=5, Maceration=6, Steam Distillation=7
3	Biochemistry class	Absent=0, Phenolic compounds=1, Terpenoids=2, Compounds with nitrogen=3
4	Compound group	Phenols and phenolic acids=1, Flavonoids=2, Tannins=3, Monoterpenoids and sesquiterpenoids=4, Diterpenes=5, Triterpenoids=6, Tetraterpenoids=7, Alkaloids=8, Cyanogenic glycosides=9
5	Phenols and phenolic acids	Absent=0, Phenylpropanoids=1, Coumarins=2, Anthraquinones=3, Procyanidins=4, Ellagic acid=5, Tannic acid=6, Gallic acid=7
6	Flavonoids	Absent=0, Naringenin=1, Kaemferol=2, Anthocyanins=3, Isoflavonoids=4, Butein=5, Catechins=6, Chlorogenic acid=7, Hypolaetin=8
7	Tannins	Absent=0, Granatin=1, Neostriatin=2, Ellagitanin=3
8	Monoterpenes	Absent=0, Poliprenol=1, Ocimene=2, Spathulenol=3, Isolatedene=4, Bergamotene=5, Pinene=6, Aristolochene=7, Cadinene=8, Germacrene=9
9	Sesquiterpenes	Absent=0, Farnesol=1, Elemene=2, Caryophyllene=3, Guaiaol=4, Tomentosin=5, Ishwarane=6
10	Diterpenes	Absent=0, Phytol=1, Geranylgeraniol=2, Geranyl terpinene=3, Geranyl linalool=4, Farnesyl=5
11	Triterpenes	Absent=0, Saponins=1, Steroids=2, Stigmasterol=3, Sitosterol=4, Squalene=5
12	Tetraterpenes	Absent=0, Carotenoids=1, 9'-cis-norbixin=2, Trans-norbixin=3, Bixin=4, Norbixin=5, Diapocarotenoids=6
13	Alkaloids	Absent=0, Atrophin=1
14	Cyanogenic glycosides	Absent=0, Saponins=1
15	Biological activity	Absent=0, Chemo preventive=1, Anti-inflammatory=2, Hepatoprotective=3, Antioxidants=4, Cytotoxic=5
16	Antimicrobial activity	Absent=0, <i>Pseudomonas aeruginosa</i> =1, <i>Escherichia coli</i> =2, <i>Staphylococcus aureus</i> =3, <i>Salmonella</i> sp=4, <i>Candida albicans</i> =6
17	Anticancer activity	Absent=0, HepG2=1, U251=2, MCF-7=3, HeLa=4, NCI-H460=5, PC-3=6, HT-29=7, A549=8, MCF-7=9

Statistical Analysis

The statistical analysis was developed with two approaches. The first through a cladistics analysis that incorporates the approach of Popper's critical rationalism through the refutation of phylogenetic hypotheses examined under a parsimonious principle [29,30]; and through non-parametric statistics using the WinClada version 1.00.08 84,85 software (free license) [31]; with the Bootstrap/Jackknife resampling methods, approaching the genotypes as populations through a random simulation until generating a parsimonious cladogram [30]. This analysis defines the stability of the clades and identifies the state of the outstanding variables. The analysis was repeated 1000 times creating the values such as support indices, consistency, and reliability in the cladograms [32]. The systematic reviews carried out in the meta-analysis were directed towards the information disseminated, to reanalyze it with approaches adapted to the present research [33]. It must be clarified that the criteria selected were those with complete, traceable data, and reproducible results to avoid biases in the study [34].

The second approach was to determine the explicative accumulated variance, the statistical weight of each variable, and its state through a descriptive multivariate analysis and multiple

correspondence analysis (MCA), with the FactoMineR [35] and factoextra [36] libraries with the Rstudio statistical package [37].

3. Results and Discussion

Figure 1 shows the different organs of *B. orellana* L. that could have an impact on the identification of the various genotypes, while Figure 2 presents the general cladogram that indicates the distribution of the *B. orellana* L. genotypes analyzed in function of the characters organ of extraction, method, biochemical class, and biological activity (Table 2). In total, 12 trees were obtained to get a consensus tree. This tree showed 149 steps or changes, a consistency index of the cladogram of 50%, and a retention index that reflects the percentage of characters that retain and conserve the change of taxa of 64%.



