

Effects of Cu²⁺ and humic acids on degradation and fate of TBBPA in pure culture of *Pseudomonas* sp. strain CDT

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ABSTRACT

Soil contamination with tetrabromobisphenol A (TBBPA) has caused great concerns; however, the presence of heavy metals and soil organic matter on the biodegradation of TBBPA is still unclear. We isolated Pseudomonas sp. strain CDT, a TBBPA-degrading bacterium, from activated sludge and incubated it with ¹⁴C-labeled TBBPA for 87 days in the absence and presence of Cu²⁺ and humic acids (HA). TBBPA was degraded to organic-solvent extractable (59.4% \pm 2.2%) and non-extractable (25.1% \pm 1.3%) metabolites, mineralized to CO_2 (4.8% ± 0.8%), and assimilated into cells (10.6% ± 0.9%) at the end of incubation. When Cu²⁺ was present, the transformation of extractable metabolites into non-extractable metabolites and mineralization were inhibited, possibly due to the toxicity of Cu²⁺ to cells. HA significantly inhibited both dissipation and mineralization of TBBPA and altered the fate of TBBPA in the culture by formation of HA-bound residues that amounted to $22.1\% \pm 3.7\%$ of the transformed TBBPA. The inhibition from HA was attributed to adsorption of TBBPA and formation of bound residues with HA via reaction of reactive metabolites with HA molecules, which decreased bioavailability of TBBPA and metabolites in the culture. When Cu²⁺ and HA were both present, Cu²⁺ significantly promoted the HA inhibition on TBBPA dissipation but not on metabolite degradation. The results provide insights into individual and interactive effects of Cu2+ and soil organic matter on the biotransformation of TBBPA and indicate that soil organic matter plays an essential role in determining the fate of organic pollutants in soil and mitigating heavy metal toxicity.

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Introduction

Tetrabromobisphenol A [4,4'-isopropylidenebis (2,6dibromophenol), TBBPA] is a brominated flame retardant widely used in electric and electronic products (He et al., 2010). TBBPA was widely detected in the environment, such as soil, water, sewage sludge, dust, as well as human tissue (Feng et al., 2012; Gorga et al., 2013; Johnson-Restrepo et al., 2008; Tang et al., 2014; Wu et al., 2016; Xu et al., 2013). The highest level of TBBPA in soils of electronic wastes (e-wastes)

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dismantling sites detected in south China amounted to 1.94 mg/(kg·dw) (Wang et al., 2015). TBBPA is toxic to aquatic organisms and caused 100% mortality of zebrafish (Danio rerio) at 1.5 and 3 μ mol/L (McCormick et al., 2010). Although TBBPA showed low toxicity to mammals, cytotoxic effects of TBBPA were found in some studies (Colnot et al., 2014; Guyot et al., 2014), *e.g.*, affecting the host immunity of mice by significantly increasing the pulmonary viral titer (Watanabe et al., 2010).

The almost ubiquitous presence of TBBPA in the environment motivated several biodegradation investigations. TBBPA can be degraded in active sludge, soil, sediments or soil-plant systems under oxic and anoxic conditions (Brenner et al., 2006; Chang et al., 2012; Gerecke et al., 2006; Li et al., 2015b; Li et al., 2014a; Sun et al., 2014). TBBPA can also be degraded in pure culture of *Ochrobactrum* sp. T or *Comamonas* sp. strain JXS-2-02 isolated from oxic and anoxic sludge, respectively (An et al., 2011; Peng et al., 2013; Zu et al., 2014).

Due to the presence of the phenolic group and its high octanol/water partition coefficient (logK_{ow} = 5.9), TBBPA sorbs onto organic matter and minerals in soil or sediment and forms non-extractable (or bound) residues (Li et al., 2015b; Luo et al., 2010; Tong et al., 2015). It has been reported that in surface sediments, bound residues accounted for 6.5%-48.0% of the total TBBPA detected (Luo et al., 2010). Humic acids (HA) are major component of soil organic matrix and can react with phenolic organic pollutants to form HA-bound residues of pollutants and their transformation products (Li et al., 2012a; Riefer et al., 2013; Vinken et al., 2005). Interactions of pollutants with HA can affect mobility, bioaccumulation, and biodegradation of the pollutants in the environment (Kastner and Mahro, 1996) and the effects may be compound-specific (Burgos et al., 2000; Li et al., 2007; Ortega-Calvo and Saiz-Jimenez, 1998; Vinken et al., 2004). It has been shown that biodegradation of TBBPA was responsible for the formation of bound residues of TBBPA and its metabolites in soil under both oxic and anoxic conditions (Li et al., 2015b; Liu et al., 2013). However, direct evidence for relationship between binding of TBBPA-derived residues to HA and biodegradation of TBBPA does not exist.

