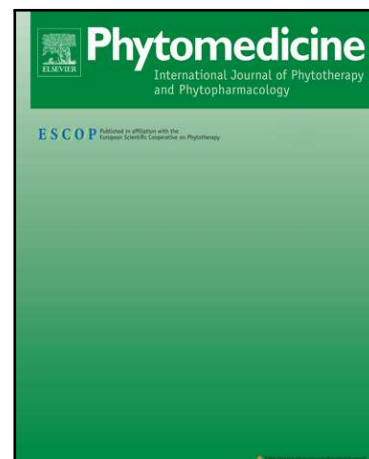


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Synergistic mutual potentiation of antifungal activity of *Zuccagnia punctata* Cav. and *Larrea nitida* Cav. extracts in clinical isolates of *Candida albicans* and *Candida glabrata*

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1 **Synergistic mutual potentiation of antifungal activity of *Zuccagnia punctata* Cav.**
 2 **and *Larrea nitida* Cav. extracts in clinical isolates of *Candida albicans* and**
 3 ***Candida glabrata***

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12
 13 **Keywords:**

14 *Zuccagnia punctata*
 15 *Larrea nitida*
 16 Bi-herbal combinations
 17 Synergism
 18 Antifungal
 19 MixLow method
 20

21 **Abbreviations**

ZpE *Zuccagnia punctata* dichloromethane extract
LnE *Larrea nitida* dichloromethane extract
 L_{ϕ} Loewe Index
 CI Combination Index
 DRI Dose Reduction Index
 IC_x, inhibitory concentration to achieve X% effect
Ca *Candida albicans*
Cg *Candida glabrata*
 NDGA Nordihydroguaiaretic acid
 DNDGA 3''- deoxy NDGA
 MNDGA 3'- O-methyl NDGA
 MixLow Mixed-effects Loewe Method
 LOD limit of detection
 LOQ limit of quantification
 S synergism
 An antagonism

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 25

26 **Abstract**

27

28 *Background:* *Zuccagnia punctata* Cav. (Fabaceae) and *Larrea nitida* Cav. (Zygophyllaceae) are
29 indistinctly or jointly used in traditional medicine for the treatment of fungal-related infections.
30 Although their dichloromethane (DCM) extract have demonstrated moderate antifungal activities
31 when tested on their own, antifungal properties of combinations of both plants have not been
32 assessed previously.

33 *Purpose:* The aim of this study was to establish with statistical rigor whether *Z. punctata* (*ZpE*) and
34 *L. nitida* DCM extract (*LnE*) interact synergistically against the clinically important fungi *Candida*
35 *albicans* and *Candida glabrata* and to characterize the most synergistic combinations.

36 *Study design:* For synergism assessment, the statistical-based Boik's design was applied. Eight
37 *ZpE-LnE* fixed-ratio mixtures were prepared from four different months of one year and tested
38 against *Candida* strains. L_ϕ (Loewe Index) of each mixture at different fractions affected (ϕ)
39 allowed for the finding of the most synergistic combinations, which were characterized by HPLC
40 fingerprint and by the quantitation of the selected marker compounds.

41 *Methods:* L_ϕ and confidence intervals were determined *in vitro* with the MixLow method, once the
42 estimated parameters from the dose-response curves of independent extracts and mixtures, were
43 obtained. Markers (four flavonoids for *ZpE* and three lignans for *LnE*) were quantified in each
44 extract and their combinations, with a valid HPLC-UV method. The 3D-HPLC profiles of the most
45 synergistic mixtures were obtained by HPLC-DAD.

46 *Results:* Three over four IC_{50ZpE}/IC_{50LnE} fixed-ratio mixtures displayed synergistic interactions at
47 effect levels $\phi > 0.5$ against *C. albicans*. The dosis of the most synergistic ($L_\phi = 0.62$) mixture was
48 $65.96 \mu\text{g/ml}$ (*ZpE* = 28%; *LnE* = 72%) containing 8 and 36% of flavonoids and lignans
49 respectively. On the other hand, one over four IC_{50ZpE}/IC_{50LnE} mixtures displays synergistic
50 interactions at $\phi > 0.5$ against *C. glabrata*. The dosis of the most synergistic ($L_\phi = 0.67$) mixture
51 was $168.23 \mu\text{g/ml}$ (*ZpE* = 27%; *LnE* = 73%) with 9.7 and 31.6% of flavonoids and lignans
52 respectively.

53 *Conclusions:* Studies with the statistical-based MixLow method, allowed for the finding of the most
54 *ZpE-LnE* synergistic mixtures, giving support to a proper joint use of both antifungal herbs in
55 traditional medicine.

56

57

58 **Introduction**

59 In 2012, the Ministry of Science, Technology and Productive Innovation of Argentina has
60 launched a National Strategic Plan (2012-2015) for developing phytomedicines containing native or
61 endemic plants based on the previous investigations conducted by research groups of this country.

62 Our group has already carried out a project devoted to the search for antifungal Latin American
63 plants (Svetaz et al. 2010) and also to the isolation of their main antifungal components (Escalante
64 et al. 2008; Pacciaroni et al. 2008; Derita et al. 2009; Vila et al. 2010; López et al. 2011; Fernández
65 et al. 2014; and others). Among the tested antifungal plants, two species, *Zuccagnia punctata* Cav.
66 (Fabaceae) (Ulibarri 1999, 2005) and *Larrea nitida* Cav. (Zygophyllaceae) showed moderate
67 antifungal properties against clinical important fungi (Svetaz et al. 2007; Agüero et al. 2010, 2011;
68 Alvarez et al. 2012; Nuño et al. 2014).

69 *Z. punctata* (common names “*jarilla macho*”, “*jarilla de la punta*”, “*laca*” or “*pus-pus*”) is a
70 monotypic species (Ulibarri 2005) currently used in traditional medicine for bacterial and fungal
71 infections. Moreover, this species can also be used for other pathologies like asthma, arthritis,
72 rheumatism and tumors (Ratera and Ratera 1980). During many years, *Z. punctata* has been
73 described as growing in Chile and Argentina (Ulibarri 1999). However, a thorough study based on
74 its botanical and bibliographic collections, allowed to demonstrate that the shared distribution with
75 Chile was erroneous and thus it was firmly established that *Z. punctata* is endemic of Argentina
76 (Ulibarri 2005).

77 Among the different extracts previously tested for antifungal properties, the dichloromethane
78 (DCM) extract (*ZpE*) was the most active one (Svetaz et al. 2007; Agüero et al. 2010). This extract
79 led to the isolation of several compounds of which 5,7-dihydroxy-3-flavonol (galangin, **1**), 5,7-
80 dihydroxiflavanone (pinocembrin, **2**) and 2',4'-dihydroxychalcone (**3**) showed antifungal properties
81 while 2',4'-dihydroxy-3'-methoxychalcone (**4**) (Fig. 1) was the most abundant (though inactive)
82 compound. The other compounds did not show significant activity in concentrations up to 250
83 µg/ml.

84 In turn, the native plant *L. nitida* (common names “*jarilla de la sierra*” and “*jarilla fina*”) (Del
85 Vitto et al. 1997) is one of the four South American species of *Larrea* genus (Timmermann et al.
86 1979) that grows in Argentina and Chile (Hunziker 2005). Antioxidant and antifungal activities
87 were previously reported for this species (Torres et al. 2003; Agüero et al. 2011), being the DCM
88 extract (*LnE*) the most active one. Its bioassay-guided fractionation led to the isolation of five
89 lignans of which only three, nordihydroguaiaretic acid (NDGA, **5**), 4-[4-(4-hydroxy-phenyl)-2,3-
90 dimethyl-butyl]-benzene-1,2-diol (DNDGA, **6**) and 3'-O-methyl-nordihydroguaiaretic acid
91 (MNDGA, **7**) (Fig. 1) showed moderate antifungal properties (Agüero et al. 2011).

92 INSERT FIG. 1

93 Although *L. nitida* is less abundant and grows in more restricted regions, it is indistinctly or
94 conjunctly used with *Z. punctata* due to their similar morphological characteristics and also to their
95 common name (*jarilla*) (Del Vitto et al. 1997). However, until now, their combinations were used
96 in an empirical basis, as no mixtures have been scientifically assessed by their type of interaction.

97 The use of bi-herbal mixtures for the treatment of a disease is a common practice in traditional
98 medicine (Wagner and Ulrich-Merzenich 2009) in the belief that they may achieve a better
99 therapeutic effect (synergism) than when used independently (Sibandze et al. 2010). However,
100 antagonistic or additive effects can be also found between the components of a plant combination
101 (Williamson 2001; Odds 2003).

102 The study of antifungal interactions between two extracts is a complex task. The fungi target, the
103 ratio between extracts in the mixture and the methodology for quantifying synergism must be
104 chosen carefully in order to achieve trustworthy conclusions. In addition, since extracts are complex
105 mixtures that usually present seasonal variations, several characterized batches must be combined in
106 order to get quantified mixtures that display the highest synergism.

107 In this paper we report the study of the antifungal behavior of four bi-herbal *ZpE-LnE*
108 combinations, each of them prepared by mixing one sample of *ZpE* with one sample of *LnE*
109 collected in a same period of a year. This was made for the four periods in which the plants were
110 collected.

111 *Candida albicans* and *Candida glabrata* were used as targets for the antifungal evaluation. The
112 choice of *C. albicans* was due to this yeast is the most common cause of opportunistic fungal
113 infections in immune compromised hosts (Pfaller and Diekema 2007). In turn, the selection of *C.*
114 *glabrata* was made for it has been identified as the second leading cause of adult candidemia
115 particularly in patients with hematologic malignancies (Malani et al. 2005; Pfaller and Diekema
116 2007).

