

1 **Effect of process parameters in Laccase Mediator System delignification of flax pulp. Part II. Impact**  
2 **on effluents properties**

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8 **Abstract**

9 Flax pulp was bleached by using an enzyme treatment with laccase (L stage) in the presence of HBT as  
10 mediator in order to replace chlorine-based pulp bleaching processes which produce toxic organochlorine  
11 compounds with biotechnological products, achieving an environmentally friendly process technology.  
12 Enzyme treatments were sequentially conducted according to a factorial design involving four variables:  
13 laccase dose, HBT dose, treatment time and oxygen pressure in the reactor. Their influence on the effluent  
14 properties (COD, color, toxicity, spectra and residual enzymatic activity) was examined to evaluate the  
15 impact of L stage on environment since no experimental data have been published concerning this matter.

16 Mathematical models accurately predicting effluent properties in terms of the previous four variables were  
17 developed. High COD levels were obtained as a result of using commercial laccase. Also, red color  
18 produced, especially at long treatment time, relates to formation of oxidation products from HBT. The  
19 residual enzymatic activity depends basically on mediator dose, and mainly activity loss is produced during  
20 the first 30 min of treatment. The toxicity of the effluents was below the limits set by the sewage regulations  
21 for Catalonia and can be ascribed to the combined effect of the laccase-mediator system.

22 **Keywords**

23 Bleaching, COD, color, Effluents, fax pulp, 1-hydroxybenzotriazole, laccase, mathematical model, oxygen  
24 pressure, residual enzymatic activity, sequential statistical plan, toxicity.



## 25 1. Introduction

26 Biotechnology possesses a high potential for cutting costs, improving end products and lessening the  
27 environmental impact of the pulp and paper industry, which has traditionally been regarded as a strong  
28 source of contamination [1]. Production of TCF pulp is growing steadily in response to environmental  
29 regulations seeking both to reduce organochlorine emissions and meeting the need to close bleaching plants  
30 via filtrate recycling circuits. Complete recycling is difficult with sequences using chlorine or its derivatives  
31 owing to their corrosiveness; this has promoted the use of TCF sequences, which allow most of the filtrate to  
32 be reused in the recovery cycle [2,3,4,5,6]. Although the effluents from TCF bleaching sequences contain no  
33 organochlorines, they must be checked for COD and color in order to facilitate circuit closure in future  
34 industrial biobleaching applications.

35 Laccase-mediator systems have facilitated the development of TCF sequences, the replacement of oxygen-  
36 and ozone-based delignification stages, and the efficient bleaching of pulp in addition to reducing kappa  
37 numbers and saving reagents. Based on the literature published recently, the use of a laccase in the  
38 presence of a mediator is a very promising choice for biobleaching pulp [7,8,9].

39 For a mediator to be industrially useful, it should be affordable and environmentally benign. The most  
40 efficient mediators for delignifying pulp are substances containing an  $-N(OH)-$  group (e.g. 1-  
41 hydroxybenzotriazole, HBT) [10,11]. However, HBT is expensive and potentially toxic by itself [12] or through  
42 its reactions products [13,14]. Effluents from laccase-mediator treatments have not seemingly been studied  
43 before; also, product safety sheets typically contain no information about their potential toxicity as assessed  
44 with the Microtox method. In addition, it is known that laccase can be inactivated by oxidized species of  
45 some mediators [15,16,17].

46 In previous studies, laccase-mediator systems were found to be efficient in the bleaching of flax pulp  
47 [18,19,20,21,22]. In a first part of the present study [23] the operating conditions for the laccase-HBT system  
48 were optimized and their influence on the pulp properties was examined, with the main objective of  
49 minimizing the reagent doses and the reaction time to make more suitable the industrial application. In this  
50 paper corresponding to a second part of the study, we examined the influence of process variables in a  
51 laccase-mediator treatment on the properties of the resulting effluents (COD, color, toxicity, residual

52 enzymatic activity and UV/vis absorbance spectra), being the first results obtained in this matter since no  
53 data have been reported before.

## 54 **2. Materials and methods**

### 55 *2.1 Raw material*

56 The raw material used was unbleached flax pulp (*Linum usitatissimum*) subjected to acid washing. Its  
57 properties are ISO brightness of 36.5 %ISO, kappa number of 10.5 and viscosity of 952 mL·g<sup>-1</sup>. The  
58 commercial enzyme was laccase from *Trametes villosa* supplied by Novozymes® (Ref. NS-51002) with an  
59 activity of 39.4 U·mL<sup>-1</sup>, and the mediator hydroxybenzotriazole (HBT) from Fluka (Ref. 54802).

### 60 *2.2 Experimental design*

61 Enzyme treatments were sequentially conducted according to a 2<sup>4</sup> factorial design with three replications at  
62 the centre point and an 8-point star, being the four variables: laccase dose ( $X_1$  from 1 to 20 U·g<sup>-1</sup>), HBT dose  
63 ( $X_2$  from 0.1 to 2 %odp), treatment time ( $X_3$  from 0.5 to 6.5 h) and oxygen pressure in the reactor ( $X_4$  from  
64 0.2 to 0.6 MPa). The statistical analysis of pulp properties was based on the results of a planned sequence  
65 of tests. The experimental design used to this end was conducted stage wise in each sequence. Thus, the  
66 results obtained in each stage were used to decide whether the next was to be performed. More information  
67 concerning the experimental design is reported in the first part of this study [23].

### 68 *2.3 Laccase mediator treatment (L stage)*

69 Enzymatic treatments were carried out in a pressurized reactor at 30 °C and 3 %odp consistency using a 50  
70 mM sodium tartrate buffering solution at pH 4. Control test was performed in the absence of enzyme and  
71 mediator, using a treatment time of 6.5 h and oxygen pressure of 95 0.6 MPa, which were the respective  
72 maximum values employed in the statistical plan [23].

### 73 *2.4 Effluent properties*

#### 74 *2.4.1 COD and color properties*

75 The COD and color values of effluents from L stage were calculated in accordance with ASTM D1252-00  
76 and ASTM D1029-00 standards, respectively. Absorbance data were taken at 600 nm for COD and 465 nm  
77 for color.

#### 78 *2.4.1 Toxicity values*

79 Effluent toxicity was determined with the Microtox method, which measured the light reduction caused by the  
80 microbe *Vibrio fischeri* in contact with toxins (UNE-EN ISO 11348-3). One equitox·m<sup>3</sup> is defined as the  
81 reciprocal of the wastewater dilution (expressed in parts per one) resulting in 50 % inhibition within 15 min  
82 under typical biotesting conditions. Toxicity measurements were color-corrected as per the recommendations  
83 of Azur Environmental, the manufacturer of the MicrotoxOmni equipment used.

#### 84 *2.4.3 Spectrophotometric curves*

85 The effluents from L stage were diluted to 1:20 and their absorbance measured between 200 nm and 400  
86 nm in a UV-vis Shimadzu 1603 spectrophotometer.

#### 87 *2.4.4 Residual enzyme activity*

88 The enzymatic activity of laccase was defined as the amount of enzyme needed to convert 1 μmol of the  
89 substrate ABTS per minute. Oxidation of ABTS was followed by an absorbance increase at 436 nm  
90 ( $\epsilon_{436}=29300 \text{ M}^{-1}\cdot\text{cm}^{-1}$ ) in a UV-vis Shimadzu 1603 spectrophotometer. The reaction mixture contained 5 mM  
91 ABTS, 100 mM sodium acetate buffer at pH 5 and between 10 and 50 μL of enzyme.

