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1 Development of analytical methodologies to assess recalcitrant pesticide
2 bioremediation in biobeds at laboratory scale

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17

18

19 ABSTRACT:

20 To assess recalcitrant pesticide bioremediation it is necessary to gradually increase the
21 complexity of the biological system used in order to design an effective biobed
22 assembly. Each step towards this effective biobed design needs a suitable, validated
23 analytical methodology that allows a correct evaluation of the dissipation and
24 bioconversion. Low recovery yielding methods could give a false idea of a successful
25 biodegradation process. To address this situation, different methods were developed and
26 validated for the simultaneous determination of endosulfan, its main three metabolites,
27 and chlorpyrifos in increasingly complex matrices where the bioconverter
28 basidiomycete *Abortiporus biennis* could grow. The matrices were culture media, bran,
29 and finally a laboratory biomix composed of bran, peat and soil. The methodology for
30 the analysis of the first evaluated matrix has already been reported. The methodologies
31 developed for the other two systems are presented in this work. The targeted analytes
32 were extracted from fungi growing over bran in semisolid media YNB (Yeast Nitrogen
33 Based) with acetonitrile using shaker assisted extraction, The salting-out step was
34 performed with $MgSO_4$ and NaCl, and the extracts analyzed by GC-ECD. The best
35 methodology was fully validated for all the evaluated analytes at 1 and 25 $mg\ kg^{-1}$
36 yielding recoveries between 72 and 109% and RSDs <11% in all cases. The application
37 of this methodology proved that *A. biennis* is able to dissipate 94 % of endosulfan and
38 87 % of chlorpyrifos after 90 days. Having assessed that *A. biennis* growing over bran
39 can metabolize the studied pesticides, the next step faced was the development and
40 validation of an analytical procedure to evaluate the analytes in a laboratory scale
41 biobed composed of 50 % of bran, 25 % of peat and 25 % of soil together with fungal
42 micelium. From the different procedures assayed, only ultrasound assisted extraction
43 with ethyl acetate allowed recoveries between 80 %-110 % with RSDs <18 %.

44 Linearity, recovery, precision, matrix effect and LODs/LOQs of each method were
45 studied for all the analytes: endosulfan isomers (α & β) and its metabolites (endosulfan
46 sulfate, ether and diol) as well as for chlorpyrifos. In the first laboratory evaluation of
47 these biobeds endosulfan was bioconverted up to 87 % and chlorpyrifos more than 79 %
48 after 27 days.

49 Keywords: Basidiomycete; Recalcitrant pesticides; Bioremediation; Method
50 validation

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54 1. Introduction

55 Environmental pollution caused by hazardous wastes containing recalcitrant
56 xenobiotic chemicals has become a major problem that threatens the sustainability of
57 the ecosystems as well as human health. Unlike the naturally occurring organic
58 compounds that are readily degraded upon introduction into the environment, some of
59 these synthetic chemicals are extremely resistant to biodegradation by native
60 microorganisms. Particularly, surface and ground waters are exposed to diffuse
61 pollution due to pesticides via percolation, runoff, drainage and drift (i.e. movement of
62 airborne spray droplets) as well as spills during equipment washing. These critical
63 points are the main sources of soil and water contamination, contributing significantly
64 to the deterioration of natural water sources [1, 2-5]. This is an issue of global strategic
65 importance as groundwater represents about 98 % of the available fresh water of our
66 planet [6]. Several field surveys and measurement campaigns on a catchment scale have
67 demonstrated that 40–90 % of surface water contamination by pesticides is attributable
68 to direct losses of the active ingredients [1, 3-5]. To protect ground and surface water
69 quality, practical solutions to minimize the pesticide entrance into hydrographic
70 networks are continuously investigated. One of the possible strategies to minimize it is
71 to degrade the pesticides before releasing the machinery washings and container triple
72 rinses to the environment.

73 As the degradation of recalcitrant pesticides occurs slowly under natural conditions,
74 the process has to be enhanced.

