



# Mineralization of high concentrations of the endocrine disruptor dibutyl phthalate by *Fusarium culmorum*

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## Abstract

Dibutyl phthalate (DBP) is a widely used plasticizer, whose presence in the environment as a pollutant raises concern because of its endocrine-disrupting toxicity. Growth kinetics, glucose uptake, biodegradation constant of DBP ( $k$ ), half-life of DBP biodegradation ( $t_{1/2}$ ) and percentage of removal efficiency ( $%E$ ) were evaluated for *Fusarium culmorum* grown on media containing glucose and different concentrations of DBP (500 and 1000 mg/l). Intermediate compounds of biodegraded DBP were identified by GC–MS and a novel DBP biodegradation pathway was proposed on the basis of the intermolecular flow of electrons of the intermediates identified using quantum chemical modeling. *F. culmorum* degraded 99% of both 1000 and 500 mg of DBP/l after an incubation period of 168 and 228 h, respectively.  $%E$  was 99.5 and 99.3 for 1000 and 500 mg of DBP/l, respectively. The  $k$  was 0.0164 and 0.0231 h<sup>-1</sup> for 500 and 1000 mg of DBP/l, respectively. DBP was fully metabolized to fumaric and malic acids, which are compounds that enter into the Krebs cycle. *F. culmorum* has a promising ability for bioremediation of environments polluted with DBP because it efficiently degrades DBP and uses high concentrations of this compound as carbon and energy source.

**Keywords** Constant of biodegradation · Dibutyl phthalate · *Fusarium culmorum* · Quantum chemical modeling · Removal efficiency

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## Introduction

Phthalates or esters of phthalic acid are xenobiotic organic compounds that are added to plastics to make them soft and flexible (plasticizers). Plasticizers are the largest group of additives in polymers. Low-molecular weight phthalates, such as dibutyl phthalate (DBP), are used to impart flexibility to thin films. It is one of the most abundantly produced and used plasticizers. It is used as a solvent in personal-care products (e.g., perfumes, lotions), elastomers, lacquers, explosives, printing inks, paper coatings, adhesives, resin solvents, and nail polish (Meeker et al. 2009). These compounds are not chemically bound to the polymer mesh, being released into the external environment during manufacture, use, disposal, leaking or evaporation from container or land-fill sites. This has become a major current environmental concern. DBP is one of the predominant phthalate esters present in atmospheric particles as well as in fresh water and sediments (Gao and Wen 2016). At phthalates production sites, concentrations in surface water range from 1 to 220 µg/l and from 7.5 to 2045 mg/kg in sediment (Green

Facts 2008). The highest levels in air occur around polyvinyl chloride (PVC) processing plants (ECB 2003). DBP can be taken up by plants and other living organisms, thereby entering the food supply in both marine and fresh environments (Muneer et al. 2001). Several studies have shown that DBP has anti-androgenic and estrogenic effects in male rats and fish and has also been reported to cause reproductive defects in humans (Chen et al. 2014; Xu et al. 2014). It has been reported that shorter ester chains such as DBP can be easier to biodegrade and mineralize than phthalates with long ester chains such as diethyl hexyl phthalate (DEHP) (Liang et al. 2008). Therefore, the occurrence of phthalate esters in the environment depends on their use, production and biodegradability. The hydrolysis and photolysis of phthalate esters are very slow. Microbial degradation plays a major role in mineralizing phthalate esters in the environment (Prasad and Suresh 2012; Pradeep et al. 2013; Ahuactzin-Pérez et al. 2014; Bouchiat et al. 2015; Meng et al. 2015). Bhardwaj et al. (2012) reported that microbial enzymes are one of the most powerful tools for biodegradation of plastics and that the activity of most enzymes is higher in fungi than in bacteria. Enzymes are able to transform pollutants and are potentially suitable to restore polluted environments (Rao et al. 2010). Esterases catalyze the hydrolysis of ester bonds of various compounds (i.e., lipids, oils, and phthalates) and break them down into the corresponding carboxylic acids (Benjamin et al. 2015). It is important to identify the pathways of biodegradation to insure that toxic compounds are not generated as a result of microbial degradation (Wackett 2014). Some species of the genus *Fusarium* have been shown to be highly efficient phthalate-degrading fungi due to their secretion of enzymes such as esterase (Kim et al. 2007; Obruca et al. 2012; Pradeep and Benjamin 2012; Chhaya and Gupte 2013; Aguilar-Alvarado et al. 2015; Bouchiat et al. 2015; Canavati-Alatorre et al. 2016). Quantum chemical calculations have been used to study biological systems from different approaches (Kaneco et al. 2006; Merz 2014; Ahuactzin-Pérez et al. 2016). Ahuactzin-Pérez et al. (2016) reported that calculation of the molecular quantum parameters was useful for proposing the DEHP biodegradation pathway by *Fusarium culmorum*. The calculation of quantum chemical parameters, such as the electrostatic potential of molecules, allows understanding of the molecular alignment and estimation of the electron transfer coefficient (ETC) (González-Pérez 2015). The quantum method, i.e., the semi-empirical parameterized model number 3 (SE-PM3), has been used to correlate molecular quantum chemical parameters and experimental data (Odunola and Semire 2007). Determination of chemical transformations helps to obtain a crucial understanding of how chemical reactions take place. In this work, the specific growth rate ( $\mu$ ), maximum biomass ( $X_{\max}$ ), biodegradation constant of DBP ( $k$ ), half-life of DBP biodegradation ( $t_{1/2}$ ) and percentage of