### 92 **3. Results and discussion**

#### 93 *3.1 Statistical analysis*

94 The four variables were normalized to three different values (-1, 0 and 1) for implementation of the factorial  
95 design. Table 1 shows the relationship between the process variables and their normalized values in the L  
96 stage. Table 2 shows the COD, color and residual enzymatic activity values in L stage effluents. The  
97 saturated model of the factorial design is showed in Eq. 1.

$$\begin{aligned}
Y_L = & b_0 + b_1X_1 + b_2X_2 + b_3X_3 + b_4X_4 + b_{12}X_1X_2 + b_{13}X_1X_3 + b_{14}X_1X_4 + b_{23}X_2X_3 + b_{24}X_2X_4 \\
& + b_{34}X_3X_4 + b_{123}X_1X_2X_3 + b_{234}X_2X_3X_4 + b_{124}X_1X_2X_4 + b_{134}X_1X_3X_4 + b_{1234}X_1X_2X_3X_4
\end{aligned}
\tag{1}$$

98 3.1.1 Statistical analysis of the COD in L stage

99 *Step 1. Study of factorial design.* The  $2^4$  design was established from tests L 1 to L 16 (see Table 2) and the  
100 probability graph for the saturated model obtained (Eq. 1 and Fig. 1 a). Those coefficients falling on the right  
101 of the significance line in the graph and departing from it were deemed significant. This allowed non-  
102 significant coefficients for the saturated model to be discarded. The new model constructed was used to  
103 determine the new coefficients for the kappa number and their significance. Only term  $X_1$  in the model (Eq. 2)  
104 was significant ( $p < 0.05$ ).

$$Y_{\text{COD-L}} = 317 + 157X_1 \tag{2}$$

105 *Step 2. Study of the variance.* In order to detect potential curvature in a model, the variance might be  
106 checked to verify if remains constant throughout the experimental field studied. By comparing the residual  
107 mean square (RMS = 347) and variance for the three centre points ( $S_c^2 = 1010$ ), the model for the kappa  
108 number was found to be homoscedastic ( $p = 0.22 > 0.05$ ), i.e. the variance was constant.

109 *Step 3. Is it influenced by quadratic terms?* In order to facilitate detection of potential curvature, the model  
110 provided by the previous  $2^4$  factorial design was expanded with the 2 coefficients previously deemed  
111 significant and that representing quadratic terms. The design matrix contained the central tests L 17 to L 19  
112 (Table 2) and a  $2^4$  design with 3 central responses. Based on the statistical analysis for curvature, the  
113 coefficient for the quadratic terms was not significant ( $p = 0.4$ ).

114 *Step 5. Verification of the final model.* COD was measured in tests L 20 to L 23 (Table 2); although these  
115 tests were unnecessary to construct the model, they were used in its analysis. The final model was  
116 constructed from 23 responses and fitted the equation Eq. 3. The coefficient of determination ( $R^2$ ) was 0.98  
117 and the probability associated to  $F_{\text{calc}}$  10.4. Table 3 shows the coefficients of the model and their  
118 significance.

$$Y_{\text{COD-L}} = b_0 + b_1 X_1 \quad (3)$$

119 Residuals distributed according to no well-defined pattern as a function of the estimated responses. In  
120 addition, the COD responses predicted by the model, and the experimental responses, distributed around a  
121 straight line, the differences between the two ranging from - 42 to 27 kg.t<sup>-1</sup> (data not shown). Also, the points  
122 in the normal probability graph Z:N(0,1) vs. residuals fitted a straight line (Fig. 1 b). Based on these results,  
123 the model was deemed statistically accurate.

### 124 3.1.2 Statistical analysis of the color in L stage

125 *Step 1. Study of factorial design.* The 2<sup>4</sup> design was established from tests L 1 to L 16 (see Table 2) and the  
126 probability graph for the saturated model obtained (Eq. 1 and Fig. 2 a). The graphical analysis of the  
127 saturated model allowed non-significant coefficients to be discarded. Those coefficients falling on the right of  
128 the significance line and departing from it were deemed significant and were used to test the model and  
129 determine its significance. All terms in the model thus obtained (Eq. 4) were significant (p < 0.05).

$$Y_{\text{color-L}} = 15.8 + 10.8X_1 + 14.7X_2 + 6.9X_3 + 10.7X_1X_2 + 5.4X_1X_3 + 6.8X_2X_3 + 5.3X_1X_2X_3 \quad (4)$$

130 *Step 2. Study of the variance.* Detecting curvature in a model requires checking whether the variance  
131 remains constant with time throughout the experimental range studied. A comparison of the RMS (3.19) and  
132 variance at the three centre points (1.00) provided p > 0.05; therefore, the model was homoscedastic (i.e. the  
133 variance was constant).

134 *Step 3. Is it influenced by quadratic terms?* Potential curvature in the model obtained by using the previous  
135 2<sup>4</sup> factorial design was examined by expanding it with the 8 coefficients previously deemed significant and  
136 that representing quadratic terms. The design matrix encompassed the central tests (L 17 to L 19) (Table 2)  
137 and a 2<sup>4</sup> design with 3 central responses, and was used to search for potential curvature in the model. Based  
138 on the results of the statistical analysis, the representative quadratic term was significant (p = 0.00) and had  
139 a coefficient of -3.2. Because the central tests did not allow the specific factor influencing the response in a  
140 quadratic manner to be identified, the statistical study had to be repeated with additional tests.

141 Step 4. *Which factors have a quadratic effect?* In order to identify the factors with a quadratic effect on the  
142 response, the study was expanded with the star points (tests L 20 and L 22 to L 26) in order to deconvolute  
143 quadratic terms (Table 2). The model obtained by using the  $2^4$  factorial design with three central tests and  
144 the eight star points was examined.

145 *Step 5. Verification of the final model.* The final model was obtained from 25 responses and fitted the  
146 equation Eq. 5; the coefficient of determination ( $R^2$ ) was 0.99 and the probability associated to  $F_{\text{calc}}$  277.  
147 Table 4 shows the coefficients of the model and their significance.

$$Y_{\text{color-L}} = b_0 + b_1X_1 + b_2X_2 + b_3X_3 + b_{12}X_1X_2 + b_{13}X_1X_3 + b_{23}X_2X_3 + b_{123}X_1X_2X_3 + b_{11}X_1X_1 \quad (5)$$

148 The residuals distributed according to no well-defined pattern with respect to the estimated responses. In  
149 addition, the color responses predicted by the model and their experimental counterparts clustered around a  
150 straight line (data not shown), and differences between the two ranged from 3 to 4  $\text{kgPtCo}\cdot\text{t}^{-1}$ . Also, the  
151 normal probability graph, in the form of  $Z:N(0,1)$  vs. residuals, fitted a straight line (Fig. 2 b). Based on these  
152 results, the model was deemed statistically accurate.

### 153 3.1.3 Statistical analysis of enzymatic residual activity

154 *Step 1. Study of factorial design.* The  $2^4$  design was established from tests L 1 to L 16 (see Table 2) and the  
155 probability graph for the saturated model obtained (Eq. 1, and Fig. 3 a). The graphical analysis of the  
156 saturated model allowed non-significant coefficients to be discarded. Those coefficients falling on the right of  
157 the significance line and departing from it were deemed significant and were used to test the model and  
158 determine its significance. All terms in the model thus obtained (Eq. 6) were significant ( $p < 0.05$ ).

$$Y_{\text{residual activity-L}} = 4.1 + 4.1X_1 - 2.1X_2 - 0.3X_3 - 2.1 X_1X_2 - 0.3X_1X_3 - 0.5X_2X_3 - 0.5X_1X_2X_3 \quad (6)$$

159 *Step 2. Study of the variance.* Detecting curvature in a model requires checking whether the variance  
160 remains constant with time throughout the experimental range studied. A comparison of the RMS (0.20) and  
161 the variance (0.03) at the three centre points provided  $p = 0.27 > 0.05$ ; therefore, the model was  
162 homoscedastic (i.e. the variance was constant).



