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# Larvicidal Activity of *Beauveria bassiana* Extracts against *Aedes aegypti* and Identification of Beauvericins

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*Beauveria bassiana* is an entomopathogenic fungus that has been well known for its capacity to act as biopesticide on various disease vectors. The analysis of organic extracts of strains CG71 and UNI40 led to identification of cyclodepsipeptides beauvericin, beauvericin A or F, beauvericin E and bassianolide by ultra-high pressure liquid chromatography-high resolution mass spectrometry in tandem mode (UHPLC-HRMS/MS). The larvicidal activity on 3<sup>rd</sup> instar of *Aedes aegypti* revealed LC<sub>50</sub> 0.9887 and 0.4653 ppm in 24 and 48 hours (CG71 methanolic extract), LC<sub>50</sub> 0.7834 ppm in 48 hours (CG71 ethyl acetate), LC<sub>50</sub> 0.7834 and 1.8149 ppm (UNI 40 for ethyl acetate and methanolic extracts, respectively) in 48 hours. These findings highlight the potential of *B. bassiana* metabolites for controlling the vector of Dengue and Zika diseases.

Keywords: cyclohexadepsipeptides, Dengue, Zika, beauvericins, UHPLC-HRMS/MS

# Introduction

The entomopathogenic *Beauveria bassiana* has a broad host range naturally attacking insects and variety of arthropod species.<sup>1-3</sup> Due to this ability, it is used as a biopesticide agent on diverse disease vectors, including crop pests, ticks and mosquitoes.<sup>4.5</sup> The attack on insect cuticle occurs after hyphae of *B. bassiana* penetrate the insect integument by enzymatic action.<sup>6</sup>

Species of entomopathogenic fungi have been described as producers of secondary metabolites, including bioactive non-ribosomal peptides and polyketides.<sup>7</sup> *B. bassiana* produces the cyclooligomer nonribosomal depsipeptides beauvericins and bassianolide,<sup>8.9</sup> beauveriolides,<sup>10</sup> tenellin, bassianin,<sup>11</sup> pyridovericin, pyridomacrolidin,<sup>12</sup> oosporein<sup>13</sup> and bassiacridin.<sup>14</sup>

In special, beauvericin is an ionophoric cyclodepsipeptide

detected in several fungi, mainly *Beauveria*, *Paecilomyces* and *Fusarium* species.<sup>15-17</sup> It forms complexes with cations, causing an increase in permeability of natural and artificial membranes.<sup>18,19</sup> Furthermore, beauvericin induces programmed cell death similar to apoptosis.<sup>20</sup> Initially it was regarded as a toxin against brine shrimp,<sup>8</sup> also showing insecticidal and antibiotic activities.<sup>21-23</sup> It contains three residues each D-2-hydroxyisovaleric acid (Hiv) and L-*N*-methylphenylalanine linked alternately.

Recently, new analogues of beauvericin were described, including beauvericin A, B,<sup>24</sup> D, E and F.<sup>25</sup> Bassianolide is another cyclooctadepsipeptide metabolite produced *B. bassiana* and *Lecanicillium lecanii*.<sup>9</sup> It is toxic to insects and probably also acts as an ionophore, such as other cyclodepsipeptides.<sup>15</sup> The structural diversity of metabolites from entomopathogenic fungi, associated with diverse biological activities, stimulates the research by on bioactive fungal extracts for controlling insect pests and disease vectors.

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Dengue is a viral infection transmitted to humans by *Aedes aegypti* mosquitoes that became a major concern of the public health sector.<sup>26</sup> There were 96 million apparent dengue infections globally in 2010, of which the Americas contributed 14% (13 million infections) of worldwide and over half developed in Brazil and Mexico.<sup>27</sup> Due to no effective antiviral agents and licensed vaccine for infection, the decrease in transmission depends only on vector control by elimination of artificial and disposable water inundated larvae breeding sites and application of insecticides.<sup>27,28</sup> Furthermore, studies reveal resistance in the vector population to various substances, in Brazil there is an irregular distribution of insecticide resistance.<sup>28</sup>