removal efficiency ( $\%E$ ) were evaluated for *F. culmorum* grown on media containing glucose and different concentrations of DBP (500 and 1000 mg/l). Molecular quantum chemical parameters, such as the lowest unoccupied molecular orbital energy ( $E_{\text{LUMO}}$ ), the highest occupied molecular orbital energy ( $E_{\text{HOMO}}$ ), the electrostatic potential ( $E\delta$ ), molecular partial positive charge ( $\delta+$ ), the molecular partial negative charge ( $\delta-$ ) and the energy of the band gap ( $E_{\text{Bg}}$ ), were calculated to determine the ETC for each compound of DBP biodegradation, which were identified by gas chromatography coupled to mass spectroscopy (GC-MS). A biodegradation pathway of DBP based on quantum chemical modeling was proposed.

## Materials and methods

### Chemicals and media

DBP was purchased from Sigma (purity grade 99%). Three different kinds of liquid culture media were used: (1) glucose yeast extract (GYE), (2) GYE + 500 mg DBP/l, and (3) GYE + 1000 mg DBP/l. Each medium contained (in g/l): glucose (10), yeast extract (5),  $\text{KH}_2\text{PO}_4$  (0.6),  $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$  (0.5),  $\text{K}_2\text{HPO}_4$  (0.4),  $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$  (0.25),  $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$  (0.05),  $\text{MnSO}_4$  (0.05), and  $\text{ZnSO}_4 \cdot 7\text{H}_2\text{O}$  (0.001). The final pH was adjusted to 6.5 using either 0.1 M HCl or 0.1 M NaOH. All media also contained 400  $\mu\text{l}$  of Tween 80 per liter. DBP (dibutyl phthalate) was added to the medium before autoclaving (boiling point 340 °C).

### Strain and culture conditions

*Fusarium culmorum* from the culture collection of the Research Centre for Biological Sciences at Universidad Autónoma de Tlaxcala, Mexico (Fc1-CICBUAT), was used. This fungus was isolated from the pulping of the recycled paper industry in which phthalates can be present as emulsifiers in paper dyes and as adhesives for paper envelopes and inks (Aguilar-Alvarado et al. 2015). This organism was grown on malt extract agar (DIFCO) at 20 °C and kept on agar plates at 4 °C until it was needed. Erlenmeyer flasks (125 ml) containing 50 ml of culture medium were autoclaved at 120 °C for 15 min, cooled to room temperature, and then inoculated with three mycelial plugs (of 10 mm diameter) taken from the periphery of colonies of *F. culmorum* (7 days old) grown on potato dextrose agar (DIFCO).

Cultures were incubated for 10 days at 28 °C on a rotary shaker operated at 120 rpm. Analyses were performed on samples taken at 12-h intervals and performed in triplicate. The concentrations of DBP employed have been used previously in fungal studies (Suárez-Segundo et al. 2013).

## Growth kinetics, glucose consumption and pH measurements

Biomass ( $X$ ) was harvested from cultures by filtration using filter paper (pore size 20–25  $\mu\text{m}$ ), and the specific growth rate ( $\mu$ ) was calculated in terms of changes in dry weight (g/l) using the Velhurst–Pearl or logistic equation (Eq. 1), as previously reported (Ahuactzin-Pérez et al. 2016):

$$\frac{dX}{dt} = \mu \left[ 1 - \frac{X}{X_{\max}} \right] X, \quad (1)$$

where  $\mu$  is the maximal specific growth rate and  $X_{\max}$  is the maximal (or equilibrium) biomass level achieved when  $dX/dt = 0$  for  $X > 0$ . The solution of Eq. (1) is as follows;

$$X = \frac{X_{\max}}{1 + Ce^{-\mu t}}, \quad (2)$$

where,  $C = (X_{\max} - X_0)/X_0$ , and  $X = X_0$ ; the initial biomass value.

Estimation of kinetic parameters in the above equations was performed using a non-linear least square-fitting program “Solver” (Excel, Microsoft).

The specific biomass yield ( $Y_{X/S}$ ) was calculated as the coefficient of the linear regression of biomass concentration versus substrate concentration in grams of biomass/grams of substrate consumed.

As the only reducing sugar present in the media, glucose was measured using the dinitrosalicylic acid reagent (DNS, SIGMA). Procedure details are described in our previous report (Ahuactzin-Pérez et al. 2016). The pH of the culture filtrate (CF) was measured every 12 h using a digital potentiometer (Hanna Instruments, México).