163 *Step 3. Is it influenced by quadratic terms?* Potential curvature in the model obtained by using the previous  
164  $2^4$  factorial design was examined by expanding it with the 8 coefficients previously deemed significant and  
165 that representing quadratic terms. The design matrix encompassed the central tests (L 17 to L 19) (Table 2)  
166 and a  $2^4$  design with 3 central responses, and was used to search for potential curvature in the model. Based  
167 on the results of the statistical analysis, the representative quadratic term was significant ( $p = 0.00$ ) and had  
168 a coefficient of 1.8. Because the central tests did not allow the specific factor influencing the response in a  
169 quadratic manner to be identified, the statistical study had to be repeated with additional tests.

170 *Step 4. Which factors have a quadratic effect?* In order to identify the factors with a quadratic effect on the  
171 response, the study was expanded with the star points (tests L 20, L 22 to L 26) in order to deconvolute  
172 quadratic terms (Table 2). The model obtained by using the  $2^4$  factorial design with three central tests and  
173 the eight star points was examined. The final model was obtained from 25 responses and fitted the equation  
174 Eq. 7; the coefficient of determination ( $R^2$ ) was 0.98 and the probability associated to  $F_{\text{calc}}$  140. Table 5  
175 shows the coefficients of the model and their significance. The quadratic coefficient of the model was 1.7,  
176 similar to the previously calculated representative quadratic term.

$$Y_{\text{residual activity-L}} = b_0 + b_1X_1 + b_2X_2 + b_3X_3 + b_{12}X_1X_2 + b_{13}X_1X_3 + b_{23}X_2X_3 + b_{123}X_1X_2X_3 + b_{22}X_2X_2 \quad (7)$$

177 *Step 5. Verification of the final model*

178 The residuals distributed according to no well-defined pattern with respect to the estimated responses. The  
179 enzymatic residual activity responses predicted by the model and their experimental counterparts clustered  
180 around a straight line (data not shown), and differences between the two ranged from -1.0 to 1.4 U·g<sup>-1</sup>. Also,  
181 the normal probability graph, in the form of Z:N(0,1) vs. residuals, fitted a straight line (Fig. 3 b). Based on  
182 these results, the model was deemed statistically accurate.

### 183 3.2 Effluent property model

184 Firstly, some control tests were carried out (Table 6) in order to establish the individual contribution of each  
185 reagent and operational conditions to the effluent properties. The control test was performed in the absence

186 of enzyme and mediator, using a treatment time of 6.5 h and oxygen pressure of 0.6 MPa, to evaluate the  
187 effect of buffer, temperature, oxygen pressure and pulp washing. The results show that no effect was found  
188 in color and toxicity, but 155 kg·t<sup>-1</sup> of COD was obtained due to sodium tartrate buffer. Two additional  
189 controls (control-HBT and control-laccase) were carried out, applying 2 % of HBT or 20 U·g<sup>-1</sup> of laccase,  
190 respectively. The most interesting result was the high COD value (352 kg·t<sup>-1</sup>) obtained with the control-  
191 laccase, suggesting the commercial laccase will be the main responsible for the high COD levels in L-  
192 treatments. HBT and laccase have no influence on color and toxicity properties.

### 193 3.2.1 Model for the COD in L stage

194 The model relating the COD of the effluent after the L stage to the process variables fitted the equation Eq.  
195 8 and predicted COD from 160 to 477 kg·t<sup>-1</sup>. Based on the model, the sole variable affecting COD was X<sub>1</sub>  
196 (laccase dose), which influenced it in a linear manner. The other three variables [viz. mediator dose (X<sub>2</sub>),  
197 treatment time (X<sub>3</sub>) and oxygen pressure inside the reactor (X<sub>4</sub>) had no effect on the COD response.

$$Y_{\text{COD-L}} = 318 + 158X_1 \quad (8)$$

198 where  $X_1 = (L - 10.5) / 9.5$ , L = laccase dose (U·g<sup>-1</sup>)

199 COD increased with increasing dose of laccase (Fig. 4 a). The combined COD for the reagents when used at  
200 their highest doses and the control (Table 6) was 521 kg·t<sup>-1</sup>. This value is similar to that obtained in tests  
201 involving the highest laccase dose. Therefore, the high COD levels of the effluents were a result of that  
202 contributed by the commercial enzyme preparation (specifically, by the additives used to maintain its  
203 activity), in fact commercial laccase represents 74 % of total COD in L stage. It would be interesting to  
204 recommend the use of new enzyme formulations of lower COD in order to facilitate industrial application of  
205 the laccase-mediator system.

### 206 3.2.2 Color model for the L stage

207 The color responses of the effluents after the L stage ranged from 1 to 77 kg·t<sup>-1</sup>. The model relating effluent  
208 color after L and the process variables conformed to Eq. 9 and predicted color values of 4 to 81 kg·t<sup>-1</sup>. Based  
209 on the model, the variables influencing color were the laccase dose (X<sub>1</sub>), mediator dose (X<sub>2</sub>) and treatment

210 time ( $X_3$ ). On the other hand, the oxygen pressure,  $X_4$ , had no effect on color. The variable most strongly  
211 influencing effluent color was the HBT dose ( $X_2$ ), with a coefficient of 14.8. In addition, the factor laccase  
212 dose,  $X_1^2$ , had a quadratic influence on the response.

$$Y_{\text{color-L}} = 20.4 + 10.9X_1 + 14.8X_2 + 7.2X_3 + 10.7X_1X_2 + 5.4X_1X_3 + 6.8X_2X_3 + 5.3X_1X_2X_3 - 4.5X_1^2 \quad (9)$$

213 where  $X_1=(L-10.5)/9.5$ . L= laccase dose ( $\text{U}\cdot\text{g}^{-1}$ );  $X_2=(M-1.05)/0.95$ ; M=HBT dose (%odp);  $X_3=(t-3.5)/3$ ;  
214 t=reaction time (h).

215 The lowest color level provided by the plan was  $4 \text{ kg}\cdot\text{t}^{-1}$  and corresponded to the lowest levels of variables,  
216  $X_1 = X_2 = X_3 = -1$  (viz. a laccase dose of  $0.5 \text{ U}\cdot\text{g}^{-1}$ , an HBT dose of  $0.1 \text{ %odp}$  and a treatment time of  $0.5 \text{ h}$ ).  
217 On the other hand, the highest color level provided by the plan was  $81 \text{ kg}\cdot\text{t}^{-1}$  and corresponded to  $X_1 = X_2 =$   
218  $X_3 = 1$  (viz. a laccase dose of  $20 \text{ U}\cdot\text{g}^{-1}$ , an HBT dose of  $2 \text{ %odp}$  and a treatment time of  $6.5 \text{ h}$ ). Therefore, the  
219 highest effluent color was obtained with the highest laccase dose, HBT dose and time studied in the plan.

220 Fig. 4 (b) shows the response surfaces of the color model as a function of the process variables. Effluent  
221 color with a low laccase or mediator doses was very low (less than  $16 \text{ kg}\cdot\text{t}^{-1}$ ), even at long time reaction.  
222 Therefore, the color of the effluents was not due to the initial color of the reagents (Table 6), but rather  
223 developed during the laccase-mediator treatment -which must have caused the dissolution or formation of  
224 colored compounds in the liquid phase.

225 When both laccase and HBT doses have the high level ( $X_1 = X_2 = 1$ ), color values are also higher even at  
226 short time. The more the time reaction is longer, the more higher the color is as the laccase and HBT  
227 increases. The increased color of the effluents obtained after the laccase-mediator treatment can be  
228 ascribed to the formation of colored oxidation products of the mediator and also to an increased content of  
229 degraded lignin in the effluents. Kappa number in L stage does not correlate with effluents color (Fig. 5) so  
230 color is mainly due to oxidation products from the mediator. HBT in a laccase-mediator system is partially  
231 converted into benzotriazole (BT) [8,21,22], which might be responsible for the increased red color of the  
232 effluents at long treatment times.

