



PROGRAMA DE DOCTORADO EN INGENIERÍA QUÍMICA Y AMBIENTAL

TESIS DOCTORAL:

Photosynthetic biodegradation of domestic and agroindustrial wastewater

Presentada por **Dimas Alberto García Guzmán** para optar al grado de Doctor por la Universidad de Valladolid

> Dirigida por: Dr. Raúl Muñoz Torre Dra. Silvia Bolado





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Memoria para optar al grado de Doctor presentada por el Lic. en Química:

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Valladolid, septiembre del 2018





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El incremento poblacional ocurrido en las pasadas décadas a nivel mundial ha generado una creciente demanda de alimentos y suministros, lo que ha causado un incremento irracional en el uso de los recursos hídricos. Esto ha generado grandes volúmenes de aguas residuales domésticas e industriales que requieren ser tratadas. Aunque todavía se utilizan tecnologías físico-químicas para el tratamiento de aguas residuales, sus altos costes de operación e impactos ambientales han fomentado el desarrollo de procesos biológicos. Sin embargo, el bajo potencial de los procesos de lodos activos para la recuperación de recursos de las aguas residuales junto con su alta demanda de energía, y el bajo rendimiento de eliminación de nutrientes de los procesos anaerobios, todavía hacen necesario el desarrollo de soluciones más sostenibles y competitivas para la gestión de aguas residuales domésticas e industriales. Con el objetivo de proponer nuevas soluciones de tratamiento de aguas residuales sostenibles ambientalmente y de bajo costo, esta tesis se centró en determinar el potencial y las limitaciones de los procesos fotosintéticos de depuración antes futuro escalado.

El estado del arte del tratamiento de aguas residuales mediante procesos fotosintéticos se presenta en la sección **Introduction**. Los objetivos, el desarrollo y las estrategias seguidas en esta tesis se resumen en la sección **Aims and Scope**.

En el Capítulo 1 (**Chapter 1**) se estudió la biodegradabilidad de un agua residual doméstica real (RDWW) en un nuevo fotobiorreactor anóxico-aerobio de algas y bacterias. Esta agua residual se eligió por su relevancia ambiental y baja relación C/N/P. Con el fin de superar esta limitación de carbono inorgánico, se acopló una columna de absorción de biogás al fotobiorreactor para proporcionar CO₂ adicional. El lavado de biogás permitió una casi completa nitrificación del NH₄⁺ a NO₃⁻ y promovió un mayor crecimiento de microalgas (con la consiguiente mejora en la asimilación de N y P). Durante los 208 días de operación del sistema, las eficiencias de remoción (RE) de COT, NT, NH₄⁺ y PT representaron 88, 82, 98 y 61%, respectivamente, mientras que la concentración de biomasa del efluente promedió 26 mg SST/L. Por lo tanto, los resultados aquí obtenidos confirmaron que la nueva configuración de fotobiorreactor de algas y bacterias aquí evaluada permite obtener unas adecuadas eficacias de eliminación de C, N y P, y la suplementación de biogás permitió superar la limitación de carbono inorgánico y soportó un proceso eficiente de nitrificación-desnitrificación. Además de la eliminación

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de CO_2 , el sulfuro de hidrógeno presente en el biogás se oxidó a sulfato utilizando el O_2 fotosintéticamente producido en el fotobiorreactor, lo que conllevó una purificación efectiva del biogás a biometano. El suministro de biogás también permitió mantener un pH estable y neutro en el fotobiorreactor, evitando así el uso de reactivos químicos costosos. Finalmente, la sedimentación y recirculación de biomasa mantuvo una concentración de biomasa estable y alta en el proceso, lo que probablemente permitió el dominio de un consorcio de *Scenedesmus* con buenas propiedades de sedimentación.

En el Capítulo 2 (Chapter 2) se investigó el predominio a largo plazo de las poblaciones de microalgas. La mayoría de los estudios realizados en condiciones de laboratorio o exteriores se han centrado en la eliminación de contaminantes claves presentes en las aguas residuales, prestándose muy poca atención a la monitorización de las dinámicas de población de microalgas. En este estudio, se evaluó la evolución de las poblaciones de microalgas durante el tratamiento de purines de cerdo (PWW) diluidos al 15% en cuatro fotobiorreactores abiertos operados a un tiempo de retención hidráulico (HRT) de 27 días y pH = 8 (controlado por adición de CO_2), e inoculados con *Chlorella* sp. (R1), Acutodesmus obliquus (R2), Oscillatoria sp. (R3) y en ausencia de inóculo (R4) durante 180 días. La ausencia de cultivos monoalgales en los cuatro fotobiorreactores durante el período experimental y las rápidas variaciones registradas en la estructura de las poblaciones de microalgas (principalmente en R1 y R2) indicaron la dificultad de mantener cultivos monoalgales durante el tratamiento de PWW en sistemas abiertos. En efecto, la caracterización morfológica de las microalgas reveló que la inoculación de un fotobiorreactor con una microalga específica durante el tratamiento de PWW no garantiza su dominancia a largo plazo. El dominio de Chlorella sp. en los cuatro fotobiorreactores abiertos confirmó la alta tolerancia de esta microalga verde a la contaminación orgánica. Las concentraciones de biomasa en estado estacionario oscilaron entre 2445-2610 mg/L en R1-R3 y 3265 mg/L en R4 (control). No se observaron diferencias significativas en las eficiencias de eliminación de COT (86-87%), CI (62-71%), NT (82-85%) y PT (90-92%). Además, las eficiencias de eliminación de Zn ascendieron a 26% en R3, 37% en R2 y 49% en R1 y R4. En general, se registró una biodegradación eficiente independientemente de las especies de microalgas inoculadas, lo que confirma la robustez de los procesos de algas y bacterias dedicados a la eliminación de carbono y nutrientes de aguas residuales agroindustriales.

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El consorcio de Chlorella sp. obtenido en el Capítulo 2 se utilizó como inóculo en el Capítulo 3 (**Chapter 3**), donde se llevó a cabo una evaluación sistemática del potencial de los fotobiorreactores abiertos de algas-bacterias para el tratamiento de PWW en condiciones interiores y exteriores. Se evaluaron las RE de materia orgánica, nutrientes y zinc de PWW, la dinámica de la estructura de microalgas y la concentración de biomasa. Se evaluó el rendimiento de cuatro fotobiorreactores abiertos alga-bacteria operados a \approx 26 días de HRT con PWW diluido 10 (×10) y 20 (×20) veces en ambiente interior y exterior durante 120 días. Los valores más altos de COT-RE, PT-RE y Zn-RE (94±1%, 100% y $83\pm 2\%$, respectivamente) se registraron en condiciones de interior con una dilución de ×10 PWW, mientras que la NT-RE más alta (72±8%) se obtuvo en condiciones de exterior también en ×10 PWW. C. vulgaris fue la especie de microalga dominante, independientemente de las condiciones ambientales y de la dilución del PWW. Finalmente, el análisis de secuenciación DGGE de las comunidades bacterianas reveló la presencia de cuatro filos, siendo Proteobacteria el filo dominante con 15 de las 23 bandas más intensas. En resumen, este trabajo demostró por primera vez que ni la eliminación de contaminantes ni la estructura de las comunidades de microalgas y bacterias bajo condiciones de interior se puede extrapolar directamente a fotobiorreactores operados en condiciones exteriores.

Finalmente, el Capítulo 4 (**Chapter 4**) tuvo como objetivo evaluar de forma sistemática durante 224 días el rendimiento de fotobiorreactores abiertos alga-bacteria y de bacterias púrpuras fotosintéticas (PPB) durante el tratamiento en condiciones de interior de PWW bajo iluminación artificial. Se investigó la influencia del tiempo del HRT en la eliminación de carbono, nitrógeno, fósforo y zinc, y la influencia en la estructura de la población del consorcio alga-bacteria y de PPB. Además, se investigó la influencia de la dilución del PWW en el rendimiento de biodegradación de un consorcio alga-bacteria y PPB en un sistema en lote en condiciones similares al experimento en continuo. Los resultados demostraron que el fotobiorreactor de algas y bacterias (PBR-AB) soportó las mayores eficiencias de eliminación de nitrógeno, fósforo y zinc (87 ± 2 , 91 ± 3 y $98\pm1\%$), mientras que las remociones de carbono orgánico más altas se registraron en el fotobiorreactor de PPB (PBR-PPB) a un HRT de 10,6 días ($87 \pm 4\%$). Las mayores concentraciones de biomasa fueron registradas en PBR-AB en comparación con PBR-PPB, hecho que fue más evidente a un HRT de 10.6 días, probablemente debido a la activa asimilación fotosintética de CO₂ por parte de las microalgas. *C. vulgaris*, la especie de

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microalga inoculada en PBR-AB, fue detectada en todas las etapas desde el día 7 hasta el día 224, junto con otras especies de *Chlorella*. Curiosamente, la disminución del HRT provocó en PBR-PPB la aparición de microalgas desde el día 98 en adelante. La disminución del HRT en PBR-PPB también determinó el predominio de *Firmicutes* y *Epsilonbacteriaeota* (46.7% y 19.8% de abundancia a un HRT de 4.1 días) y una disminución de la contribución de *Proteobacteria* hasta una abundancia de 12.1%. Se observó también un lavado de la especie *Rhodopseudomonas* seguido del dominio de la especie *Rhodoplanes* en PBR-PPB. Este estudio reveló que el consorcio de algasbacterias soportó un rendimiento superior en términos de eliminación de materia orgánica, lo cual no se observó durante los ensayos de biodegradación de PWW en lote. En general, la disminución del HRT, en lugar del tipo de iluminación, provocó cambios significativos en la estructura de las poblaciones de microalgas y bacterias.

Los resultados obtenidos en esta tesis representan una herramienta valiosa para comprender mejor la dinámica y la estructura de las poblaciones microbianas durante el tratamiento fotosintético de aguas residuales. Estos hallazgos apoyarán la optimización de los procesos de biodegradación fotosintética para mejorar la recuperación de carbono y nutrientes de los efluentes agroindustriales a través de la maximización de la productividad de biomasa. La biomasa generada durante el tratamiento de estas aguas residuales podría utilizarse como materia prima en la producción industrial de bioproductos comerciales, como biocombustibles y biofertilizantes.

iv

Abstract

The increase in human population occurred in the past decades has resulted in an increasing demand of food and supplies, which has caused a higher use of water resources. This has generated larger volumes of domestic and industrial wastewaters that need to be treated. Although physical/chemical technologies are still used to treat wastewater, their high operating costs and environmental impacts have fostered the development of biological processes. However, the poor resource recovery potential of activated sludge processes along with their high energy demand, and the low nutrient removal performance of anaerobic process, still require the development of more cost-competitive and sustainable solutions for the management of domestic and industrial wastewaters. In order to propose new low-cost and sustainable wastewater treatment solutions, this thesis focused on determining the potential and limitations of photosynthetic processes prior to their future scale-up.

The state-of-the-art of wastewater treatment by photosynthetic processes is presented in the **Introduction** section. The objectives, development and strategies followed in this thesis are summarized in **Aims and Scope** section.

In **Chapter 1** the biodegradability of a real domestic wastewater (RDWW) was studied in a novel anoxic-aerobic algal-bacterial photobioreactor. RDWW was chosen based on its environmental relevance and low C/N/P ratio. In order to overcome this carbon limitation a biogas absorption column was coupled to the photobioreactor to provide additional CO₂. Biogas scrubbing supported an almost complete nitrification of the NH₄⁺ to NO₃⁻ and promoted microalgae growth (with the subsequent enhancement in N and P assimilation). Over the 208 days of operation, the removal efficiencies (REs) of TOC, TN, NH₄⁺ and TP accounted for 88, 82, 98 and 61%, respectively, while the TSS effluent concentration averaged 26 mg/L. Therefore, the results herein obtained confirmed that the novel algal-bacterial photobioreactor configuration here evaluated supported consistent removal rates of C, N and P, and biogas supplementation overcame the inorganic carbon limitation and supported an efficient process of nitrificationdenitrification. In addition to CO₂ removal, the hydrogen sulfide present in biogas was oxidized to sulfate using the O₂ photosynthetically produced in the photobioreactor, which entailed an effective biogas upgrading to biomethane. Biogas supplementation also

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Abstract

allowed maintaining a stable and neutral pH in the photobioreactor, thus avoiding the use of costly chemical reagents. Finally, a stable and high biomass concentration was maintained in the process via biomass settling and recirculation, which likely induced the dominance of a *Scenedesmus*-based consortium with enhanced settling properties.

The long-term dominance of microalgae was investigated in Chapter 2. Most studies conducted under laboratory or outdoors conditions focused on the removal of key pollutants present in wastewater, while little attention has been paid to the monitoring of the dynamics of microalgae population. The dynamics of microalgae population during the treatment of piggery wastewater (PWW) diluted at 15% in four open photobioreactors operated at a HRT of 27 days and pH = 8 (controlled by CO_2 addition), and inoculated with Chlorella sp. (R1), Acutodesmus obliquus (R2), Oscillatoria sp. (R3) and in the absence of inoculum (R4), were evaluated for 180 days. The absence of monoalgal cultures in the four photobioreactors throughout the experimental period and the rapid variations in microalgae population structure here recorded (mainly in R1 and R2) revealed the difficulty to maintain monoalgal cultures during the treatment of PWW in open systems. Indeed, the morphological microalgae characterization revealed that inoculation of a photobioreactor during PWW treatment with a specific microalga does not guarantee its long-term dominance. The higher dominance of Chlorella sp. in the four open photobioreactors confirmed the high tolerance of this green microalga to organic pollution. Steady state biomass concentrations ranged from 2445-2610 mg/L in R1-R3 to 3265 mg/L in R4 (control). Interestingly, no significant differences were recorded in the removal efficiencies (REs) of TOC (86-87%), IC (62-71%), TN (82-85%) and TP (90-92%). In addition, Zn-REs accounted for 26% in R3, 37% in R2, and 49% in R1 and R4. Overall, an efficient biodegradation occurred regardless of the microalgae species inoculated, which confirmed the robustness of algal-bacterial processes devoted to carbon and nutrient removals from livestock wastewaters.

The robust *Chlorella* sp. consortium obtained in **Chapter 2** was utilized as inoculum in **Chapter 3**, where a systematic evaluation of the potential of open algal-bacterial photobioreactors for the treatment of PWW under indoor and outdoor conditions was carried out. The REs of organic matter, nutrients and zinc from PWW, along with the dynamics of biomass concentration and structure were assessed. The performance of four open algal-bacterial photobioreactors operated at ≈ 26 days of HRT with 10 (×10) and 20

(×20) times diluted PWW under indoor and outdoor conditions was evaluated for 120 days. The highest TOC-RE, TP-RE and Zn-RE (94±1%, 100% and 83±2%, respectively) were achieved indoors in ×10 PWW, while the highest TN-RE (72±8%) was recorded outdoors in ×10 PWW. *Chlorella vulgaris* was the dominant species regardless of the ambient conditions and PWW dilution. Finally, DGGE-sequencing of the bacterial community revealed the occurrence of four phyla, *Proteobacteria* being the dominant phylum with 15 out of the 23 most intense bands. In summary, this work demonstrated for the first time that neither pollutant removal nor the structure of microalgae and bacterial communities under indoor conditions can be directly extrapolated to outdoors photobioreactors.

Finally, Chapter 4 aimed at systematically evaluating for 224 days the performance of open algal-bacterial and Purple Photosynthetic Bacteria (PPB) photobioreactors for the indoor treatment of PWW under artificial illumination. The influence of the hydraulic retention time (HRT) on the removal of carbon, nitrogen, phosphorous and zinc, and on the structure of the algal-bacterial and PPB population was investigated. In addition, the influence of PWW dilution on the biodegradation performance of an algal-bacterial consortium and PPB was assessed batchwise. The results demonstrated that the algalbacterial photobioreactor (PBR-AB) supported the highest removal efficiencies of nitrogen, phosphorus and zinc $(87\pm2, 91\pm3 \text{ and } 98\pm1\%)$, while the highest organic carbon removals (87±4%) were achieved in the PPB photobioreactor (PBR-PPB) at a HRT of 10.6 days. The higher biomass concentrations recorded in PBR-AB compared to PBR-PPB, which were more evident at a HRT of 10.6 days, were likely due to the active photosynthetic CO₂ assimilation by microalgae. C. vulgaris, the microalga species inoculated in PBR-AB, was detected in all stages from day 7 to day 224, along with other Chlorella species. Interestingly, the stepwise decrease in HRT in PBR-PPB induced the occurrence of microalgae from day 98 onwards. The decrease in HRT in PBR-PPB also mediated the dominance of Firmicutes and Epsilonbacteraeota (46.7 % and 19.8 % of abundance at a HRT of 4.1 days) and decreased the contribution of Proteobacteria to 12.1%. A wash-out of *Rhodopseudomonas* followed by the dominance of *Rhodoplanes* was observed in PBR-PPB. This study revealed that algal-bacterial consortia supported a superior performance in terms of nutrients and zinc removal, although PBR-PPB exhibited a slightly better capacity to remove organic matter, which was not observed during batch PWW biodegradation tests. Overall, the stepwise decrease in HRT, rather

than the type of illumination, caused significant changes in the structure of microalgae and bacterial population.

The results obtained in this thesis represent a valuable tool to better understand the dynamics and structure of microbial populations during photosynthetic wastewater treatment. These findings will support the optimization of photosynthetic biodegradation processes to enhance carbon and nutrient recovery from livestock effluents via maximization of biomass productivity. The biomass generated during livestock treatment could be used as a feedstock in the industrial production of commercial bioproducts such as biofuels and biofertilizers.

List of publications

List of publications included in the thesis

The following publications are presented as part of the present thesis. Three of them are published in international journals indexed in ISI Web of Knowledge (Papers I, II and III). Paper IV has been submitted for publication.

Paper 1. García, D., Alcántara, C., Blanco, S., Pérez, R., Bolado, S., Muñoz, R., 2017. Enhanced carbon, nitrogen and phosphorus removal from domestic wastewater in a novel anoxic-aerobic photobioreactor coupled with biogas upgrading. Chemical Engineering Journal. 313, 424 - 434. doi:10.1016/j.cej.2016.12.054

Paper 2. García, D., Posadas, E., Blanco, S., Acién, G., García-Encina, P., Bolado, S., Muñoz, R., 2018. Evaluation of the dynamics of microalgae population structure and process performance during piggery wastewater treatment in algal-bacterial photobioreactors. Bioresource Technology Journal. 248 (Pt B), 120 -126. doi:10.1016/j.biortech.2017.06.079

Paper 3. García, D., Posadas, E., Grajera, C., Blanco, S., Martínez, S., Acién, G., García-Encina, P., Bolado, S., Muñoz, R., 2017. Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions. Bioresource Technology Journal. 245 (Pt A), 483 - 490. doi.10.1016/j.biortech.2017.08.135

Paper 4. García D., de Godos, I., Dominguez, C., Turiel, S., Bolado, S., Muñoz, R.
2018. A systematic comparison of the potential of a microalgal-bacterial consortium and purple phototrophic bacteria for the continuous treatment of piggery wastewater.
Submitted for puplication to Bioresource Technology (Submitted: 27/08/2018).

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Contribution to the papers included in the thesis

Paper 1. In this work, I was responsible of the design, start-up and operation of the experimentation under the supervision of Dr. Raúl Muñoz. I performed the mass balance calculations, results evaluation and manuscript writing under the supervision of Dr. Cynthia Alcántara, Dr. Silvia Bolado and Dr. Raúl Muñoz. Dr. Saúl Blanco and Dr. Rebeca Pérez were responsible of the characterization of the microalgal and bacterial populations, respectively, where I contributed in the data analysis and discussion.

Paper 2. I was responsible of the design, start-up and operation of the experimental setup under the supervision of Dr. Raúl Muñoz. I performed the mass balance calculations, results evaluation and manuscript writing under the supervision of Dr. Esther Posadas, Dr. Silvia Bolado and Dr. Raúl Muñoz. Dr. Gabriel Acién was responsible of the design and construction of the control system unit of the experimental set-up. Dr. Saúl Blanco was responsible of the characterization of the microalgal populations, where I contributed in the data analysis and discussion.

Paper 3. In this work, I was responsible of the design, start-up and operation of the experimental set-up in collaboration with MSc Eng. Carlos Grajeda under the supervision of Dr. Raúl Muñoz. I conducted the mass balance calculations, results evaluation and manuscript writing under the supervision Dr. Esther Posadas, Dr. Silvia Bolado and Dr. Raúl Muñoz. Dr. Gabriel Acién was responsible of the design and construction of the control system unit of the experimental set-up. Dr. Saúl Blanco and Dr. Sonia Martínez were responsible of the characterization of the microalgal and bacterial populations, respectively, where I contributed in the data analysis and discussion.

Paper 4. During this research, I was in charge of the design, start-up and operation of the experimental set-up in collaboration with Christian Dominguez under the supervision of Dr. Raúl Muñoz. I carried out the mass balance calculations, results evaluation and manuscript writing under the supervision of Dr. Ignacio de Godos, Dr. Silvia Bolado and Dr. Raúl Muñoz. Dr. Sara Turiel was responsible of the characterization of the microalgal populations, where I contributed in the data analysis and discussion.

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Introduction

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1.1 Wastewater pollution

Water is the main component of the hydrological cycle, is essential for sustainable development and it possesses a large social, economic and environmental value (Figure 1) (United Nations, 2017). Water demand worldwide is foreseen to increase in the next decades, with agricultural sector expected to be responsible for 70% of water extraction (UN WWAP, 2017). Additionally, water availability nowadays is affected by human pollution caused by the intensive agricultural, industrial and urban use.



Figure 1. Wastewater in the water cycle (Source: UN WWAP, 2017).

The rapid and sustained human growth will entail a population of 11.2 billion people in 2100 (United Nations, 2015). Consequently, this increase in population density involves the increase on the use of watershed and aquifers (Ihp, 2008) and the uncontrolled discharge of nitrogen and phosphorus from human activities to water bodies (UNWater, 2015). In this context, wastewater composition has changed over the years. For instance, seventy years ago industrial activity increased the number of heavy metals and synthetic organic compounds discharged to the environment, with 10000 new organic compounds being annually introduced to natural water bodies (Metcalf and Eddy, 2003). Table 1 shows the main wastewater pollutants as a function of the type of wastewater and their associated effects.

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	Main	Wastewater		
Pollutant	representative parameters	Domestic	Agroindustrial	Possible effects of the pollutant
		XXX		Aesthetic problems
C	s Total suspended solids		\leftrightarrow	Sludge deposits
Suspended solids				Pollutant adsorption
				Protection of pathogens
D ¹ 1 1 1 1	D'. 1	xxx	\leftrightarrow	Oxygen consumption
Biodegradable organic matter	Biochemical oxygen demand			Death of fish
organic matter				Septic conditions
				Eutrophication
	Nitrogen			Toxicity to fish (ammonia)
Nutrients	Nitrogen, Phosphorus	XXX	\leftrightarrow	Illness in new-born infants
	Thosphorus			(nitrate)
				Pollution of groundwater
Pathogens	Coliforms	XXX	\leftrightarrow	Water-borne diseases
		x	↔	Toxicity (various)
Non-	Pesticides, some detergents, others			Foam (detergents)
Biodegradable				Reduction of oxygen transfer
organic matter				(detergents)
orguine mutter				Non-biodegradability
				malodours (i.e. phenols)
	Specific	x	\leftrightarrow	Toxicity
				Inhibition of biological
Metals	elements (As,			treatment Restrictions in agriculture
wictais	Cd, Cr, Cu, Hg,			use of sludge
	Ni, Pb, Zn, etc.)			Contamination of
				groundwater
		xx		Excessive salinity – harm to
	T 1 1 1 1		↔	crops (irrigation)
Inorganic	Total dissolved solids,			Toxicity to plants
dissolved solids	conductivity			(some ions)
				Problems with soil
				permeability (sodium)
x : small	xx : medium	xxx: large	\leftrightarrow variable	empty: typically not
		č		important

(Source: modified from von Sperling, 2007).

The Europe Union (EU) is currently the second largest pig producer in the World after China, with an average yearly production of $154 \cdot 10^6$ pig head over the last 10 years (Figure 2) (statista, 2018). European pig farming generates $215 - 430 \cdot 10^6$ m³/year (4-8 L/day/pig) of piggery wastewater (PWW) containing high concentrations of organic matter and nutrients (Table 2) (de Godos et al., 2009a).



Figure 2. Pig population in European Union during the last 10 year (statista, 2018).

The estimated average organic matter and nutrient load from EU piggery wastewater in 2017 amounted to 8.923.000 tn chemical oxygen demand (COD)/year, 890.000 tn nitrogen (N)/year and 223.000 tn phosphorous (P)/year (statista, 2018). Likewise, an excessive input of residual phosphorus and nitrogen can cause eutrophication in rivers and lakes (García Ferrero, 2013; Olguín, 2003). In addition, PWW can contain heavy metals such as zinc and copper, typically used as growth promoters in swine nutrition (Abe et al., 2012; De la Torre et al., 2000). Table 2 shows piggery wastewater composition.

Parameters	Range (mg/L)	
	Min	Max
Total Solid (TS)	6000	7000
Total Nitrogen (TN)	3000	4700
Total Phosphorus (TP)	900	1300
BOD ₅	10000	20000
COD	38000	42000
Organic matter	35000	40000
Heavy metals (HMs)	variable	

 Table 2. Piggery wastewater composition.

Source: (García Ferrero, 2013).

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1.2 Photosynthetic wastewater treatment

Wastewater treatment (WWT) regulations are becoming stricter worldwide to prevent the development of septic conditions, mal odors, eutrophication and toxicity in natural water bodies. Many technologies have been engineered and up-scaled in the past century to deal with the health and environmental concerns derived from the uncontrolled discharge of untreated wastewater. Among them, photosynthetic wastewater treatment (either based on microalgae or on purple photosynthetic bacteria) has emerged as a promising platform for wastewater treatment coupled with resource recovery (Metcalf and Eddy, 2003).

Algal-bacterial consortia in photobioreactors can actively reduce the levels of organic matter and nutrients in wastewater via aerobic carbon oxidation and nutrient assimilation into biomass (de Godos et al., 2014; Liu et al., 2017). Likewise, the heavy metals present in PWW are typically removed by biosorption (Abe et al., 2012; Muñoz et al., 2006; Romera et al., 2007), while photosynthesis promotes the uptake of the greenhouse gas carbon dioxide (CO₂) and supplies the oxygen required for the aerobic oxidation of organic matter (Figure 3) (Cho et al., 2013).



Figure 3. Algal-bacterial symbiosis during organic matter oxidation in photobioreactors. Source: (Muñoz and Guieysse, 2006).

However, microalgae metabolism requires proper physical, chemical and biological conditions in order to support an effective pollutant removal. Thus, temperature (Choi et al., 2013; Rawat et al., 2011), pH (Arbib et al., 2013) (Markou et al., 2014) and light irradiance (Cuellar-Bermudez et al., 2015; Lavoie and de la Noüe, 1985) are key environmental parameters determining microalgae growth during photobioreactor operation. Additionally, the aqueous concentrations of dissolved oxygen (DO) (Diameter and Sewers, 2000) (Muñoz and Guieysse, 2006) and carbon dioxide (Heubeck et al.,

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2007; Park and Craggs, 2010; Posadas et al., 2015a) and the hydraulic retention time (HRT) (Gutiérrez et al., 2015; Hao et al., 2013; Sutherland et al., 2014) also govern the performance of microalgae during wastewater treatment in algal-bacterial photobioreactors.

For instance, Shriwastav et al., (2014) investigated the effects of nutrient levels on the growth and nutrient uptake, as well as the effects of cultivation conditions, on the physiology of *Chlorella Sorokiniana*. Evans, et al., (2017) evaluated the potential of *Chlorella vulgaris* for the treatment of municipal primary settled in term of NH₃, PO₄ and COD removal (Evans et al., 2017). Likewise, *Chlorella* sp. GD was used for the production of biomass and lipids using piggery as cultivation medium (Kuo et al., 2015). Prandini et al., (2016) used *Scenedesmus* sp. to enhance nutrient removal from swine wastewater digestate coupled to biogas upgrading in two interconnected 16.9 L glass closed-photobioreactors.

On the other hand, Purple Photosynthetic Bacteria (PPB) have the potential to become a key player in the coming generation of wastewater treatment technologies. Indeed, PPB have the ability to transform a great variety of inorganic and organic pollutant into harmless minerals or valuable products, which can be recycled. Indeed, PPB can support the bioconversion of residual organic matter into polyhydroxyalkanoate (Hassan et al., 1997), Polypoly- β -hydroxybutyrate (Khatipov et al., 1998; Luongo et al., 2017), biohydrogen (Basak and Das, 2007) and phototrophic bacterial biomass (Azad et al., 2001). PPB have a very versatile metabolism, able to grow either in the presence or absence of molecular oxygen (O₂) (Rittmann and McCarty, 2012) and phototrophically based on infrared light (Puyol et al., 2017). PPB photosynthesis is carried out with reduced sulfur compounds or simple organic compounds as electron donor according (Figure 4). PPB can grow autotrophically using infrared light as the energy source for CO₂ fixation and inorganic electrons donor such as H₂, Fe²⁺, S²⁻ or S₂O₃²⁻ (cyclic anoxygenic photosynthesis) (Goldman et al., 1972; Overmann and Garcia-Pichel, 2013).



Figure 4. Mechanism of PPB photosynthesis. H₂A refers to the electron donor (Goldman et al., 1972).

Moreover, PPB can also grow heterotrophically in presence and absence of light. Figure 5 shows the schematic mechanistic model of PPB metabolism during domestic wastewater treatment (Puyol et al., 2017).



Figure 5. Schematic summary of PPB metabolism during domestic wastewater treatment. Key: N₂ase: Nitrogenase complex. TCA-c: Tri-carboxylic acid cycle. DF: Dark fermentation. VFA: volatile fatty acids. e⁻: electrons. Dashed lines: electron flow. Dotted line: proton flow. S_{AC} Concentration of acetate (mg COD/L). S_{IC} Concentration of inorganic carbon (mol C/L). S_S Concentration of biodegradable soluble fraction but excluding acetate, alcohols and some sugars (mg COD/L).*: Model components. Source (Puyol et al., 2017).

1.3 Microalgae-based wastewater treatment

Algal-bacterial symbiosis constitutes a cost-effective technology for the treatment of domestic wastewater (Table 1) and piggery wastewater because of their high concentration of biodegradable organic matter and nutrients (Table 2) (de Godos et al., 2010). This technology has attracted a significant attention in the past decade based on the global quest for environmentally friendly technologies for wastewater treatment (Muñoz et al., 2012), the mitigation of greenhouse gases (Dassey and Theegala, 2013) and recent interest in biomethane production (Xia and Murphy, 2016). The lower energy demand of microalgae-based wastewater treatment compared to conventional activated sludge systems, along with the ability of microalgae to fix CO₂ phosynthetically, significantly increase the environmental sustainability of this technology (Cheah et al., 2016; Dassey and Theegala, 2013). (Unnithan et al., 2014). The in-situ oxygen supply mediated by microalgae photosynthesis during organic matter oxidation represents however the main advantage of this green technology (Oswald et al., 1957). Microalgaebased processes can support an effective wastewater treatment coupled to the recovery of microalgae rich in proteins, carbohydrates and lipids that can be used as a feedstock for the production of added-value products using a biorefinery approach (Olguín et al., 2013). However, despite microalgae cultivation in wastewater entails significant economic and environmental advantages over conventional CO₂-supplemented mass production of microalgae in mineral salt media, controversy still exists in literature about the possibility of maintaining monoalgal cultures during microalgae-based wastewater treatment.

