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3 **Chronic impact of tetracycline on the biodegradation of an** 4 **organic substrate mixture under anaerobic conditions**

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24 **Abstract**

25 The study evaluates the chronic impact of the antibiotic *tetracycline* on the biodegradation of
26 organic substrate under anaerobic conditions. The experiments involved an anaerobic

1 sequencing batch reactor fed with a synthetic substrate mixture including glucose, starch and
2 volatile fatty acids, and operated in a sequence of different phases with gradually increasing
3 tetracycline doses of 1.65 – 8.5 mg/L, for more than five months. *Tetracycline* exerted a
4 terminal/lethal effect at 8.5 mg/L on the microbial community under anaerobic conditions,
5 which caused the inhibition of substrate/COD utilization and biogas generation and leading to
6 a total collapse of the reactor. The microbial activity could not be recovered and re-started
7 within a period of more than 10 days, even after stopping *tetracycline* dosing. At lower doses,
8 substrate utilization was not affected but a reduction of 10-20% was observed in the
9 biogas/methane generation, suggesting that substrate utilization of *tetracycline* to the
10 biomass was limiting their bioavailability. During the experiments, *tetracycline* was partially
11 removed either through biodegradation or conversion into its by-products. The adverse long-
12 term impact was quite variable for fermenting heterotrophic and methanogenic fractions of
13 the microbial community based on changes inflicted on the composition of remaining/residual
14 organic substrate.

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17 **Keywords:** tetracycline; chronic inhibition; anaerobic biodegradation; methanogenesis; COD
18 removal

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1 1. INTRODUCTION

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3 Antibiotics, as one of the most important pharmaceutical group, have different usage areas
4 such as human and veterinary medicine, growth promoters in livestock, and agriculture.
5 Since these active compounds are not totally metabolized in human bodies and cannot be
6 eliminated completely in sewage treatment systems (Ternes *et al.*, 2004) they are found in
7 receiving water bodies. While antibiotic concentrations in raw domestic wastewater are
8 usually reported in the range from 100 ng/L to 6 µg/L (Giger *et al.*, 2003; Santos *et al.*, 2009)
9 their concentration in hospital and pharmaceutical industry effluents can reach 100 - 500
10 mg/L level (Kummerer, 2001, Larsson *et al.*, 2007), and an effective control and removal of
11 these compounds would provide greatly beneficial stability in domestic sewage treatment. As
12 antibiotics inhibit biological activities directly, they are likely to exert adverse/inhibitory effect
13 on the biodegradation of organic compounds in the wastewaters and this way, they
14 negatively affect the efficiency while by-passing conventional aerobic biological treatment
15 processes (Joss *et al.* 2006). Anaerobic treatment is an alternative for the removal of these
16 compounds in pharmaceutical industry waste streams because of high COD content and
17 persistent character (Oktem *et al.*, 2008).

18

19 Tetracycline (TET) is one of the most extensively used antibiotics in human activities (Figure
20 1). It is generally used for the treatment of respiratory tract infections and has a reversible
21 inhibitory effect. It is a broad-spectrum active compound, which inhibits bacterial protein
22 synthesis by binding the 30S ribosomal subunit to prevent the association of the aminoacyl-
23 tRNA to the ribosomal acceptor-A site (Chopra and Roberts, 2001). It causes structural
24 change in 16S rRNA (Loftin *et al.*, 2005). The behavior of this compound on sewage
25 treatment plants has been reported in the literature: It remains non-biodegradable, but it is
26 easily sorbed onto sewage sludge and therefore it is mostly discharged to the environment
27 through biosolids (Kummerer, 2001; Prado 2009). In another study, the authors found out
28 that tetracycline presented good adsorbability with 72 mg/g of the Langmuir maximum

1 adsorption capacity ($C_{s,max}$) (Prado *et al.*, 2009). The compound and its derivatives are
2 commonly used as promoter in animal growth, and therefore most of the studies about TET
3 degradation have focused on the anaerobic digestion of manure, which contains this
4 compound (Arikan *et al.*, 2006; Stone *et al.*, 2009; Wu *et al.*, 2011; Hu *et al.*, 2011). These
5 studies showed that TET in manure could be biodegraded in a range from 70% (Wu *et al.*,
6 2011) to more than 90% (Hu *et al.*, 2011) under anaerobic conditions. On the contrary,
7 Gartiser *et al.* (2007) determined TET as non-biodegradable under the anaerobic conditions
8 in the water matrix. On the other hand, limited information was found about the effect of TET
9 on anaerobic wastewater treatment systems: Arikan *et al.* (2008) reported a 30% inhibition in
10 methane production with 9.8 mg/L of TET dosing while the same level of inhibition was
11 observed with a much higher TET concentration of 28 mg/L in the study conducted by Stone
12 *et al.* (2009).

13

14 In order to evaluate the inhibitory impact of a selected compound in a biological system two
15 different experimental approaches are commonly applied: chronic and acute tests. The short-
16 term, acute tests usually involve a microbial population not previously exposed to the
17 inhibitor. Under anaerobic conditions, the methanogenic activity has been successfully
18 interpreted to yield the magnitude of observed inhibition induced by the tested chemical (Ince
19 *et al.*, 2009). Only a few studies have focused the removal and inhibition of antibiotics in
20 anaerobic systems, and some of them use the enzyme analogy for the evaluation of the
21 inhibitory action (Amin *et al.*, 2006; Fountalakis *et al.*, 2008). In a recent study, Cetecioglu *et al.*
22 (2012) evaluated the acute inhibition impact of three antibiotics including tetracycline on
23 the methanogenic activity of acclimated biomass fed with acetate. The significant effect was
24 mainly on process stoichiometry, preventing complete utilization of substrate removed in
25 metabolic reactions; almost complete methane inhibition was observed for antibiotic doses
26 above 500 mg/L. Although acute tests provide valuable information about inhibitory impact of
27 a contaminant, they only give a partial image of inhibition, while long-term chronic
28 experiments with continuous feeding of the inhibitor may indicate changes in the

1 biodegradation pattern accounting for adaptation and/or resistance of the microbial
2 community as argued by Kummerer (2004). Indeed, acute and chronic experiments
3 complement on another in providing information on the full response of the microbial
4 community in biological treatment systems under different conditions. The chronic
5 experiments are the indispensable part of the evaluation as they reflect the continuous
6 impact of lower antibiotic concentrations, similar to those encountered in full-scale treatment
7 systems.

8
9 In this context, the main objective of this study was to evaluate the chronic impact of
10 tetracycline on the biodegradation of a synthetic substrate under anaerobic conditions as well
11 as to evaluate degradation and distribution of tetracycline itself. For this purpose, a
12 sequencing batch reactor system operated with semi-continuous tetracycline feeding
13 throughout the experiments enabled to interpret the chronic inhibitory impact of the selected
14 antibiotic on process performance. Accordingly, both methane/biogas production together
15 with the biodegradation characteristics of both, tetracycline and the synthetic substrate
16 mixture (glucose, starch and volatile fatty acids) were used as the main evaluation
17 parameters. The results obtained from the chronic study were compared with those from
18 previous acute inhibition tests performed in similar experimental conditions (Cetecioglu, 2011
19 and Cetecioglu *et al.*, 2012).