75 A newly explored route to diminish pesticide spillage into waters or in places where
76 it can lixiviate to ground water is the biobeds technology. Biobeds are environmentally
77 friendly solutions for the remediation of impacted zones as the pesticides convertor and
78 degrading agents are saprophytes microorganisms. This methodology has been used in

79 Europe for several years and it is now being evaluated in South America as a friendly
80 alternative for remediation of recalcitrant compounds [7, 8]. Ligninolytic fungi are
81 biodegradators of particular interest because they have shown to degrade and mineralize
82 a large variety of recalcitrant compounds due to the nonspecificity of their enzyme
83 machinery [9, 10]. White rot fungi produce a number of extracellular oxidative enzymes
84 including laccases, lignin peroxidases and manganese peroxidases, which are normally
85 involved in the breakdown of the plant structural material lignin [11] and are also
86 responsible for the degradation of xenobiotics.

87 Challenging targets to assess biobeds suitability are organochlorine (OC)
88 agrochemicals. They are persistent compounds which have been accumulating in the
89 biosphere after decades of massive application in agriculture. Among them, Endosulfan,
90 is an actually banned OC which had been used in many countries until recent years,
91 leaving highly contaminated zones as an unsolved problem. In addition, there are also
92 huge amounts of already synthesized endosulfan that have to be stored, waiting for a
93 final destination. At the same time, massive use of highly toxic endosulfan, has been
94 substituted by chlorpyrifos situation that renders an even more complicated scenario.

95 The proof of the efficacy of a bioremediation process is sustained on validated,
96 highly sensitive analytical methods that assess the efficiency of the whole process.
97 Agrochemical biodegradation has to be assayed in the laboratory, with microorganisms
98 growing in conventional culture media, in order to select the microorganisms capable of
99 dissipating them. Once the microorganisms have been selected and the efficiency of the
100 transformation process evaluated, trial biobeds made of soil, peat and bran are assayed
101 at a lab scale. Dissipation kinetics within the bioreactor is established, and the residual
102 water that lixivates through it is evaluated for pesticide non detection [12].

103 In a previous communication our group described ligninolytic fungi capable of
104 degrading recalcitrant pesticides using endosulfan as a model compound in culture
105 media, highlighting the importance of having analytical methods that can assure the
106 effective dissipation of the pesticides into harmless products [13]. This work presents
107 the advances in the development of a bioreactor using endosulfan and chlorpyrifos as
108 model compounds, based in the Swedish biobed design [12] and native Basidiomycetes
109 fungi as bioconvertors.

110 For each of the three different complex matrices employed: culture media, bran and
111 the biobmix where the fungal mycelium grows and the biotransformation is performed,
112 different methodologies were developed. Linearity, recovery, precision, matrix effect
113 and LODs/LOQs, were determined for each of the endosulfan isomers (α & β) and some
114 of its metabolites (endosulfan sulfate, endosulfan ether and diol) as well as for
115 chlorpyrifos. Although chlorpyrifos metabolites, particularly TCP (3,5,6-
116 trichloropyridinol) [14], is toxicologically important from an environmental point of
117 view [15], only the dissipation of the parent compound as a model has been followed in
118 this study. At this point, our interest was focused in the improvement of the biobed
119 design and future work will need to study TCP degradation in the final biobed.

120

121 2. Materials and methods

122

123 2.1. Standards and reagents

124

125 Analytical grade organic solvents, pesticide residues free were purchased from Merck
126 (Darmstadt, Germany). Pesticide standards and the internal standard were from Dr.
127 Ehrenstorfer (Augsburg, Germany, 99 %). The culture media were provided by Difco.

128 Bran and land peat were commercially available. Magnesium Sulfate and Sodium
129 Sulfate p.p.a from Sigma-Aldrich (St.Louis, MO, USA); Sodium Chloride from Merck
130 (Darmstadt, Germany), Silica: MN Kiesel 60 from J.T. Baker (State of Mexico).
131 Stock solutions were prepared from the standard substances at 1000 and 2000 mg L⁻¹ in
132 ethyl acetate. Working standard mixtures were prepared by appropriately diluting the
133 stock solutions with ethyl acetate. All solutions were stored at 4°C.

134

135 2.2 Apparatus and experimental conditions.

136

137 Gas chromatographic (GC) analyses were performed using a Shimadzu GC 17A
138 equipped with an ECD detector and a PTV injector using internal standard method. All
139 compounds were resolved on a capillary column Mega 68 (30m, 0.32 mm ID, 0.25 µm
140 film thickness) Mega Legnano (Italy). The experimental conditions were as follow:
141 PTV, 60°C (0.3min), then 5°C min⁻¹ to 280°C (40min). Oven temperature, 100°C
142 (3min), 100-180°C at 10°C min⁻¹, 180°C (15min), then 180-270°C at 5°C min⁻¹, 270°C
143 (10 min). Detector temperature, 280°C.