## Identification of DBP biodegradation intermediates by GC–MS

Intermediate products of biodegradation of DBP were analyzed by GC–MS, using a GC/MSD Agilent 890A gas chromatograph coupled to a mass spectrometer (5975C VL MSD, USA), equipped with a triple-axis detector and a capillary Agilent column (HP-5MS, USA) with 5% phenyl methyl siloxane. 1 ml of each supernatant was previously placed at 95 °C for 2 min in a water bath to denature protein. A 2- $\mu\text{l}$  aliquot of the supernatant from cultures was injected with an injection source (GC ALS) at a port temperature of 300 °C with the purge valve on (split mode), using a split ratio of 20:1 and split flow of 21.14 ml/min. Helium was the carrier gas and a flow rate of approximately 1.057 ml/min was used. The chromatographic conditions were; an initial oven temperature of 90 °C for 5 min, followed by a temperature ramp to 290 °C at 20 °C/min, and then held

for 15 min using a syringe size of 10  $\mu\text{l}$  in split mode to 300 °C at 20:1 in the injector. Data were analyzed with the MSD ChemStation software (Agilent Technologies), and intermediates were identified using the Nist MS 2.0 library (Ahuactzin-Pérez et al. 2016). Spectral match factors were greater than 90%.

## Percentages of biodegradation and removal efficiency, and half-life of DBP

Percentage of biodegradation was determined by measuring the disappearance of DBP in the CF. The biodegradation percentage of DBP ( $\%B_{\text{DBP}}$ ) in a sample was calculated as follows:

$$\%B_{\text{DBP}t} = \%C_0 - \%C_t, \quad (3)$$

where  $C_0$  = initial concentration of DBP in the test solution,  $C_t$  = concentration of DBP in the test solution at time  $t$  (h).

Percentage of removal efficiency ( $\%E$ ) of DBP was calculated as follows:

$$\%E = 1 - (C_f/C_0) \cdot 100, \quad (4)$$

where  $C_f$  = concentration of DBP at the end time ( $t = 240$  h).

The half-life ( $t_{1/2}$ ) of DBP corresponded to the time interval of the concentration of DBP to decrease to half of its initial value (Benjamin et al. 2015). It was calculated according to the following equation:

$$\text{At initial time } t_0, \ln C = -kt + \ln C_0, \quad (5)$$

where  $C$  = concentration of DBP in mg/l,  $t$  = time (h) and  $k$  = degradation constant:

$$t_{1/2} = 0.693/k, \quad (6)$$

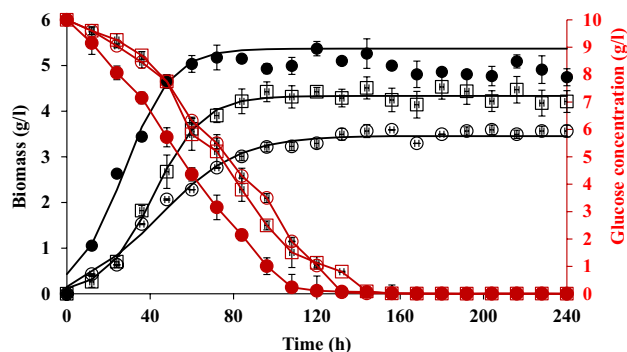
## Quantum chemical investigation

A DBP biodegradation pathway was proposed based on the compounds identified by GC–MS as intermediates from DBP and the quantum chemical parameters of each compound (according to the principles of molecular orbital theory).  $E_{\text{LUMO}}$ ,  $E_{\text{HOMO}}$ ,  $\delta+$  and  $\delta-$  were calculated using the SE-PM3 as previously reported (Ahuactzin-Pérez et al. 2016).  $E_{\text{Bg}}$ ,  $E\delta$  and ETC were calculated as follows;

$$\begin{aligned} E_{\text{Bg}} &= E_{\text{HOMO}} - E_{\text{LUMO}}, \\ E\delta &= |(\delta-) - (\delta+)|, \\ \text{ETC} &= E_{\text{Bg}}/E\delta. \end{aligned} \quad (7)$$

## Data analysis

All reported data are the mean and standard deviation of three replicates. Statistical analysis of data was performed using one-way ANOVA at the significance level of 0.05 and Tukey



**Fig. 1** Glucose consumption and biomass production by *Fusarium culmorum* on GYE medium (white circle), 500 (white square) and 1000 (black-filled circle) mg of DBP/l in submerged fermentation. Biomass curves were fitted (lines) using the logistic equation (Eqs. 1 and 2). Error bars are the SEM from triplicate experiments

post-test using the Graph Pad Prism® program (San Diego, CA, USA).

## Results

### Growth kinetics, glucose consumption and pH of the cultures

Biomass production and glucose consumption by *F. culmorum* on media supplemented with 500 and 1000 mg of DBP/l are shown in Fig. 1. *F. culmorum* produced a higher amount of biomass in medium supplemented with 1000 mg of DBP/l than in medium containing 500 mg of DBP/l. The organism reached the stationary phase after 72 and 84 h of growth in media supplemented with 1000 mg of DBP/l and 500 mg of DBP/l, respectively. Glucose was most rapidly consumed and the greatest amount of biomass was produced in the medium containing 1000 mg of DBP/l. Glucose was completely consumed by *F. culmorum* after 120 and 144 h in medium added with 1000 mg of DBP/l and in medium supplemented with 500 mg of DBP/l, respectively. Higher  $X_{max}$ ,  $\mu$  and  $Y_{X/S}$  were shown in media supplemented with 1000 mg of DBP/l than in medium containing 500 mg of DBP/l (Table 1). Cultures of both DBP concentrations showed a similar pH pattern during the fungal growth. The pH of the cultures dropped from the starting pH 6.5–5.1 after 24 h. pH value of the cultures increased after 36 h of growth, reaching a pH value of 8.4 at the termination of the fermentation (Fig. 2).