117 Among the several mathematical methods to quantify synergism that have been proposed over
118 the last few decades (Berembaum 1989; Merlin 1994; Greco et al. 1995; Tallarida 2001; Chou
119 2006; Boik et al. 2008), here we chose the Mixed-effects Loewe (MixLow) method (Boik et al.
120 2008) to determine the Loewe Index (L_{ϕ}) previously defined as Combination Index (CI) within the
121 Median-effect method (Chou 2006). Both the MixLow and the Median-effect methods share the
122 following characteristics: (i) assess the data from single ray fixed-ratio experiments; (ii) allow the
123 identification of the optimal concentration (within the fixed-ratio) that will give the maximum
124 synergy; (iii) present the results in a graphical form. However, MixLow method has the advantage
125 over the Median-effect that it allows for the statistically comparison of the combinations' effects by

126 providing accurate dose-response curves' parameters and confidence intervals for L_{ϕ} that are vital
127 to fully assess whether drugs in a mixture interact synergistically, antagonistically or additively.

128 As a result of this study, we could determine the L_{ϕ} of each mixture at different effects (ϕ), and
129 the composition of the extracts in the two most synergistic *ZpE-LnE* combinations against each
130 fungus. The content of the selected markers in both combinations was determined by HPLC-UV
131 following the International Conference on Harmonization guidelines (ICH 1996). The 3D-HPLC
132 profiles of both synergistic combinations were provided too.

133 **Materials and methods**

134 *Plant material*

135 Aerial parts (AP) of *Z. punctata* and *L. nitida* (www.theplantlist.org) were collected over four
136 periods of 2013 (February, May, September and November) at Las Flores and Bauchaceta towns,
137 Agua Negra pathway, Iglesia district, San Juan province (Argentina), respectively. The plants were
138 identified by Dr. Martin Hadad, Instituto de Biotecnología (IBT), Universidad Nacional de San Juan
139 (UNSJ) and each batch was deposited at the herbarium of the IBT-UNSJ and identified as *ZpLF*
140 AT-IBT 02 (abbreviated as *ZpE* Feb), *ZpLF* AT-IBT 05 (*ZpE* May), *ZpLF* AT-IBT 09 (*ZpE* Sept),
141 and *ZpLF* AT-IBT 11 (*ZpE* Nov), *LnBau* AT-IBT 02 (*LnE* Feb), *LnBau* AT-IBT 05 (*LnE* May),
142 *LnBau* AT-IBT 09 (*LnE* Sept) and *LnBau* AT-IBT 11 (*LnE* Nov).

143 *Preparation of extracts*

144 *ZpEs* and *LnEs* were obtained by dipping fresh AP (500g) in cold DCM (3 x 1l) at room
145 temperature for 40 s. The solutions were evaporated under vacuum to give semisolid yellow
146 residues (*ZpE* Feb, 65 g, 13% w/w yield; *ZpE* May, 55 g, 11%; *ZpE* Sept, 60 g, 12%; *ZpE* Nov, 50
147 g, 10%. *LnE* Feb, 50 g, 10%; *LnE* May, 42.5 g, 8.5%; *LnE* Sept, 45 g, 9% and *LnE* Nov, 49 g,
148 9.8%).

149 *Source of markers*

150 *ZpE*'s markers galangin (**1**) and pinocembrin (**2**) and *LnE*'s marker NDGA (**5**) were purchased
151 from Sigma-Aldrich (St. Louis, MO, USA). Instead, 2',4'-dihydroxychalcone (**3**), 2',4'-dihydroxy-
152 3'-methoxychalcone (**4**) and MNDGA (**7**) were obtained in our laboratory from *ZpE* or *LnE* as
153 previously described (Svetaz et al. 2007; Agüero et al. 2010, 2011). The purities of reference
154 compounds were $\geq 95\%$ as determined by HPLC-DAD.

155 *Preparation of extracts' and markers' solutions*

156 Acetonitrile (ACN) solutions of *ZpE* and *LnE* from Feb, May, Sept and Nov and of **1-7** were
157 prepared at an appropriate concentration, filtered through a 30 mm, 0.45 μm Target Nylon
158 Membrane Syringe Filter (Scientific Instrument Services, Ringoes, NJ) prior to injection into the
159 HPLC system.

160 *HPLC-ESI-MS/MS analyses*

161 HPLC-ESI-MS/MS analyses of all batches of *ZpE* and *LnE* were carried out by using a
162 MicroTOFQ II apparatus (Bruker Daltonics, MA, USA), equipped with an ESI ion source with
163 nitrogen as nebulizing gas (4 psi) and drying gas (8 l/min, 200 °C); capillary 4500 V and end plate
164 offset at 500 V. Mass accuracy was verified by infusing a 10 mM solution of Na-formiate (Sigma-
165 Aldrich) dissolved in MeOH:H₂O (50:50) and a C18-RP column Thermo Scientific (USA) Hypersil
166 Gold (50 x 2.1 mm, 3 μm). Firstly, pure compounds were characterized by direct infusion to ESI
167 using a syringe pump (Harvard Apparatus 11 Plus) recording both MS and MS/MS spectra.
168 Because of the improved performance reached using direct infusion to ESI, we report only MS and
169 MS/MS data obtained in the negative mode. Reference compounds and sample solutions were
170 introduced in the HPLC (5 μl) using an autosampler (Agilent HiP-ALSSL+) at 25 °C. Flow rate
171 was set to 0.20 ml/min and 0.25 ml/min for *ZpE* and *LnE* samples and their own reference
172 compounds, propelled by an Agilent 1200 series G1312B SL binary pump, using ultra-pure water
173 or HPLC-grade ACN with 0.1% formic acid (solutions A and B respectively) following different
174 methods: for *ZpE* extracts and reference compounds, the gradient started with 20% B, changing to
175 100% B within 10 min, then the composition was held 2 min, returning to 20% B in 3 min and
176 keeping this condition for additional 5 min to achieve the column stabilization before the next run
177 (total run time was 20 min). Eluted compounds were monitored at 254 nm. For *LnE* extracts and
178 reference compounds, the program started with 40% B, changing to 45% B within 10 min, and to
179 100% B the following 5 min, held by 4 min and returning to 40% B in 1 min (total run time 20
180 min). Eluted compounds were monitored at 280 nm. The HPLC flow was introduced into the Mass
181 spectrometer *via* an ESI source. For the analyses of chromatograms and mass spectra, the Data
182 Analysis 4.0 SP1 software (Bruker Daltonik GmbH, Germany) was used. Compounds **1-7** were
183 identified by comparing their retention times (Rt), HRMS and MS/MS with those of reference
184 compounds (Agüero et al. 2010, 2011). Markers' content was quantified in the extracts of the
185 different periods using the HPLC-UV chromatograms. Five calibration curves of pure compounds
186 **1-5** were prepared with five appropriate dilutions of stock ACN solutions by triplicate. To monitor
187 the samples, the wavelengths were chosen according to absorption maxima of markers: 267 nm (**1**),
188 289 nm (**2**), 341 nm (**3** and **4**) and 280 nm (**5**).

189 *Method validation*

190 Analytical method's linearity, limit of detection and quantification (LOD and LOQ) and inter-
 191 day / intra-day precision were validated following the ICH guidelines (ICH 1996). Recovery was
 192 used to evaluate the accuracy of the method.

193 *Strains used for the antifungal assessment of extracts and mixtures*

194 For the antifungal evaluation of independent extracts and their combinations, clinical isolates of
 195 *C. albicans* (CCC 125) and *C. glabrata* (CCC 115) provided by Centro de Referencia en Micología
 196 (CEREMIC, CCC, Rosario, Argentina), were used. Strains were grown on Sabouraud-
 197 chloramphenicol agar slants for 48 h at 30 °C, maintained on Sabouraud-Dextrose Agar (SDA,
 198 Oxoid) and sub-cultured every 15 days to prevent pleomorphic transformations. Inocula were
 199 adjusted to $1-5 \times 10^3$ cells with colony forming units (CFU)/ml according to the Clinical and
 200 Laboratory Standards Institute (CLSI 2008).

201 *Synergism studies design*

202 The nature of the interaction between *ZpE* and *LnE* in the presence of either *C. albicans* or *C.*
 203 *glabrata* inoculum, was analyzed by the L_ϕ (CI) defined by Eq. (1).

$$L_\phi = \frac{ICx \text{ ZpE in mixture}}{ICx \text{ ZpE alone}} + \frac{ICx \text{ LnE in mixture}}{ICx \text{ LnE alone}} \quad (1)$$

204 Where ICx is the inhibitory concentration used to achieve X% effect (measured as diminution of
 205 fungal growth) of each extract alone and in the mixture. The estimation of L_ϕ (Boik et al. 2008) was
 206 a three-step (a-c) process (Scheme 1): (a) preparation of dose-response curves for each extract alone
 207 and for their combinations at a wide range of concentrations. Estimation of parameters that define
 208 the shape of each dose-response curve by using a non-linear regression model (outlined in Scheme
 209 1, a_1 , a_2); (b) use of the estimated parameters for calculating L_ϕ ; (c) generation of confidence
 210 intervals for L_ϕ .

211 INSERT SCHEME 1

213 *Dose-response curves of the independent extracts against Candida strains*

214 Broth microdilution techniques were performed in 96-well microplates according to the
 215 M27-A3 document of CLSI for yeasts (CLSI 2008). Each plate was designed as follows (Boik
 216 et al. 2008): 10 different concentrations of each extract (wells 3-12, Fig. 2), by sextuplicate
 217 (wells A-F, Fig. 2) were prepared [T wells (= 60), Fig. 2] from DMSO (maximum
 218 concentration $\leq 1\%$) 0stock solutions of each extract, diluted with RPMI-1640. In addition, 36
 219 control wells [12 treatment control growth wells (TC) containing culture medium plus
 220 inoculum; 4 medium control wells (MC) (culture medium plus water) and 20 blanks-by-tray

221 control wells (BT) (extracts plus water)] were included in the plate (Fig. 2). Each assay was
 222 repeated thrice. So, 18 replicates of each concentration (i.e. (3A-3F) x3; (4A-4F) x3, and so
 223 on) were prepared.