Coexistence of TBBPA and heavy metals has been often detected in soil (Wager et al., 2012; Wu et al., 2016; Zhou et al., 2013), because electric and electronic products contain both brominated flame-retardants and heavy metals. Copper (Cu), a common heavy metal used in electronic products (Stuhlpfarrer et al., 2016; Yang et al., 2016), can be toxic to microorganisms and enzymes (Ochoa-Herrera et al., 2011; Wang et al., 2007) and can inhibit biodegradation of organic pollutants, e.g., fluorine and biphenyl, in both pure bacterial culture and soil (Nam et al., 2015; Wannoussa et al., 2015). In addition, heavy metals can affect conformational structure and hydrophobicity of HA (Nuzzo et al., 2013; Stevenson, 1977) and therefore the interaction between HA and organic pollutants. In addition, complexation of Cu^{2+} with functional groups of HA may reduce the concentration of free Cu²⁺ (Klucakova, 2014; Li et al., 2015a) and therefore reduce Cu toxicity to microorganisms. The co-effects of HA and Cu on TBBPA biodegradation are still unknown.

In this study, we isolated a TBBPA-degrading bacterium from an oxic activated sludge and studied the dissipation, degradation, metabolism, and bound-residue formation of TBBPA in pure culture of this bacterium in the absence and presence of Cu^{2+} and HA, in order to provide insights into the individual and combined effects of Cu^{2+} and HA on the biotransformation and fate of TBBPA in soil environment.

1. Materials and methods

1.1. Chemicals and humic material

Uniformly ¹⁴C-ring-labeled TBBPA (¹⁴C-TBBPA) was synthesized from uniformly ¹⁴C-ring-labeled phenol *via* BPA (Liu et al., 2013). ¹⁴C-TBBPA had a specific activity of 1.22 × 10⁹ Bq/mmol and a radiochemical purity of 99%. Non-labeled TBBPA was purchased from Tokyo Chemical Industry Co., LTD with purity of 99% (Tokyo, Japan). Other chemicals of chromatographic or analytical grade were purchased from Nanjing Chemical Reagent, LTD (Nanjing, China).

HA were extracted from a pine forest soil in Aachen, Germany (Vinken et al., 2005). To prepare an HA stock solution, HA powder was added to a mineral salt medium (MSM) at pH 7.2 (see below), and then 4 mol/L NaOH was added dropwise to the medium until the HA were completely dissolved. This preparation procedure prevented oxidation of HA by air under alkaline condition.

1.2. MSM medium

The MSM had the following composition (per liter): 1 g of NaCl, 0.5 g of KCl, 0.1 g of MgCl₂·6H₂O, 0.015 g of CaCl₂·2H₂O, 2 g of NH₄Cl, 0.2 g of KH₂PO₄, 0.15 g of Na₂SO₄, sodium phosphate buffer (pH 7.2; 20 mmol/L; autoclaved separately), and 0.5% of the trace element solution SL11 (Brune et al., 1995).

1.3. Isolation and identification of bacterial strains

Bacterial strains were isolated from an activated sludge sample collected from the Dachang sewage treatment plant in Nanjing, China. The sludge was washed with a phosphate buffer (20 mmol/L, pH 7.2) three times. Aliquot of the washed sludge (10 mL) with a mixed liquor suspended solids (MLSS) $(1.33 \times 10^4 \text{ mg/L})$ was added to MSM (40 mL) containing 100 mg/L TBBPA in a 150-mL flask, and incubated at 30°C on a rotary shaker (130 r/min). About 10% (V/V) of the culture was inoculated into 45 mL fresh MSM containing 100 mg/L TBBPA once every week in the first month and every other week in the next two months to enrich TBBPA-degrading bacteria. Aliquots of 100 µL of the final enrichment were spread onto MSM-agar plates containing 100 mg/mL TBBPA. All agar plates were incubated at 30°C. Colonies with distinct morphology were picked out and streak plate method was used to get pure strains. Pure strains were inoculated to standard lysogeny broth (LB) medium, and were preserved in a mixture of glycerol (50%, in pure water) and LB medium (1:1, V/V).