233 3.2.3 Model for residual enzyme activity after the L stage

234 Table 2 shows the residual activity responses obtained after the L stage and the respective operational  
235 conditions used in the laccase-mediator treatment (L). It was only determined the residual activity in two of  
236 the tests involving the lowest laccase dose ( $0.5 \text{ U}\cdot\text{g}^{-1}$ , L 5 and L 9) and found it to be  $0.6$  and  $0.9 \text{ U}\cdot\text{g}^{-1}$ ,  
237 respectively. Although the activity was seemingly retained or even increased after the treatment, the  
238 differences were very small and the average activity was similar to the detection limit of the assessment  
239 method used. For this reason, the model was constructed under the assumption that the residual activity  
240 after the tests involving the low enzyme dose ( $0.5 \text{ U}\cdot\text{g}^{-1}$ ; L 1, L 3, L 5, L 7, L 9, L 11, L 13 and L 15) would be  
241  $0 \text{ U}\cdot\text{g}^{-1}$ . The activity loss was calculated from Eq. 10. The responses ranged from 0 to  $18 \text{ U}\cdot\text{g}^{-1}$ .

$$\text{activity loss (\%)} = \frac{\text{initial laccase dose} - \text{residual activity}}{\text{initial laccase dose}} \quad (10)$$

242 Analysis of the mathematical model for the residual enzymatic activity

243 The model relating residual activity in the pulp after the L stage to the process variables conformed to Eq. 9  
244 and predicted residual activity values from  $13.1$  to  $-1.9 \text{ U}\cdot\text{g}^{-1}$ . Although some predicted values were negative,  
245 they were taken to be  $0 \text{ U}\cdot\text{g}^{-1}$ . Based on the model, the variables influencing residual activity were the  
246 laccase dose, ( $X_1$ ), mediator dose ( $X_2$ ) and treatment time ( $X_3$ ). On the other hand, the oxygen pressure  
247 inside the reactor ( $X_4$ ), had no influence on the residual activity response. The model factor mediator dose,  
248  $X_1^2$ , exhibited a quadratic influence on the response.

$$Y_{\text{residual activity-L}} = 2.4 + 4.0X_1 - 2.2X_2 - 0.4X_3 - 2.1X_1X_2 - 0.5X_2X_3 - 0.6X_1X_2X_3 + 1.7X_1^2 \quad (11)$$

249 where  $X_1=(L-10.5)/9.5$ . L= laccase dose ( $\text{U}\cdot\text{g}^{-1}$ );  $X_2=(M-1.05)/0.95$ ; M=HBT dose (%odp);  $X_3=(t-3.5)/3$ ;  
250 t=reaction time (h).

251 Fig. 6 (a) shows the residual activity response as a function of the variables of the plan. The responses  
252 corresponding to low laccase doses were excluded as they were close to the detection limit of the method  
253 used to determine residual activity. The highest residual activity was  $13 \text{ U}\cdot\text{g}^{-1}$  and corresponded to  $X_1 = 1$  and

254  $X_2 = -1$  (viz. a laccase dose of  $20 \text{ U}\cdot\text{g}^{-1}$  and an HBT dose of 0.1 %odp). The lowest residual activity provided  
255 by the plan corresponded to  $X_1 = -1$  and  $X_2 = 2$  (viz.  $1 \text{ U}\cdot\text{g}^{-1}$  for laccase and 2 %odp for HBT). Therefore, the  
256 enzyme can lose virtually its whole activity during the laccase-mediator treatment depending on the  
257 particular reagent doses used. Residual activity after the laccase-mediator treatment therefore essentially  
258 depends on the laccase and mediator doses used. By contrast, the variable treatment time has little  
259 influence; residual activity decreases with increasing time, but the effect is only significant with a high  
260 mediator dose.

261 A study of the activity loss as the difference between the initial and residual activity revealed that the  
262 absolute value of the loss at a given mediator dose was greater in the tests involving an enzyme dose of 20  
263  $\text{U}\cdot\text{g}^{-1}$  ( $X_1 = 1$ ) than in those involving a dose of  $10 \text{ U}\cdot\text{g}^{-1}$  ( $X_1 = 0$ ), even though the relative activity was  
264 independent of the initial laccase dose (Fig. 6 b). Using the lowest HBT dose, 0.1 %odp, in combination with  
265 a treatment time of 0.5 h sufficed to reduce the enzyme activity considerably. The residual activity with an  
266 initial laccase dose of 20 and  $10 \text{ U}\cdot\text{g}^{-1}$  was 13 and  $6 \text{ U}\cdot\text{g}^{-1}$ , respectively. At a fixed laccase dose, increasing  
267 the mediator dose decreased the residual activity. At the highest HBT dose studied, 2 %odp, the residual  
268 activity fell below  $3 \text{ U}\cdot\text{g}^{-1}$ . This decrease with increase in mediator dose was substantial even when the dose  
269 was raised from 0.1 to 1 %odp. Higher mediator doses resulted in rather low residual activity, so further  
270 increasing the HBT dose caused no substantial change in residual activity (Fig. 6 a).

271 The residual enzymatic activity was found to depend essentially on the laccase and mediator doses; by  
272 contrast, the treatment time had little effect on it -in fact, most of the enzyme activity was lost within the first  
273 30 min of treatment. The relative residual activity was essentially a function of the mediator dose. Although  
274 low mediator doses sufficed to decrease the residual activity by 40 %, an HBT dose of 2 %odp reduced the  
275 activity by up to 90 %. HBT radicals formed during the treatment inactivate laccase by oxidizing aromatic  
276 amino acids on its protein surface [24]. In the presence of pulp, laccase-mediator systems exhibit reduced  
277 inactivation of the enzyme [16,17, 25] by effect of the free radicals formed reacting with the pulp and  
278 delaying inactivation of laccase as a result. Under these conditions, the loss of enzyme activity as the  
279 mediator dose is increased can be ascribed to an increased formation of oxidation radicals at an early stage  
280 of the process that react with the pulp, but also facilitate inactivation of the enzyme.

### 281 3.4 Toxicity analysis

282 The results of the toxicity analysis of the effluents from the L stage are shown in Table 7. Toxicity responses  
283 ranged from undetectable to 12 equitox·m<sup>-3</sup>. Tests which involved the highest enzyme and mediator doses  
284 (20 U·g<sup>-1</sup> and 2 %odp, respectively), were those providing the highest toxicity values.

285 As per Spain's Water Bill (RDL 1/2001), qualitative and quantitative restrictions on the composition of  
286 sewage are imposed by the competent hydraulic authority; by exception, sewage disposed of at any point in  
287 the public network or through collectors managed by regional or local governments must be authorized by  
288 the competent local or regional body. Decree 130/2003, which passed the regulations on public sewage  
289 services currently in force in Catalonia Spain), set the limit for discharges of inhibitory matter (IM) at 25  
290 equitox·m<sup>-3</sup>. The toxicity values obtained in this study fell below such a limit. In any case, toxicity levels  
291 changed with the particular treatment conditions (see Table 7).

292 The presence of color and/or turbidity can be the source of interferences and overestimated toxicity in some  
293 effluents [26]. Therefore, as recommended in the Microtox instruction manual, our toxicity values were  
294 corrected against absorbance measurements of the samples. Such corrections reduced the initially  
295 estimated toxicity levels for the most strongly colored samples by about 20 %. The effluent from the control  
296 treatment, which involved treating the pulp with sodium tartrate buffer containing no enzyme or mediator,  
297 exhibited no toxicity (Table 7). Nor did the effluent from treatment L 1, which used low reaction doses and a  
298 short time. A HBT solution containing 620 mg·L<sup>-1</sup>, which was the mediator concentration providing a 2 %odp  
299 dose at 3 % consistency in the absence of pulp, resulted in an effluent toxicity value of 3 equitox·m<sup>-3</sup>. An  
300 identical solution additionally containing the amount of laccase used to obtain an enzyme dose of 20 U·g<sup>-1</sup>  
301 provided an identical residual activity; therefore, the enzyme introduces no toxicity in the effluent. The toxicity  
302 levels obtained by using the laccase-mediator system in the presence of pulp (L stage) were higher than  
303 those of the initial toxicity exhibited by the mediator. As can be seen from Table 7, increasing the laccase  
304 and/or mediator dose increased effluent toxicity. Therefore, the toxicity of the effluents from L cannot be  
305 exclusively ascribed to the mediator (HBT) as it can also result from HBT oxidation and degradation by-  
306 products (e.g. benzotriazole), which are more toxic than the mediator itself.