224 INSERT FIG. 2

225 In each plate, *ZpE* and *LnE* concentrations ranges were selected in order to cover growth
 226 percentages within 95-5% range. To better distribute the points in *x*-axis, and therefore obtain
 227 a better adjustment of the sigmoidal curve, equi-spaced concentrations in the logarithmic
 228 scale (ln) were prepared (Table S1 of Supplementary Material). An inoculum suspension (100
 229 μ l) was added to each well (final well volume = 200 μ l) and plates were incubated 24 h at 28-
 230 30 °C in a moist dark chamber. After incubation, the wells' absorbance values were recorded
 231 at 405 nm with a VERSA Max microplate reader (Molecular Devices, Sunnyvale, CA, USA)
 232 and introduced in the MixLow package of R software (Boik and Narasimhan 2010). Dose-
 233 response curves modeled as a sigmoidal function of responses $Y_{d,t,w}$ (*y*-axis) and extracts'
 234 concentrations (*c*) (*x*-axis) (Eq. 2 and 3) were generated.

$$Y_{d,t,w} = \exp(\mu + b_t) (1 - \phi_{d,w,t}) + \epsilon_{d,t,w} \quad (2)$$

235 where

$$\phi_{d,t,w} = 1 - \frac{1}{1 + \left(\frac{\exp(c_{d,t,w})}{\exp(\psi_{d,0.5})}\right)^{\gamma_d}} \quad (3)$$

236 $\phi = E[Fa]$, where *Fa* (fraction affected) represents the fraction of fungi affected by an extract at a
 237 concentration *c*, and $E[\bullet]$ is the expected value; $c_{d,t,w}$ refers to the log of the extract's concentration
 238 for the *d* (*d*th extract), *t* (*t*th tray), *w* (*w*th well) that is a known value. Each sigmoidal curve is
 239 parameterized by three constants: $\psi_{d,0.5}$, γ_d and μ (Scheme 1, *a*₁). The parameter $\psi_{d,0.5}$ is the log
 240 concentration of each extract that produce a $\phi = 0.50$ (50% inhibition relative to controls). By
 241 convention, $\exp(\psi_{d,0.5})$ is called the IC_{50} ; γ_d is a shape parameter and μ is the mean of control wells
 242 from all trays (Boik et al. 2008).

243 *Dose-response curves of mixtures against Candida strains*

244 Mixtures of *ZpE* and *LnE* were prepared by mixing the mass that produces an equipotent effect
 245 of each partner, in the fixed ratio IC_{50ZpE}/IC_{50LnE} and submitted to antifungal evaluation in 96-
 246 wells microplates with the same experimental design of independent extracts. In brief, within each
 247 plate, 10 different concentrations of each fixed-ratio were prepared (Table S2 of Supplementary
 248 material). Each sigmoidal curve is parameterized by three constants: $\psi_{m,0.5}$, γ_m and μ (Scheme 1,
 249 *a*₂).

250 *Quantification of synergism*

251 Once obtained the estimated parameters from the dose-response curves of the independent
 252 extracts ($\psi_{d,0.5}$ and γ_d) and mixtures ($\psi_{m,0.5}$ and γ_m), the MixLow package can calculate the L_ϕ at
 253 different ϕ effects using Eq. (4).

$$254 \quad L_\phi = \sum_{d=1}^n \tau_d \exp \left(\log \left(\left(\frac{\phi}{1-\phi} \right)^{1/\gamma_m} \right) + \psi_{m,0.5} - \log \left(\left(\frac{\phi}{1-\phi} \right)^{1/\gamma_d} \right) - \psi_{d,0.5} \right) \quad (4)$$

255 where τ_d is the fraction of the mixture that is composed of extract d .

256 The confidence intervals for the L_ϕ at different ϕ are calculated with Eq. (5).

$$257 \quad \exp(\log(L_\phi) \pm t_{df,1-\frac{\alpha}{2}} \text{SE}(\log L_\phi)) \quad (5)$$

258 *Dose Reduction Index (DRI)*

259 The DRI is a measure of how many times the dose of each drug in a synergistic combination is
 260 reduced at a given effect level compared with the doses of each independent drug. The DRI value
 261 for each drug is calculated using Eq. 6 (Chou 2006). A greater DRI indicates a greater dose
 262 reduction for a given effect level.

$$\text{DRI} = \frac{\text{ICx extract alone}}{\text{ICx extract in mixture}} \quad (6)$$

263 *3D-HPLC profile*

264 The most synergistic mixtures were dissolved in ACN, filtered by a membrane filter and
 265 subjected to HPLC analysis in a Hewlett Packard 1050 (Palo Alto, CA), coupled to a DAD detector
 266 (HP/Agilent series 1050 DAD), with a quaternary pump and autosampler (HP/Agilent series 1050)
 267 and Luna C18-RP column (Phenomenex) of 25 cm x 4.6 mm, 5 μm of particle size. The isocratic
 268 solvent phase was composed of ultra-pure water supplemented with formic acid 0.1% (40%) and
 269 HPLC-grade ACN (60%). The flow rate was 0.5 ml/min and the injection volume, 5 μl . Peaks in
 270 the extracts, monitored at $\lambda = 254$ nm, were assigned based on the R_t of reference compounds.

271 *Statistical analysis*

272 Statistical analysis was performed with GraphPad PrismTM 4.0 (GraphPad software Inc., La
 273 Jolla, CA). The data were analyzed using both, Kruskal-Wallis test (non-parametric ANOVA) and
 274 Dunn test to verify the difference between treatments; p values < 0.05 were considered significant.

275 **Results and discussion**276 *Markers for HPLC analyses*

277 The choice of markers **1-7** (Fig. 1) was based on our previous reports. Flavonoids **1-3** and
278 lignans **5-7** were considered active markers (EMEA 2007) due to they previously showed to be the
279 main anti-*Candida* constituents of *ZpE* and *LnE* respectively (Svetaz et al. 2007; Agüero et al.
280 2010, 2011) and **4** was considered an analytical marker (EMEA 2007) since, although inactive, it
281 was present in high amounts in *ZpE* (Agüero et al. 2010).

282 *HPLC fingerprints of ZpE and LnE batches*

283 HPLC fingerprints of *ZpE* and *LnE* batches (Feb, May, Sept and Nov), are shown in Fig. 3; **1-4**
284 and **5-7** were unequivocally identified (Rt, HRMS and MS/MS data) (Table 1) in each batch of
285 *ZpE*, and *LnE* respectively (Agüero et al. 2011).

286 INSERT FIG. 3 and TABLE 1

287 *Quantitative assessment of markers in ZpE and LnE batches*

288 The quantification of **1-4** and **5-7** was made by HPLC-UV in each batch of *ZpE* or *LnE*
289 respectively. HPLC method was first validated for linearity, LOD and LOQ, precision and accuracy
290 following ICH guidelines (ICH 1996), with compounds **1-5**. Compounds **6** and **7** of *LnE* were
291 quantified by the relative response factors (Gao et al. 2009) based on the calibration curve of **5**.

292 *Linearity and calibration curves:* Linearity of pure **1-5** calibration curves was established by
293 calculating the slope, intercepts and R^2 coefficient. The regression equation and R^2 (0.99-1) showed
294 good linearity response in the ranges detailed in Table 2. LOD and LOQ were calculated as $3.3 \sigma/S$
295 and $10 \sigma/S$ respectively, being σ the response standard deviation, and S the slope of each marker.

296 *Precision:* Intra- and inter- day variability test was determined for three times within 1 day and 3
297 separated days at three different concentrations, respectively. Variations were expressed by the
298 relative standard deviations (RSD) (Table 2), confirming the precision of the proposed method.

299 *Accuracy:* Three concentrations of pre-analyzed sample solutions were spiked with known
300 quantities of the standards and injected in triplicate to perform recovery studies. The percentage
301 recovery for **1-5** were between 90.6-106.6% (RSD < 4%, n = 3), confirming the accuracy of the
302 proposed method.

303 INSERT TABLE 2

304 Results of quantitative analyses of markers in the eight extracts are shown in Table 3.

305 INSERT TABLE 3

306 The non-parametric Kruskal-Wallis test found significant differences between the medians of
307 markers content (data not shown) for all the periods ($p < 0.05$). The Dunn test showed statistically
308 significant differences ($p < 0.05$) in: (a) the content of **1**, **2** and **4** between *ZpE* Feb and *ZpE* May;

(b) the content of **3** between *ZpE* Sept and *ZpE* Nov; (c) the content of **5** and **7** between *LnE* Feb and *LnE* Nov and (d) the content of **6** between *LnE* May and *LnE* Nov. This variation in markers' composition among batches justified the preparation of mixtures with each of them, since the interactions would be different.

Statistically supported synergism assessment by using fixed-ratio mixtures

Each *ZpE* or *LnE* batch was tested against the yeasts *C. albicans* and *C. glabrata* in 96-wells microplates prepared as explained in Materials and Methods (Fig. 2).

With the absorbance values obtained from 18 replicates of each concentration, a sigmoidal dose-response curve for each *ZpE* or *LnE* batch was constructed and analyzed with the MixLow method, which gave the estimated parameters $\psi_{d,0.5}$ [and thus $\exp(\psi_{d,0.5}) = IC_{50}$], γ_d and μ , which are shown in Table S3. Table 4 (columns 3 and 4) shows the IC_{50} values of each extract against each fungus.