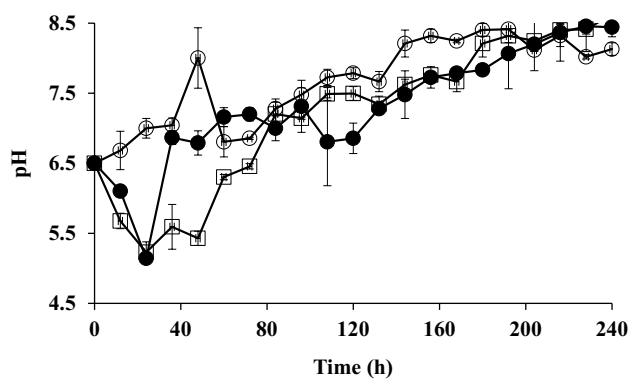
### Percentages of biodegradation and removal efficiency of DBP

The disappearance of DBP (in percentage) during the fermentation is shown in Fig. 3a. *F. culmorum* degraded 99%

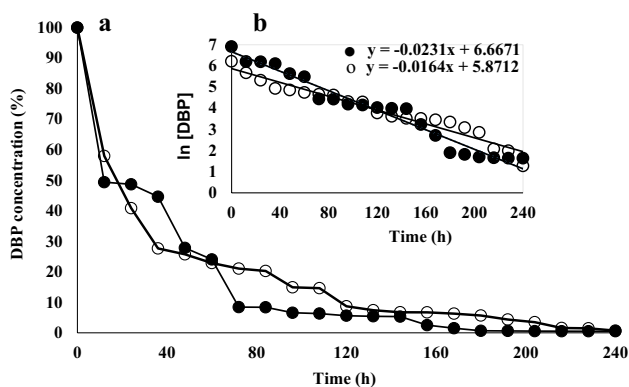
**Table 1** Growth kinetic parameters of *Fusarium culmorum* grown in GYE medium and in DBP supplemented media under submerged fermentation conditions

Growth kinetic parameters	Culture media		
	GYE	DBP (mg/l)	
		500	1000
$X_{max}$ (g/l)	3.4 <sup>c</sup> (0.10)	4.3 <sup>b</sup> (0.06)	5.0 <sup>a</sup> (0.21)
$\mu$ (h <sup>-1</sup> )	0.05 <sup>c</sup> (0.001)	0.08 <sup>b</sup> (0.006)	0.10 <sup>a</sup> (0.006)
$Y_{X/S}$ (gX/gS)	0.34 <sup>c</sup> (0.01)	0.43 <sup>b</sup> (0.01)	0.49 <sup>a</sup> (0.02)

Values are expressed as mean  $\pm$  SD ( $n = 3$ ); means within the same column not sharing common superscript letters (a–c) differ significantly at 5% level. Kinetic parameters ( $X_{max}$  and  $\mu$ ) of the logistic equation were evaluated using a non-linear least squares fitting program



**Fig. 2** pH of *Fusarium culmorum* grown on GYE medium (white circle), 500 (white square) and 1000 (black-filled circle) mg of DBP/l in submerged fermentation. Error bars are the SEM from triplicate experiments



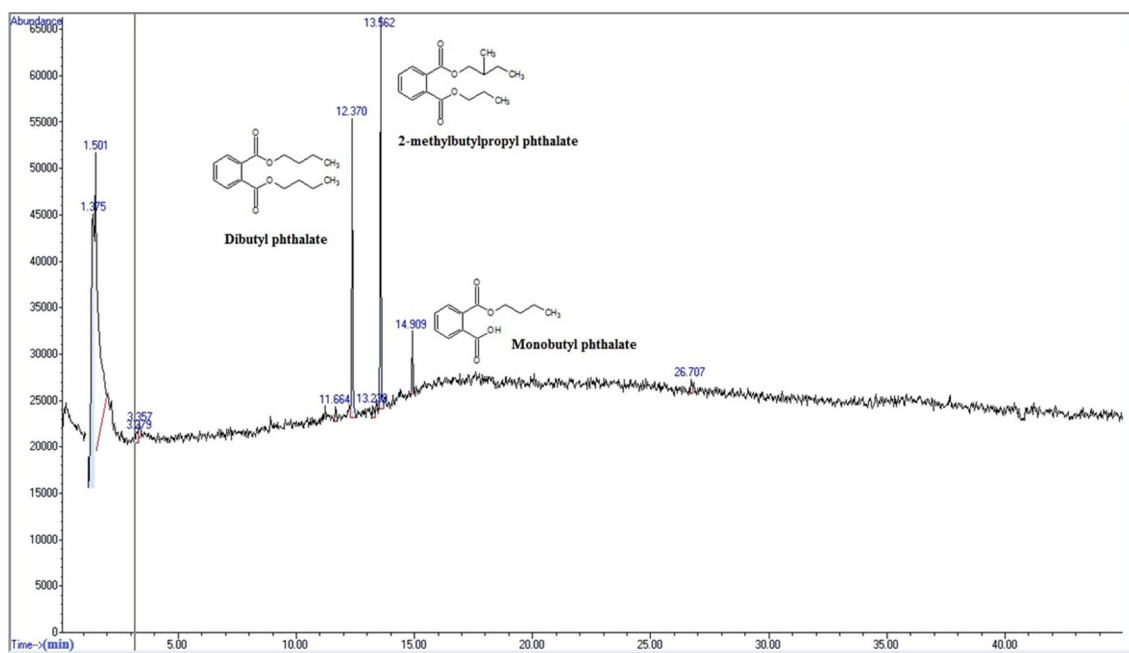
**Fig. 3** Biodegradation percentage of DBP by *Fusarium culmorum* (a) and DBP concentration (log-transformed) plotted across time used to determine  $k$  (b). Media with 500 (white circle) and 1000 (black-filled circle) mg of DBP/l added