As stated in section 1.2, algal growth during wastewater treatment is influenced by abiotic factors such as pH, salinity, temperature, light (quality and quantity) and the concentration of nutrients, ions, toxicants, dissolved oxygen and carbon dioxide (Gupta and Diwan, 2017). Likewise, microalgae activity is also influenced by biotic factors such as the presence of pathogens (bacteria and viruses), predation by zooplankton and competition for essential nutrients with other microorganisms (Lau et al., 1995). Finally, algal productivity depends on operational parameters such as culture agitation, HRT, addition of external inorganic carbon and harvest frequency (de la Noüe et al., 1992).

Microalgae can grow under heterotrophic and phototrophic conditions and require the supply of essential elements such as carbon, hydrogen, oxygen, nitrogen, phosphorus,

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sulfur and iron (Rittmann and McCarty, 2012). Microalgae capture sunlight during oxygenic photosynthesis, being responsible for the production of approximately half of atmospheric oxygen on earth and absorbing massive amounts of carbon dioxide (Safi et al., 2014). Carbon dioxide, whose concentration has increased from 280 to 390 ppm_v since the pre-industrial era till 2013, is the main greenhouse gas emitted nowadays (Cheah et al., 2015; Wayne Chew et al., 2017). Microalgae are able to biosequester CO_2 from flue gases generated in power plants, which can contribute to significantly reduce greenhouse gas emissions (Cheah et al., 2015). Likewise, the superior heavy metal biosorption in algal-based processes compared to conventional activated sludge processes has been recently demonstrated (Muñoz et al., 2012). Additionally, microalgae-based wastewater treatment is typically envisaged as a platform for algal biofuel feedstock production (Alcántara et al., 2015c).

Despite the above-mentioned advantages, this technology exhibits some disadvantages such as its high land requirements and the poor control of microalgae population in large-scale open systems. Likewise, microalgae-based WWT has some limitations for full-scale implementation (Muñoz et al., 2012) derived from the limited number of cost-effective biomass harvesting techniques that can cope with the poor sedimentation of most species (Su et al., 2011). Process operation at short HRT (<5 days) can limit ammonium nitrification (Alcántara et al., 2015a; Chaumont, 1993).

The mechanisms of carbon, nitrogen, phosphorous and heavy metal removal underlying WWT in algal-bacterial photobioreactors are described in the following sections.

1.3.1 Carbon Removal

Carbon in the form of carbon dioxide from the atmosphere, flue gas, biogas or organic matter biodegradation can be fixed via photosynthesis into microalgae biomass (Cai et al., 2013). Likewise, domestic, agroindustrial and livestock wastewaters contain high concentrations of organic matter and inorganic carbon that will be fixed into algalbacterial biomass during WWT via aerobic carbon oxidation and photosynthesis. Equations 1 and 2 exemplify these reactions through the aerobic oxidation of glucose and the production of carbohydrates [1] and [2] (Rittmann and McCarty, 2012):

$$C_6 H_{12} O_6 + 3 O_2 \rightarrow 3(C H_2 O)h + 3 C O_2 + 3 H_2 O$$
^[1]

The aerobic biodegradation of one mol of glucose by heterotrophs (h) consumes three mol of oxygen, which we can obtain by converting three mol of carbon dioxide into oxygen by microalgae according to Equation [2]. Empirical observations in our lab have consistently demonstrated that the main mechanisms of carbon removal in WWT open-photobioreactors are bioassimilation and carbon dioxide stripping (Posadas et al., 2014).

$$CO_2 + H_2O + Light \rightarrow (CH_2O)p + O_2$$
[2]

Equation 2 clearly shows that the production of one mole of oxygen and one mole of phototrophic biomass (p) is associated to the assimilation of one mole of CO₂.

Biogas upgrading can also support CO₂ removal: In this context, biogas production by anaerobic digestion reduces the emissions of methane (CH₄) and nitrous oxide (N₂O) potentially produced during the storage and utilisation of untreated animal manure as fertiliser. The global warming potential of methane is higher than CO₂ and N₂O by factors of 23 and 296, respectively (Seadi et al., 2008). Biogas is mainly composed of methane (CH₄ \approx 40-75%), (CO₂ \approx 25-50%), hydrogen sulfide (H₂S \approx 0.005-2%) and ammonia (NH₃ < 1%) (Toledo-Cervantes et al., 2017a). In this context, biogas supplementation to photobioreactors improve biomass productivity, nutrient removal and generates a high quality biomethane (Figure 6) (Posadas et al., 2017; Serejo et al., 2015). Photosynthetic biogas upgrading is based on the fixation of CO₂ via photosynthesis by microalgae and the oxidation of H₂S to sulfate by sulfur oxidizing bacteria using the oxygen photosynthetically produced (Toledo-Cervantes et al., 2016).



Figure 6. Schematic of the experimental set-up devoted to the simultaneous biogas upgrading and diluted centrate treatment (Source: Alcántara et al., 2015b).

According to the Directive 2003/55/EC (European Parliament, 2003), the upgraded biogas must comply with specific quality requirements to be injected into natural gas grid:

of $CH_4 \ge 95\%$, $CO_2 \le 2\%$, $O_2 \le 0.3\%$ and negligible amounts of H_2S . Table 3 compiles the composition of biomethane obtained in microalgae-based processes.

Type of Biogas	Type of PBR	L/G ratio	CO2 (%)	O2 (%)	N2 (%)	CH4 (%)	References
+Raw	20L Bag	-	N.R	N.R	N.R	88.25	(Xu et al., 2015)
*Synthetic	HRAP+AC	1	0.4	0.03	2.4	97.2	(Toledo-Cervantes et al., 2016)
*Synthetic	HRAP+AC	0.5	0.4	0.7	2.7	96.2	(Toledo-Cervantes et al., 2017b)
*Synthetic	HRAP+AC	1	N.R	N.R	N.R	99.6	(Marín et al., 2018)

Table 3. Compilation of the composition of the biomethane produced in microalgaebased upgrading processes.

*Synthetic biogas CH₄:CO₂:H₂S=70%:29.5%:0.5%

+Raw biogas CH₄:CO₂:H₂S= $58.67\pm3.45\%$: $37.54\pm2.93\%$: $0.79\pm0.06\%$ N.R- Not reported.

AC: absorption column

1.3.2 Nutrient removal

Microorganisms typically prefer ammonium over nitrate and nitrite as a nitrogen source for cells synthesis based on its lower oxidation state, although when ammonium is not available many prokaryotic cells can use oxidized forms of nitrogen to support protein synthesis (Rittmann and McCarty, 2012) (Figure 7). Nitrogen in the form of NH₄⁺ and organic nitrogen is typically present at high concentrations in domestic and livestock wastewaters, while the presence of nitrate in industrial discharges. Several mechanisms, ranging from bioassimilation to abiotic processes, support nitrogen removals of 90-98% during microalgae-based wastewater treatment (Alcántara et al., 2015b).



Figure 7. Biogeochemical and physical-chemical (pc) processes underlying the speciation of nitrogen in aquatic systems. (Source: Tsuyuzaki, 2014).

1.3.2.1 Assimilatory nitrogen removal

During microalgae-based wastewater treatment, nitrogen removal via assimilation is limited by the concentration of total carbon (organic+inorganic) present in the wastewater. Overall, biomass yields are lower when nitrate or nitrite are used as a nitrogen source than those obtained with ammonium (Alcántara et al., 2015c). Table 4 compiles the removal efficiencies of nitrogen in algal-bacterial photobioreactors treating wastewater.

Figure 8 shows the steps followed by the reduction of oxidized nitrogen to ammonium and its further incorporation into amino acids. Nitrate and nitrite reduction is carried out by the enzymes nitrate and nitrite reductase, respectively. Nitrate reductase uses the reduced form of nicotinamide adenine dinucleotide (NADH) to transfer two electrons, resulting in the conversion of nitrate into nitrite. Nitrite is reduced to ammonium by the enzyme nitrite reductase and ferrodoxin (Fd). All forms of inorganic nitrogen are ultimately reduced to ammonium prior to their assimilation into aminoacids within the cytoplasm. Finally, glutamate (Glu), adenosine triphosphate (ATP) and the enzyme glutamine synthase facilitate the incorporation of ammonium into the amino acid Lglutamine.



Figure 8. Simplified schematic of the assimilation of nitrate. (Source: Cai et al., 2013).

1.3.2.2 Dissimilatory nitrogen removal

More specifically, nitrification is carried out by nitrifying bacteria in a two-step process (Rittmann and McCarty, 2012). *Nitrosomonas* carry out the first step where NH_4^+ is oxidized to NO_2^- according to the energy-yielding reaction depicted (equation [3]). Species from the genera *Nitrosococcus*, *Nitrosopira*, *Nitrosovibrio* and *Nitrosolobus* are also able to oxidize NH_4^+ to NO_2^- .

$$\frac{1}{6}NH_4^+ + \frac{1}{4}O_2 = \frac{1}{6}NO_2^- + \frac{1}{3}H^+ + \frac{1}{6}H_2O$$
[3]

The second step of nitrification is the aerobic oxidation of NO₂⁻ to NO₃⁻ (equation [4]),

which is typically carried out by the genera Nitrobacter and Nitrospira.

$$\frac{1}{2}NO_2^- + \frac{1}{4}O_2 = \frac{1}{2}NO_3^-$$
[4]

On the other hand, denitrification consists of the dissimilatory reduction of NO_3^- or NO_2^- to N_2 , where NO_3^- or NO_2^- are the electron acceptors for energy generation during the oxidation of organic matter or reduced inorganic compounds. The denitrification process occurs in four sequential steps where NO_3^- is reduced first to nitrite NO_2^- (equation [5]), then nitric oxide (NO) (equation [6]), nitrous oxide (N₂O) (equation [7]) and finally to N₂ gas (equation [8]).

$$NO_{3}^{-} + 2e^{-} + 2H^{+} = NO_{2}^{-} + H_{2}O$$
Nitrate reductase [5]

$$NO_{2}^{-} + e^{-} + 2H^{+} = NO + H_{2}O$$
Nitrite reductase [6]

$$2NO + 2e^{-} + 2H^{+} = N_{2}O + H_{2}O$$
Nitric oxide reductase [7]

$$N_{2}O + 2e^{-} + 2H^{+} = N_{2(q)} + H_{2}O$$
Nitrous oxide reductase [8]

Overall, the reduction from NO_3^- to N_2 requires 10 electron equivalents per mol of nitrogen (N₂): four electrons in the first step and two electron in the following three steps (Rittmann and McCarty, 2012). During wastewater treatment, nitrification and denitrification processes can occur either simultaneously in the same tank or in two separate tanks. In the particular case of microalgae-based wastewater treatment, several studies have consistently shown the feasibility to conduct denitrification in an anoxic tank interconnected via an internal recirculation to a photobioreactor supporting NH_4^+ oxidation (Alcántara et al., 2015a; de Godos et al., 2014).

1.3.2.3 Abiotic nitrogen removal

Microalgae photosynthesis increases the pH in the cultivation broth as a result, among others, of CO₂ uptake (equation [9]). This rise in the pH is severe under carbon limiting conditions (Alcántara et al., 2015c). This pH increase mediated by algal photosynthesis results in an enhanced NH_4^+ stripping (in the form of NH_3 with is the dominant form at high pH). Thus, abiotic nitrogen removal in open ponds is controlled by parameters that determine algae activity solar radiation, temperature, HRT or sludge retention time, agitation and superficial area (García et al., 2000). Table 4 compiles the removal efficiencies of nitrogen in algal-bacterial photobioreactors treating wastewater.

 $CO_2(l) + H_2O(l) \leftrightarrow H_2CO_3 \leftrightarrow HCO_3^- + H^+ \leftrightarrow CO_3^{2-} + 2H^+$[9]

Type of Wastewater	Type of PBR	Volumen (L)	HRT (days)	TN (%)	TP (%)	References
Piggery	HRAP	464	10	88	*	(de Godos et al., 2009a)
Piggery	Tubular biofilm	7.5	7	100	90	(de Godos et al., 2009b)
Piggery	Enclosed jacketed glass	3.5	4.4	37	*	(de Godos et al., 2010)
Piggery	HRAP	400	40	86	*	(Aguirre et al., 2011)
Fish processing	HRAP	3	5	> 95	73	(Riaño et al., 2011)
Fish processing	HRAP	3	10	> 99	76	(Riaño et al., 2012)
Centrate & domestic	Biofilm	31	10	70	85	(Posadas et al., 2013)
Raw domestic	Biofilm	31	10	92	96	(Posadas et al., 2014a)
Fish farm & domestic	HRAP	180	10	83	94	(Posadas et al., 2014b)
Domestic	Raceway	800	6	98	6	(Posadas et al., 2015a)
Vinasse wastewater	HRAP	180	7	74	78	(Posadas et al., 2015b)
Synthetic wastewater	Enclosed jacketed glass	3.5	2	79	*	(Alcántara et al., 2015a)
Piggery	Vertical column	4	4	99	94	(Lee and Han, 2016)
Centrate	HRAP	180	4	86	92	(Posadas et al., 2017)

Table 4. Compilation of removal efficiencies of nitrogen and phosphorus in algalbacterial photobioreactors treating wastewater.

*- Not determined

1.3.3 Phosphorus removal

1.3.3.1 Biological phosphorus removal

Phosphorus is an essential macronutrient that spurs the growth of microalgae and cyanobacteria, leading to accelerated eutrophication of lakes (Rittmann and McCarty, 2012). Table 4 compiles the removal efficiencies of phosphorus in algal-bacterial photobioreactors treating wastewater. There are two biotic mechanisms to remove phosphorus in algal-bacterial processes devoted to WWT (Rittmann and McCarty, 2012).

a.) Regular phosphorous uptake into biomass: during regular phosphorus uptake the algal-bacterial biomass typically contains 0.5 to 1% P dry weight basis. This phosphorous is used to build-up cell membranes and genetic material.

b.) Enhanced biological phosphorus uptake: Certain heterotrophic bacteria present in the algal-bacterial mixed liquor during WWT are capable of sequestering high levels of phosphorus as intracellular polyphosphate (poly P), which is an energy storage material. These bacteria are collectively called polyphosphate accumulating organisms (PAOs), the most well-known group of PAOs being '*Candidatus Accumulibacter phosphatis*' (Yuan et al., 2012). If such microorganisms are selected, induced to store poly P, and wasted when rich in poly P, the net removal of P through biomass uptake can be increased significantly. When the enhanced biological phosphorus removal is successful, the biomass wasted can contain up to 2 to 5 times higher P content than regular biomass (Rittmann and McCarty, 2012).

1.3.3.2 Abiotic phosphorus removal

As indicated above, the microalgae photosynthesis shifts the equilibrium represented by equation [9], increasing the pH and dissolved oxygen (DO) levels in the cultivation broth. Phosphate precipitation occurs at high pH in the presence of Ca^{+2} and Mg^{+2} via hydroxyapatite formation and surface adsorption via formation of hydrogen bonds with the extracellular polysaccharides secreted by microalgae (equation [10]) (González-Fernández and Muñoz, 2017) (Liu et al., 2017).

 $3 HPO_4^{2-} + 5 Ca^{2+} + 4 OH^- \rightarrow Ca_5(OH)(PO_4)_3 + 3 H_2O.....[10]$

1.3.4 Heavy metals removal

Microalgae support heavy metals removal from wastewater through a combination of passive and active mechanisms. In this context, the main mechanisms of metal uptake in microalgae are physical adsorption, ion exchange, transport across microbial cell wall and bioassimilation, complexation and precipitation (Javanbakht et al., 2014). (Chojnacka et al., 2005). During WWT in algal-bacterial photobioreactors, bacterial biosorption also contributes to HMs removal (Mohamed, 2015). Table 5 compiles the removal efficiencies of Zn (heavy metal typically used in pig nutrition) by microalgae during WWT.

Type of Wastewater	Type of PBR	Algae type	Zn initial concentration (mg/L)	Zn removal (%)	References
Acid wastewater	Batch test	Cyanidium caldarium	41.5	95	(Ahlf, 1988)
Zinc ore treatment plant	Batch test	Oscillatoria anguistissima	2.2 8.5 75.0	66 93 92	(Ahuja et al., 1999)
Zn Solutions	Batch test	Scenedesmus obliquus	0.5 1.5 4.5 8.0	59 52 45 41	(Omar, 2002)
		Scenedesmus quadricauda	0.5 1.5 4.5 8.0	61 49 43 37	
Synthetic wastewater & real mine dump leachate	Porous Substrate Bioreactor (PSBR)	Stichococcus bacillaris	2.0 3.0	80 80	(Li et al., 2015)
Zn Solutions	Batch test	Chlorella minutissima UTEX2341	2 (mM) 4 (mM) 6 (mM)	62 46 38	(Yang et al., 2015)
Zn Solution	Batch test	Chlorophyceae spp	3.0	88	(Saavedra et al., 2018)

Table 5. Compilation of the removal efficiencies of Zn by microalgae.

1.4 Purple Phototrophic Bacteria-based wastewater treatment

Purple phototrophic bacteria (PPB) are ubiquitous in fresh and marine water, soil, wastewater and activated sludge (Myung et al., 2004). PPB can use multiple sources of energy and carbon for growth and maintenance. For instance, PPB can utilize different type of carbon as a carbon source for the synthesis of new cellular material, heterotrophic microorganisms being able to utilize organic carbon as an energy source while autotrophic organisms utilize carbon dioxide exclusively as a carbon source (United States Environmental Protection Agency, 2013). PPB are classified according to their ability to use infrared light as the energy source as phototroph and chemotrophs (Overmann and García-Pichel, 2013). In addition, PPB require nutrients and trace metals for cell synthesis (Rittmann and McCarty, 2012). PPB-based WWT processes have emerged as an alternative to conventional wastewater treatment with a superior potential for energy and nutrient recovery (Hülsen et al., 2016b). In this context, PPB have attracted recent interest due to their capacity to transform organic matter and nutrients with high substrate-to-product conversion yields, in the absence of oxygen, and under a wide range of infrared light wavelengths (Basak and Das, 2007; Hülsen et al., 2016b; Puyol et al., 2017). For instance, the performance of PPB has been tested for the treatment of domestic

(Hülsen et al., 2016a, 2016b, 2014; Zhang et al., 2003), piggery (Myung et al., 2004), rubber sheet (Kantachote et al., 2005), pharmaceutical (Madukasi et al., 2010), palm oil mill (Hassan et al., 1997) and fishery wastewaters (de Lima et al., 2011). PPB are considered a versatile platform for WWT based on their high grow rates, ability to assimilate organic matter and to remove nutrients by denitrification and poly-P formation (Hülsen et al., 2014). The recovery of organic matter and nutrients from WW is typically carried out in a photobioreactor that utilizes infrared light as a source for PPB growth.

The next sections describe the main mechanisms of carbon, nutrient and heavy metal removal by PPB.

1.4.1 Carbon removal

The main characteristic of PPB is their high metabolic versatility. Indeed, these microorganisms are unique in their ability to employ all known modes of metabolism to assimilate carbon (Figure 5) (Puyol et al., 2017). In addition to the obvious function of providing carbon for cell material construction, CO₂ assimilation and organic carbon oxidation play a central role in maintaining the redox balance of PPB (Hunter et al., 2009). On the other hand it should be stressed that although PPB possess simplified, and sometimes unique, photosystems compared to higher plants and green algae, their carbon metabolism pathways are rather complex (Tang et al., 2011).

1.4.1.1 Autotrophic CO₂ fixation

Photosynthesis supports the assimilation of CO_2 autotrophically into cellular material with the aid of light, with the most well-known CO_2 assimilation (Tang et al., 2011). Many metabolites, intermediates or products produced by these autotrophic carbon fixation pathways are essential for building cellular material (Hügler and Sievert, 2011; Tang et al., 2011).

1.4.1.2 Carbohydrate metabolism

Most heterotrophic organisms use carbohydrates as a carbon sources to build up cell material and provide reductive power. Three are carbohydrate catabolic pathways in photosynthetic organisms: the Emden–Meyerhof–Parnas (EMP) pathway (glycolysis), Entner–Doudoroff (ED) pathway, and PP pathway (phosphogluconate pathway) (Tang et al., 2011).

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1.4.2 Biological nutrient removal

PPB can assimilate NH_4^+ and reduce nitrate to organic N for assimilatory purposes. The genes for nitrate assimilation are expressed when ammonia and other forms of fixed nitrogen are limiting and nitrate is available. PPB are also able to fix N₂ (Hunter et al., 2009). The three main mechanism involved in nitrogen removal are reviewed in the following sections.

1.4.2.1 Nitrogen assimilation

Purple phototrophic bacteria have the ability to reduce nitrate to ammonium for assimilatory purpose. This is an 8-electron reduction step where nitrate is reduced to nitrite and nitrite is further reduced to ammonium (Figures 7, 8). See section 1.3.2.1

1.4.2.2 Denitrification

Denitrification is the biological reduction of nitrate to nitrite, nitric oxide, nitrous oxide and finally to gaseous nitrogen oxide using an electron donor (Dworkin et al., 2006). There are four enzymes required for the generation of N₂ from nitrate, which entail the production of three intermediates: nitrite (NO₂⁻), nitric oxide (NO) and nitrous oxide (N₂O) (Figure 7). Therefore, denitrification involves the enzymes nitrate reductase (Nar), nitrite reductase (Nir), nitric oxide reductase (Nor) and nitrous oxide reductase (Nos). For instance, the genomes of the three *Rhodobacter sphaeroides* and five *Rhodopseudomonas palustris* strains available in literature clearly show the variability in the distribution of denitrification machinery among photosynthetic denitrifiers, the phylogenetic group with the largest number of denitrifiers being proteobacteria with denitrification being particularly prominent among α -proteobacteria (Dworkin et al., 2006).

1.4.2.3 Nitrogen Fixation

The large reservoir of nitrogen gas in our atmosphere is made biologically available by diazotrophic (N_2 -fixing) bacteria as a result of its ability to reduce dinitrogen to ammonium. The ability to fix dinitrogen is found only among representatives of two of the primary kingdoms of living organisms: prokaryotic archaea and bacteria. The reduction of nitrogen gas to ammonium requires 8 electrons as shown in equation [11] (Hunter et al., 2009).

$$N_2 + 8e^- + 8H^+ + 16 - 24 ATP \rightarrow 2NH_3 + 12H_2 + 16 - 24 ADP + 24 P_i$$
 [11]

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The reduction of N₂ to NH₃ is catalyzed by enzyme called nitrogenasea, and the genes required for nitrogen fixation are commonly called nif genes. Nitrogenases consist of two dissociable metalloproteins, dinitrogenase (e.g., NifDK) and dinitrogenase reductase (e.g., NifH). All N₂-fixing bacteria contain a Mo-nitrogenase, and in addition, some species possess one or two non-Mo nitrogenases (Hunter et al., 2009).

1.4.3 Biological phosphorus removal

Phosphorus has been considered a key element leading to eutrophication in aquatic habitats and also a vital element to life (Liang et al., 2010). A wide variety of species of microorganisms, including bacteria, yeasts, fungi, and microalgae, are capable of inorganic polyphosphate (polyP) accumulation above their structural levels (Hiraishi et al., 1991). The microbial synthesis of intracellular polyphosphate in PPB is primarily catalyzed by the enzyme polyphosphate kinase (PPK), which transfers the terminal phosphate of ATP to polyphosphate (Liang et al., 2010). Photosynthetic purple bacteria exhibit a very rich portfolio of membrane lipids including phospholipids not commonly found in bacteria (such as phosphatidylcholine and glycolipids typically found in plant chloroplasts such as sulfoquinovosyl diacylglycerol, betaine and ornithine lipids). These latter lipids lack phosphorus in their structure, presumably allowing PPB to outcompete other organisms in a phosphorus-depleted environment (Hunter et al., 2009).

1.4.4 Heavy metals removal

Similar than microalgae, the mechanisms utilized by PPB for the removal of heavy metals (HMs) are biosorption and bioaccumulation. Some PPB may use more than one mechanisms for the removal of metals. For instance, Panwichian et al., (2011) studied the potential of two PPB strains, *Rhodobium marinum* NW16 and *Rhodobacter sphaeroides* KMS24, to remove HMs from contaminated shrimp pond water. HM were detected in the sediment of shrimp ponds at 0.30 mM Pb²⁺, 0.89 mM Zn²⁺, 0.0067 mM Cd²⁺ and 0.54 mM Cu²⁺ (Panwichian et al., 2010). The removal efficiencies of Pb, Zn, Cd and Cu of the two PPB strains tested accounted for 97.29, 91.84, 90.79 and 90.52%, respectively. Likewise, *Rhodobium marinum* NW16 and *Rhodobacter sphaeroides* KMS24 had been used to remove heavy metals and salts from sediments and water collected from contaminated areas to decrease their phytotoxicity (Panwichian et al., 2012).

1.5 Influence of operational and environmental parameters on the activity of algal-bacterial consortia and PPB

1.5.1 pH

The design and operation of biological WWT processes must consider the optimum pH conditions required for microbial growth (Rittmann and McCarty, 2012). The pH of microalgae cultivation broth are typically neutral to alkaline (Rittmann and McCarty, 2012). This can be related to the availability of inorganic carbon in the form of bicarbonate, carbonate and carbon dioxide. When algae grow and extract carbon dioxide from water, pH tends to increase according the following equations [12] and [13]:

$$H_2CO_3 \to CH_2O + O_2 \tag{12}$$

 $HCO_3^- + H_2O \rightarrow CH_2O + O_2 + OH^-$ [13]

Where, the first reaction illustrates the fact that algal growth results in oxygen production and consumption of carbonic acid, thus causing the pH to rise, according to equation 9. The second reaction indicates that, additionally, if bicarbonates is the carbon source, the hydroxyl ions produced tend to increase the pH of the cultivation broth. On other hand, the bacterial release of CO_2 as a result of organic matter respiration and NH_4^+ nitrification maintain the pH below inhibitory levels. Values of pH between 8.5 and 9.0 have been shown to decrease algal growth, although, some microalgae species can grow at pH values as high as 10-11 (Rittman and Mccarty, 2012). In this regard, pH influences the biosorption capacity of biomass by modifying metal ion solubility and biosorbent total charge, since protons can be absorbed or released (Romera et al., 2007).

On the other hand, most species of bacteria exhibit the optimum range of growth at pH between 6 and 8. In addition, some bacteria, particularly chemolithotrophs that oxidize sulfur or iron for energy production, thrive best under highly acidic conditions. This characteristic enhances their chance for survival, since the end product of their energy metabolism is generally a strong acid (such as sulfuric acid) that prevents competition. PPB are a physiologically versatile group of purple bacteria that can grow well both phototrophically and in darkness. Photosynthesis in purple bacteria occurs at temperatures up to 57 °C and down to 0 °C, at pH values as low as 3 or as high as 11, and at salinities up ~32% NaCl (Hunter et al., 2009). Khatipov et al. (1998), who studied the accumulation of poly-β-hydroxybutyrate by the PPB *Rhodobacter sphaeroides* on various

carbon and nitrogen substrates, observed an increase in pH caused by a decrease in H_2 production and an increase in poly- β - hydroxybutyrate accumulation on lactate under nitrogen-deprived conditions.

1.5.2 Dissolved oxygen

 O_2 is consumed by heterotrophic microorganisms and produced by microalgae during WWT in a algal-bacterial processes (Oswald et al., 1957). According with Rittmann and Mccarty, (2012), 2 mg/L of dissolved oxygen is generally sufficient to maintain an active aerobic organic matter oxidation and nitrification. Hence, Wang et al., (2015) founded that low dissolved oxygen concentration (0.09 - 0.19 mgO₂/L) inhibited nitrite oxidizing bacteria, resulting in stable nitritation. On the other hand, higher dissolved oxygen concentration (< 20 mgO₂/L) combined with an intense irradiance could inhibit microalgae and bacterial activity due to photo-oxidation (Molina et al., 2001). In this context, Molina et al., (2001) determined that microalgae photosynthetic activity was determined as the rate of change in dissolved oxygen concentration. However, the dissolved oxygen concentration in the cultivation of microalgae-based WWT processes remains typically low as result of the active consumption by bacterial and heterotrophic microalgae.

On the other hand, many purple sulfur bacteria and purple non-sulfur bacteria are extremely sensitive to oxygen (Talaiekhozani and Rezania, 2017). Thus, oxygen partial pressure is a major factor regulating the formation of the photosynthetic apparatus and the cell differentiation of most facultative purple phototrophic bacteria capable of respiratory and photosynthetic modes of energy transduction. Two species forming the photosynthetic apparatus under both aerobic and anaerobic conditions are exceptions to this generalization: *Rhodopseudomonas sulfidophil* and Rhodospirillum *centenum* (Yurkov and Beatty, 1998). Thus, the competitive success of purple bacteria in nature requires light and anoxic conditions (Hunter et al., 2009). However, Meng et al., (2017), who worked with *Rhodopseudomonas* during synthetic wastewater treatment in batch tests, reported the highest COD (93%) and NH₃-N removal (83%) under DO of 4–8 mg/L.

1.5.3 Temperature

Temperature plays an important role in algal-bacterial processes devoted to WWT (Mobin and Alam, 2014). Temperatures influence the rate of all biochemical reactions involved in bacterial and microalgal growth, with higher rates observed when temperature increases. However, at very high temperatures, key enzymes are denaturalized and the organism may not survive. Despite some extremophile microalgae can grow at 0 °C and others at 90 °C, the optimal range for microalgae activity lies in 20-30 °C (Rittmann and McCarty, 2012)(Singh and Singh, 2015). Talbot et al., (1991) compared microalgae growth at different temperature and observed no significant difference between *P. bohneri* and *A. falcatus* at 25, 30 and 35 °C in batch test using an artificial medium. Ho et al., (2012) reported that 28 °C improved CO₂ fixation of indigenous *Scenedesmus obliquus* in a tubular glass photobioreactor of 1L. Studies carried out under outdoors conditions in two identical HRAPs with temperatures ranging between 11 °C in January and 25 °C in July reported Chlorophyll *a* concentrations higher (3.0 to 4.0 mg/L) during the spring and summer than during autumn and winter (1.0 to 2.0 mg/L) (García et al., 2000).

PPB are able to grow at temperatures ranging from 0 to 57 °C (Hunter et al., 2009). The temperatures ranges prevailing during wastewater treatment using purple phototrophic bacteria are similar to those recorded when using algal-bacterial systems. For instance, Kantachote et al., (2005) reported 30 °C as optimal temperature for cell growth during the treating rubber sheet wastewater enrichment with PPB (*Rhodopseudomonas blastica*). Likewise Madukasi et al., (2010) used a temperature between 20 and 30 °C during the treatment of pharmaceutical wastewater by a wild PPB strain named Z08 and identified as *Rhodobacter-sphaeroides* isolated from soil.

1.5.4 Photosynthetically active radiation

Light in the range 400 to 700 nm of the electromagnetic spectrum provides the energy for photosynthesis and plays an central role in the performance of algal-bacterial processes (Figure 9) (MacDonald, 2003). The solar radiation available for photosynthesis inside the cultivation broth of the microalgae based photobioreactor controls the rate of carbon fixation and therefore biomass productivity and WWT performance (Frouin and Pinker, 1995) According with Ogbonna and Tanaka (2000), light is the most critical factor in

photobioreactor and very often the factor limiting algal photosynthesis. Cheloni et al., (2014) showed the important role of light intensity and spectral composition on Cu uptake and the effects on the green alga *Chlamydomonas reinhardtii*. High irradiations increased cellular Cu concentrations, but mitigated the Cu-induced decrease in chlorophyll fluorescence, oxidative stress and lipid peroxidation at high Cu concentrations, indicating that Cu and high irradiations interact in an antagonistic manner.