Figure 1. Leaf: (a), flower (b), capsule (c), and seed (d) of *Bixa Orellana* L.

The parsimonious distribution of the genotypes (Figure 2) is not indicative of a strict genealogical relationship, since there are no morphological and genetic characters; however, it helps to understand the adaptive specialization [38,39] of plants in face of the difference in environmental conditions different from those in their habitat. In general, reproductive isolation, selective pressure, and lack of variability create unique survival characters reflected in the content and diversity of secondary metabolites [40].

The biochemical and biological activity variables showed that the genotype with origin from Africa1 was located as the closest one to the basal state, hypothetically indicating due to the variables analyzed that it could have greater similarity with a genotype from the original habitat (Figure 1).

Nine clades derive from Africa1. The first formed by Nigeria2 and India3, genotypes closer to the root, which share the presence of alkaloids as plesiomorphic characters. Even when the publications do not record the time of introduction to Nigeria and India, it is presumed that they could have had some reproductive isolation, absence of variability, and agroclimatic conditions

different from their geographic origin (Central and South America). Various authors mention how reproductive isolation and absence of biological variation in some organisms promotes unique characters that can be used in different sectors of society, such as the case of enzymes responsible to produce secondary metabolites with biological effect of medical, agricultural or industrial interest [41].

The second clade derives starting with India3, formed by genotypes USA1, India4, Brazil2 and Brazil3, which made up an independent evolutionary route characterized by sharing the seed as organ of extraction and which is a derived state.

The USA1 genotype shares with the rest of the genotypes from this group the presence of geranylgeraniol (apomorphic state), highlighting that it has the derived characters *cis*-norbixin and *trans*-norbixin, which are related with the biological activity against *Staphylococcus aureus*, also a derived state. Brazil2 and Brazil3 are sibling genotypes, and presumed to be those of greatest “evolved” specialization from this group, characterized by the presence of flavonoids that is classified as an ancestral state. When it comes to apomorphic states, the presence of terpenes, ocimene, spathulenol, isodene, bergamotene stands out, as well as bixin and norbixin.

The genotype Brazil3 shows anticancer activity against the cell lines U251, MCF-7, NCI-H460, PC-3 and HT-29 and presents as a plesiomorphic state (ancient or primitive character). The distinction of the proliferative activity in a genotype that is in the origin center of the species proves that the specific conditions of this place favored the presence of the compounds mentioned and contributed to the anticancer activity. However, when agroclimatic conditions change outside its place of origin, it loses chemical variability.

In the case of Ecuador1, it forms an independent clade and shows an evolutionary divergence possibly due to reproductive isolation. A plesiomorphic state stands out in the group, which is the extraction method by vapor sweeping, different from the rest of the genotypes, while the presence of ocimene, pinene, germacrene, farnesol and caryophyllene, as well as the activity against *Staphylococcus aureus*, are new characters or derived states (apomorphic). In this group, the extraction method marked the difference in the compounds detected.