of both 1000 mg of DBP/l and 500 mg of DBP/l after an incubation period of 168 and 228 h, respectively (Eq. 3) (Fig. 3a). The logarithm of DBP concentration versus time was plotted, and was used to calculate the first-order biodegradation constant of DBP (Fig. 3b). %E was 99.5 and 99.3 for 1000 and 500 mg of DBP/l, respectively, (Eq. 4). The  $k$  was around 50% higher for the medium supplemented with 1000 mg of DBP/l ( $0.0231 \text{ h}^{-1}$ ) than that  $k$  containing 500 mg of DBP/l ( $0.0164 \text{ h}^{-1}$ ) (Eq. 5).  $t_{1/2}$  was 42 and 30 h for 500 and 1000 mg of DBP/l, respectively, (Eq. 6) (Fig. 3b).

### Proposed DBP biodegradation pathway

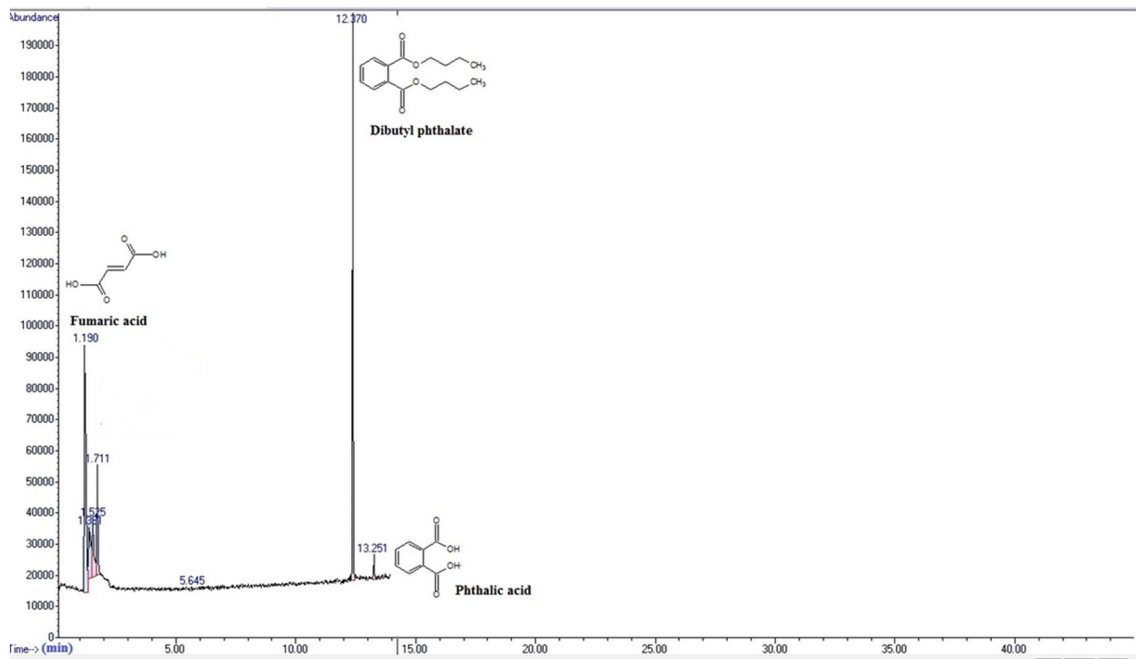
The DBP biodegradation pathway by *F. culmorum* was proposed on the basis of the intermolecular flow of electrons of the identified intermediate using quantum chemical modeling. Compounds identified by GC–MS (Figs. 4, 5) are shown by dashed lines and the rest of the compounds in the biodegradation pathway were proposed using quantum chemical modeling (Fig. 6). ETC values (Eq. 7) for the compounds of the DBP biodegradation pathway are shown in the table (Online Resource 1). The DBP biodegradation pathway was achieved by a DBP equilibrium isomerization reaction, with the formation of 2-methylbutylpropyl phthalate (MBPP). DBP is demethylated to produce MBPP, and the reaction moves from a less excited state to an excited state because the ETC value for DBP was lower than the ETC value for MBPP, which means that the electrons