Figure 9. The electromagnetic spectrum (source: http://schoolbag.info/physics/physics_math)

During photosynthesis (Figure 10), inorganic compounds and light energy are converted into organic matter by photoautotrophs (Richmond, 2004). Microorganisms obtain their energy for growth and maintenance from Redox reactions triggered by the photon captured from solar light. Microalgae use oxidation-reduction reactions to convert the light energy into ATP and NAD (Rittman and Mccarty, 2012).



Figure 10. Products of the light and dark reactions of photosynthesis (adapted from Hall & Rao, 1999). Source (Richmond, 2004).

According with Cogdell and Thornber, (1980) PPB can absorb radiation in the wavelength range between 800 and 900 nm due to the ability of the harvesting complexes present in the bacteriochlorophyll (Figure 9). All PPBs have in-vivo bacteriochlorophyll and carotenoids that carry out photosynthesis without oxygen production and their photosynthesis rates depends not only on the infrared light intensity but on the supply of external electron donors such as sulfide, molecular hydrogen or organic substances (Kantachote et al., 2005; Madukasi et al., 2011). Bertling et al., (2006) demonstrated the feasibility of PPB growth supported by a Laser Diode (in infrared spectrum) irradiating

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at 30 W/m². Likewise, Hülsen et al., (2013) used infrared selected PPB as a platform for the biological removal of carbon, nitrogen and phosphorous under anaerobic conditions with removal efficiencies of COD of $63 \pm 5\%$, NH₄⁺-N of 99.6% and PO₄³⁻-P of 88% from primary settled domestic wastewater in 24 h.

1.5.5 Evaporation rates

Evaporation is an important parameter during the wastewater treatment in open photobioreactors based on its pollutant concentration effect. In this context, Guieysse et al., (2013) have highlighted that evaporation has a critical impact on the economics and sustainability of algae mass cultivation in open systems. The high evaporation rates in open photobioreactors treating fish processing wastewater exerted a negative impact on the biomass growth (Riaño et al., 2012). The main factors impacting on water evaporation in photobioreactors are solar radiation, temperature, relative humidity, vapour pressure deficit, atmospheric pressure and wind (Kumar et al., 2016). Posadas et al., (2015) reported evaporation losses ranging from 4.4 ± 1.4 to 7.3 ± 0.2 L m⁻².d⁻¹ as a result of the operational conditions in a 180 L HRAP. The main cause of the high water evaporation losses was the high turbulence as a consequence of its pilot scale design, where the paddlewheel engine was oversized.

1.5.6 Hydraulic retention times

HRAP are typically operated at 2-6 days HRT and similar values have been reported in enclosed photobioreactors (Muñoz and Guieysse, 2006). Aguirre et al., (2011) set 40 days of HRT during the treatment of PWW with a COD of 3000 mgO₂/L using and HRAP of 400L. However, de Godos et al., (2009a) operated with 10 days of HRT in a 464L HRAP treating 20 and 10 folds diluted swine manure.

1.5.7 C/N/P ratio

The most important limitation of biological processes during urban and livestock wastewater treatment is the low C:N:P ratio of these types of wastewaters (Arbib et al., 2013), which ideally should match with the typical biomass composition represented by the Redfield ratio C:N:P: 106/16/1 (Richmond, 2004). Most research treating domestic, industrial, agroindustrial and livestock effluents in microalgae-based processes were operated under carbon limitation (Arbib et al., 2013).

1.6 Photobioreactors

Photobioreactors have emerged as an alternative platform to treat wastewater using algalbacterial consortia or phototrophic purple bacteria. Photobioreactors used for wastewater treatment have similar design and operational criteria than conventional photobioreactor used for mass cultivation. For instance, a high surface/volume ratio to maximize light utilization efficiency (and therefore oxygen production), adequate mixing and degassing, good scalability, low hydrodynamic stress on the algal-bacterial flocs, control over the environmental conditions and low construction and operation costs rank among the top characteristics desired in photobioreactors devoted to WWT (Muñoz and Guieysse, 2006; Richmond, 2004). However, the most import issue to design a photobioreactor are related to light supply, which represents the most costly parameter to optimize (Adessi and De Philippis, 2014).

1.6.1 Closed photobioreactors

Closed photobioreactors are characterized by the regulation and control of nearly all the environmental parameters as well as by the following fundamental benefits: a reduced contamination risk, no CO₂ losses, reproducible cultivation conditions, controllable hydrodynamics and temperature, and a flexible technical design (Pulz, 2001). Closed photobioreactors support higher photosynthetic efficiencies as a result of their higher illuminated area to volume ratio and a better control on operational parameters than open systems (Muñoz and Guieysse, 2006). The main disadvantages of these photobioreactors derive from the high operational cost to maintain the optimal temperature to avoid a significant productivity drop (Béchet et al., 2010). Closed photobioreactors are currently used for microalgae mass cultivation in the following configurations: tubular (Figure 11) and flat-plate (Figure 12). Tubular photobioreactors (vertical, horizontal and helical) are the only type of closed systems used at large scale. The optimum diameter of the tubes in TPBRs range from 3 to 12 cm since smaller diameters could cause biomass clogging, while the recommended linear velocity in the tubes ranges from 20 to 50 cm \cdot s⁻¹, which is normally controlled by the use of airlift systems or centrifugal pumps (Christenson and Sims, 2011) (González-Fernández and Muñoz, 2017).





Figure 11. Tubular photobioreactor configurations (Credit: Image: Photobioreactor PBR 4000 G IGV Biotech)

Figure 12. Vertical-plate photobioreactor (Credit: Image: Photobioreactor PBR 500 P IGV Biotech)

Flat-plate photobioreactors provide an optimized light distribution. In addition, their simpler construction compared to tubular reactors allows the use of less expensive plastic materials. Flat plate systems are characterized by an open gas transfer area, thus reducing the need for a dedicated degassing unit (Dasgupta et al., 2010)(Carvalho and Meireles, 2006).

1.6.2 Open photobioreactors

High rate algal ponds, also named raceway ponds, are most common open photobioreactors, whose design and operational parameters depend on the influent characteristics and environmental conditions such as temperature, irradiance and evaporation losses (Alcántara et al., 2015c; Muñoz and Guieysse, 2006). HRAPs have been used since 1950's and are typically constructed based on a design criterion of 11 m^2 per population equivalent (Oswald et al., 1957)(European Union Commission (2001). HRAPs are 2-3 m wide and 0.1- 0.3 m depth shallow open ponds built in a raceway configuration, lined with PVC, clay or asphalt to avoid infiltration. The surface of these photobioreactors range from 1000 to 10000 m² in large-scale applications (Muñoz and Guieysse, 2006). Three open photobioreactor configurations have been up-scaled: raceways ponds, circular ponds and thin layer systems. The latter exhibit limitations such as the sedimentation of cells at points of lower turbulence, strong evaporative losses, high rates of CO₂ desorption and considerable requirement of energy for continuously pumping the culture to the head of the inclined surface (Richmond, 2004). Figures 13 and 14 depicts the most common open photobiorector configurations used for wastewater treatment.





Figure 6. HRAP with a thick dividing section. In this picture, the water flows in a clockwise direction powered by two paddle wheels. (Credit: SARDI: South Australian Research and Development Institute)

Figure 14. Circular raceway for microalgae cultivation of 5 m^2 . (Source: Wang et al., 2018)

The main disadvantage of these photobioreactor configuration is the large treatment area needed to complete phosphorus removal compared to the area needed to remove of carbon and nitrogen (Alcántara et al., 2015c). Under optimal conditions, HRAPs can treat up to $35 \text{ g} \cdot \text{BOD} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$ (175 g $\cdot \text{BOD} \cdot \text{m}^{-3} \text{d}^{-1}$ in a 0.2 m deep pond) compared to 5–10 g $\cdot \text{BOD} \cdot \text{m}^{-2} \cdot \text{d}^{-1}$ (5-10 g BOD $\cdot \text{m}^{-3} \text{d}^{-1}$ in a 1 m deep pond) in waste stabilization ponds (Muñoz and Guieysse, 2006; Racault and Boutin, 2005).

1.7 Biomass valorization

Biomass from domestic and agroindustrial wastewater treatment can be used as a feedstock to produce biofertilizers based on its high carbon and nutrients concentration. Indeed, Cabanelas et al., (2013) reported contents of 40-60%, 4-7% and 0.1-1% of carbon, nitrogen and phosphorus, respectively in biomass grown in a photobioreactor inoculated with *Chlorella vulgaris* and supplemented with CO₂ during the treatment of effluent from primary settler. In this context, limitation in nitrogen or phosphorus could result in an increase in the carbon content of the harvested biomass (mainly composed of proteins (40%-60%), lipids (5%-60%) and carbohydrates (8%-30%)) (Markou et al., 2014)(Tijani et al., 2015). Algae-based fuels or biofuels offer many advantages compared to conventional fossil fuels (Balat, 2011). Production of bioethanol from algal biomass is one alternative to reduce the demand of fuels from petroleum and its associated environmental pollution (Balat, 2011). Bioethanol has received in the past decades an increasing attention worldwide. However, the cost of bioethanol production is still higher

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than that of fossil fuels (Sarkar et al., 2012). Biodiesel production is another alternative algae-based fuel which can be used as substitute or additive of conventional diesel (Vivek and Gupta, 2004). However, the lipid content of biomass grown in WWT plants is usually low. In this context, Álvarez-Díaz et al., (2015) increased the lipid (from 35.8% to 49%) content of ω -3 eicosapentaenoic acid in Scenedesmus obliquus treating domestic wastewater, which opened the door to develop wastewater-to-biodiesel strategies during microalgae-based WWT. Biohydrogen is another biofuel produced from or by microalgae (Chandrasekhar et al., 2015). Biological production of hydrogen (Biohydrogen) using microalgae as a feedstock or biocatalyst include direct biophotolysis, indirect biophotolysis, photo-fermentations, dark-fermentation, microbial electrolysis cell, multistage integrated process and water-gas shift (Levin et al., 2004) (Holladay et al., 2009). Finally, microalgae can be converted into biogas via anaerobic digestion (Converti et al., 2009). In this context, the biogas produced from algal biomass contain a high-energy value and the energy recovery is comparable to that of the extraction from cell lipids. Due to its simplicity and extensive experience in design and operation of anaerobic digesters, biogas is becoming the most popular biofuel produced from microalgae (Wayne Chew et al., 2017).

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Aims and Scope
Aims and Scope

2.1 Justification

During the past decades, the intense growth of human population has generated large amounts of domestic and livestock wastewaters, which must be treated before discharge into water bodies. In this context, multiple biological and physical/chemical technologies have been engineered to treat wastewater. Algal-bacterial processes have received an increasing attention as a cost-efficient platform for water reclamation as a result of their low-cost oxygenation, simplicity, nutrient recovery potential, carbon dioxide mitigation potential and production of a valuable feedstock biomass. This solar driven technology is based on the combination of heterotrophic, mixotrophic and phototrophic metabolisms and UV photolysis, and can be coupled with biogas upgrading. However, this green technology only exhibits consistent C, N and P removal efficiencies when treating effluents with high C/N/P ratios ($\approx 100/18/1$), which are not typically encountered in domestic and livestock wastewaters. Despite the advantages of algal-bacterial processes in wastewater treatment, more research is needed to optimize wastewater treatment and to guarantee a consistent biomass composition in order to support the development of algae-based biorefineries for the production of biofuels, biofertilizers or high value bioproducts. On the other hand, purple photosynthetic bacteria (PPB) have emerged as an alternative photosynthetic platform for wastewater treatment based on their superior resource recovery potential, higher tolerance to organic pollution, lower dependence to temperature and more versatile metabolism. In brief, microalgae and PPB based wastewater treatment has the potential to provide a simple, low-cost and environmentally friendly solution for wastewater sanitation in developing countries, where environmental and social conditions are more favorable to these solar based technologies and the acceptance of the bioproducts generated.

This thesis aimed at tackling the main problems hindering the widespread implementation of algal-bacterial photobioreactors during wastewater treatment: i) the low concentration of inorganic carbon in domestic wastewater, ii) the lack of studies confirming the potential of denitrification-nitrification of algal-bacterial photobioreactors using real domestic wastewater, iii) the poor knowledge of the dynamics of the microalgae population structure in algal-bacterial photobioreactors, iv) the representativeness of the results obtained indoors and v) the lack of alternatives to microalgae as a platform for photosynthetic wastewater treatment.

2.2 **Objectives**

The overall goal of the present thesis was the assessment of the potential of photosynthetic treatment of domestic and livestock wastewater using algal-bacterial consortia and purple photosynthetic bacteria. The work conducted was focused on evaluating the potential of novel bioreactor configurations and photosynthetic microorganisms, and providing insights on the long-term feasibility of alga biorefineries based on wastewater treatment. More specifically:

Objective 1. Systematic evaluation of the performance of novel photobioreactor configurations capable of overcoming the limitations of conventional High Rate Algae Ponds during domestic wastewater treatment.

Objective 2. Evaluation of the long-term dominance of specific microalgae and cyanobacteria during piggery wastewater treatment.

Objective 3. Evaluation of relevance of the results obtained indoors under artificial irradiation.

Objective 4. A comparative evaluation of the performance of algal-bacterial consortia and purple photosynthetic bacteria.

Aims and Scope

2.3 Development of the thesis

In order to fulfill the specific objectives above stated, four series of experiments were conducted over the past 4 years:

The performance of an innovative anoxic-aerobic photobioreactor coupled with biogas upgrading and operated with biomass recycling was assessed (**Chapter 1**).

In this context:

- ✓ This work was devised to treat urban domestic wastewater with low inorganic carbon concentration at short hydraulic retention times.
- Biogas addition was used to overcome inorganic carbon limitation and promote a stable nitrification-denitrification process.
- ✓ The process incorporated a biomass settling step followed by recycling to the anoxic tank to promote a total denitrification and improve the biomass settling characteristics.

The long-term dynamics of the structure of microalgae population and wastewater treatment performance was evaluated during piggery wastewater treatment in open photobioreactors (**Chapter 2**).

- This study was carried out to estimate microalgae production at large-scale in the context of microalgae-based biorefineries.
- Piggery wastewater was used as a nutrient source due to the high organic matter and nutrients concentrations, able to support high biomass productivities.
- ✓ The dynamics of microalgae population structure were evaluated in four photobioreactors under similar operational conditions and inoculated with two microalgae species, a cyanobacterium and compared with a system not inoculated.

Piggery wastewater (PWW) treatment in open algal-bacterial photobioreactors was also comparatively evaluated under indoor and outdoor conditions (Chapter 3).

✓ A comparative study was carried to evaluate microalgae evolution and process performance in both operational conditions, since most studies have been conducted under laboratory conditions. ✓ Two PWW dilutions were used in order to evaluate the tolerance to organic pollution and its influence on microalgae dominance during PWW treatment.

Finally, the potential of purple photosynthetic bacteria for piggery wastewater treatment under infrared illumination was comparatively tested against that of an algal-bacterial consortium under LED illumination in open photobioreactors under indoor conditions (**Chapter 4**).

- This study was conducted to elucidate the best photosynthetic platform during PWW treatment.
- ✓ The performance of both microbial groups at different hydraulic retention times was evaluated both in term of biomass productivity, effluent quality and potential for nutrient recovery.

Chapter 1

 Enhanced carbon, nitrogen and phosphorus removal from domestic wastewater in a novel anoxic-aerobic photobioreactor coupled with biogas upgrading García, D., Alcántara, C., Blanco, S., Pérez, R., Bolado, S., Muñoz, R. 2017. Chemical Engineering Journal. 313, 424 - 434. doi:10.1016/j.cej.2016.12.054. Accepted 13 December 2016



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Abstract

This work evaluated the performance of an innovative anoxic-aerobic algalbacterial photobioreactor coupled with biogass upgrading for the treatment of domestic wastewater via nitrificationdenitrification. The process, which incorporated a biomass settling step followed by recycling to the anoxic tank, was operated at a hydraulic retention time of 2 days, a sludge retention time of \approx 11 days under a 12h/12h light/dark irradiation cycle at 392 μ E m⁻²·s⁻¹. An increase in the removal efficiency of TN from 38% to 81%, NH₄⁺ from 39% to 97%, and P-PO $_4^{3-}$ from 59% to 64% were recorded when additional CO₂ was supplied to the photobioreactor via biogas scrubbing, which supported an almost complete nitrification of the NH4⁺ to NO_3^- and promoted microalgae subsequent growth (with the

enhancement in N and P assimilation). TOC removal remained constant at $90\pm2\%$ regardless of the addition of CO₂, while the effluent biomass concentration averaged 26±12 mgTSS/L. A DGGEsequencing analysis of the bacterial community revealed the occurrence of 10 phyla, Proteobacteria being the dominant phylum. Finally, the morphological characterization of the microalgae population dynamics revealed a gradual dominance of the genus Scenedesmus, which accounted for 94-100 % at the end of the experiment.

Keywords

Biogas upgrading; Biomass recycling; Inorganic carbon limitation; Microalgae; Nitrification-denitrification.

1. Introduction

The steady increase in human population [1] and industrial activity is generating large amounts of wastewaters and greenhouse gases [2], which represent two of the major environmental challenges sustainability to global nowadays. Domestic and industrial wastewaters and anaerobic digestion effluents are characterized by their high loads of carbon (C), nitrogen (N) and phosphorus (P), which must be treated before discharge into natural water bodies to avoid oxygen depletion, toxicity issues and eutrophication [3]. A wide range of biological and physical/chemical technologies is currently available for carbon and nutrient removal in wastewater treatment plants (WWTPs). Unfortunately, these technologies often entail high investment and operating costs and do not allow for a cost-effective recovery of nutrients due to the low C/N and C/P ratios of most domestic and industrial wastewaters [4][5].

In this context, algal-bacterial processes can support both a low-cost process oxygenation and an enhanced nutrient recovery. The oxygen produced by microalgae during photosynthesis can support the oxidation of organic pollutants and ammonium by aerobic heterotrophs and nitrifiers, respectively, which thus reduces the operating costs and environmental impacts associated with conventional mechanical aeration in WWTPs [6]. On the other hand, the ability of algal-bacterial consortia to assimilate both organic carbon (inherently present in most wastewaters) inorganic carbon (from and the biological oxidation of organic carbon, alkalinity in wastewater or residual carbon dioxide (CO_2) externally supplied) result in larger biomass productivities and therefore enhanced nutrient recoveries[7]. However, despite the above-mentioned advantages, algalbacterial devoted processes to wastewater treatment still present severe technical limitations that hinder the fullscale implementation of this technology, such as nutrient supply and recycling, gas transfer and exchange [8].

In this regard, although photoautotrophic algal metabolism can enhance N and P recovery in anoxic-aerobic algalbacterial photobioreactor (AA-ABPh), the alkalinity present in raw wastewaters is low to support a complete nutrient recovery/removal and residual CO2 sources (such as flue gas) are not always available on-site. In addition, the low hydraulic retention times (HRT)

required in algal-bacterial processes to compete with activated sludge systems would limit the development of nitrifying bacterial communities that would eventually support nitrificationdenitrification processes during the treatment of wastewaters with low C/N ratios. Finally, the poor sedimentation ability of the microalgae generated in the process often results in effluent total suspended solid concentrations (TSS) above the maximum European Union (EU) discharge limit (50 mg/L), which limits the scale-up of microalgae-based wastewater treatment [9]. In this context, AA-ABPh operated with autofloculated biomass settling and recycling constitutes an innovative technology capable of overcoming the above mentioned limitations. This technology was successfully evaluated for the treatment of synthetic wastewaters at moderate HRTs but experienced severe inorganic carbon limitations, which ultimately restricted the treatment potential of this innovative technique. Therefore, there is an urgent need to develop novel operating strategies to overcome above mentioned the inorganic carbon limitation and to evaluate the performance of this innovative technology using real domestic wastewater (RDWW) at low HRTs.

This work was devised to evaluate the treatment of RDWW in an innovative AA-ABPh configuration operated with biomass settling and recycling at a HRT of 2 days coupled to the simultaneous upgrading of synthetic biogas (in a separate and interconnected column). In this system, the supply of biogas (eventually available on-site from the anaerobic digestion of the algal-bacterial biomass produced in the process) will provide the additional inorganic C source required to boost nutrient removal by assimilation and bacterial nitrification to sustain an efficient nitrificationdenitrification process [10][11]. The influence of photosynthetic biogas the mechanisms upgrading on underlying C, N and P removal in the anoxic tank and photobioreactor treating RDWW was assessed using a mass А balance approach. detailed characterization of the dynamics of microalgal and bacterial population structure conducted was using morphological and molecular identification tools.

2. Materials and methods

2.1 Microorganisms and culture conditions

The anoxic and aerobic tanks were inoculated with 3.2 g/L of total

suspended solid (TSS) of an mixture of a microalgal-cyanobacterial consortium (from now on referred to as microalgae) from a high rate algal ponds (HRAP) treating diluted vinasse [12] and aerobic activated sludge from Valladolid WWTP (Spain). Domestic wastewater was collected from a nearby sewer located at Department of Chemical Engineering and Environmental Technology of Valladolid University. The average composition of the RDWW treated continuously was: 176±26 mg/L of dissolved total organic carbon (TOC), 152±34 mg/L of dissolved inorganic carbon (IC), 106±9 mg/L of total nitrogen (TN), 93±9 mg/L of Nammonium (N-NH₄⁺), 39 ± 12 mg/L of sulfate (SO₄²⁻) and 33±8 mg/L of Pphosphate ($P-PO_4^{3-}$) (Table 1).

2.2 Experimental set-up

Three operational stages (SI, SII and SIII) were carried out in the experimental set-up below described to evaluate the influence of biogas supply and internal recycling rate on WWT performance. The experimental set-up consisted of an anoxic tank, with a total working volume of 0.9 L. interconnected to а photobioreactor with a total working volume of 2.7 L. Both reactors were operated as completely mixed flow reactors. The anoxic tank was

maintained in the dark and magnetically stirred at 320 rpm. The photobioreactor was continuously illuminated for 12 hours/day (05:00h to 17:00h) by LED in lamps arranged а concentric configuration providing an average light intensity of $392\pm19 \ \mu mol/m^2 \cdot s$ at the outer wall of the photobioreactor (Figure 1). Air was introduced during the dark period in order to maintain the dissolved O₂ concentration in the photobioreactor above non-limiting concentrations [13][14] (Table 1). The temperature in photobioreactor the jacketed was maintained constant at 25±1°C using a recirculating water stream cooled down in a water bath (Huber, Germany). The temperature in the photobioreactor was controlled to systematically assess the influence of biogas supply and internal recirculation flow rate on process performance without the interference of any other varying parameter. In this context. the 12h/12hlight/dark illumination regime would entail an increase in process temperature (as a result of the heat dissipated from the LEDs) during the illuminated period that might interfere with the activity of the most sensitive microbial communities (i.e nitrifiers). The temperature in the anoxic tank was not controlled but remained constant at 25 °C as a result of the constant room temperate and high

internal recycling flow rate from the photobioreactor. Likewise, magnetic agitation in the photobioreactor was maintained constant and 320 rpm. During stage II (SII) and stage III (SIII), the photobioreactor was interconnected to a 0.3 L external absorption column (\emptyset = 2.54 cm; height = 60 cm) in order to provide an extra source of CO2 via photosynthetic biogas upgrading (Figure 1). The use of a separate biogas absorption column supported both a higher CO₂ gas-liquid mass transport and a lower O₂ stripping. Hence, the direct

scrubbing of biogas in the typically shallow photobioreactors would entail a low volumetric CO₂ gas-liquid mass transfer coefficient, while an external absorption column can be designed independently of the light path (depth) of the photobioreactor. On the other hand, process operation with an external absorption column interconnected to the HRAP with a recycling cultivation broth stream allows a control of the oxygen potentially stripped out to the upgraded biogas.



Figure 1. Schematic diagram of the anoxic-aerobic algal-bacterial photobioreactor set-up coupled with an absorption column for CO2 supplementation via biogas upgrading.

2.3 Experimental design

The HRT and the Sludge Retention Time (SRT) were maintained at 2 days

(HRT_{anoxic} = 0.5 day, HRT_{aerobic} = 1.5 days) and \approx 11 days, respectively, during the whole experiment (Table 1). These

short HRTs are required to make algalbacterial processes competitive with activated sludge systems. The experimental design was based on the operational limitations identified in previous studies in this AA-ABPh configuration [13][14]. The experiment was divided in three operational stages.

The internal recycling (IR) flow rate from the photobioreactor to the anoxic tank was maintained at 2.8 L/d during SI (maintained for 78 days) and SII (maintained for 74 days), while this parameter was increased by 30% (3.6 L/d) during SIII (maintained for 56 days) in order to evaluate the maximum denitrifying capacity in the anoxic tank to ultimately boost the dissimilatory N removal in the experimental system. The external recycling flow rate (ER) from the settler to the anoxic tank was maintained at 0.5 L/d during the three operational stages (Table 1). The pH in the photobioreactor was maintained between 7.0 and 8.6 during SI by daily addition of 1.2 mL of chlorhydric acid (37%), while biogas upgrading into the absorption column supplied CO₂ to overcome the IC limitation encountered during SI and to maintain the pH below inhibitory values for bacterial activity (< 9) during SII and SIII. The synthetic biogas mixture supplied was composed

of methane (70%), carbon dioxide (29.5%) and hydrogen sulfide (0.5%)(Abello Linde, Barcelona, Spain). Biogas was supplied to the absorption column at 2.6 L/d (1.8 ml/min) through a 10 µm metallic sparger located at the bottom of the column co-currently with a recycling algal-bacterial broth stream drawn from the photobioreactor at a liquid to biogas ratio (L/G) of 10 (v/v)(Figure 1). Liquid samples of 100 ml were taken twice per week from the RDWW, anoxic tank, aerobic tank, settled biomass and effluent to determine concentrations of TOC, IC, TN, N-NH₄⁺, N-NO₂⁻, N-NO₃⁻, SO₄²⁻, P-PO₄³⁻ and TSS. pH, temperature and dissolved concentration (DO)oxygen were measured daily in both bioreactors. Likewise, the C, N and P content of the algal bacterial biomass was measured under steady state at each operational stage. The sludge volumetric index (SVI) and the maximum biomass settling rate, which were used to monitor the settling characteristics of the algal-bacterial biomass [15], were determined in the anoxic and aerobic bioreactors under steady state at each operational stage. The microalgae population structure was assessed at the end of SI, SII and SIII using biomass samples from the photobioreactor preserved with lugol acid at 5% and formaldehyde at 10%,

and stored at 4 °C prior to analysis. Biomass samples from the photobioreactor were collected at the end of each steady state, and immediately stored at -20 °C in order to evaluate the richness and composition of the bacterial communities [14]. Finally, the composition of the biogas at the inlet and outlet of the absorption column was determined twice a week during SII and SIII.

2.4 Mass balance

The dilution effect in the anoxic tank caused by the internal and external recirculations was considered bv calculating a virtual concentration for parameter in the influent each wastewater to the anoxic tank. Hence, the actual C, N and P removals in the denitrification reactor were assessed using the virtual concentrations (Vi) for dissolved IC, TOC, N-NH4⁺ and P-PO4³⁻ at the entrance of the anoxic tank according to Eq. (1):

$$V_i\left(\frac{mg}{L}\right) = \frac{\left(C_{i\,feed} \cdot Q_{feed}\right) + \left(C_{i\,aerobic} \cdot Q_{RI}\right) + \left(C_{i\,aerobic} \cdot Q_{RE}\right)}{Q_{feed} + Q_{RI} + Q_{RE}} \tag{1}$$

where C_i feed and C_i aerobic represent the dissolved concentrations of the parameter "*i*"= TOC, IC, TN, N-NH₄⁺ and P-PO₄³⁻ in the RWW and the photobioreactor, respectively, while Q_{feed} represents the RDWW flow rate, Q_{RI} the internal recirculation flow rate and Q_{RE} the external recirculation flow rate. An overall mass balance to the anoxicaerobic photobioreactor was conducted for TOC, IC, TN, N-NH₄⁺ and P-PO₄³⁻ based on their average concentrations under steady state conditions for each operational stage. The validity of the instrumental and analytical methods was thus assessed by means of the mass recovery factors for the parameter "*i*" according to Eq. (2) [14]:

$$M_{i} Recovery (\%) = \frac{(M_{irem})anox + (M_{irem})photobior + (M_{i})effl}{(M_{i})RWW} \cdot 100$$
(2)

where $(M_i \text{ rem})_{anoxic}$ and $(M_i \text{ rem})_{photobior}$ represent the mass flow rate (g/d) of the parameter i = TOC, IC, TN, N-NH4⁺ and P-PO4⁻³ removed in the anoxic tank and photobioreactor, respectively. M_i effl and M_{iRWW} represent the mass flow rate (g/d) of the parameter in the treated effluent and RWW, respectively.

The removal efficiencies herein reported for each tank refer to the individual contribution of the anoxic and photobioreactor units to the overall removal of the inlet loading for each monitored parameter.

2.5 Analytical procedures

The light intensity was measured as photosynthetically active radiation (PAR) using a LI-250A light meter (LI-COR Biosciences, Germany). Biogas composition was determined using a Bruker 430 GC-TCD (Palo Alto, USA) equipped with a CP-Molsieve 5A (15 m \times 0.53 mm \times 15 μ m) and a CP-Pora BOND Q (25 m \times 0.53 mm \times 15 µm) columns. The injector, detector and oven temperatures were maintained at 150 °C, 175 °C and 40 °C, respectively. Helium was used as the carrier gas at 13.7 $cm^{3}/min[16].$ TOC, IC and TN concentrations were measured using a TOC-V CSH analyzer equipped with a TNM-1 module (Shimadzu, Japan). N-NH4⁺ was measured using the Nessler analytical method [15] in a U-2000 spectrophotometer (Hitachi, Japan), while NO_2^- and NO_3^- were determined by the cadmium reduction column method [15]. $P-PO_4^{-3}$ and SO_4^{2-} were analyzed by high performance liquid chromatography-ion chromatography (HPLC-IC) with a Waters 515 HPLC

pump coupled with a Waters 432 ionic conductivity detector and equipped with an IC-Pak Anion HC (150 mm \times 4.6 mm) waters column. A 510 pH meter (EUTECH Instrument, The Netherlands) was used for pH determination. DO concentration and temperature were recorded using an OXI 330i oximeter (WTW, Germany). The determination of the TSS concentration, SVI and settling rate were conducted according to Standard Methods [15]. The analysis of the C and N biomass content was carried out using a LECO CHNS-932 elemental analyzer with pre-dried and grinded algal-bacterial biomass. The content of P in the biomass was measured using a 725-ICP Optical Emission Spectrophotometer (Agilent, USA) at 213.62 The nm. identification. quantification and biometry measurements of microalgae were conducted by microscopic examination (OLYMPUS IX70, USA) of the algalbacterial cultivation broths according to Phytoplankton Manual [17].

