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1 **Exploring substrate promiscuity of chlorophenol hydroxylase against biphenyl**
2 **derivatives**

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14

15 **Abstract**

16 A 2,4-dichlorophenol hydroxylase, whose gene was derived from the metagenomic
17 library of polychlorinated biphenyls (PCBs)-contaminated soil had been found to
18 exhibit broad range of activity for single ring aromatic contaminants including
19 chlorophenols (CPs) and their homologues. In this study, we intended to explore its
20 activity to aromatic bicyclic compounds such as biphenyl and its derivatives which are
21 also important persistent environmental contaminants. Results demonstrated that the
22 enzyme exhibited broad substrate specificity to selected biphenyl derivatives including
23 hydroxylated biphenyls, halogenated biphenyls, PCBs and hydroxylated PCBs, which
24 extended its substrate promiscuity apart from CPs and their homologues. The enzymatic
25 activities against these aromatic bicyclic compounds were congener dependent and the
26 position and type of the substituent on biphenyl derivatives greatly affected the
27 substrate priority of this enzyme. The hypothesis of the catalysis preference of the
28 enzyme on the aromatic ring was preliminarily proposed on the basis of the analyses of
29 the enzymatic activities against biphenyl derivatives. The high activity and removal
30 ability of this enzyme against selected aromatic contaminants would make it a very
31 promising catalyst for biphenyl derivatives bioremediation.

32 **Keywords:** Biphenyl derivatives; Chlorophenol hydroxylase; Hydroxylation; Substrate
33 promiscuity; Bioremediation

34

35 1. Introduction

36 Enzyme promiscuity is the engine of evolutionary innovation and it has attracted
37 significant attention from chemists and biochemists.¹⁻⁴ It is increasingly being perceived
38 as immensely useful phenomenon which can dramatically enhance utility of biocatalysis
39 in biotechnology.² Broad specificity of an enzyme in terms of catalysis of the same
40 reaction with range of substrates is called substrate promiscuity (also known as substrate
41 ambiguity or broad substrate specificity).³ Apart from catalytic promiscuity and
42 condition promiscuity, substrate promiscuity is one of the most important parts of
43 enzyme promiscuity and it might lead to improvements in existing catalysts and results
44 in far larger ranges of organic compounds which can be obtained by biocatalysis.^{3,5-7}

45 Substrate promiscuity have been reported for numerous enzyme classes including
46 cytochrome P450s,^{8,9} kinase,^{10,11} phosphatases,¹² acylaminoacyl peptidase,¹³ DNA
47 methyltransferase,¹⁴ cyclic dipeptide prenyltransferase,¹⁵ glutathione S-transferases,¹⁶
48 laccases¹⁷ and lipases.¹⁸ Among these enzymes, oxidoreductase such as cytochrome
49 P450 superfamily (EC 1.14) and laccases (EC 1.10.3.2) have been increasingly used in
50 the enzymatic-catalyzed degradation of polycyclic aromatic hydrocarbons (PAHs)
51 contaminants due to their high degree of substrate promiscuity.¹⁹ PAHs contaminants
52 such as biphenyl and its derivatives including hydroxylated biphenyls, halogenated
53 biphenyls, PCBs and hydroxylated PCBs (OH-PCBs) are found to be persistent
54 pollutants with high toxicity, bioaccumulation and widespread distribution in the
55 environment.²⁰ Enzymatic degradation of these compounds formally could only be

56 conducted by biphenyl dioxygenases.²¹ Recent literatures reported that successfully
57 biotransformation of these compounds could also be achieved by monooxygenase
58 cytochrome P-450.²²

59 2,4-dichlorophenol (2,4-DCP) hydroxylase (EC 1.14.13.20) is another
60 monooxygenase which have been reported to display high degree of substrate
61 promiscuity against chlorophenols (CPs) and their homologues.²³ This enzyme and
62 multifunction biocatalysis cytochrome P-450, are classified in the same category (EC
63 1.14) in enzyme commission number. It catalyzes the FAD-dependent oxidative
64 hydroxylation of 2,4-DCP and its homologues, in the presence of O₂ and
65 NADPH/NADH as an electron donor, into the corresponding 3,5-dichlorocatechol/CPs,
66 NADP⁺/NAD⁺, and H₂O.²³ Since the hydroxylation activities of this enzyme against
67 chlorophenol congeners were in general much higher than those of the reported
68 cytochrome P-450s and laccases, there has been substantial interest in expanding the
69 substrate scope of 2,4-DCP hydroxylase apart from CPs and their homologues.

70 Our previous research found that 2,4-DCP hydroxylase exhibited broad substrate
71 spectrum against chlorophenols (CPs) and excellent CPs removal ability at both mild
72 and low temperatures, which might make this catalyst more attractive for
73 bioremediation and industrial use.²³ However, the use of this enzyme in the
74 biotransformation was only observed in the biodegradation of single ring aromatic
75 contaminants including above mentioned CPs and their homologues. Limited research
76 has been carried out on its biodegradation of PAHs so far. To explore further the
77 substrate promiscuity of 2,4-DCP hydroxylase, we sought to investigate its ability to

78 degrade biphenyl and its derivatives in this study. Since enzymatic degradation of
79 compounds with higher substituent group was usually reported to be less effective, and
80 many biphenyl derivatives were not commercially available, only lower chlorinated and
81 hydroxyl (each bearing at most two substitutes at different position on the aromatic
82 ring) substitutional bicyclic aromatic compounds were used in this study (structures and
83 names shown in Fig. 1). Cofactors, such as FAD, required for the hydroxylase activities
84 of biphenyl and its derivatives were also investigated because this enzyme exhibits a
85 high sequence and structural similarity to FAD-dependent hydroxylase.²⁴

86 **2. Material and methods**

87 **2.1. Material**

88 Eight biphenyl and its derivatives: Biphenyl, 4-Chlorobiphenol,
89 4,4'-Dichlorobiphenyl, 4-Hydroxybiphenyl, 4,4'-Dihydroxybiphenyl,
90 4-Hydroxy-3-chlorobiphenyl, 4-Hydroxy-2-chlorobiphenyl,
91 4-Hydroxy-4'-chlorobiphenyl and 2,4-DCP of analytical grade were purchased from
92 J&K Scientific Ltd. (Shanghai, China). Other chemicals of analytical grade were
93 obtained from Sigma. Recombinant *Escherichia coli* DH5 α containing the TfdB-JLU
94 gene for 2,4-DCP hydroxylase expression was from our lab. BugBuster protein
95 extraction reagent was from Novagen (Nottingham, UK).

