



Importance of Methane Oxidation for Microbial Degradation of the Herbicide Bentazone in Drinking Water Production

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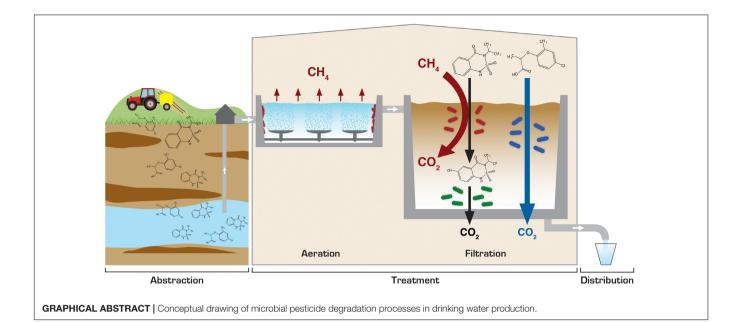
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Citation:

Hedegaard MJ, Schliemann-Haug MA, Milanovic N, Lee CO, Boe-Hansen R and Albrechtsen H-J (2020) Importance of Methane Oxidation for Microbial Degradation of the Herbicide Bentazone in Drinking Water Production. Front. Environ. Sci. 8:79. doi: 10.3389/fenvs.2020.00079 Bentazone is a herbicide, which is frequently detected in groundwater due to its mobility and persistence in aquifers. Groundwater is used as a drinking water source all over the world, and sustainable methods to remove pesticides at low concentrations are urgently needed since pesticide contaminations can adversely affect human health. The aim of this study was to investigate whether microbial bentazone degradation was associated with methane oxidation in full-scale drinking water treatment plants. To this end, we investigated bentazone biodegradation in microcosms with water and filter material from rapid sand filters, or biomass from aeration systems, and we investigated the statistical relation between the presence of methane and bentazone in groundwater abstraction wells. An array of evidence supported an association between bentazone degradation and methane oxidation in the biological treatment process. The biodegradation potential of bentazone was associated with the presence of methane in the raw water at 14 different water works. In contrast, no association was observed with any of the other investigated inorganic energy sources, e.g., ammonium. Addition of acetylene inhibited methane oxidation and the bentazone degradation in filter material from two investigated waterworks. Biomass from the aeration tanks degraded bentazone, but only while oxidizing methane. Bentazone removal rates and methane removal rates correlated significantly across all the experiments with biomass or filter material, with an overall transformation yield of 15 \pm 1 \times 10⁻⁵ mole_{BTZ}/mole_{CH4}. This demonstrated that the bentazone degradation was conducted by the same type of process in all the investigated communities, governed by methane oxidation. Furthermore, based on more than 10.000 water analyses from waterworks abstraction wells in Denmark, bentazone was detected significantly less frequent in wells with high methane concentrations (>1 mg/L) than in wells without methane. This suggests that biological treatment of bentazone contamination in drinking water may be achieved using methanotrophs.

Keywords: co-metabolic biodegradation, methane oxidation, pesticides, bentazone, rapid sand filters, drinking water treatment, groundwater



INTRODUCTION

Groundwater constitutes a major drinking water source worldwide (IWA, 2014). Unfortunately, pesticides are frequently detected in groundwater, especially those which have been extensively used, are highly mobile and persistent (Benner et al., 2013). In Denmark, traces of pesticides were detected in 40.8% of the groundwater abstraction wells (2018) (GEUS and Danish Ministry of Climate Energy and Utilities, 2019). In the European Union (EU), the guideline value for pesticides and their degradation products is 0.1 µg/L for a single compound in drinking water, or 0.5 μ g/L for the sum of all pesticides (European Commission, 1998). One of the most frequently detected pesticides in groundwater is the herbicide bentazone (Kolpin et al., 2000; Schipper et al., 2008; GEUS and Danish Ministry of Climate Energy and Utilities, 2019), and bentazone was found in 31.7% of 164 groundwater samples in EU, with the average concentration of 0.116 µg/L and maximum concentration of 10.55 µg/L (Loos et al., 2010).

Pesticides can be removed from drinking water by several processes such as activated carbon filtration, reverse osmosis or ozonation followed by biological activated carbon filtration (Camel and Bermond, 1998; van der Hoek et al., 1999). However, if pesticide contaminations can be managed through simple biological water treatment, with aeration of anaerobic groundwater followed by rapid sand filtration, it would be of large practical and commercial interest, since this treatment process is more environmentally sustainable (Godskesen et al., 2011).

Biological rapid sand filters are widely used to remove iron, manganese and ammonium in drinking water production (Mouchet, 1992; Lytle et al., 2007; Tekerlekopoulou et al., 2013; Lee et al., 2014) and may also remove and mineralize some organic trace contaminants, such as pharmaceuticals and pesticides (Zearley and Summers, 2012; Hedegaard and Albrechtsen, 2014; Vandermaesen et al., 2019). As an example, full-scale rapid sand filters at a groundwater-based waterworks removed the herbicide mecoprop (MCPP) to below detection limit (Hedegaard et al., 2014).