144 Orbital shaker: SL1 Stuart (Staffordshire,UK). Ultrasonic bath:
145 Elma®Transsonic T460/H. Centrifuge IEC: HNS-II (U.S.A.).

146

147 2.3 Microbiological matrix preparation for validation study

148

149 2.3.1 Matrix A (Bran)

150 The prepared inoculum was added to a mixture of 5 mL of semisolid medium
151 YNB (Yeast Nitrogen Base) and 2.2 g of bran which was previously homogenized in a
152 water bath for 90 min at 45°C. Then, it was incubated at 28±2°C for 20 days.

153 Inoculum: A portion of 1 cm diameter of the external growth of preinoculum was added
154 to 10 mL of Malt Extract medium and incubated at 28 ± 2 °C for 10 days.

155 Preinoculum: *Abortiporus biennis* was cultured in solid media Potato Dextrose Agar
156 (PDA) at 28 ± 2 °C for 5 days.

157

158 2.3.2 Matrix B (Biobed)

159 The laboratory scale biobed was prepared by adding 20 times matrix A to 1 kg
160 of a mixture containing 50 % bran, 25 % soil and 25 % peat and incubated for 30 days
161 at room temperature.

162

163 2.4 Recovery tests

164

165 2.4.1 Recovery tests for matrix A

166 For recovery studies the matrix was prepared spiking at two levels: 1 and 25 mg
167 kg^{-1} with chlorpyrifos, endosulfan and its metabolites. Levels 1 and 25 mg kg^{-1} were
168 prepared in quintuplicate by adding 0.3 and 3.5 mL respectively of a solution containing
169 100 mg L^{-1} of endosulfan α and β , endosulfan ether, endosulfan diol, endosulfan sulfate
170 and chlorpyrifos to matrix A.

171

172 2.4.2 Recovery tests for matrix B

173 For recovery studies the matrix was prepared spiking at three levels: 1, 25 and
174 50 mg kg^{-1} with endosulfan and its metabolites and 1, 30 and 60 mg kg^{-1} with
175 chlorpyrifos. Level 1 mg kg^{-1} was prepared in quintuplicate by adding 0.05 mL of a
176 solution containing 200 mg L^{-1} of chlorpyrifos, endosulfan α and β , endosulfan ether,
177 endosulfan diol and endosulfan sulfate to grown fungi in 10 g of matrix B (laboratory

178 biobed); level 25 and 50 mg L⁻¹ for endosulfan and its metabolites were prepared by
179 adding 0.12 mL and 0.25 mL respectively of a 2025 mg L⁻¹ solution and level 25 and 60
180 mg L⁻¹ for chlorpyrifos were prepared by adding 0.12 mL and 0.30 mL respectively of
181 a 2020 mg L⁻¹ solution.

182

183 2.5 Extraction and clean up methods comparison

184

185 2.5.1 Bran (Matrix A)

186 The evaluated methods were:

187 1. Extraction with ethyl acetate and homogenization in Stomacher® and clean up using
188 open column packed with silica and sodium sulfate.

189 2. Extraction with ethyl acetate and homogenization in orbital shaker for 24 h and clean
190 up using open column packed with silica and sodium sulfate.

191 3. Extraction with acetonitrile and water, homogenization in orbital shaker for 2 h and
192 salting out with NaCl and MgSO₄.

193 The comparison of the recovery results obtained with the three extraction
194 methods tested was performed spiking at level 50 mg L⁻¹ of endosulfan and 60 mg L⁻¹
195 of chlorpyrifos.

196 2.5.1.1. General aspects for all the extraction procedures for Matrix A:

197 The internal standard (IS) used was methyl bromophos, the final aliquot was
198 evaporated with a gentle stream of nitrogen, redissolved with IS and diluted to volume
199 with ethyl acetate to obtain a final vial concentration of 1 mg L⁻¹ of the IS and the
200 evaluated. The chromatographic analysis was performed in a GC-ECD under the
201 conditions explained in “Apparatus and experimental conditions”.

202

203 Extraction and clean up methods for matrix A are summarized in table 1 so that they can
204 be easily compared.

205

206 Table 1.

207

208

209 2.5.2 Laboratory scale Biomix (Matrix B)

210 The compared methods were:

211 1. Extraction with acetonitrile and water, homogenization in orbital shaker and salting-
212 out with magnesium sulfate and sodium chloride.