INSERT TABLE 4

Ten different concentrations of each $IC_{50}ZpE/IC_{50}LnE$ fixed-ratio mixture were prepared according to Materials and Methods (Table 4, column 5). Each fixed-ratio concentrations' set can be regarded as a ray, in a so-called "ray design". All sets were tested against either *C. albicans* or *C. glabrata* in 96-wells plates following the same design used for the independent extracts. After incubation, the absorbance values were introduced in the MixLow package of R software generating sigmoidal dose-response curves and the corresponding parameters $\psi_{m,0.5}$, γ_m and μ for all combinations (Table S3). Fig. 4 and 5 (A,B,C,D) show the dose-response curves obtained for *ZpE* and *LnE* on their own and in fixed-ratio combinations against *C. albicans* and *C. glabrata* respectively in the four periods of the year. In Fig. 4 and 5 (A'B'C'D') the affected fraction of yeasts' population at a giving drug concentration ϕ was plotted vs the Loewe Index L_ϕ . In these graphs, full lines indicate the L_ϕ at different ϕ for a given mixture and dotted lines indicate the ninety-five percent confidence intervals of the index. $L_\phi = 1$ indicates additivism (Ad) or no interaction; $L_\phi > 1$ indicates antagonism (An) and $L_\phi < 1$ indicates synergism (S) (Boik et al. 2008; Liu et al. 2013). Within values of $L_\phi < 1$, the lower L_ϕ , the greater synergism (Chou 2006). As previously suggested (Chou 2006), $L_\phi = 0.91-1.09$ represents near (n)Ad; $L_\phi = 0.85-0.90$, slight (sl)S; 0.70-0.84, moderate (m)S; 0.30-0.69, medium (med)S; 0.10-0.29 strong (s)S and < 0.1 , very strong (vs)S; 1.10-1.20 slight (sl)An; 1.21-1.45 moderate (m)An; 1.46-3.30 medium (med)An; 3.40-10, strong (s)An and > 10 very strong (vs)An.

INSERT FIG. 4 AND 5

In Fig. 4 (A'-D'), it is clear that, when acting against *C. albicans*, all *ZpE-LnE* mixtures showed some degree of synergism in the following ϕ ranges: $0.02 \leq \phi \leq 0.74$ (Feb); $0.02 \leq \phi \leq 0.94$ (May); $0.50 \leq \phi \leq 0.94$ (Sept.) and $0.06 \leq \phi \leq 0.94$ (Nov.). In turn, when acting against *C. glabrata* (Fig. 5, A'-D'), *ZpE-LnE* Feb and May mixtures showed synergism in the ϕ ranges: $0.40 \leq \phi \leq 0.94$ and $0.02 \leq \phi \leq 0.49$ respectively, while *ZpE-LnE* Sept and Nov mixtures showed antagonism ($L_\phi > 1$) at all ϕ . For the sake of clarity, the L_ϕ values and confidence intervals for $\phi = 0.50, 0.80, 0.90$ and 0.95 extracted from Fig. 4 and 5 (A'-D') are recorded in Table 5.

INSERT TABLE 5

Table 5 clearly shows that the *ZpE-LnE* May mixture was the most synergistic against *C. albicans*, displaying the lowest L_ϕ values = 0.62, 0.65 and 0.68 [(med)S] at $\phi = 0.80-0.95$. It also showed (m)S with $L_\phi = 0.73$ at $\phi = 0.50$ against this clinically important fungus.

Instead, *C. glabrata* was less sensitive to the tested mixtures. The only combination that showed synergism was that of Feb, which achieved a $L_\phi = 0.67$ [(med)S] when $\phi = 0.95$, and $L_\phi = 0.71$ or 0.77 [(m)S] at $\phi = 0.90$ or 0.80 . Interesting enough, all mixtures of May, Sept and Nov showed (n)Ad or (sl) or (m)An, but none of them showed S.

Composition of synergistic mixtures at different effects levels and DRI values

For the most synergistic mixtures (*ZpE-LnE* May for *Ca* and *ZpE-LnE* Feb for *Cg*), the concentrations of each extract in each mixture were calculated using Eq. (2) and (3) (see Materials and Methods) at $\phi = 0.50, 0.80, 0.90$ and 0.95 . Table 6 shows that for achieving an effect level of $\phi = 0.95$ against *C. albicans*, the sum of *ZpE* and *LnE* in the mixtures should be $65.96 \mu\text{g/ml}$ (column III), composed by $18.84 \mu\text{g/ml}$ of *ZpE* (column IV) and $47.12 \mu\text{g/ml}$ of *LnE* (column V). Instead, for achieving lower ϕ (0.9, 0.8 and 0.5), the sum of *ZpE* and *LnE* doses were $57.70 \mu\text{g/ml}$ (*ZpE*: 16.48 and *LnE*: $41.22 \mu\text{g/ml}$); 49.91 (*ZpE* 14.26 and *LnE* $35.65 \mu\text{g/ml}$) and $38.94 \mu\text{g/ml}$ (*ZpE* 11.80 and *LnE* $27.14 \mu\text{g/ml}$) respectively. Instead, for achieving an effect level of $\phi = 0.95$ against *C. glabrata*, the sum of *ZpE* and *LnE* concentrations in the mixtures should be $168.23 \mu\text{g/ml}$ (column III) composed by $45.47 \mu\text{g/ml}$ of *ZpE* (column IV) and $122.76 \mu\text{g/ml}$ of *LnE* (column V). In addition, for lower ϕ (0.90, 0.80 and 0.50), their *ZpE* and *LnE* doses are also higher ($158.56, 148.69$ and $133.22 \mu\text{g/ml}$) than those needed for inhibiting *C. albicans*.

INSERT TABLE 6

DRI values were calculated for different ϕ from Eq. 6. As shown in Table 6 (columns VII and VIII), *ZpE* in combination with *LnE* improved 2.66 and 1.29-fold (= DRI values) the capability of inhibition of 50% or 80% of *C. albicans* growth compared to when acting on their own (IC_{50} and IC_{80} decreased from 31.40 to $11.80 \mu\text{g/ml}$ and from 45.91 to $14.26 \mu\text{g/ml}$ respectively, columns I

and IV). Interestingly, IC₉₀ and IC₉₅ diminished 3.48 and 3.73-fold (IC₉₀, from 57.33 to 16.48 µg/ml and IC₉₅ from 70.36 to 18.84 µg/ml), indicating that a significant reduction of the doses of ZpE in the combination respective to the independent extract, is required for a complete inhibition of the fungus. From the point of view of LnE, IC₅₀, IC₈₀, IC₉₀ and IC₉₅ against *C. albicans* decreased 2.71, 2.76, 2.83 and 2.89-fold respectively.

Against *C. glabrata*, ZpE Feb within the combination, improved 1.85 and 1.86-fold the capability of 90 or 95% growth inhibition, compared to when acting on its own (IC₉₀ and IC₉₅ decreased from 79.43 to 42.86 µg/ml and from 84.69 to 45.47 µg/ml respectively). Additionally, LnE Feb improved 6.02 and 7.83-fold the capability of inhibition of 90 or 95% of *C. glabrata* growth in combination with ZpE Feb, compared to when acting on its own (IC₉₀ and IC₉₅ of LnE decreased from 697.03 µg/ml and 960.28 µg/ml to 115.70 and 122.76 µg/ml respectively).

Markers content in most synergistic mixtures

Table 7 shows the concentrations of each marker in May and Feb mixtures at different effect levels (ϕ) against *C. albicans* and *C. glabrata*, respectively.

INSERT TABLE 7

Against *C. albicans*, the most synergistic ZpE-LnE mixture contains 8% of flavonoids **1-4** and 36% of lignans **5-7**. Within Zp markers, a predominance of 2,4-dihydroxychalcone **3** (4.5%) followed by 2,4-dihydroxy-3-methoxy chalcone **4** (2.6%) and pinocembrin **1** + galangin **2** (0.9%) was observed. Within Ln markers, a prevalence of MNDGA **7** (18%) followed by NDGA **5** (12%) and then DNDGA **6** (6%) was observed. However, although the proportion of markers are the same for all ϕ , the doses of the whole mixtures are different for achieving the different effects $\phi = 0.50$, 0.80, 0.90 and 0.95 (38.94, 49.91, 57.70 and 65.96 µg/ml respectively) (Fig. 6). Obviously, the dosis of 65.96 µg/ml for achieving a $\phi = 0.95$ (95% inhibition of *C. albicans*) (with markers proportion described above) should be chosen for preparing a phytomedicine with activity against *C. albicans* infections.

Against *C. glabrata*, ZpE-LnE the most synergistic mixture contains 9.7% of flavonoids **1-4** and 31.6% of lignans **5-7**. Within Zp markers, a predominance of **3** (4.7%) was observed, closely followed by **4** (4.4%) and with lower concentrations of **1** + **2** (0.90%). Within Ln markers, a prevalence of **7** (17%) followed by **5** (8%) and then **6** (7%) was observed. The whole combinations' doses for achieving $\phi = 0.50$, 0.80, 0.90 and 0.95 were 133.22, 148.69, 158.56 and 168.23 µg/ml, much higher than the doses needed for inhibiting *C. albicans* (Fig. 6). The dosis of 168.23 µg/ml (with the proportion of markers described above) for $\phi = 0.95$ should be chosen when preparing a phytomedicine that is able to control *C. glabrata* infections. Fig. 6 comparatively shows the doses needed for inhibiting each fungus for all $\phi = 0.50$, 0.80, 0.90 and 0.95.

408 INSERT FIG. 6

409 It is clear that the dosis of the fixed ratio mixture of *ZpE+LnE* necessary to achieve the higher
 410 synergism at the best effect level $\phi = 0.95$ against *C. glabrata* (168.23 $\mu\text{g/ml}$) is 2.55 times higher
 411 than the dosis required to inhibit *C. albicans* (65.96 $\mu\text{g/ml}$). The ratios between the amount of
 412 flavonoids markers **1-4**, and lignans markers **5-7** will be 0.22 (5.39 $\mu\text{g/ml}/23.63$ $\mu\text{g/ml}$) against *C.*
 413 *albicans* and 0.30 (16.35 $\mu\text{g/ml}/53.24$ $\mu\text{g/ml}$) against *C. glabrata*, clearly showing that lignans are
 414 required in higher proportions than flavonoids against both fungi, although this difference is less
 415 pronounced when acting against *C. glabrata*.