Furthermore, *Aedes* mosquitoes also transmit Zika virus; attracting worldwide attention in 2016, when the vector caused widespread epidemic cases in Brazil.<sup>29</sup> This disease has been recognized in Brazil since late 2014, but in 2015 coincided with an increase in the number of cases of microcephaly. The more affected Brazilian states were Bahia, Paraíba, and Pernambuco, where in the first trimester of pregnancy coincided with reports of cases of febrile and allergy compatible with Zika virus disease, suggesting an association between Zika virus infection during early pregnancy and the occurrence of microcephaly.<sup>30</sup>

Consequently, there is an urgent need to develop new methods for *A. aegypti* mosquitoes infections control. So, fungal extracts are an alternative because they constitute a rich source of bioactive metabolites.

The paper reports the activity of two strains of *Beauveria* bassiana extracts on mortality of *Aedes aegypti* larvae and identification of beauvericins and bassianolide using ultra-high pressure liquid chromatography-high resolution mass spectrometry in tandem mode (UHPLC-HRMS/MS). Additionally, the fragmentation data of beauvericin E and A of F is provided.

# Experimental

#### Origin and preservation of B. bassiana

The entomopathogenic fungal strains *B. bassiana* CG 71 and UNI 40 were kindly provided by Prof Dr Luis Francisco Angeli Alves from Universidade Estadual do Oeste do Paraná (Cascavel, PR, Brazil) and Prof Dr Pedro Manuel Oliveira Janeiro Neves from Universidade Estadual de Londrina (Londrina, PR, Brazil), respectively. The CG 71 and UNI 40 strains are preserved in the culture collection of Universidade Estadual de Londrina, respectively.<sup>31</sup> Both strains are also preserved at -80 °C in the Coleção Micológica de Lavras, Universidade Federal de Lavras

(Lavras, MG, Brazil) under the accession numbers CML 2677 (CG 71) and CML 2678 (UNI 40).

#### Morphological characterization

The strains were preliminary identified as *Beauveria bassiana* based on the macro and micromorphological characteristics of colonies, conidia and conidiophores after cultivation for 7 days at room temperature on malt extract agar (Himedia, Mumbai, India).<sup>32</sup>

#### DNA sequencing and phylogenetic analyses

Strains CG 71 and UNI 40 were cultivated on malt extract agar for five days at 24 °C. The mycelium was scrapped from the colonies, macerated under liquid nitrogen, and subjected to genomic DNA extraction with the Wizard Genomic Deoxyribonucleic Acid (DNA) Purification Kit (Promega, Madison, USA). A fragment of the second largest subunit of the ribonucleic acid (RNA) polymerase gene (RPB2) was amplified from the extracted DNA using the primers 5F and 7cR, and the polymerase chain reaction (PCR) conditions of Liu *et al.*<sup>33</sup> The purified PCR products were sent for DNA sequencing by a commercial service.

Consensus sequences were assembled using SeqAssem ver. 07/2008 (SequentiX - Digital DNA Processing, Germany). Blast searches against the GenBank database suggested that both strains belonged to Beauveria bassiana. DNA sequences of RPB2 from reference strains of Beauveria bassiana and related taxa<sup>34</sup> were used to compose a multiple sequence alignment using multiple sequence comparison by log-expectation (MUSCLE), as implemented by Mega 6 software.35 Maximum likelihood (ML) phylogenetic analysis was conducted using Mega 6. 1000 bootstrap pseudo-replicates were employed for estimating node support. The general time reversible model with gamma distribution of rates across sites (GTR+G) model of evolution, estimated using JModeltest,<sup>36</sup> was used in the ML analysis. Isaria farinosa (ARSEF 4029) was used as the outgroup.