cannot easily move from DBP to MBPP. MBPP was likely metabolized by enzymatic ester hydrolysis to phthalic acid (PA). This reaction take place from an excited state to a less excited state because the ETC value for MBPP was higher than the ETC value for PA, which means that the electrons can easily move from MBPP to PA. Alternatively, DBP biodegradation seems to progress by enzymatic ester hydrolysis of DBP with the formation of monobutyl phthalate (MBP), and the reaction moves from a less excited state to an excited state because the ETC value for DBP was lower than the ETC value for MBP. This means that the electrons cannot easily move from DBP to MBP. MBP degradation would be continued by enzymatic ester hydrolysis with the production of PA. MBP had an ETC value higher than the ETC value of PA, which shows that the de-esterification occurs from an excited state to a less excited state, meaning that the electrons can easily move from MBP to PA. PA would be then metabolized to two different compounds: fumaric acid (FA) and but-1-e-yne (BE) by a retro Diels–Alder reaction and by the electron resonance effect. This reaction occurs for both compounds from a less excited state to an excited state because the ETC value for PA was lower than the ETC value for both FA and BE. This means that the electrons cannot easily move from PA to both compounds FA and BE. FA can enter into Krebs cycle to be metabolized to  $\text{CO}_2$  and  $\text{H}_2\text{O}$ . On the other hand, BE would be oxidized to (2Z)-2-butene-1,4-diol (BD), and the reaction would occur from an excited state to a less excited state because the ETC value for BE was higher than the ETC value for BD, meaning that



**Fig. 4** Chemical compounds produced during DBP degradation by *Fusarium culmorum*, as analyzed by GC/MS at a total run time of 60 min. Dibutyl phthalate (RT = 12.370 min), 2-methylbutylpropyl phthalate (RT = 13.562 min), monobutyl phthalate (RT = 14.909 min)





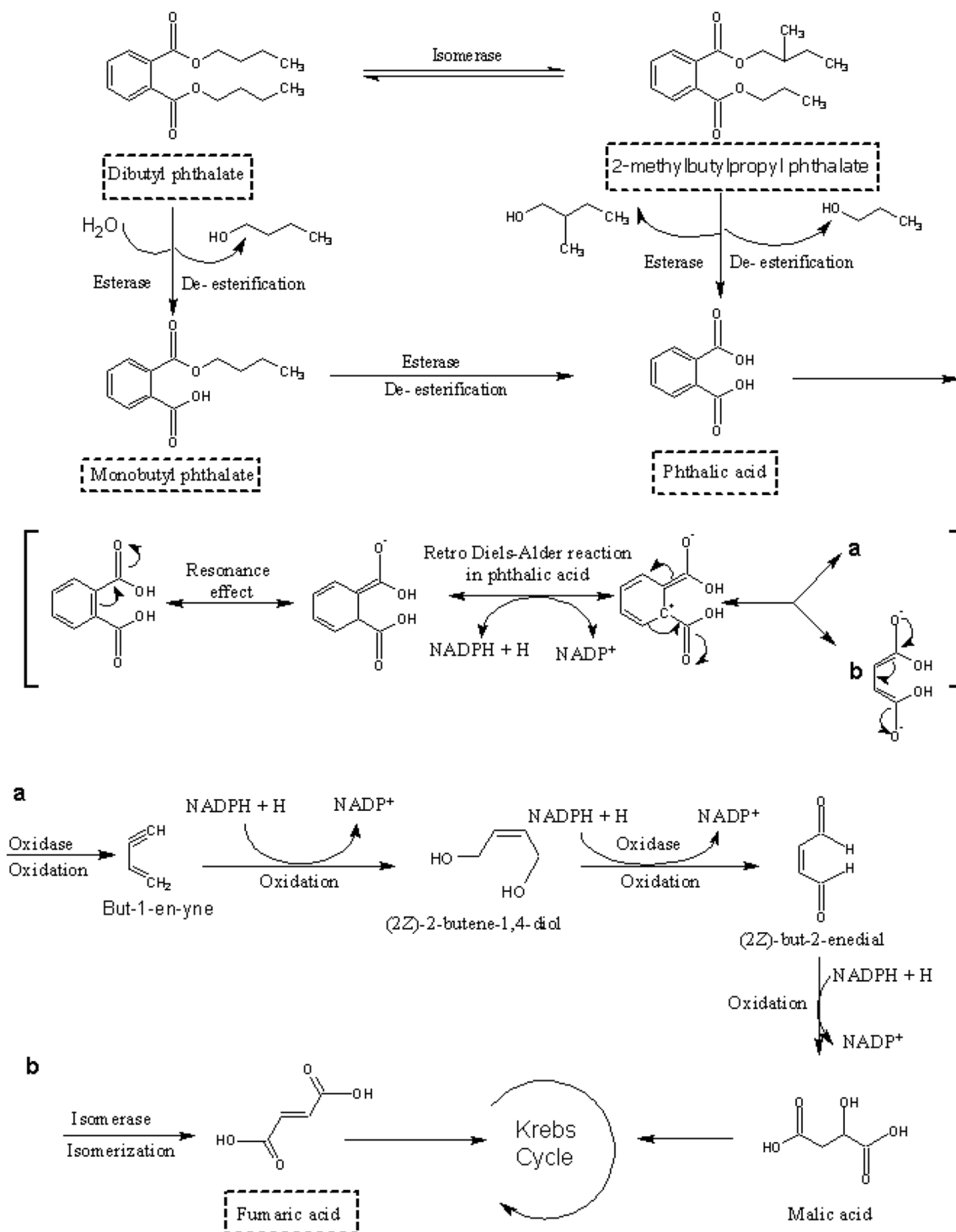
**Fig. 5** Chemical compounds produced during DBP degradation by *Fusarium culmorum*, as analyzed by GC/MS at a total run time of 60 min. Fumaric acid (RT = 1.190 min), dibutyl phthalate (RT = 12.370 min), phthalic acid (RT = 13.251 min)