416 *Declaration of herbal extracts and their most effective synergistic combination characteristics*

417 Considering that the most synergistic combinations are those that have an effect level $\phi = 0.95$
 418 (95% fungal growth inhibition or higher), it can be declared that, in accordance to EMA guidelines
 419 (EMA 2010), each dosis (1 ml) of the most synergistic mixture against *C. albicans*, should contain
 420 65.96 μg of the whole mixture, composed of 18.84 μg of *ZpE* dry extract from *Z. punctata* Cav.
 421 aerial parts (9:1) and 47.12 μg of *LnE* dry extract from *L. nitida* Cav. aerial parts (12:1),
 422 corresponding to 5.39 μg of *ZpE* flavonoids markers and 23.63 μg of *LnE* lignans markers. In turn,
 423 each dosis (1 ml) of the most synergistic mixtures against *C. glabrata* should contain 168.23 μg of
 424 the whole mixture, composed of 45.47 μg of *ZpE* dry extract from *Z. punctata* Cav. aerial parts
 425 (8:1) and 122.76 μg of *LnE* dry extract from *L. nitida* Cav. aerial parts (10:1), corresponding to
 426 16.35 μg of *ZpE* flavonoids markers and 53.24 μg of *LnE* lignans markers.

427 *3D HPLC chromatograms*

428 The 3D HPLC profiles of the most synergistic mixtures against *C. albicans* (*ZpE-LnE* May) and
 429 *C. glabrata* (*ZpE-LnE* Feb) are shown in Fig. 7A and 7B respectively. This chromatogram was
 430 monitored at 254 nm, which allows to visualize both groups of compounds, lignans (λ_{max} 280 nm)
 431 and flavonoids (λ_{max} 267, 289 and 341 nm).

432 INSERT FIG. 7

433 *Conclusions*

434 Synergism studies with the statistical-based MixLow method, allowed us to give support to the
 435 jointly use of *ZpE* and *LnE* in traditional medicine. As the results of this study, we could determine
 436 that 3 over 4 fixed-ratio mixtures, whose composition were estimated by a valid method, showed
 437 synergism against *C. albicans* while only one showed synergism against *C. glabrata* (at $\phi > 0.5$).

438 Of them, one herbal-quantified *ZpE-LnE* preparation acting against *C. albicans* and one acting
 439 against *C. glabrata* are both of great interest for the development of an antifungal phytomedicine.

440

441 **Conflict of interest**

442 The authors declare they have no conflict of interest.

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449

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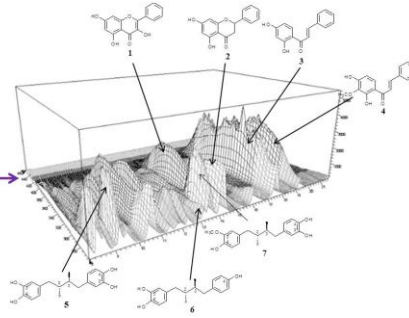
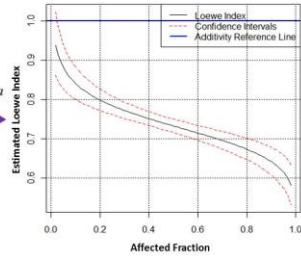
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ACCEPTED MANUSCRIPT

<i>Zuccagnia punctata</i> Cav. (Fabaceae)	
4 batches	
ZpE Feb	ZpE May
ZpE Sept	ZpE Nov
<i>Larrea nitida</i> Cav. (Zygophyllaceae)	
4 batches	
LnE Feb	LnE May
LnE Sept	LnE Nov

Candida strains



ACCEPTED MANUSCRIPT

Table 1

Identification data of markers **1-4** (*ZpE*) and **5-7** (*LnE*): Retention time (Rt), HRMS values and MS/MS fragments. Molecular Formulas (MF) and Molecular Weights (MW) are included.

Markers	Rt (min)	MF	MW	HRMS (error in ppm)				MS/MS fragments
				Feb	May	Sept	Nov	
<i>ZpE</i>								
1	12.9	C ₁₅ H ₁₀ O ₅	270.05	269.0451 (1.8)	269.0457 (-0.5)	269.0446 (3.7)	269.0446 (3.6)	269.0451, 169.0653, 171.0443, 195.0431, 143.0496
2	12.9	C ₁₅ H ₁₂ O ₄	256.07	255.0655 (3.1)	255.0656 (2.7)	255.0650 (4.9)	255.0649 (4.9)	255.0655, 227.0704, 151.0033, 123.0083, 107.0145
3	14.0	C ₁₅ H ₁₂ O ₃	240.08	239.0704 (3.9)	239.0709 (1.9)	239.0696 (4.8)	239.0703 (4.7)	239.0704, 197.0625, 169.0606, 148.0176, 135.0100
4	14.2	C ₁₆ H ₁₄ O ₄	270.09	269.0813 (2.2)	269.0814 (1.9)	269.0807 (4.4)	269.0812 (2.6)	269.0813, 254.0580, 150.9983, 106.0055, 94.0056
<i>LnE</i>								
5	6.2	C ₁₈ H ₂₂ O ₄	302.15	337.1207 (1.6)	337.1204 (2.5)	337.1213 (-0.3)	337.1201 (3.3)	302.1470, 273.1486, 122.0366
6	9.6	C ₁₈ H ₂₂ O ₃	286.16	321.1256 (2.0)	321.1249 (4.3)	321.1253 (3.1)	321.1253 (3.0)	123.0407, 122.0366, 108.0211
7	10.1	C ₁₉ H ₂₄ O ₄	316.17	351.1366 (0.8)	351.1358 (3.1)	351.1362 (1.9)	351.1363 (1.5)	300.1357, 149.0596, 135.0440, 122.0367

The Rt was determined by three individual analysis (n=3). The detected compounds had the greatest responses under the negative mode and so, the [M-H]⁻ was used as the precursor ion for **1-4**, and [M+Cl]⁻ for **5-8**.

Table 2Linear regression data, precision, LOD and LOQ of compounds **1-5**.

Comp.	λ (nm)	Linear regression data			Precision, RSD (%)			LOD (mg/ml)	LOQ (mg/ml)
		Regressive equation	R^2	Linear range (mg/ml)	Conc. (mg/ml)	Intra- day (n=3)	Inter- day (n=3)		
1	267	$y=39148.63x - 2097.40$	0.99	0.10-0.50	0.10	0.15	0.99	0.03	0.09
					0.30	0.24	1.00		
					0.50	0.98	1.20		
2	289	$y=33843.54x - 104.24$	1.00	0.05-0.60	0.05	0.22	0.83	0.01	0.04
					0.20	0.19	0.95		
					0.60	0.28	0.99		
3	341	$y=9045.88x - 89.18$	0.99	0.50-2.00	0.50	0.24	0.90	0.15	0.46
					1.00	0.78	1.16		
					2.00	0.58	1.53		
4	341	$y=10600.88x - 162.71$	1.00	0.10-1.00	0.10	0.57	0.82	0.01	0.03
					0.50	0.48	0.93		
					1.00	0.30	1.20		
5	280	$y=4149.35x - 304.55$	1.00	0.30-2.00	0.30	1.87	1.90	0.10	0.30
					0.50	0.61	0.80		
					2.00	0.68	0.99		

Table 3

Quantitative assessment (g of compound /100 g extract) of the seven markers **1-7** in the four batches of each *ZpE* and *LnE* (of February, May, September and November of one year) by HPLC-UV method. Values are the mean \pm Standard Deviation calculated from three replicates.

Markers	Feb	May	Sept	Nov
<i>ZpE</i>				
1	03.36 \pm 0.01	03.10 \pm 0.01	03.11 \pm 0.01	03.26 \pm 0.03
2	03.36 \pm 0.01	03.10 \pm 0.01	03.11 \pm 0.01	03.26 \pm 0.03
3	16.39 \pm 0.05	15.93 \pm 0.07	15.51 \pm 0.09	16.54 \pm 0.10
4	16.22 \pm 0.09	09.64 \pm 0.06	15.15 \pm 0.01	10.37 \pm 0.21
<i>LnE</i>				
5	10.39 \pm 0.01	16.80 \pm 0.05	12.48 \pm 0.11	19.09 \pm 0.08
6	09.14 \pm 0.02	08.53 \pm 1.03	09.66 \pm 0.02	10.24 \pm 0.01
7	23.84 \pm 0.07	24.81 \pm 0.10	27.37 \pm 0.09	28.46 \pm 0.02

Table 4

IC₅₀ values (µg/ml) of each extract alone obtained from the dose-response curves against *C. albicans* (*Ca*) and *C. glabrata* (*Cg*). For synergism studies, the ratio of concentrations tested (IC₅₀ *ZpE*/IC₅₀ *LnE*) is recorded. Each mixture was prepared from extract of the same period.

Fungal strain	Period	IC ₅₀ <i>ZpE</i> (µg/ml).	IC ₅₀ <i>LnE</i> (µg/ml).	Ratio tested (IC ₅₀ <i>ZpE</i> /IC ₅₀ <i>LnE</i>)
<i>Ca</i>	Feb	27.33 ± 0.01	53.14 ± 0.02	0.51
	May	31.40 ± 0.02	73.50 ± 0.01	0.43
	Sept	31.90 ± 0.03	57.20 ± 0.01	0.56
	Nov	28.63 ± 0.01	36.88 ± 0.01	0.78
<i>Cg</i>	Feb	65.79 ± 0.01	271.70 ± 0.03	0.24
	May	66.47 ± 0.01	164.72 ± 0.01	0.40
	Sept	39.56 ± 0.01	152.16 ± 0.01	0.26
	Nov	58.73 ± 0.01	154.31 ± 0.01	0.38

Amphotericin B was used as standard drug: IC₅₀ against *C. albicans* = 0.25 µg/ml; against *C. glabrata* = 0.25 µg/ml.

Table 5

L_{ϕ} values (combination indexes) and confidence intervals and type of interaction of the mixtures *ZpE-LnE* from different months of a year, against *C. albicans* (*Ca*) and *C. glabrata* (*Cg*) at fractions affected (ϕ) = 0.50, 0.80, 0.90 and 0.95 (inhibitory percentages 50, 80, 90 and 95 %).