# Culture of *B. bassiana*, extraction and chromatographic fractionation

The *B. bassiana* CG71 and UNI 40 strains were cultivated on potato dextrose agar (PDA) at 26 °C for 7 days, and pre-inoculated in 60 mL of supplemented liquid medium (ML) divided into 4 Erlenmeyer flasks (50 mL) containing 30.0 g glucose; 20.0 g malt extract; 2.0 g bactopeptone; 1.0 g yeast extract; 0.5 g KCl;

0.5 g MgSO<sub>4</sub>.7H<sub>2</sub>O; 0.5 g KH<sub>2</sub>PO<sub>4</sub>; 1 L water, as reported by Bunyapaiboonsri *et al.*<sup>37</sup> The flasks were incubated on a rotary shaker (200 rpm) for 8 days. These seed cultures (240 mL of each strain) were used to inoculate 11 liters ML (55 Erlenmeyer flasks of 1 L with 200 mL of medium for each strain). After incubation at room temperature for 20 days in static mode, the mycelium was separated by reduced pressure filtration and the liquid phase was submitted to liquid-liquid fractionation with ethyl acetate (EtOAc). After lyophilization, the mycelial masses were combined and extracted totally with methanol (MeOH). The organic solvents were removed by vacuum distillation at 55 and 60 °C using a rotary evaporator.

The ethyl acetate extracts CG71a (4.7600 g) and UNI40a (4.9623 g), and the methanolic extracts CG71m (18.8054 g) and UNI40m (14.6802 g) were fractionated through Sephadex LH-20 (1.75 mm  $\times$  0.3 Ø) using MeOH as eluent. In both extracts of CG71 and UNI40 and some fractions were detected beauvericin, beauvericin A or F, beauvericin E and bassianolide.

#### Cyclodepsipeptides identification

The UHPLC-HRMS/MS analyzes were performed on an Accela High Speed LC coupled to a LTQ Orbitrap Velos FT-MS (ThermoFisher Scientific, San Jose, CA, USA), equipped with an electrospray source (HESI-II) and operated at a resolution of 60,000 FWHM, with positive ionization and 25 eV at HRMS/MS mode. The chromatographic system was fitted with an RP Luna C<sub>18</sub> column (150 × 4.6 mm, 5 µm, Phenomenex) applying as mobile phase H<sub>2</sub>O/CH<sub>3</sub>OH (30:70  $\rightarrow$  0:100, v/v, both buffered with 0.1% of formic acid) in 15 min at a flow rate of 0.4 mL min<sup>-1</sup>. The software Thermo Xcalibur 2.1 (ThermoFisher Scientific) was used to control the full system. The sample injection (auto injector) volume was 0.5 µL.

#### Bioassay to control Aedes aegypti larvae Rockefeller strain

The bioassay was performed at the Laboratory of Biocontrol of Arthropods (LABIART), Veterinary Institute, Department of Animal Parasitology, Universidade Federal Rural do Rio de Janeiro (UFRRJ). The extracts tested were CG71a, CG71m, UNI40a and UNI40m.

Aedes aegypti of the Rockefeller strain were maintained in colonies at  $27 \pm 3$  °C, and 80% humidity on a 12-h light/ dark cycle. Four replications of ten (10) third instar larvae (L-3) of Aedes aegypti susceptible in the insecticide were tested in five concentrations: 25, 50 100, 150, and 200 µL of each extract dissolved in dimethyl sulfoxide 10% (DMSO). Two control groups were tested, one containing the solvent used in extracts and the other containing only pure water. The analyzes of the results of larval mortality were assessed at 15 minutes and at successive counts of 15, 30 minutes, 1, 2, 3 and 24 and 48 hours. The lethal concentrations  $LC_{50}$  and their respective confidence intervals were calculated through Probit analyses, using the R software (R Core Team).

## **Results and Discussion**

The ethyl acetate and methanol extracts of entomopathogenic fungi *Beauveria bassiana* UNI40 and CG71 were analyzed by HRMS data analysis, obtained in an Orbitrap mass analyzer. The metabolites beauvericin (1), beauvericin A (2) or F (2a), beauvericin E (3) and bassianolide (4) were identified (Figure 1).