### 307 3.5 UV/vis absorbance spectra for the effluents

308 Fig. 7 shows the spectra for the effluents from the laccase-mediator treatment as obtained following 1:20  
309 dilution. The control effluent, obtained in the absence of enzyme and mediator, exhibited absorbance below  
310 245 nm that was due to the sodium tartrate buffer used (control curve). A HBT solution containing 620 mg·L<sup>-1</sup>,  
311 which was the mediator concentration providing an HBT dose of 2 %odp at 3 % consistency in the  
312 absence of pulp, exhibited an absorbance peak at 305 nm and another at 280 nm (HBT curve). An enzyme  
313 solution containing the concentration needed to obtain a dose of 20 U·g<sup>-1</sup> exhibited no appreciable  
314 absorbance (laccase curve). The spectral signals observed over the wavelength range 200 to 290 nm may  
315 correspond to the mediator, dissolved lignin, and HBT degradation and oxidation products. Because some  
316 signals were overlapped and due to various products, the effluents from the laccase-mediator treatment  
317 precluded determining the amount of dissolved lignin from absorbance measurements as described in Tappi  
318 222 om-02. For this reason, the absorbance signals could only be used for qualitative purposes.

319 As can be seen from Fig. 7, the control signal exhibited the same spectral profile as in the treatments  
320 involving a low mediator dose (0.1 %odp), irrespective of the particular laccase dose and treatment time  
321 used. With a mediator dose of 1 %odp, the spectrum contained three peaks; those at 280 and 305 nm can  
322 be assigned to the mediator –an the HBT solution exhibited both-; on the other hand, the signals between  
323 240 and 300 nm can be assigned to dissolved lignin, degradation products and mediator oxidized forms -in  
324 fact, the signal decreased with decreasing laccase dose and time. The absorbance signals obtained in the  
325 treatments using an HBT dose of 2 %odp were higher than those provided by lower mediator doses. The  
326 peaks at 280 and 305 nm were always lower than those exhibited by the HBT solution. Based on these two  
327 signals, the laccase-mediator treatment reduced the concentration of HBT.

328 Fig. 8 shows the spectra from effluents of L treatments at 2 %odp of HBT and different doses of laccases  
329 and reaction time. The signal at 305 nm was higher with a low laccase dose and a short treatment time. On  
330 equal times, a high laccase dose (20 U·g<sup>-1</sup>) resulted in decreased absorbance values. Therefore, increasing  
331 the laccase dose reduced the concentration of HBT, possibly through more marked oxidation of the mediator  
332 by the enzyme. On the other hand, the signals between 240 and 290 nm were stronger with a high enzyme  
333 dose; this may have been the result of an increased concentration of lignin, and HBT degradation and/or  
334 oxidation products. With a laccase dose of 20 U·g<sup>-1</sup>, the signal at 305 nm peaked at a treatment time of 0.5 h;  
335 therefore, increasing the time to 6.5 h might reduced the concentration of HBT in the system. The signals

336 between 240 and 290 nm were stronger in the long treatment (6.5 h); therefore, prolonging the treatment  
337 increased the concentrations of HBT degradation and/or oxidation products.

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343 starting pulp and Novozymes (Denmark) for supply the laccase.

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404 **FIGURE CAPTIONS**

405 Figure 1. Probability graphs of the model for COD in the L stage. (a) Semi-normal graph of the saturated  
406 model. (b) Residuals vs. Z in the normal law.

407 Figure 2. Probability graphs for the color model in the L stage. (a) Semi-normal graph of the saturated model.  
408 (b) Residuals vs. Z in the normal law.

409 Figure 3. Probability graphs for the residual enzymatic activity in the L stage. (a) Semi-normal graph of the  
410 saturated model. (b) Residuals vs. Z in the normal law.

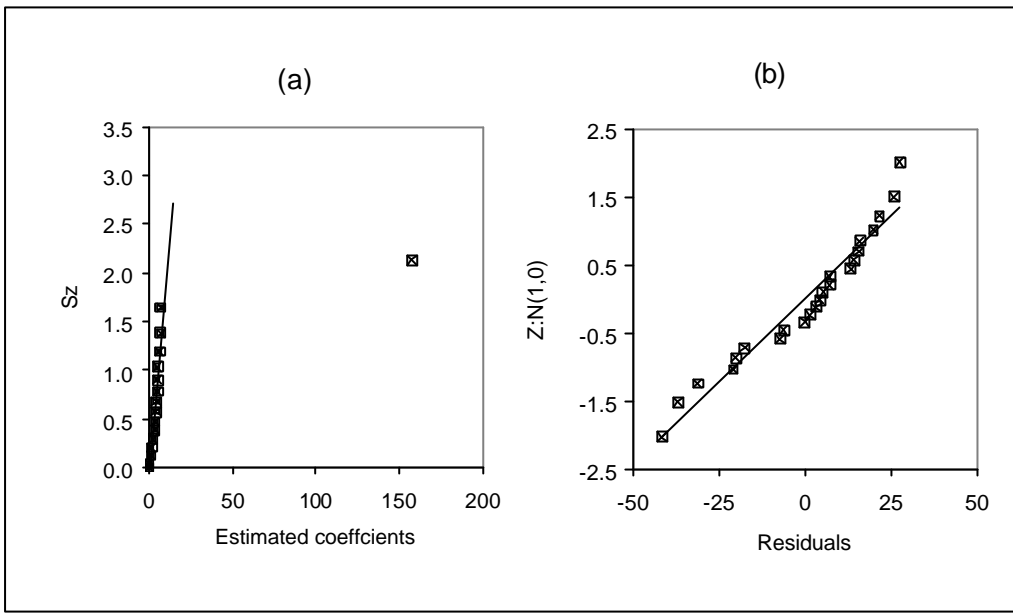
411 Figure 4. Effluent properties, COD (a) and color (b), as a function of the variables of the statistical plan for  
412 the L stage.

413 Figure 5. Relationship between kappa number and effluents color in L stage.

414 Figure 6. Residual activity as a function of the variables of the statistical plan for the L stage.

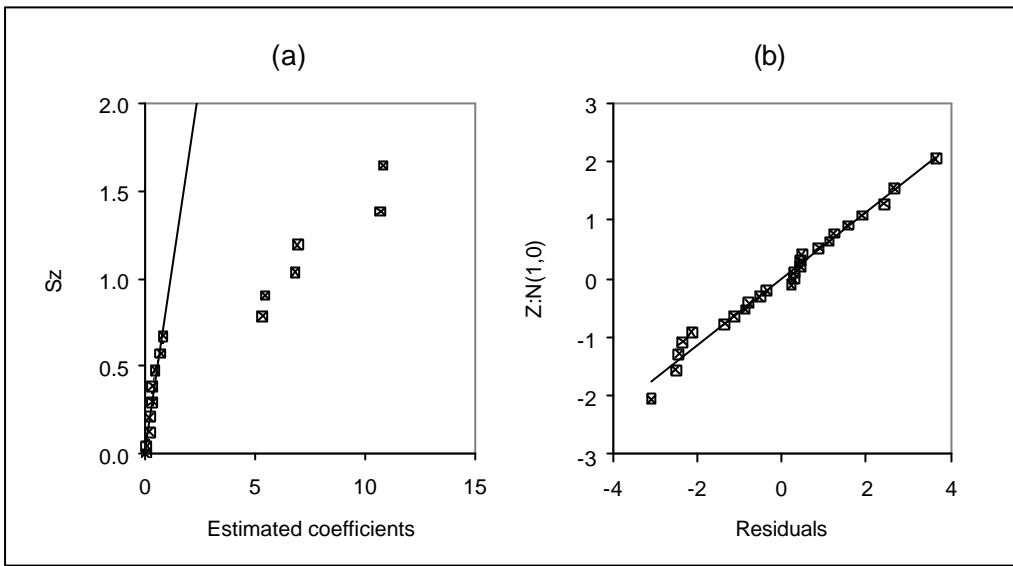
415 Figure 7. Spectra for an HBT solution, a laccase solution and the effluents (following 1:20 dilution) from the L  
416 stage. Control curve: spectra from effluent from control treatment (without enzyme and mediator). HBT  
417 curve: spectra from HBT solution (2 %odp, 3 % consistency). Laccase curve: spectra from laccase solution  
418 ( $20 \text{ U}\cdot\text{g}^{-1}$ ). L curves: spectra from L treatments at 0.1 %, 1 % or 2 % HBT.