213 2. Extraction with ethyl acetate and homogenization in ultrasonic bath (ultras) for 15
214 min.

215 3. Extraction with acetonitrile and homogenization in ultrasonic bath for 15 min.

216 4. Extraction with ethyl acetate and homogenization in shaker with iron pellets
217 assistance.

218 5. Extraction with ethyl acetate and homogenization in ultrasonic bath (3 cycles of 15
219 min. each one).

220 6. Extraction with ethyl acetate in soxhlet apparatus.

221

222 2.5.2.1 General aspects for all the extraction procedures for Matrix B:

223 Sample humidity was adjusted with a solution of NaCl (10 %). The internal
224 standard (IS) used was methyl bromophos. The final aliquot was evaporated with a
225 gentle stream of nitrogen and redissolved with a solution of the IS in ethyl acetate and
226 diluted to volume with ethyl acetate in a 25 mL volumetric flask obtaining a final vial

227 concentration of 1 mg L^{-1} of the IS. The chromatographic analysis was performed in a
228 GC-ECD under the conditions explained in “Apparatus and experimental conditions”.
229 Extraction methods for matrix B are summarized in table 2 so that they can be easily
230 compared.

231 Table 2.

232 (*) Addition of 1 iron pellet per g of sample.

233 (**) 3 cycles of 15 min each one and manual agitation between cycles.

234

235 2.6 Sample preparation for endosulfan/its metabolites and chlorpyrifos analysis.

236

237 2.6.1 Final method for Bran (Matrix A)

238

239 Matrix A was finally analyzed using method 3 summarized under “Extraction
240 and clean up methods comparison, Bran (Matrix A)”:

241 The whole flask of rice straw with the fungi grown in semisolid media was extracted
242 with 15 mL acetonitrile and 15 mL water in an orbital shaker apparatus for two hours.

243 Then, 20g of MgSO_4 and 2g NaCl were added and placed in an orbital shaker for 5

244 hours. Finally it was centrifuged 10 min at 3000 rpm and an aliquot of 0.5 mL of the

245 extract was evaporated with a gentle stream of nitrogen, redissolved with IS and diluted

246 to volume with ethyl acetate to obtain a final vial concentration of 1 mg L^{-1} of the IS

247 and analytes evaluated. The chromatographic analysis was performed in a GC-ECD

248 under the conditions described in “Apparatus and experimental conditions”.

249

250 2.6.2 Final Method for the laboratory scale Biomix (Matrix B)

251

252 Matrix B was finally analyzed using method 5 summarized in “Laboratory scale
253 Biomix analysis (Matrix B)”:

254

255 10 g of biobed mix and inoculum (Microbiological matrix preparation for
256 validation study: Matrix B (Biobed)) were placed in a 250 mL Erlenmeyer flask and 10
257 mL solution of 10 % NaCl and 50 mL of ethyl acetate were added and extracted in an
258 ultrasonic bath (3 cycles of 15 min each one). The final aliquot (0.5 mL) was evaporated
259 with a gentle stream of nitrogen and redissolved with a solution of the IS in ethyl
260 acetate and diluted to volume with ethyl acetate in a 25 mL volumetric flask obtaining a
261 final vial concentration of 1 mg L^{-1} of the IS and analytes evaluated. The
262 chromatographic analysis was performed in a GC-ECD under the conditions explained
263 in “Apparatus and experimental conditions”.

264

265 2.7. Methods validation

266 2.7.1 Matrix A

267

268 The following parameters were evaluated for the analytical method validation:
269 linearity, recoveries, repeatability (RSD_r), within-laboratory reproducibility (RSD_{wR}),
270 LOQs and LODs. All the tests were performed at two levels, five replicates in three
271 different days. Solvent and matrix matched calibration curves were compared and
272 matrix effects were quantified. Matrix effects were evaluated at two different amounts
273 of matrix 0.5 and 0.02 g mL^{-1} of extract which correspond to the dilutions needed to
cover the wide range of concentrations 1 to 25 mg L^{-1} .

274 2.7.2 Matrix B

275 The parameters evaluated for the analytical method validation exposed in the
276 item “Methods validation, Matrix A” were evaluated in the same conditions. Matrix

277 effects were evaluated at three different amount of matrix 0.2, 0.07 and 0.03 g mL⁻¹ of
278 extract which corresponds to the dilutions needed to cover the wide range of
279 concentrations 1 to 50 and 60 mg L⁻¹ for endosulfan and chlorpyrifos respectively.