#### Cyclodepsipeptides fragmentation

Mass spectrometry has played essential role in the structure determination of cyclodepsipeptides originated from diverse fungal species, grains and foods. The fragmentation is a main step in sequencing and primary stage usually is bond rupture in the ester or amide group.<sup>38</sup> Interpretation and elucidation of the amino acid sequence using collision-induced dissociation (CID) depends essentially on the sequence of specific ions present ( $a_n$ ,  $b_n$ ,  $c_n$  and  $x_n$ ,  $y_n$ ,  $z_n$ ). The fragment ions produced in this process can be divided in two series.<sup>39</sup>

Beauvericin is a cyclohexadepsipeptide isolated in 1969 from *B. bassiana*, derived from two dipeptidol monomers alternately, D-hydroxyisovalerate (D-HiV) and *N*-methyl-L-phenylalanine (N-Me-Phe), while bassianolide isolated in 1977 from *B. bassiana* and *Verticillium lecanii* is a cyclooctadepsipeptide also formed by two dipeptidol monomers units of D-hydroxyisovalerate (D-HiV) and *N*-methyl-L-leucine (N-Me-Leu).<sup>18</sup>

The cyclodepsipeptides beauvericin, beauvericin A or F, beauvericin E and bassianolide were detected in two strains of *B. bassiana* (CG71 and UNI 40) extracts and fractions semi-purified according to the observed accurate masses for protonated molecule and the corresponding cationized molecules, displayed at their full scan spectra (Figure 2) and selected ion chromatogram (Figure 3). Moreover, the product ion spectra allowed the comparison of the compounds fragmentation profile to the data found in the literature (Table 1).<sup>24,25,40-44</sup>

The electrospray ionization tandem mass spectra (ESI-MS/MS) and fragmentation mechanism proposal for beauvericin A or F are depicted in Figures 4 and 5,

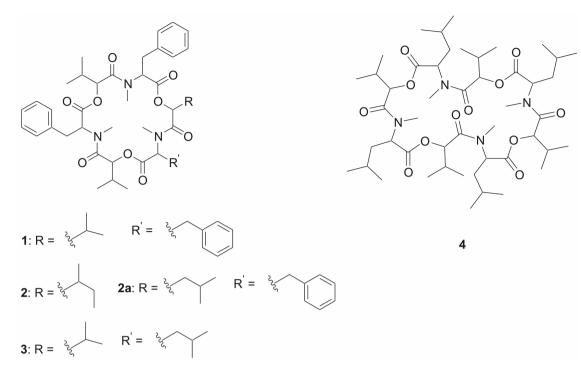


Figure 1. Structures of compounds 1-4 identified from the extracts of B. bassiana.

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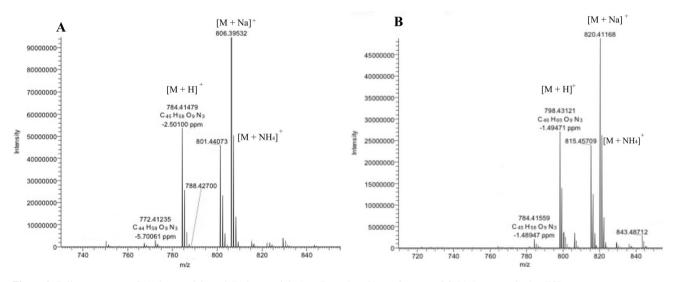


Figure 2. Full scan spectra of (A) beauvericin and (B) beauvericin A or F, produced by B. bassiana CG 71. Data acquired at ESI.

respectively. The *N*-methyl-L-phenylalanine (-161 Da) loss gave rise to the fragment ion m/z 637.3486. While the peaks m/z 555.3066 and 294.1701 are detected due to the isopropylideneketene losses (82 Da), the ion m/z 376.2121 occurs through the rupture of the amide bond. It is worth to highlight that this new fragmentation proposal is fundamental for the confirmation of the beauvericin A or F presence in the organics extracts of entomopathogenic fungi.