20

21 **2. MATERIALS AND METHODS**

22

23 **2.1. The experimental approach**

24 The experiments were essentially designed for evaluating the chronic inhibitory impact of
25 tetracycline on the metabolic activities of a microbial culture sustained in a reactor operated
26 at steady state, under anaerobic conditions. An anaerobic sequencing batch reactor (ASBR)
27 was run in a daily “fill and draw” mode using a synthetic substrate mixture including volatile
28 fatty acids, glucose and starch. The operation of ASBR included a start-up period of around

1 150 days for acclimation and establishment of steady state conditions. Then, its performance
2 was observed during the next 154 days under steady state conditions, to make sure that
3 these conditions prevailed before semi-continuous exposure to TET dosing, *i.e.* chronic
4 impact of TET could be observed on an acclimated microbial community with a well-defined
5 culture history. A sequence of five different phases were included in the experimental
6 observation: During the first phase, *phase A*, (till day 77) ASBR was operated with feeding of
7 just the selected synthetic substrate without TET addition whereas during the following three
8 phases it was operated with semi-continuous feeding of the substrate/TET mixture: In *phase*
9 *B* (days 78-90), the daily TET dose was maintained at 1.65 mg/L; the antibiotic dose was
10 gradually increased to 5.7 mg/L in *phase C* (days 91-114) and to 8.5 mg/L in *phase D* (days
11 115-143). The TET dosing was stopped in the last phase (*phase E*) in order to observe a
12 possible recovery of the reactor performance during the next 10 days (days 144-154). A
13 second ASBR, which was operated in parallel for the entire period under identical conditions,
14 but without antibiotic dosing, served as control reactor. The sequence of different phases
15 was primarily designed to observe the tolerance and possible failure of the microbial
16 community under semi-continuous exposure to TET; this was the reason why TET
17 concentration was gradually increased once the expected microbial response was observed.
18 In fact, at the highest tested TET dosing, the observed response was the metabolic collapse
19 of the microbial culture, which did not recover after TET dosing was stopped. This approach
20 enabled to observe different responses of the system at selected/gradually increased TET
21 doses, which constituted the basis of the evaluation.

22

23 The evaluation of ASBR performance was mainly based on daily measurements of soluble
24 COD and volatile fatty acid (VFA) concentrations determined both in the influent and effluent
25 streams; they were accompanied with parallel daily measurements of biogas production and
26 composition assessing main fractions such as CH₄, CO₂ and H₂. Specific methanogenic
27 activity tests (SMA) were also conducted on biomass sustained under different TET feeding

1 regimes, for assessing the methanogenic activity of the acclimated microbial community
2 under inhibitory conditions.

3

4 **2.2. Operation of Anaerobic Sequencing Batch Reactor systems**

5 Two Anaerobic Sequencing Batch Reactors (ASBRs) with 1 L total volume were set-up and
6 operated at 35 °C under dark conditions to prevent photo degradation. The reactors were
7 operated with a 24-hour cycle consisting of fill (10 min), react (23 h), settle (45 min) and
8 decant (5 min). The reactors were mixed continuously using a magnetic stirrer at 90 rpm.
9 The systems were inoculated by an anaerobic sludge taken from the stock reactor treating a
10 synthetic substrate with a total COD of 4400 mg/L including the following ingredients: starch,
11 2090 mg COD/L; glucose, 1350 mg COD/L; sodium acetate, 240 mg COD/L; sodium
12 butyrate, 330 mg COD/L; sodium propionate, 490 mg COD/L. The MLVSS concentration of
13 the reactors was 4500 mg/L. The total COD of the synthetic substrate used for the reactors
14 was adjusted to 2250 mg/L; it was a similar mixture, mainly composed of starch and glucose:
15 starch, 1045 mg COD/L; glucose, 675 mg COD/L; which accounted for more than 76% of the
16 COD feeding; it also contained 120 mg/L of acetate, 165 mg/L of butyrate and 245 mg
17 COD/L of propionate, corresponding to the remaining 24% of daily COD loading. Trace
18 element solution which is adapted from a previous study (Amin *et al.*, 2006) as mg/L
19 ($\text{FeCl}_2 \cdot 4\text{H}_2\text{O}$, 2; $\text{CoCl}_2 \cdot 6\text{H}_2\text{O}$, 2; MnCl_2 , 0.32; CuCl_2 , 0.024; ZnCl_2 , 0.05; H_3BO_3 , 0.05;
20 $(\text{NH}_4)\text{Mo}_7\text{O}_{24} \cdot 4\text{H}_2\text{O}$, 0.09; Na_2SeO_3 , 0.068; $\text{NiCl}_2 \cdot 6\text{H}_2\text{O}$, 0.05; EDTA, 1; resazurine, 0.5; HCl
21 (36%) 0.001 mL), vitamins as mg/L (4-aminobenzoic acid, 0.04; D(+)-biotin, 0.01; nicotinic
22 acid, 0.1; calcium D(+)-pantothenate 0.05; pyridoxine dihydrochloride, 0.15; thiamine, 0.1 in
23 NaP buffer (10 mM, pH 7.1) and 0.05 mg/L B12) solution were added to the wastewater. The
24 pH of the reactors at the start of each cycle was observed to vary from 6.8 to 7.2 mainly due
25 to the alkalinity level of around 1000 mg/L CaCO_3 for sustaining the operation stability of the
26 anaerobic reactor.

27

1 The reactors were operated with an organic loading rate (OLR) 1.4 g/L.d for the first 10 days
2 of operation and then increased to 2.25 g/L.d in a stepwise manner. The solid retention time
3 was approximately 50 days throughout the study for both ASBRs and was calculated based
4 on VSS loss in the effluent and removed during sampling of the excess sludge. The hydraulic
5 retention time of the reactors was 2.8 days.

6
7 Temperature, pH and gas production were monitored daily in situ. Duplicate samples were
8 collected from the reactors for chemical and microbiological analysis.

9 10 **2.3. Specific Methanogenic Activity Test**

11 Methanogenic activity tests were performed using the pressure transducer technique
12 (Colleran *et al.*, 1992) to determine the chronic effect of TET on methanogenic pathway. The
13 pressure increase in sealed vials fed with non-gaseous substrates as acetate, propionate,
14 butyrate was monitored. The hand-held pressure transducer (Lutron PM-9107, U.S.A.) was
15 capable of measuring a pressure in a range of 5 to 7000 mbar, corresponding to 0.01 mol
16 biogas in 60 mL headspace. The biomass seed was adjusted to 2000 mg/L VSS, so that
17 each serum bottle was inoculated with 120 mg VSS at the start of operation in 60 mL active
18 volume. The sludge taken from TET fed ASBR at the end of each period to use an inoculum
19 in test bottles. The aim of this test was to compare the chronic effect of TET on the
20 methanogenic activity. Acetate and VFA mixture (acetate, butyrate, propionate)
21 concentrations in a range of 1000-5000 mg/L were initially tested in order to reach maximum
22 potential methane production (PMP) rate during the batch tests. Among those 4000 mg/L of
23 acetate concentration and 3000 mg/L of VFA concentration were found to be optimum. The
24 basal medium in the batch experiments was prepared based on OECD311 protocol under
25 strict anaerobic conditions (2006). During the 6-day test duration, the bottles were stored at
26 35 ± 2 °C and shaken daily by hand. Headspace pressure was measured every day by hand-
27 held pressure transducer.

28

1 **2.4. Analytical Methods**

2 Methane content in the biogas and VFA concentrations were measured using gas
3 chromatograph (Perichrom, France and Agilent Technologies 6890N, USA, respectively).
4 Suspended solids (SS), volatile suspended solids (VSS), total suspended solids (TS), total
5 volatile suspended solids (TVS) and soluble COD were determined according to Standard
6 Methods (APHA, 2005).

7

8 **2.5. Measurement of Tetracycline in water and sludge samples**

9 A mass balance could also be established for tetracycline through measurements in the
10 influent, effluent and biomass samples. For the sludge samples, 20 mg of freeze-dried
11 sludge was weighted in 15 mL centrifuge tube and 10 mL of the extraction buffer (5% (w/v)
12 sodium acetate, 100 mM EDTA in a methanol:water (1:1) solution adjusted to pH 8 with
13 sodium hydroxide) was added to each tube. The tube was sonicated for 15 min and then
14 centrifuged at 1370Xg at 25°C during 10 min. The supernatant was transferred to 60 mL
15 glass tube. The extraction protocol was performed 3 times for each sample and the obtained
16 supernatant was evaporated at 25°C under nitrogen stream to remove the organic solvent
17 and diluted with MilliQ water to 500 mL and filtered. Further sample clean-up was performed
18 by solid phase extraction (SPE) using OASIS HLB cartridges (6mL, 200 mg, Waters, USA).
19 Each cartridge was conditioned with 5 mL methanol followed by 5 mL HPLC grade water.
20 Sample was loaded into the cartridge at a rate of approximately 1 mL/min. The cartridge was
21 washed by 10 mL HPLC grade water and then dried by vacuum during 30 min. Sample was
22 then eluted by 6 mL of methanol. The extract was evaporated to less than 50 µL under
23 nitrogen streams and then reconstituted to 1 mL with 1:1 methanol:water mixture. Before
24 analysis, 10 ppb of chlorotetracycline as internal standard was added. The concentration of
25 TET in the samples was quantified by internal standard calibration curve, in order to correct
26 for possible matrix effects.