96 **2.2. Methods**

97 **2.2.1. Protein expression and purification**

98 The recombinant *E. coli* was cultivated in LB medium containing 30 μg
99 kanamycin/ml and 34 μg chloramphenicol/ml at 37°C. Protein expression was induced
100 at 18°C by the addition of 0.2 mM isopropyl- β -D-1-thiogalactoside (IPTG) (Fisher
101 Scientific, Fairlawn, NJ) at an OD₆₀₀ of 0.4. After 15 h incubation, the cell pellets were
102 harvested by centrifugation at 12000 rpm and washed with 50 mM sodium phosphate
103 buffer, pH 8.0. For the preparation of crude extract, cells (1.11 g of *E. coli* cell paste
104 from 400 ml fermentation culture) were suspended in 4 ml pH 8.0 Bugbuster Protein
105 Extraction Reagent with 1 mM dithiothreitol (DTT) and 0.6 mM PMSF
106 (phenylmethylsulfonyl fluoride) to yield a higher specific activity compared to
107 ultrasonication. The protein extraction was performed for 10 min at 20°C at 150 rpm.
108 Then, the lysate was centrifuged at 12,000 rpm for 10 min using a Thermo Sorvall WX
109 Ultracentrifuge (Fisher Scientific, Fairlawn, NJ, USA) at 4°C. The supernatant was
110 transferred onto a Hislink™ column (Promega, Madison, WI, USA), rinsed with wash
111 buffer (10 mM HEPES (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid), 10 mM
112 imidazole, pH 8.0), and eluted with elution buffer (10 mM HEPES, 1 M imidazole, pH
113 8.0). The protein supernatant was loaded onto a nickel-nitrilotriacetic agarose resin
114 (Qiagen, Germany) equilibrated with the same buffer. After washing with 5 column
115 volumes of the wash buffer (40 mM imidazole), the bound enzyme was eluted with the
116 elution buffer (200 mM imidazole). The fractions containing 2,4-DCP hydroxylase
117 activity were pooled and concentrated by ultrafiltration by using Amicon Ultra-15
118 centrifugal filter units (Millipore, USA) and then were diafiltered with 50 mM sodium
119 phosphate buffer, pH 7.5, containing 10% (v/v) glycerol. A total of 4 ml protein solution

120 (300 $\mu\text{g ml}^{-1}$) was obtained after 4000 rpm centrifugation. Samples were stored at -80°C
121 for further analysis.

122 **2.2.2. Enzymatic removal of biphenyl and its derivatives**

123 The experiments on conversion of biphenyl and its derivatives were performed in a
124 500 μl eppendorf tube. The reaction mixture was placed into the air-bath constant
125 temperature oscillation incubator (HZQ-F160, Beijing donglian har Instrument
126 manufacture co.,ltd). Unless otherwise indicated, standard reactions were performed by
127 incubating purified enzyme (final concentration 12 $\mu\text{g ml}^{-1}$) with 0.1 mM biphenyl and
128 its derivatives (dissolved in acetone), 0.2 mM NADPH (nicotinamide adenine
129 dinucleotide phosphate) in 50 mM sodium phosphate buffer (pH 7.5) and 5 μM FAD
130 (Flavin adenine dinucleotide) at 25 and 0°C (immersed in ice water) with mild shaking
131 for 1 h. After the reaction, samples were quickly moved to 100°C hot water to
132 deactivate the enzyme. All the removals were performed for three times and statistical
133 significance was determined by one-way analysis of variance (ANOVA) followed by
134 Dunnett's test.

135 **2.2.3. Cofactor requirement for hydroxylase activity and removal of biphenyl** 136 **and its derivatives**

137 2,4-Dichlorophenol hydroxylase is bright yellow and its visible absorption
138 spectrum is typical of a flavoprotein. The prosthetic group is FAD since FAD alone
139 reconstituted active enzyme from apoenzyme. The FAD requirement experiment at 25

140 and 0°C were conducted the same as that of the enzymatic biphenyl and its derivatives
141 removal method described above. Experiments without addition of FAD were used as
142 control.

143 **2.2.4. Enzyme assay and characterization**

144 The enzyme assay for biphenyl and its derivatives during the reaction was
145 determined by monitoring the decrease in absorbance at 340 nm ($\epsilon_{340} = 6,220 \text{ M}^{-1} \text{ cm}^{-1}$)
146 following the substrate-dependent oxidation of NADPH. Unless otherwise indicated,
147 standard enzyme activity assays were performed by incubating the purified enzyme with
148 0.1 mM biphenyl or its derivatives, 5 μM FAD and 0.2 mM NADPH in 50 mM sodium
149 phosphate buffer (pH 7.5) at 25 or 0°C in 500 μl eppendorf tube. The total volume of
150 the reaction mixture is 200 μl . One unit of activity was defined as the amount of enzyme
151 required to consume 1 μmol NADPH per min at 25°C. The kinetic parameters of the
152 purified enzyme for biphenyl and its derivatives at 25°C were obtained using NADPH
153 at 0.2 mM, 5 μM FAD and varying biphenyl or its derivatives from 0.5 to 200 μM . The
154 kinetic constants were calculated from Lineweaver–Burk plots via non-linear regression
155 using GraphPad Prism 5 (GraphPad, San Diego, CA).²⁴ Protein concentrations were
156 determined by the BCA method (Novagen® BCA Protein Assay Kit) using bovine
157 serum albumin as the standard. Biphenyl and its derivatives removal were measured
158 after 1 h reaction using UV spectrometry. The removal of biphenyl and its derivatives
159 was calculated by dividing the concentration of the amount of reduction of NADPH by
160 the amount of the initial NADPH.

161 3. Results and discussion

162 3.1. Substrate specificity of 2,4-DCP hydroxylase against biphenyl and its 163 derivatives

164 Although 2,4-DCP hydroxylase exhibited broad substrate to certain chlorophenol
165 congeners and derivatives, its substrate specificity to biphenyl and its derivatives at low
166 and moderate temperature has yet to be investigated.²³ In our study, 2,4-DCP
167 hydroxylase activities to biphenyl and its derivatives were investigated at 25 and 0°C.
168 The reported possibility of product inhibitory effect was not evaluated in this study. To
169 explore the substrate specificity and catalysis preference of this enzyme, seven biphenyl
170 derivatives used in this study represent chloro- and hydroxyl-substituent at different
171 positions on the aromatic bicyclic molecules: the single substituent (4-Chlorobiphenol
172 and 4-Hydroxybiphenyl), double substituents on the same ring
173 (4-Hydroxy-2-chlorobiphenyl and 4-Hydroxy-3-chlorobiphenyl), and double
174 substituents on the different rings (4,4'-Dichlorobiphenyl, 4,4'-Dihydroxybiphenyl and
175 4-Hydroxy-4'-chlorobiphenyl). 2,4-DCP hydroxylase shows a broad substrate
176 specificity and satisfactory activities to certain biphenyl and its derivative at 25°C (Fig.
177 2). The relative enzymatic activity (expressed as a percentage of the maximum enzyme
178 activity against its regarded natural substrate 2,4-DCP at 25°C without FAD) to
179 4-Hydroxy-2-chlorobiphenyl, 4-Chlorobiphenol, 4-Hydroxybiphenyl and
180 4,4'-Dichlorobiphenyl was 273%, 235%, 131% and 96%, respectively at 25°C with
181 FAD. Superior to laccase which do not accept nonhydroxylated biphenyl substrates, the
182 enzymatic activities for 4-Chlorobiphenol (235%) and 4,4'-Dichlorobiphenyl (96%)

183 were satisfactory. Temperature has been found to have greatly effect on enzymatic
184 activities. Fig. 2 shows that the relative enzymatic activities against the detected
185 substrate at 0°C were in general significantly lower than those at 25°C. The differences
186 between these enzymatic activities were statistically significant ($p < 0.05$).