Genomic DNA was extracted using the protocol described in the Fast® DNA Spin Kit for Soil (MP Biomedicals, LLC) handbook. The V6-V8 regions of the bacterial 16S ribosomal ribonucleic acid (rRNA) genes were amplified by Polymerase Chain Reaction (PCR)

analysis using the universal bacterial primers 968-F-GC and 1401-R (Sigma-Aldrich, St. Louis, MO, USA; [18]). The PCR mixture contained 1 µL of each primer (10 ng µL-1 each primer), 25 µL of BIOMIX ready-to-use 2 reaction mix (Bioline, Ecogen), $2 \mu L$ of the extracted DNA and Milli-Q water up to a final volume of 50 µL. The PCR thermocycling program consisted of 2 min of pre-denaturation at 95°C, 35 cycles of denaturation at 95°C for 30 s, annealing at 56°C for 45 s, and elongation at 72°C for 1 min, with a final 5-min elongation at 72°C. The denaturing gradient gel electrophoresis (DGGE) analysis of the amplicons was performed with a D-Code universal mutation system (Bio Rad Laboratories) using 8% (w/v)with polyacrylamide gels a urea/formamide denaturing gradient of 45 to 65%. DGGE running conditions were applied according to Roest et al. (2005) [19]. The gels were stained with GelRed Nucleic Acid Gel Stain (biotium) for 1 h. The most relevant bands were excised from the DGGE gel in order to identify the bacteria present in the samples, resuspended in 50 μ L of ultrapure water and maintained at 60 °C for 1hour to allow DNA extraction from the gel. A volume of 5 μ L of the supernatant was used for reamplification with the original primer set. Before

sequencing, PCR products were purified with the GenElute PCR DNA Purification Kit (Sigma-Aldrich, St. Louis, MO, USA). DGGE profiles were compared using the GelCompar IITM software (Applied Maths BVBA, Sint-Martens-Latem, Belgium). After image normalization, bands were defined for each sample using the bands search algorithm within the program. The peak heights in the densitometric curves were also used to determine the Shannon-Wiener diversity index (H). Therefore, this index reflects both the sample richness (relative number of DGGE bands) and evenness (relative intensity of every band). It ranges from 1.5 to 3.5 (low and high species evenness and richness, respectively) and can be calculated according to Eq. (3)[20]:

 $H = -\Sigma \left[P_i ln(P_i) \right] \tag{3}$

Where H is diversity index and P_i is the importance probability of the bands in a lane ($P_i = n_i/n$, where n_i is the height of an individual peak and n is the sum of all peak heights in the densitometric curves). Similarity indices of the compared profiles were calculated from the densitometric curves of the scanned DGGE profiles by using the Pearson product–moment correlation coefficient [21].The taxonomic position of the

sequenced DGGE bands was obtained using the RDP classifier tool (50% confidence level) [22]. The closest cultured and uncultured relatives to each band were obtained using the BLAST search tool at the NCBI (National Centre for Biotechnology Information) [23]. The sequences generated from this work are deposited in GenBank under accession numbers KU854389-KU854421.

2.6 Statistical analysis

The data displayed in Table 1, Figure 2 and Figure 7(c) correspond to the mean \pm standard deviation of the target parameters during steady state at each operational stage. A one-way ANOVA analysis was performed to assess any significant difference between the settling rate of the biomass from the anoxic reactor and the photobioreactor using Excel (Microsoft, USA). A Pearson correlation analysis was conducted to determine the similarity indexes the population among established during steady state operation.

Stage Parameter / Reactor		Wastewater	SI		SII		SIII		
			Anoxic	Aerobic	Anoxic	Aerobic	Anoxic	Aerobic	
Operational period (days)		n.a	78		74		56		
HRT (days) n.a		n.a	0.5	1.5	0.5	1.5	0,5	1.5	
SRT (days)		n.a	12.5 ± 3.5		11 ± 0.9		10.5 ± 0.5		
Light (µmol/m ² .s) n.a		n.a	n.a	367±57	n.a	412±15	n.a	395±21	
RWW feeding rate (L/d)		n.a	1.8		1.8		1.8		
Internal recycling rate (L/d)		n.a	2.8		2.8		3.6		
External recycling rate (L/d)		n.a	0.5			0.5		0.5	
pH (units)	Light	n.a	7.4±0.3	8.6±0.6	6.9±0.1	8.0±0.7	6.8±0.2	8.9±0.9	
	Dark	n.a		7.1±0.3		7±0.2		7.0±0.4	
Dissolved oxygen	Light	n.a	0	23.3±4.1	0	22.0±2.0	0	23.0±1.8	
(mg/L)	Dark	n.a		6.0±0.6		3.1±1.2		3.9±0.7	
TOC (mg/L)		176±26	26±5	20±4	30±4	19±1	23±2	18±2	
IC (mg/L)		152±34	48±5	2±1	56±9	15±5	47±4	13±4	
TN (mg/L)		106±9	74±10	66±8	32±4	21±3	28±2	18±4	
$N-NH_4^+$ (mg/L)		93±9	64±4	54±8	32±5	3±1	33±2	3±1	
$N-NO_2^-$ (mg/L)		n.a	0.01 ± 0.01	5.6±4.0	0.03±0.05	3.1±3.8	0.41±0.68	$1.1{\pm}1.8$	
$N-NO_3$ (mg/L)		n.a	0.04±0.03	0.9±0.9	0.14±0.15	8.9±5.5	0.73±1.25	13.0±3.2	
$P-PO_4^{3-}$ (mg/L)		33±8	26±5	16±7	18±3	9±2	18±1	11±2	
SO ₄ ²⁻		39±12	29±7	32±6	51±15	76±11	47±7	55±4	
TSS (mg/L)		n.a	1519±252	1216±260	3113±361	2854±324	2480±309	2047±186	
SVI (mL/g)		n.a	95	161	128	169	80	97	
Air flow (mL/min)		n.a	0	6	0	4	0	2	
Biogas (L/day)		n.a	n.a			2.6		2.6	
n.a : Not applicable									

Table 1. Operational conditions and physical/chemical characterization of the real wastewater and cultivation broth in the anoxic tanks and

3. Results and Discussion

The mass balance calculations over the 208 days of operation showed recoveries for TOC, IC, TN and P-PO $_4^{3-}$ of 100±1%, 99±4%, 100±5% and 100±14%, which validated the analytical and instrumental methodologies used in this study. This mass balance approach allowed to better understand the symbiosis between microalgae and bacteria in this novel anoxic-aerobic algal-bacterial photobioreactor configuration [14], by quantifying the extent of the mechanisms involved in C, N and P removal in each reactor.

The overall removal efficiency of organic matter measured as TOC under steady state operation averaged 89±2% along the 3 operational stages at 2 days of HRT due to the high photosynthetic oxygenation capacity and denitrification activity of the system. In this context, while the DO concentration in the anoxic tank remained close to 1 mg O₂/L (thus supporting an efficient denitrification since the O_2 carried out by the internal recycling was lower that O₂ demand of the RDWW), the DO in the photobioreactor fluctuated from 15 and $32 \text{ mg O}_2/\text{L}$ during illuminated periods to 1.5 and 7 mg O_2/L during dark periods in the photobioreactor (supplementary

material Figure S1). These oxygen concentrations were sufficient to satisfy the bacterial demand from NH4⁺ and TOC oxidation in the photobioreactor. The organic matter removal recorded in this study was similar to that typically achieved in conventional activated sludge systems (COD-REs of 85-90%) and in conventional HRAPs treating domestic wastewater (COD-REs 81-88%). The high light intensity used in this lab-scale study to boost microalgae photosynthetic activity (392 ± 19) μ mol/m²·s) supported an efficient overall steady state IC removal (95±4%) mainly based microalgae on assimilation, nitrification representing a minor IC removal mechanism ($\approx 4.1\%$ of the total IC input).

On the other hand, the average TN removal during SI under steady state operation accounted for $38\pm6\%$ with average NH₄⁺ removals of $39\pm9\%$. This low TN-RE was due to the low efficiency of NH₄⁺ nitrification during SI as a result of a severe IC limitation. Biogas supplementation in SII and SIII overcame this limitation and promoted steady state removals of TN and NH₄⁺ of $81\pm3\%$, $97\pm2\%$, respectively, at a HRT of 2 days. NH₄⁺ nitrification in the photobioreactor was the key step to ensure an efficient nitrogen removal in

the anoxic tank via denitrification, despite NH₄⁺ oxidation during SI was limited by the active photosynthetic IC uptake by microalgae. Comparable TN-REs ranging from 68% to 85% and N-NH₄⁺-REs of 80-93% are typically achieved in CO₂-supplemented HRAPs treating domestic wastewater but at HRTs of 3-7 days, with nitrogen assimilation into biomass and NH₃ stripping identified as the main nitrogen removal mechanisms [24]. Lower TN-REs ranging from 57% to 73% are often achieved in HRAPs during the treatment of domestic sewage without CO₂ supplementation at HRTs of 3-10 days, which highlights the superior of performance our two-stage photobioreactor [25]. In addition, the values hereby obtained for nitrogen removal were comparable with the removal efficiencies of $\approx 80\%$ typically reported in nitrification-denitrification activated sludge plants, although conventional WWTPs operate at 0.5-1 day of HRT [26].

Finally, average orthophosphate removal efficiencies of $59\pm17\%$ were recorded under steady state operation in SI. However, the supplementation of biogas resulted in an enhanced biomass growth and therefore in a slight increase in P-PO₄³-REs up to steady state values of

 $67\pm13\%$ and $60\pm6\%$ in SII and SIII, respectively. Bioassimilation into algalbacterial biomass was likely the main phosphorous removal mechanism since pH values fluctuated from 6.8 to 9.4 during illuminated periods and from 6.4 to 8.1 during the dark periods in the photobioreactor (supplementary material Figure S1). The average pH values recorded along the entire experiment were likely not sufficient to support phosphate precipitation, which has been shown to occur at pHs > 9.0 [6][27]. The $P-PO_4^{3}$ -REs here obtained (59 - 67%) were similarly those typically reported in HRAPs (50% to 75%) at significantly higher HRTs (3 - 7 days) and activated sludge processes at HRTs of 0.5 - 1 days [24]. However, the volumetric PO_4^{3-} removal rates achieved were superior based on the fact that the phosphate concentration in the RWW used in this study (33 \pm 8 mg P/L) was \approx 5 times higher than the P-PO₄³⁻ concentrations typically present in medium strength WW (\approx 7 mg P/L)[28] (Table 1).

Finally, the high robustness of this process configuration should be highlighted based on the consistent effluent concentrations of TOC, IC, NH_4^+ , TN, PO_4^{3-} despite the inherent variations of these parameters in RDWW.

3.1 Carbon and nutrient removal in the anoxic reactor

The overall removal efficiencies of TOC in the anoxic reactor accounted for 77±4% under steady state operation, with values of 77±4%, 76±6% and 79±3% for SI, SII and SIII, respectively (Figure 2a). This heterotrophic TOC removal (organic matter acting as electron donor) resulted in steady state concentrations of 27±5 mg/L in the anoxic tank regardless of the operational stage [29]. On other hand, a negative IC removal efficiency of -14±13% was recorded during SI as a result of CO₂ production from TOC oxidation in the anoxic tank (mainly driven by the use of O_2 electron which as acceptor, represented 67% of the total e⁻ acceptor consumption during SI) and the absence of a significant CO₂ stripping due to the overall CO₂ limitation in the process (Figure 2b). The slightly higher aqueous CO₂ concentration in the anoxic tank during SII and SIII mediated by biogas scrubbing supported a desorption of CO₂ from the anoxic tank, resulting in IC REs of 29±12% and 30±6%, respectively.

TN-REs in the anoxic tank increased from 18±8% in SI to 50±6% and 50±7% in SII and SIII, respectively (Figure 2c). This increase in TN removal was likely induced by the enhanced nitrification in the photobioreactor mediated by biogas supplementation, which ultimately promoted N-NO2⁻ and N-NO3⁻ reduction in the anoxic tank using the organic matter present in the influent wastewater. In fact, N-NO₂⁻ and N-NO₃⁻ represented the main e⁻ acceptors in SII and SIII, with a contribution to TOC oxidation of 56% and 60%, respectively. The steady state removals of N-NH4⁺ remained low at 6±14%, 9±11% and 2±7% during SI, SII and SIII, respectively. NH₄⁺ removal in the anoxic tank was due to biomass assimilation mediated by heterotrophic TOC removal, which remained constant regardless of the operational stage (Figure 2d). $N-NO_2^-$ concentrations in the anoxic tank under steady state operation were negligible (0.01±0.01 mg/L, 0.03±0.05 and 0.41±0.68 in SI, SII and SIII, respectively).

Enhanced carbon, nitrogen and phosphorus removal from domestic wastewater in a novel anoxic-aerobic photobioreactor coupled with biogas upgrading



Figure 2. Removal efficiency of (a) TOC, (b) IC, (c) TN, (d) N-NH₄⁺ and (e) P-PO₄³⁻ in the anoxic tank (\square), aerobic photobioreactor (\square) and overall system (\square) during the steady states achieved in the three operational stages evaluated. Vertical bars represent the standard deviation from replicate measurements during steady state operation.

Likewise, N-NO₃⁻ concentrations recorded in the anoxic tank in SI, SII and SIII were 0.04±0.03mg/L, 0.14±0.15mg/L and 0.73±1.25mg/L, respectively (Figure 3). These findings confirmed that both NO₂⁻ and NO₃⁻ derived from the photobioreactor and settler via the internal and external recirculations were efficiently reduced.



Figure 3. Time course of nitrite (triangles) and nitrate (squares) concentrations in the anoxic tank (black) and photobioreactor (white) during the entire experiment. Vertical dashed lines separate the different operational stages evaluated.

Negative overall P-PO₄³-REs of -17±31% were recorded in the anoxic tank under steady state operation, with P-PO4³⁻ removals of -14±36%, -18±29% and -20±25% during SI, SII and SIII, respectively (Figure 2e). These negative $P-PO_4^3$ -REs indicated that P was released by the algal-bacterial consortium in the absence of an eacceptor (nitrite, nitrate and dissolved oxygen) during SI, SII and SIII, respectively. In this context, recent studies have reported the ability of microalgae to accumulate non-structural P-PO₄³⁻ under aerobic conditions, which is then released in the absence of eacceptor (similarly to phosphate accumulating organisms, PAOs) [30][31]. In addition, the DGGEsequencing analysis revealed the

presence of heterotrophic bacteria with the ability to accumulate energy in the form of polyphosphate under excess of e⁻ acceptor and use this energy under anoxic conditions with the subsequent release of PO₄³⁻ to the culture medium. Hence. PAOs from the genus Acinetobacter (SI, SII and SIII), Luteolibacter (SI, SII and SIII), Thauera (SII and SIII), Pseudomonas (SIII), and Aeromonas (SI and SIII) were identified (supplementary materials Table S1) [29][32].

3.2 Carbon and nutrient removal in the photobioreactor

The TOC-REs under steady state operation in the photobioreactor averaged 12±5% regardless of the operational stage as a result of the

efficient removal of organic matter in the anoxic tank (Figure 2a). The consistent concentrations of TOC 19 ± 3 mg/L in the effluent over the entire experiment allowed us to estimate the fraction of non-biodegradable organic matter in the influent RWW to 11%. IC-REs in the photobioreactor under steady state condition accounted for 86±27% as a result of the intensive photosynthetic activity during the illuminated period along the three operational stages in the photobioreactor (Figure 2b). IC was almost completely depleted during the SI (supplementary material Figure S2b). The occurrence of IC limitation during SI supported the addition of biogas in order to supply an additional CO₂ source. Even under CO₂ supplementation, high IC-REs of $63\pm10\%$ and $62\pm7\%$ were SII recorded during and SIII. IC respectively. The enhanced availability mediated by biogas upgrading entailed an increase in the concentration of algal-bacterial biomass.

Low TN-REs of $20\pm7\%$ were recorded under SI steady state, which increased up to $30\pm7\%$ and $32\pm9\%$ in SII and SIII, respectively, as a result of the higher biomass production induced by biogas supplementation (Figure 2c). Likewise, while IC limitation mediated low N-NH₄⁺-REs (33±20%) during SI, the increase in nitrification activity supported by CO₂ supplementation increased N-NH4⁺-REs up to 89±11% and 96±7% in SII and SIII, respectively (Figure 2d and supplementary material Figure S2d). N-NO₂⁻ was the dominant form of oxidized nitrogen $(N-NO_2)^-$ = $5.6\pm4.0 \text{ mg/L} vs \text{ N-NO}_3^- = 0.9\pm0.9 \text{ mg/L}$ during SI (Table 1 and Figure 3). CO₂ supplementation via biogas upgrading promoted nitrification, which resulted in a decrease in N-NO₂⁻ concentration to 3.1 ± 3.8 mg/L concomitant with an increase in N-NO₃⁻ concentration up to 8.9 ± 5.5 mg/L in SII [29]. Likewise, an complete nitrification almost was achieved during steady SIII, with N- NO_3^- and $N-NO_2^-$ of 13.0±3.2 mg/L and 1.1±1.8 mg/L of N-NO₂⁻, respectively.

The overall steady state $P-PO_4^{3-}$ -REs in the photobioreactor accounted for $80\pm39\%$, with values of $73\pm49\%$, $85\pm28\%$ and $81\pm29\%$ in SI, SII and SIII, respectively (Figure 2e). P assimilation into algal-bacterial biomass was the most likely removal mechanism in the photobioreactor based on the range of pH values recorded during illuminated periods (6.8-9.4) and (6.4-8.1) during the dark periods in SII and SIII.

3.3 Biomass concentration and sludge volumetric index

TSS concentration in the anoxic tank increased from 1519±252 mg TSS/L in SI to 3113±361 mg TSS/L and 2480±309 mg TSS/L during SII and SIII, respectively (Table 1 and Figure 4a). Likewise, biomass concentration in the aerobic tank under steady state operation accounted for 1216±260 mg TSS/L, 2854±324mg TSS/L and 2047±186 mg TSS/L in SI, SII and SIII, respectively.

The fact that TOC removal remained similar along the three operational stages clearly showed that the increase in biomass concentration recorded during SII and SIII was mediated by the enhanced of autotrophic growth microbial communities (microalgae, cyanobacteria and nitrifying bacteria). Finally, effluent TSS concentrations during steady state gradually decreased from 163±83 mg TSS/L in SI, to 81±45 mg TSS/L in SII and 26±12 mg TSS/L in SIII (Figure 4a). The value obtained under steady state in SIII enabled compliance with the European Directive 97/271/CEE [33].

The sludge volumetric index recorded at the end of SI in the anoxic tank and photobioreactor accounted for 95 mL TSS/g and 161 mL TSS/g, respectively (Table 1 and Figure 4b). Surprisingly, the enhanced sedimentation observed during SII, based on the decrease in the effluent TSS concentrations, was not correlated with the SVI in the anoxic tank (128 mL TSS/g) or in the photobioreactor (169 mL TSS/g). These high SVI were likely due to the presence of the filamentous bacteria Caldilineae in SI and SII and *Clostridium* in SI, SII and SIII. However, the decrease in SVI recorded during SIII in both the anoxic tank and photobioreactor (80 mL TSS/g and 97 mL TSS/g, respectively) was correlated with low effluent TSS concentrations (Figures 4a and 4b). Overall, SVI of 50 - 100 mL/g in activated sludge plants are considered an indication of a good biomass settling [29]. Low SVI were also reported by al. (2015)Alcántara et in а photobioreactor designed with а continuous biomass recycling. Park et al. (2011) also reported an increase in microalgae settleability by 20% when implementing biomass recycling strategies in HRAPs, which confirmed the key of role of this operational strategy to enhance biomass settling [14][34]. The settling rates of biomass present in the anoxic tank accounted for 1.86 m/h, 1.20 m/h and 1.44 m/h in SI, SII and SIII, respectively. Settling rates of 1.56m/h, 1.02 m/h and 1.47 m/h were recorded for the biomass present in the

photobioreactor in SI, SII and SIII, respectively. An analysis of variance confirmed that the biomass present in the anoxic tank exhibited higher settling rates than the biomass in the photobioreactor in SI and SII. The results here obtained were comparable with those reported by de Godos et al. (2014) and higher than the rates obtained by Alcántara et al. (2015) using a similar AA-ABPh [13][14]. Similarly, 80% of algal biomass present in a HRAP treating domestic WW at 4 days of HRTs exhibited rates higher than 0.4 m/h (Gutierrez et al. 2016) [35].



Figure 4. Time course of (a) TSS concentration in the anoxic tank (\blacklozenge) and aerobic tank (\circ) and effluent (×, secondary axis), and (b) **SVI** in the anoxic tank (\blacksquare) and photobioreactor (\Box) during the steady states achieved in the three operational stages evaluated. Vertical dashed lines separate the different operational stages.

3.4 Dynamics of microalgae and bacteria population

Morphological characterization of microalgae population structure revealed a gradual dominance of the genus *Scenedesmus*, which accounted for 46 % of total microalgae population in the absence of biogas supply, and for 94-100% when CO₂ was supplemented to the wastewater treatment process (Figure

5). Desmodesmus spinosus, Pseudanabaena Leptolyngbya sp., benthonica and Acutodesmus obliquus represented 38%, 30%, 23% and 8% of the total microalgae population in SI, respectively. In SII, Leptolyngbya benthonica and Pseudanabaena sp were gradually replaced by Desmodesmus spinosus and Acutodesmus obliquus, which accounted for 50% and 44% of the total population, respectively. Finally, microalgae population in SIII became dominated by Desmodesmus spinosus (76%) and Scenedesmus tenuispina (24%). Scenedesmus species is commonly found in HRAPs treating

domestic WW [36] because of their tolerance to high nitrogen and organic matter concentrations [37][38]. This that biomass study suggests sedimentation and recycling can contribute the enrichment to of monoalgal microalgae species with good settling properties. Previous studies in pilot HRAPs conducted with biomass recycling promoted the dominance of unialgal cultures [34]. In this context, biomass settling and recycling also resulted in the dominance of Micractinium sp and Scenedesmus sp in HRAPs treating RWW with an external CO₂ supplementation [27].



Figure 5. Microalgae population structure in the photobioreactor during the entire operational period: *Chlorella* \square , *Pseudanabaena sp.* \square , *Leptolyngbya benthonica* \square , *Nitzschia palea* \square , *Scenedesmus tenuispina* \square , *Desmodesmus spinosus* \blacksquare and *Acutodesmus obliquus* \blacksquare .

DGGE analysis of the bacterial community in the photobioreactor revealed the occurrence of 10 phyla and

33 bands (Figure 6 and supplementary material Table S1). *Proteobacteria*, which are ubiquitous in the environment,

was the dominant phylum with 17 bands of the 33 sequenced. The phylum Proteobacteria was the most dominant with 9, 9, 6 and 12 bands detected in the inoculum, SI, SII and SIII, respectively (Figure 6 and supplementary material Table S1)[39]. The analysis also identified the phyla Acidobacteria, Verrucomicrobia, *Firmicutes* and Actinobacteria with two bands each, the phyla Chloroflexi, Cyanobacteria/Chloroplast,

Gemmatimonadetes, Ignavibacteriae, Candidatus Saccharibacteria with one band and 3 unclassified bacteria. Bacteria from the phyla Proteobacteria, Acidobacteria, Actinobacteria and Firmicutes were likely responsible for the degradation of organic matter in both the anoxic and photobioreactor tanks. Bacteria from the above mentioned phyla are typically found in activated sludge WWTP, autotrophic nitrifying and denitrifying bioreactors and HRAPs. More specifically, denitrifying bacteria such as Pseudomonas (SIII), Litorilinea (SI and SII), Gp4 (SII) and Thauera (SII and SIII) were identified (Figure 6 and supplementary material Table S1). Likewise, nitrifying bacteria belonging to the family Xhantomonadaceae (SI, SII and SIII) and genus Aeromonas (SI and SIII), Aquamicrobium (SI, SII and SIII), Luteliobacter (SI, SII and SIII), Thauera

(SII and SIII) *and Gp4* (SII) were detected as a result on the increased availability of CO₂.



Figure 6. DGGE profile of the bacterial community present in the anoxic-aerobic algalbacterial photobioreactor in the inoculum (S0), stage I (SI), stage II (SII) and stage III (SIII). Horizontal arrows and numbers indicate the most abundant bacterial communities. The name of the samples and the Shannon-Weiner diversity indexes are also depicted in the upper part of the gel.

The Shannon-Weiner diversity index (H) for the inoculum (S0) and the population established in the different operational stages showed a high bacterial diversity.

The Shannon-Weiner diversity index typically ranges from 1.5 to 3.5, higher H values corresponding to a higher species richness and evenness [16][20]. In this study, H indexes of 3.4, 3.5, 3.2 and 3.2 were estimated in the inoculum and in the microbial populations established during SI, SII and SIII, respectively (Figure 6). HRAPs treating WW typically exhibit H indexes ranging from 3.0 to 3.5, which confirms the high robustness and functionality of the microbiology present in algal-bacterial processes. Both the H index and the DGGE band profile clearly showed that biogas supplementation in SII and SIII stabilized the bacterial community. The analysis of the similarity indexes showed a lower similarity between the inoculum (S0) and the population in SI (25.4%), than the similarity between the populations in SI and II (63.2 %) in SII and SIII (41.30%), which indicated a functional specialization to the host environment during the experiment [40].

3.5 Biogas upgrading

 CO_2 supplementation via biogas upgrading was crucial to ensure an efficient nitrification in the photobioreactor and further denitrification in the anoxic tank. CO_2 removal from biogas in the absorption column averaged 92±2% and 93±2 %

Π III. during steady state and respectively (Figures 7a). No significant absorption of CH₄ was detected in the biogas absorption column here evaluated, likely due to the low solubility of CH₄. which prevented the uncontrolled release of this GHG from the experimental system.

H₂S was completely removed from biogas regardless of the operational stage as a result of its higher solubility compared to CO_2 (Figure 7b). The results obtained here were in agreement with the REs of 80% for CO₂ and 100% for H₂S reported by other authors in HRAPs devoted to biogas upgrading using a similar L/G ratio of ≈ 10 (v/v) While CO₂ supplied was [12][41]. assimilated by nitrifying bacteria and microalgae, H₂S was rapidly oxidized to SO_4^{2-} using the O_2 photosynthetically produced in the photobioreactor. In this context, the removal efficiencies of SO₄²⁻ in the anoxic tank accounted for $92\pm54\%$ and 16±50% during steady state II and III, respectively (Figure 7c). On the other hand, SO42-REs of -140±58% and -83±60% recorded were in the photobioreactor during steady state II and III, respectively, as a result of SO₄²⁻ production from H₂S oxidation (Figure 7b and 7c). The DGGE sequencing analysis revealed the presence of the H₂S

degrading strain *Pseudomonas* frederiksbergensis NR_117177, which supported the biological oxidation of H₂S in the system (Supplementary material Table S1) [42][43].



Figure 7. Time course of the inlet (\blacklozenge) and outlet (Δ) concentrations, and removal efficiency ($__$) of CO₂ (a) and H₂S (b), in the absorption column during stage II and III, and (c) removal efficiency of SO₄²⁻ in the anoxic tank (\blacksquare), aerobic photobioreactor (\blacksquare) and overall system (\blacksquare) during the steady states achieved in the three operational stages evaluated. Vertical bars represent the standard deviation from replicate measurements during steady state operation.

4. Conclusion

The novel anoxic-aerobic algal-bacterial photobioreactor coupled with a biogas upgrading unit here evaluated exhibited consistent C, N and P removal efficiencies. CO₂ supplementation from biogas was required to overcome the overall IC limitation recorded in SI, and

supported both an efficient nitrificationdenitrification process and an enhanced N and P removal by assimilation during SII and SIII. This innovative process configuration also supported an efficient biogas upgrading, with CO_2 and H_2S removal efficiencies of 85 and 100 %, respectively. Continuous biomass settling and recycling promoted the enrichment of an unialgal culture by the end of the experiment. Finally, DGGEsequencing analysis confirmed that biogas supplementation promoted the development of nitrifying, denitrifying and H₂S degrading bacteria during SII and SIII.

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SUPPLEMENTARY MATERIAL

Enhanced Carbon, Nitrogen and Phosphorus removal from domestic wastewater in a novel anoxic-aerobic photobioreactor coupled with biogas upgrading.

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CONTENT

Figure S1; Figure S2; Table S1;



Figure S1. Time course of pH and dissolved oxygen concentration in the photobioreactor during the entire experiment. The thick solid and dotted lines represent the dissolved oxygen concentration during the illuminated and dark periods, respectively. The pH in the illuminated and dark periods is represented by solid and dashed lines, respectively.

Enhanced carbon, nitrogen and phosphorus removal from domestic wastewater in a novel anoxic-aerobic photobioreactor coupled with biogas upgrading



Figure S2. Time course of the concentration of TOC (a), IC (b), N-NH₄⁺ and P-PO₄³⁻ in the RWW (\blacktriangle), anoxic tank (\diamond) and photobioreactor (\bullet) during the entire experiment. Vertical dashed lines separate the different operational stages evaluated. IC concentration during SII and SIII accounted for the CO₂ transferred from the biogas upgrading column.
Table S1. RDP classification of the DGGE bands sequenced and corresponding matches (BLASTN) using the NCBI database with indication of the similarity percentages and sources of origin. The presence/absence of each band in each sample together with its intensity are also shown.