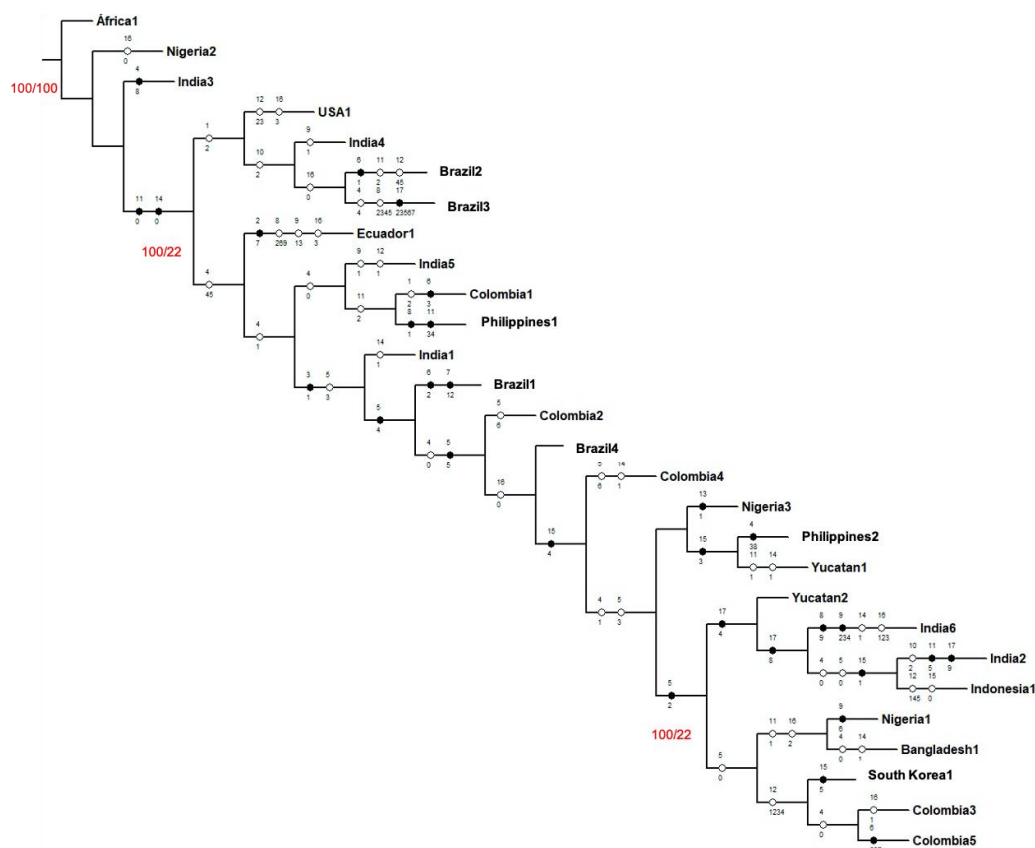


Figure 2. Cladogram of genotypes of *Bixa orellana* L. with different geographic origins, based on the plant organ used, extraction method, biochemical characters, and biological activity. White spots represent apomorphic variations and black spots plesiomorphic variations. The values separated by the diagonal line represent the Bootstrap/Jackknife indices, with L= 149, Ci= 50 and Ri=64.

The six remaining clades derive from Ecuador1. India5, Colomba1 and Philippines1 form a group characterized by the new or derived characters represented by the farnesol compounds, saponins and carotenoids. The genotypes from Colombia and Philippines present three ancestral characters constituted by anthocyanins and polyprenol, as well as stigmaesterol and sitoesterol. Although both genotypes do not have a geographic grouping, they do present it by group of compounds, which indicates a possible displacement of the genotypes from the center of origin towards the Philippines and India, where the original compounds could be conserved, or else, there was an influence of similar agroecological conditions that impacted the production of these secondary metabolites.

Also, Ecuador1 is derived from the group constituted by India1, Brazil1, Colombia2, Brazil4 and Colombia4. It should be mentioned that the genotypes from Brazil are again those that present the highest number of plesiomorphic states (kaempferol, granatin, neostictinin, procyanidines, ellagic acid, as well as antioxidant activity). When it comes to apomorphic characters, the presence of saponins, tannic acid, and anthraquinones stands out, which are present in the genotypes from Colombia and India. In this group, the geographic grouping is clear regarding the states that highlight the ancestral characteristics of the genotypes from Brazil, zone registered as origin center of *B. orellana*.

Brazil4 is attributed with antioxidant activity and can be linked to the presence of ellagic acid, situation that agrees with other [42], who determined that this compound acts as chemo-protector against different types of cancer and which shows strong anti-proliferative activity against colon, lung and prostate cancer cells.

From Colombia4, 4 subgroups are derived, an independent one formed by Nigeria3, Philippines2 and Yucatán1 which are characterized by presenting three plesiomorphic states represented by tannins, alkaloids and atropines.

From this group, the genotype located in Yucatán is the most evolved and it is proven by the derived states present, such as the presence of saponins and the hepatoprotective activity. The evolution can be due to the agroclimatic characteristics or the manipulation of the crops in the zone, since in Yucatán there are commercial crops of *B. orellana* L. that have been genetically improved to reach higher seed production, which can be a factor that impacts the production of secondary metabolites [43].

The other sibling arm of Colombia4 groups, on the one hand, Yucatán2, India6, India2 and Indonesia1, which, despite not having a geographic grouping was characterized by the presence of the highest number of plesiomorphic states, among which germacrene, elemene, caryophyllene and squalene stand out, as well as chemo-preventive activity.

The anticancer activity against the cell lines HeLa, A549 and MCF-7 is a simplesiomorphic state, since it also presents in Brazil3, which is a genotype close to the root.