The inspection of the MS/MS data showed that the peaks at m/z 523.2809 in beauvericin and m/z 682.4630 in bassianolide spectra represent the loss of an 2-hydroxy-

3-methyl-1-buten-1-one molecule (100 Da), the identical D-HiV monomer (Figures 6 and 7). Additionally, the peaks at m/z 623.3333 (Figure 6) and 782.5159 (Figure 7) show the loss of *N*-methyl-L-phenylalanine (-161 Da) and *N*-methyl-L-leucine (-127 Da), respectively. Similar fragments containing phenylalanine residues (m/z 262) are described for beauvericin and E, which then lost H<sub>2</sub>O to give m/z 244 (Table 1).

Figure 8 shows fragmentation mechanism proposal for beauvericin E (precursor ion m/z 736.4722). Differently from the mechanism of the other cyclodepsipeptides produced by the fungus, the MS/MS data from beauvericin E showed

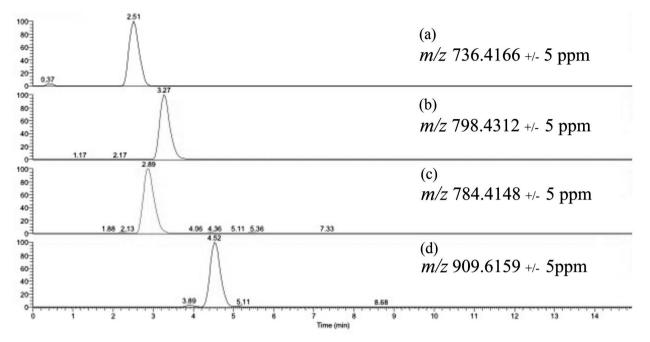


Figure 3. Selected ion chromatogram of (a) beauvericin E, (b) beauvericin A or F, (c) beauvericin and (d) bassianolide.

Table 1. Characterization	(mass accuracy) of named ion	ns cyclodepsipeptides and MS/MS	S parameters of <i>Beauveria bassiana</i> extracts

	Beauvericin	Error / ppm	Beauvericin A or F	Error / ppm	Beauvericin E	Error / ppm	Bassianolide	Error / ppm
Molecular formula	C45H57N3O9		$C_{46}H_{59}N_3O_9$		$C_{41}H_{57}N_3O_9$		$C_{48}H_{84}N_4O_{12}$	
Calcd. $[M + H]^+$	784.4175		798.4332		736.4175		909.6168	
Calcd. $[M + NH_4]^+$	801.4441		815.4598		753.4441		926.6433	
Calcd. [M + Na] <sup>+</sup>	806.3996		820.4153		758.3996		931.5988	
Found $[M + H]^+$	784.4148	-2.50	798.4312	-1.49	736.4166	-0.26	909.6159	-0.85
Found $[M + NH_4]^+$	801.4407		815.4171		753.4433		926.6425	
Found [M + Na]+	806.3953		820.4117		758.3986		932.6007	
Precursor ion $(m/z)$	784.4174	0.84	798.4323	-0.12	736.4722		909.6159	-0.85
Product ions $(m/z)$	623.3333	1.09	637.3486	0.40	708.4216		782.5159	-1.25
	523.2809	1.30	555.3066	0.29	575.3326		682.4630	2.76
	362.1967	1.42	376.2121	0.57	475.2802		555.3640	3.37
	262.1441	1.39	294.1701	0.51	362.1963		427.3168	-5.84
	244.1336 (M-H <sub>2</sub> O)	1.45			262.1438 244.1332 (M-H <sub>2</sub> O)		328.2121	-7.41
Collision energy / eV	25		25				25	
Tube lens offset / V	70		70				70	

Calcd .: mass calculated.

CO loss, corresponding to the fragment ion m/z 708.4216 [M + H – CO]<sup>+</sup>. The other peaks m/z 575.3326, 475.2802, 362.1963 and 262.1438 occur due to the rupture of ester or amide bonds. The losses of *N*-(2-phenylethylidene)-methanamine (-133 Da), 2-hydroxy-3-methyl-1-buten-1-one (-100 Da), 2-amino-4-methyl-1-penten-1-one

(-113 Da) correspond to the peaks mentioned above. This fragmentation mechanism proposal evidences the presence of beauvericins E in the organic extracts of *B. bassiana* CG 71 and UNI 40 and may assist cyclodepsipeptide characterizations according to the additional structural information.