Taxonomic placement	Band		SAM	IPLES		Closest relatives in Blast	Similarity	Source of origin	
(50% confidence level)	n°	S 0	SI	SII	S III	Name (accession number)	(%)		
Phylum Proteobacteria	1		Х			Uncultured bacterium (LN562529)	99	Refuse dump Nest 11 layer 3	
						Uncultured bacterium (KM290982)	97	Earthworm compost	
						Uncultured bacterium (LK392805)	97	Activated sludge treating municipal wastewater	
						Uncultured bacterium (KJ940485)	97	Activated sludge	
						Uncultured bacterium (FJ660585)	97	Activated sludge	
	2	XXX		XX	XX	Uncultured proteobacterium (AY755362)	91	Bioreactor	
						Uncultured bacterium (HQ827981)	90	Carpet-like mucilaginous cyanobacterial blooms in a hypereutrophic lake	
Class Gammaproteobacteria	3		XX		XX	Aeromonas sp. (HG763857)	92	Wheat rhizosphere from saline areas	
						Aeromonas sp. (KR189722)	91	Raw water	
Order Xanthomonadales									
Family Xanthomonadaceae	4	XX	XXX		XXX	Uncultured bacterium (GQ480132)	94	Activated sludge from wastewater treatment plant	
						Uncultured bacterium (KM290397)	94	Earthworm compost	
	5		XXX		XX	Arenimonas donghaensis (NR_043790)	92	Seashore sand	
						Uncultured bacterium (HM023338)	92	Surface water around 1m depth	
						Uncultured bacterium (JX271974)	92	Activated sludge in lab-scale reactor at 0 cm depth with dissolved oxygen between 0.7 mg/l and 0.9 mg/l	
						Uncultured bacterium (HQ640559)	92	Partial nitrifying-ANAMMOX municipal wastewater reactor	
						Uncultured bacterium (JX271974)	91	Autotrophic nitrifying bioreactor	
	6		XX	XX		Uncultured gamma proteobacterium (KF827303)	91	Biofilm	
						Aquimonas voraii (NR_042968)	91	Warm spring	
Genus Pseudofulvimonas	Genus Pseudofulvimonas 7 XXX	Х	Х	Uncultured bacterium (KP797893)	100	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludge			
						Uncultured Xanthomonadaceae (EU305597)	99	Wastewater plant	
						Uncultured bacterium (KF911229)	97	Composting process	
Order Aeromonadales	8		XX			Aeromonas sanarellii (NR_116584)	95	Culture collection	
						Aeromonas sp. (KR189870)	95	Raw water	
						Aeromonas sp. (KC571192)	95	Activated sludge	
			1			Aeromonas sanarellii (JF920492)	95	Wastewater treatment plant - treated wastewater	

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Family Aeromonadaceae								
Genus Aeromonas	9	XXX				Aeromonas caviae (KJ946377)	99	Wastewater treatment plant influent
						Uncultured bacterium (KP797890)	99	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludg
						Uncultured Aeromonas sp. (KJ651209)	99	Sludge in MBR for piggery wastewater
	10		XX		Х	Uncultured Xanthomonadaceae (KC994700)	98	Microalgae photobioreactor
						Uncultured bacterium (KM290640)	97	Earthworm compost
						Uncultured bacterium (FJ529949)	97	Autotrophic nitrifying bioreactor
						Uncultured Xanthomonadaceae (EF020325)	97	
Order Pseudomonadales								
Family Moraxellaceae								
Genus Acinetobacter	11		XX	XX	XX	Acinetobacter genomo sp. (KR611801)	98	Soil and water ecosystems
						Uncultured bacterium (LK393051)	98	Supernatant of activated treating municipal wastewater
						Acinetobacter harbinensis (KC843488)	98	River
Family Pseudomonadaceae								
Genus Pseudomonas	12				XX	Pseudomonas sp. (FJ889639)	99	Soil
						Uncultured bacterium (KM292305)	99	Sludge with earthworm
						Pseudomonas sp. (KJ742898)	99	Soil
						Pseudomonas mandelii (KJ726566)	99	River water
						Pseudomonas mandelii (CP005960)	99	Culture collection
						Pseudomonas sp. (KF286228)	99	Shallow aquifer
						Pseudomonas frederiksbergensis (NR_117177)	99	Soil from a gasification site
Class Betaproteobacteria Order Rhodocyclales								
Family Rhodocyclaceae	13	XX				Uncultured bacterium (AB488374)	92	Rrice paddy soil
						Uncultured bacterium (JN391796)	92	Biofilm in aerobic tank of hybrid reactor
			1			Uncultured bacterium (AB608667)	92	Rice paddy soil
			1			Uncultured bacterium (AB378590)	92	[13C]DNA fraction isolated from paddy soil ammended with nitrate and [13C]succinate
Genus Thauera	14	XXX		XX	XX	Uncultured bacterium (HG380609)	96	Wastewater
			1			Uncultured bacterium (KP641119)	96	Hhigh-rate denitrifying reactor treated with synthetic wastewater
			1			Thauera sp. (EF205258)	96	Sediment soils
		1		1	1	Thauera phenylacetica (NR_027224)	96	Anaerobic sewage sludge
		1	1	1	1	Uncultured bacterium (KP136297)	96	Wastewater from high-rate denitrifying reactor

	-	-						T
						Uncultured bacterium (KP054189)	96	Nitrification and denitrification reactors
Order Burkholderiales	15	XX			XXX	Uncultured bacterium (KM01625)	99	Anaerobic bioreduction of nitrate in hydrogen-based membrane biofilm reactor
						Uncultured bacterium (AB487629)	97	Rice paddy soil
						Uncultured bacterium (AB672290)	97	Rice paddy soil
Class Alphaproteobacteria					1			
Order Rhodobacterales								
Family Rhodobacteraceae					1			
Genus Roseibaca	16	XXX			XX	Uncultured bacterium (KM823738)	95	River sediment
						Uncultured alpha proteobacterium (EF367342)	94	Mixed biomass developed in SBR lab scale reactors inoculated with activated sludge from a municipal wastewater treatment in Portugal
						Rhodobacteraceae bacterium (AM403177)	93	marine aquaculture
Order Rhizobiales			1	1				
Family Phyllobacteriaceae		1	1	1	1			
Genus Aquamicrobium	17	XXX	XX	XX	XX	Uncultured bacterium (HG380597)	99	Wwastewater
						Uncultured bacterium (FJ167453)	99	Denitrifying bioreactor
						Aquamicrobium aestuarii (NR_108709)	99	Beach sand sample of the Yellow Sea
						Uncultured bacterium (HQ843719)	99	Nitrifying bioreactor
						Uncultured bacterium (JN087889)	99	Nitrifying bioreactor under inorganic carbon limitation
						Aquamicrobium ahrensii (AM884148)	98	Experimental biofilters for the treatment of animal rendering plant waste gases
						Uncultured Aquamicrobium sp. (EU305598)	98	Wastewater plant (WWTP)
Phylum Acidobacteria								
Class Acidobacteria-Gp4								
Genus <i>Gp4</i>	18			XX		Uncultured bacterium (HQ592595)	99	Activated sludge
						Uncultured bacterium (EU244112)	99	River
						Uncultured bacterium (KP054179)	98	Nitrification and denitrification reactors
Genus Blastocatella	19		XX	XX	Х	Unidentified bacterium (AF097780)	99	Activated sludge
						Unidentified marine bacterium (KC003380)	99	Surface seawater samples from China offshore waters and South China Sea
Phylum Verrucomicrobia								
Class Verrucomicrobiae								
Order Verrucomicrobiales								
Family Verrucomicrobiaceae								
Genus Luteolibacter	20		XX	Х	Х	Uncultured bacterium (FJ354094)	97	17th Street Canal, water, greater than 3 micron
						Uncultured bacterium (KM294491)	96	Sludge with earthworm
						Luteolibacter sp. (KC921164)	95	Soil

Image: A constraint of the sector of the s			1				Uncultured bacterium (KM294491)	94	Sludge with earthworm
Physical Problem Physical Problem<								94	Suuge with carthworld
Desc CostrainedNo </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>Uncultured Verrucomicrobium sp. (AY157106)</td> <td>93</td> <td>NOX activated sludge system</td>							Uncultured Verrucomicrobium sp. (AY157106)	93	NOX activated sludge system
Order Cohendration1213141	Phylum <i>Firmicutes</i>								
Image: Constraint of the sector of the sec	Class Clostridia								
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Genus Costrutury M							Clostridium sp. (JF312678)	92	Bioreactor
Image: Problem in the second secon	Family Peptostreptococcaceae								
Image: A strandImage: A stran	Genus Clostridium XI	23		XXX	XXX	XX	Uncultured bacterium (KP797907)	98	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludge
Phylum ActinobacteriaII							Uncultured bacterium (KJ808497)	98	Activated sludge
Class Actimates information Image: Marking and Constraint of Marking Mar							Uncultured bacterium (KC551590)	98	Activated sludge
Subclass ActionabactivitiesII </td <td>Phylum Actinobacteria</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td>	Phylum Actinobacteria								
Order ActionaryCentales2AXXVXVXVxVxUncultured bacteriam (IF497778)99Activated slageConstructionIIIIIVxVxUncultured bacteriam (IF497778)96Microalgas from HRAP treating diluted vinases with wastewater treatment plant activated slageImage: Image: Imag									
And And And And And And And And Inclusion									
Image: Marking Markin	Order Actinomycetales	24	XX			XX	Uncultured bacterium (JF497778)	99	Activated sludge
Image: Constraint of the state of the st							Uncultured bacterium (HQ860566)	97	Stream water
LocLo								96	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludge
Phylum Chloroflexi Image: Marking Caldilineage Image: Marking Caldilineage <thimage: caldilineage<="" marking="" th=""> Image: M</thimage:>		25		XXX	XX	XXX	Uncultured bacterium (JF497778)	97	Activated sludge
Class Caldilineale Image: Marking Caldilineales Image: Ma							Uncultured bacterium (HQ860566)	95	Stream water
Order CaldinealesImage: Marking CaldinaceaeImage: M									
Family CaldilineaceaeImage: Marking Caldilin									
Genus Litorilinea 26 XXX XX XXX XXX Vacultured bacterium (HQ014651) 97 Wastewater treatment plant Image:	Order Caldilineales								
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Image: Marking	Genus <i>Litorilinea</i>	26	XXX	XX	XXX		Uncultured bacterium (HQ014651)	97	Wastewater treatment plant
Phylum Cyanobacteria/Chloroplast Image: Constraint of the second sec							Uncultured bacterium (JN391658)	97	Activated sludge in aerobic tank of activated sludge reactor
Class Cyanobacteria Image:							Uncultured bacterium (AB576907)	97	Denitrifying PGE pellet samples
Family Family IV Image: Im	Phylum Cyanobacteria/Chloroplast								
Genus GplV 27 XXX Uncultured bacterium (KF912972) 97 Soil Phylum Ignavibacteriae I I I I Uncultured cyanobacterium (HQ032346) 96 Tailing pound Class Ignavibacteriae I									
Image: Section of the section of th									
Image: Constraint of the constraint	Genus <i>GpIV</i>	27	XXX	х	Х	Х	Uncultured bacterium (FJ612380)	97	Lake water
Phylum Ignavibacteriae Image: Class Ignavibacteria Image: Class Ignavibacteria Image: Class Ignavibacteria Image: Class Ignavibacteria							Uncultured bacterium (KF912972)	97	Soil
Class Ignavibacteria							Uncultured cyanobacterium (HQ032346)	96	Tailing pound
	Phylum Ignavibacteriae		1						
Order Ignavibacteriales	Class Ignavibacteria								
	Order Ignavibacteriales		1	1					

Family Ignavibacteriaceae								
Genus Ignavibacterium	28	XXX	Х	Х		Bacterium enrichment (KM210546)	97	Activated sludge from the aerobic tank was enriched with ammonium chloride and sodium sulfate
						Uncultured bacterium (FJ710747)	97	Anaerobic ammonium oxidation reactor
Phylum Gemmatimonadetes								
Class Gemmatimonadetes								
Order Gemmatimonadales								
Family Gemmatimonadaceae								
Genus Gemmatimonas	29	XXX				Uncultured Gemmatimonadetes (HM022183)	98	Thermomineral spring
						Uncultured Gemmatimonas sp. (EU283561)	98	Lake
						Uncultured bacterium (HM357106)	97	Algal mat
						Uncultured bacterium (GQ441328)	97	Marine microbial mats from a sandy intertidal beach
Phylum Candidatus Saccharibacteria					1			
Genus Saccharibacteria-genera-incertae-sedis	30		XX		XX	Uncultured bacterium (JX105655)	93	Ornamental fish aquaria
						Uncultured bacterium (KR306129)	92	Sludge-amended soil with spent mushroom compost day 28
Unclassified Bacteria	31			XXX		Uncultured bacterium (JQ427509)	95	Soil
						Uncultured bacterium (JQ978896)	95	Atrazine-contaminated soil added with inorganic nitrogen and DAT1
						Uncultured bacterium (JQ056068)	94	Soil
	32		XX			Uncultured bacterium (HE647661)	99	CAS wastewater treatment pilot plant
						Environmental 16s rDNA sequence (CU466684)	99	Evry wastewater treatment plant anoxic
						Uncultured bacterium (AB286410)	98	Activated sludge
	1		1			Uncultured bacterium (KM340821)	97	Earthworm cast
	33	XX	XX			Uncultured bacterium (KP797908)	91	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludge

(84 **)**

Chapter 2





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Abstract

The dynamics of microalgae population during piggery wastewater (PWW) treatment in four open photobioreactors operated at 27 days of hydraulic retention time, and inoculated with Chlorella sp. (R1), Acutodesmus obliquus (R2), Oscillatoria sp. (R3) and in the absence of inoculum (R4), were evaluated for 6 months. In addition, the algal-bacterial biomass concentration, removal of organic matter, nutrients and heavy metals were also assessed. The results revealed a high diversity and rapid variations in the structure of microalgae populations, Chlorella sp. being dominant in R4 throughout most of the operational period. Steady state average biomass concentration ranged from 2445-2610 mg/L in R1-R3 to 3265 mg/L in R4. No significant differences were recorded in the removal efficiencies (REs) of total organic carbon (86-87%), inorganic carbon (62-71%), total nitrogen (82-85%) and total phosphorous (90-92%). Finally, Zn-REs accounted for 26% in R3, 37% in R2, and 49% in R1 and R4.

Keywords

Algal-bacterial processes; Biomass production; Heavy metal removal; Microalgae dynamics; Piggery wastewater treatment.

1. Introduction

The current global energy and climate change crisis has triggered the quest for alternative green energy sources with a low carbon dioxide (CO₂) footprint (González-Fernández et al., 2012a). In this context, microalgae have emerged as a promising renewable energy platform due to their ability to transform sunlight directly into gas biofuels (i.e H₂) or an organic biomass feedstock that can be further bioconverted into multiple liquid and gas biofuels (Richmond, 2004). Thus, microalgal biomass can be anaerobically digested yielding biogas $(CH_4 + CO_2)$ and a nutrient rich digestate (Ehimen et al., 2011; González-Fernández et al., 2012b). In addition, while the lipid fraction of microalgae can transesterified into biodiesel be al.. 2011). (Vimalarasan et the carbohydrate fraction can be fermented into bioethanol (Naik et al., 2010) or biohydrogen (Chandrasekhar et al., 2015). Microalgae exhibit multiple advantages over conventional energy crops such as high areal productivities (50-100 tn/ha·y), cultivation in nonarable land (preventing competition with food) and high lipid or carbohydrate fractions depending on the cultivation conditions. Likewise, microalgae can be cultivated in fresh, marine or

wastewaters (Cheah et al., 2016).

In this context, nutrient-rich wastewaters represent a valuable feedstock to reduce of the costs microalgae and cyanobacteria (from now on referred to as microalgae) cultivation, which will ultimately increase the costcompetitiveness of microalgae-based biofuels (Acién et al., 2016). Algalbacterial symbiosis can combine a lowcost mass production of biomass with the treatment of wastewater to levels required for discharge into natural water bodies. Indeed, both domestic, industrial livestock wastewaters and have successfully supported microalgae cultivation (Muñoz et al., 2003; Muñoz 2006). and Guieysse, During microalgae-based wastewater treatment, both the organic carbon, nitrogen and phosphorous present in the residual effluent are assimilated into algalbacterial biomass. Heavy metals and pathogens are also efficiently removed during microalgae growth as a result of adsorption and pH-mediated mechanisms. Despite microalgae cultivation entails in wastewater significant economic and environmental advantages over axenic mass production of microalgae in mineral salt media, controversy still exists in literature about the possibility of maintaining monoalgal

cultures with a constant biomass composition during microalgae-based wastewater treatment. This is central to the development of microalgae-based biorefineries for biofuel production, whose viability depends on the supply of a biomass with a consistent year-round composition and characteristics. Hence, while most studies conducted under laboratory or outdoors conditions focused on the removal of key pollutants present in wastewater, little attention has been paid to the monitoring of the dynamics of microalgae population.

Pig production is a key economic sector in many countries in Europe, accounting for 148.7 million pigs heads and 44.3% of the total European livestock (EU, 2015; MAGRAMA, 2015) in 2015. European pig farming generates 217-434 million m^3/y (4-8 L/day/pig) of piggery wastewater containing high concentrations of organic matter and nutrients (De Godos et al., 2009). The estimated average organic matter and nutrient load present in EU piggery wastewaters in 2015 amounted to 8.923.000 tn chemical oxygen demand (COD)/y, 890.000 tn nitrogen (N)/y and 223.000 tn phosphorous (P)/y (EU, 2016). In addition, piggery wastewater can contain high concentrations of heavy metals such as Zinc and Copper,

typically used as growth promoters in swine nutrition (Abe et al., 2012; De la Torre et al., 2000).

The experimental work herein conducted evaluated the dynamics of microalgae population during piggery wastewater treatment in four open continuous photobioreactors inoculated with two green microalgae species, a cyanophyta, and without inoculum. In addition, the influence of the microalgae inoculum on the steady state organic matter, nutrient and heavy metal removal was assessed.

2. Materials and methods

2.1 Microalgae

Chlorella Fott minutissima and Nováková was obtained from an indoor high rate algal pond (HRAP) treating centrate at the Dept. of Chemical Environmental Engineering and Technology from Valladolid University (Spain). Acutodesmus obliguus and Oscillatoria sp were kindly provided by of the Department Chemical Engineering from Almeria University (Spain).

2.2 Piggery wastewater

Fresh centrifuged piggery wastewater (PWW) was collected at a nearby farm at Cantalejo (Spain) and stored at 4°C. The

average composition of the piggery wastewater diluted at 15% was: 1340±34 mg/L of total suspended solids (TSS), 1375±121 mg/L of total organic carbon (TOC), 314±55 mg/L of inorganic carbon (IC), 393±26 mg/L of total nitrogen (TN), 9.4±0.4 mg/L of total phosphorus (TP) and 0.7±0.2 mg/L of zinc (Zn). Nitrate (NO_3^{-}), nitrite (NO_2^{-}), (Cu) and arsenic copper (As) concentrations remained below detection limit (Table 1).

2.3 Experimental set-up

The experimental set-up consisted of four 15.8 cm deep 3 L open photobioreactors illuminated at 2800 μ mol/m²·s for 12 hours a day (08h00 to 20h00) by LED lamps arranged in a horizontal configuration 20 cm above the photobioreactor surface (Figure1). The photobioreactors were immersed in a water bath to prevent the high temperatures imposed by the LEDs irradiation. Immersion water pumps were used to mix the algal-bacterial cultivation broth in the reactors. The photobioreactors were fed with piggery wastewater diluted at 15% using an auto control 205U7CA multi-channel cassette pump (Watson-Marlow, UK). The pH in the cultivation broth was automatically maintained at 8.0 via CO₂ addition (CARBUROS **METALICOS-**Barcelona, Spain) using a Crison multimeter M44 control unit (Crison Instruments, Spain).



Figure 1. Schematic diagram of the algal-bacterial photobioreactor set-up using carbon dioxide supplementation for pH control.

2.4 Experimental design

Photobioreactors 1, 2 and 3 (namely R1, R2 and R3, respectively) were inoculated with Chlorella minutissima Fott and Nováková, Acutodesmus obliquus and Oscillatoria sp., respectively, at an initial TSS concentration of 220 mg/L (corresponding to initial cell concentrations of 1.750, 0.295 and $0.332 \cdot 10^9$ cells/L, respectively). Photobioreactor 4 (R4) was not inoculated and served as control. The photobioreactors, which were initially filled with tap water, were operated at a hydraulic retention time (HRT) of ≈ 27 days (estimated based on the influent PWW) for 176 days. Photobioreactors effluents overflowed separately as a function of the evaporation rates. Liquid samples of 30 mL were weekly drawn from the influent PWW and effluent of R1, R2, R3 and R4 to determine the concentrations of TOC, IC, TN, NO2⁻,

 NO_3^- , TP and TSS. Effluent samples were filtered through 1 µm glass fiber filters prior analysis. Likewise, the microalgae population structure in R1, R2, R3 and R4 was weekly assessed from biomass samples preserved with lugol acid at 5% and formaldehyde at 10%, and stored at 4 °C prior to analysis (only 8 samples from each photobioreactor were analyzed). The dissolved oxygen and temperature of the cultivation broths were measured twice per day, while the influent and effluent flowrates were daily recorded in all photobioreactors to monitor water evaporation losses. Finally, the C, N and P content of the algal bacterial biomass was measured under steady state at the end of the experiment.

The C, N and P removal efficiencies (RE) were calculated according to Eq. (1):

(1)

$$RE(\%) = \frac{(C_{feed} \times Q_{feed}) - (C_{eff} \times Q_{eff})}{C_{feed} \times Q_{feed}} \times 100$$

where C_{feed} and C_{eff} represent the dissolved concentrations of TOC, IC, TN, TP and Zn in the PWW and photobioreactors effluents, respectively, while Q_{feed} and Q_{eff} represent the PWW and effluent flow rates. The process was considered under steady state when the TSS concentrations in the photobioreactors remained stable for at least four consecutive samplings (~ 1 month). The results were here provided as the average \pm standard deviation from duplicate measurements along one month of steady state (days 150-176).

2.5 Analytical procedures

A Crison M44 multimeter and a Crison PH 28 meter were used for the on-line measurement of the pH. Dissolved oxygen (DO) and temperature (T) were recorded using an OXI 330i oximeter (WTW, Germany). A LI-250A light meter (LI-COR Biosciences, Germany) was used to measure the light intensity as photosynthetically active radiation (PAR). TOC, IC and TN concentrations were determined using a TOC-V CSH analyzer equipped with a TNM-1 module (Shimadzu, Japan). Nitrate and nitrite were analyzed by high performance liquid chromatography-ion conductivity (HPLC-IC) in a Waters 515 HPLC pump coupled with a Waters 432

ionic conductivity detector and equipped with an IC-Pak Anion HC ($150 \text{ mm} \times 4.6$ mm) column. TSS and TP concentrations were determined according to Standard Methods (APHA, 2005). The analysis of the C, N and P content in the algalbacterial biomass was carried out using a LECO CHNS-932 elemental analyzer with pre-dried and grinded algalbacterial biomass. The concentration of Zn, Cu and As was determined using a 725-ICP Optical Emission Spectrophotometer (Agilent, USA) at 213.62. The identification and quantification of microalgae were conducted by microscopic examination (OLYMPUS IX70, USA) according to Phytoplankton Manual (Sournia, 1978).

Parameter	PWW	R 1	R2	R3	R4
Evaporation (%)	n.a	60±6	60±7	60±6	60±8
Temperature (°C)	n.a	30±2	30±2	30±2	30±2
Dissolved Oxygen (mg/L)	n.a	0.8	1.1	1.3	0.9
TOC (mg/L)	1375±121	459±31	452±31	482±27	490±37
IC (mg/L)	314±55	285±14	242±34	227±33	294±27
TN (mg/L)	393±26	174±11	166±15	165±12	149±10
Nitrite (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
Nitrate (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
TP (mg/L)	9.4±0.4	2.4±0.3	2.1±0.2	1.9±0.5	1.8±0.3
Zinc (mg/L)	0.7±0.2	0.9±0.2	1.1±0.1	1.3±0.3	0.9±0.3
Copper (mg/L)	< 0.6	< 0.6	< 0.6	< 0.6	< 0.6
Arsenic (mg/L)	< 0.6	< 0.6	< 0.6	< 0.6	< 0.6
TSS (mg/L)	1340±34	2610±191	2569±69	2445±222	3265±133
n.a : Not applicable					

Table 1. Physical/chemical characterization of the diluted swine manure and cultivation broth in the photobioreactors at steady state.

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3. Results and Discussion

3.1 Dynamics of microalgae population

Chlorella sp., the inoculated microalgae species in R1, was detected throughout most of the experimental period in this photobioreactor and dominant at days 37 and 86 (at concentrations of $0.5 \cdot 10^9$ and $0.9 \cdot 10^9$ cells/L, respectively). Acutodesmus obliquus was also identified in R1 and became the dominant species by day 58. Finally, Aphanothece sp. was detected for the first time by day 58 and was dominant from day 122 to the end of the operation of R1 (Figure 2a). Similarly, the inoculated microalga species in R2 (Acutodesmus obliquus) was identified along the entire photobioreactor operation, with a significant dominance by days 37, 58 and 122 at cell concentrations of $1.3 \cdot 10^9$, $1.8 \cdot 10^9$ and 0.3.10⁹ cells/L, respectively. Chlorella sp. was identified in R2 from the first operational days and remained at similar cell concentrations throughout the entire experiment (from $0.3 \cdot 10^9$ to $0.7 \cdot 10^9$ cells/L). Finally, Aphanothece sp. became dominant in R2 by the end of operation, with final cell concentrations $2.9 \cdot 10^9$ cells/L of (Figure 2b). Oscillatoria sp. was replaced by Chlorella sp. and Acutodesmus obliquus

in R3 from the first operational days (after the inoculation a change in color from green to red was noticed), *Chlorella* sp. being the dominant species throughout the entire operation with a maximum concentration of $8.2 \cdot 10^9$ cells/L by day 58 (Figure 2c). The higher pollution-tolerance of Chlorella sp. to PWW, combined with the high temperature and irradiations prevailing in this study, could have caused this rapid replacement of Oscillatoria sp (Talbot et al., 1991). Despite R4 was not inoculated, Chlorella sp. and Aphanothece sp. were present in the photobioreactor from the first days, *Chlorella* sp. being the dominant species along the 6 months of experiment. The gradual increase in number of cells of Aphanothece sp. in R1, R2 and R4 of suggest the influence the **PWW** characteristics of the on microalgae population (Figure 2).

The higher dominance of *Chlorella* sp. in the four photobioreactors confirmed the high tolerance of this green microalgae to the pollutants and concentrations typically present in PWW (Kim et al., 2016; Kuo et al., 2015; Yuan et al., 2013). Indeed, the high abundance of *Acutodesmus obliquus* and *Chlorella* sp. (both belonging to the Chlorophyta phylum) along the experimental period

in R1, R2 and R3 matched the microalgae pollution-tolerance classification reported by Palmer et al. (1969), who ranked Scenedesmus and *Chlorella* 4th and 5th, respectively. It can be hypothesized that organic pollution exhibited a higher influence on microalgae population structure than other environmental parameters such as water hardness, light intensity, pH, DO or temperature (Palmer, 1969). On the other hand, Aphanothece sp., which was not previously classified as a pollution tolerant microalga, was mainly identified at the end of experiment in R1 and R2 (Palmer, 1969). However, Aphanothece microscopica nägeli and Aphanothece Clathrata successfully supported the removal of organic matter and nitrogen from parboiled rice wastewater (REs of 83.4 and 72.7% for COD and N-TKN, respectively) in a 4.5 L tubular photobioreactor operated batchwise for 24 hours (Queiroz et al., 2007). Likewise, Bastos et al. (2014) reported COD and N-TKN REs of 97 and 78%, respectively, in a 4L batch tubular reactor treating parboiled rice wastewater for 24 hours.

The lack of monoalgal cultures in the four photobioreactors throughout the experimental period and the rapid variations in microalgae population structure here recorded (mainly in R1 and R2) revealed the difficulty to maintain monoalgal cultures during the treatment of PWW in open systems (Posadas et al., 2015). In this context, a lower microalgae diversity was observed at higher biomass concentrations, which was in agreement with Park et al. (2011). In addition, the current morphological microalgae characterization revealed that the inoculation of a photobioreactor during PWW treatment with a specific microalga does not guarantee its longterm dominance (Serejo et al., 2015). Finally, it should be stressed that the different microalgae cells concentration in the inoculum of the photobioreactors $(1.750, 0.295 \text{ and } 0.332 \cdot 10^9 \text{ cells/L for}$ R1, R2 and R3, respectively) only affected the time required to reach steady state and the initial treatment performance, but it did not modify the conclusions here obtained since the performance of the systems was analyzed at constant under steady state.



Figure 2. Time course of microalgae population structure in (a) R1, (b) R2, (c) R3 and (d) R4. Acutodesmus obliquus (2), Aphanothece sp. (2), Chlorella sp. (2), Oscillatoria sp. (2), and total number of microalgae cells (•).

3.2 Biomass concentration and productivity

Biomass concentration in R1, R2 and R3 increased from 220 mg TSS/L to 530, 680, and 660 mg TSS/L, respectively, during the first 38 days. A moderate increase from 0 to 200 mg TSS/L was also recorded in R4 (Figure 3). A significant biomass concentration increase occurred in R1, R2 and R3 from day 38 to 93, when TSS the concentrations of 2440, 2140 and 2500 mg TSS/L, respectively, were measured. However, a lower biomass growth rate was observed during this period in R4, where concentrations up to 1200 mg TSS/L were recorded (Figure 3). Biomass concentration in R2 and R3 remained constant from day 93 onwards at average concentrations of 2569±69 and 2445±222 mg TSS/L, respectively. Biomass concentration in R1 fluctuated from day 93 to 150, to finally stabilize at 2610±191 mg TSS/L, which was similar to the concentrations reached in R2 and R3 (Figure 3). On the other hand, biomass concentration exponentially increased in R4 from day 93, to reach average value of 3265±133 mg TSS/L by the end of the experiment. Surprisingly, the highest algal-bacterial biomass concentration under steady state was achieved in the non-inoculated photobioreactor despite its longer lag

phase. Likewise, the highest TOC, IC, TN, TP and Zn REs (below discussed) were obtained in R4, which highlighted higher robustness of the native microalgae species acclimated to the environmental and operational conditions prevailing during PWW treatment (Figures 2 and 3, Table 1) (Olguín et al., 2013). In addition, the results clearly showed a similar biomass growth pattern in the photobioreactors inoculated with a specific photosynthetic microorganisms in comparison with the control unit R4.

The high biomass concentrations here recorded were supported by the high carbon and nutrients concentrations in the diluted PWW and by the high water evaporation rates in the systems, which accounted for 60 % of the influent PWW in all photobioreactors as noticed by Guieysse et al. (2013) (Table 1). Hence, biomass productivities under steady state averaged 6.2±0.5, 6.1±0.2, 5.8±0.6 and 7.8 ± 0.3 g/m²·d in R1, R2, R3 and R4, respectively. These productivities were comparable to those obtained during the secondary domestic treatment of wastewater treatment in pilot raceways at high HRT in Almeria (Spain), and were likely limited by the long HRT needed to ensure satisfactory organic

matter and nutrients removals (Posadas et al., 2015).

Finally, the comparison between the evolution of the total number of microalgae cells in the cultures and the TSS concentrations (Figures 2 and 3) showed no direct correlation as a result of the dominant role of bacteria in the process, which itself was influenced by the high biodegradable organic matter load. In this regard, an accurate empirical determination of the individual bacteria and microalgae populations would bring valuable insights about the mechanisms underlying organic matter and nutrient removal during PWW treatment.



Figure 3. Time course of TSS concentration in R1 (Δ), R2 (\Diamond), R3 (\Box) and R4 (\circ)

3.3 Carbon and nutrient removal

A comparable bioremediation performance in terms of TOC, IC, TN and TP removal was recorded regardless of the microalgae inoculated in the photobioreactor (Figure 4 and Table 1). In this context, the dominant microalgae species prevailing in the photobioreactor did not influence process performance. In this particular study, the high light irradiances and the optimum temperature for microbial activity supported an effective PWW treatment. Thus, despite the low DO concentrations in the cultivation broth ($\leq 1.3 \text{ mg/L}$), TOC-REs accounted for 86 ± 1 , 87 ± 5 , 86 ± 1 and 86 ± 1 % in R1, R2, R3 and R4, respectively, which resulted in average TOC concentrations in the effluent at the end of the operational period of 459 ± 31 , 452 ± 31 , 482 ± 27 and 490 ± 37 mg/L, respectively (Figure 4 and Table 1). Please note that the high water evaporation rates typically encountered in open photobioreactors resulted in moderately high effluent TOC concentration despite the high removal efficiencies achieved. The results herein obtained confirmed the consistent removal of organic matter from PWW by algal-bacterial processes and were in agreement with the study conducted by De Godos et al. (2009), who reported COD removal efficiencies of 76±11% in a 464 L high rate algal ponds (HRAP) during the treatment of 20 and 10 folds diluted PWW. Similarly, IC-REs of 63±3, 69±4, 71±4 and 62±3 % were recorded at the end of the process in R1, R2, R3 and R4, respectively, which resulted in IC concentrations in the cultivation broth of the photobioreactors of 285±14, 242±34, 227±33 and 294±27 mg/L, respectively (Figure 4 and Table 1). These high IC-REs were promoted by the intensive photosynthetic activity during the illuminated period over the 176 days of operation. However, carbon removal stripping by (prior mineralization of the organic carbon to CO_2) was the main mechanism accounting for carbon fate, since only 37, 38, 36 and 48 % of the total carbon removed was recovered in the harvested biomass in R1, R2, R3 and R4, respectively, under steady state conditions. This estimation was based on the carbon content of the biomass under

steady state (as described below) and did not account for the CO_2 input for pH control.

TN-REs of 82±1, 83±3, 83±1 and 85±1 % were recorded under steady state in R1, R2, R3 and R4, respectively, which resulted in TN concentrations in the photobioreactor effluent of 174±11, 166±15, 165±12 and 149±10 mg/L, respectively (Figure 4 and Table 1). These high TN effluent concentrations in spite of the effective nitrogen removal efficiencies were due to the high evaporation rates in the photobioreactors. The TN-REs here recorded were similar to those reported by De Godos et al. (2009), who measured average total kjeldahl nitrogen (TKN) removals of 86±6% during PWW treatment in an open HRAP, and higher than the TN-REs of 63% obtained during the treatment of PWW under laboratory conditions in a 500 ml conical flasks incubated on a rotatory shaker at 27 °C 150 under and rpm continuous illumination (Abou-Shanab et al., 2013). Likewise, Posadas et al., (2017) operated an outdoors HRAP supporting TN-REs of 80-86% during the treatment of centrate. Nitrogen removal by stripping was the main mechanism in our study, since only 26, 26, 23 and 31 % of the total nitrogen removed was recovered in

the harvested biomass in R1, R2, R3 and R4, respectively.