In many areas, groundwater contains methane. For instance, in Denmark the average methane concentration in groundwater is 0.67 mg/L, with a maximum concentration of 61 mg/L (Maksimavičius and Roslev, 2020), while in United States the average is 0.44 mg/L, and the concentration can be as high as 150 mg/L (Kulongoski and McMahon, 2019). Methane in the raw water might sustain growth of methanotrophs in the rapid sand filters, and so, methanotrophs are frequently found in biological rapid sand filters along with e.g., ammonium oxidizing bacteria (Albers et al., 2015; Gülay et al., 2016; Palomo et al., 2016; Maksimavičius and Roslev, 2020). Both methanotrophs and ammonium oxidizing bacteria are known to perform co-metabolic degradation (e.g., Colby et al., 1977; Forrez et al., 2009; Semrau et al., 2010; Xu et al., 2017), of some organic trace contaminants, which are degraded along with the primary substrate (e.g., methane, ammonium or organic compounds) (Dalton and Stirling, 1982; Alexander, 1994). This process has gained attention, since it allows degradation of trace contaminants even at concentrations that would not sustain specific degraders (µg/L) (e.g., Semprini et al., 1990; Semrau et al., 2010). Taking advantage of these co-metabolic properties of certain bacteria may pose a possible strategy for removing some organic trace contaminants using biological rapid sand filters (Benner et al., 2013; Papadopoulou et al., 2019).

Our previous work has shown how a methanotrophic enrichment culture co-metabolically transformed bentazone to 6-OH-, 8-OH-, isopropyl-OH-, and di-OH-bentazone which accumulated in this enrichment culture (Hedegaard et al., 2018). However, these hydroxylated transformation products could not be detected during bentazone degradation in filter material from a waterworks, whereas the presence of further degraded transformation products demonstrated the importance of other metabolic pathways in these systems (Hedegaard et al., 2019). Thus, it is still unknown whether the initial bentazone degradation step in filter material depends on methane oxidation or other biological processes and to what extent this is valid in water treatment system. In order to take advantage of bentazone degradation, it is essential to know to which extent this process depends on methane oxidation in water treatment.

The aim of this study was to investigate if bentazone degradation was associated with methane oxidation in biological drinking water treatment. To achieve this, we investigated:

- To which extent the bentazone biodegradation potential in filter material from 14 different waterworks was associated with the presence of any of the different inorganic energy sources for autotrophic microorganisms in the raw water.
- Whether inhibition of monooxygenases affected bentazone degradation in filter material and whether active methane oxidation was essential for bentazone degradation in biomass from the waterworks aeration system.
- Whether bentazone removal rates were associated with methane removal rates in biological material collected from waterworks.
- Whether methanotrophs could contribute to bentazone degradation in full-scale environmental systems, thus, whether presence of bentazone was associated with absence of methane in groundwater abstraction wells.

Experiments were conducted in microcosms with material and water from full-scale waterworks.

MATERIALS AND METHODS

Chemicals

Mineralization and removal at low concentrations $(1 \ \mu g/L)$ was investigated using [carbonyl-¹⁴C]-bentazone (Izotop, Institute of Isotopes Co., Ltd., Hungary). The radiochemical purity of [carbonyl-¹⁴C]-bentazone was 100% (chemical purity 99.77%) according to the manufacturer. Stock solutions were prepared in sterile MilliQ water, and the concentration was validated by HPLC-DAD immediately before the experiment. Bentazone (chemical purity 99.1%, Dr. Ehrenstorfer GmbH) was dissolved in sterile MilliQ water at least 1 day prior to the experiment. Acetylene was added from a gas flask (see **Supplementary Information**).

Waterworks

Across Denmark we selected 14 groundwater-based waterworks which represented different raw water quality (**Table 1**). At all the selected waterworks the water treatment included aeration, and filtration in primary and secondary rapid sand filters. In most waterworks, the water was aerated by stair aerators, but in some plants aeration tanks or intensive tray aerators (INKA chambers) were used to assure sufficient stripping of methane and/or hydrogen sulfide. Filter material was collected from the prefilter's top layer (10 cm) by an aluminum bucket on an extendable shaft, disinfected with 1% hypochloride. At two waterworks receiving methane-rich raw water, biomass samples were collected by scraping the sides of the aeration tanks with the aluminum bucket. Water was collected from the inlet to the clean water tanks.

Microcosms and Sampling

Bentazone degradation was investigated in batch experiments (for specific conditions in each experiment, e.g., number of replicates, controls, methane, acetylene, and bentazone concentrations, see Table 2). Either 100 g filter material or 5 mL biomass from aeration tanks was collected and transferred to 300 mL serum bottles along with 100 mL water within 2 days. The serum bottles had been acid-washed and heated to 555°C for 12 h. Controls with filter material were autoclaved three times (20 min, 1 bar and 121°C, the microcosms cooled for approx. 30 min inside the autoclave - to less than 80°C - before autoclaving was repeated). The serum bottles were sealed with autoclaved Teflon coated rubber stoppers and aluminum lids. In microcosms with methane present, 31 mL of the headspace was replaced by methane using a syringe with a needle through the rubber stopper. To mimic waterworks conditions microcosms were incubated in the dark at 10°C and placed on a shaker table (100-20 rpm). We have previously shown that methane oxidation was successfully inhibited by acetylene in a methanotrophic enrichment culture (Hedegaard et al., 2018). Therefore, in inhibited controls we added acetylene to a concentration of 26 mg/L in the water phase by replacing 8 mL of the headspace with acetylene using a syringe with a needle through the rubber stopper (Table 2).