Fungal strain	Combination	L_{ϕ} values at ϕ (inhibition percentage)							
		0.50 (50%)	Int (clasif)	0.80 (80%)	Int (clasif)	0.90 (90%)	Int (clasif)	0.95 (95%)	Int (clasif)
<i>Ca</i>	<i>ZpE-LnE</i> Feb	0.92 ± 0.02	Ad (n)	0.96 ± 0.02	Ad (n)	0.99 ± 0.03	Ad (n)	1.01 ± 0.04	Ad (n)
	<i>ZpE-LnE</i> May	0.73 ± 0.01	S (m)	0.68 ± 0.02	S (med)	0.65 ± 0.03	S (med)	0.62 ± 0.04	S (med)
	<i>ZpE-LnE</i> Sept	0.95 ± 0.02	Ad (n)	0.77 ± 0.02	S (m)	0.71 ± 0.03	S (m)	0.67 ± 0.04	S (med)
	<i>ZpE-LnE</i> Nov	0.89 ± 0.01	S (sl)	0.88 ± 0.01	S (sl)	0.87 ± 0.02	S (sl)	0.87 ± 0.02	S (sl)
<i>Cg</i>	<i>ZpE-LnE</i> Feb	0.91 ± 0.01	Ad (n)	0.77 ± 0.02	S (m)	0.71 ± 0.02	S (m)	0.67 ± 0.02	S (med)
	<i>ZpE-LnE</i> May	0.98 ± 0.01	Ad (n)	1.00 ± 0.01	Ad (n)	1.03 ± 0.02	Ad (n)	1.05 ± 0.02	Ad (n)
	<i>ZpE-LnE</i> Sept	1.21 ± 0.01	An (m)	1.13 ± 0.01	An (sl)	1.09 ± 0.02	Ad (n)	1.06 ± 0.02	Ad (n)
	<i>ZpE-LnE</i> Nov	1.20 ± 0.01	An (m)	1.15 ± 0.01	An (sl)	1.12 ± 0.02	An (sl)	1.09 ± 0.02	Ad (n)

Int: type of interaction; Clasif: classification of Synergism (S), Antagonism (An) and Additivism (Ad) in: slight (sl), moderate (m), medium (med), (n): near. *Ca*: *C. albicans* CCC 125; *Cg*: *C. glabrata* CCC 115.

Table 6

IC_x (Inhibitory Concentration to achieve X% of effect) of ZpE and LnE alone and in the May or February combination against *C. albicans* (*Ca*) and *C. glabrata* (*Cg*) respectively, and DRI values, at $\phi = 0.5, 0.8, 0.90$ and 0.95 (inhibitory percentages 50, 80, 90 and 95%).

Fungal strain	ϕ (X% effect)	I	II	III (= IV+V)	IV	V	VI	VII (I/IV)	VIII (II/V)
		IC _x ZpE alone (µg/ml)	IC _x LnE alone (µg/ml)	IC _x ZpE + IC _x LnE in the combination (µg/ml)	IC _x ZpE in the combination (µg/ml)	IC _x LnE in the combination (µg/ml)	L _{ϕ} = CI	DRI ZpE	DRI LnE
<i>Ca</i>	0.50 (50%)	31.40	73.50	38.94	11.80	27.14	0.73±0.01	2.66	2.71
	0.80 (80%)	45.91	98.28	49.91	14.26	35.65	0.68±0.02	1.29	2.76
	0.90 (90%)	57.33	116.48	57.70	16.48	41.22	0.65±0.03	3.48	2.83
	0.95 (95%)	70.36	136.22	65.96	18.84	47.12	0.62±0.03	3.73	2.89
<i>Cg</i>	0.50 (50%)	65.79	271.70	133.22	26.12	107.09	0.91±0.01	2.52	2.54
	0.80 (80%)	74.09	492.32	148.69	40.19	108.50	0.77±0.02	1.84	4.54
	0.90 (90%)	79.43	697.03	158.56	42.86	115.70	0.71±0.02	1.85	6.02
	0.95 (95%)	84.69	960.28	168.23	45.47	122.76	0.67±0.24	1.86	7.83

Table 7

Concentrations ($\mu\text{g/ml}$) of markers **1-7** in the most synergistic *ZpE-LnE* mixtures against *C. albicans* and *C. glabrata* respectively, at different ϕ .

ϕ	1 + 2	3	4	Total <i>ZpE</i> flavonoids	5	6	7	Total <i>LnE</i> lignans	Whole mixture
<i>ZpE-LnE</i> best synergistic mixture (May) against <i>C. albicans</i>									
0.50	0.34 ± 0.03	1.77 ± 0.02	1.07 ± 0.01	3.18 (8%)	4.67 ± 0.01	2.37 ± 0.04	6.90 ± 0.03	13.94 (36%)	38.94
0.80	0.44 ± 0.02	2.27 ± 0.01	1.37 ± 0.01	4.08 (8%)	5.99 ± 0.02	3.04 ± 0.04	8.85 ± 0.04	17.88 (36%)	49.91
0.90	0.51 ± 0.02	2.63 ± 0.01	1.58 ± 0.05	4.72 (8%)	6.92 ± 0.02	3.52 ± 0.03	10.23 ± 0.04	20.67 (36%)	57.70
0.95	0.58 ± 0.01	3.00 ± 0.04	1.81 ± 0.02	5.39 (8%)	7.92 ± 0.03	4.02 ± 0.05	11.69 ± 0.05	23.63 (36%)	65.96
<i>ZpE-LnE</i> best synergistic mixture (Feb) against <i>C. glabrata</i>									
0.50	1.28 ± 0.02	5.84 ± 0.03	5.80 ± 0.03	12.92 (9.7%)	10.12 ± 0.01	8.92 ± 0.02	23.04 ± 0.08	42.15 (31.6%)	133.22
0.80	1.35 ± 0.02	6.57 ± 0.03	6.52 ± 0.02	14.44 (9.7%)	11.27 ± 0.02	9.92 ± 0.02	25.86 ± 0.10	47.05 (31.6%)	148.69
0.90	1.44 ± 0.01	7.01 ± 0.01	6.96 ± 0.04	15.41 (9.7%)	12.02 ± 0.01	10.58 ± 0.01	27.58 ± 0.10	50.18 (31.6%)	158.56
0.95	1.53 ± 0.01	7.44 ± 0.02	7.38 ± 0.04	16.35 (9.7%)	12.75 ± 0.03	11.23 ± 0.01	29.26 ± 0.09	53.24 (31.6%)	168.23

Markers of *ZpE* are: **1** (galangin), **2** (pinocembrin), **3** (2',4'-dihydroxychalcone) and **4** (2',4'-dihydroxy-3'-methoxychalcone). Markers of *LnE* are: **5** (NDGA), **6** (DNDGA) and **7** (MNDGA). Compounds **1** and **2** had same Rt, so they could not quantify separately.

Scheme 1. Whole process of MixLow method: (a) Preparation of three dose-response curves: (a₁) one curve for each extract alone, confectioned with the data of each extract (*ZpE* and *LnE*). Estimation of parameter values; (a₂) Preparation of a dose-response curve for a fixed-ratio mixture of both extracts. Estimation of parameter values. (b) Use estimated parameters in calculating the Loewe Index (L_{\square}). (c) Generation of confidence intervals for L_{\square} , and obtaining an Affected fraction vs L_{\square} graph.

Fig. 1. Isolated compounds from *ZpE* (1-4), and for *LnE* (5-7).

Fig. 2. Design of a 96-well microplate used for analysis of antifungal activity of the extracts alone and mixtures.

Fig. 3. HPLC-UV chromatograms of four batches of each *Zuccagnia punctata* extract (*ZpE*) (left) at 254 nm and *Larrea nitida* extract (*LnE*) (right) at 280 nm. Marker compounds of *ZpE* and their Retention times (Rt, min) are: 5,7-dihydroxy-3-flavonol (galangin **1**) 12.9 min; 5,7-dihydroxy flavanone (pinocembrin **2**) 12.9 min; 2',4'-dihydroxychalcone (**3**) 14.0 min; 2',4'-dihydroxy-3'-methoxy chalcone (**4**) 14.2 min. Marker compounds of *LnE* and their Rt are: nordihydroguaiaretic acid (NDGA **5**) 6.2 min; 4-[4-(4-hydroxyphenyl)-2,3-dimethyl-butyl]-benzene-1,2-diol (DNDGA **6**) 9.6 min; 3'-methyl nordihydroguaiaretic acid (MNDGA **7**) 10.1 min.

Fig. 4. (A-D): Dose-response curves of single and combined *ZpE* and *LnE*, prepared with plants collected in four periods of one year against *C. albicans*. Mixtures *ZpE-LnE* were tested at the fixed ratio IC_{50ZpE}/IC_{50LnE} at 10 equi-spaced concentrations in the ln scale. (A'-D'): Affected fraction (ϕ) vs Estimated Loewe Index (L_{ϕ}) (full line) with 95% confidence interval (dotted lines). Full line at $L_{\phi} = 1$ represents additivity line. Lines below or above the additivity show synergism or antagonism, respectively.

Fig. 5. (A-D): Dose-response curves of single and combined *ZpE* and *LnE*, prepared with plants collected in four periods of one year against *C. glabrata*. Mixtures *ZpE-LnE* were tested at the fixed

ratio IC_{50ZpE}/IC_{50LnE} at 10 equi-spaced concentrations in the ln scale. (A'-D'): Affected fraction (ϕ) vs Estimated Loewe Index (L_ϕ) curves (full line) with 95% confidence interval (dotted lines). Full line at $L_\phi = 1$ represents additivity. Lines below or above $L_\phi = 1$ denote synergism or antagonism, respectively.

Fig. 6. Doses of the whole synergistic *ZpE-LnE* combinations and their content in marker compounds for achieving an effect level $\phi = 0.50, 0.80, 0.90$ and 0.95 against *C. albicans* and *C. glabrata*.

Fig. 7. 3D-HPLC profile of the two-herbal combination formed by *Z. punctata* and *L. nitida* in the concentration that produces 95% of growth inhibition. The representative peaks of the markers of both herbs were indicated. Peaks number **1-4** belong to *Z. punctata* and peaks number **5-7** belong to *L. nitida*. (A) *ZpE-LnE* mixture of May. (B) *ZpE-LnE* mixture of February. Detection was at 254 nm.