419 Figure 8. Spectra for an HBT solution and the effluents (following 1:20 dilution) from the L stage at an HBT  
420 dose of 2 %odp, and different laccase doses (1 or  $20 \text{ U}\cdot\text{g}^{-1}$ ) or reaction time (0.5 or 6.5 h).



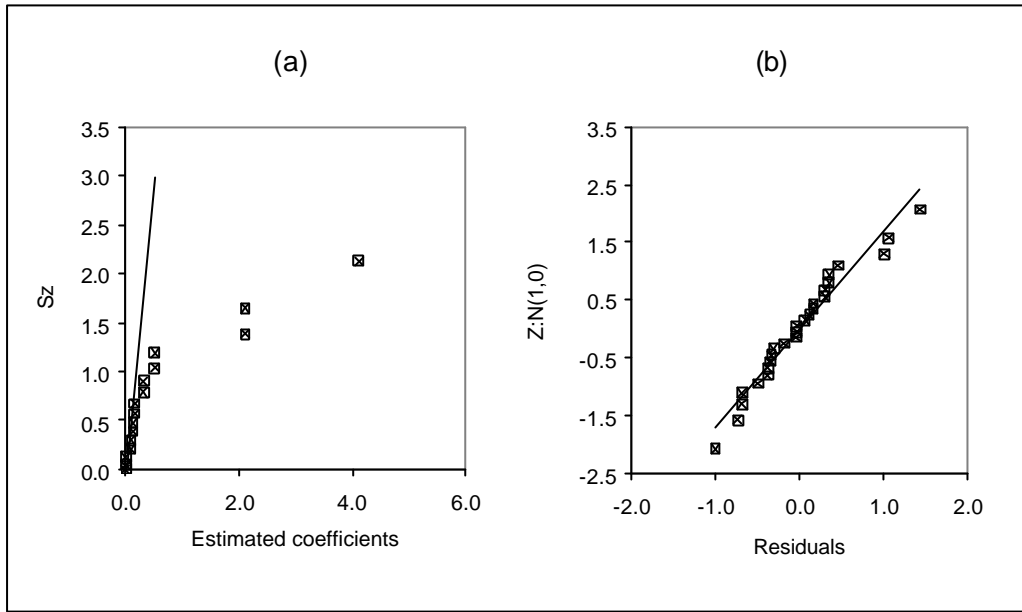
422

423 Figure 1.



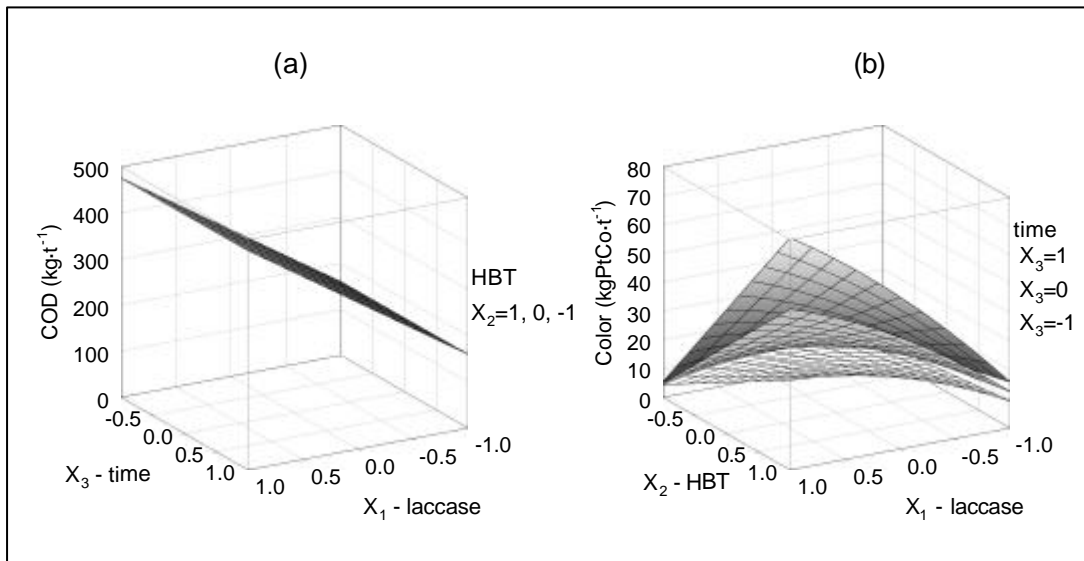
425

426 Figure 2.



428

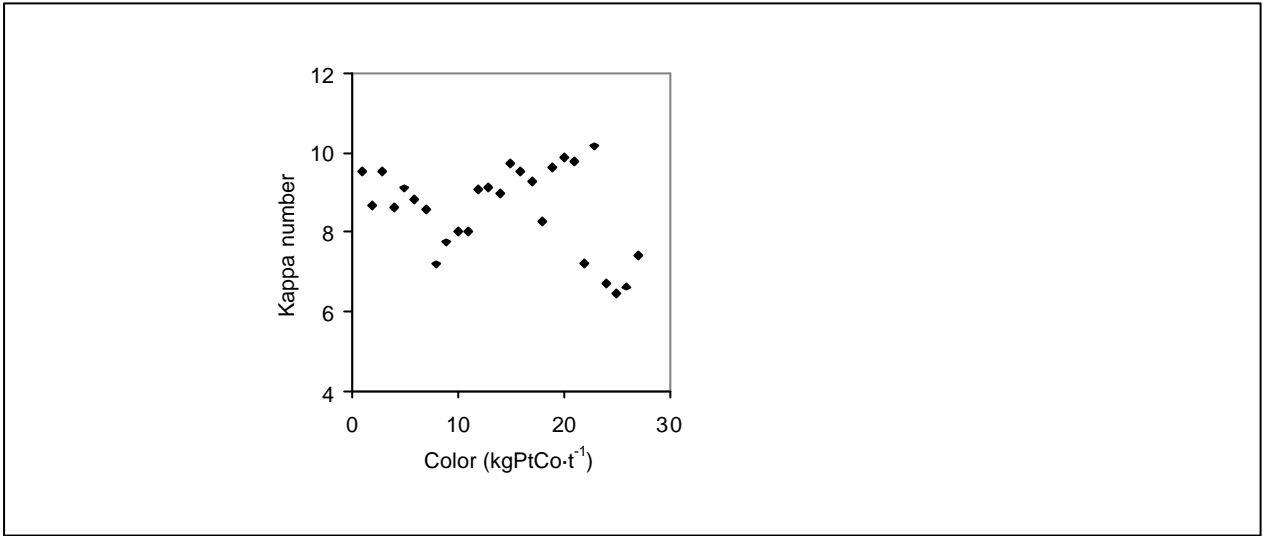
429 Figure 3.