Bentazone was added to the microcosms at time zero using a syringe with a needle through the rubber stopper. During the experiment water samples were collected by a syringe with a needle through the rubber stopper. All the collected water samples were replaced by an equivalent volume of sterile air. The collected water samples were filtered through a 0.22 μ m nylon GF filter (Frisenette Aps, Q-max[®] GPF Syringe Filters,

 TABLE 1 | Concentration of water quality parameters for investigated waterworks.

| ww | NH_4^+ | Fe | Mn | Cu | CH ₄ |
|------------------|-----------|------------|-------------|--------|-----------------|
| | (mg-N/L) | (mg/L) | (mg/L) | (μg/L) | (mg/L) |
| Brandsbjerg | 1.3–4.2 | 0.5–1.7 | 0.3–0.7 | - | 1.4 |
| Bregnerød | 0.2 | 2.5 | 0.28 | - | 0.068 |
| Brønderslev | 0.06-0.88 | 0.16-0.44 | 0.08–0.3 | - | 0.11 |
| Føllenslev | 0.53 | 3.5–4.0 | 0.21 | - | 0.14 |
| Gilleleje | 0.49 | 0.73 | 0.05 | <0.2 | 10.6 |
| Hareskov | 0.63 | 5.8 | 0.19 | - | 0.04 |
| Herning | 0.14-0.21 | 0.95–5.5 | 0.15-0.26 | - | 0.088 |
| Kerteminde | 0.01-0.07 | 0.013-0.1 | 0.002-0.008 | - | < 0.01 |
| Lønstrup | 0.02-0.05 | 0.04-0.06 | 0.01–0.6 | - | < 0.01 |
| Sjaelsø Plant I | 0.35 | 2.29 | 0.09 | <0.2 | 0.24 |
| Sjaelsø Plant II | 0.92-1.26 | 0.38–2.6 | 0.012-0.054 | - | 17.9 |
| Stenholt | >1.0 | 0.04 | 0.005 | <0.2 | 20.6 |
| Strib | 0.02-0.1 | 0.002-0.09 | 0.001-0.01 | - | < 0.01 |
| Ullerup | 0.87 | 0.12 | 0.004 | 1.96 | 14.9 |

 CH_4 is the maximum methane concentration in the raw water.

TABLE 2 | Experimental set-up.

| | Microcosms | Autoclaved controls | Inhibited controls | Methane added | 14C-bentazone | Bentazone (HPLC) |
|------------------|-------------------|------------------------------|----------------------------|----------------------------|----------------------------|----------------------------|
| | (number) | (number) | (number, concentration) | (number, concentration) | (number, concentration) | (number, concentration) |
| Screening of be | ntazone degradati | on potential at 14 different | t waterworks | | | |
| Brandsbjerg | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Bregnerød | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Brønderslev | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Føllenslev | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Gilleleje | 4 | 1 | - | - | 4 (0.5–1 µg/L) | 1 (1 mg/L) |
| Hareskov | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Herning | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Kerteminde | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Lønstrup | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Sjaelsø Plant I | 4 | 1 | - | - | 4 (0.5–1 µg/L) | 1 (1 mg/L) |
| Sjaelsø Plant II | 4 | 1 | - | - | 4 (0.5–1 µg/L) | 1 (1 mg/L) |
| Stenholt | 4 | 1 | - | - | 4 (0.5–1 µg/L) | 1 (1 mg/L) |
| Strib | 2 | 1 | - | - | 3 (0.5–1 µg/L) | - |
| Ullerup | 4 | 1 | - | - | 3 (0.5–1 µg/L) | 1 (1 mg/L) |
| Activity of mono | oxygenases | | | | | |
| Sjaelsø Plant II | 2 | - | 2 (26 mg/L) | 4 (5 mg/L) | - | 4 (1 mg/L) |
| Stenholt | 2 | - | 2 (26 mg/L) | 4 (5 mg/L) | - | 4 (1 mg/L) |
| Biomass from a | eration tanks | | | | | |
| Ullerup | 3 | - | - | 3 (5 mg/L)* | 3 (0.5–1 µg/L) | 2 (1 mg/L) |
| Stenholt | 3 | - | - | 3 (5 mg/L)* | 3 (0.5–1 µg/L) | 2 (1 mg/L) |

Left side of the table: Number of microcosms, autoclaved controls and inhibited controls with acetylene included in the experiments. Right side of the table: The number of the included microcosms to which methane, ¹⁴C-bentazone and/or bentazone was added and calculated initial concentration. *Methane added after 21 days.

diameter 25 mm). At the beginning of the experiments the oxygen concentration was >10 mg/L, and when concentrations decreased to below 5 mg/L during the experiments, pure oxygen was added. pH remained at 7.5–8 during the experiments.

Analytical Approaches Methane, Oxygen, ¹⁴C-Bentazone and HPLC-DAD Measurements

All below mentioned methods are described in details in Hedegaard et al. (2018). Methane concentration was measured every 1–3 days in all microcosm with methane, and based on these measurements the methane consumption was calculated. A sample of 50 μ L was collected from the headspace of the microcosms and analyzed for methane by GC-FID. Aqueous oxygen saturation was monitored by Oxygen-Sensitive Minisensors and a fiber optic oxygen meter (Fibox 3, Loligo Systems ApS). The concentration of ¹⁴C-bentazone in the water phase and the ¹⁴CO₂ produced from mineralization was quantified by a double vial system (Janniche et al., 2010). Water samples for bentazone analysis by HPLC-DAD were preserved with acetic acid and analyzed within few days.