27

1 Wastewater samples were filtered and diluted by 1:50 according to their expected
2 concentration in the reactor, using the methanol:water mixture. Analysis of both, wastewater
3 and sludge extracts was performed by ultra-high-performance liquid chromatography coupled
4 to quadrupole-linear ion trap tandem mass spectrometry following the method developed by
5 Gros *et al.* (2012). The Waters Acquity Ultra-Performance™ liquid chromatograph system
6 was equipped with two binary pumps (Milford, MA, USA) and an Acquity BEH T3 column
7 (50mm x 2.1mm i.d., 1.7 µm particle size) was used for chromatographic separation.
8 Compounds were analyzed under positive ionization mode. The optimized separation
9 conditions were as follows: solvent (A) acetonitrile, solvent (B) water with 0.1% formic acid at
10 a flow rate of 0.5mL/min. The gradient elution was: initial conditions 5% A; 0–3 min, 70%A;
11 3.0–3.5 min, 100% A; 3.5–5.0 min, 100% A; from 5.0 to 5.1 return to initial conditions; 5.1 to
12 6, equilibration of the column. A sample volume of 5µL was injected in the UPLC instrument,
13 coupled to a 5500 QTRAP hybrid triple quadrupole-linear ion trap mass spectrometer
14 (Applied Biosystems, Foster City, CA, USA) with a turbo Ion Spray source. Tetracycline and
15 the corresponding internal standard were analyzed by positive ionization mode in Multiple
16 reaction monitoring (MRM) as indicated by Gros *et al.* (2012). Limit of detection (LOD) and
17 limit of quantification (LOQ) of the measurement were 0.58 and 1.94 ng/mL, respectively.
18 The recovery of the sludge sample was 117.2±29.0%.

19

20 **2.6. Statistical Analysis**

21 To determine the statistical significance of TET inhibition, COD removal efficiencies of the
22 ASBRs were compared using ONE WAY ANOVA test, which was followed by running a
23 Post-hoc Dunnett's test and student's T-test, respectively. Graphpad Prism 4 software was
24 used for all statistical analysis.

25

26 **3. RESULTS**

27

28 **3.1. COD removal**

1 Efficient COD removal was observed during *phase A* in the TET reactor: Soluble COD in the
2 effluent was reduced from an initial COD concentration of 2200 mg/L at the beginning of
3 each cycle to 73 ± 19 mg/L, corresponding to an efficiency higher than 96% (see Figure 2).
4 Similar COD removal could be maintained in the control reactor for the entire monitoring
5 period. It should be noted that the synthetic substrate is composed of organics compounds
6 that are all totally biodegradable in nature; based on similar studies conducted with these
7 compounds as single substrates or substrate mixtures, it would be acceptable to assume that
8 under the operation conditions selected for the reactors they would be totally removed so
9 that the low soluble COD level measured in the effluent is essentially residual soluble
10 microbial products generated in the course of biochemical reactions (Germirli Babuna *et al.*,
11 1998, Amin *et al.*, 2006).

12

13 Semi-continuous TET dosing of 1.65 mg/L in *phase B* and 5.7 mg/L in *phase C* did not seem
14 to exert a noticeable effect on the overall COD removal: As illustrated in Figure 2, the effluent
15 soluble COD basically maintained the same level as before, with an average level of 71 ± 28
16 mg/L and only slightly decreased to 57 ± 3 mg/L. However, TET dosing increased to 8.5
17 mg/L in the following operation phase (*phase D*) resulted in a significant upset in the reactor
18 performance: The soluble COD value in the effluent increased to more than 2000 mg/L
19 corresponding to overall COD reduction of only 9% after 134th day (Figure 2). At the end of
20 *phase D*, TET dosing was stopped in order to observe any possible recovery in the reactor
21 performance in the final *phase e*. However, the metabolic activity of the biomass could not be
22 re-activated to induce noticeable substrate utilization and the reactor operated was
23 terminated on day 154.

24

25 **3.2. Biogas generation**

26 Biogas generation is the inherent complement of COD removal under anaerobic conditions; it
27 is now regarded as the scientific yardstick for evaluating the magnitude of related metabolic
28 activities. During the initial ASBR operation without TET dosing in *phase a*, complete COD

1 removal was also accompanied with a biogas generation of 1046 ± 28 mL/day,
2 corresponding to an average biogas yield, Y_{BG} of 0.46 L/g COD removed. The generated
3 level remained quite stable throughout the phase and almost coincided with the level
4 monitored in the control reactor (Figure 3). The methane percentage in the biogas was
5 determined as $62.5 \pm 3\%$, indicating an average specific methane production yield, Y_{CH_4} of
6 0.32 L/g COD removed. This level is in conformity with the default value reported by
7 Tchobanoglous *et al.* (2003). Analysis of the biogas composition revealed that the other main
8 component of the biogas was CO_2 ($37.5 \pm 3.8\%$), with no detectable H_2 formation. Cetecioglu
9 (2011) mentioned that methane percentage was $58.0 \pm 1.7\%$ in the short-term operation up to
10 250 mg/L TET fed anaerobic system. The slight difference in the level of methane generation
11 is obviously due to different carbon sources utilized in the two studies, without noticeable
12 impact of TET inhibition.

13

14 In *phase B* involving semi-continuous TET dosing of 1.65 mg/L, biogas generation persisted
15 at a slightly lower level of 951 ± 12 mL/day, i.e. with a 10% decrease. A similar decrease
16 down to around 60% was also observed in the methane content, corresponding to a methane
17 production yield of 0.25 L/g COD removed. A further decrease in the biogas generation
18 started in day 91, the beginning of *phase C*, with the application of a higher TET dose of 5.7
19 mg/L: The daily biogas level was reduced to 864 ± 21 mL/day, 82% of the level in the control
20 reactor, while the methane content of the biogas remained approximately the same (58%). It
21 should be noted that *phases B* and *C* were characterized with complete COD removal as in
22 the early phase of the ASBR operation without the antibiotic addition (*phase A*). The
23 observed decrease in the biogas/methane generation despite full COD removal confirms
24 results obtained in the acute test with TET under anaerobic conditions, similarly preventing
25 complete utilization of substrate removed in the corresponding metabolic reactions
26 (Cetecioglu *et al.*, 2012). In the following *phase D* characterized by a higher semi-continuous
27 TET dose of 8.5 mg/L, the significant adverse effect of the reactor performance was also
28 observed for biogas generation, which dropped from 853 mL/day to 71 mL/day between days

1 115 and 143, tandem with a similar decrease in substrate removal. While the methane
2 generation exhibited a parallel decrease, the methane remained in the range of 0.2 L CH₄/g
3 COD removed (Figure 3b).