On the other hand, steady state TP-REs of 90±2, 91±1, 92±2 and 92±2 % were recorded in R1, R2, R3 and R4, respectively, which supported effluent TP concentrations of 2.4±0.3, 2.1±0.2, 1.9 ± 0.5 and 1.8 ± 0.3 mg/L, respectively (Figure 4, Table1). The TP-REs as (P- PO_4^{3-}) herein obtained were similar to those reported by Posadas et al., (2017) during the treatment of centrate in an outdoors HRAP (84 - 92%). Likewise, the TP-REs reported were in agreement with Franchino et al. (2016), who recorded phosphate REs > 90% during the treatment of 5 and 10 times diluted digestate in 250 ml flasks. Phosphorous assimilation into algal-bacterial biomass was the main removal mechanism in the photobioreactors based on the moderate

pН values prevailing in the photobioreactors during the entire experiment (pH=8), which did not significant phosphate support a precipitation (García et al., 2017). Thus, a phosphorus mass balance revealed that 93, 93, 96 and 100 % of the total phosphorus removed was recovered in the harvested biomass for R1, R2, R3 and R4, respectively.

Overall, it is worth noting that a similar macroscopic bioremediation performance was recorded in the photobioreactors in spite of the different microalgae population structures under steady state (and during most of the experiment period), which suggest that bacteria played a dominant role during of high the treatment strength wastewaters such as piggery effluents.



Figure 4. Average removal efficiencies of TOC (), IC (), TN () and TP () under steady state. Vertical bars represent the standard deviation from replicate measurements during steady state operation.

Finally, comparable carbon, nitrogen and phosphorus contents were measured in the harvested biomass under steady state regardless of the initial inoculum, with average values of 50 ± 1 , 7.8 ± 0.3 and 0.75±0.06 % for C, N and P, respectively (Figure 5). These elemental biomass compositions were similar to those reported by Posadas et al. (2013) during domestic wastewater treatment in a 15 L algal-bacterial biofilm photobioreactor (42±2, 7±1 and 1.3±0.3 % for C, N and P, respectively), despite the different C/N/P ratio in both wastewaters (C/N/P of 100/15.6/0.6 in PWW and 100/18/5 in domestic wastewater). Likewise, these

results were in agreement with those obtained by Cabanelas et al. (2013), who reported a C, N and P content in the harvested biomass of ≈ 44 , 7.5 and 0.5 %, respectively, in a photobioreactor inoculated with Chlorella vulgaris and supplemented with CO₂ during the treatment of effluent from primary settler. In this context, the results herein obtained confirmed the similar algalbacterial biomass composition regardless of the microalgae species present in the cultivation broth or operational conditions.



Figure 5. C (\mathbb{X}), N (\square) and P (\blacksquare) content in the biomass present in the photobioreactors under steady state.

3.4 Heavy metals removal efficiency

The overall steady state Zn-REs in R1, R2, R3 and R4 accounted for 49±6, 37 ± 6 , 26 ± 5 and 49 ± 5 %, respectively, which resulted in average effluent Zn concentrations of 0.9 ± 0.2 , 1.1 ± 0.1 , 1.3 ± 0.3 and 0.9 ± 0.3 mg/L, respectively, at the end of the operational period (Table 1). These values were similar (Zn-REs of 37%) to those reported by Abe et al. (2008) during PWW treatment in wetlands. The fact that the highest Zn-REs occurred in the photobioreactors with the highest biomass concentrations (R1 and R4) and the lowest Zn-RE in R3 (at the lowest biomass concentration) suggested that Zn removal was mediated by biosorption onto the algal-bacterial biomass present in the photobioreactor

(Table 1) (Kaplan et al., 1987; Muñoz et al., 2006). This showed the high tolerance of species such as *Chlorella* sp. to heavy metal contamination (Muñoz et al., 2006). Higher Zn-REs by biosoportion would be expected at higher pHs according to Muñoz et al. (2006), who observed an increase in Zn accumulation into the algal-bacterial biomass from 5.0 to 11.7 mg Zn/g biomass when pHs was raised from 7 to 9, respectively. The determination of copper and arsenic removal efficiencies was not possible based on the low concentrations of these metals in the PWW (below the detection limit of the instrument = 0.6 mg/L).

4. Conclusion

This research revealed the difficulty to maintain monoalgal cultures during **PWW** treatment in openphotobioreactors operated under similar environmental and operational conditions. The high abundance of Chlorella sp. in most photobioreactors confirmed the high tolerance of this microalga the pollutants. to The acclimation of native species to the characteristics of the PWW resulted in highest biomass concentrations. An efficient PWW treatment occurred regardless of the microalgae species inoculated, which confirmed the robustness of algal-bacterial processes devoted to carbon and nutrient removals from livestock wastewaters. Finally, the heavy metals can be removed by biosorption into the algal-bacterial biomass produced during **PWW** bioremediation.

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SUPPLEMENTARY MATERIALS

Evaluation of the dynamics of microalgae population structure and process performance during piggery wastewater treatment in algalbacterial photobioreactors

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CONTENT

Figure S1 Figure S2 Table S1 Table S2

Evaluation of the dynamics of microalgae population structure and process performance during piggery wastewater treatment in algal-bacterial photobioreactors



Figure S1. Time course of the evaporation rates in R1 (Δ), R2 (\Diamond), R3 (\Box) and R4 (\circ) for the entire experiment.

Evaluation of the dynamics of microalgae population structure and process performance during piggery wastewater treatment in algal-bacterial photobioreactors



Figure S2. Time course of the concentration of TOC (**a**), IC (**b**) and TN (**c**) in the PWW (\bullet), R1 (Δ), R2 (\Diamond), R3 (\Box) and R4 (\circ) for the entire experiment.

HRT (d)	Qfeed (L/d)	Qeff (L/d)	% Evaporation	V PBR (L)	Height PBR (cm)	Area PBR (m ²)
27	0.112	0.044	60	3.0	15.8	0.019

 Table S1. Operational parameters and photobioreactor dimensions.

Parameter	PWW	R1	R2	R3	R4
TOC (mg/L)	1375	459	452	482	490
TOC (mg/d) in		153	153	153	153
TOC (mg/d) out		20.4	20.1	21.4	21.8
% TOC RE		87	87	86	86
IC (mg/L)	314	285	242	227	294
IC (mg/d) in		35	35	35	35
IC (mg/d) out		12.7	10.8	10.1	13.1
% IC RE		64	69	71	63
TN (mg/L)	393	174	166	165	149
TN (mg/d) in		44	44	44	44
TN (mg/d) out		7.7	7.4	7.3	6.6
% TN RE		82	83	83	85
Nitrite (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
Nitrate (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
TP (mg/L)	9.4	2.40	2.10	1.90	1.80
TP (mg/d) in		1.0	1.0	1.0	1.0
TP (mg/d) out		0.1	0.1	0.1	0.1
% TP RE		90	91	92	92
Zinc (mg/L)	0.7	0.9	1.1	1.3	0.9
Zn (mg/d) in		0.08	0.08	0.08	0.08
Zn (mg/d) out		0.04	0.05	0.06	0.04
% Zn RE		49	37	26	49
Copper (mg/L)	< 0.6	< 0.6	< 0.6	< 0.6	< 0.6
Arsenic (mg/L)	< 0.6	< 0.6	< 0.6	< 0.6	< 0.6
TSS (mg/L)	1340	2610	2569	2445	3265
Productivity (g/m ² /d)		6.1	6.0	5.7	7.6
С%		48.4	50.5	50.3	49.1
C (mg/d) biomass		56.1	57.7	54.7	71.2
% Carbon recovered		36	37	35	47
N %		7.9	8.2	7.4	7.6
N (mg/d)		9.2	9.4	8.0	11.0
biomass).2	2.4	0.0	11.0
% Nitrogen recovered		26	26	22	30
P%		0.7	0.8	0.8	0.7
P (mg/d)		0.8	0.9	0.9	1.0
biomass % Phosphorus		88	91	93	100

 Table S2. Mass balance calculations of PWW treatment under steady state conditions.

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Chapter 3





Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions
Comparative evaluation of piggery wastewater treatment in algalbacterial photobioreactors under indoor and outdoor conditions.

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Abstract

This work evaluated the performance of four algal-bacterial open photobioreactors operated at ≈ 26 days of hydraulic retention time during the treatment of 10 (\times 10) and 20 (\times 20) times diluted piggery wastewater (PWW) under indoor (I) and outdoor (O) conditions for four months. The removal efficiencies (REs) of organic matter, nutrients and zinc from PWW, along with the dynamics of biomass concentration and structure of algalbacterial population were assessed. The highest TOC-RE, TP-RE and Zn-RE 100% $(94\pm1\%)$ and $83\pm2\%$. respectively) were achieved indoors in

 $\times 10$ PWW, while the highest TN-RE $(72\pm8\%)$ was recorded outdoors in $\times10$ PWW. Chlorella vulgaris was the dominant species regardless of the ambient conditions and PWW dilution. DGGE-sequencing Finally, of the bacterial community revealed the occurrence of four phyla, Proteobacteria being the dominant phylum with 15 out of the 23 most intense bands.

Keywords

Algal-bacterial processes; Heavy metal biosorption; Microalgae-bacteria dynamics; Nutrient removal; Piggery wastewater biodegradation.

1. Introduction

Europe, with an annual production of 23.5 million th of pork meat, was the second largest pig producer in the world in 2015 (Statista, 2016). Europe's pig production accounted for 149 million heads, which represented approx. 44.3 % of the total European livestock in 2015 (EU. 2015; MAGRAMA, 2015). However, this relevant economic sector annually generates 217-434 million m³ of piggery wastewater (PWW) (4-8 L/d·pig) containing high concentrations of organic matter, nutrients, solids and heavy metals (De Godos et al., 2009; Franchino et al., 2016). The treatment of strength such high wastewaters represents both a technical challenge and a severe economic burden for the livestock sector. In this context, next generation PWW treatment technologies should allow complying with European wastewater regulations (1999/31/EC) (Council Directive, 1999) while producing added-value bioproducts out of the organic matter and nutrients **PWW** present in (2008/98/EC) (European Commisssion, 2008).

Algal-bacterial symbiosis has emerged as a promising platform for resource recovery and recycling from PWW in rural areas (where space is often not limiting). Algal-bacterial symbiosis has been successfully applied in photobioreactors for the treatment of domestic wastewater (García et al., 2017a; Oswald et al., 1957), digestates (Anbalagan et al., 2016; Wang et al., 2013), livestock effluents (Tigini et al., 2016), parboiled rice wastewater (Bastos et al., 2009), olive oil mill wastewater and wastewater from the pulp and paper industry (Muñoz and Guieysse, 2006). The use of microalgae during PWW treatment can support a cost-effective removal of organic matter, nutrients, heavy metals, pathogens and emerging pollutants as a result of their dual autotrophic and heterotrophic metabolisms, photosynthetic O₂ release and ability to increase the pH of the cultivation broth (García et al., 2017a; Muñoz and Guieysse, 2006). The ability of microalgae to grow on both wastewater alkalinity and the carbon dioxide (CO₂) released during organic matter oxidation entails 2-3 folds larger productivities (compared to activated sludge systems) of a biomass that can be used as a feedstock for the production of biofertilizers or bioenergy. In addition, the lower energy demand of microalgaebased wastewater treatment, along with the CO₂ fixation ability of microalgae, significantly increase the environmental sustainability of this technology (Cheah

et al., 2016; Dassey and Theegala, 2013). Despite the merits of algal-bacterial processes for PWW treatment and the intensive research conducted in this field in the past 10 years, very few studies have been carried out outdoors under the periodically fluctuating and high solar irradiations and temperatures (De Godos et al., 2009; García et al., 2017b; Posadas et al., 2017). In this context, the absence of comparative studies systematically assessing the representativeness of the results obtained indoors (under artificial irradiation and temperature controlled environments) compared to those supported by outdoors photobioreactors severely limits the use of most data available in literature for the design and operation of full-scale microalgae-based systems.

This work aimed at systematically evaluating the potential of open algalbacterial photobioreactors for the treatment of PWW under indoor and outdoor conditions. The removal of carbon, nitrogen, phosphorus and heavy metals was assessed at two PWW dilutions under solar and artificial illumination. Finally, the influence of both PWW dilution and environmental conditions on the structure of the microalgae and bacteria communities was investigated.

2. Materials and methods

2.1 Algal-bacterial inoculum and piggery wastewater

An acclimated Chlorella vulgaris culture, obtained from an indoor open algal-bacterial photobioreactor treating 15% diluted PWW at the Department of Chemical Engineering and Environmental Technology at Valladolid University (Spain), was used as inoculum. Fresh PWW was collected from a nearby farm at Cantalejo (Spain) and stored at 4 °C. The PWW was centrifuged for 10 minutes at 10000 rpm before dilution to reduce the concentration of suspended solids. The average composition of the 10 and 20 folds diluted PWW is shown in Table 1.

2.2 Experimental system

The indoors experimental set-up 3 consisted of two L open photobioreactors (15.8 cm depth, 15.5 cm internal diameter) illuminated at $1417\pm82 \ \mu mol/m^2 \cdot s$ for 12 hours a day (08h00 to 20h00) by LED lamps arranged in a horizontal configuration 60 cm above the photobioreactor surface under indoor conditions (Fig. 1, Table 1). Likewise, two similar open photobioreactors were located outdoors Department Chemical at the of Engineering and Environmental

Technology at Valladolid University (Spain). The average photosynthetic active radiation (PAR) in these systems at 11h00 was 1394 \pm 171 µmol/m² \cdot s (Fig. 1, Table 1). This value was comparable to the daily average PARs provided by the official AEMET meteorological station located at the University of Valladolid during the experimental period $(1210\pm126 \ \mu mol/m^2 \cdot s)$. The temperature of the indoor and outdoor photobioreactors was partially controlled using a water bath to prevent the high temperatures induced by both LEDs and solar irradiation. The algal-bacterial cultivation broth in the photobioreactors was gently mixed via water immersion

pumps. The indoors and outdoors photobioreactors were fed with both 10 and 20 times diluted PWW using an auto control 205U7CA multi-channel cassette pump (Watson-Marlow, UK). PWW were selected based dilutions on previous investigations carried out with this kind of wastewater and aiming to avoid microbial inhibition as а consequence of PWW toxicity (De Godos et al., 2009; García et al., 2017b; González et al., 2008). Pure CO_2 was added to the cultivation broth of the phobioreactors to automatically maintain the pH at 8.0 using a Crison multimeter M44 control unit (Crison Instruments, Spain).



Figure 1. Schematic diagram of the algal-bacterial photobioreactor set-up equipped with carbon dioxide supplementation for pH control under indoor and outdoor conditions.

2.3 Experimental design and sampling procedure

The indoors photobioreactors fed with 10 and 20 times diluted PWW (namely I-10 and I-20, respectively) and the outdoors photobioreactors fed with 10 and 20 times diluted PWW (namely O-10 and O-20, respectively) were inoculated with a fresh Chlorella vulgaris culture at an initial TSS concentration of \approx 680 mg/L (corresponding to an initial microalgae cell concentration of $\approx 1.06 \cdot 10^9$ cells/L, respectively). The photobioreactors, which were initially filled with tap water, were operated at an average hydraulic retention time (HRT) of ≈ 26 days for 120 days (from May-2016 to Sept-2016). A higher HRT than in conventional HRAPs (3-10 days) was chosen in this research to guarantee an effective carbon and nutrients removal, and to prevent toxicity effects on microbial population due the high loads of organic matter and nutrients of the PWW treated in this study (Aguirre et al., 2011; De Godos et al., 2009). The effluent from the photobioreactors overflowed separately as a function of the evaporation rates. Liquid samples from the influent PWWs and effluents of the photobioreactors

$$RE(\%) = \frac{(C_{feed} \times Q_{feed}) - (C_{eff} \times Q_{eff})}{C_{feed} \times Q_{feed}} \times 100$$

were taken weekly to determine the concentration of total organic carbon (TOC), inorganic carbon (IC), total nitrogen (TN), nitrate (NO3-), nitrite (NO_2) , total phosphorus (TP), zinc (Zn) and total suspended solid (TSS). Likewise, the structure of the microalgae population in the photobioreactors was periodically assessed from biomass samples preserved with lugol acid at 5% and formaldehyde at 10%, and stored at 4 °C prior to analysis. A cultivation broth sample from the photobioreactors was also collected under steady state (day 120) and immediately stored at -20 °C to evaluate the richness and composition of the bacterial communities (Alcántara et al., 2015). Dissolved oxygen (DO) concentration and temperature in the photobioreactors were measured twice per day (11h00 and 17h00), while the influents and effluents flowrates were daily recorded to monitor water evaporation losses (Table 1). Finally, the C, N and P content of the algal-bacterial biomass present in the photobioreactors was measured under steady state.

The removal efficiencies of C, N, P and Zn were calculated according to Eq. (1):

(1)

where C_{feed} and C_{eff} represent the dissolved concentrations of TOC, IC, TN, TP and Zn in the influent PWWs and photobioreactors effluents, respectively, while Q_{feed} and Q_{eff} represent the PWWs and effluents flow rate, respectively. The mass flow rate of C-CO₂ injected to control the pH was negligible compared to the input mass flow rate of C in the influent PWW (data not shown). The process was considered under steady state when the TSS concentrations in the photobioreactors remained stable for at least four consecutive samplings (~ 1 month). The results obtained were here provided as the average ± standard deviation from duplicate measurements along the one-month steady state period (days 91-120).

2.4 Analytical procedures

pH was in-situ measured using a Crison M44 multimeter and a Crison PH 28 meter. An OXI 330i oximeter was used to measure the DO and temperature (WTW, Germany). A LI-250A light meter (LI-COR Biosciences, Germany) was used to measure the light intensity as PAR. TOC, IC and TN concentrations were determined using a TOC-V CSH analyzer equipped with a TNM-1 module (Shimadzu, Japan). Nitrate and analyzed nitrite were by high performance liquid chromatography-ion

conductivity (HPLC-IC) (Posadas et al., 2013). N-NH₄⁺ was not analyzed based on the fact that no inhibition was expected at a pH of 8.0, where the ammonium share is greater than > 90%of the total nitrogen (Metcalf and Eddy, 2003). TP and TSS concentrations were determined according to Standard Methods (APHA, 2005). The analysis of the C, N and P biomass content in pregrinded algal-bacterial dried and biomass was carried out using a LECO CHNS-932 elemental analyzer. Zinc was determined using a 725-ICP Optical Emission Spectrophotometer (Agilent, USA) at 213.62 nm. The concentrations of arsenic and copper were not determined based on the results obtained by (García et al., 2017b), who observed that the concentration of these heavy metals in this PWW always remained below the detection limit (< 0.6 mg/L).

The identification and quantification of microalgae were conducted bv microscopic examination (OLYMPUS IX70, USA) according to Sournia (1978). Molecular analysis of the bacterial populations was carried out according with Frutos et al. (2015). The genomic desoxyribonucleic acid (DNA) was extracted using the protocol described in the Fast® DNA Spin Kit for Soil (MP Biomedicals, LLC) handbook.

The genes in the V6-V8 regions of the bacterial 16S ribosomal ribonucleic acid (rRNA) were amplified by Polymerase Chain Reaction (PCR) analysis using the universal bacterial primers 968-F-GC and 1401-R (Sigma-Aldrich, St. Louis, MO, USA) (Nübel et al., 1996). The denaturing gradient gel electrophoresis (DGGE) analysis of the amplicons was performed with a D-Code universal mutation system (Bio Rad Laboratories) using 8 % (w/v) polyacrylamide gels with a urea/formamide denaturing gradient of 45 to 65 %. DGGE running conditions were applied according to Roest et al. (2005). Sequences were deposited in GenBank Data Library under accession numbers MF380643 al MF380665. The Shannon-Wiener diversity index (H) was determined using the peak heights in the densitometric curves. This index, which reflects both the sample richness and evenness and ranges from 1.5 to 3.5 (low and high species evenness and richness,

respectively), can be calculated according to Eq. (2) (MacDonald, 2003): $H = -\Sigma \left[P_i ln(P_i) \right]$ (2)

Where *H* is diversity index and P_i is the importance probability of the bands in a lane ($P_i = n_i/n$, where n_i is the height of an individual peak and *n* is the sum of all peak heights in the densitometric curves). Similarity indices of the compared profiles were calculated from the densitometric curves of the scanned DGGE profiles by using the Pearson product-moment correlation coefficient (Häne et al., 1993). The taxonomic position of the sequenced DGGE bands was obtained using the RDP classifier tool (50 % confidence level) (Wang et al., 2007). The closest cultured and uncultured relatives to each band were obtained using the BLAST search tool at the NCBI database (National Centre for Biotechnology Information) (McGinnis and Madden, 2004).

Parameters		PWW (×10)	PWW (×20)	I-10	I-20	O-10	O-20
Operation period (days)		*	*	120	120	120	120
HRT (days)		*	*	≈26	≈26	≈26	≈26
pH (units)		*	*	8.0	8.0	8.0	8.0
PAR (µmol/m ² .s)		*	*	1417±82	1417±82	1394±171	1394±171
Torrestore (CC)	11h00	*	*	21±4	21±4	26±5	26±5
Temperature (°C)	17h00	*	*	24±4	24±4	35±5	35±5
	11h00	*	*	3.5±2.2	6.1±4.4	1.9 ± 0.8	3.9+1.7
Dissolved Oxygen (mg/L)	17h00	*	*	2.3±1.7	5.8±4.8	1.3+0.3	2.7±1.2
Evaporation rates (%)		*	*	27	27	44	44
TOC (mg/L)		963±71	497±33	80±5	91±5	150±11	133±8
IC (mg/L)		160±15	82±2	188±3	191±5	191±9	241±6
TN (mg/L)		341±27	170±3	201±6	162±5	168±6	158±7
Nitrate (mg/L)		< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
Nitrite (mg/L)		< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
TP (mg/L)		4.9±0.2	2.5±0.1	0.07±0.01	0.04 ± 0.01	1.67 ± 0.08	0.70 ± 0.07
Zinc (mg/L)		0.66±0.04	0.37±0.02	0.16±0.02	0.15±0.05	0.35 ± 0.03	0.32 ± 0.08
TSS (mg/L)		291±3	156±3	1284±71	720±16	1328±28	655±13
*Not applicable							

Table 1. Operational conditions and physical/chemical characterization of the piggery wastewater (PWW) and cultivation broth in the photobioreactors.

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3. Results and Discussion

3.1 Biodegradation of carbon, nitrogen and phosphorous

The range of temperatures, DO and PAR along with pH control resulted in a successful PWW treatment regardless of the environmental and operational conditions imposed (Table 1, Fig. 2). The higher evaporation rates under outdoor conditions (44% of the influent PWW flowrate compared to 27% under indoor conditions) resulted in a significant deterioration of the quality of the treated effluent (Table 1).

The TOC-REs accounted for 94 ± 1 , 87±2, 91±1 and 85±1% in I-10, I-20, O-10 and O-20, respectively, which resulted in average TOC concentrations in the effluent at the end of the operational period of 80 ± 5 , 91 ± 5 , 150 ± 11 and 133 ± 8 mg/L, respectively (Table 1, Fig. 2). These high REs were supported by the DOs $> 1 \text{ mg O}_2/\text{L}$ in the cultivation broth of the four photobioreactors mediated by an intense photosynthetic activity (Table 1). The slightly higher TOC-REs during the treatment of 10 times diluted PWW regardless of the environmental conditions can be explained by the differences in microbial population structure and biomass concentration

encountered in the photobioreactors under steady state. Thus, a higher share, diversity and concentration of bacteria compared to microalgae was present in the photobioreactors supplied with 10 times diluted PWW as revealed by the TSS measurements, DGGE analyses and microalgae population characterization. (Table 1). On the other hand, the higher DO concentrations recorded in the indoor photobioreactors regardless of the organic loading rate applied were likely caused by the lower temperatures (mediating a higher O₂ solubility and a lower bacterial activity) and by the constant PAR (resulting in a higher microalgal activity) (Posadas et al., 2015). The results here obtained were in agreement with the organic matter removal efficiencies reported by De Godos et al. (2009) under outdoor conditions in a 464 L HRAP operated at 10 days of HRT (COD-REs of 76±11% during the treatment of 10 and 20 folds diluted PWW). Likewise, Aguirre et al. (2011) recorded COD-REs \geq 90 % during the treatment of raw PWW in a 400 L **HRAPs** under outdoor environmental conditions at a HRT of 40 days. Finally, IC-REs of 13±3, -72±2, 33 ± 2 and $-67\pm13\%$ were recorded under steady state in I-10, I-20, O-10 and O-20, respectively, which resulted in average IC concentrations in the effluent of

188±3, 191±5, 191±9 and 241±6 mg/L, respectively (Table 1, Fig. 2). The negative IC-REs recorded in I-20 and O-20 resulted from the accumulation of inorganic carbon mediated by the high TOC oxidation activity in the systems, which was in agreement with the results obtained by (Posadas et al., 2013). The higher IC-REs during the treatment of 10 times diluted PWW were likely supported by the higher biomass concentrations (\approx 1300 mg TSS/L in I-10 and O-10 compared to \approx 700 mg TSS/L in I-20 and O-20). In this context, process operation at high biomass concentrations in photobioreactors under high PARs can prevent microalgae photoinhibition and thus induce high photosynthetic activities. Carbon removal by stripping (prior mineralization of the organic carbon to main CO_2) was the mechanism accounting for carbon removal in I-10 and O-10, since only 49 and 37 % of the total carbon removed was recovered in form of harvested biomass. the respectively.

The TN-REs accounted for 56 ± 4 , 30 ± 14 , 72 ± 8 and $48\pm9\%$ in I-10, I-20, O-10 and O-20, respectively, which resulted in average TN concentrations in the effluent under steady state conditions of 201 ± 6 , 162 ± 5 , 168 ± 6 and 157 ± 7 mg/L,

respectively (Table 1, Fig. 2). The higher temperatures prevailing outdoors likely lowered NH₃ solubility and therefore increased N removal by stripping (Fig. 2) (Metcalf and Eddy, 2003). In addition, the systems supporting higher biomass concentrations (I-10 and O-10) mediated higher N removals compared to process operation with 20 folds diluted PWW (Fig. 2). The TN-REs here achieved were similar to those obtained by Aguirre et al. (2011) who reported TN-REs ranging from 65 to 85% during PWW treatment in 400 L HRAPs operated at HRTs of 40-80 days, but lower than those reported by García et al. (2017b) (82 - 85%) during the treatment of 15% diluted PWW in open photobioreactors at a HRT of ≈ 27 days operated indoors. The nitrogen mass balances conducted revealed that stripping was the main N removal mechanism in I-10, O-10 and O-20, with nitrogen assimilation into biomass accounting for only 44, 25 and 37% of the total nitrogen removed, respectively. However, nitrogen assimilation was the main N removal mechanism in I-20 (85% of TN removed) likely due to the lower temperatures TN and concentrations prevailing in the cultivation both.

The TP-REs accounted for 99 ± 3 , 99 ± 5 , 81 ± 8 and $84\pm13\%$ in I-10, I-20, O-10 and O-20, respectively, which resulted in

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average TP concentrations in the effluent of 0.07 ± 0.07 , 0.04 ± 0.08 , 1.67 ± 0.08 and 0.70 ± 0.07 mg/L, respectively, under steady state (Table 1, Fig. 2). The fact that higher TP-REs were recorded indoors regardless of the PWW dilution and biomass concentration at a constant pH requires further investigation. The TP-REs herein obtained were similar to those reported by García et al. (2017b) during the treatment of 15% diluted PWW in indoors open photobioreactors (90 to 92%). Phosphorous assimilation into algal-bacterial biomass was likely the main removal mechanism based on the moderate pH values prevailing in the photobioreactors during the entire experiment (pH=8.0), which did not significant support а phosphate precipitation (García et al., 2017a). Thus, a phosphorus mass balance revealed that 100, 99, 100 and 100% of the total removed phosphorus was recovered in the harvested biomass in I-10, I-20, O-10 and O-20 respectively.



Figure 2. Average removal efficiencies of TOC (\square), IC (\blacksquare), TN (\blacksquare) and TP (\blacksquare) under steady state. Bold numbers indicate the steady state removal efficiencies, while vertical bars represent the standard deviation from replicate measurements during steady state operation.

3.2 Heavy metal removal

The overall steady state Zn-REs in I-10, I-20, O-10 and O-20 accounted for 83 ± 2 , 71 ± 9 , 70 ± 5 and $51\pm13\%$, respectively, which resulted in average Zn concentrations at the end of the operational period of 0.16 ± 0.02 , 0.15±0.05, 0.35±0.03 and 0.32±0.08 mg/L, respectively (Table 1). In this context, the higher abundance of microalgae and cyanobacteria induced by laboratory conditions likely supported the higher Zn-REs recorded indoors, while the higher biomass

concentrations in the HRAPs treating 10 folds diluted PWW explained the superior Zn-REs in I-10 and O-10 compared to I-20 and O-20, respectively. The latter suggests that biosorption was the main mechanism governing Zn removal (Javanbakht et al., 2014; Kaplan et al., 1987). The Zn-REs herein obtained were higher than those reported by García et al. (2017b) during PWW treatment in 3 L indoors HRAPs operated at a HRT of \approx 27 days (26 to 49%).

3.3 Concentration, productivity and elemental composition of the algal-bacterial biomass

The algal-bacterial biomass concentration in I-10, I-20 and O-20 initially decreased from 680 mg TSS/L to 127, 177 and 170 mg TSS/L, respectively, during the first 28 days of operation, while biomass concentration slightly increased from 680 to 800 mg TSS/L in O-10 during process start-up (Fig. 3). The previous acclimation of microalgae to the pollutants loading rate and environmental conditions imposed to I-20 explain this increase in biomass concentration. Biomass concentration increased exponentially afterwards in I-10 and O-10 up to steady state values of 1284±71 mg TSS/L and 1328±28 mg TSS/L, respectively. Similarly, biomass concentration in I-20 and O-20 increased up to steady state values of 720±16 and 620±79 mg TSS/L, respectively (Fig. 3). Indeed, the steady state biomass concentrations in systems supplied with 10 folds diluted PWW were ~2 times higher than those recorded in the photobioreactors fed with 20 folds diluted PWW (Table 1). On the other hand, the higher water evaporation rates in the outdoor photobioreactors resulted in slightly lower biomass productivities: 5.6, 3.1, 4.8 and 2.4 g/m²·d in I-10, I-20, O-10 and O-20, respectively. The lower productivities biomass recorded outdoors could be also explained by the effects pernicious on microbial metabolism caused by the high and fluctuating temperatures and irradiations. These biomass productivities were comparable to those reported by García et al. (2017b) under indoor conditions during the treatment of 15% diluted PWW in 3 L HRAPs operated at ≈ 27 days of HRT (5.8 to 7.8 $g/m^2 \cdot d$).

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Figure 3. Time course of TSS concentration in I-10 (\bullet), O-10 (\circ), I-20 (\blacktriangle) and O-20 (Δ).

The C, N and P content of the biomass cultivated indoors averaged 49.3±0.6, 8.6 ± 0.4 and $0.52\pm0.06\%$ (which entailed a C/N/P of 100/17/1), respectively, and $46.0{\pm}0.02, \hspace{0.2cm} 8.2{\pm}0.02 \hspace{0.2cm} and \hspace{0.2cm} 0.58{\pm}0.02\%$ (C/N/P of 100/17/1), respectively, when cultivated outdoors. These elemental compositions were in agreement with those reported by Cabanelas et al. (2013), who observed a C, N and P content in the harvested biomass of ≈ 44 , 7.5 and 0.5%, respectively, in a photobioreactor inoculated with Chlorella vulgaris and supplemented with CO₂ during the treatment of settled domestic wastewater.