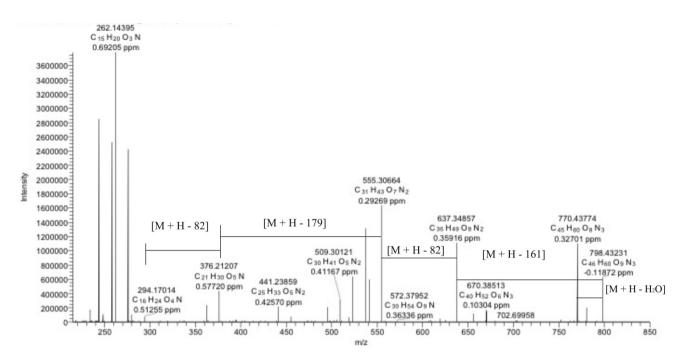


Figure 4. ESI-MS/MS spectra of beauvericin A or F, precursor ion *m/z* 798.4323.

#### Phylogenetic analyses

Strains CG 71 and UNI 40 were grouped in a wellsupported clade with reference strains *Beauveria bassiana* in the phylogenetic analyzes conducted with partial RPB2 DNA sequences (Figure 9), thus confirming their identity.

#### Larvicidal activity

Table 2 shows larvicidal activity of the *B. bassiana* CG71 and UNI40 extracts against  $3^{rd}$  instar of *Aedes aegypti*. The methanolic extract of strain CG71 revealed the highest activity with LC<sub>50</sub> values of 0.9887 and 0.4653 ppm after 24 and 48 hours, respectively. The ethyl acetate extract of CG71 displayed LC<sub>50</sub> of 29.1777 and 1.2309 ppm after 24 and 48 hours, respectively.

The UNI 40 strain showed LC<sub>50</sub> values of 0.7834 and 1.8149 ppm for the ethyl acetate and methanolic extracts after 48 hours of incubation, respectively. These extracts showed no larvicidal activity in the first 24 hours of incubation. Besides the cyclodepsipeptides detected in both CG 71 and UNI 40 extracts, other peptides with peaks at m/z above 1000 were detected in the ethyl acetate and methanolic extracts of UNI 40.

New studies using fungal organic extracts aim to identify new bioactive secondary metabolites, such as *Fusarium* sp. extract with trypanocidal activity, which revealed beauvericin as the responsible for the toxicity of *Fusarium* sp. to *Trypanosoma cruzi*.<sup>45</sup>

Beauvericin has been reported to be toxic on *Aedes aegypti* larvae with  $LC_{50}$  26 ppm,<sup>24</sup> and 10 and 20 µg mL<sup>-1</sup> showed 39 and 86% of mortality in 48 hours, respectively.<sup>46</sup> The presence of beauvericin, beauvericin A or F, beauvericin E and bassianolide in the extracts of two *B. bassiana* strains may explain the excellent larvicidal activity observed.

The cyclodepsipeptides have shown insecticidal activity against several species, such as fifth instar larvae of the silkworm *Bombyx mori*,<sup>9</sup> *Calliphora erythrocephala*, *Aedes aegypti*, *Lygus* spp., *Spodoptera frugiperda*, *Schizaphis graminum*<sup>47</sup> and *Sitophilus* spp.<sup>48</sup> However, the mycelium and conidia of a *B. bassiana* strain, pathogenic to *Tribolium castaneum* and *Sitophilus oryzae*, showed no toxicity against rats and mice.<sup>49</sup>