187 It is very interesting that the substrate specificity to different substrates was quite
188 different. The specificity pattern of the enzyme for biphenyl derivatives was correlated
189 with both the relative positions of the chlorine or hydroxyl substituent on the biphenyl
190 rings and with the number of chlorine or hydroxyl substituent on the rings. Thus, we
191 would like to propose a preliminary assumption on the metabolic pathways for
192 degradation of biphenyl and its derivatives in the enzymatic hydroxylazation step prior
193 to the tedious and precise detection. Data analysis was conducted on the basis of the
194 enzymatic activities against biphenyl derivatives to estimate the position preference of
195 this enzyme. The substrate specificities at 25 and 0°C were similar, as such, the
196 assumption was proposed on the basis of the results at 25°C (Table 1). Enzymatic
197 activities observed for the biphenyl and its derivatives were quite different. Almost no
198 enzymatic activity was observed when biphenyl was used as substrate (Table 1).
199 However, the enzymatic activities were greatly improved when biphenyl derivatives
200 with substituent group were used as substrates, which suggested that suitable
201 substitution on the biphenyl is of significant for stimulating the enzymatic activity. The
202 result also shows that the activities of the enzyme were related to the substitution type
203 and patterns of specific biphenyl derivatives. The enzyme activities against single
204 substitute substrates (235% for 4-Chlorobiphenol and 131% for 4-Hydroxybiphenyl) are

205 in generally higher than those of double substitute substrates with the exception for that
206 of 4-Hydroxy-2-chlorobiphenyl (273%), which exhibited the highest activity in the
207 detected substrates. The higher enzyme activity of 4-Chlorobiphenol (235%) compared
208 to that of 4-Hydroxybiphenyl (131%), as well as 4,4'-Dichlorobiphenyl (96%)
209 compared to those of 4-Hydroxy-4'-chlorobiphenyl (76%) and 4,4'-Dihydroxybiphenyl
210 (45%) suggested that enzyme activities against chloro-substituted substrates are higher
211 than those of hydroxyl-substituted substrates when the substituent groups are on the
212 same positions on the aromatic ring of biphenyl. Significant enzymatic activity
213 differences were observed when the substrates with double substituent on one aromatic
214 ring (273% for 4-Hydroxy-2-chlorobiphenyl and 12% for 4-Hydroxy-3-chlorobiphenyl).
215 This result suggested that the enzyme might have a strict specificity for attacking at
216 position 3 (*ortho*-position to 4 hydroxyl group) on one aromatic ring of biphenyl during
217 the hydroxylation reaction since when this position was occupied, only few activities
218 were left. Similar as that of cytochrome P450-catalyzed aromatic hydroxylation, the
219 result was generally consistent with the rules of electrophilic aromatic substitution (EAS)
220 effect.²⁵ Hydroxylazation of biphenyl and its derivatives is a typical EAS reaction.
221 Hydroxyl and chloro are important substituents for EAS. And these two substituents
222 will have different effects on the electron distributions in the biphenyl ring system.

223 It is well known that both the speed and the regioselectivity of EAS are affected by
224 the substituents already attached to the aromatic ring.²⁵ In terms of speed, some groups
225 promote the reaction rate of hydroxylazation, while other groups decrease it.
226 Substituents can generally be divided into two classes regarding electrophilic

227 substitution: activating and deactivating towards the aromatic ring. Activating
228 substituents or activating groups such as hydroxyl will stabilize the cationic
229 intermediate formed during the substitution by donating electrons into the ring system,
230 by either inductive effect or resonance effects. This well explained the promotion of
231 enzymatic activities against OH-biphenyl compared to that of biphenyl. On the other
232 hand, deactivating substituents such as chloro would destabilize the intermediate cation
233 and thus decrease the reaction rate. They do so by withdrawing electron density from
234 the aromatic ring. The increase of enzymatic activities against chloro-substitute
235 biphenol derivatives compared to that of biphenyl is surprising since chloro is a
236 deactivating group for aromatic ring. The deactivating effect might be offset by other
237 factors. In the enzymatic reaction, regioselectivity of EAS and substrate might also play
238 an important role in influencing the reaction rate. It might be that there is some
239 interaction between chloro-substituent on the biphenyl derivatives and the active site of
240 enzyme, which help the hydroxylase to direct the substrates more efficiency during
241 hydroxylazation. Interaction with enzyme might also change the balance of resonance
242 and polar effects, strengthen the weak rate-enhancing resonance effect, or weaken the
243 strong rate-retarding polar effect. The in general higher activities of single substitute
244 substrates than those of double substitute substrates might be due to that the enzyme has
245 a sterically permissive active site that is not overly restrictive to the motion of single
246 substitute substrates.²⁶

247 This result was coincident with the previous reports in terms of preferred
248 hydroxylation position since most of the other favoprotein hydroxylases that

249 hydroxylating the primary substrate either *ortho* or *para* to the existing hydroxyl groups.
250 The slightly activities of 4-Hydroxy-3-chlorobiphenyl suggested that other positions on
251 the phenol ring might also be hydroxylated. Despite the certainty of product formation,
252 hydroxylation might not be the only pathway for the reaction, further study to identify
253 the product is needed to be done.