280 The different parameters evaluated were calculated as explained below:

281 RSD_r presented are an estimation of the precision of the method. They were calculated
282 as the relative standard deviation (coefficient of variation) of the five replicates from the
283 recovery test measurement of each analyte, obtained using the same method on the same
284 sample in a single laboratory over a short period of time, during which differences in
285 the materials and equipment used and the analysts involved did not occur.

286 RSD_{wR} presented are within laboratory reproducibility. They were calculated as
287 the relative standard deviation (coefficient of variation) of all the replicates from the
288 recovery test measurement of each analyte obtained using the same method in a single
289 laboratory, by different analysts, and over a period in which differences in the materials
290 and equipment occurred.

291 LOQs limits of quantitation (quantification) presented are the lowest
292 concentration of the analyte that has been validated with acceptable accuracy
293 (recoveries between 70-120% and RSDs < 20%) by applying the complete analytical
294 method as recommended by DG-SANCO [16].

295 LODs limits of detection presented are the lowest concentration of the analyte that
296 presents a response with a signal to noise ratio equal or above 3 by applying the
297 complete analytical method.

298

299

300

301 3. Results and Discussion

302

303 The global strategy followed to accomplish the biobed design at laboratory scale is
304 based in the developing of fit for purpose analytical methodologies for very complex
305 and heterogeneous matrices, in which the pesticides and their metabolites have to be
306 determined.

307 The Swedish biobed is composed of a mixture of peat, straw and soil (1:2:1). As the
308 selected fungi did not grow properly over straw to yield enough biomass, we looked for
309 other fungi grow promoting substrates. It has been reported that straw can be replaced
310 by other agricultural byproducts with equal or superior biotransformation capacities
311 [17].

312 Looking for straw alternative substrates, good biomass amounts were obtained with
313 the fungi growing over cereal bran, which is a more nutritive substratum. Therefore, the
314 bioconversion of the targeted pesticides by fungi growing over bran was assayed and
315 the best analytical methodology for this matrix was selected and validated. The
316 pesticides to evaluate the performance of the bioreactor were chosen based on their use
317 in agriculture their persistence and their toxicity [18-20].

318

319 3.1. Methods comparison

320 3.1.1. Bran (Matrix A)

321

322 Three methods were evaluated aiming to find a good methodology to analyze
323 chlorpyrifos, endosulfan and its metabolites in the microbiological media prepared in
324 the laboratory with bran as the major component of the system. The combination of
325 different extraction parameters (solvents, homogenization methods and different
326 shaking systems to assist the extraction) and clean up strategies allowed the selection of

327 a method with recoveries between 70-120 % for endosulfan, its studied metabolites and
328 chlorpyrifos. Method 3 presents the best recoveries of the analytes as shown in Table 3.
329 The other methodologies evaluated did not accomplish DG-SANCO guidelines criteria
330 for pesticide residue analysis [16], obtaining recoveries under our laboratory conditions
331 $< 70\%$ or $> 200\%$ with RSDs between 1 and 10 %.

332 Table 3.

333

334 3.1.2. Laboratory scale biomix (Matrix B)

335 As for matrix A, different methods were tested. This matrix required six method
336 comparisons due to its increased chemical complexity as soil, peat and bran were
337 together in the biomix (Table 4).

338 Method 5 presented the best results probably due to the formation of a biphasic
339 system through water addition and ultrasound assisted extraction which also made the
340 extraction time shorter. Also, ethyl acetate proved to have better extractability of the
341 analytes than acetonitrile (used in Methods 1 and 3). Traditional methods such as
342 soxhlet (Method 6) provide low recoveries which would lead to a overestimation of the
343 bioconversion. Performing a second extraction (Method 1) gave a large load of co-
344 extractives and unacceptable recoveries above 120 %.

345

346 Table 4.

347

348 3.2. Method validation.