3.4 Time course of the microalgae population structure

Chlorella vulgaris, which achieved a maximum cell concentration of $1.74 \cdot 10^9$ cells/L by day 92, represented the dominant photosynthetic species in I-10

throughout the entire experimental period. Pseudanabaena sp. was also identified in I-10 from day 92 onwards at concentrations of $\approx 0.30 \cdot 10^9$ cells/L (Fig. 4a). A similar microalgae population dynamics was recorded in I-20, with C. vulgaris representing the dominant species with maximum cell concentrations of $2.95 \cdot 10^9$ by days 42 and 70. However, Pseudanabaena sp. became dominant by day 120 with a concentration of $0.46 \cdot 10^9$ cells/L as a result of the gradual decrease in C. vulgaris population from day 70 (Fig. 4b). C. vulgaris was also dominant in the photobioreactors operated outdoors regardless of the PWW dilution applied. However, the maximum concentration of C. vulgaris in O-10 was recorded in the inoculum $(0.52 \cdot 10^9 \text{ cells/L})$, with a gradual decrease afterwards. Pseudanabaena sp. was identified by days 105 and 120 at concentrations of $0.15 \cdot 10^9$ and $0.08 \cdot 10^9$ cells/L. respectively, in O-10. In addition. Aphanothece sp. was also detected from day 70 to 120 in O-10, but at negligible concentrations (Fig. 4c). Finally, the maximum cell concentration of C. *vulgaris* in O-20 was $1.73 \cdot 10^9$ cells/L by day 25, with Acutodesmus obliquus (identified by day 25) and Aphanothece sp. (identified by days 56 and 72) detected at negligible concentrations, and *Pseudanabaena* sp. identified by day 92 at a concentration of $0.59 \cdot 10^9$ cells/L (Fig. 4d). The high tolerance of C. vulgaris to organic and heavy metals pollution likely supported the observed dominance of this microalga regardless of the operational and environmental conditions. Thus, C. vulgaris ranked 11/80 in the ranking of pollution-tolerant microalgae species published by Palmer (1969), while the Chlorella ranked 5/60 at a genus level. Process inoculation with C. vulgaris at a high concentration, along with the high tolerance of this microalga to organic pollution, guaranteed its longterm dominance and an effective PWW treatment. Pseudanabaena sp., which

belongs to the order of Oscillatoriales, was also identified at relevant concentrations under steady state (Acinas et al., 2009). The tolerance of Pseudanabaena sp to organic pollution herein recorded was in agreement with the observations of García et al. (2017a), who identified *Pseudanabaena* sp. during the treatment of domestic wastewater in enclosed an photobioreactor at a HRT of 2 day, and by Serejo et al. (2015) during the treatment of digested vinasse in a 180 L HRAP. This study also suggested that the high and fluctuating temperatures and irradiations prevailing under outdoors operation resulted in both a reduced population of microalgae and cyanobacteria compared to indoors in lower cultures. and biomass productivities (Fig. 4a, 4b). Finally, the microalgae highest concentration recorded during the treatment of 20 times regardless diluted PWW of the environmental conditions was likely caused by the lower toxicity at increasing PWW dilutions.



Figure 4. Time course of the microalgae population structure in I-10 (a), I-20 (b), O-10 (c) and O-20 (d). Acutodesmus obliquus ⊠, Aphanothece sp. , Chlorella vulgaris , Pseudanabaena sp. , and total numbers of cells (●).

3.5 Bacteria population structure

The DGGE analysis of the microbial communities present in the open photobioreactors revealed the occurrence of 4 phyla and 23 bands (Fig. 5). Proteobacteria, which is ubiquitous in the environment, was the dominant phylum (15 out of 23 bands sequenced) in the inoculum and in all photobioreactors (bands 1-15) (Fig. 5) (Shin et al., 2015). Despite not present in the inoculum, the phylum *Bacteroidetes* was identified under steady state in all photobioreactors (bands 16-20). The phylum Firmicutes was identified in the inoculum (bands 21 and 22) and in O-20, while the phylum Cyanobacteria/Chloroplast

corresponded to band 23 in the inoculum and in I-10, I-20 and O-10 (Fig. 5). In this context, the open nature of the with photobioreactors, along the different environmental conditions and characteristics of the PWW fed, likely induced the enrichment of photobioreactor-specific bacterial populations different from the inoculum. Bacteria from the phyla Proteobacteria, Bacteriodetes and Firmicutes were likely the responsible for the biodegradation of organic matter in the photobioreactors. Thus, bacteria from the phylum *Proteobacteria* belonging to the genus Psychrobacter (I-10, I-20, O-

10 and O-20), the class Betaproteobacteria (I-10, I-20, O-10 and O-20) and the genus Thauera (I-10, I-20) and O-10) have been identified in synthetic wastewater, swine effluents, anaerobic digesters treating feedstock from cheese manufacturing, wastewater from dye industry and anoxic biotrickling filters treating BTEX, which confirmed the capacity of these microorganisms biodegrade to the organic pollutants present in PWW (Akmirza et al., 2017; Lucas et al., 2013). Similarly, bacteria from the phylum **Bacteriodetes** have been identified during the anoxic removal of BTEX, Laboratory-scale partial nitrifying-ANAMMOX reactor and municipal wastewater treatment (Fig. 5) (Akmirza et al., 2017; Biswas and Turner, 2012). Finally, bacteria from the *Firmicutes* phylum (syntrophic microorganisms) were detected in a SBR reactor treating swine waste (Loureiro, 2008; Rivière et al., 2009).

The Shannon-Wiener diversity indexes (H) of the inoculum, I-10, I-20, O-10 and O-20 were 2.66, 2.69, 2.72, 2.63 and 2.17, respectively (Fig. 5). The photobioreactors operated in this study exhibited a relatively low-medium bacterial diversity (H \approx 2.6) likely due to the extreme environmental conditions

applied and to the high toxicity of the wastewater treated. The analysis of the similarity indexes (76.3% between I-10 and I-20 and 76.3% between O-10 and O-20) showed high similarities between the respective indoor and outdoor photobioreactors. On the other hand, low similarity indexes were recorded between I-10 and O-10 (14.2%) and I-20 and O-20 (41.6%). Thus, these results confirmed that temperature and irradiation under indoor and outdoor conditions can result in significantly different bacterial population structure. These results were in agreement with the findings reported by Ferrero et al. (2012),who observed that environmental parameters such as temperature or the impinging irradiation can play a more important role than organic matter and nutrients loading in the structure of the bacterial community.



Figure 5. Bacterial DGGE profile of the microbial communities in the inoculum (Inoc.) and in the open algal-bacterial photobioreactors I-10, I-20, O-10 and O-20. Horizontal arrows and numbers indicate the most abundant bacterial communities. The name of the samples and the Shannon-Wiener diversity indexes (H) are also shown in the upper part of the gel profiles.

4. Conclusions

This work demonstrated for the first time that neither pollutant removal nor the structure of microalgae and bacterial communities under indoor conditions can be directly extrapolated to outdoors photobioreactors. Unexpectedly, the lowest PWW dilution always resulted in a superior PWW treatment performance. The dominance of Chlorella vulgaris in all photobioreactors regardless of the environmental conditions and PWW dilution confirmed the high pollutiontolerance of this species. The DGGE analysis revealed a high dominance of the Proteobacteria phylum in all photobioreactors, and the key influence of temperature and irradiation on the final bacterial population structure.

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SUPPLEMENTARY MATERIAL

Comparative evaluation of piggery wastewater treatment in algalbacterial photobioreactors under indoor and outdoor conditions

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Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions



Figure S1. Time course of the temperature at 11h00 (a) and 17h00 (b) in I-10 (\bullet), I-20 (\blacktriangle), O-10 (\circ) and O-20 (Δ) throughout the experimental period.

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions



Figure S2. Time course of the dissolved oxygen concentration at 11h00 (a) and 17h00 (b) in I-10 (\bullet), I-20 (\blacktriangle), O-10 (\circ) and O-20 (\bigtriangleup) throughout the experimental period.

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions



Figure S3. Time course of the average PAR under indoor (\blacksquare) and outdoor (\square) conditions.

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions



Figure S4. Time course of the concentration of TOC (a), IC (b) TN (c) and TP (d) in the 10 fold diluted PWW (\blacksquare), I-10 (\bullet) and O-10 (\circ).

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions



Figure S5. Time course of the concentration of TOC (a), IC (b) TN (c) and TP (d) in the 20 fold diluted PWW (\blacksquare), I-20 (\blacktriangle) and O-20 (Δ).

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions

Table S1. Operational parameters and photobioreactor dimensions under indoor conditions.

HRT (d)	Qfeed (L/d)	Qeff (L/d)	Evaporation %	V PBR (L)	Height PBR (cm)	Area PBR (m ²)
27	0.113	0.082	27	3.0	15.8	0.019

PBR: Photobioreactor

PWW (×10)	PWW (×20)	I-10	I-20
963	497	80	91
		108.3	55.9
		6.6	7.5
		94	87
160	82	188	191
		18.0	9.2
		15.4	15.7
		14	-70
341	170	201	162
		38.4	19.1
		16.5	13.3
		57	30
< 0.5	< 0.5	< 0.5	< 0.5
< 0.5	< 0.5	< 0.5	< 0.5
4.9	2.5	0.07	0.04
		0.55	0.28
		0.01	0.00
		99	99
0.67	0.37	0.16	0.15
		0.08	0.04
		0.01	0.01
		83	70
291	156	1284	720
		5.55	3.11
		48.82	49.72
		51.48	29.40
		49	70
		8.88	8.37
		9.36	4.95
		43	85
		0.56	0.47
			0.47
		0.39	0.20
		100	100
	963 160 341 <0.5 <0.5 4.9 0.67	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	963 497 80 108.3 6.6 94 94 160 82 188 160 82 188 160 82 188 160 82 188 160 82 188 160 82 188 160 82 188 160 82 188 170 201 38.4 16.5 57 57 < 0.5 < 0.5 < 0.5 < 0.5 < 0.5 < 0.5 < 0.5 < 0.5 < 0.5 < 0.55 0.07 0.01 99 0.67 0.37 0.16 0.08 0.01 0.08 0.01 83 291 156 1284 5.55 48.82 $5.1.48$ 49 43 9.36 43 9.36

Table S2. Mass balance calculations of PWW treatment under steady state in the indoor

 photobioreactors.

Comparative evaluation of piggery wastewater treatment in algal-bacterial photobioreactors under indoor and outdoor conditions

 Table S3. Operational parameters and photobioreactor dimensions under outdoors

conditions.	PBR:	Photobioreactor	

HRT (d)	Qfeed (L/d)	Qeff (L/d)	Evaporation %	V PBR (L)	Height PBR (cm)	Area PBR (m ²)
25	0.122	0.068	44	3.0	15.8	0.019

Parameter	PWW (×10)	PWW (×20)	O-10	O-20
TOC (mg/L)	963	497	150	133
TOC (mg/d) in			117	61
TOC (mg/d) out			10.2	9.1
% TOC RE			91	85
IC (mg/L)	160	82	191	241
IC (mg/d) in			20	10
IC (mg/d) out			13	16
% IC RE			33	-65
TN (mg/L)	341	170	168	158
TN (mg/d) in			41.6	20.7
TN (mg/d) out			11.5	10.8
% TN RE			72	48
Nitrite (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5
Nitrate (mg/L)	< 0.5	< 0.5	< 0.5	< 0.5
TP (mg/L)	4.9	2.5	1.67	0.7
TP (mg/d) in			0.60	0.31
TP (mg/d) out			0.114	0.048
% TP RE			81	84
Zinc (mg/L)	0.67	0.37	0.35	0.32
Zn (mg/d) in			0.08	0.05
Zn (mg/d) out			0.02	0.02
% Zn RE			71	52
TSS (mg/L)	291	156	1328	655
Productivity (g/m ² /d)			4.78	2.36
С %			46.0	46.0
C (mg/d) biomass			41.7	20.6
% Carbon recovered			37	46
N %			8.19	8.22
N (mg/d) biomass			7.43	3.68
% Nitrogen recovered			25	37
P%			0.57	0.59
P (mg/d) biomass			0.52	0.26
% Phosphorus recovered			100	100

Table S4. Mass balance calculations of PWW treatment under steady state in the outdoor photobioreactors.

Taxonomic placement (50% confidence level)	Band nº	Inoc.	I-10	I-20	O-10	O-20	Closest relatives in Blast Name (Accession number)	Similarity (%)	Source of origin
Phylum Proteobacteria	1	-	-	-	Х	-	Uncultured bacterium (JQ320097)	87%	Soil polluted with BDE209 and Cd
Class Gammaproteobacteria	2	-	XX	XX	XX	-	Uncultured bacterium (JQ300186)	90%	Soil
Order Pseudomonadales									
Family Moraxellaceae									
Genus Psychrobacter	3	-	-	-	-	XX	Uncultured <i>proteobacterium</i> (JQ218906)	96%	Marine macro-alga
							Psychrobacter sp. Bsw21512 (GQ358937)	96%	Seawater
							Uncultured bacterium (JF332609)	96%	Duodenal biopsy
	4	-	-	-	-	XX	Uncultured <i>Psychrobacter</i> sp. (JQ999390)	97%	Lake Vostok accretion ice (Antarctica)
							Uncultured bacterium (JF332609)	97%	Duodenal biopsy
							<i>Psychrobacter</i> sp. (KY406049)	96%	Soil sample from penguin breeding colony
	5	-	XX	XX	X	XXX	<i>Psychrobacter piscatorii</i> (NR_112807)	99%	Wastewater
							<i>Psychrobacter psychrophilus</i> (DQ337513)	99%	Swine effluent holding pit
	6	-	XX	XX	X	XXX	<i>Psychrobacter</i> sp. Mixed culture X14-2 (KR029412)	99%	Bioaerosol emitted from wastewate treatment plant
							Uncultured bacterium (JF332609)	99%	Duodenal biopsy
							Uncultured bacterium (KR514346)	99%	Bovine reproductive tract
	7	-	-	-	XXX	-	Uncultured Psychrobacter	100%	Lake Vostok accretion ice

Table S5. RDP classification of the DGGE bands sequenced and corresponding matches (BLASTN) using the NCBI database with indication of the similarity percentages and sources of origin. The presence/absence of each band in each sample, together with its intensity, are also shown.

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Taxonomic placement (50% confidence level)	Band n°	Inoc.	I-10	I-20	O-10	O-20	Closest relatives in Blast Name (Accession number)	Similarity (%)	Source of origin
							sp. (JQ999390)		(Antarctica)
							<i>Psychrobacter</i> sp. Mixed culture X14-2 (KR029412)	100%	Bioaerosol emitted from wastewater treatment plant
							Psychrobacter sp. KHH8 (KT368953)	100%	-
	8	-	-	-	-	XX	<i>Psychrobacter</i> sp. Mixed culture X14-2 (KR029412)	97%	Bioaerosol emitted from wastewater treatment plant
							Uncultured Psychrobacter sp. (JQ999390)	97%	Lake Vostok accretion ice (Antarctica)
Class Betaproteobacteria	9	XX	XXX	XX	XXX	-	Uncultured bacterium (GU390196)	86%	Anaerobic digester treating feedstock
	10	XXX	XXX	XXX	-	XXX	Acinetobacter sp. HPC497 (AY854128)	89%	Wastewater from dye industry
	11	XXX	XXX	-	XXX	-	Uncultured bacterium (KU991981)	87%	Anoxic removal of BTEX compounds
Order Rhodocyclales									
Family Rhodocyclaceae									
Genus Thauera	12	-	XX	XX	XXX	-	Uncultured bacterium (HG380609)	97%	Wastewater
							Uncultured beta proteobacterium (AF450463)	97%	Full-scale aerated-anoxic wastewater treatment plant
							Uncultured <i>Thauera</i> sp. (KX914731)	97%	Activated sludge
	13	-	XX	-	XXX	-	<i>Thauera</i> sp. (MF155554)	98%	Wastewater treatment plant
							Uncultured bacterium (HG380609)	98%	Wastewater
							Uncultured beta proteobacterium (AF450463)	98%	Full-scale aerated-anoxic wastewater treatment plant
Class Alphaproteobacteria	14	Х	XX	XX	XX	-	Uncultured bacterium (KT200337)	86%	Algal-bacterial biomass from an air- lift bioreactor treating toluene,

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Taxonomic placement (50% confidence level)	Band nº	Inoc.	I-10	I-20	O-10	O-20	Closest relatives in Blast Name (Accession number)	Similarity (%)	Source of origin
									inoculated with activated sludge, Pseudomonas putida and Chlorella Sorokiniana
Order Rhizobiales	15	XX	-	-	-	X	Iron-reducing bacterium enrichment culture clone fec_1_F2 (FJ802355)	89%	Danube River sediment
Phylum Bacteroidetes	16	-	-	Х	-	-	Uncultured bacterium (KU991981)	90%	Anoxic removal of BTEX compounds
							Uncultured bacterium (KU650792)	90%	Anaerobic full-scale reactors
							Uncultured bacterium (JN087868)	90%	Nitrifying bioreactor under inorganic carbon limitation
	17	-	XX	-	XX	-	Uncultured bacterium (HQ640531)	98%	Laboratory-scale partial nitrifying- ANAMMOX reactor
							Uncultured bacterium (KU650792)	98%	Anaerobic full-scale reactors
							Uncultured bacterium (JN087868)	98%	Nitrifying bioreactor under inorganic carbon limitation
	18	-	XXX	XX	XXX	-	Uncultured bacterium (HQ640531)	98%	Laboratory-scale partial nitrifying- ANAMMOX reactor
							Uncultured bacterium (KU650792)	98%	Anaerobic full-scale reactors
							Uncultured bacterium (JN087868)	98%	Nitrifying bioreactor under inorganic carbon limitation
Class Cytophagia									
Order Cytophagales	19	-	-	-	X	-	Uncultured bacterium (HQ640531)	100%	Laboratory-scale partial nitrifying- ANAMMOX reactor
							Uncultured bacterium (KU650792)	99%	Anaerobic full-scale reactors
Class Sphingobacteria									
Order Sphingobacteriales	20	-	XX	Х	XX	-	Uncultured bacterium (KU650792)	99%	Anaerobic full-scale reactors

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Taxonomic placement (50% confidence level)	Band nº	Inoc.	I-10	I-20	O-10	O-20	Closest relatives in Blast Name (Accession number)	Similarity (%)	Source of origin
							Uncultured bacterium (JN087868)	99%	Nitrifying bioreactor under inorganic carbon limitation
							Uncultured bacterium (KU991981)	99%	Anoxic removal of BTEX compounds
Phylum Firmicutes									
Class Clostridia									
Order Clostridiales	21	XX	-	-	-	-	Uncultured bacterium (GQ132773)	86%	SBR reactor treating swine waste; reactor 2, day 809; temperature: 35 deg C; ammonia: 1,800 mg N/L; solids loading rate: 2.2 g VS/L/day
	22	XX	-	-	-	XX	Uncultured bacterium (KP797907)	87%	Microalgae from HRAP treating diluted vinasse with wastewater treatment plant activated sludge
Phylum Cyanobacteria/Chloroplast									
Class Chloroplast Family Chloroplast									
Genus Chlorophyta	23	XX	XX	XXX	XXX	-	Uncultured bacterium (KT200344)	98%	Algal-bacterial biomass from an air- lift bioreactor treating toluene, inoculated with activated sludge, <i>Pseudomonas putida</i> and <i>Chlorella</i> <i>Sorokiniana</i>
							Plastid <i>Chlorella</i> sp. UMPCCC 1110 (KM218897)	98%	Water
							Uncultured <i>Chlorella</i> (KC994689)	96%	Microalgae photobioreactor

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Chapter 4





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Abstract

This study evaluated the performance of two open photobioreactors operated with a consortium of microalgae-bacteria (PBR-AB) and purple photosynthetic bacteria (PBR-PPB) during the continuous treatment of diluted (5%) piggery wastewater (PWW) at multiple hydraulic retention times (HRT). At a HRT of 10.6 days, PBR-AB supported the highest removal efficiencies of nitrogen, phosphorous and zinc (87±2, 91 ± 3 and $98\pm1\%$), while the highest organic carbon removals were achieved in PBR-PPB (87±4%). The decrease in

HRT from 10.6, to 7.6 and 4.1 day caused a gradual deterioration in organic material and nitrogen removal, but did not influence the removal of phosphorus and Zn. The decrease in HRT caused a severe wash-out of microalgae in PBR-AB and played a key role on the structure of bacterial population in both photobioreactors. In addition, batch biodegradation tests at multiple PWW dilutions (5, 10 and 15%) confirmed the slightly better performance of algalbacterial systems regardless of PWW dilution.

Keywords

Algal-bacterial processes; Photobioreactor; Photosynthetic biodegradation; PPB; Swine manure.

1. Introduction

The large volumes of wastewater yearly generated from domestic, industrial and agricultural activities demand a rapid and cost-effective wastewater treatment prior discharge into natural water bodies. More specifically, an insufficient treatment of agroindustrial effluents, which rank among the highest strengths wastewaters, can cause severe episodes of eutrophication in surface waters and pollution of groundwater (Mateo-sagasta and Burke, 2012). Only in the European Union, 215-430 Mm³ of piggery wastewaters (PWW) with [COD] > 50 g L^{-1} or [TN] > 5 g L^{-1} are annually generated (García et al., 2018; statista, 2018). Conventional agricultural (WWT) wastewater treatment technologies (e.g. activated sludge, trickling filters) are highly energy intensive and entail a significant loss of valuable nutrients. In this context, photosynthetic treatments have emerged as a cost-effective alternative to conventional WWT based on their potential to support a superior nutrients and carbon recovery from agricultural wastewaters.

Photosynthetic WWT has been traditionally based on the cultivation of microalgae, which produce O₂ and assimilate nutrients using the visible spectrum of sunlight as energy source, in symbiosis with heterotrophic and nitrifying bacteria. More specifically, microalgal-bacterial consortia can support an efficient removal of organic matter, nutrients, heavy metals and pathogens as a result of their dual autotrophic and heterotrophic metabolism (Rittmann and McCarty, 2012). This symbiosis results in a low consumption and carbon energy footprint since the CO₂ generated during organic matter oxidation is photosynthetically fixed (Cheah et al., 2016; Dassey and Theegala, 2013). Algal-bacterial processes have been successfully tested for the treatment of domestic wastewater (García et al., 2017a; Oswald et al., 1957), centrates (Posadas et al., 2017), vinasse (Serejo et al., 2015), digested livestock effluents (Franchino et al., 2016; Tigini et al., 2016), parboiled rice wastewater (Bastos et al., 2009) and PWW (de Godos et al., 2010; García et al., 2018, 2017b).

On the other hand, purple phototrophic bacteria (PPB) can also use solar radiation (the infrared spectrum) as during energy source anoxygenic photosynthesis, which requires electron donors such as organic matter and nutrients to built-up PPB biomass (Bertling et al., 2006). PPB can support high rates of organic matter and nutrient assimilation and exhibit a high tolerance towards wastewater toxicity. Furthermore, there is an increasing interest in PPB-based WWT since PPB able are to synthesize polyhydroxybutyrates (PHB) and polyphosphates, and possess a more versatile metabolism than microalgae (Hülsen et al., 2014). PPB have been recently used to treat domestic wastewaters (Hülsen et al., 2016a, 2016b, 2014; Zhang et al., 2003), PWW (Myung et al., 2004), rubber sheet wastewater (Kantachote et al., 2005), pharmaceutical wastewater (Madukasi et al., 2010) and fish industry effluent (de Lima et al., 2011) with promising results. However, while microalgae-based WWT has been evaluated both indoors and outdoors in open and enclosed photobioreactors from lab scale to industrial facilities (Craggs et al., 2012; de Godos et al., 2016), PPB-based WWT has been only evaluated indoors under lab scale conditions. In this context, there

is a lack of comparative studies systematically assessing the treatment capacity of both phototrophic microorganisms.

This work aimed at systematically evaluating the performance of open algal-bacterial and PPB photobioreactors for the indoor treatment of PWW under artificial illumination. The influence of the hydraulic retention time (HRT) on the removal of carbon, nitrogen, phosphorous and heavy metals, and on the structure of the algal-bacterial and PPB population was investigated. In addition, the influence of PWW dilution on the biodegradation performance of an algal-bacterial consortium and PPB was assessed batchwise.

2. Materials and methods

2.1 Inocula and piggery wastewater A Chlorella vulgaris culture obtained from an outdoors high rate algal pond (HRAP) treating centrate was used as inoculum in the algal-bacterial photobioreactor. The PPB inoculum used was obtained from a batch enrichment in diluted PWW (17%) under continuous infrared light illumination at 50 W/m². Fresh PWW was collected from a nearby swine farm at Cantalejo (Spain) and stored at 4 °C. The PWW

was centrifuged for 10 min at 10000 rpm before dilution to reduce the concentration of TSS. The average composition of the 5% diluted PWW is shown in Table 1.

2.2 Batch PWW biodegradation tests

An algal-bacterial (AB) batch test was conducted in three gas-tight glass bottles of 1.1 L illuminated by LED lamps at $1380\pm 24 \,\mu mol/m^2 \cdot s \,(302.2 \,W/m^2)$ for 12 hours a day. Similarly, a purple phototrophic bacteria (PPB) batch test was carried out in 1.1 L gas-tight glass bottles illuminated by IR lamps at 45±1 W/m^2 for 12 hours a day. Both light intensities were selected to simulate the natural sun radiation conditions of PAR and IR (García et al., 2017b; Hülsen et al., 2016a). The bottles were initially filled with 400 mL of 5, 10 and 15% diluted PWW and inoculated with fresh biomass at 760 mg TSS/L. The algalbacterial inoculum, obtained from the algal-bacterial photobioreactor (PBR-AB), was composed of Chlamydomonas kessieri, sp., Chlorella Chlorella vulgaris and Scenedesmus acutus, which represented 14, 23, 43 and 20% of the algal population, respectively. Similarly, the PPB inoculum, obtained from the PPB photobioreactor (PBR-PPB), was mainly composed of bacteria from the phyla *Proteobacteria*, *Synergistetes*, *Firmicutes*, which represented 83.8, 5.3 and 3.6 % of the bacterial population, respectively.

All bottles were flushed with N₂ for 10 minutes establish to an initial environment totally deprived from O₂. The tests were incubated at 30 °C (using thermostatic water baths) under continuous magnetic agitation (200 rpm). Liquid samples of 20 mL were periodically taken to determine the pH and concentration of total suspended solids (TSS), total organic carbon (TOC), inorganic carbon (IC) and total nitrogen (TN). In addition, gas samples from the headspace of the bottles were daily drawn using gas-tight syringes to by GC-TCD determine the gas concentration of N₂, CO₂ and O₂.

2.3 PWW biodegradation in continuous photobioreactors

The experimental set-up consisted of two 3L open photobioreactors (0.15 m deep and 0.02 m² of superficial area). The PBR-AB was illuminated at 1393 \pm 32 µmol/m²·s for 12 hours a day (04h00 to 16h00) using visible LED lamps arranged in a horizontal configuration 60 cm above the surface of the PBR (Fig. 1). The PBR-PPB was illuminated at 48 \pm 4W/m² for 12 hours a day (04h00 to

16h00) by IR LED lamps arranged in a horizontal configuration 20 cm above the surface of the PBR (Fig. 1). The PBR-AB was jacketed and connected to a cooling water bath to maintain similar temperatures in both PBRs. The cultivation broths of PBR-AB and PBR-PPB were mixed via two water immersion pumps. Both photobioreactors were initially filled with tap water, inoculated with fresh biomass at 275 mg TSS/L and fed with 5% diluted PWW using a 205U7CA multi-channel cassette pump (Watson-Marlow, UK) at HRTs of 10.6, 7.6 and 4.1 days in stage I, II and III, respectively (Table 1).



Figure 1. Schematic diagram of the algal-bacterial photobioreactor (PBR-AB) and purple photosynthetic bacteria photobioreactor (PBR-PPB) treating diluted PWW.

Liquid samples from the influent PWW and the effluents of the photobioreactors were drawn weekly to determine the concentrations of TOC, IC, TN, NH_4^+ , NO_3^- , NO_2^- , TP, Zn and TSS. Likewise, the microalgae population structure in the photobioreactors was weekly assessed from biomass samples preserved with lugol acid at 5% and formaldehyde at 10%, and stored at 4 °C prior to analysis. Cultivation broth samples from the photobioreactors were also collected under steady state conditions and immediately stored at -20 °C to evaluate the richness and composition of the bacterial communities. Dissolved oxygen concentration (DO) and pH in the cultivation broth of the photobioreactors were daily measured, while the influents and effluents flow rates were daily recorded to monitor water evaporation losses. Finally, the C, N and P content of the biomass was monitored in both PBRs under steady state conditions.

The removal efficiencies of C, N, P and Zn were calculated according to Eq. (1):

$$RE(\%) = \frac{(C_{inf} \times Q_{inf}) - (C_{eff} \times Q_{eff})}{C_{inf} \times Q_{inf}} \times 100$$
(1)

where C_{inf} and C_{eff} represent the concentrations of TOC, IC, TN, TP and Zn in the influent PWW and PBR effluents, respectively, while Q_{inf} and Q_{eff} represent the influents and effluents flow rates, respectively. The process was considered under steady state when the TSS concentrations in the photobioreactors remained constant for at least four consecutive samplings. The results here provided correspond to the average ± standard deviation from duplicate measurements drawn weekly along one month of steady state.

2.4 Analytical procedures

A 510 pH meter (EUTECH Instrument, The Netherlands) was used to measure the pH, while a CellOX® 325 oximeter was used to measure the dissolved (WTW, oxygen and temperature Germany). PAR was measured with a LI-250A light meter (LI-COR Biosciences, Germany), while the intensity of infrared radiation was determined with a PASPort light meter (PASCO airlink®, California. USA). TOC, IC and TN concentrations were determined using a TOC-V CSH analyzer equipped with a TNM-1 module (Shimadzu, Japan). NO2⁻ and NO_3^- concentrations were analyzed by HPLC-IC with a Waters 515 HPLC pump coupled with a Waters 432 ionic conductivity detector and equipped with an IC-Pak Anion HC (150 mm \times 4.6 mm) Waters column (García et al., $N-NH_4^+$ 2017a). TP. and TSS concentrations were determined according to Standard Methods (APHA, 2005). The analysis of the C, N and P content in pre-dried and grinded biomass was carried out using a LECO CHNS-932 elemental analyzer. Zinc was determined using a 725-ICP Optical Emission Spectrophotometer (Agilent, USA) at 213.62 nm.

Microalgae were identified and quantified by microscopic examination

(OLYMPUS IX70, USA) according with phytoplankton manual (Sournia, 1978). Molecular analysis of the bacterial populations in PBR-AB and PBR-PPB was conducted according to García et al., (2017b). The genomic desoxyribonucleicacid (DNA) was extracted from inocula and photobioreactors effluents, respectively, by FastDNASpin Kit for Soil (MP Biomedicals, LLC, USA) according to the manufacturer's protocol. An aliquot of 300 ng DNA of each sample was provided to Fundació per al Foment de la Investigació Sanitária i Biomédica de la Comunitat Valenciana (FISABIO, Valencia, Spain) for 16S Amplicon sequencing by Illumina Miseq Platform 926F (50 using AAACTYAAAKGAATTGACGG-30)

and 1392wR (50-ACGGGCGGTGWGTRC-30) primer set (Engelbrektson et al., 2010; Hülsen et al., 2016b). Raw paired reads were first trimmed by Trimmomatic to remove short (less than 190bp) and low quality reads (lower than Phred-33 of 20). Data were analyzed according to Hülsen et al., (2016b) and trimmed paired reads were then assembled by Pandaseq with default parameters. The adapter sequences were removed by FASTQ Clipper of FASTX-The joined high Toolkit. quality sequences were analysed by QIIME using open-reference v1.8.0 OTU strategy picking by uclustat 1% phylogenetic distance and assigned taxonomy by uclust against green genes database. OTUs with only one read were filtered from the OTUs table by command filter_otus_from_otu_table.py in QIIME. An in house script was used to find a centroid normalized OTUs table with 40,000 reads per sample. Finally, the Shannon-Wiener diversity index (H)was determined using the peak heights in the densitometric curves. This index, which represents both the sample richness and evenness and ranges from 1.5 to 3.5 (low and high species evenness and richness. respectively), was according calculated to Eq. (2)(MacDonald, 2003):

$$H = -\sum [P_i ln(P_i)] \tag{2}$$

where P_i is the importance probability of the bands in a lane ($P_i = n_i/n$), where n_i is the height of an individual peak and n is the sum of all peak heights in the densitometric curves). **Table 1.** Operational conditions and physical/chemical characterization of the diluted PWW and cultivation broth in PBR-AB and PBR-PPB during steady state along the three operational stages.