5 3.3. Effluent VFA composition

6 Monitoring of the presence and composition of the volatile fatty acids in the process effluent
7 provided additional information on the chronic impact of *tetracycline* on the metabolic
8 activities under anaerobic conditions. The analysis in the effluent covered, aside the three
9 VFAs in the influent, isobutyrate, isovalerate and n- valerate. VFAs were not detected in the
10 effluent until *phase D* i.e during the first 116 days of operation where the semi-continuous
11 TET dosing was started and gradually increased to 5.7 mg/L, confirming complete removal of
12 the available substrate. It also confirms the inhibitory impact of semi-continuous TET dosing
13 in the selected range of 1.65 – 5.7 mg/L, which partially blocked the utilization of the
14 substrate removed in the metabolic activities, as evidenced by the observed reduction in
15 biogas/methane generation. However TET dosing, when increased to 8.5 mg/L in *phase D*,
16 seriously impaired and inhibited propionic and acetic acid utilization pathways as shown in
17 Figure 4: The observations indicated that after the first day in *phase D* (day 116), acetate and
18 propionate accumulation in the system began and levels were measured as 27 and 28 mg/L,
19 respectively.

20 Acetic acid concentration in the effluent increased to 110 mg/L at the end of *Phase D* and
21 reached to 457 mg/L at the end of the operation, *Phase E*.

22 Propionic acid accumulation has also similar trend however, the concentration was higher
23 than acetic acid. Its concentration was measured as 750 mg/L at the end of *Phase D*.
24 However propionic acid concentration decreased in the *Phase D* and it was detected as 385
25 mg/L at the end of operation.

1 Also butyric and valeric acids were observed in the effluent of the TET reactor at *Phases D*
2 and *E*. While butyric acid concentration varied between 4 and 20 mg/L, valeric acid
3 concentration increased from 14 mg/L to 70 mg/L slowly until the end of operation.

4

5 **3.4. Fate of tetracycline during ASBR operation**

6

7 TET was measured in the effluent and in the biomass in order to ascertain its fate and
8 possible biodegradation in the anaerobic reactor in each phase of treatment. Measurement
9 indicated that TET concentrations in the effluent always remained significantly lower than the
10 corresponding influent doses as seen in Figure 5a: In phase b the effluent TET concentration
11 was 0.55, around one third of the influent level. In the following phase (*phase C*) the TET
12 value in the effluent was slightly lowered to 0.44 mg/L, while the influent dosing was
13 increased to 5.7 mg/L. When the influent TET concentration was increased to final level of
14 5.5 mg/L, the corresponding effluent level initially remained the same (0.47 mg/L), then it was
15 reduced down to 0.06 mg/L (*phase D_2*), to finally reach a higher value of 1.36 mg/L in
16 *phase D_3* as illustrated in Figure 5a.

17

18 One of the possible explanations for the observed discrepancy between TET influent and
19 effluent TET levels in the anaerobic reactor is physical removal by means of sorption onto
20 biomass. In fact, a number of similar studies on activated sludge systems reported sorption
21 as the dominant mechanism for the removal of antibiotics (Kim *et al.*, 2005; Prado, 2009). As
22 shown in Figure 5a, TET sorption onto biomass did not exhibit a ascending trend, i.e. a
23 continuous increase in the TET fraction in the sludge: This level was initially 0.17 mg/L in
24 *phase B*; it dropped down to an almost negligible level of 0.05 mg/L in the following phase
25 and then it increased to 1.78 mg/L with a gradual descent to 1.25 mg/L by the end of *phase*
26 *D*, when the influent TET dose was adjusted to 8.5 mg/L. In the following phase (*phase E*),

1 where the TET dosing was stopped, TET fraction in the biomass was desorbed and the
2 concentration was measured as 0.32 mg/L.

3

4 The results outlined above and displayed in Figure 5a cannot be directly used for mass
5 balance, which would indicate the extent of TET biodegradation, mainly because the TET
6 fraction sorbed onto sludge would accumulate the same way as biomass, leaving the reactor
7 only as part of the excess sludge. Therefore, the observed TET concentration in the sludge,
8 TET_S , should be corrected by a factor of (HRT/SRT) in order to obtain the effective TET
9 concentration in the biomass, TET_{SE} , and the corrected value incorporated in the mass
10 balance (Hocaoglu and Orhon, 2010):

11

$$12 \quad TET_{SE} = TET_S (HRT/SRT)$$

13

14 This expression allows calculating of the extent of TET degradation efficiency (TET deg_eff),
15 corrected for entrapment and accumulation in the biomass:

16

$$17 \quad TET \text{ deg_eff} = (TET_I - TET_E - TET_{SE}) / TET_I \times 100$$

18

19 where, TET_I = influent TET dose; TET_E = measured TET concentration in the effluent.

20

21 Efficiency of TET reduction in different phases of reactor operation, using the expression
22 defined above is illustrated in Figure 5b. It basically indicates a TET reduction pattern that
23 started with more than 50%, increased to more than 90% in phase c, and sustained around
24 40% at the end of *phase D*, where the metabolic activities and the COD removal efficiency
25 were practically stopped. The reduction profile may be attributed to total biodegradation of
26 TET under anaerobic conditions, a novel result not previously reported, or to its partial
27 biodegradation and conversion to its major by-products

1 3.5. Assessment of specific methanogenic activity

2 Assessment of the specific methanogenic activity of the biomass (SMA) in batch reactors has
3 been a useful experimental approach for the appraisal of adverse/inhibitory effects (Ince *et*
4 *al.*, 2009); this approach was previously adopted evaluate the acute impact of *tetracycline*
5 along with two other antibiotics on the biodegradation of acetate and VFA mixture under
6 anaerobic conditions (Cetecioglu, 2011 and Cetecioglu *et al.*, 2012). The SMA test is
7 designed differently for acute and chronic impacts: While the acute SMA test involves a
8 series of parallel batch reactors inoculated with the same (control) biomass and the selected
9 substrate but with increasing doses of the antibiotic, the chronic SMA test is run with biomass
10 seeding from different phases of the reactor operation under semi-continuous impact of the
11 antibiotic and fed with the same substrate dose.

12 In this study, the SMA test was similarly performed with biomass seeding taken from the end
13 of different operation phases of the TET reactor, namely from phases A, B, C, D and E; it
14 should be noted that semi-continuous TET dosing was adjusted to 1.65, 5.7 and 8.5 mg/L in
15 the first three phases and stopped in the last phase. The SMA test was run twice, the first
16 one with acetate and the second/parallel one with a VFA mixture – i.e. an acetate-butyrate-
17 propionate mixture.

18 In the first SMA test, all batch reactors, each with an effective volume of 60 mL, were started
19 with 4000mg/L of acetate as the sole carbon source so that the reactors all included the
20 same initial acetate dose of 4250 mg COD/L or around 255 mg acetate COD. The test was
21 run for 8 days (192 hours). The observed cumulative biogas production (CBP) and
22 cumulative methane production (CMP) profiles in the SMA test are given in Figure 6,
23 showing that each specific profile reached a different plateau after around 168 hours
24 depending on the operation phase which yielded the biomass seed. The CMP value of the
25 biomass representing the initial phase without TET dosing was determined as 77 mL,
26 corresponding to 0.30L/g COD, a value quite in agreement with the level associated with
27 semi-continuous operation. As shown in Figure 6, the CMP test detected a loss of activity in

1 the biomass taken from phase b (1.65 mg/L TET dosing), as the collected methane volume
2 was reduced down to 55 mL, corresponding to around 30% decrease. The CMP levels were
3 gradually reduced to 47mL (39% decrease) and finally to 23 mL (71% decrease) in reactors
4 seeded with biomass from phases *C* and *D*. However, a significant recovery of the
5 methanogenic activity was observed in connection with the last phase (*phase E*) where the
6 TET dosing was stopped, evidence with an increase in the corresponding CMP value from
7 23 mL to 42mL, i.e. 0.16L methane/g COD_{removed}. The methane content of the biogas was
8 also decreased gradually depending on TET concentration. While the methane percent was
9 65%, 60% and 58% at *Phases B, C* and *D*, respectively, the value increased to 63% again at
10 *Phase E*, in which TET addition was terminated.