The derived character has to do with the presence of geranylgeraniol, carotenoids, bixin and norbixin, in addition to the activity against *P. aeruginosa*, *E. coli* and *S. aureus*. The biological and anticancer activity is determined by the variety of phytochemical compounds present in *B. orellana* L. and by the capacity of geranylgeraniol to induce apoptosis in A549 cells [26,44,45]. In this group, it can be inferred that there was a flow of plants from Yucatán towards India and Indonesia and the antiproliferative activity was conserved.

The last two groups derive from the branch coming from Colombia4. The node formed by Nigeria1 and Bangladesh share the apomorphic states represented by saponins and the biological activity against *E. coli*. Only an ancestral state is present (tomentosin).

The genotypes SouthKorea1, Colombia3 and Colombia5 are the last group and share four apomorphic states integrated by carotenoids, cis-norbixin, trans-norbixin and bixin. In addition, they present butein, catechins and chlorogenic acid, as well as cytotoxic activity as ancestral characters. The genotypes located in Colombia were considered the most evolved, compared to Africa1, Nigeria2 and India3, whose evolution can be due to pressure processes, such as the manipulation, edaphoclimatic conditions, or the genetic flow between genotypes. It should be highlighted that the activity found in the genotypes present in this bioprospective study is consistent with that found authors [1,46,47], who determine that the tannins, quinones and terpenoids have biological activity; in addition, lipophylic flavonoids can be disruptive for the cell membrane [48].

Table 3 shows the apomorphic characters present in the genotypes studied, observing sinapomorphic characters (shared characters) among the genotypes USA1, India2 and India4, such as geranylgeraniol, while India4 and India5 share farnesol; Brazil2 and India5, steroids; Colombia2 and Colombia4, tannic acid; Colombia4, India6 and Bangladesh1, saponins; India5 and Indonesia1, carotenoids; Brazil2 and Indonesia1, bixin and norbixin; Brazil3 and Ecuador1, ocimene; USA1 and Ecuador1, activity against *S. aureus*; and India6 and Colombia3 activity against *P. aeruginosa*. On the other hand, cis-norbixin, trans-norbixin, spathulenol, isodene, bergamotene, germacrene, farnesol and caryophyllene are autapomorphic states (unique characters) because they are present in a single taxon or genotype.

Table 4 also shows that apomorphic characters (new or derived) are related with the antimicrobial activity against *P. aeruginosa*, *E. coli*, *S. aureus* and biochemical class, primarily the carotenoids bixin, norbixin, 9'-cis-norbixin, transnorbixin, saponins and monoterpenes. The plesiomorphic characters are related more with the hepatoprotective, chemo preventive and anticancer activity against the cell lines MCF-7, NCI-H460, PC-3, HT-29, HeLa and A549; as well as the presence of flavonoids (naringenin, kaempherol, anthocyanins, procyanidines, ellagic acid), triterpenes (stigmasterol and sitoesterol), tannins (granatin, neostriectinin), sesquiterpenes (elemene, caryophyllene) and coumarins. It stands out that some compounds from *B. orellana* in this bioprospective analysis act on microorganisms that can cause public health problems, highlighting characters for a possible program for genetic improvement.

Table 3. Apomorphic characters (new or derived) from the heuristic clade with biochemical and biological activity variables in 28 genotypes of *Bixa orellana* L. The diagonal indicates the character and its state of relevant character.

Genotype	Character/character state	Biochemistry	Biological activity/antimicrobial activity/anticancer activity
Nigeria2			
India3			
USA1	10/2, 12/2, 12/3, 16/3,	Geranylgeraniol, cis-norbixin, trans-norbixin	<i>Staphylococcus aureus</i>
India4	9/1, 10/2,	Farnesol, geranylgeraniol	
Brazil2	11/2, 12/4, 12/5	Steroids, bixin, norbixin	
Brazil3	4/4, 8/2, 8/3, 8/4, 8/5	Mono and sesquiterpenoids, ocimene, spathulenol, isodene, bergamotene	
Ecuador1	8/2, 8/6, 8/9, 9/1, 9/3, 16/3	Ocimene, pinene, germacrene, farnesol, caryophyllene	<i>Staphylococcus aureus</i>
India5	9/1, 11/2, 12/1	Farnesol, saponins, carotenoids	