349

350 3.2.1. Matrix A

351 The selected method yielded good results for endosulfan, its studied metabolites
352 and chlorpyrifos (Table 5); recoveries were between 72-109 % at all concentration
353 levels studied. The method was precise: its repeatability (RSD_r) was below 11% and its
354 within-laboratory reproducibility (RSD_{wR}) is below 17 % for all analytes at all the
355 evaluated levels, according to DG-SANCO guidelines criteria for pesticide residue
356 analysis [16]. The determined LOQs and LODs were adequate for the purpose of the
357 method: the quantification of chlorpyrifos and endosulfán, residues as well as its
358 metabolites in a fungi growing in cereal bran in semisolid media. The values for LOQs
359 and LODs are 1 mg kg⁻¹ and 0.3 mg kg⁻¹ respectively. It was not possible to perform the
360 sub sampling and sample comminution steps prior to solvent addition in order to
361 achieve good accessibility of the solvent to the analytes because the fungi grows
362 heterogeneously.

363

364

Table 5.

365

366 The fungal mycelium and the components of bran are responsible for marked
367 matrix effects when the amount of matrix injected is higher, being very important for
368 experiments where the initial endosulfan concentration was 1 mg/kg and at 25 mg kg⁻¹
369 (Table 6). Matrix-matched calibration curves were used for quantitation at all
370 concentration levels because the influence of the amount of matrix on the analyte
371 response was very heterogeneous. The recommendation is to perform the quantitation
372 step using matrix matched calibration curves when assessing pesticide dissipation and
373 metabolites generation with a blank extract prepared in the same way as the samples.

374

375

Table 6.

376

377 One of the advantages of this method is that as the sample contained (fungi +
378 semisolid culture medium + bran) is completely extracted, no sub sampling is
379 necessary, so all problems related to representativeness are avoided. Also this
380 characteristic improves the results in terms of repeatability and reproducibility.

381

382 3.2.2. Matrix B

383 The results obtained with the best method for chlorpyrifos, endosulfan, and the
384 different metabolites assayed are presented in Table 7. Recoveries were between 71-110
385 % at all concentration levels studied. The method was precise: its repeatability (RSD_r)
386 was below 17 % and its within-laboratory reproducibility (RSD_{wR}) is below 18 % for
387 all analytes at all the evaluated levels, according to DG-SANCO guidelines criteria for
388 pesticide residue analysis [16]. The determined LOQs and LODs were adequate for the
389 purpose of the method: the quantification of endosulfan, chlorpyrifos residues and its
390 metabolites for a fungi growing in a laboratory biobed. The values for LOQs and LODs
391 are 1 mg kg⁻¹ and 0.3 mg kg⁻¹ respectively.

392

393

Table 7.

394

395 Matrix effects vary with the different analytes (Table 8), for example,
396 endosulfan sulfate showed the highest matrix effect at all concentrations but for
397 endosulfan ether it was zero. The recommendation is to always quantify with matrix
398 matched calibration curves when assessing endosulfan, its metabolites or chlorpyrifos
399 dissipation. These curves are generated with a blank extract prepared in the same way as
400 the samples injecting the same amount of matrix co-extractives.

401

402

Table 8.

403

404 Ultrasound assisted extraction allowed better solvent accessibility to the
405 analytes. The method has few steps and is easy to perform in the laboratory considering
406 the complexity of the matrix composed of a microbiological organism grown in a
407 biobed degrading recalcitrant pesticides.

408 A global strategy to study bioremediation of contaminants at laboratory scale, based
409 on the precise knowledge of the concentration of pesticides present in the biobed is
410 presented. Two very employed; semipersistent to persistent pesticides were used as
411 models. The extraction and clean up methodologies used for the sample preparation of
412 matrix A and B are adequate for the extraction of chlorpyrifos, endosulfan and its
413 metabolites as it was shown during the validation steps. These methods allow the study
414 of the proposed bioreactor and provide tools for studying biotransformation processes of
415 endosulfan, its metabolites and chlorpyrifos by GC-ECD. Exploratory trials of
416 endosulfan and chlorpyrifos dissipation at laboratory scale biobeds yielded 87 % for
417 endosulfan and 79% for chlorpyrifos degradation after 27 days. Figure 1 shows the
418 chromatogram of the residual pesticides in the biobed after 9 and 27 days of
419 bioconversion respectively, proving the viability of the proposed bioreactor to degrade
420 the agrochemicals under study.

421 Further work is in progress seeking to optimize the conditions of the bioreactor. The
422 results obtained are a step forward in the search of an environmentally friendly tool to
423 diminish the impact of recalcitrant compounds in affected areas.