The isolated strains were identified by morphology and 16S rRNA gene analysis. Cell morphologies were observed by optic microscopes (normal and phase-contrast microscope). 16S rRNA gene sequences were amplified by PCR using the forward primer 27F (AGAGTTTGATCCTGGCTCAG) and the reverse primer 1492R (TACGGCTACCTTGTTACGACTT). The PCR reaction conditions and product purification were described by Li et al. (2014b). The obtained 16S rRNA gene sequences were annotated in GenBank database through NCBI Blast for taxonomic classification.

1.4. Cell suspension

The bacterial strain was inoculated into LB medium and TBBPA degradation was induced by adding 100 mg/L TBBPA into the medium. The bacteria were cultivated in fresh LB medium at 30°C for 17.5 hr until the end of the exponential growth phase to harvest cell suspension. Cultures were then centrifuged repeatedly at 4000 r/min and washed with MSM (pH 7.2). The washing process was repeated three times and the cells were resuspended in MSM (OD_{600} =6.6) for further experiments.

1.5. Incubation of TBBPA in cell suspension

Incubation experiments were performed in 50-mL glass bottles. Bacterial cells were incubated in MSM containing 2.0 mg/L ^{14}C -TBBPA (8333 Bq/bottle) at 30°C with shaking (130 rpm) in the dark. Extra yeast extract (100 mg/L) and casamino acids (100 mg/L) were added to each bottle to facilitate the bacterial growth and biodegradation as determined in pre-experiments. HA (1.0 mg/mL), Cu (10 μ mol/L CuCl₂), or both were added to the culture to study their effects on the degradation of TBBPA. Treatments with sterilized cells (autoclaved for 20 min at 121°C) were set as controls. CO₂ generated in the bottle was trapped in 1.0 mL of NaOH (1 mol/L) and radioactivity was periodically determined by liquid scintillation counter (LSC, see below). All experiments were performed in triplicate.

At 0, 8, 15, 24, 31, 41, and 87 days, individual samples were taken out and acidified with 6 mol/L HCl to adjust pH to 2 to release the dissolved CO_2 from the medium. The medium (containing cells) was extracted four times with ethyl acetate (10 mL each time) and repeated sonication (1 kW, 25 kHz, 5 min), vigorous vortexing (4 min), and centrifuging (2000 r/min, 5 min). The supernatants from each step of extraction were combined and the radioactivity was determined by LSC (see below). The extracts were then evaporated to dryness on a rotary evaporator, and re-dissolved in 400 μL methanol. The concentrated extracts were analyzed by high-performance liquid chromatography coupled to a radio flow scintillation detector (HPLC-LSC) (see below). The cultures after organic extraction were centrifuged at 5500 r/min for 10 min. The radioactivity in the cell free supernatants was determined by LSC (see below) and represented the radioactivity in hydrophilic non-extractable residues of TBBPA. The pellets were resuspended in 3 mL MSM and used for determination of radioactivity by LSC (see below).

The flasks of the treatments with HA addition, the medium was firstly extracted with ethyl acetate and then adjusted to pH 12 with 4 mol/L NaOH. The medium was centrifuged at 5500 r/min for 10 min to separate cells. The supernatant, containing HA, was acidified with 6 mol/L HCl to pH 1 and centrifuged again at 5500 r/min for 20 min. The radioactivity retained in the supernatant was determined by LSC (see

below) and represented the radioactivity in hydrophilic non-extractable residues of TBBPA. The pellet HA was re-dissolved in 3 mL NaOH (0.1 mol/L); the radioactivity was determined by LSC (see below) and represented the radioactivity of HA-bound residues derived from TBBPA.