Colombia1	11/2	Saponins	
Philippines1			
India1	14/1	Saponins	
Brazil1			
Colombia2	5/6	Tannic acid	
Brazil4			
Colombia4	5/6, 14/1	Tannic acid, saponins	
Nigeria3			
Philippines2			
Yucatán1	11/1	Saponins	
Yucatán2			
India6	14/1, 16/1, 16/2, 16/3	Saponins	<i>Pseudomonas aeruginosa</i> , <i>Escherichia coli</i> , <i>Staphylococcus aureus</i>
India2	10/2	Geranylgeraniol	
Indonesia1	12/1, 12/4, 12/5	Carotenoids, bixin, norbixin	
Nigeria1			
Bangladesh1	14/1	Saponins	
South Korea1			
Colombia3	16/1		<i>Pseudomonas aeruginosa</i>
Colombia5			

Table 4. Apomorphic and plesiomorphic characters identified in the branches of the heuristic clade based on biochemical variables of biological activity of *Bixa orellana* L. The diagonal indicates the character and its state of relevant character.

Branch	Apomorphic character	Branch	Plesiomorphic character
1	Seeds, mono and sesquiterpenoids, ocimene, spathulenol, isodene, bergamotene, farnesol, saponins, geranylgeraniol, cis-norbixin, trans-norbixin, bixin, norbixin, <i>Staphylococcus aureus</i>	1	Alkaloids Naringenin, U251, MCF-7, NCI-H460 PC-3 and HT-29 cell lines
2	Phenols and phenolic acids, mono and sesquiterpenoids, diterpenes, ocimene, pinene, germacrene, farnesol, caryophyllene, <i>Staphylococcus aureus</i>	2	Steam distillation
3	Seed, phenols and phenolic acids, farnesol, steroids, carotenoids	3	Anthocyanins, phenylpropanoids, stigmasterol, sitosterol
4	Phenols and phenolic acids, anthraquinones, tannic acid, saponins	4	Phenolic compounds, procyanidins, ellagic acid, kaempferol, granatin, neostictinin, antioxidant
5	Saponins	5	Tannins, alkaloids, atrophin, hepatoprotective
6		6	

7	Geranylgeraniol, carotenoids, bixin, norbixin, <i>Pseudomonas aeruginosa</i> , <i>Escherichia coli</i> , <i>Staphylococcus aureus</i>	7	Coumarins, germacrene, elemene, caryophyllene, squalene, chemo preventive, HeLa cell lines, A549, MCF-7
8	Saponins, <i>Staphylococcus aureus</i>	8	Coumarins, caryophyllene
9	Carotenoides, cis-norbixin, trans-norbixin, bixin, <i>Pseudomonas aeruginosa</i>	9	Coumarins, butein, catechins, chlorogenic acid, cytotoxic

It is important to highlight that the non-detection of a compound does not mean it is absent, since it could have been undetected because of the plant organ used, extraction method, or seasonal time of sample harvesting. For future studies, the proposal is to elucidate the “absent” compounds to secure the grouping; something to remember is that since it is a meta-analysis, comparative biases between taxa can be originated due to the various methods of sampling, extraction and analysis. Over-studied and under-studied taxa can bring biases to the analysis.

Multivariate Analysis

The multivariate analysis allowed identifying the variables that explain the highest explicative variance, in addition to exploring the correlations and reducing the dimension of the analysis with new indices (Córdoba et al., 2012). It was determined that in four principal components (PC), the accumulated value is 86.31% (Table 5).

Table 5. Characteristic values and proportion accumulated for four principal components of the analysis of 28 genotypes of *Bixa orellana* L. with different geographic origins, based on the organ of the plant used, extraction method, biochemical characters, and biological activity.

PC	Eigenvalues	Variance	Cumulative variance	%
1	0.1046	0.0963	0.0963	9.63
2	0.0963	0.0887	0.1851	28.14
3	0.0717	0.0727	0.2578	53.92
4	0.0717	0.0660	0.3239	86.31

Table 6 indicates the variance by dimension that suggests that the antiproliferative activity is statistically significant for the cell lines U251, MCF-7, HeLa, PC-3, NCI-H460 and HT-29 in CP1 and CP4, where the presence of atropine, bixin, norbixin, naringenine, anthraquinones, alkaloids, coumarins, phenylpropanoids, tannins and some triterpenes is also significant, which suggests a relationship between the presence of some of these compounds or their synergy for *B. orellana* to present biological activity against the cell lines mentioned and some bacteria. In the rest of the dimensions, no statistical significance is present for the anticancer activity.