424

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497 Captions for Figures and Tables

498 Table 1. Extraction and clean up methods comparison for matrix A.

499 Table 2. Extraction methods comparison for matrix B.

500 Table 3. Recoveries (Rec (%)) comparison for the evaluated methods to determine

501 endosulfan and chlorpyrifos and their respective RSDs (%) in matrix A.

502 Table 4. Recoveries (%) and their respective RSDs (%) for the 6 evaluated methods in

503 matrix B.

504 Table 5. Recoveries (%), repeatability (RSDs %) and within-laboratory reproducibility

505 (RSDwR %) of the evaluated pesticides in matrix A.

506 Table 6. Matrix effect for chlorpyrifos, endosulfan and its metabolites at 1 and 25 (mg

507 kg^{-1}) by GC-ECD in matrix A.

508 Table 7. Recoveries (%) and RSDs (%) of the evaluated pesticides for the analytical

509 method developed in matrix B; levels 1, 25 and 50 mg kg^{-1} for endosulfan and510 metabolites and levels 1, 25 and 60 mg kg^{-1} for chlorpyrifos

511 Table 8. Matrix effects for endosulfan, chlorpyrifos and its metabolites at 1, 25, 50 mg

512 kg^{-1} by GC-ECD in matrix B. 1, 25 and 50 mg kg^{-1} for endosulfan and its metabolites513 and 1, 25 and 60 mg kg^{-1} for chlorpyrifos.

514 Figures:

515 Figure 1. GC-ECD chromatogram of the chlorpyrifos and endosulfan residues in labs

516 biobeds after 9 and 27 days, showing their degradation. Black trace: residues after 9

517 days; purple trace: residues after 27 days.

518

519

520 Table 1.

521

	Method 1	Method 2	Method 3
522			
523 Sample (g)	15	15	15
524 Solvent, volume (mL)	AcOEt, 40	AcOEt, 40	MeCN, 15
525 Water addition (mL)	-	-	15
526 Agitation/time (h)	Stomacher/0.08	Shaker/24	Shaker/2
527 Clean up: NaSO ₄ /Silica column (g)	20 / 30	20 /30	-
528 Elution: solvent, volume (mL)	AcOEt, 100	AcOEt, 100	-
529 Salting out: NaCl/MgSO ₄ (g)	-	-	2/20
530 Agitation/time(h)	-	-	Shaker/5
531 Centrifugation (min)/(rpm)	-	-	10/3000
532 Final aliquot (mL)	0.1	0.1	0.5

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535 Table 2.

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537		Method1	Method2	Method3	Method4	Method5	Method 6
538	Sample (g)	15	15	15	10	10	10
539	Humidity (%)	40	40	40	70	70	40
540	Solvent (mL)	MeCN(30)	AcOEt(30)	MeCN(30)	AcOEt(50)	AcOEt(50)	AcOEt(250)
541	Water (mL)	15	-	-	*	-	-
542	Extraction	shaker	ultras	ultras	shaker	ultras	soxhlet
543	Time (h)	2	0.15	0.15	5	0.45**	2.30
544	MgSO ₄ /NaCl	20(g)/2(g)	-	-	-	-	-
545	Solvent (mL)	MeCN(30)	-	-	-	-	AcOEt(50)
546	Shake (h)	5	-	-	-	-	-
547	Centrifugation (min/rpm)		10/3000	-	10/3000	10/3000	-
548	Aliquot (mL)	0.5	0.5	0.5	0.5	0.5	0.5

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551 Table 3.

552

553	554	Endosulfan α			Endosulfan β			Chlorpyrifos		
		Rec	RSDr	RSDwR	Rec	RSDr	RSDwR	Rec	RSDr	RSDwR
555		(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)
556	Method 1	41	3	6	43	4	9	201	20	10
557	Method 2	63	2	3	65	1	1	205	12	6
558	Method 3	83	3	4	91	3	4	76	8	10

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579 Table 4.

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581		Endosulfan α			Endosulfan β			Chlorpyrifos		
		Rec	RSDr	RSDwR	Rec	RSDr	RSDwR	Rec	RSDr	RSDwR
582		(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)
584	Method 1	170	-	-	169	-	-	220	-	-
585	Method 2	111	1	1	115	2	1	172	0	0
586	Method 3	55	1	2	62	1	2	60	2	3
587	Method 4	122	8	7	119	10	8	116	8	7
588	Method 5	94	10	11	92	10	11	75	5	7
589	Method 6	67	10	15	65	10	16	60	8	13

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592 Table 5.