1.6. Determination of radioactivity

Radioactivity in all liquid samples was quantified on a liquid scintillation counter (LS6500; Beckman Coulter, USA). For the radioactivity of 14 CO₂, 1 mL NaOH-trap was mixed with 3 mL scintillation cocktail (Gold Star multipurpose; Meridian Biotechnologies Ltd., UK). The radioactivity of organic extracts was determined by adding 3 mL scintillation cocktail to 1 mL extract. For determining the radioactivity of hydrophilic non-extractable fraction (10 mL out of total fraction), cells and HA (3 mL), the volume of scintillation cocktail was 10 mL.

1.7. HPLC-LSC analysis

HPLC-LSC analysis was performed with an Eclipse XDB-C18 column (250 mm × 4.6 mm, 5 μ m; Agilent Technology, USA) at 30°C on an Agilent HPLC 1200 series device equipped with an online radio flow detector (FlowStar LB 513; Berthold Technologies, Germany). The mobile phase contained water (22%) and methanol (78%), running isocratically at 1 mL/min for 20 min. The flow rate of the scintillation cocktail (Gold Flow multipurpose; Meridian Biotechnologies Ltd., Epsom, UK) was controlled by a pump (Liquid Scintillator Pump LB 5036; Berthold Technologies, Germany) and was 1 mL/min.

1.8. Data analysis

F-test and ANOVA were used to analyze individual and mutual treatment effects of Cu or HA. A statistical probability of p < 0.05 was considered significant.

2. Results and discussion

2.1. Identification of TBBPA-degrading strain

One bacterial strain was isolated from the sludge samples. The strain was named as strain CDT and was used in further study on degradation of TBBPA. The 16S rDNA sequence analysis revealed that strain CDT was most related to *Pseudomonas pseudoalcaligenes* strain W-20 in NCBI Genbank with a similarity of 97.12%. Based on these factors, strain CDT was classified into the genus *Pseudomonas* and was named as *Pseudomonas* sp. strain CDT. Similar to previously found TBBPA-degrading microbes, like the genera *Ochrobactrum, Comamonas, Achromobacter,* and *Sphingomonas* (An et al., 2011; Li et al., 2012b; Peng et al., 2012), *Pseudomonas* sp. strain CDT also belongs to the phylum Proteobacteria and is gram-negative.

2.2. Dissipation of TBBPA

At the end of 87-day incubation, $47.3\% \pm 6.0\%$ of the initial TBBPA was recovered from the culture of *Pseudomonas* sp. strain CDT alone (Fig. 1), which was similar to the results of Li

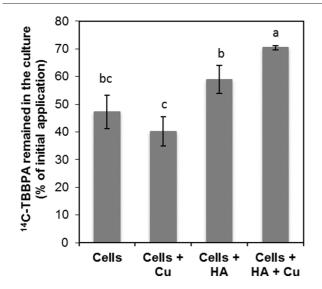


Fig. 1 – TBBPA remaining in the culture of Pseudomonas sp. strain CDT at the end of incubation with active cells and in the presence of Cu^{2+} (10 μ mol/L CuCl₂), HA (1000 mg/L), as well as with both Cu^{2+} and HA (concentrations were the same as above). Data are mean values of three individual experiments. Error bars indicate standard deviation. Different letters indicate significant difference among treatments as tested with Tukey's test.

et al. (2016) who inoculated the TBBPA-degrading bacterium Ochrobactrum sp. T into sediments. In the presence of Cu²⁺, the amount of recovered TBBPA was 40.2% ± 5.2%, which was not significantly different from treatment with active cells. In the presence of HA, the remaining TBBPA in the culture significantly increased to 58.9% \pm 5.0%, indicating that HA inhibited the initial degradation step of TBBPA by strain CDT in the culture. This is likely attributed to the adsorption and binding of TBBPA on HA, which decreased the bioavailability of TBBPA to the bacteria. This inhibitory effect of HA contrasted with stimulation of the dissipation of nonylphenol (Li et al., 2007) and phenanthrene (Ortega-Calvo and Saiz-Jimenez, 1998) by HA in bacterial culture. Compared to TBBPA, nonylphenol and phenanthrene are more volatile, thus adsorption to HA in the culture increased the chance of pollutants to stay in the culture. Pollutants could then partially desorb from HA to become bioavailable, thus their degradation by bacteria could be increased by the presence of HA (Li et al., 2007; Ortega-Calvo and Saiz-Jimenez, 1998). On contrary, TBBPA has a much lower vapor pressure (1.19 \times 10⁻⁵ Pa at 20°C) (Liu et al., 2016), which prevents TBBPA to volatilize from the medium, thus adsorption of TBBPA onto HA in the culture may decrease the bioavailability of TBBPA and further suppressed the initial degradation of TBBPA.