Table 6. Characteristic vectors of the analysis of 28 genotypes of *Bixa orellana* L. with different geographic origins, based on the organ of the plant used, the extraction method, bio-chemical characters and biological activity.

Variable	Variable states	CP1	CP2	CP3	CP4
Extraction organ	Leaves	0.40600	0.02947	0.00653	0.00442
	Seeds	0.16010	0.09092	0.00378	0.02480
Extraction method	Aqueous and ethanolic extracts	0.08108	0.00111	0.04358	0.00607
	Solvent system	0.00741	0.00181	0.14477	0.01652
	Methanol	0.00011	0.01826	0.00144	0.00843
	Ethanol	0.07360	0.03881	0.00227	0.04915

	Petroleum ether	0.01204	0.00217	0.00390	0.01648
	Maceration	0.00519	0.29127	0.06014	0.00480
	Steam distillation	0.00519	0.52403	0.09163	0.00072
Biochemistry class	Phenolic compounds	0.00470	0.01508	0.00001	0.00259
	Terpenoids	0.18442	0.20710	0.05018	0.17695
	Compounds with nitrogen	0.01603	0.00538	0.00219	0.16758
Compound group	Phenols and phenolic acids	0.11318	0.00859	0.00795	0.01836
	Flavonoids	0.29988	0.12431	0.16761	0.13349
	Tannins	0.26420	0.12683	0.20160	0.15101
	Monoterpenoids and sesquiterpenoids	0.00000	0.00000	0.00000	0.00000
	Diterpenes	0.00000	0.00000	0.00000	0.00000
	Triterpenoids	0.18557	0.06410	0.08623	0.09697
	Tetraterpenoids	0.04881	0.00190	0.00214	0.01425
	Alkaloids	0.26420	0.12683	0.20160	0.15101
	Cyanogenic glycosides	0.00000	0.00000	0.00000	0.00000
Phenols and phenolic acids	Phenylpropanoids	0.15877	0.06654	0.12130	0.13140
	Coumarins	0.21741	0.02699	0.00046	0.35100
	Anthraquinones	0.26599	0.07780	0.04298	0.14762
	Procyanidins	0.00656	0.00017	0.00004	0.06603
	Ellagic acid	0.00002	0.00002	0.00005	0.02223
	Tannic acid	0.00108	0.00116	0.00310	0.04040
	Gallic acid	0.00000	0.00000	0.00000	0.00000
Flavonoids	Naringenin	0.40034	0.02715	0.02587	0.09716
	Kaemferol	0.00272	0.00312	0.00331	0.10585
	Anthocyanins	0.00059	0.00027	0.01002	0.00397
	Isoflavonoids	0.00000	0.00000	0.00000	0.00000
	Butein	0.00013	0.00063	0.00117	0.08565
	Catechins	0.00000	0.00943	0.00020	0.03638
	Chlorogenic acid	0.01195	0.00235	0.03611	0.05006
	Hypolaetin	0.00000	0.00000	0.00000	0.00000
Tannins	Granatin	0.04169	0.03330	0.06398	0.01294
	Neostriatinin	0.00000	0.00000	0.00000	0.00000
	Ellagitanin	0.00000	0.00000	0.00000	0.00000
Monoterpenes	Poliprenol	0.04622	0.01371	0.20700	0.28666
	Ocimene	0.00098	0.77755	0.17288	0.00777
	Spathulenol	0.00098	0.77755	0.17288	0.00777
	Isoledene	0.00000	0.00000	0.00000	0.00000
	Bergamote	0.00000	0.00000	0.00000	0.00000
	Pinene	0.00098	0.77755	0.17288	0.00777