the electrons can easily move from BE to BD. BD would also be oxidized to (2Z)-but-2-enedial (BDL). This reaction moves from a less excited state to an excited state because the ETC value for BD was lower than the ETC value for BDL, meaning that the electrons cannot easily move from BD to BDL. Finally, BDL is oxidized to malic acid (MA), and the reaction occurs from an excited state to a less excited state because the ETC value for BDL was higher than the ETC value for MA. This means that the electrons can easily move from BDL to MA. MA can enter the Krebs cycle to be metabolized to CO<sub>2</sub> and H<sub>2</sub>O (Fig. 4). In quantum chemical modeling, an enzyme should be used when a reaction occurs from a less excited state to an excited state. However, if a reaction occurs from an excited state to a less excited state, an enzyme may or may not be required.

## Discussion

*Fusarium culmorum* had higher biomass production and growth kinetic parameters in DBP-supplemented media, which shows that DBP was used by the fungus as a carbon and energy source. Biomass was much higher than the expected biomass in normal conditions. It could be due to the use of glucans (from mature hypha) by the fungus to growth (increasing biomass production). Sánchez et al. (2004) reported that hyphal ageing in the fungus involves glucan storage that is available, after cell wall lysis, as a

carbon source for the growth of the fungus (at the end of the exponential phase). However, further studies on alkali-soluble cell-wall glucan (*S*-glucan) and alkali-insoluble cell-wall glucan (*S*-glucan) ratio need to be carried to corroborate this suggestion. For example, the *S/R* glucan ratio increased about 3- to 4-fold during fruit body formation in basidiomycetes (Schwalb, 1978). We previously reported glucose consumption and specific growth rate of *F. culmorum* in GYE medium (medium lacking dibutyl phthalate) (Ahuactzin-Pérez et al. 2016). In the present research, *F. culmorum* actively grew in media supplemented with DBP since the glucose uptake was higher in DBP-supplemented media compared to medium lacking DBP (Ahuactzin-Pérez et al. 2016). The use of phthalate as carbon source depends on the microorganism used and complexity of the phthalate structure. It has been reported than phthalates with shorter ester chains such as DBP can be easier to biodegrade and mineralize than phthalates with long ester chains such as DEHP (Liang et al. 2008). In the present study, *F. culmorum* reached the stationary phase after 72 h of growth in medium supplemented with 1000 of DBP/l. Ahuactzin-Pérez et al. (2016) reported that *F. culmorum* grown on DEHP (500 and 1000 mg/l) attained the stationary phase after 4 days of growth. Córdoba-Sosa et al. (2014) reported that the edible mushroom *Pleurotus ostreatus* grown on different concentrations of DEHP (750, 1200 and 1500 mg/l) reached to the stationary phase after 15 days of growth. In this study, *F. culmorum* degraded 99% of both 1000 mg of DBP/l and



**Fig. 6** DBP biodegradation pathway by *Fusarium culmorum* proposed using quantum chemical modeling. The dashed line shows compounds identified by GC-MS

500 mg of DBP/l after 168 and 228 h, respectively. These results show that DBP induced esterase production, which is in agreement with the increase in biomass in 1000 mg of DBP/l (Fig. 1). The significance of esterase in phthalate ester degradation and its induction by these compounds have

already been reported. Córdoba-Sosa et al. (2014) reported that high concentrations of DEHP (1500 mg/l) induced esterase production, enhancing the growth of *P. ostreatus*. Moreover, Jin et al. (2016) reported that the DBP-degrading bacterium *Gordonia* sp. revealed the presence of putative