Bassianolide was described as a virulence factor of *B. bassiana* against *Galleria mellonella*, *Spodoptera exigua*, *Helicoverpa zea*<sup>50</sup> and *Atta sexdens sexdens*.<sup>51</sup> Beauvericin was cytotoxic (IC<sub>50</sub> 0.5  $\mu$ M) on a lepidopteran *Spodoptera frugiperda* (SF-9) cell line<sup>52</sup> and on the Colorado potato beetle (LC<sub>50</sub> 633 ppm).<sup>15</sup>

# Conclusions

The cyclodepsipeptides beauvericin, beauvericin A or F, beauvericin E and bassianolide were identified in the *B*. *bassiana* CG71 and UNI 40 extracts. The methanolic extract of CG71, ethyl acetate and methanolic extracts of UNI40 showed larvicidal activity against  $3^{rd}$  instar of *Aedes aegypti* with LC<sub>50</sub> values of 0.4653, 0.7834 and 1.8149 ppm after 48 hours of incubation, respectively. These results suggest

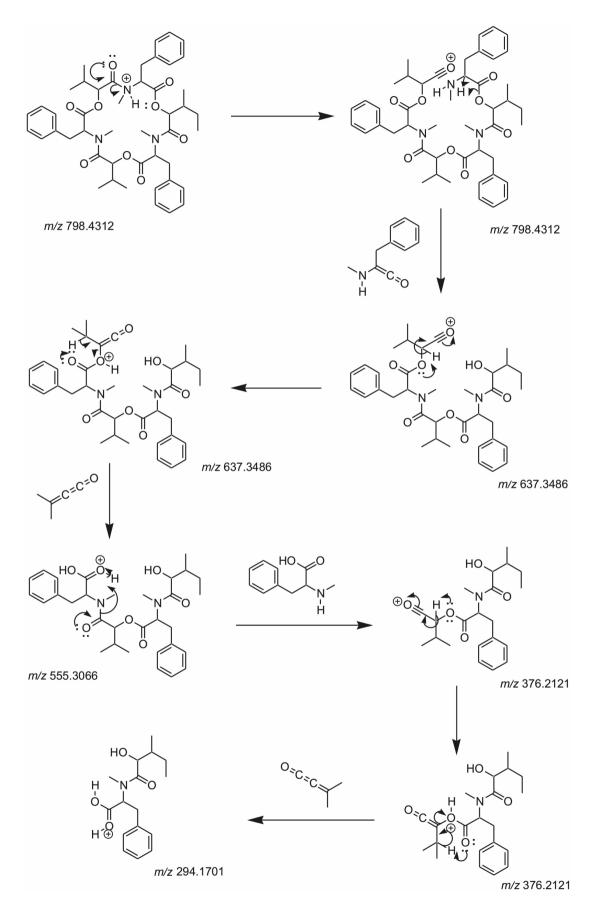


Figure 5. Plausible fragmentation mechanism for beauvericin A or F in collision-induced dissociation (CID) spectra.

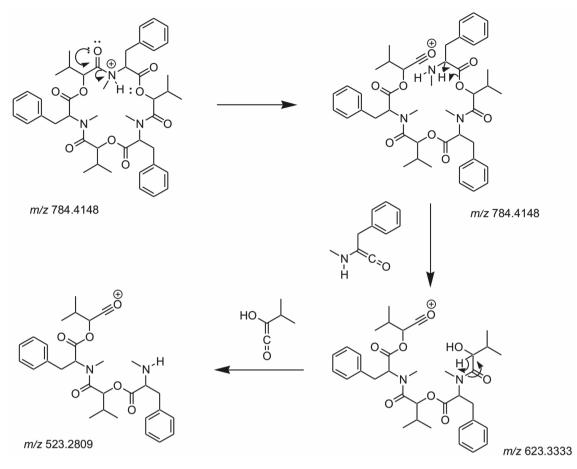


Figure 6. Plausible fragmentation mechanism for beauvericin in collision-induced dissociation (CID) spectra.