254 Moreover, introducing the second substituent on the other aromatic ring of
255 biphenyl derivative resulted in decreasing the enzyme activity compared to those with
256 substituent only on one aromatic ring. For example, the enzymatic activity to
257 4-Hydroxybiphenyl was 131%. However, only 76% (for 4-Hydroxy-4'-chlorobiphenyl)
258 and 45% (for 4,4'-Dihydroxybiphenyl) activity was remained, respectively after
259 introducing the other substituent on the other aromatic ring of 4-Hydroxybiphenyl.
260 Similarly, the enzymatic activity to 4-Chlorobiphenol was 235%. And only 96% activity
261 was remained for 4,4'-Dichlorobiphenyl, after introducing the other chlorine on the
262 other aromatic ring of 4-Chlorobiphenol. As far as the enzymatic activities of OH-PCBs
263 were concerned, the enzymatic activities preference of OH-PCBs is in the order of
264 4-Hydroxy-2-chlorobiphenyl > 4-Hydroxybiphenyl > 4-Hydroxy-4'-chlorobiphenyl >
265 4-Hydroxy-3-chlorobiphenyl. The reactivity order result suggested that the secondary
266 substituent groups might be very important for the substrate orientation when acting on
267 the active site of the enzyme. When the biphenyl ring has two substituent groups, each
268 exerts an influence on subsequent substitution reactions. Both chloro and
269 hydroxyl-substituents are *ortho-para* director for aromatic compounds. The highest
270 enzymatic activity against 4-Hydroxy-2-chlorobiphenyl might be that the two

271 substituents (hydroxyl and chloro) on the one ring of biphenyl have the same directing
272 effect for hydroxylazation reaction, and thus greatly improve the reaction rate. The final
273 result of the electrophilic aromatic substitution seemed hard to predict. The lowest
274 enzymatic activity against 4-Hydroxy-3-chlorobiphenyl might be caused by steric
275 hindrance between substituent and electrophile.

276 To sum up, three hypotheses on the catalysis preference of the enzyme on the
277 aromatic ring was concluded based on the presence results: (1) position 3 in one ring of
278 biphenyl is the preferred position for hydroxylation; (2) the presence of substituent in
279 *para*-position in one ring of biphenyl greatly improve the enzyme activity; (3) the
280 activity of chloride substituent is better than that of hydroxyl group. Our assumption
281 needs to be further confirmed by other experiments. Notably, the premise of our
282 assumption is that the main reaction is hydroxylation reaction. Moreover, hydroxylation
283 only occurs on the free position of the biphenyl structure instead of dechlorination.²⁷
284 Many recent studies have shown the multiple functions of oxygenase.^{28,29} As such, the
285 oxygen consumption³⁰ and products derived from each substrate should be investigated
286 to determine whether other side reactions occur or not.

287 Apparent kinetic parameters (Michaelis-Menten constant, K_m ; catalytic constant,
288 k_{cat} , and catalytic efficiency, k_{cat}/K_m) for the hydroxylation were calculated from
289 Lineweaver-Burk and Eadie-Hofstee plots. Our kinetic results shown in Table 2 fit well
290 with our enzyme specificity result. The higher the substrate activity, the lower the
291 corresponding K_m , suggesting that 2,4-DCP hydroxylase exhibits high affinity against
292 its favourable biphenyl derivatives. The K_m of 2,4-DCP hydroxylase against

293 4-Hydroxy-2-chlorobiphenyl ($4.2 \mu\text{M}$) and 4-Chlorobiphenol ($5 \mu\text{M}$) are even
294 comparable with that of its preferred nature substrate 2,4-DCP ($5 \mu\text{M}$). Also k_{cat} and
295 $k_{\text{cat}}/K_{\text{m}}$ values were in the order of 4-Hydroxy-2-chlorobiphenyl > 4-Chlorobiphenol >
296 4-Hydroxybiphenyl > 4,4'-Dichlorobiphenyl > 4-Hydroxy-4'-chlorobiphenyl >
297 4,4'-Dihydroxybiphenyl > Biphenyl > 4-Hydroxy-3-chlorobiphenyl.

298 **3.2. Enzymatic removal of biphenyl and its derivatives**

299 The removal of biphenyl and its derivatives should be double-checked by using
300 high-performance liquid chromatography to confirm the removal of biphenyl and its
301 derivatives, not just by measuring NADPH consumption detected by UV detection. The
302 product derived from each substrate was not determined in this study. The results
303 indicated that the high enzymatic activities for biphenyl and its derivatives generally
304 resulted in corresponding high biphenyl and its derivatives removal. For example, the
305 high activities of 4-Hydroxy-2-chlorobiphenyl (273%), 4-Chlorobiphenol (235%) and
306 4-Hydroxybiphenyl (131%) resulted in corresponding high substrate removal which was
307 81.92%, 73.08% and 84.69%, respectively at 25°C . Notably, although the enzymatic
308 activities for certain biphenyl and its derivatives were similar (235% relative activity for
309 4-Chlorobiphenol) or even higher (273% relative activity for
310 4-Hydroxy-2-chlorobiphenyl) than that of 2,4-DCP (235% relative activity) at 25°C ,
311 their removal were lower (73.08% removal for 4-Chlorobiphenol and 81.92% removal
312 for 4-Hydroxy-2-chlorobiphenyl) than that of 2,4-DCP (92.38% removal), as shown in
313 Fig. 3. As far as the enzymatic removal of double substitutions biphenyl derivatives

314 were concerned, most of the detected derivatives were more resistant to 2,4-DCP
315 hydroxylase degradation than 4-Hydroxy-2-chlorobiphenyl. The results in Fig. 3 also
316 indicated that the removal of biphenyl and its derivatives were less efficient at 0°C than
317 that at 25°C. Since the removal of biphenyl and its derivatives was not as good as that of
318 2,4-DCP in one hour. We intended to prolong the reaction time of enzymatic removal of
319 biphenyl and its derivatives to see if these contaminants could be further degraded. Fig.
320 4 shows that further increasing the reaction time to 24 h resulted in remarkable
321 improvement of the biphenyl and its derivatives removal at 25°C. However, no obvious
322 increase of biphenyl and its derivatives removal was observed when further increasing
323 the reaction time at 0°C (data not shown). The differences between these removals were
324 statistically significant ($p < 0.03$). Although the enzymatic removal rates of biphenyl
325 and its derivatives was lower than those of CPs which required only one hour to achieve
326 their maximum removal,²³ this enzymatic process is still attractive for industrial use.

327 **3.3. Cofactor requirement for hydroxylase activity and removal of biphenyl** 328 **and its derivatives**

329 Previous studies reported that the reactions catalyzed by specific hydroxylases
330 require FAD as a cofactor to stimulate their substrates.³¹ However, hydroxylases, such
331 as the hydroxylase from *Arthrobacter*, do not exhibit any demonstrable FAD
332 requirement.³¹ TfdBs display high sequence and structural similarity to FAD-dependent
333 hydroxylases and contain FAD as a prosthetic group.³² The FAD requirement for
334 hydroxylase activities and removal of biphenyl and its derivatives was investigated to

335 specify if FAD is the essential cofactor for this enzyme.