	Aristolochene	0.00000	0.00000	0.00000	0.00000
	Cadinene	0.00000	0.00000	0.00000	0.00000
	Germacrene	0.00098	0.77755	0.17288	0.00777
Sesquiterpenes	Farnesol	0.02186	0.38639	0.25792	0.23356
	Elemene	0.00098	0.77755	0.17288	0.00777
	Caryophyllene	0.00098	0.77755	0.17288	0.00777
	Guaiol	0.00000	0.00000	0.00000	0.00000
	Tomentosin	0.00000	0.00000	0.00000	0.00000
	Ishwarane	0.00031	0.02955	0.35986	0.01434
Diterpenes	Phytol	0.00247	0.01756	0.09808	0.08027
	Geranyl geraniol	0.00183	0.00469	0.34644	0.04852
	Geranyl terpinene	0.01807	0.02596	0.47671	0.14103
	Geranyl linalool	0.00011	0.01908	0.33076	0.00082
	Farnesyl	0.00011	0.01908	0.33076	0.00082
Triterpenes	Saponins	0.10829	0.00292	0.00052	0.02278
	Steroids	0.00186	0.01413	0.00282	0.00832
	Stigmasterol	0.10291	0.07116	0.16746	0.11540
	Sitosterol	0.19469	0.00139	0.00820	0.14306
	Squalene	0.12040	0.00217	0.00390	0.01648
Tetraterpenes	Carotenoids	0.40393	0.02718	0.03930	0.09930
	9'-cis-norbixin	0.01080	0.00040	0.00393	0.01509
	Trans-norbixin	0.01080	0.00040	0.00393	0.01509
	Bixin	0.48865	0.04809	0.03452	0.02605
	Norbixin	0.48865	0.04809	0.03452	0.02605
	Diapocarotenoids	0.00000	0.00000	0.00000	0.00000
Alkaloids	Atrophen	0.21741	0.02699	0.00046	0.35100
Cyanogenic glycosides	Saponins	0.00000	0.00000	0.00000	0.00000
Biological activity	Chemo-preventive	0.00568	0.00568	0.02083	0.06208
	Anti-inflammatory	0.00000	0.00000	0.00000	0.00000
	Hepatoprotective	0.00544	0.00644	0.00785	0.02204
	Antioxidants	0.02597	0.05981	0.07794	0.05050
	Cytotoxic	0.00000	0.00000	0.00000	0.00000
Antimicrobial activity	<i>Pseudomonas aeruginosa</i>	0.16510	0.00000	0.14469	0.08732
	<i>Escherichia coli</i>	0.08361	0.01917	0.35559	0.02388
	<i>Staphylococcus aureus</i>	0.05017	0.21822	0.00890	0.03508
	<i>Salmonella sp</i>	0.13572	0.04070	0.02404	0.00446
	<i>Candida albicans</i>	0.17932	0.01001	0.03820	0.13449
Anticancer activity	HepG2	0.00000	0.00000	0.00000	0.00000
	U251	0.45618	0.02193	0.05048	0.22606
	MCF-7	0.47323	0.02313	0.03783	0.16672

HeLa	0.43024	0.00843	0.00906	0.00554
NCI-H460	0.04561	0.02193	0.05048	0.22606
PC-3	0.45618	0.02193	0.05048	0.22606
HT-29	0.45618	0.02193	0.05048	0.22606
A549	0.00048	0.00056	0.03869	0.15758
B16F10	0.00000	0.00000	0.00000	0.00000

The values highlighted in bold letters are statistically significant by principal component.

4. Conclusions

There is scientific evidence for the use of *B. orellana* L. as agent with anticancer activity, primarily against the cell lines U251, MCF-7, HeLa, NCI-H460, PC-3, A549 and HT-29, as well as biological activity against *S. aureus*, *E. coli* and *P. aeruginosa*. The antimicrobial and anticancer activity is related primarily with biochemical compounds such as geranylgeraniol, ellagic acid, carotenoids (bixin and norbixin), naringenin, and alkaloids. The conditions of reproductive isolation of the genotypes from Brazil, Yucatán, India and Indonesia provided the ideal agroclimatic conditions to produce compounds with biological activity because they produce those metabolites. This analysis can be used as reference for additional studies, genetic improvement programs, and revaluation of the species.

Author Contributions: For research articles with several authors, a short paragraph specifying their individual contributions must be provided. The following statements should be used “Conceptualization, JCI, and LMRP; methodology, JCI, LAGL; software, JFAM; validation, RMSH. and IAS; formal analysis, ESO; investigation, CHAA, CSM; writing—original draft preparation, JCI, LAGL.; writing—review and editing, JCI, LMRP, LALG. All authors have read and agreed to the published version of the manuscript.” Please turn to the CRediT taxonomy for the term explanation. Authorship must be limited to those who have contributed substantially to the work reported.

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