When both Cu^{2+} and HA were present in the culture, the inhibition of TBBPA dissipation was higher (70.4% ± 0.7% of initial ¹⁴C-TBBPA recovered) than in the presence of HA alone (Fig. 1), indicating that Cu^{2+} increased the inhibition effect of HA on the dissipation of TBBPA in the culture. This may be explained by the increased adsorption of TBBPA on HA in the presence of Cu^{2+} , which decreased the concentration of free TBBPA available to bacteria. Zhao et al. (2013) found that Cu^{2+} significantly increased tetracycline adsorption on soils, because of the formation of [tetracycline-Cu]⁺ complexation that strongly adsorbs on soils. TBBPA has two hydroxyl groups with pK_{a1} and pK_{a2} being 7.5 and 8.5 (Kuramochi et al., 2008). At pH 7.2 of the medium, TBBPA existed mostly in neutral species, so forming [Cu-TBBPA]⁺ to promote the adsorption of TBBPA on HA played a less important role in our study. But Cu²⁺ as a multivalent cation can function as an ion bridge between soil organic matter and organic pollutants (MacKay and Canterbury, 2005), which also increased adsorption of TBBPA on HA.

2.3. Mineralization of TBBPA

At the end of incubation in active culture, only $(2.3 \pm 0.1)\%$ of the initial ¹⁴C-TBBPA was mineralized to ¹⁴CO₂ (Fig. 2), while no $^{14}\mathrm{CO}_2$ was detected in the sterilized culture, indicating that the mineralization was resulted from the complete degradation of ¹⁴C-TBBPA by strain CDT. This much lower mineralization rate than the dissipation rate of TBBPA (52.7% \pm 0.6%) (Fig. 1) indicated that compared to TBBPA parent compounds, strain CDT are less likely to degrade TBBPA metabolites. To increase the mineralization efficiency of TBBPA, different bacterial strains capable of degrading TBBPA metabolites, like 2,4-dibromophenol (2,4-DBP), 2,4,6-tribromophenol (TBP), and bisphenol A (BPA), could be added. Previous studies have found that through cooperation of many microorganisms, the degradation of complex organic compounds could be accelerated when their population sizes reached to a certain level and the function and structure of microbial consortia were stable (Cho et al., 2002; Li et al., 2016; Wongwilaiwalin et al., 2010). In the presence of Cu²⁺, the mineralization of TBBPA at the end of the incubation became significantly lower (1.2% \pm 0.2%) (Fig. 2). The inhibition of Cu²⁺ on mineralization of TBBPA was possibly due to the toxic effects of Cu²⁺ on microbial metabolism and related enzyme activity, because Cu2+ can suppress the activity of fermentative, aerobic glucose-degrading heterotrophic, and nitrifying bacteria (Ochoa-Herrera et al., 2011), as well as the removal efficiency of ammonia and organic constituents in bacterial culture (Chen et al., 2013, 2014; Nam et al., 2015) possibly by chelating sulfhydryl groups on proteins.

HA significantly inhibited the mineralization of TBBPA (only $0.7\% \pm 0.1\%$ ¹⁴CO₂ generated at the end of incubation) (Fig. 2), additionally to its inhibitory effect on TBBPA dissipation in the culture (Fig. 1). However, when both Cu²⁺ and HA were present, $0.6\% \pm 0.1\%$ of TBBPA was mineralized at the end of incubation (Fig. 2), which was almost the same as that in the presence of HA alone. This differed from the synergistic inhibitory effects of Cu²⁺ and HA on TBBPA dissipation (Fig. 1) and suggested that the interactions between Cu²⁺ and HA and between Cu²⁺ and TBBPA discussed above influenced more likely the initial degradation/dissipation of TBBPA than the mineralization (complete degradation) of TBBPA by strain CDT, possibly due to the adsorption of Cu²⁺ on HA, which decreased the effective concentration of Cu²⁺ in the medium.