336 Our previous results showed that FAD is very important in improving 2,4-DCP
337 hydroxylase activity against 2,4-DCP and the optimum FAD concentration is 5 μM .²³
338 So 5 μM was selected as the final FAD concentration for the following FAD
339 requirement investigation. The result shows that addition of FAD resulted in a general
340 significant increase in the hydroxylase activity in the range of 1.05-fold to 2.63-fold
341 (Fig. 5a), and 1.12-fold to 8.80-fold (Fig. 5b) at 25 and 0°C, respectively, for different
342 biphenyl and its derivatives with the exception of 4-Hydroxy-3-chlorobiphenyl.
343 Moreover, the enzymatic activity incremental effects were substrate dependent. Notably,
344 the FAD requirement for hydroxylase activity at 0°C seemed to be higher than that at
345 25°C because the enzymatic activity improvements were in general higher at 0°C.

346 The result of cofactor requirement for removal of biphenyl and its derivatives are
347 shown in Fig. 6. The addition of FAD also resulted in a general improvement of
348 biphenyl and its derivatives removal in the range of 1.24-fold to 3.74-fold (Fig. 6a), and
349 1.10-fold to 6.21-fold (Fig. 6b) at 25 and 0°C, respectively. Notably, the result in Fig. 6
350 also demonstrated that the removal improvement with the addition of FAD for
351 4-Hydroxy-2-chlorobiphenyl (3.74-fold at 25°C and 3.2-fold at 0°C) was fairly high
352 than those of other biphenyl derivatives, suggesting the high FAD requirement for this
353 substrate. The differences between the enzymatic activities shown in Fig. 6 and the
354 removal shown in Fig. 5 were statistically significant ($p < 0.05$).

355 The bright yellow color and its visible absorption spectrum (Fig. S1) suggested that
356 certain amount of FAD bound to the enzyme after the protein purification. Also FAD

357 concentration in the supernatant after heat-denaturing of protein was determined
358 according to method in the literature.³³ The concentration of the free FAD released was
359 assumed to be equivalent to the concentration of the FAD-bound enzyme. The free FAD
360 released from the enzyme and its concentration was calculated on the basis of the free
361 FAD molar absorption coefficient (ϵ_{450} of $11.3 \text{ mM}^{-1} \text{ cm}^{-1}$).³³ And the FAD
362 concentration measured after the heat-denature experiment was $0.59 \mu\text{M}$. This result
363 well explained the existence of activity and substrate transformation ability of the
364 enzyme. Cofactor requirement results also showed that further addition of FAD in the
365 reaction mixture, led to in general improvement of enzymatic activities as well as
366 substrates transformation ability. In general, one flavin per enzyme active site is
367 required. All flavoprotein aromatic hydroxylases contain one molecule of FAD per
368 subunit and that the 2,4-DCP hydroxylase is a tetrameric protein. So the molar ratio of
369 FAD/protein should be 4:1. It is notable that the enzyme concentration in the reaction
370 mixture is $0.19 \mu\text{M}$, and the molar ratio of FAD/protein ratio without FAD addition was
371 approximately 3:1, which is lower than 4:1. Our result might suggest that
372 supplementing the flavin cofactor FAD in the reaction mixture may be possible to
373 reconstitute the flavoprotein.

374 **4. Conclusions**

375 In the present study, substrate promiscuity of 2,4-DCP hydroxylase against
376 biphenyl derivatives was explored. The enzyme activities of certain biphenyl derivatives
377 are comparable with that of its regarded natural substrate 2,4-DCP. The high removal

378 ability of this enzyme against certain biphenyl derivatives as well as CPs would make it
379 a potentially catalyst in the bioremediation of aromatic contaminants. This enzyme
380 would also be a promising template candidate for PAHs bioremediation-catalyst
381 reconstruction through directed evolution and protein engineering. The preliminary
382 assumption we proposed on the metabolic pathways for degradation of biphenyl and its
383 derivatives in the enzymatic hydroxylazation step would provide a good reference value
384 for screening new potential substrates and enzyme reconstruction. Further enzymatic
385 and reaction mechanism studies may improve our understanding of biphenyl derivatives
386 degradation pathway and help optimize our efforts to remediate biphenyl
387 derivatives-contaminated environment.

388 **Acknowledgment**

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392 **References**

- 393 1 O. Khersonsky and D.S. Tawfik, *Annu. Rev. Biochem.*, 2010, **79**, 471-505.
- 394 2 L. Jiang and H. Yu, *Biotechnol. Lett.*, 2014;**36**, 99-103.
- 395 3 B. Arora, J. Mukherjee and M. N. Gupta, *Sustain. Chem. Process.*, 2014, **2**, 1-9.
- 396 4 C. Pandya, J. D. Farelli, D. Dunaway-Mariano and K. N. Allen, *J. Biol. Chem.*, 2014, **289**,
397 30229-30236.

- 398 5 M. López-Iglesias and V. Gotor-Fernández. *Chem. Rec.*, 2015, DOI: 10.1002/tcr.201500008.
- 399 6 K. Hult and P. Berglund, *Trends Biotechnol.*, 2007, **25**, 231-238.
- 400 7 S. D. Copley, *Trends Biochem. Sci.*, 2015, **40**, 72-78.
- 401 8 K. Auclair and V. Polic, *Adv. Exp. Med. Biol.*, 2015, **851**, 209-228.
- 402 9 V. Polic and K. Auclair. *Bioorg. Med. Chem.*, 2014, **22**, 5547-5554.
- 403 10 S. Lutz, J. Lichter and L. Liu, *J. Am. Chem. Soc.*, 2007, **129**, 8714-8715.
- 404 11 Y. Li, H. Yu, Y. Chen, K. Lau, L. Cai, Cao H, V. K. Tiwari, J. Qu, V. Thon, P. G. Wang and X.
405 Chen, *Molecules*, 2011, **16**, 6396-6407.
- 406 12 H. Huang, C. Pandya, C. Liu, N. F. Al-Obaidi, M. Wang, L. Zheng, S. Toews Keating, M. Aono, J.
407 D. Love, B. Evans, R. D. Seidel, B. S. Hillerich, S. J. Garforth, S. C. Almo, P. S. Mariano, D.
408 Dunaway-Mariano, K. N. Allen, J. D. Farelli, *Proc. Natl. Acad. Sci. U S A.*, 2015, **112**,
409 E1974-1983.
- 410 13 E. A. Brunialti, P. Gatti-Lafranconi and M. Lotti, *Biochimie*, 2011, **93**, 1543-1554.
- 411 14 J. Aranda, M. Roca and I. Tuñón. *Org. Biomol. Chem.*, 2012, **10**, 5395-5400.
- 412 15 J. M. Schuller, G. Zocher, M. Liebhold, X. Xie, M. Stahl, S. M. Li and T. Stehle, *J. Mol. Biol.*,
413 2012, **422**, 87-99.
- 414 16 J. D. Hayes, J. U. Flanagan and I. R. Jowsey, *Annu. Rev. Pharmacol. Toxicol.*, 2005, **45**, 51-88.
- 415 17 R. Chandra and P. Chowdhary, *Environ. Sci. Process. Impacts.*, 2015, **17**, 326-342.
- 416 18 A. S. de Miranda, L. S. Miranda and R. O. de Souza, *Biotechnol. Adv.*, 2015, **33**, 372-393.
- 417 19 A. Sjödin, R. S. Jones, S. P. Caudill, L. Y. Wong, W. E. Turner and A. M. Calafat, *Environ. Sci.*
418 *Technol.*, 2014, **48**, 753-760.
- 419 20 L. Passatore, S. Rossetti, A. A. Juwarkar and A. Massacci, *J. Hazard. Mater.*, 2014, **278**, 189-202.