Scheme 1

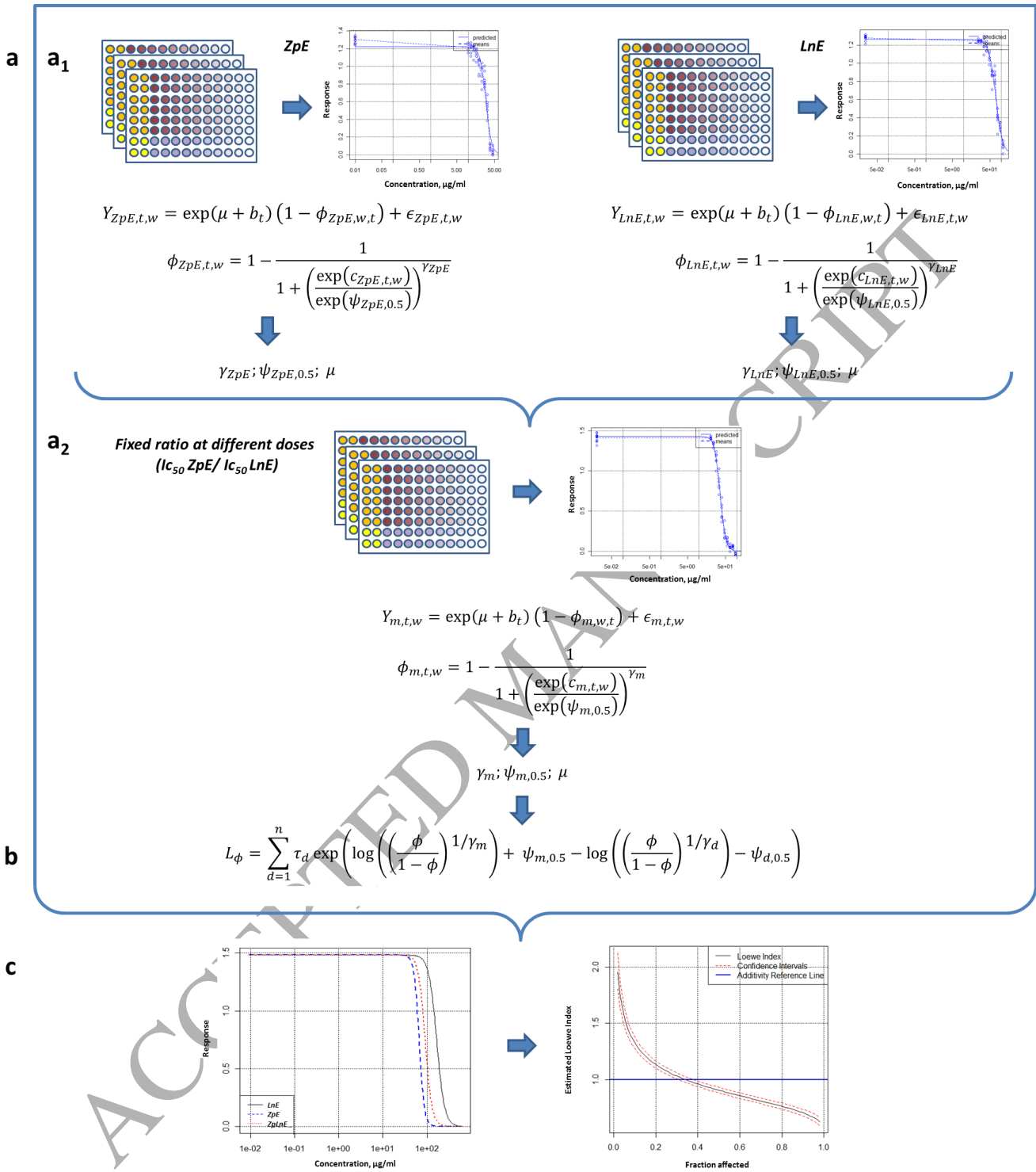
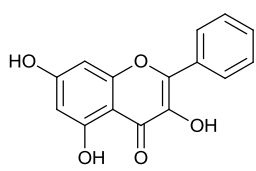
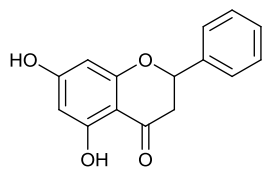


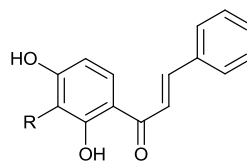
Fig. 1



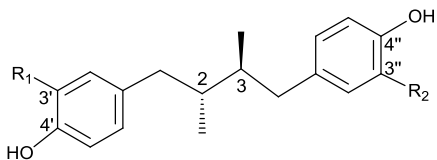
1
galangin



2
pinocembrin



3 R= H; 2',4'-dihydrochalcone
4 R= OCH₃; 2',4'-dihydroxy-3'-methoxychalcone



5 R₁= OH R₂= OH; nordihydroguaiaretic acid
6 R₁= OH R₂= H; 4-[4-(4-hydroxyphenyl)-2,3-dimethylbutyl]-benzene-1,2-diol
7 R₁= OCH₃ R₂= OH; 3'-methyl nordihydroguaiaretic acid