2.4. Fate of TBBPA

During the incubation with strain CDT, TBBPA was taken up by cells, transformed to intermediates, including organic

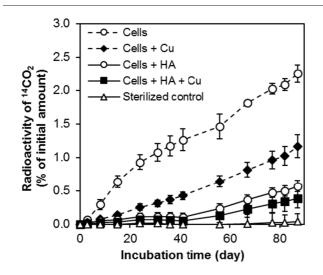


Fig. 2 – Radioactivity of ¹⁴CO₂ during the incubation of ¹⁴C-TBBPA mineralized in active and sterile culture with or without the presence of Cu²⁺ (10 μ mol/L CuCl₂), HA (1 g/L) as well as with both Cu²⁺ and HA (concentrations were the same as above). Data are mean values of three individual experiments. Error bars indicate standard deviation.

extractable and non-extractable metabolites, and finally to a limited extent mineralized to CO_2 . The major fate of TBBPA during degradation is transformation into organic extractable metabolites, accounting for 52.4%–71.2% of all transformed ¹⁴C-TBBPA in all treatments with active cells (Fig. 3). The metabolites detected in the organic extracts during the incubation were all more polar (retention time 3–8 min) than the parent TBBPA (Fig. 4). The structures of the metabolites were not identified, due to the low chemical concentration of metabolites generated. Previous studies on aerobic degradation of TBBPA in pure bacterial culture, nitrifying activated sludge, and soil showed that polar metabolites with retention

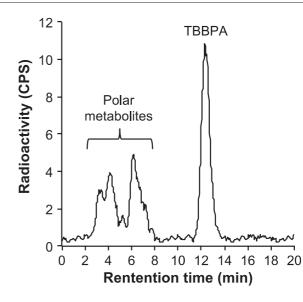


Fig. 4 – Typical radio-chromatogram of organic extracts after 87-day incubation of ¹⁴C-TBBPA in active culture of strain CDT. Polar metabolites (retention time: 3–8 min) of TBBPA (retention time 12.5 min) were detected.

times shorter than TBBPA were debromination products and compounds with a single benzene ring and polar groups, such as hydroxyl and carbonyl groups, which originated from cleavage and oxidation of the alkyl chain of TBBPA (An et al., 2011; Li et al., 2014a; 2015b; Peng et al., 2013; Sun et al., 2014; Zu et al., 2014). Continuous increase of bromide concentration in the culture of strain CDT (data not shown) proved that debromination occurred in the culture, while large amounts of radioactivity remained in the aqueous phase after organic solvent extraction (Fig. 3) indicated that hydrophilic metabolites were generated during the degradation of TBBPA by strain CDT.

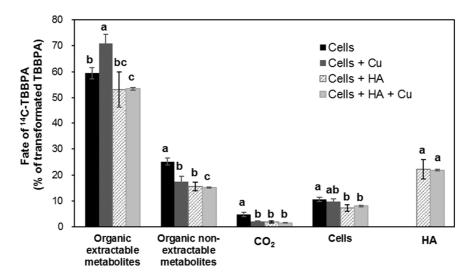


Fig. 3 – Relative radioactivity amounts of bio-transformed ¹⁴C-TBBPA recovered in the form of organic solvent-extractable metabolites, non-extractable metabolites (in water fraction), ¹⁴CO₂, cells, and associated to HA at the end of incubation of ¹⁴C-TBBPA in the culture of strain CDT. Data are mean values of three individual experiments. Error bars indicate standard deviation. Bars with the different letters indicate significant differences among treatments (cells, cells + Cu²⁺, cells + HA).

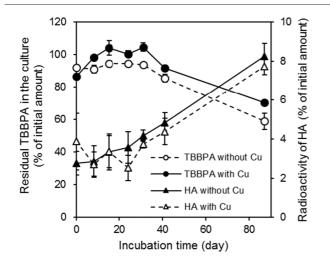


Fig. 5 – Amount of TBBPA in the culture (left y-axis) and radioactivity associated to HA (right y-axis) during incubation of ¹⁴C-TBBPA in active culture of strain CDT with or without Cu²⁺ (10 μ mol/L CuCl₂). Data are mean values of three individual experiments. Error bars indicate standard deviation.