hydrolase/esterase genes involved in phthalate esters. Gao and Chi (2015) reported that intracellular and extracellular esterases play key roles in the degradation of DBP. Ahuactzin-Pérez et al. (2016) found that *F. culmorum* degraded DEHP (initial concentration 1000 mg/l) within 224 h. In general, *F. culmorum* degraded higher concentrations of DBP and showed higher efficiency of removal than other organisms. Studies about DBP degradation (initial concentration 50 mg/l) by *Mycobacterium* sp. revealed that 80% was degraded after 5 days (Ren et al. 2016). Tang et al. (2016) found that *Rhizobium* sp. LMB-1 had a removal capacity of nearly 100 mg of DBP/l within 60 h. Kumar and Maitra (2016) found that *Methylobacillus* sp. degraded 70% of DBP (initial concentration 2000 mg/l) after 192 h. Lee et al. (2007) reported that the white rot fungus *Polyporus brumalis* nearly eliminated a concentration of 350 mg of DBP/l within 12 days and that the mycelial growth of the fungus was inhibited. Jin et al. (2016) found that *Gordonia* sp. degraded DBP (initial concentration 750 mg/l) after 32 h. It has been reported that *Agrobacterium* sp. completely degraded DBP within 48 h when the initial concentration was lower than 200 mg/l (Wu et al. 2011). In the present study, the  $k$  of DBP for *F. culmorum* was 0.0164 and 0.0231 h<sup>-1</sup> for 500 and 1000 mg of DBP/l, respectively. Tang et al. (2016) studied DBP degradation (initial concentration 100 mg/l) by *Rhizobium* sp. LMB-1 and found that  $k$ , was 0.040 h<sup>-1</sup>. Ahuactzin-Pérez et al. (2016) reported that *F. culmorum* had a  $k$  of DEHP (initial concentration of 1000 mg/l) of 0.024 h<sup>-1</sup>. Hu et al. (2015) reported that  $k$  was 0.026 h<sup>-1</sup> for DBP (initial concentration: 100 mg/l) in a bioaugmented reactor using *Micrococcus* sp. It has been reported that the  $k$  of DBP (initial concentration: 2000 mg/l) using microalgal species was 0.0169, 0.0035, 0.0034 h<sup>-1</sup> for *Cylindrotheca closterium*, *Dunaliella salina* and *Chaetoceros muelleri*, respectively (Gao and Chi 2015). In the present study, %E was 99.5 and 99.3 for 500 and 1000 mg of DBP/l, respectively. Tang et al. (2016) studied DBP degradation (initial concentration 100 mg/l) by *Rhizobium* sp. LMB-1 and found that %E was 71.5. Ahuactzin-Pérez et al. (2016) found that the %E of DEHP (initial concentration 1000 mg/l) by *F. culmorum* was 99.8. Hu et al. (2015) reported that the %E was 85 for DBP (initial concentration 100 mg/l) in a bioaugmented reactor using *Micrococcus* sp. Microalgal species showed a %E of DBP by *D. salina*, *C. muelleri* and *C. closterium* of 40.0, 47.1 and 93.1, respectively (Gao and Chi 2015). In the present study,  $t_{1/2}$  was 42 and 30 h for 500 and 1000 mg of DBP/l, respectively. Ahuactzin-Pérez et al. (2016) reported that *F. culmorum*  $t_{1/2}$  was 28 h when the initial concentration was 1000 mg of DEHP/l. Tang et al. (2016) studied DBP degradation (initial concentration 100 mg/l) by *Rhizobium* sp. LMB-1 and found that  $t_{1/2}$  was 17 h. Fang et al. (2010) reported  $t_{1/2}$  by *Enterobacter* sp. to be about 21 h when DBP concentration was

500 mg/l. Xu et al. (2005) studied DBP biodegradation by *Pseudomonas fluorescens* at initial concentrations of 2.5–10 mg/l and found that  $t_{1/2}$  increased from approximately 14–24 h, respectively. Other methods of phthalate degradation are quite inefficient and slow. For example, it has been reported that the photodegradation and hydrolysis rates of phthalates are very slow under natural conditions. For example, butyl benzyl phthalate has an aqueous photolysis  $t_{1/2}$  of > 100 days, and dimethyl phthalate and DEHP have hydrolysis rates (at neutral pH)  $t_{1/2}$  of about 3 and 2000 years, respectively (Gao and Wen 2016). It is suggested that *F. culmorum* fully metabolized DBP with fumaric and malic acids as final products, which can enter the Krebs cycle. This fungus mineralized DBP, which is in accordance with alkaline pH in cultures at the end of the fermentation, since alkaline pH is observed when CO<sub>2</sub> dissolves in H<sub>2</sub>O because it dissociates into bicarbonate ions. MBP and phthalic acid (PA) were reported as intermediate compounds found in degradation of DBP in a batch reactor bioaugmented with *Micrococcus* sp. (Hu et al. 2015). Kumar and Maitra (2016) also found MBP, PA and pyrocatechol as intermediate compounds during DBP degradation by *Methylobacillus* sp. Metabolites of degradation of DBP (initial concentration 1000 mg/l) by *Rhizobium* sp. were found to be diethyl phthalate (DEP), PA, dimethyl phthalate, monomethyl ester and tartaric acid, suggesting that the former metabolite can enter the Krebs cycle (Tang et al. 2016). On the other hand, the white rot fungus *P. brumalis* produced MBP, DEP and phthalic acid anhydride as DBP degradation products (Lee et al. 2007). Patil et al. (2006) studied DBP degradation by *Delfia* sp. and found MBP, phthalate and protocatechuate (PC) as intermediates, suggesting that this bacterium can mineralize PC by a meta-cleavage pathway. In this study, *F. culmorum* completely metabolized DBP (500 and 1000 mg/l), forming end products (fumaric and malic acids) that can enter the Krebs cycle. *F. culmorum* has a promising ability for bioremediation of environments polluted with DBP because it efficiently degrades DBP and uses high concentrations of this compound as carbon and energy source. For further understanding of DBP biodegradation processes, future studies need to be carried out on isolation, purification, characterization and eventual expression for large-scale production of the enzymes involved in the DBP-degradation process.

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**Author contributions** MA-P; did the experimental work (Ph.D. thesis). ST-B and ES-J; drew the DBP biodegradation pathway. JG-D; did the GC-MS analysis. MG-P; did the quantum chemical studies. MCG-R; co-supervised MAP Ph.D. thesis and checked the MS. CS; co-supervised MAP Ph.D. thesis and wrote the MS.



## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest in the publication.

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