Data From Groundwater Abstraction Wells

Data were collected from the public Danish "Jupiter database" which contains all historical data from drinking water wells

(accessed at the 2nd to the 5th of December 2016) (GEUS and Danish Ministry of Energy Utilities and Climate, 2016). We included data from waterworks wells where water samples for methane and bentazone-analysis were collected at the same time, and the data was divided into four groups depending on the presence/absence of methane and presence/absence of bentazone. A well was considered as "Containing methane" if at least one sample showed a methane concentration higher than 1 mg/L and/or "Containing bentazone" if the concentration of bentazone in a least one sample was above the detection limit (0.01 μ g/L).

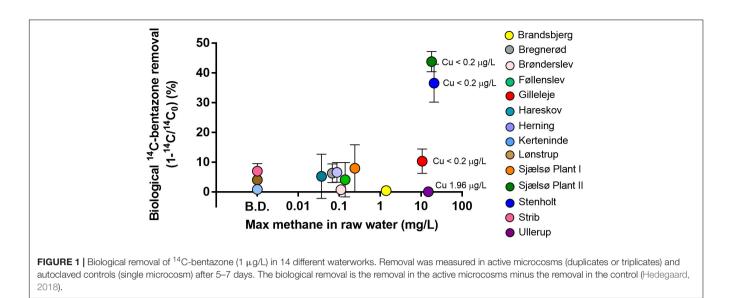
Statistics

The data were treated with the statistical software GraphPad Prism 6, performing: Correlation tests (Spearman), linear regression models, including tests for whether slopes and intercepts were significantly different (equivalent to ANCOVA), and Fisher's exact test.

RESULTS AND DISCUSSION

Methane in the Raw Water and Removal of Bentazone in Filter Material

Fourteen waterworks with various raw water methane concentrations ranging from below detection limit (<0.01 mg/L) to 21 mg/L were screened for a bentazone degradation potential



and its relation to raw water chemistry. The removal of bentazone was studied in microcosms mimicking the conditions in the sand filters (initial concentration at approx. 1 μ g/L), and the difference in removal of ¹⁴C-bentazone in autoclaved controls and biologically active microcosms expressed the biological removal potential in the different filter materials. In all 10 waterworks with less than 2 mg methane/L in the raw water the biological removal of bentazone was low (<8%) during 5-7 days of monitoring. The biological removal of bentazone did not lead to mineralization (no ¹⁴CO₂ was detected during the experiments) at any of these waterworks (Figure 1). At Ullerup waterworks the ¹⁴C-bentazone removal was insignificant and at Gilleleje waterworks it was low (10%) although methane was present (15 and 11 mg methane/L, respectively). However, at the two waterworks, with the highest methane concentrations (17-21 mg/L) (Sjaelsø Plant II and Stenholt), bentazone removal was substantially higher (37-44%) than at all other waterworks (Figure 1).

Rapid sand filters represent a complex microbial environment (Albers et al., 2015; Gülay et al., 2016), where several abiotic and biological processes occur simultaneously. Besides methanotrophs, other groups of bacteria in these filters have co-metabolic properties, e.g., both ammonium and manganese oxidizing bacteria are associated with degradation of trace organic contaminants (Forrez et al., 2009). The raw water chemistry is governing the composition of the microbial community in the rapid sand filters (Albers et al., 2015; Gülay et al., 2016), and the presence of methane in the raw water allows the presence of methanotrophs in the filter sand (Maksimavičius and Roslev, 2020). Therefore, we investigated for relations between the presence of inorganic energy sources for autotrophic microorganisms in the raw water (methane, ammonium, iron, and manganese) and the bentazone degradation potential. We divided the waterworks into four groups based on the presence of a compound in the raw water (for methane, ammonium and iron the limit was set to >1 mg/L, for manganese >0.1 mg/L) and whether the filter sand showed

a substantial bentazone degradation potential (>10%). All nine waterworks with low methane concentrations (<1 mg/L) also showed low bentazone degradation potential (<10%). Two waterworks had high methane concentrations (>1 mg/L) but did not show substantial bentazone degradation (<10%), while three waterworks with high methane concentrations (>1 mg/L) showed substantial bentazone degradation (>10%). The association between presence of methane in the raw water and the potential for biological degradation of bentazone in the filter sand was significant (Fisher's exact test, P = 0.0275). In contrast, we observed no association between bentazone degradation potential and presence of other inorganic energy sources such as ammonium, iron or manganese.

Filter material from most waterworks removed microbiologically a fraction of bentazone (up to 8%), independent of the methane concentration in the raw water. So a minor fraction of the bentazone degradation was not associated with methane oxidation (**Figure 1**).

The dynamics of the bentazone removal in filter material from five of the waterworks (Sjaelsø Plant I and II, Gilleleje, Ullerup and Stenholt) was studied at both high concentrations (1 mg/L bentazone measured by HPLC-DAD), and low concentrations (1 μ g/L, investigated as ¹⁴C-activity) for 14 days. In both approaches, bentazone removal in the filter material from Sjaelsø Plant II and Stenholt was much larger than in filter material from the remaining waterworks (**Supplementary Figure S1**). However, even though degradation clearly followed a linear trend through the entire period when measured by HPLC (**Supplementary Figure S1A**), the dissolved ¹⁴C-activity did not decrease substantially after 1 day at Sjaelsø Plant II, or after 7 days at Stenholt waterworks (**Supplementary Figure S1B**), probably due to formation of transformation products (Hedegaard et al., 2019).