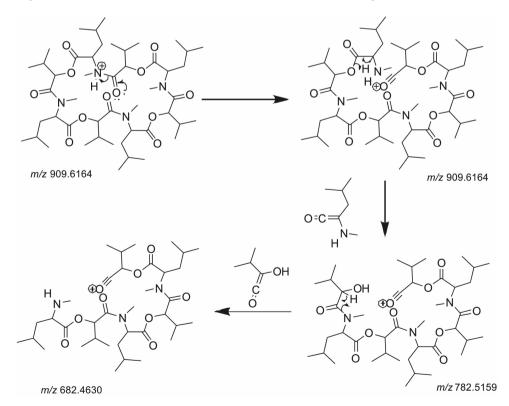
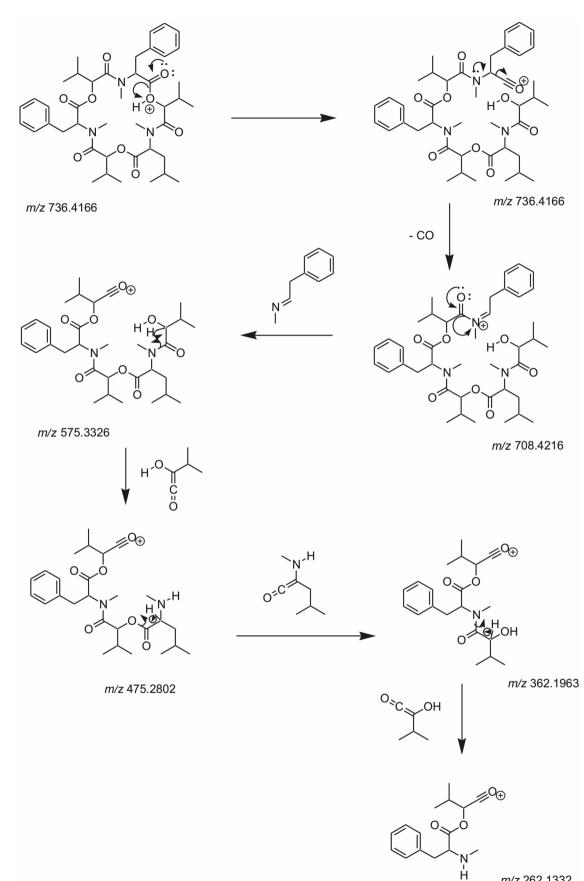


Figure 7. Plausible fragmentation mechanism for bassianolide in collision-induced dissociation (CID) spectra.

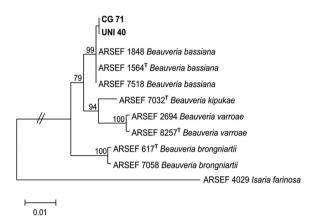


*m/z* 262.1332



Extract	LC <sub>50</sub> (24 hours) / ppm	95% CI	LC <sub>50</sub> (48 hours) / ppm	95% CI
CG71a	29.1777	24.2879-35.5307	1.2309	1.1736-1.2950
CG71m	0.9887	0.9710-1.0076	0.4653	0.4650-0.4656
UNI401a	no activity		0.7834	0.7460-0.8251
UNI401m	no activity		1.8149	1.7527-1.8906

Table 2. Larvicidal activity Beauveria bassiana UNI40 and CG71 extracts against 3rd instar of Aedes aegypti



**Figure 9.** Maximum likelihood phylogenetic tree of partial RPB2 sequences of CG 71 and UNI 40, and reference strains of *Beauveria bassiana* and related species. Bootstrap values equal or higher than 70% are showed by the nodes. *Isaria farinosa* (ARSEF 4029) was used as the outgroup. T identifies ex-type strains.

that cyclodepsipeptides are probably the active principles responsible for the larvicidal action. Therefore, further studies must be undertaken in order to isolate and confirm the compounds responsible for this activity. Therefore, they may be considered as potential insecticidal components in the formulations for the Dengue and Zika vector. The liquid chromatography coupled with electrospray ionization tandem mass spectrometry (LC-ESI-MS/MS) methodology used proved to be an excellent tool for the identification of cyclodepsipeptides in fungal extracts, and contributed with new fragmentation studies for beauvericin A or F and E.

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