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594		1 (mg kg ⁻¹)			25 (mg kg ⁻¹)		
595		Rec(%)	RSDr(%)	RSDwR(%)	Rec(%)	RSDr(%)	RSDwR(%)
596	Endosulfan ether	83	3	3	86	3	8
597	Endosulfan alcohol	78	11	16	106	8	14
598	Endosulfan α	80	3	11	89	5	10
599	Endosulfan β	80	4	14	96	7	8
600	Endosulfan sulfate	90	3	12	81	1	17
601	Chlorpyrifos	72	5	13	109	7	9

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620 Table 6.

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622 1 (mg kg⁻¹) 25 (mg kg⁻¹)

623 (%) (%)

624 Endosulfan ether -49 11

625 Endosulfan alcohol 36 52

626 Endosulfan α -39 21627 Endosulfan β -34 -1.6

628 Endosulfan sulfate 27 49

629 Chlorpyrifos -7 25

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633 Table 7.

634

635		1(mg kg ⁻¹)			25(mg kg ⁻¹)			50/60(mg kg ⁻¹)		
		Rec	RSDr	RSDwR	Rec	RSDr	RSDwR	Rec	RSDr	RSDwR
636		(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)	(%)
637										
638	Endosulfan ether	76	5	11	81	15	13	83	17	18
639	Endosulfan alcohol	71	8	7	90	13	17	110	15	15
640	Endosulfan α	80	6	10	84	13	12	94	15	16
641	Endosulfan β	77	5	17	87	14	12	101	18	16
642	Endosulfan sulfate	90	9	11	97	13	10	106	15	16
643	Chlorpyrifos	102	8	7	110	15	16	105	12	10

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660 Table 8.

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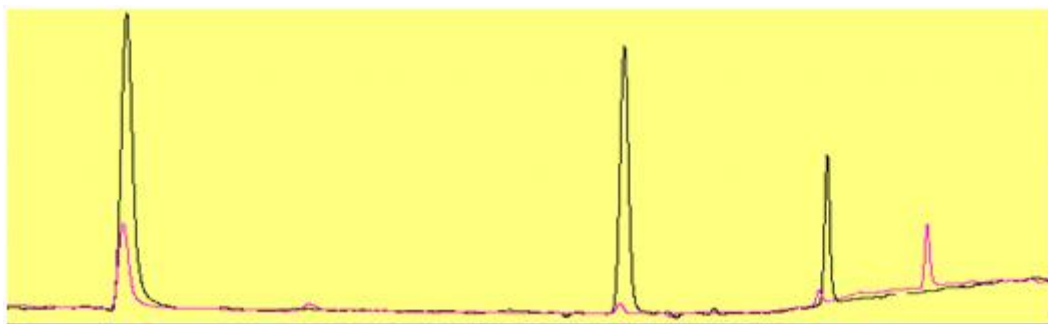
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672 Figure 1.

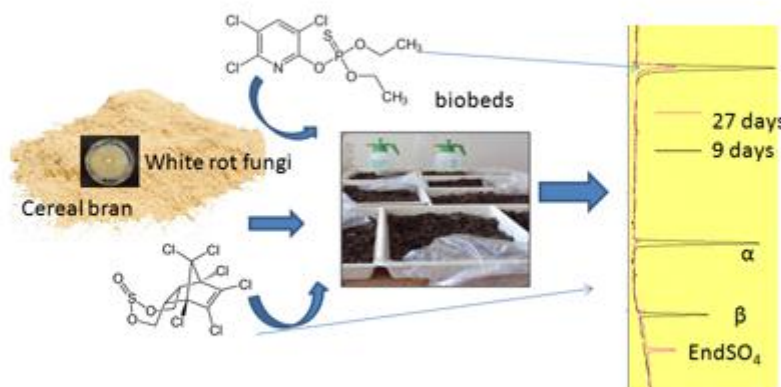


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674 Highlights

- 675
- 676 • Xenobiotic conversion in biobeds can only be proved with validated analytical methods.
 - 677 • Cereal bran is suitable for the growth of native bioconverter Basidiomycetes.
 - 678 • Fit for purpose methodologies to assess biobeds performance were developed
 - 679 • Laboratory biobeds dissipated 79 % chlorpyrifos and 80 % endosulfan after 27 days

680 Graphical abstract



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