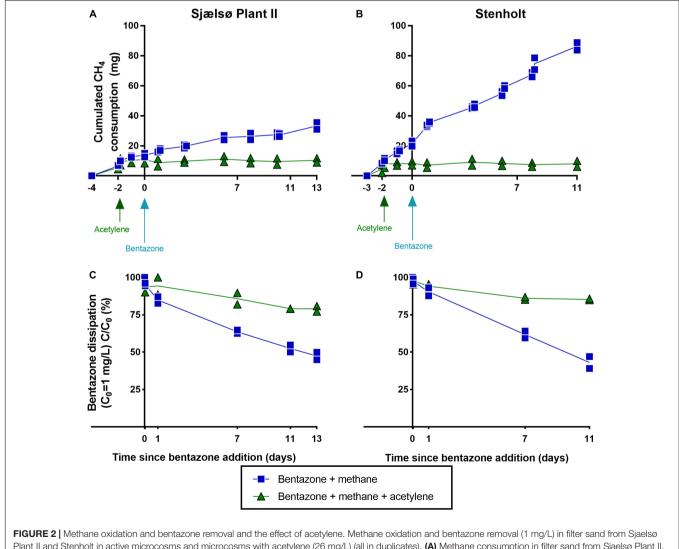
In spite of the substantial bentazone removal in filter material from Stenholt, degradation did not lead to ¹⁴CO₂-production, in contrast to the filter material from Sjaelsø Plant II (**Supplementary Figure S1C**). Hence, the microbial

community in the filter material from Stenholt only conducted the initial transformation of bentazone, while it was unable to mineralize the metabolites completely. In contrast, some of the initial bentazone was completely mineralized in filter material from Sjaelsø Plant II (**Supplementary Figures S1A,C**; Hedegaard et al., 2019).

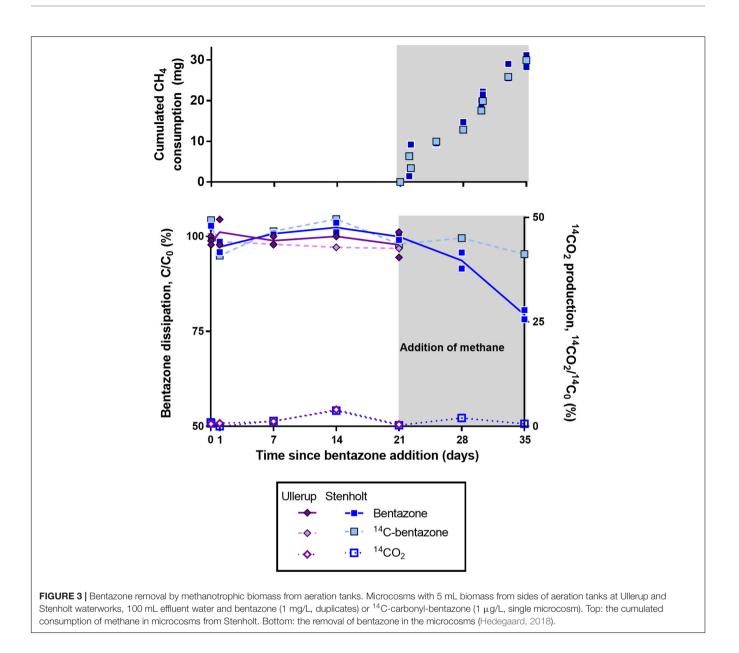
Inhibition of Methane Oxidation and the Effect on Bentazone Removal in Filter Material

Acetylene inhibits methane monooxygenases (MMO), as well as ammonium monooxygenases (AMO) (Prior and Dalton, 1985; Sullivan and Chase, 1996). To investigate whether the degradation of bentazone in filter material was conducted by monooxygenases, acetylene was added to microcosms with filter material from the two waterworks with the largest bentazone degradation potential (Sjaelsø Plant II and Stenholt). Before addition of acetylene, the microcosms from each waterworks all consumed methane at an equal rate (**Figures 2A,B**). Acetylene efficiently inhibited methane consumption immediately when added to the microcosms (at time: -2 days), while microcosms without acetylene addition continued to consume methane (**Figures 2A,B**), and thus microcosms with acetylene worked as a control without methane oxidation. Bentazone was added to all microcosms (at time 0 days), and in microcosms from both waterworks significantly less bentazone was removed in presence of acetylene than without (Sjaelsø Plant II: P = 0.00043; Stenholt: P < 0.0001) (**Figures 2C,D** and **Table 3**). Thus, the major fraction of bentazone removal in filter material was caused by the activity of monooxygenase.

However, a minor fraction of bentazone was removed in the presence of acetylene inhibiting monooxygenases, which suggests



Plant II and Stenholt in active microcosms and microcosms with acetylene (26 mg/L) (all in duplicates). (A) Methane consumption in filter sand from Sjaelsø Plant II. (B) Methane consumption in filter sand from Stenholt. (C) Bentazone removal in filter sand from Sjaelsø Plant II. (D) Bentazone removal in filter sand from Stenholt (Hedegaard, 2018).



other degradation pathways not associated with monooxygenases to contribute to the bentazone removal in filter material.