11

12 In the second set of SMA tests, all batch reactors, each with an effective volume of 60 mL,
13 were fed acetate, butyrate and propionate mixture; as 3000 mg/L of each VFA. The initial
14 VFA dose for each reactor corresponds to 13080 mg COD/L. The CBP and CMP profiles
15 during 8 days are given in Figure 7 and each profile reached the specific plateau at around
16 7th day like acetate fed SMA test reactors. The CMP value of *Phase A* without TET dosing
17 was observed as 312 mL and this value is equivalent to 0.39L/g COD, which is quite higher.
18 As seen in Figure 7, CMP value of *Phase B* (1.65 mg/L TET dosing) decreased dramatically
19 to 93 mL, corresponding to around 0.12L/g COD. A 70% reduction in CMP value was
20 observed. The CMP values gradually decreased to 68 mL and 14 mL in *Phase C* and *D*,
21 respectively. A 75% recovery was also observed at the last phase (*Phase E*) in which TET
22 dosing was stopped and the CMP values reached to 52 mL. Differently from acetate fed SMA
23 test bottles, the methane content of VFA fed set was quite stable as 50%.

24

25 The results explained above compares two sets of batch SMA tests, one conducted with
26 acetate and the other with selected VFA mixture and show that the biogas methane
27 generation in the latter test conducted with the VFA mixture always remained clearly below

1 the corresponding levels obtained with acetate tests. This observation may be interpreted as
2 lack of available acetate for acetoclastic methanogens and therefore, failure of heterotrophic
3 fermenters to convert propionate and butyrate into acetate due to adverse impact of TET
4 dosing. This observation supports the findings related to the semi-continuously fed ASBR
5 that the chronic damage of TET dosing was more effective and finally lethal on heterotrophic
6 fermenters as compared with methanogens.

8 **4. DISCUSSION**

10 **4.1. Difference from acute impact**

11 The chronic impact of *tetracycline* on substrate biodegradation under anaerobic conditions
12 was severe and occurred in the range of 5.7 – 8.5 mg/L dosing level, a much lower level as
13 compared with the acute impact: In parallel tests, an initial TET dose of 50 mg/L was
14 observed as the threshold of a noticeable acute impact on anaerobic biodegradation; the
15 inhibitory effect of TET addition became detrimental when the initial dose was increased to
16 500 mg/L (Cetecioglu *et al.*, 2012). These results indicate that while short-term assays may
17 be useful in assessing the effect of inhibitors received as pulse discharges, they will not be
18 sufficient to reflect the real inhibition mechanism on microbial communities in the long range
19 (Kummerer, 2004).

21 **4.2. Terminal/lethal effect**

22 *Tetracycline* exerted a terminal/lethal effect at 8.5 mg/L on the microbial community under
23 anaerobic conditions, stopping substrate/COD utilization and biogas generation and leading
24 to total collapse of the reactor. The microbial activity could not be recovered and re-started
25 within a period of more than 10 days; even after the *tetracycline* dosing was stopped. From a
26 practical perspective, this level is obviously too high for domestic sewage, but quite relevant
27 for pharmaceutical plants, hospitals, *etc.*, where anaerobic treatment becomes appropriate,
28 due to high organic content of the effluents (Amin *et al.*, 2006, Larsson *et al.*, 2007).

1

2 **4.3. Nature of impact - different with variable doses**

3 The effect of *tetracycline* at the lower dose of 5.7 mg/L was quite different: In this phase of
4 the study (*phase B*) substrate utilization was not impaired and full COD removal was
5 achieved but around 20% reduction was observed in the biogas/methane generation. The
6 same mechanism with a lower decrease of biogas generation in the range of 10% was also
7 measured when the *tetracycline* dose was 1.65 mg/L in the previous phase. These results
8 confirm similar findings reported in the literature, where the *stoichiometric disturbance* – *i.e.*
9 substrate removed but partially utilized for methane generation - was attributed to the
10 blockage of certain enzymatic steps in related metabolic reactions (Fountoulakis *et al.*, 2008;
11 Cetecioglu *et al.*, 2012). In a simplified way, it may be interpreted as the substrate binding
12 effect of TET, in accordance with *uncompetitive inhibition* analogy.

13

14 **4.4. Accumulation/biodegradation**

15 Semi-continuous dosing resulted in the accumulation of *tetracycline* in the biomass with a
16 gradual increase to around 1.5 mg/L throughout the observation period as indicated in
17 previous studies conducted with erythromycin (Amin *et al.*, 2006) and tylosin (Shimada *et al.*,
18 2008). The observed *tetracycline* profile in the biomass suggested equilibrium established on
19 the basis of simultaneous adsorption/desorption mechanisms. The interesting/novel aspect
20 of the evaluation was that the major fraction (>80%) of the *tetracycline* introduced into the
21 anaerobic reactor could be fully or partially biodegraded along with the organic substrate.
22 Figure 5 shows an appreciable overall removal of TET even in *phase D*, where the COD
23 removal efficiency was significantly dropped. Since TET is a xenobiotic much more difficult to
24 degrade compared to the substrate mixture fed to the reactor, it is more likely that the main
25 mechanism for the observed TET removal in this study is formation of metabolites rather
26 than biodegradation. This issue deserves more emphasis in future studies on the subject.

27

28 **4.5. Microbial dynamics**

1 The experimental results were quite interesting from the viewpoint of substrate/biomass
2 interactions and microbial population dynamics. In other words, the experiments were
3 designed in such a way to yield the impact of *tetracycline* not only on the overall anaerobic
4 biodegradation efficiency, but also on changes inflicted on the nature of remaining/residual
5 organic substrate and on the activity of different significant components of the microbial
6 community. It should be remembered that the organic substrate mixture was mainly
7 composed of starch and glucose (76%), but it also included 120 mg COD/L of acetate, 245
8 mg COD/L of propionate and 165 mg COD/L of butyrate corresponding to a total VFA level of
9 530 mg/L, i.e. only 23% of the total COD level in the influent. Latest developments in the
10 modeling of anaerobic systems such as ADM, identify exactly similar changes and
11 conversions between substrate components and correlate them with groups of
12 microorganisms capable of performing these metabolic activities, without having to go into a
13 detailed molecular analysis (Batstone *et al.*, 2002). Adoption of a similar basis of evaluation
14 for the substrate mixture selected for the study would suggest that the mixed microbial
15 culture sustained at steady-state in the reactor would inherently include regular *anaerobic*
16 *heterotrophs* to hydrolyse starch into simple sugars; *fermenting heterotrophic*
17 *microorganisms* converting glucose/simple sugars mainly to acetate and propionate;
18 *propionate degraders*, utilizing propionate for the generation of acetate, and finally
19 methanogens processing mainly acetate for the production of biogas. Interpretation of the
20 results obtained enabled to visualize the impact of *tetracycline* on these metabolic activities
21 and therefore, on related components of the microbial community throughout the observation
22 period: (i) The initial impact of *tetracycline* when increased to 8.5 mg/L (*phase D*) was more
23 focused on *methanogens*, evidenced by the gradual increase of VFAs in the effluent; at the
24 end of *phase D*, the total VFA level was increased to around 820 mg COD/L, significantly
25 higher than the corresponding influent level. (ii) The inhibitory impact was specifically
26 pronounced for propionate degraders. (iii) A recovery of methanogenic functions was
27 observed after the *tetracycline* dosing was stopped, with gradual depletion of accumulated
28 VFAs, suggesting adaptation/resistance mechanisms in the corresponding fraction of the

1 biomass; this result was also in conformity with a similar recovery in the specific
2 methanogenic activity. (iv) The adverse effect of *tetracycline* dosing on regular/fermenting
3 heterotroph gradually increased and finally inflicted lethal and non-reversible damage,
4 stopping VFA generation and the overall COD removal. This interpretation was also fully
5 supported by a comprehensive microbial community analysis based on DNA and RNA based
6 molecular microbial techniques; the results of the molecular analysis will be reported in detail
7 as the following part of the study.