ACCEPTED MANUSCRIPT

Fig. 2

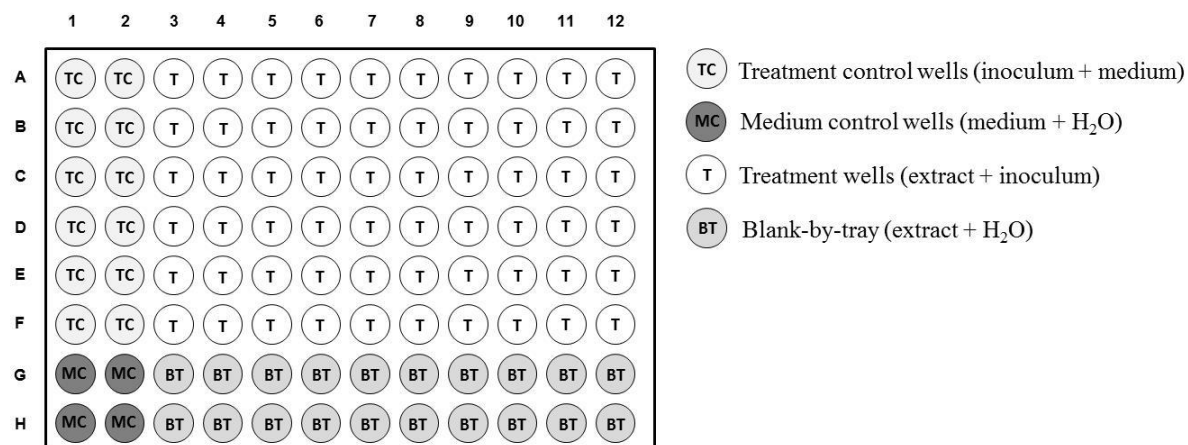


Fig. 3

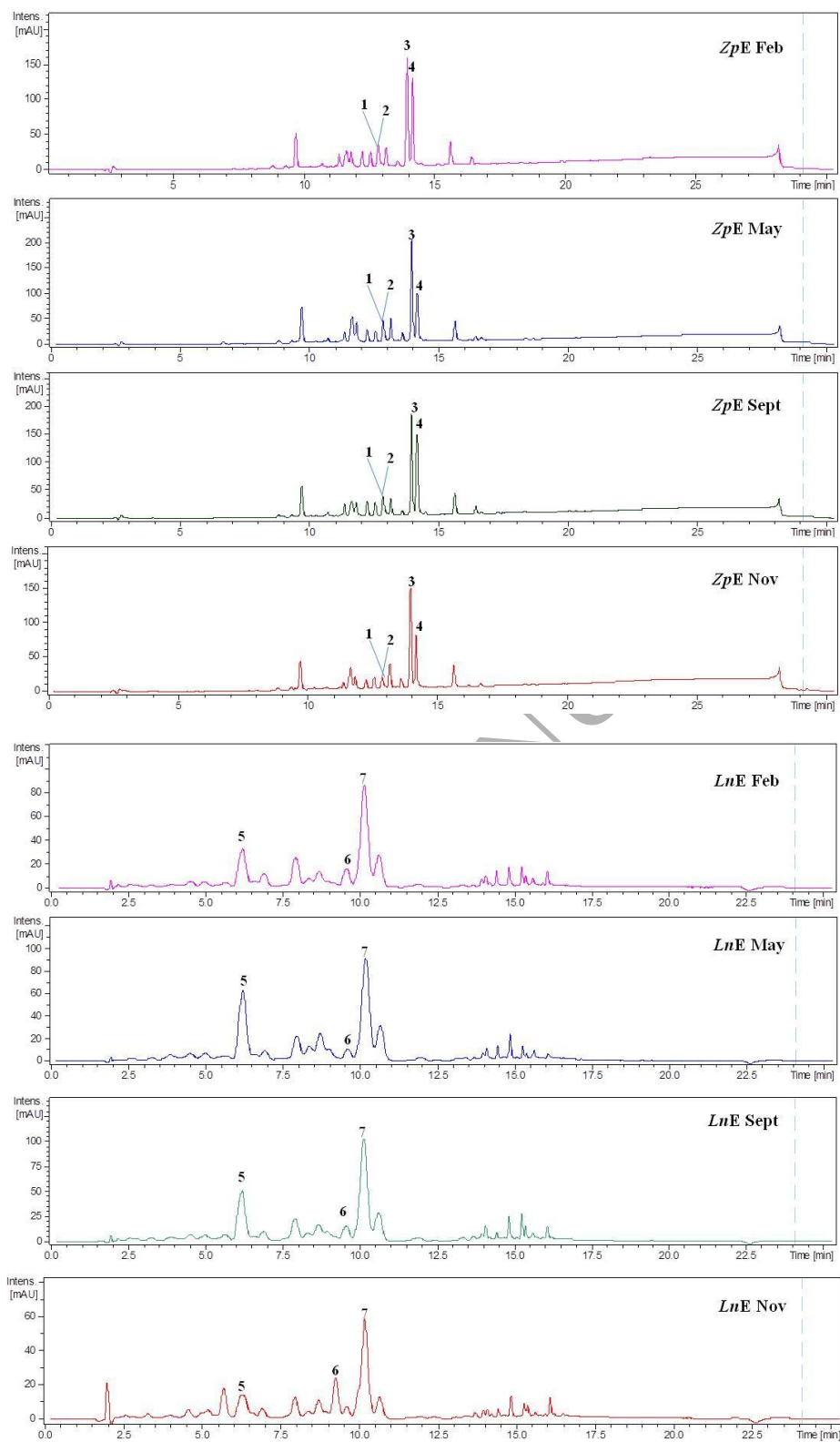


Fig. 4

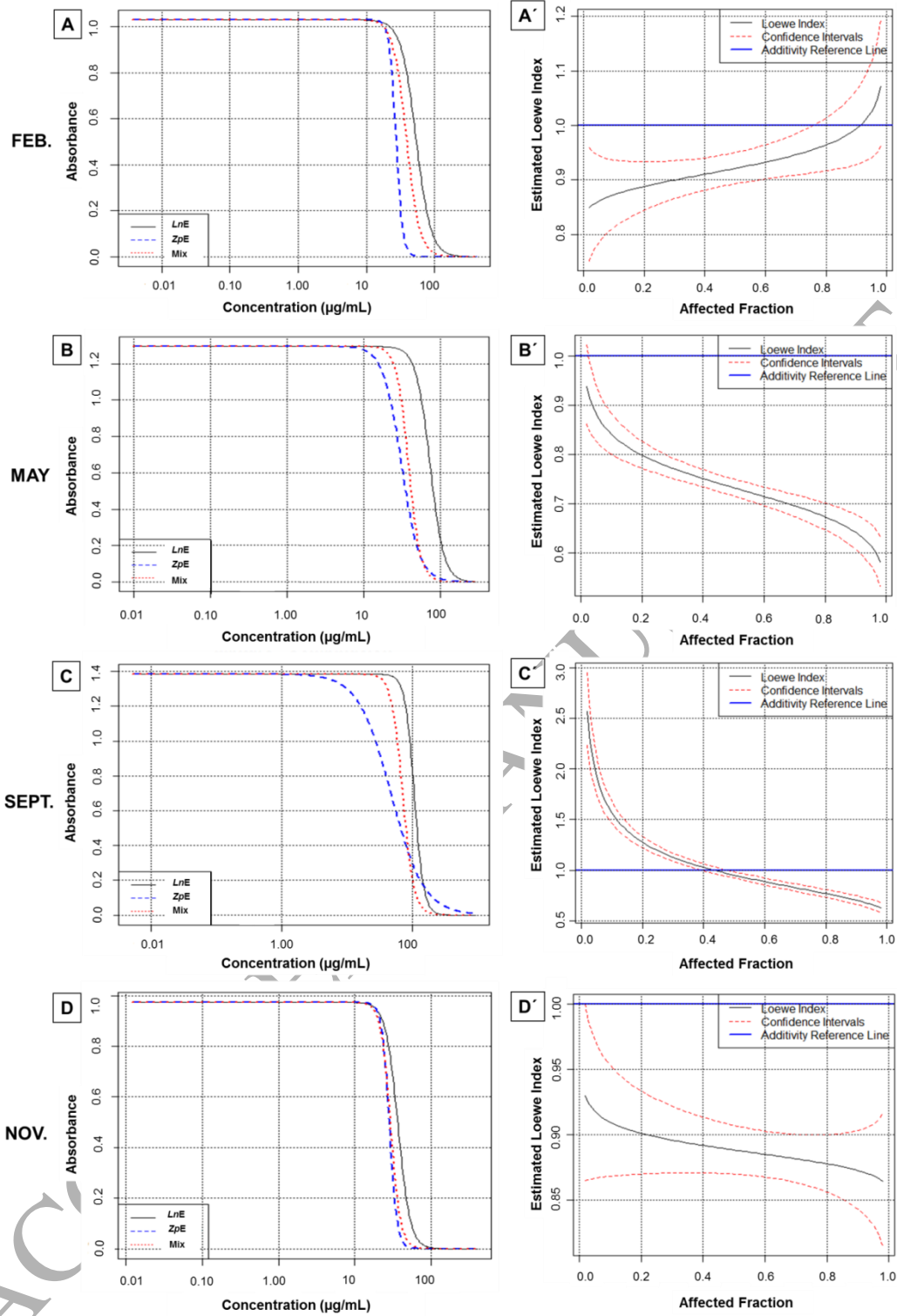


Fig. 5

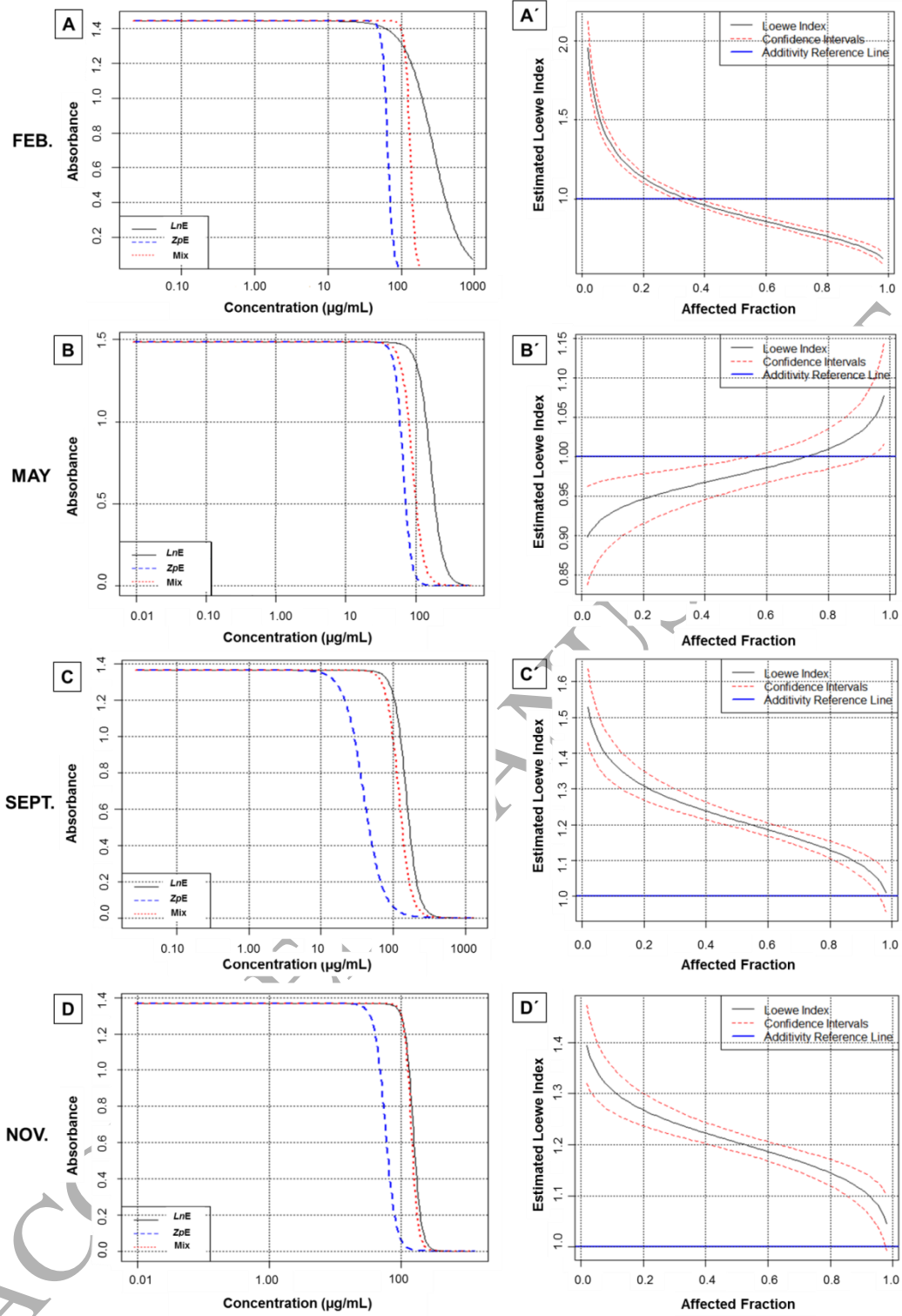


Fig. 6

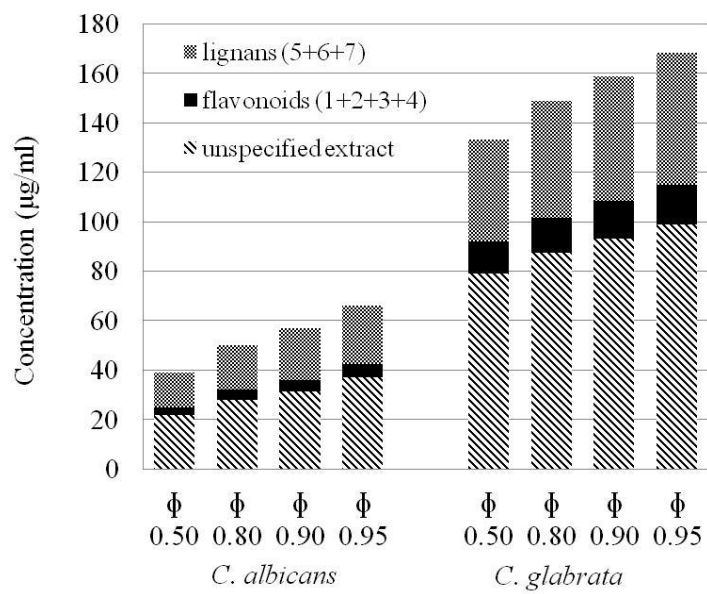


Fig. 7