Bentazone Removal by Methanotrophic Biomass From Aeration Tanks

In aeration systems, some methane is stripped off, or biologically oxidized. Consequently, the methane concentrations are higher in the aeration systems than in the following rapid sand filters (Maksimavičius and Roslev, 2020). Therefore, we investigated if biomass scraped from the surfaces of the aeration systems could degrade bentazone, and whether this removal was associated with methane oxidation.

In absence of methane, biomass from the aeration tanks at Ullerup and Stenholt waterworks did not degrade bentazone within an incubation period of 21 days (Figure 3). After this period the experiment with biomass from Ullerup was stopped, while we added methane to microcosms with biomass from Stenholt waterworks. Methane consumption started immediately when methane was added to the microcosms (Figure 3 top), and the bentazone removal started simultaneously - the biomass removed 21% of the bentazone (initial concentration 1 mg/L) within the next 14 days (Figure 3). Subsequently, bentazone transformation only occurred during active methane oxidation. The biomass neither removed ¹⁴C-activity from the water phase nor produced ¹⁴CO₂, so the biomass apparently transformed bentazone into transformation products accumulating in the water phase. A similar transformation was observed with a methanotrophic enrichment from a rapid sand filter (Hedegaard et al., 2018). Bentazone transformation by biomass from the aeration tanks at Stenholt thus clearly depended on methane oxidation.

| | Acetylene | ē | Mass | Time (days) | r _{BTZ} (nmole _{BTZ} /h/microcosm) | rcH4 (µ mole _{CH4} /h/microcosm) | T _y ,BTZ/CH4 (mole _{BTZ} /mole _{CH4}) | Свт2/сн4 (mole _{вт2} /mole _{Сн4}) | SP (CH4/BTZ) | References |
|--------------------|------------|-----|-------|----------------|---|--|--|---|-----------------|------------------------|
| Enrichment culture | Exp. 1 | 1 | 20 g | - | $5.39 \pm 1.26 \ (r^2 = 0.65), \ n = 12$ | $39.4 \pm 4.70 \ (r^2 = 0.88), \ n = 12$ | 14×10^{-5} | 9.6×10^{-3} | 20 | Hedegaard et al., 2018 |
| from SPII | | (+) | 20 g | - | $2.90 \pm 1.08 \ (r^2 = 0.42), n = 12$ | $17.6 \pm 5.60 \ (r^2 = 0.58), n = 8$ | 17×10^{-5} | 9.6×10^{-3} | 58 | |
| | Exp. 2 | I | 10 g | 15 | $0.37 \pm 0.05 \ (r^2 = 0.77), n = 21$ | $6.11 \pm 0.47 \ (r^2 = 0.81), n = 42$ | 6.1×10^{-5} | 9.6×10^{-3} | 158 | Hedegaard et al., 2018 |
| | | + | 10 g | 15 | $0.04 \pm 0.04 \ (r^2 = 0.06), \ n = 20$ | ND ($r^2 = 0.12$), $n = 45$ | NC | | | |
| Filter sand | SPII | Ι | 100 g | 13 | $0.66 \pm 0.04 \ (r^2 = 0.96), \ n = 12$ | $4.35 \pm 0.27 \ (r^2 = 0.91), \ n = 28$ | 9.4×10^{-5} | 13×10^{-3} | 142 | Present study |
| | | + | 100 g | 13 | $0.25 \pm 0.04 \ (r^2 = 0.79), n = 12$ | $0.82 \pm 0.35 \ (r^2 = 0.21), \ n = 22$ | NC | | | |
| | | | | | P = 0.00043 | P < 0.0001 | | | | |
| | Stenholt | I | 100 g | 11 | $0.87 \pm 0.04 \ (r^2 = 0.98), n = 10$ | $15.6 \pm 0.60 \ (r^2 = 0.98), n = 30$ | 4.2×10^{-5} | 13×10^{-3} | 319 | Present study |
| | | + | 100 g | 11 | $0.22 \pm 0.03 \ (r^2 = 0.85), \ n = 10$ | $0.71 \pm 0.39 \ (r^2 = 0.15), \ n = 20$ | NC | | | |
| | | | | | P < 0.0001 | P < 0.0001 | | | | |
| Aeration tank | Stenholt – | Ι | 5 mL | 11 | $0.26 \pm 0.04 \ (r^2 = 0.92), n = 6$ | $5.5 \pm 0.3 \ (r^2 = 0.97), n = 18$ | 4.6×10^{-5} | 13×10^{-3} | 288 | Present study |

| Transformation | Yield of | Bentazone | as a |
|-----------------------|----------|-----------|------|

Bentazone Degradation Related to Methane Oxidation

Function of Methane Oxidation Transformation yields express moles of degraded trace contaminant per mole of methane consumed (Anderson and Mccarty, 1997). With data from all experiments in our present and previous studies, including filter material from two different waterworks, biomass from an aeration tank and a methanotrophic enrichment culture (Hedegaard et al., 2018), we investigated for an association between bentazone degradation rate and rate of methane removal (estimated by linear regression models, data in Table 3) across all the experiments. The bentazone removal rate correlated significantly with the methane removal rate across the six experiments (P = 0.0583) (Figure 4), resulting in an overall transformation yield, $T_{y,BTZ/CH4}$, of $15 \pm 1 \times 10^{-5}$ mole_{BTZ}/mole_{CH4} (Table 3). Hence, all the investigated methanotrophic microbial communities transformed similar amounts of bentazone molecules per oxidized methane molecules, and this process did not depend on the origin of microbial community, such as where in the treatment process it was collected. The minor variation (up to a factor of four) in transformation yields (Table 3), could be due to differences in the composition of methanotrophic community, e.g., the number of soluble methane monooxygenase (sMMO) versus particulate methane monooxygenase (pMMO) expressing methanotrophs (Semrau et al., 2010; Jagadevan and Semrau, 2013). This indicated, that methane oxidation governed the bentazone removal in the different materials from waterworks.