In the presence of Cu^{2+} , the amount of organic solventextractable metabolites significantly increased, while the amount of non-extractable, viz. hydrophilic metabolites significantly decreased (Fig. 3). Considering that Cu^{2+} did not significantly affect TBBPA dissipation (Fig. 1), this indicated that Cu^{2+} probably inhibited the transformation of organic extractable metabolites to hydrophilic metabolites and finally to CO_2 .

When HAs were present in the culture, the amount of TBBPA derived hydrophilic metabolites, CO_2 , and incorporated into cells

significantly decreased, and large amounts of TBBPA-derived residues bound to HA (Fig. 3), indicating an alteration of the fate of TBBPA in the culture. At the end of incubation period, the radioactivity was distributed as follows: organic solventextractable metabolites (59.4% ± 2.2%), non-extractable hydrophilic metabolites (25.1% \pm 1.3%), CO₂ (4.8% \pm 0.8%), and bacterial cells (10.6% \pm 0.9%) (Fig. 3). The formation of HA-bound residues was owing to the reaction of reactive metabolites of TBBPA with HA macromolecules (Tong et al., 2015) and therefore increased continually during the incubation of TBBPA (Fig. 5). As bound residues per se have lower bioavailability (Gevao et al., 2000), formation of HA-bound residues significantly inhibited the further transformation of the metabolites to CO_2 (Li et al., 2015; Liu et al., 2013). The combined effects of Cu²⁺ and HA on the occurrence of metabolites in the culture were similar to that of HA (Fig. 3), suggesting that Cu²⁺ had minor effects on the transformation of TBBPA when HA were present in the culture.

2.5. Conclusions

In summary, TBBPA was degraded by strain CDT to organic-solvent extractable and non-extractable metabolites, and finally mineralized to a lesser extent to CO_2 (Fig. 6). Cu^{2+} significantly inhibited the mineralization and transformation from organic extractable metabolites into non-extractable metabolites, but had no significant effects on initial degradation (dissipation) of TBBPA. HA markedly altered the fate of TBBPA by reacting with reactive metabolites, forming large amounts of HA-bound residues, whereas the dissipation, metabolite formation, and mineralization of TBBPA were significantly inhibited. Coexistence of Cu^{2+} and HA significantly promoted the inhibitory effects of HA on initial

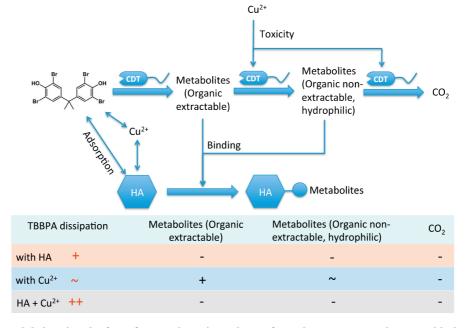


Fig. 6 – Conceptual model showing the fate of TBBPA in active culture of *Pseudomonas* sp. strain CDT with the influences of Cu²⁺ and humic acids (HA). During the incubation, TBBPA was taken up by cells, transformed to intermediates, including organic extractable and non-extractable metabolites, and finally mineralized to CO₂. Symbol "+", "–", and "~" indicate an increase, decrease, and no significant change in absolute (TBBPA parent compounds) or relative amounts (transformation products) of the compounds to the total transformation in the culture at the end of incubation.

degradation of TBBPA but had little or no effects on further transformation of TBBPA, owing to the complexation of Cu^{2+} and TBBPA, ion bridging function of Cu^{2+} , and neutralization of negative charge of HA by Cu^{2+} , which increased adsorption of TBBPA on HA thus decreased the bioavailability of TBBPA to strain CDT. In contrast, the presence of HA mitigated the effects of Cu^{2+} on transformation of TBBPA by strain CDT, owing to decrease of free Cu^{2+} concentration in the medium by complexation.

The results provide insights into individual and interactive effects of Cu²⁺ and HA on the biotransformation of TBBPA in pure culture systems and indicate the complexity of the interactive effects of heavy metals and soil organic matter on fate of organic pollutants in the environment. Pure culture provides a simple system for studying the effects of environmental factors on fate and biotransformation of organic pollutants and the underlying mechanisms, especially formation of bound residues in the environment, which is essential for risk assessment of organic pollutants in environments co-contaminated with heavy metals.

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