431

432 Figure 4.

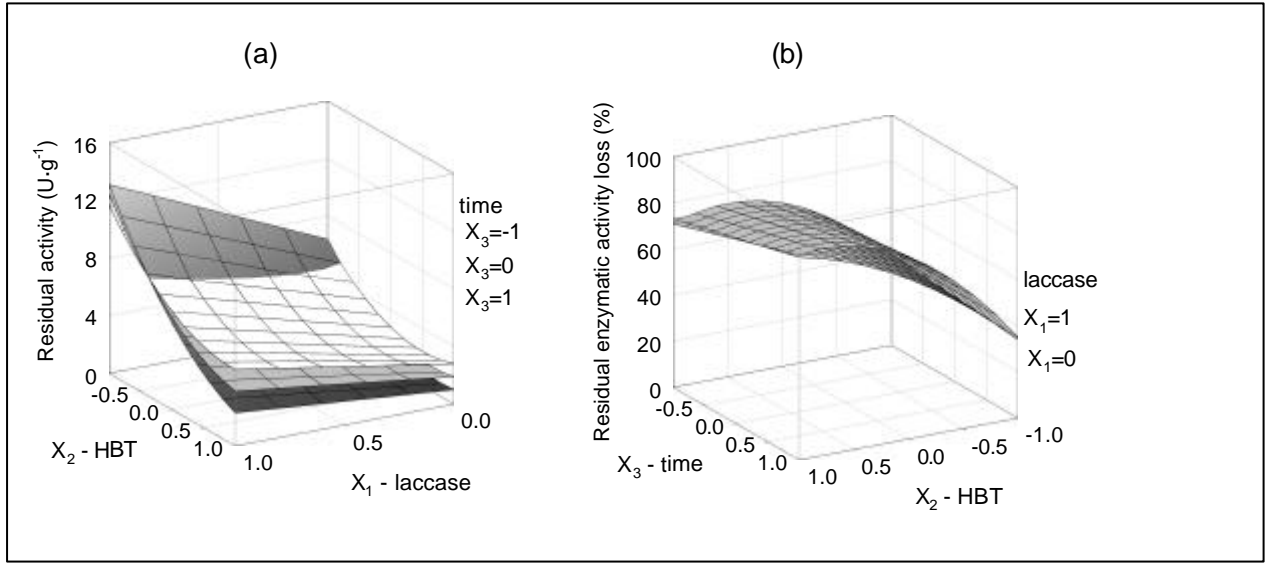
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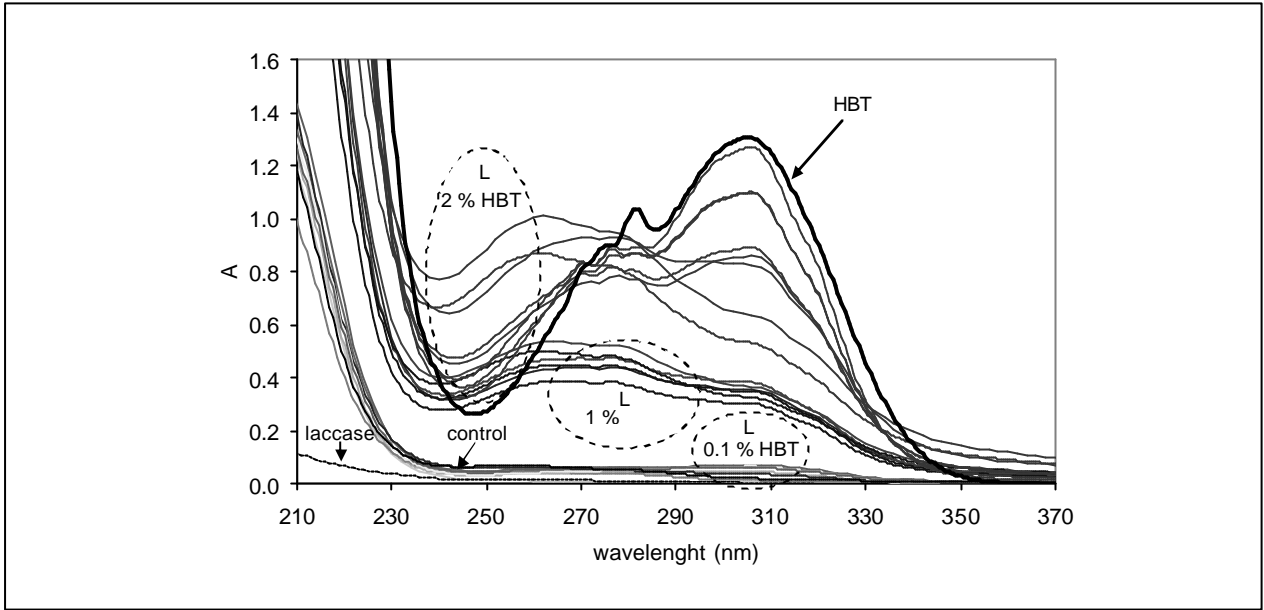
435 Figure 5.





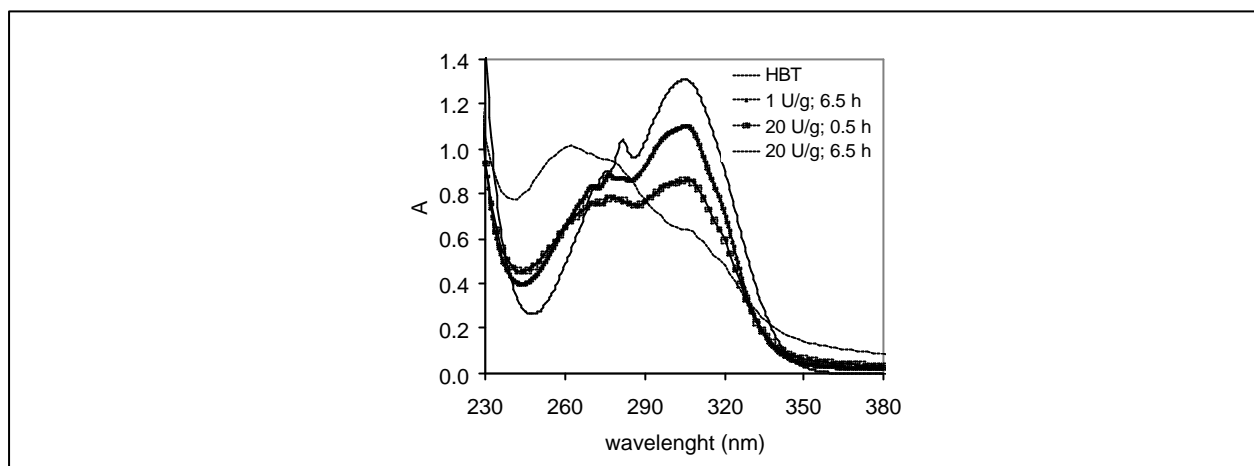
437

438 Figure 6.



441 Figure 7.

442



443

444 Figure 8.

445 **TABLES**

446 Table 1. Normalized values of the process variables in L stage.

447

|   |         | Normalized values |      |     |
|---|---------|-------------------|------|-----|
| Variables                                     | Factors | -1                | 0    | 1   |
| Laccase dose ( $\text{U}\cdot\text{g}^{-1}$ ) | $x_1$   | 1                 | 10.5 | 20  |
| HBT dose (%odp)                               | $x_2$   | 0.1               | 1.05 | 2   |
| Time (h)                                      | $x_3$   | 0.5               | 3.5  | 6.5 |
| Oxygen pressure (MPa)                         | $x_4$   | 0.2               | 0.4  | 0.6 |