To compare bentazone removal in water treatment systems with co-metabolic degradation of other trace contaminants conducted by MMO, the normalized substrate preferences were calculated [as suggested by Hedegaard et al. (2018)]. The normalized substrate preferences ranged from 142 to 319 in the present experiments, and was thus within the range reported for methanotrophic co-metabolic degradation of chlorinated ethenes (3-400) (Anderson and Mccarty, 1997; Hedegaard et al., 2018). A similar magnitude in the preference of MMO for methane over other trace contaminants, indicated that the bentazone removal in water treatment systems was similar to co-metabolic degradation of other trace contaminants conducted by MMO (Hedegaard et al., 2018).

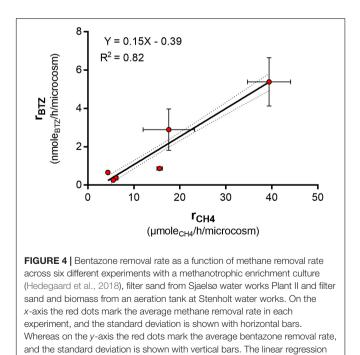
Association Between Methane and Bentazone in Groundwater Abstraction Wells

When aerobic rain water infiltrates from the soil surface toward methane-rich aquifers, fronts will be produced in the subsurface between the downward infiltrating aerobic water and the deeper, anaerobic, methane containing groundwater, which creates growth conditions for methanotrophs (Amaral and Knowles, 1995; Kotelnikova, 2002). Laboratory investigations with aquifer material has shown the recalcitrance of bentazone in aquifers (Albrechtsen et al., 2001), also in methanogenic groundwater systems (Albrechtsen et al., 2001; Luo et al., 2019). However, when bentazone contaminated water infiltrates toward a methane-rich aquifer, we suggest that methanotrophs in the methane and oxygen counter gradients can degrade

bentazone, which imply that presence of bentazone is less probable in anoxic methane-rich aquifers (Figure 5). To validate this assumption, we investigated whether bentazone was detected less frequently in methane-rich waterworks abstraction wells compared to wells without/small amounts of methane ($\leq 1 \text{ mg/L}$). The groundwater methane concentration is high (>1 mg/L) in 12% of the municipalities in Denmark (Maksimavičius and Roslev, 2020). Data from 15,437 wells showed that in 14,295 wells neither methane (>1 mg/L) nor bentazone was detected (Figure 5). 682 wells (4.4%) contained substantial amounts of methane (>1 mg/L), but no bentazone (<0.01 μ g/L), whereas bentazone was detected in 457 wells (>0.01 μ g/L) without methane (≤ 1 mg/L). Only three wells showed simultaneous presence of methane and bentazone (Figure 5). Thus, a twosided Fischer's exact test demonstrated a significant correlation between the two parameters (P < 0.0001), and bentazone was detected significantly less frequently in wells with methane (>1 mg/L) (Figure 5). These results indicate that methanotrophs may protect the methane-rich groundwater against bentazone contaminations. Unfortunately, it is not possible from the data to clarify how geological settings may influence the absence of bentazone in methane-rich groundwater. For instance, methane-rich groundwater could be associated with confined aquifers, where there is a better protection against pesticide contamination.

GENERAL DISCUSSION

The majority of bentazone degradation in the filter material was inhibited by acetylene, which suggests that bentazone biotransformation in the filter material is connected with the



model is shown as a full line with 95% confidence intervals (dotted lines).

activity of monooxygenases. Unfortunately, it was not possible to distinguish between the activity of AMO and MMO (Bédard and Knowles, 1989; Lontoh et al., 2000) since both can be inhibited by acetylene and may be present in rapid sand filters (Albers et al., 2015; Gülay et al., 2016). However, the biological bentazone degradation potential in filter materials from 14 waterworks was associated with the presence of methane in the raw water, but neither with the presence of manganese, iron nor ammonium. Additionally, in the biomass from the waterworks aeration tanks the bentazone removal only occurred simultaneously with methane oxidation. Hence, several lines of evidence supported that the majority of bentazone transformation was associated with methane oxidation in biological water treatment processes.

Surprisingly, in one filter material (Ullerup) bentazone was not biologically removed (Figure 1 and Supplementary Figure S1), despite high concentrations of methane in the raw water. This may be explained by the high copper concentrations (1.92 μ g/L) in the inlet to the filter, compared to the other waterworks ($<0.2 \mu g/L$) (Table 1). The copper concentration regulates the type of MMO being expressed: whether it is pMMO or sMMO (Semrau et al., 2010). sMMO is only expressed at low copper to biomass ratios (Semrau et al., 2013; Sirajuddin and Rosenzweig, 2015), and is the least specific of the two enzymes - capable of oxidizing aromatic compounds, while pMMO is limited to degrading alkanes of up to five carbon-atoms (Burrows et al., 1984; Trotsenko and Murrell, 2008; Semrau et al., 2010). Thus the 10 times higher copper concentrations at Ullerup waterworks than at the other waterworks could indicate a dominant formation of pMMO, which is suspected to be less efficient than sMMO to degrade bentazone.