8

9 **5. Conclusions**

10

11 In the light of evaluations presented in the previous sections, the significant findings of the
12 study on the chronic impact of tetracycline may be outlined as follows:

13

14 The results suggested that the nature of the *adverse impact was quite variable* as a function
15 of the inhibitor dose: At low levels, available substrate was removed but only partially utilized
16 for biogas/methane generation, presumably due to the blockage of certain enzymatic steps in
17 related metabolic reactions; at higher doses, it induced total collapse of the microbial activity
18 and metabolic functions. For the selected conditions of the study the terminal dose for
19 tetracycline inhibition was 8.5 mg/L.

20

21 The effect on microbial dynamics was *selective*, exerting markedly different inhibitory impact
22 on various steps of substrate utilization and metabolic reactions associated with the activities
23 of the microbial community sustained in the reactor. *A cumulative impact* was observed for
24 the sequence of biochemical processes converting different substrate fractions into acetate,,
25 possibly affected by adsorption and progressive accumulation of *tetracycline* on the biomass.
26 However the adverse effect was quite different and *reversible* for the methane generation
27 process,, upsetting the utilization of available/generated VFAs at first and with subsequent
28 partial recovery of methanogenic activity when the inhibitor addition was stopped. This

1 aspect was further investigated by means of parallel molecular studies reflecting changes on
2 the composition and nature of the microbial community. Related results are quite
3 comprehensive and will not reported at this stage.

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13

ACCEPTED MANUSCRIPT

Highlights

- ❖ Chronic impact of tetracycline was lethal at 8.5 mg/L on the microbial community
- ❖ At lower doses, substrate removal was not impaired but biogas volume was reduced.
- ❖ Tetracycline was partially biodegraded.
- ❖ Impact was cumulative on fermenting heterotrophs due to TET adsorption/accumulation.
- ❖ Impact was reversible for methanogens with partial recovery of biogas generation.

Figure Captions

Figure 1. Chemical structure of Tetracycline

Figure 2. COD removal efficiency in the tetracycline reactor

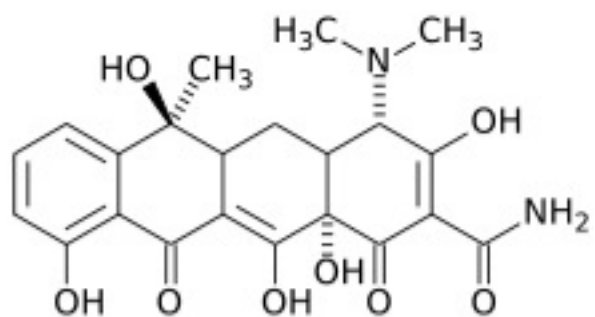
Figure 3. (a) Effect of tetracycline on biogas generation, (b) Stoichiometry of CO₂ and CH₄ generation

Figure 4. Fate of volatile fatty acids under the inhibitory impact of tetracycline

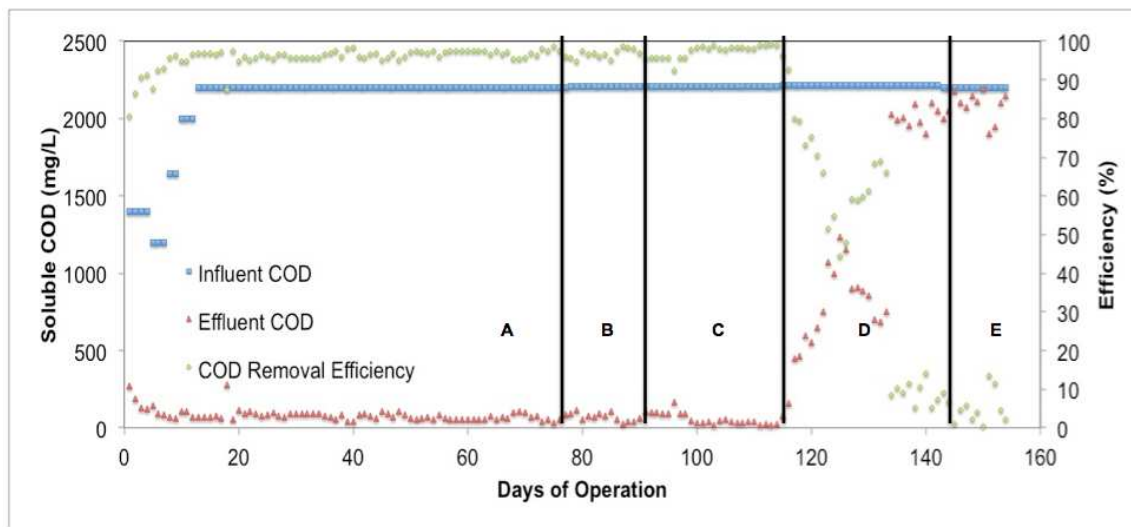
Figure 5. (a) Tetracycline concentration in liquid/solid phases; (b) Biodegradation profile of tetracycline

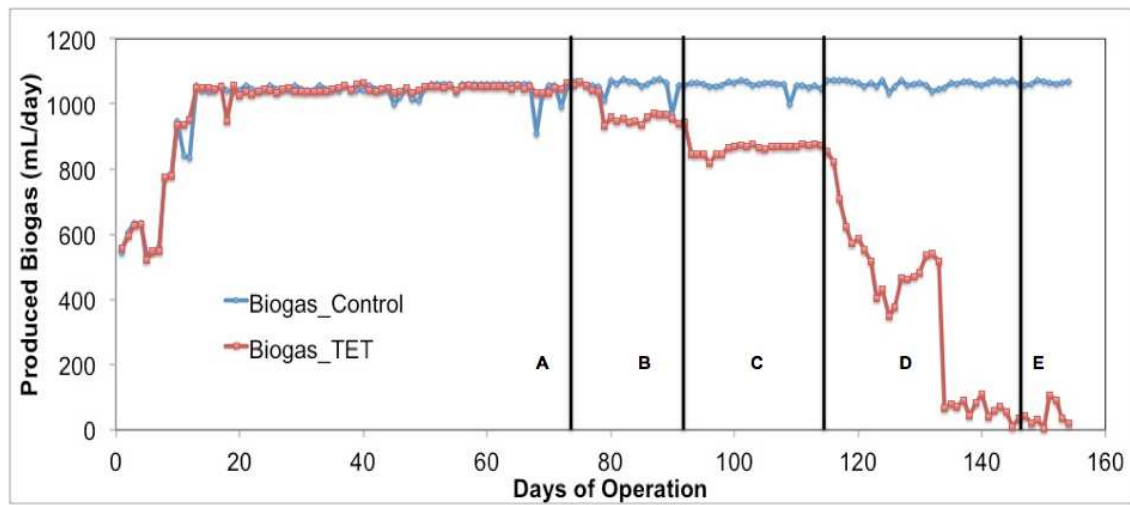
Figure 6. Specific methanogenic activity induced by acetate feeding at different phases of reactor operation (a) biogas production; (b) methane production

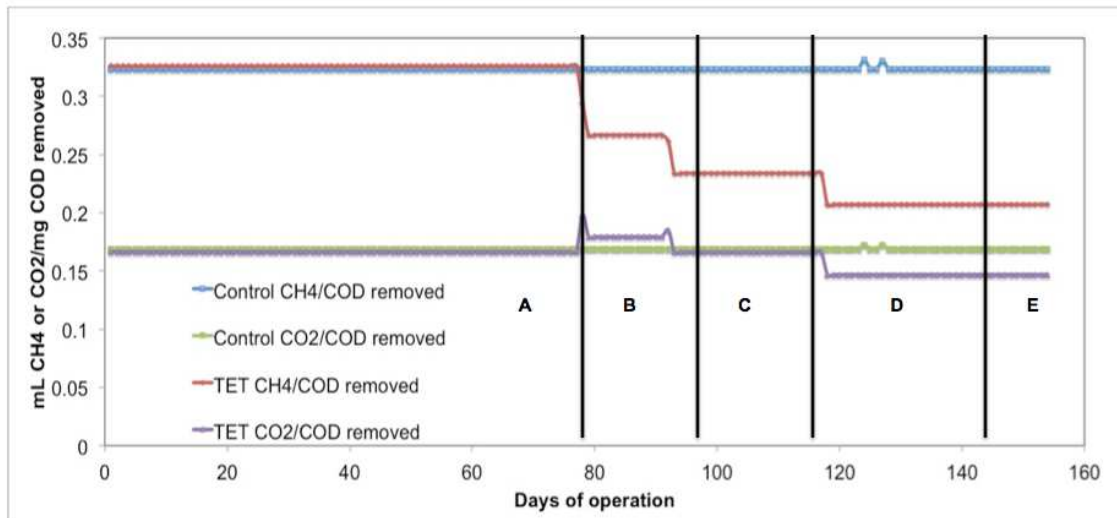
Figure 7. Specific methanogenic activity induced by VFA feeding at different phases of reactor operation (a) biogas production; (b) methane production