448

449 Table 2. COD, color and residual enzymatic activity of effluents after L stage.

| X <sub>1</sub> | X <sub>2</sub> | X <sub>3</sub> | X <sub>4</sub> | Ref. | COD<br>(kg·t <sup>-1</sup> ) | Color<br>(kgPtCo·t <sup>-1</sup> ) | Residual enzymatic<br>activity (U·g <sup>-1</sup> ) |
|----------------|----------------|----------------|----------------|------|------------------------------|------------------------------------|---|
| -1             | -1             | -1             | -1             | L 1  | 180                          | 1                                  | 0   |
| 1              | -1             | -1             | -1             | L 2  | 484                          | 1                                  | 12.3  |
| -1             | 1              | -1             | -1             | L 3  | 154                          | 7                                  | 0   |
| 1              | 1              | -1             | -1             | L 4  | 482                          | 27                                 | 6.4   |
| -1             | -1             | 1              | -1             | L 5  | 129                          | 0                                  | 0   |
| 1              | -1             | 1              | -1             | L 6  | 456                          | 1                                  | 12.4  |
| -1             | 1              | 1              | -1             | L 7  | 153                          | 10                                 | 0   |
| 1              | 1              | 1              | -1             | L 8  | 490                          | 74                                 | 2.5   |
| -1             | -1             | -1             | 1              | L 9  | 140                          | 1                                  | 0   |
| 1              | -1             | -1             | 1              | L 10 | 480                          | 1                                  | 11.8  |
| -1             | 1              | -1             | 1              | L 11 | 176                          | 5                                  | 0   |
| 1              | 1              | -1             | 1              | L 12 | 484                          | 28                                 | 4.9   |
| -1             | -1             | 1              | 1              | L 13 | 186                          | 2                                  | 0   |
| 1              | -1             | 1              | 1              | L 14 | 440                          | 2                                  | 13.2  |
| -1             | 1              | 1              | 1              | L 15 | 160                          | 14                                 | 0   |
| 1              | 1              | 1              | 1              | L 16 | 481                          | 79                                 | 2.2   |
| 0              | 0              | 0              | 0              | L 17 | 340                          | 20                                 | 2.4   |
| 0              | 0              | 0              | 0              | L 18 | 301                          | 19                                 | 2.1   |
| 0              | 0              | 0              | 0              | L 19 | 277                          | 18                                 | 2.4   |
| 1              | 0              | 0              | 0              | L 20 | 491                          | 28                                 | 4.9   |
| 0              | 1              | 0              | 0              | L 21 | 346                          | -                                  | -   |
| 0              | 0              | 1              | 0              | L 22 | 320                          | 30                                 | 1.7   |
| 0              | 0              | 0              | -1             | L 23 | 334                          | 24                                 | 2.6   |
| 0              | -1             | 0              | 0              | L 24 | -                            | 3                                  | 7.4   |
| 0              | 0              | -1             | 0              | L 25 | -                            | 11                                 | 4.3   |
| 0              | 0              | 0              | 1              | L 26 | -                            | 23                                 | 2.6   |

450

451

452 Table 3. Coefficients of the COD model (Eq. 3) in the L stage.

| <b>Coefficients</b> | <b>Estimated coefficients</b> | <b>Standard error</b> | <b>t value</b> | <b>Significance</b>   |
|---------------------|-------------------------------|-----------------------|----------------|-----------------------|
| b <sub>0</sub>      | 319                           | 4.22                  | 75.4           | 4.7·10 <sup>-27</sup> |
| b <sub>1</sub>      | 158                           | 4.91                  | 32.2           | 2.3·10 <sup>-19</sup> |

453

454

455 Table 4. Coefficients of the final color model (Eq. 5) in L stage.

| <b>Coefficients</b> | <b>Estimated coefficients</b> | <b>Standard error</b> | <b>t value</b> | <b>Significance</b> |
|---------------------|-------------------------------|-----------------------|----------------|---------------------|
| b <sub>0</sub>      | 20.4                          | 0.76                  | 26.8           | 0.00                |
| b <sub>1</sub>      | 10.9                          | 0.52                  | 20.9           | 0.00                |
| b <sub>2</sub>      | 14.8                          | 0.52                  | 28.5           | 0.00                |
| b <sub>3</sub>      | 7.2                           | 0.50                  | 14.3           | 0.00                |
| b <sub>12</sub>     | 10.7                          | 0.54                  | 20.0           | 0.00                |
| b <sub>13</sub>     | 5.4                           | 0.54                  | 10.2           | 0.00                |
| b <sub>23</sub>     | 6.8                           | 0.54                  | 12.7           | 0.00                |
| b <sub>123</sub>    | 5.3                           | 0.54                  | 9.9            | 0.00                |
| b <sub>11</sub>     | -4.5                          | 0.92                  | -4.9           | 0.00                |

456

457

458 Table 5. Coefficients of the final model (Eq. 7) for enzymatic residual activity in the L stage.

| <b>Coefficients</b> | <b>Estimated coefficients</b> | <b>Standard error</b> | <b>t value</b> | <b>Significance</b>   |
|---------------------|-------------------------------|-----------------------|----------------|-----------------------|
| b <sub>0</sub>      | 2.4                           | 0.24                  | 10.1           | 7.7·10 <sup>-9</sup>  |
| b <sub>1</sub>      | 4.0                           | 0.17                  | 24.4           | 3.0·10 <sup>-15</sup> |
| b <sub>2</sub>      | -2.2                          | 0.16                  | -13.7          | 5.7·10 <sup>-11</sup> |
| b <sub>3</sub>      | -0.4                          | 0.16                  | -2.7           | 1.6·10E <sup>-2</sup> |
| b <sub>12</sub>     | -2.1                          | 0.17                  | -12.4          | 2.9·10 <sup>-10</sup> |
| b <sub>23</sub>     | -0.5                          | 0.17                  | -3.0           | 8.0·10 <sup>-3</sup>  |
| b <sub>123</sub>    | -0.6                          | 0.17                  | -3.4           | 2.9·10 <sup>-3</sup>  |
| b <sub>22</sub>     | 1.7                           | 0.29                  | 5.9            | 1.4·10 <sup>-5</sup>  |

459



460

461 Table 6. Effluent properties of control treatment after L stage. Individual contribution of laccase and HBT.

| <b>Ref.</b>                               | <b>COD (kg·t<sup>-1</sup>)</b> | <b>Color (kgPtCo·t<sup>-1</sup>)</b> | <b>Toxicity (equitox·m<sup>-3</sup>)</b> |
|---|--------------------------------|--------------------------------------|--|
| Control                                   | 155                            | No significative                     | No detectable                            |
| Control - HBT (2 %odp)                    | 14                             | No significative                     | 3  |
| Control - Laccase (20 U·g <sup>-1</sup> ) | 352                            | No significative                     | No detectable                            |

462

463

464 Table 7. Toxicity of HBT alone, combination of HBT and laccase and the effluents from the L stage.

|                 | x <sub>1</sub> | x <sub>2</sub> | x <sub>3</sub> | x <sub>4</sub> | Ref.          | Toxicity<br>(UT - equitox·m <sup>3</sup> ) |
|-----------------|----------------|----------------|----------------|----------------|---------------|--|
| <b>Reagents</b> | -              | 1              | -              | -              | HBT           | 3  |
|                 | 1              | 1              | -              | -              | HBT + laccase | 3  |
| <b>L stage</b>  | -              | -              | -              | -              | control       | Not detectable                             |
|                 | -1             | -1             | -1             | -1             | L 1           | Not detectable                             |
|                 | -1             | 1              | 1              | -1             | L 7           | 6  |
|                 | 0              | 0              | 0              | 0              | L 18          | 8  |
|                 | 0              | 0              | 1              | 0              | L 22          | 7  |
|                 | 1              | 0              | 0              | 0              | L 20          | 9  |
|                 | 0              | 1              | 0              | 0              | L 21          | 8  |
|                 | 1              | 1              | 1              | -1             | L 8           | 12   |
|                 | 1              | 1              | -1             | -1             | L 4           | 12   |

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