- 420 21 K. Furukawa, H. Suenaga and M. Goto, *J. Bacteriol.*, 2004, **186**, 5189-5196.
- 421 22 V. B. Urlacher, M. Girhard, *Trends Biotechnol.*, 2012, **30**, 26-36.
- 422 23 H. Ren, Y. Zhan, X. Fang and D. Yu, *RSC Advances*, 2014, **4**, 62631-62638.
- 423 24 Y. Lu, Y. Yu, R. Zhou, W. Sun, C. Dai, P. Wan, L. Zhang, D. Hao and H. Ren, *Biotechnol. Lett.*,
- 424 2011, **33**, 1159-1167.
- 425 25 K. H. Mitchell, C. E. Rogge, T. Gierahn and B. G. Fox, *Proc. Natl. Acad. Sci. U S A*, 2003, **1**,
- 426 3784-3789.
- 427 26 J. P. Uetrecht and W. Trager, In *Drug Metabolism: Chemical and Enzymatic Aspects: Textbook*
- 428 *Edition*; Informa Healthcare: New York, 2007; pp 91.
- 429 27 C. Aeppli, M. Tysklind, H. Holmstrand and Ö Gustafsson, *Environ. Sci. Technol.*, 2013, **47**,
- 430 790-797.
- 431 28 S. Eswaramoorthy, J. B. Bonanno, S. K. Burley and S. Swaminathan, *Proc. Natl. Acad. Sci.*, 2006,
- 432 **103**, 9832-9837.
- 433 29 M. M. Huijbers, S. Montersino, A. H. Westphal, D. Tischler, W. J. van Berkel, *Arch. Biochem.*
- 434 *Biophys.*, 2014, **544**, 2-17.
- 435 30 C. Binda, R. M. Robinson, J. S. Martin Del Campo, N. D. Keul, P. J. Rodriguez, H. H. Robinson,
- 436 A. Mattevi and P. Sobrado, *J. Biol. Chem.*, 2015, **290**, 12676-12688.
- 437 31 T. Liu and P. J. Chapman, *FEBS Lett.*, 1984, **173**, 314-318.
- 438 32 K. Makdessi and U. Lechner, *FEMS Microbiol. Lett.*, 1997, **157**, 95-101.
- 439 33 A. Aliverti, B. Curti and M. A. Vanoni, *Methods Mol. Biol.*, 1999, **131**, 9-23

Table 1

Catalysis preference analysis of the enzyme against biphenyl derivatives.

| Substrate | Substituent group | Occupied position | Relative activity (%) ^a |
|-----------------------------|-------------------|-------------------|------------------------------------|
| Biphenyl | NA ^b | – | 16 ± 1 |
| 4-Chlorobiphenol | Cl | 4 | 235 ± 9 |
| 4-Hydroxybiphenyl | OH | 4 | 131 ± 7 |
| 4,4'-Dichlorobiphenyl | Cl | 4,4' | 96 ± 8 |
| 4,4'-Dihydroxybiphenyl | OH | 4,4' | 45 ± 3 |
| 4-Hydroxy-4'-chlorobiphenyl | OH, Cl | 4,4' | 76 ± 6 |
| 4-Hydroxy-2-chlorobiphenyl | OH, Cl | 4,2 | 273 ± 15 |
| 4-Hydroxy-3-chlorobiphenyl | OH, Cl | 4,3 | 12 ± 1 |

^a Relative activity is expressed as a percentage of the maximum enzyme activity towards its regarded natural substrate 2,4-DCP at 25°C without addition of FAD. The specific activity is given as percentage of the activity towards 2,4-DCP, which corresponded to 1.55 U (mg protein)⁻¹ at 25°C.

^b–, Not determined.

Table 2

Kinetic parameters of 2,4-DCP hydroxylase towards biphenyl and its derivatives.

| Substrate | K_m (μM) | k_{cat} (min^{-1}) | k_{cat}/K_m ($\text{min}^{-1} \mu\text{M}^{-1}$) |
|-----------------------------|-------------------------|--|---|
| Biphenyl | 73.6 ± 3.2 | 7.0 ± 0.3 | 0.095 ± 0.006 |
| 4-Chlorobiphenol | 5.0 ± 0.2 | 102.3 ± 5.7 | 20.5 ± 0.9 |
| 4-Hydroxybiphenyl | 9.2 ± 0.5 | 56.9 ± 4.2 | 6.2 ± 0.2 |
| 4,4'-Dichlorobiphenyl | 12.3 ± 0.7 | 41.8 ± 2.9 | 3.4 ± 0.1 |
| 4,4'-Dihydroxybiphenyl | 26.1 ± 1.4 | 19.5 ± 0.8 | 0.75 ± 0.03 |
| 4-Hydroxy-4'-chlorobiphenyl | 15.5 ± 0.7 | 33.0 ± 1.1 | 2.1 ± 0.2 |
| 4-Hydroxy-2-chlorobiphenyl | 4.2 ± 0.1 | 118.5 ± 5.2 | 28.2 ± 1.8 |
| 4-Hydroxy-3-chlorobiphenyl | 98.7 ± 4.8 | 5.2 ± 0.2 | 0.053 ± 0.004 |

Figure legend

Fig. 1. Structures and names of the biphenyl derivatives investigated in the study.

Fig. 2. Specific activity of 2,4-DCP hydroxylase against biphenyl and its derivatives at 25°C (black column) and 0°C (gray column). Relative activity is expressed as a percentage of the maximum enzyme activity against its regarded natural substrate 2,4-DCP at 25°C, which corresponded to 1.55 U (mg protein)⁻¹.

Fig. 3. 2,4-DCP hydroxylase removal of biphenyl and its derivatives at 25°C (black column) and 0°C (gray column). The removal of biphenyl and its derivatives after 1 h was calculated by dividing the concentration of the amount of reduction of NADPH by the amount of the initial NADPH.