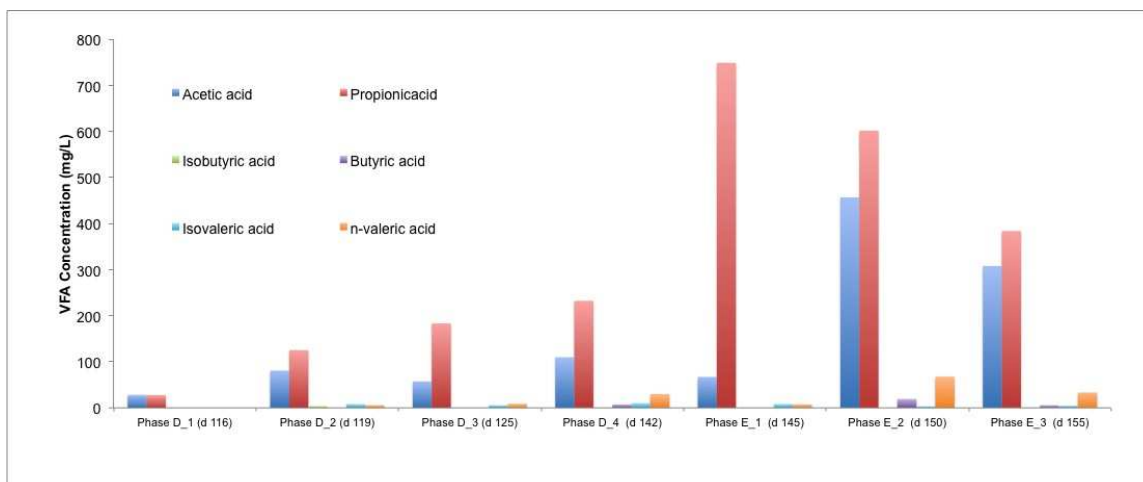


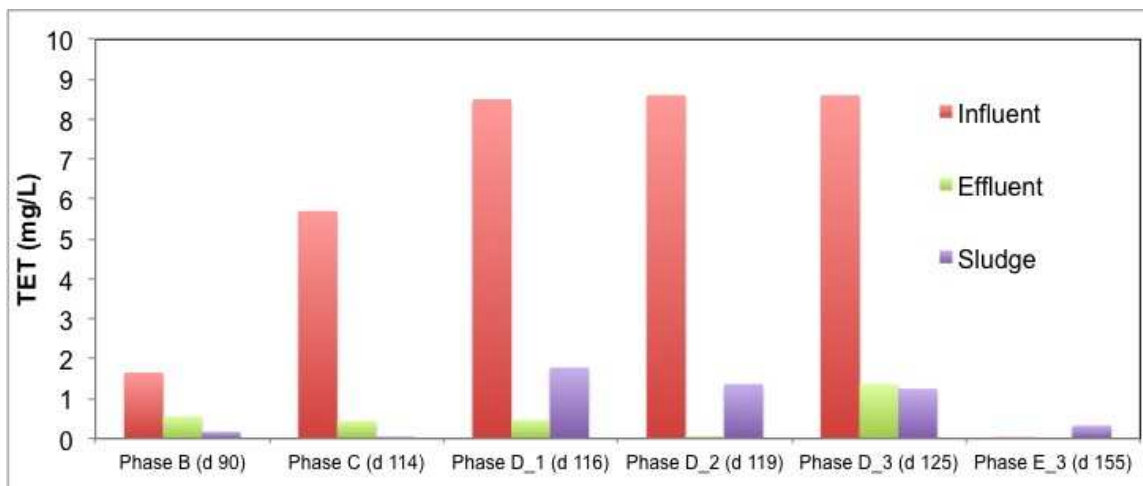
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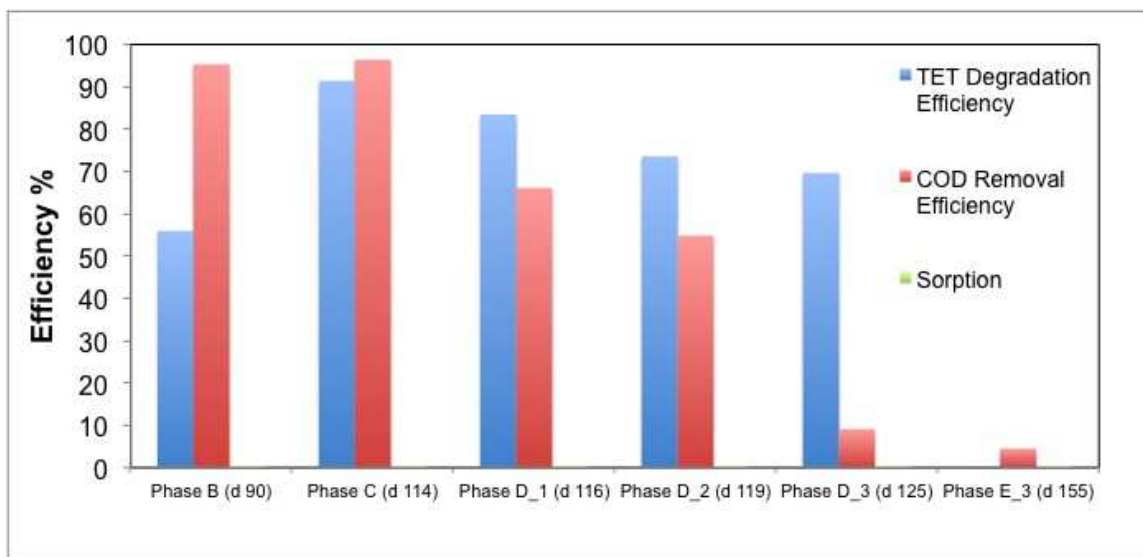