Most waterworks had a minor potential for biological bentazone degradation, even in absence of methane in the raw water, however, this removal did not lead to mineralization of bentazone. Similarly, Vandermaesen et al. (2019) found that bentazone mineralization only rarely occurred in filter material in drinking water treatment plants. We also observed that some bentazone was removed in microcosms although methane oxidation was inhibited by acetylene. Thus, bentazone removal in filter material could not be ascribed to methanotrophic activity only. This conclusion is in agreement with our previous identification of a degradation pathway for bentazone in filter material. This degradation pathway is based on N-methylation and is independent of methane oxidation (Hedegaard et al., 2019).

CONCLUSION

These investigations showed:

• The potential for biological bentazone removal was associated with the presence of methane in the raw water at 14 different water works, while it was not associated with the presence of other potential inorganic energy sources, ammonium, iron, and manganese.

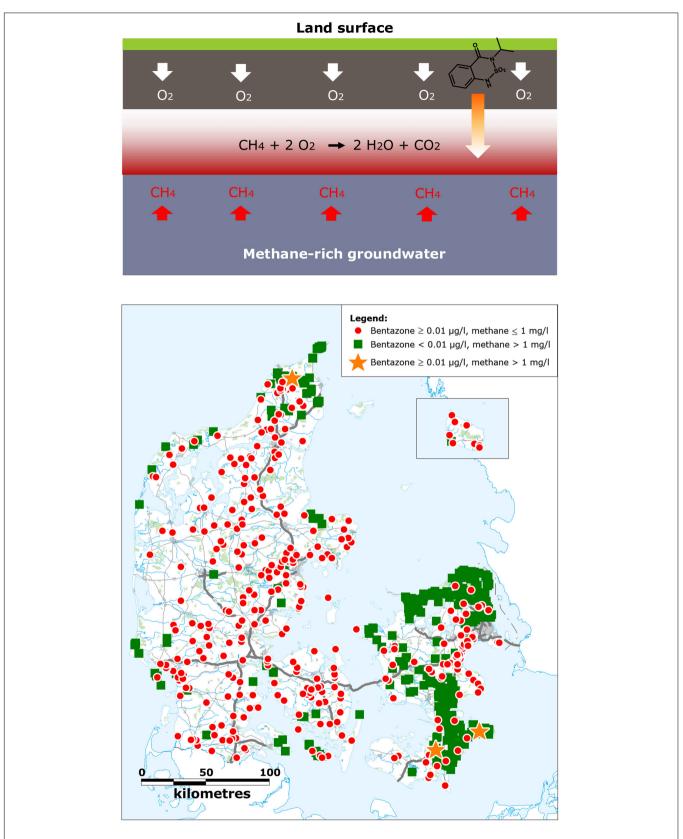


FIGURE 5 | Association between presence of methane (>1 mg/L) and bentazone in groundwater. Top: Conceptual drawing of how methane-rich groundwater can be protected from bentazone contaminations by methanotrophs in methane-oxygen counter gradients. Bottom: Presence of methane (>1 mg/L) and bentazone in active waterworks wells in Denmark.

- Acetylene inhibited *both* the methane oxidation *and* the bentazone removal in filter material from two waterworks. Thus, the majority of bentazone degradation was related to monooxygenase activity.
- Biomass collected from the aeration tanks at a waterworks receiving methane-rich groundwater, degraded bentazone, but only while oxidizing methane.
- The ratio between bentazone removal rates and methane removal rates, were similar across different systems including enrichment cultures (other studies), filter material from different waterworks and biomass from aeration tanks. with the overall transformation yield of 15 \pm 1 \times 10^{-5} mole_{BT7}/mole_{CH4}. Hence, all the investigated microbial communities transformed а similar bentazone amount of molecules per oxidized methane molecules.
- Bentazone was detected significantly less in groundwater abstraction wells with methane than in wells without methane (P < 0.0001), which indicated that methanotrophs living in methane and oxygen counter gradients, protected methane-rich groundwater from bentazone contaminations.

Thus, an array of evidence showed that bentazone degradation was connected to methane oxidation in drinking water treatment systems, indicating that MMO initiated the bentazone degradation. Therefore we suggest that bentazone degradation may be enhanced by stimulating the growth of methanotrophs in water treatment systems.

DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

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AUTHOR CONTRIBUTIONS

MH, RB-H, and H-JA had the idea for the study. MH, MS-H, NM, CL, and H-JA designed the study. MH, MS-H, NM, and CL carried out the experiments and analyzed the samples. Data from the Jupiter database was provided by RB-H. MS-H, NM, and CL did the initial data treatment of the experiments. MH did the analysis of data from Jupiter and overall data treatment, and wrote the manuscript. The major data interpretation was done by MH and H-JA, with contributions from MS-H, NM, RB-H, and CL. All authors commented on draft versions of the manuscript and provided feedback and ideas.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fenvs. 2020.00079/full#supplementary-material

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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