Fig. 4. Effect of reaction time on biphenyl and its derivatives removal. Gray columns stand for the biphenyl and its derivatives removal, respectively after 1 h reaction at 25°C with 5 μM FAD. Black column stand for the improvement of biphenyl and its derivatives removal, respectively after 24 h reaction.

Fig. 5. Cofactor requirement for hydroxylase activity against biphenyl and its derivatives (a) at 25°C and (b) at 0°C. Black and gray column stand for the reaction without 5 μM FAD and with FAD, respectively. Digits with underline above the column stand for the growth factors of hydroxylase activity against different biphenyl and its derivatives by addition of FAD, respectively.

Fig. 6. Cofactor requirement for biphenyl and its derivatives removal (a) at 25°C and (b) at 0°C. Black and gray column stand for the reaction without 5 μM FAD and with FAD, respectively. Digits with underline above the column stand for the growth factors against different biphenyl and its derivatives removal by addition of FAD, respectively.

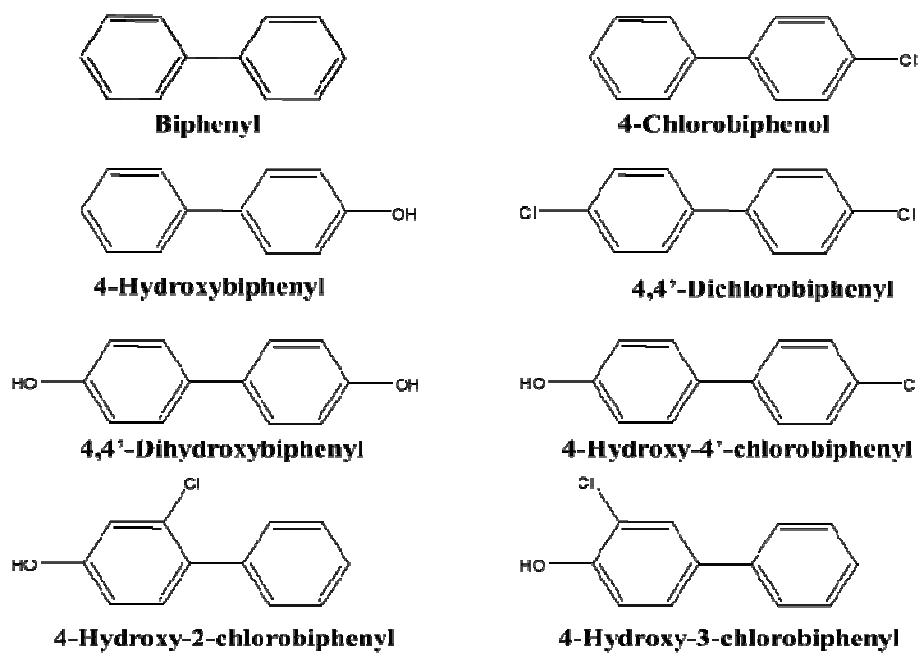


Fig. 1

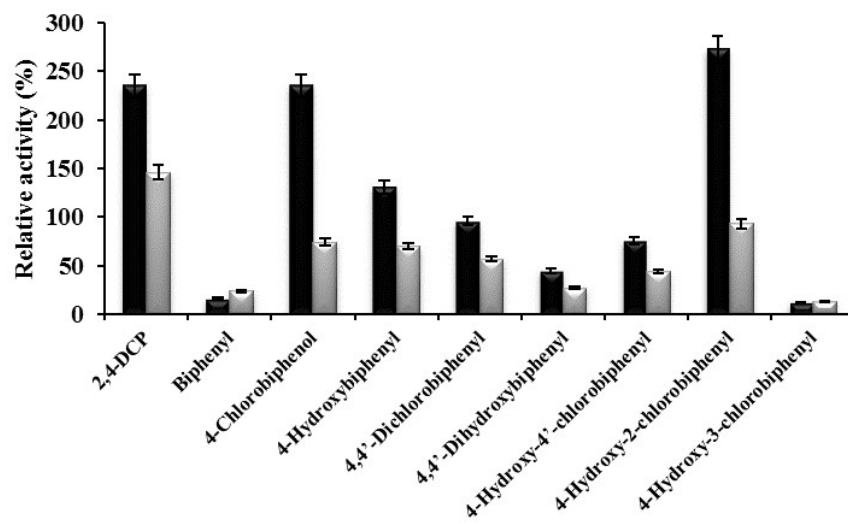


Fig. 2

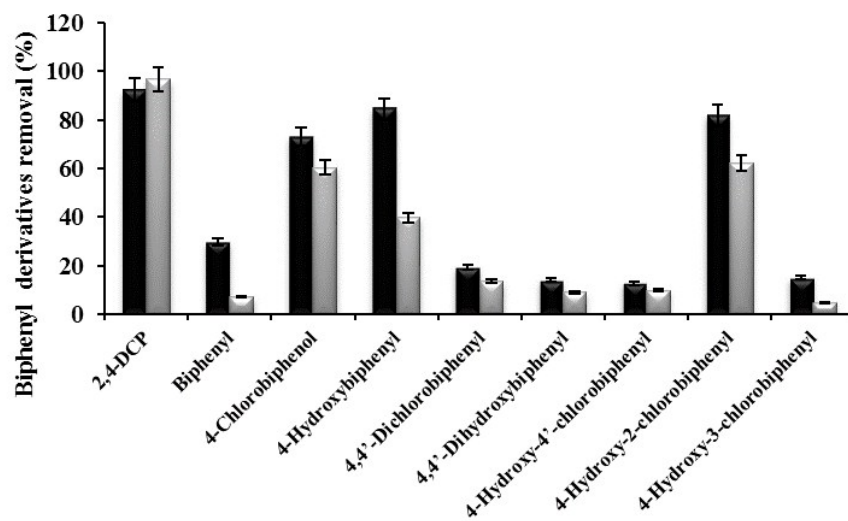


Fig. 3

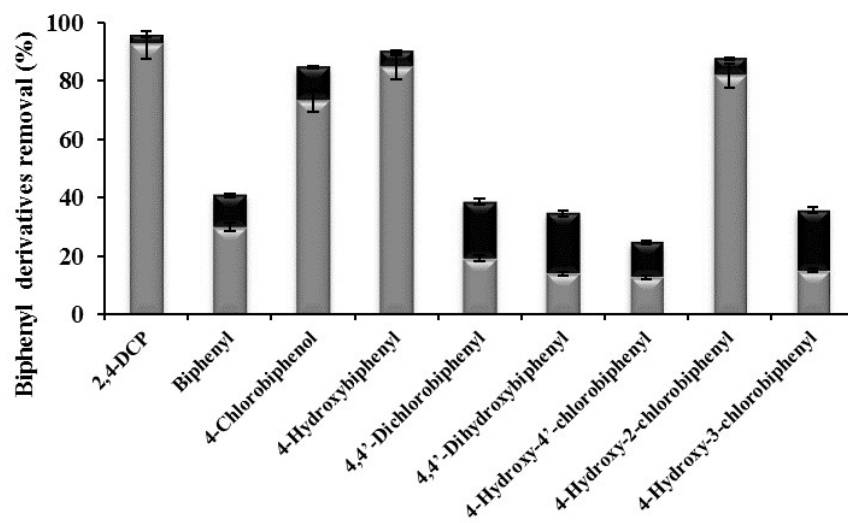
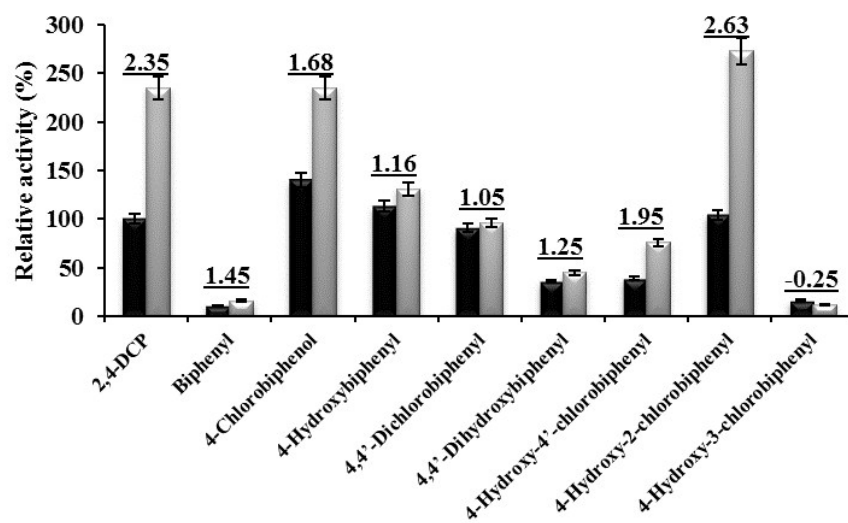


Fig. 4

a



b

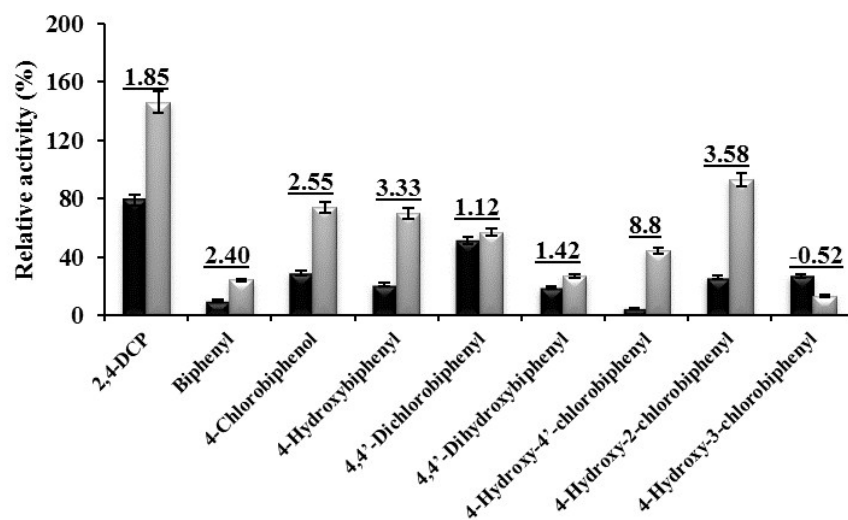
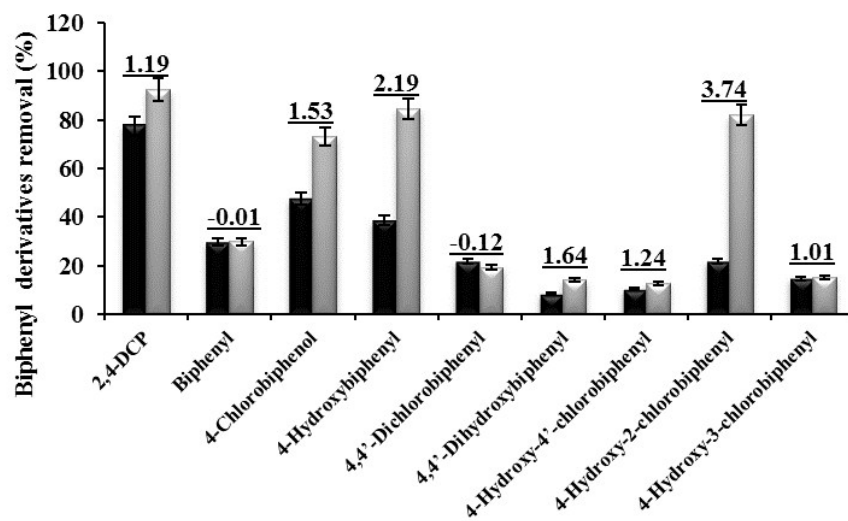


Fig. 5

a



b

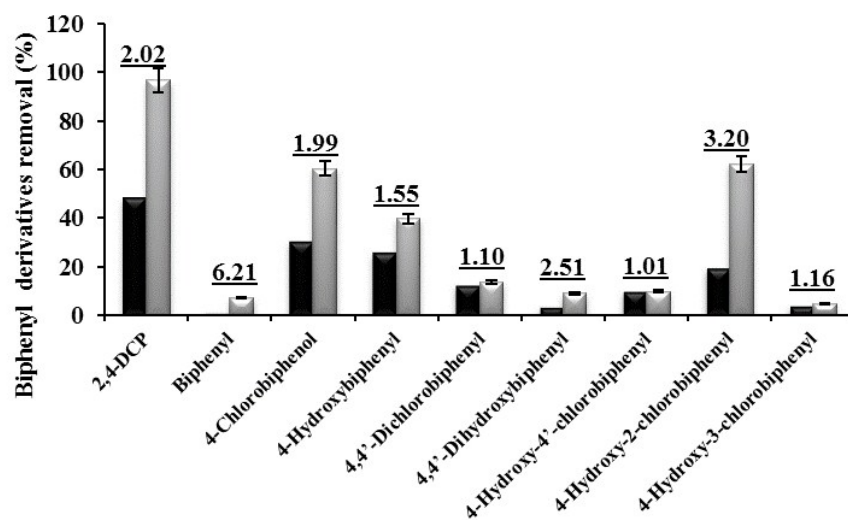


Fig. 6