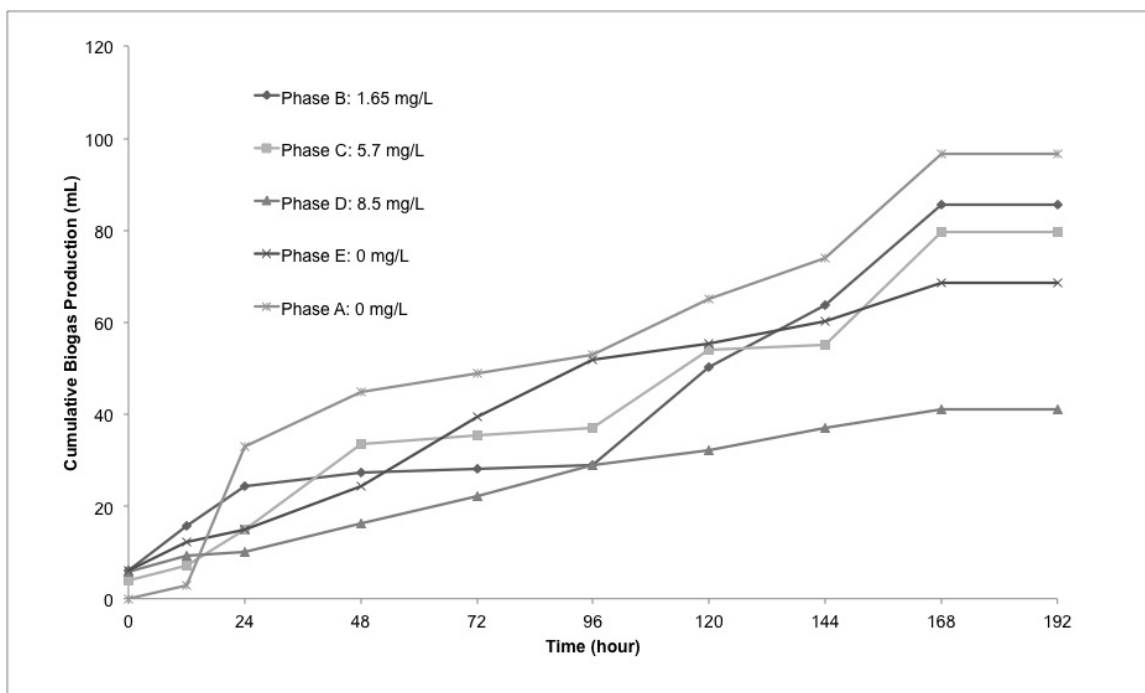


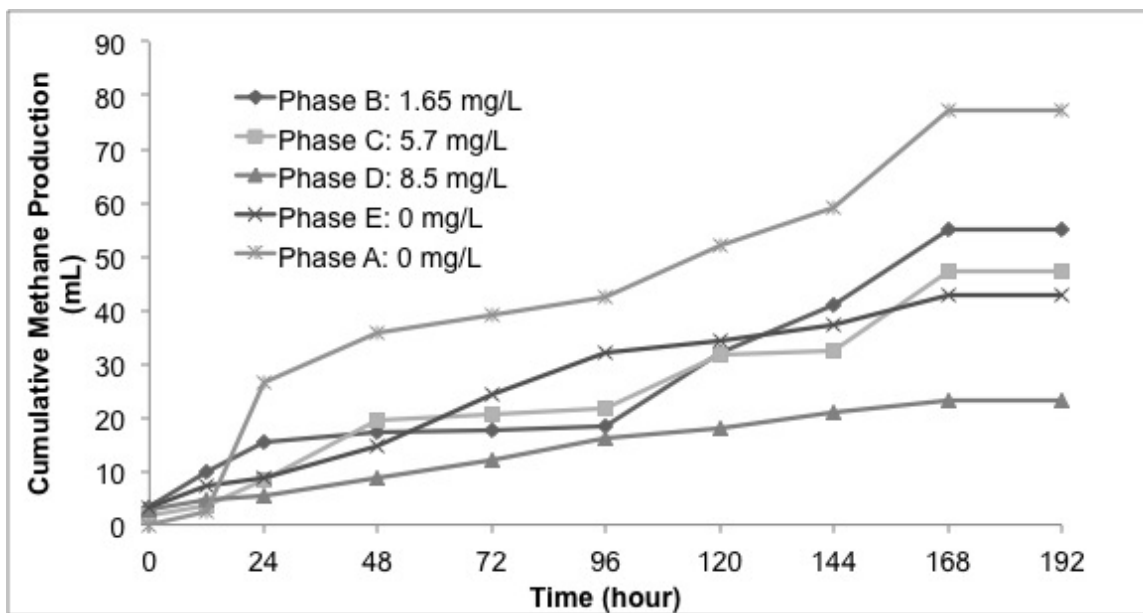


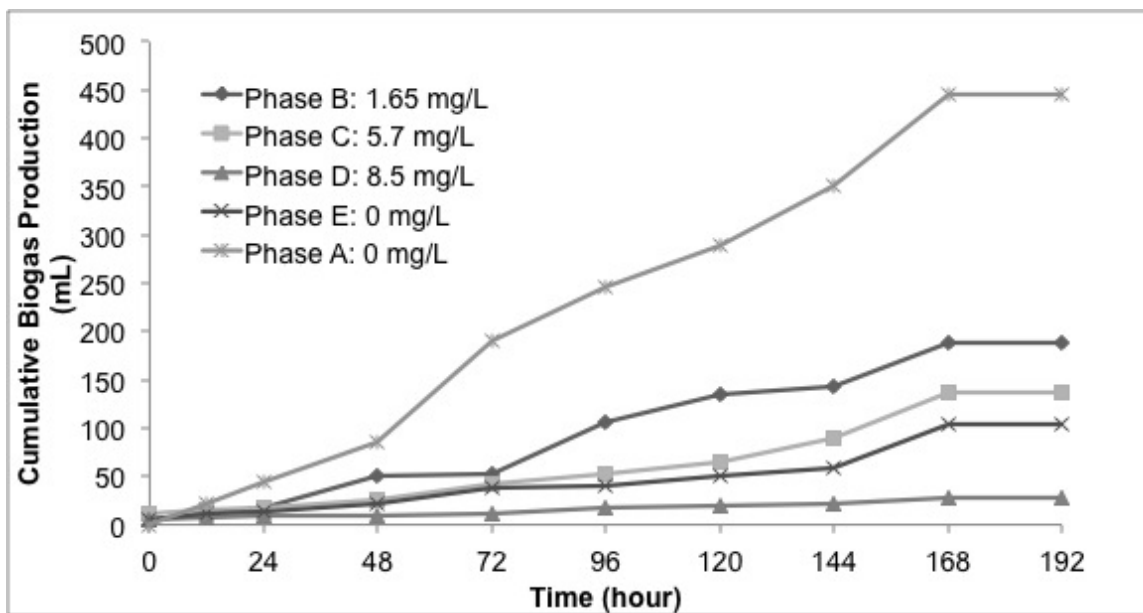


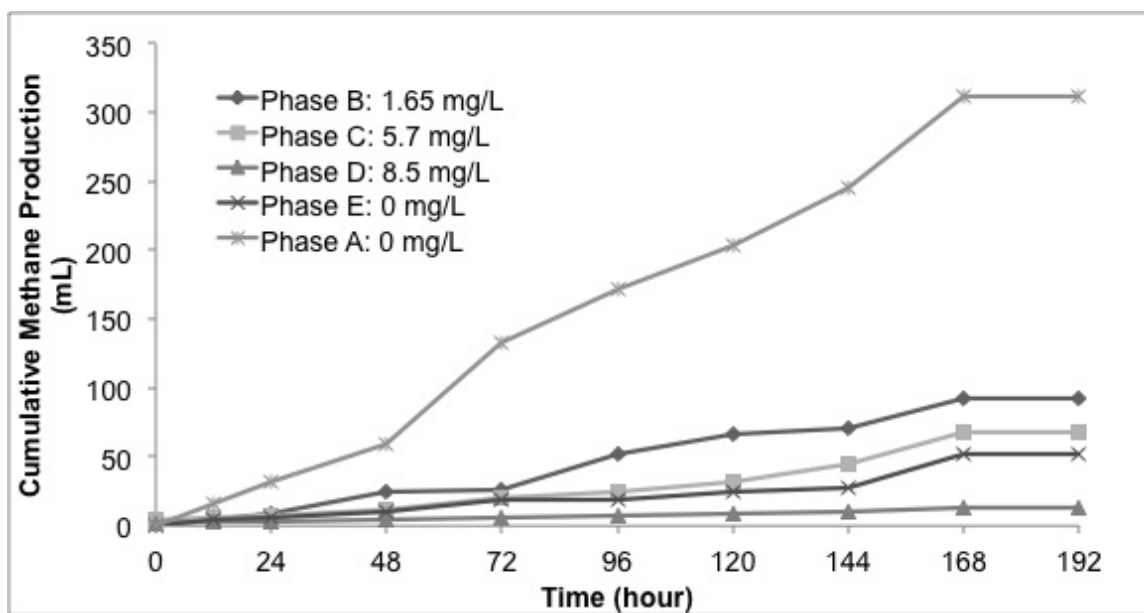












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