Parameters		PWW	Sta	ge I	Stage II		Stage III		
			PBR-AB	PBR-PPB	PBR-AB	PBR-PPB	PBR-AB	PBR-PPB	
Operational days		-	84		63		77		
HRT (days)		-	pprox 10.6		pprox 7.6		≈ 4.1		
pH (units)		-	8.7±0.1	8.6±0.1	8.7±0.1	8.7±0.1	8.5±0.1	8.5±0.3	
PAR (µmol/m ² .s)		-	1388±39	-	1379±33	-	1407±12	-	
IR (W/m ²)		-	-	50±6	-	46±1	-	48 ± 8	
Temperature (°C)	09h00	-	32.8±0.9	$30.4{\pm}1.4$	31.3±1.0	$28.4{\pm}1.5$	32.1±0.9	30.5±1.4	
	16h00	-	31.2±1.5	30.5±1.5	30.3±1.5	29.0±1.5	30.7±1.6	30.3±1.5	
Dissolved oxygen (mg/L)		-	0.04 ± 0.02	0.05 ± 0.02	0.03 ± 0.02	0.02 ± 0.01	0.03±0.02	0.03±0.02	
Evaporation rates (%)		-	72±7	56±8	42±5	36±5	21±2	18±3	
*TOC (mg/L)		574±16	309±18	181±54	199±9	136±3	246±31	156±31	
*IC (mg/L)		58±4	169±17	142±3	144 ± 10	137±7	117±13	122±15	
*TN (mg/L)		166±9	68±5	65±4	85±3	84±6	118±9	110±10	
*Ammonium (mg/L)		179±5	41±3	60±7	80±5	83±6	118 ± 14	118±6	
*Nitrate (mg/L)		< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	
*Nitrite (mg/L)		< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5	
*TP (mg/L)		5.65 ± 0.41	1.62 ± 0.61	1.35 ± 0.51	1.62 ± 0.43	1.75 ± 0.27	1.22±0.27	1.22±0.57	
*Zinc (mg/L)		0.78 ± 0.07	0.07 ± 0.03	0.15 ± 0.02	0.08 ± 0.02	0.12 ± 0.02	0.08±0.02	0.06 ± 0.01	
*TSS (mg/L)		237±63	2640±161	873±114	1005 ± 54	853±51	638±35	553±118	
- Not applicable									
* Average value under steady stage conditions									

3. Results and Discussion

3.1 Batch PWW biodegradation tests

Overall, slightly higher TOC and TN removals were recorded in algalbacterial tests during batch PWW treatment compared to the tests conducted with PPB. The final TOC-REs in AB tests carried out with 5, 10 and 15% diluted PWW (namely AB-5, AB-10 and AB-15, respectively) accounted for 62, 46, 64%, respectively, compared to 52, 45 and 50% in PPB tests at comparable dilutions (namely PPB-5, PPB-10 and PPB-15, respectively). AB and PPB tests experienced a rapid decrease in organic matter concentration during the first 165 hours, with no variation significant in TOC concentration until the end of the experiment (Fig. 2a). The higher organic carbon removal in AB tests can be attributed to differences in microbial population structure and the occurrence of aerobic conditions. Hence, the aerobic bacteria present in the AB consortium were likely capable of utilizing a wider spectrum of organic compounds as electron donors than PPB (Golomysova et al., 2010). The results herein obtained in AB tests were in agreement with the study conducted by de Godos et al., (2010), who recorded TOC-REs of 55, 42 42, and 46% during the

biodegradation of 8 fold diluted PWW with E. viridis. S. obliquus, С. sorokiniana and Chlorella spp, respectively, in symbiosis with activated sludge. However, Hülsen et al., (2018) reported lower total and soluble COD-REs (< 20% and < 40%, respectively) during the batch treatment of PWW by PPB.

The final removals of TN accounted for 47, 48 and 66% in AB-5, AB-10 and AB-15, respectively, and 43, 43 and 55% in PPB-5, **PPB-10** and PPB-15. respectively (Fig. 2b). An active removal of nitrogen was detected during the first 165 hours of assay in both test series, which suggested the assimilatory nature of the N removal mechanism (correlated to TOC removal). The results herein obtained in AB tests were similar to those reported by de Godos et al., (2010), who recorded TN-REs ranging from 25 to 46% during the batch biodegradation of 8 fold diluted PWW. However, Hülsen et al., (2018) reported TN and NH_4^+ REs < 10% and < 40%, respectively, during the batch treatment of PWW by PPB.

On the other hand, the final biomass concentrations in AB-5, AB-10 and AB-15 accounted for 750, 1520, 2000 mg TSS/L, respectively, and 820, 1290 and

1460 mg TSS/L in PPB-5, PPB-10 and PPB-15, respectively (Fig. 2c). In this context, the higher TOC-REs and TN-REs recorded in AB tests supported the higher biomass concentrations here observed compared to PPB tests (García et al., 2017a).



Figure 2. Time course of the concentration of TOC (**a**), TN (**b**) and TSS (**c**) in the algal-bacterial (open symbols) and purple photosynthetic bacteria (solid symbols) systems during the biodegradation of piggery wastewater diluted at 5% (circle), 10% (triangle) and 15% (square).

3.2 *PWW biodegradation in* continuous photobioreactors

3.2.1 Carbon, nitrogen and

phosphorous removal

A gradual deterioration in the REs of carbon and nitrogen was recorded when decreasing the HRT in both PBRs. Indeed, TOC-REs in PBR-AB averaged 84±4, 79±3 and 66±3% in SI, SII and SIII, respectively, which resulted in steady state TOC concentrations in the effluent of 309±18, 199±9 and 246±31 mg/L, respectively (Fig.3a, Table 1). The results herein obtained confirmed the consistent removals of organic matter from PWW by algal-bacterial consortia and were in agreement with García et al., (2017b), who reported **TOC-REs** ranging from 85 to 94% in 3 L HRAPs during the treatment of 20 and 10 folds diluted PWW at a HRT of 27 days. Likewise, Hernández et al., (2013) reported COD-REs of 62±2 % in an outdoors 5 L HRAP treating PWW at 10 days of HRT. On the other hand, the TOC-REs in PBR-PPB accounted for 87±4, 84±3 and 77±5% in SI, SII and SIII, respectively, which entailed average TOC concentrations in the effluent lower than those detected in PBR-AB: 181±54, 136±3 and 156±31 mg/L, respectively (Fig.3a, Table 1). The TOC-REs here achieved were higher

than the average organic matter removals recorded by González et al., (2017) (REs ~65%) during the treatment of an anaerobic effluent in a 32 L membrane photobioreactor operated with native PPB. The high TOC-REs herein recorded in PBR-PPB could be attributed to the higher metabolic versatility of PPB, which degraded organic matter both aerobically (mediated by O_2 diffusion from the open atmosphere) and anaerobically (Hunter et al., 2009). In this context, Golomysova et al., (2010) highlighted the key role of the acetate assimilatory pathway of PPB during WWT. At this point, it must be also stressed that volatile fatty acids typically represent the main fraction of the soluble COD in PWW (González-Fernández and García-Encina, 2009). On the other hand, the steady state IC-REs in PBR-AB accounted for 5 ± 25 , -74 ± 28 and -52±24% during SI, SII and SIII, respectively, which resulted in average IC concentrations in the effluents of 167±17, 144±10 and 177±10 mg/L, respectively (Fig.3b, Table 1). Likewise, the IC-REs in PBR-PPB averaged -15±45, -87±72 and -66±18% in SI, SII and SIII, respectively, resulting in steady state IC concentrations in the effluents of 142±3, 137±7 and 122±15 mg/L, respectively (Fig.3b, Table 1). These negative IC-REs recorded in both PBRs

evidenced the accumulation of inorganic carbon mediated by the active microbial TOC oxidation. This finding was in agreement with previous observations in HRAPs treating PWW (García et al., 2017b). Overall, the higher IC-REs during the treatment of PWW in PBR-AB were likely mediated by the higher biomass concentrations and the oxygenic nature of the photosynthesis prevailing in PBR-AB (Table 1). A carbon mass balance showed that bioassimilation was the main mechanism responsible for carbon removal in PBR-AB and PBR-PPB during SII and SIII, with C recoveries in the form of biomass ranging between 58 and 72% of the total carbon removed. Carbon removal by stripping (prior mineralization of the organic carbon to CO₂) was the main mechanism accounting for carbon removal in PBR-PPB during SI, with a contribution of 62% of the total carbon removed.

The TN-REs in PBR-AB under steady state averaged 87±2, 69±3 and 47±1% in SI, SII and SIII, respectively, which corresponded to average TN concentrations in the effluent of 68±5, 85±3, and 118±9 mg/L, respectively (Fig. 3c, Table 1). Similar results were found in PBR-PPB, where TN-REs accounted respectively for 83±2, 65±6 and $48\pm3\%$, respectively, resulting in average TN concentrations in the effluent of 65±4, 84±6, and 110±10 mg/L in SI, SII and SIII, respectively (Fig. 3c, Table 1). Steady state NH₄⁺-REs of 93±1, 72±2 and 49±3% in PBR-AB and of 86±1, 68±5 and 48±4% in PBR-PPB were recorded during SI, SII, and SIII, respectively (Fig. 3d, Table 1). The TN-REs here achieved in PBR-AB were similar to those reported by García et al., (2018), who recorded TN removals of 82-85% during the treatment of 20 fold diluted PWW in indoor algalbacterial open photobioreactors at a HRT of \approx 27 days. However, these TN-REs were higher than the removals of $37\pm8\%$ obtained by de Godos et al., (2010) during PWW treatment in a 3.5L indoor enclosed photobioreactor operated at a HRT of 4.4 days. The nitrogen mass balance conducted under steady state revealed that stripping was the main N removal mechanism in both PBRs during the three stages, with assimilation into biomass in PBR-AB accounting only for 15, 21 and 24% of the TN removed, in SI, SII and SIII, respectively. Similarly, assimilation in PBR-PPB nitrogen accounted for only 9, 19 and 29% of the TN removed in SI, SII and SIII, respectively.

Finally, TP-REs of 91±3, 84±4 and 83±3% were recorded in PBR-AB in SI, SII and SIII, respectively, which resulted in average TP concentrations in the effluent of 1.6±0.6, 1.6±0.4, and 1.2±0.3 mg/L, respectively (Fig. 3e, Table 1). On the other hand, the TP-REs in PBR-PPB accounted for 89 ± 3 , 81 ± 1 and $82\pm9\%$ in SI, SII and SIII, respectively, which entailed effluent TP concentrations of 1.3±0.5, 1.7±0.3, and 1.2±0.5 mg/L, respectively (Fig. 3e, Table 1). Interestingly, high **TP-REs** were recorded in both PBRs regardless of the HRT and biomass concentration. The TP-REs obtained in PBR-AB were similar to those reported by García et al., (2018) during the treatment of 15% diluted PWW in indoor algal-bacterial

open photobioreactors (REs ~ from 90-92%. These values were also higher than the TP-REs of 58% reported by Myung et al., (2004) during the treatment of PWW by PPB. Phosphorus assimilation into biomass was likely the main P removal mechanism based on the moderate pH values (8.5-8-7) prevailing in both PBRs (pH = 8.5-8.7), which were not likely to support a significant phosphate precipitation (Table 1) (García et al., 2017a). Indeed, a phosphorus mass balance to PBR-AB revealed that 97, 89 and 51% of the total phosphorus removed was recovered as biomass during SI, SII and SIII, respectively. Similar P recoveries (60-81%) were estimated in PBR-PPB.



Figure 3. Steady state removal efficiencies of TOC (**a**), IC (**b**), TN (**c**), NH_4^+ (**d**) and TP (**e**) in PBR-AB (\boxdot) and PBR-PPB (\boxdot) in the three operational stages evaluated. Upper bold numbers indicate the steady state removal efficiencies, while vertical bars represent the standard deviation from replicate measurements during steady state operation.

3.2.2 Zinc removal

Zn-REs of 98 ± 1 , 94 ± 2 and $91\pm2\%$ were attained in PBR-AB in SI, SII and SIII, respectively, which mediated very low concentrations of Zn in the effluent under steady state conditions (0.07 \pm 0.03, 0.08 \pm 0.02 and 0.08 \pm 0.02 mg/L, respectively) (Table 1). On the other hand, the Zn -REs in PBR-PPB accounted for 93 ± 1 , 90 ± 2 and $92\pm2\%$ in SI, SII and SIII, respectively, which resulted in Zn effluent concentrations of 0.15±0.02, 0.12±0.02 and 0.06±0.01 mg/L, respectively (Fig. 3e, Table 1). The moderate pH prevailing in both PBRs (8.5-8-7) suggest that biosorption likely the main mechanism was governing Zn removal, although Zn-REs were not correlated with biomass concentrations (Javanbakht et al., 2014; Kaplan et al., 1987). The Zn-REs herein achieved were higher than those reported by García et al. (2017b) during PWW treatment in 3 L indoors HRAPs operated at a HRT of ≈ 27 days (71-83%).

3.2.3 Concentration, productivity and composition of biomass

Biomass concentration in PBR-AB increased during SI from ≈ 237 mg TSS/L up to steady state concentrations of 2640±161 mg TSS/L by days 63-84. A rapid decrease of biomass concentration to 1005±54 mg TSS/L in SII and to 683±35 mg TSS/L in SIII occurred as a result of the stepwise decrease in HRT in PBR-AB (Fig. 4a, Table 1). On the other hand, the biomass concentration in PBR-PPB experienced a gradual increase during the first 28 days of operation and stabilized at 873±114 mg TSS/L. Despite the decrease in HRT from 10.6 to 7.6 days did not result in a significant variation in TSS concentration in PBR-PPB (853±51 mg TSS/L in SII), a gradual decrease to 553±118 mg TSS/L occurred during SIII mediated by the decrease in HRT to 4.1 days (Fig.4a). The higher biomass concentrations recorded in PBR-AB compared to PBR-PPB, which were more evident during SI, were likely due active photosynthetic CO₂ to the assimilation by microalgae. These differences in biomass concentrations between both PBRs could be also explained by the slightly higher evaporation rates recorded in PBR-AB induced by its slightly higher temperatures. Hence, water evaporation (estimated as the ratio between the flow rate of water evaporation and the influent flow rate) in SI, SII and SIII accounted for 72, 42 and 21% in PBR-AB, and 56, 36 and 18% in PBR-PPB, respectively.

The lowest areal biomass productivities were recorded in SI in both PBRs, accounting for 10.4 and 5.4 g/m²·d in PBR-AB and PBR-PPB, respectively. The decrease in HRT resulted in increased biomass productivities up to 11.5 and 10.8 g/m²·d in stage II in PBR-AB and PBR-PPB, respectively. Finally, process operation at a HRT of 4.1 days was characterized by the highest biomass productivities: 18.4 g/m²·d in PBR-AB and 16.6 g/m²·d in PBR-PPB. The C, N and P content of the algalbacterial biomass averaged 48 ± 4 , 7.5 \pm 1.4 and 0.62 \pm 0.15%, and 52 \pm 1, 8.4 \pm 0.5 and $0.69 \pm 0.05\%$ in the PPB biomass, with no clear correlation with the HRT (Fig. 4b). The algal-bacterial biomass composition was similar to the values reported by Cabanelas et al., (2013), who determined a C, N and P content in the harvested biomass of \approx 44, 7.5 and 0.5%, respectively, in а photobioreactor with inoculated С. vulgaris and supplemented with CO₂ during the of settled domestic treatment wastewater.



Figure 4. a) Time course of TSS concentration in PWW (\Box), PBR-AB (\circ) and PBR-PPB (Δ) during the entire experiment. **b**) C (\boxtimes), N (\boxdot) and P (\blacksquare) content in the biomass present in the photobioreactors under steady state.

3.2.4 Microalgae population structure

C. vulgaris represented the dominant species in PBR-AB from day 1 to day 84 in SI, with a maximum concentration of $1.6 \cdot 10^{10}$ cells/L bv dav 35 (corresponding to 90 % of the total cell number) (Fig. 5a). Interestingly, a severe decrease in the total number of microalgae cells was observed from day 35 to day 56, which remained stable at $1.8\pm0.2\cdot10^{9}$ cells/L by the end of SI. Chlorella kessieri was always detected from day 42 to day 84, while Scenedesmus acutus and Tetradesmus obliquus were detected for 10 and 7 weeks during SI, respectively. On the other hand, no microalgae was detected in PBR-PPB during SI (Fig. 5b). During SII, C. vulgaris and C. kessieri were present in PBR-AB throughout the experimental period from day 91 to С. vulgaris day147. achieved а maximum concentration of $0.24 \cdot 10^9$ cells/L by day 105 (corresponding to 55 % of the total cell number). However, the cell concentration maximum was recorded by day 119. where $0.51 \cdot 10^9$ cells/L and eight microalgae species were detected. Sc. acutus and Tet. obliquus (acutudesmus) were not detected from days 98 and 126 onwards, respectively. On the other hand, C. vulgaris and Chlorococcum sp. were

detected 6 and 2 times during SII in PBR-PPB, respectively. However, the maximum microalgae cell concentration in PBR-PPB was only 0.01.10⁹ cells/L, which was recorded by day 147 as a result of the occurrence of four microalgae species. Finally, C. vulgaris and C. kessieri were always present in SIII from day 154 to day 224 in PBR-AB, while *Chlorella minutissima* was identified from day 175 onwards. maximum However. the cell concentration of microalgae in PBR-AB during SIII was only 0.27.10⁹ cells/L (day 217). Finally, six microalgae species were detected in PBR-PPB during SIII. Aphanothece saxicola was detected by day 161 and day 217, while C. vulgaris, C. kessieri and С. minutissima were detected during SIII up to days 154, 161 and 182, respectively. The N₂ fixing cyanobacteria, *Cyanobium* spp. and *Pseudanabaena rosea* were dominant by the end of SIII, when the maximum microalgae concentration $(0.19 \cdot 10^9 \text{ cells/L} \text{ by day } 217) \text{ was}$ achieved (Richmond, 2004).

C. vulgaris, the microalga species inoculated in PBR-AB, was detected in all stages from day 7 to day 224, along with other *Chlorella* species. The high tolerance of microalgae from the genus *Chlorella,* which ranked 5th in the

ranking of pollution tolerant microalgae species established by Palmer (1969), supported the dominance of this microalgal species in PBR-AB regardless of the HRT. In addition, the monitoring of the microalgae population structure clearly showed that the decrease in HRT induced a gradual wash-out of microalgae, which mediated a significant decrease in the number of cells from 1.8 cells/L \cdot 10⁹ (day 84) to 0.17 cells/L \cdot 10⁹ (day 244) under steady state in PBR-AB. This study also

revealed that inoculation of the PBR-AB with a specific photosynthetic microorganism does not guarantee its long-term dominance during PWW (Serejo et al., treatment 2015). Interestingly, the stepwise decrease in HRT in PBR-PPB induced the occurrence of microalgae from day 98 onwards. Finally, it should be stressed that no direct correlation between the structure of microalgae population in the PBRs and the structure of bacterial population was found (Fig. 4, 5).



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Figure 5 Time course of the microalgae population structure in PBR-AB (a) and PBR-PPB (b) during the three operational stages.

3.2.5 Bacteria population structure

The bacterial analysis of the microbial communities present in PBR-AB revealed the occurrence of the following phyla along the three operational stages: Actinobacteria. Chloroflexi, Epsilonbacteraeota, *Cyanobacteria*, Firmicutes, Patescibacteria and Proteobacteria among others phyla. *Proteobacteria* and *Cyanobacteria* represented the main phyla present in the inoculum of PBR-AB, with shares of 67.1 and 26.9, respectively. All phyla were detected in SI and SII in PBR-AB under steady state, although the phylum Actinobacteria was not present in SIII. The decrease in HRT induced a severe swift in the structure of the bacterial community, represented by Firmicutes (43.8%), *Epsilonbacteraeota* (10.7%), (9.3%)Chloroflexi Proteobacteria (10.9%) and Cyanobacteria (13.2%) at the end of the experiment (Table 2). On the other hand, the bacterial analysis in PBR-PPB revealed the occurrence of the phyla Acidobacteria, Chloroflexi,

Epsilonbacteraeota, Firmicutes. Patescibacteria, Proteobacteria and *Synergistetes* among others. Proteobacteria **Synergistetes** and accounted for 83.8 and 5.3% of the bacteria in the inoculum of PBR-PPB. Epsilonbacteraeota, *Firmicutes* and Proteobacteria were dominant along the three stages, while Patescibacteria, which was present in SI and SII, was not detected in SIII. The decrease in HRT in PBR-PPB mediated the dominance of Firmicutes and *Epsilonbacteraeota* (46.7 % and 19.8 % of abundance) and decreased the contribution of Proteobacteria to 19.8% in SIII (Table 2). Overall, the HRT seems to play a key role on the bacterial population structure in both PBR-AB and PBR-PPB during PWW treatment. Firmicutes is one of two dominant phyla in the large intestine of human and pig (Ban-Tokuda et al., 2017). Firmicutes can degrade volatile fatty acids, which typically account for 80% of the TOC in the soluble fraction of PWW (Ferrero et al., 2012).

Dhyllum		PBR-A	B (%)		PBR-PPB (%)				
Phyllum	Inoc.	SI	SII	SIII	Inoc.	SI	SII	SIII	
Acidobacteria						0.5	2.5	1.5	
Actinobacteria		1.3	3.4						
Chloroflexi		13.2	11.3	9.3			3.7	5.4	
Cyanobacteria	26.9	25.7	3.1	13.2					
Epsilonbacteraeota		18.8	8.2	10.7		46.6	19.9	19.8	
Firmicutes	1.0	4.7	38.2	43.8	3.6	5.3	29.3	46.7	
Patescibacteria		5.5	0.6	1.0	0.0	4.5	6.5		
Proteobacteria	67.1	19.1	21.6	10.9	83.8	30.5	24.6	12.1	
Synergistetes					5.3				
Other	5.0	11.8	13.5	11.1	7.3	12.6	13.5	14.4	
Total nº of Cells	34903	18964	13683	28884	44503	29644	19355	14765	

Table 2. Taxonomic report of the bacteria present in PBR-AB and PBR-PPB.

Proteobacteria was the main phylum with 67.1 and 83.8 % of the species present in the inocula of PBR-AB and PBR-PPB, respectively (Fig. 6). Interestingly, Alphaproteobacteria was the only class found in both photobioreactors. Blastomonas, which are aerobic and catalase/oxidasepositive, was the dominant genus in the inoculum of PBR-AB with 49.7% of the total number of bacteria (Castro et al., 2017). However, Blastomonas was not found in PBR-AB during SI, SII and SIII. Rhodoplanes was detected in PBR-AB during SII (18.7% of the total number of bacteria) and SIII (1.5%), while *Rhodobacter* was only present in SIII (6.5%), despite both genera belong to PPB (Hunter et al., 2009)(Hiraishi and Ueda, 1994)(Fig. 6a). On the other hand, Rhodopseudomonas was the dominant genus in the inoculum of PBR-PPB with

81.7% of the total number of bacteria, but disappeared from SII onwards (Fig. 6b). Rhodopseudomonas are purple nonsulfur phototropic bacteria (Hiraishi and Ueda, 1994) that can metabolize organic substrates (Cheah et al., 2016). The phototrophic Rhodoplanes accounted for 9.2, 11.4 and 7.5% of the total number of bacteria in SI, SII and SIII, respectively (Hunter et al., 2009) (Hiraishi and Ueda, 1994). In this context, a wash-out of Rhodopseudomonas followed by the dominance of Rhodoplanes was also observed by Chitapornpan et al., (2013) **PPB-based** in а membrane photobioreactor during the treatment of food processing wastewater.

Finally, the Shannon-Wiener diversity indexes calculated in both photobioreactors indicated an increase in diversity compared to the inocula, which

remained similar during process operation. Thus, the values of *H* in the inoculum, SI, SII and SIII were, respectively, 0.74, 1.89, 1.64 and 1.69 in PBR-AB, and 0.53, 1.32, 1.68 and 1.48 in PBR-PPB. This low-bacterial diversity in both photobioreactors was likely due to the high toxicity of the PWW treated and the low HRT used in this study.



Figure 6. Relative abundance (%) of genera belonging to the phylum *Proteobacteria* in the inocula and cultivation broth of PBR-AB (**a**) and PBR-PPB (**b**) along the three operational stages. The abundance was calculated based on the total number of bacteria.

4. Conclusions

This work constitutes, to the best of our knowledge, the first comparative evaluation of the potential of microalgae and PPB during continuous PWW treatment in open photobioreactors. This research revealed a similar treatment performance of both photosynthetic microorganisms in terms of carbon,

nutrients and zinc removal. The PBR-PPB exhibited a slightly better capacity to remove organic matter, which was not observed during batch PWW treatment. Interestingly, a superior carbon and nutrient recovery was recorded in PBR-AB. The stepwise decrease in HRT, rather than the type of illumination used, caused significant changes in the structure of microalgae and bacterial population.

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SUPPLEMENTARY MATERIAL

A systematic comparison of the potential of a microalgal-bacterial consortium and purple phototrophic bacteria for the continuous treatment of piggery wastewater

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Figure S1. Average removal efficiencies of TOC, IC and TN under steady state for AB-5 ((1), AB-10), AB-15 ((1), PPB-5 ((1), PPB-10), PPB-15 ((1), Bold numbers indicate the steady state removal efficiencies.



the batch treatment of PWW diluted at 5% (circle), 10% (triangle) and 15% (square).



Figure S3. Time course of the headspace concentrations of N₂, CO₂ and O₂ in AB (**a**, **b** and **c**) and PPB (**d**, **e** and **f**) tests during the batch treatment of PWW diluted at 5% (AB-5 and PB-5), 10% (AB-10 and PB-10) and 15% (AB-15 and PB-15).

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Figure S4. Time course of the evaporation rates in PBR-AB (\circ) and PBR-PPB (Δ).



Figure S5. Time course of the temperature at 09h00 (**a**) and 16h00 (**b**) in PBR-AB (\circ) and PBR-PBB (Δ).



Figure S6. Time course of the dissolved oxygen concentration (**a**) and pH (**b**) in PWW (\Box), PBR-AB (\circ) and PBR-PBB (Δ).
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Figure S7. Time course of the average PAR and IR in PBR-AB (\circ) and PBR-PBB (Δ), respectively.



Figure S8. Time course of the removal efficiencies of TOC (a), IC (b), TN (c), N-NH₄⁺ (d) and TP (e) in PBR-AB (\circ) and PBR-PBB (Δ). Vertical dashed lines separate the different operational stages evaluated.

PBR	HRT (d)	Q feed (L/d)	Q eff (L/d)	Evaporation rate (%)	V PBR (mL)	Height PBR (cm)	Area PBR (m ²)
AB	10.6	0.282	0.079	72	2000	15	0 0 2 0
PPB	10.0	0.282	0.124	56	3000	15	0.020

Table S1. Operational parameters and photobioreactors dimensions in Stage I.

PBR: Photobioreactor AB: Algal-bacterial PPB: Purple phototrophic bacteria

Table S2. Mass balance calculations of PWW treatment under steady state conditions inPBR-AB and PBR-PPB in Stage I.

Parameters	PWW	PBR-AB	PBR-PPB
TOC (mg/L)	592	309	181
TOC (mg/d) in	392	166.9	166.9
TOC (mg/d) out		24.4	22.5
% TOC RE		85	87
IC (mg/L)	57	169	142
IC (mg/d) in		16.1	16.1
IC (mg/d) out		13.3	17.6
% IC RE		17	-10
TN (mg/L)	160	68	65
TN (mg/d) in		45.1	45.1
TN (mg/d) out		5.4	8.1
% TN RE		88	82
Nitrite (mg/L)		< 0.5	< 0.5
Nitrate (mg/L)		< 0.5	< 0.5
TP (mg/L)	5.20	1.61	1.35
TP (mg/d) in		1.47	1.47
TP (mg/d) out		0.13	0.17
% TP RE		91	89
Zinc (mg/L)	0.85	0.07	0.015
Zn (mg/d) in		0.24	0.24
Zn (mg/d) out		0.006	0.002
% Zn RE		98	99
TSS (mg/L)	183	2640	873
Productivity (g/m2/d)		10.42	5.42
С %		44.8	50.7
C (mg/d) biomass		93	55
% Carbon recovered		64	38
N %		6.97	8.71
N (mg/d) biomass		14.53	9.43
% Nitrogen recovered		37	25
P%		0.62	0.72
P (mg/d) biomass		1.29	0.78
% Phosphorus recovered		97	60

PBR	HRT (d)	Q feed (L/d)	Q eff (L/d)	Evaporation rate (%)	V PBR (mL)	Height PBR (cm)	Area PBR (m ²)	
AB	7.6	0.206	0.230	42	2000	15	0 0 2 0	
PPB	/.0	0.590	0.396 0.253 36 300	3000	15	0.020		

Table S3. Operational parameters and photobioreactors dimensions in Stage II.

PBR: Photobioreactor AB: Algal-bacterial PPB: Purple phototrophic bacteria

Table S4. Mass balance calculations of PWW treatment under steady state conditions inPBR-AB and PBR-PPB in Stage II.

Parameter	PWW	PBR-AB	PBR-PPB
TOC (mg/L)	566	199	136
TOC (mg/d) in		224.1	224.1
TOC (mg/d) out		45.7	34.5
% TOC RE		80	85
IC (mg/L)	54	144	137
IC (mg/d) in		21.4	21.4
IC (mg/d) out		33.1	34.7
% IC RE		-55	-62
TN (mg/L)	162	85	84
TN (mg/d) in		64.2	64.2
TN (mg/d) out		19.5	21.3
% TN RE		70	67
Nitrite (mg/L)		< 0.5	< 0.5
Nitrate (mg/L)		< 0.5	< 0.5
TP (mg/L)	6.00	1.62	1.75
TP (mg/d) in		2.38	2.38
TP (mg/d) out		0.37	0.44
% TP RE		84	81
Zinc (mg/L)	0.78	0.2	0.12
Zn (mg/d) in		0.31	0.31
Zn (mg/d) out		0.05	0.03
% Zn RE		85	90
TSS (mg/L)	263	1005	853
Productivity (g/m2/d)		12	11
С %		52	53
C (mg/d) biomass		121	115
% Carbon recovered		72	65
N %		9.08	8.72
N (mg/d) biomass		20.96	18.85
% Nitrogen recovered		47	44
P%		0.77	0.72
P (mg/d) biomass		1.78	1.56
% Phosphorus recovered		89	81

PBR	HRT (d)	Q feed (L/d)	Q eff (L/d)	Evaporation (%)	V PBR (mL)	Height PBR (cm)	Area PBR (m ²)
AB	4 1	0 721	0.577	21	3000	15	0 0 2 0
PPB	4.1	0.731	0.599	18	3000	15	0.020

Table S5. Operational parameters and photobioreactors dimensions in Stage III.

PBR: Photobioreactor AB: Algal-bacterial PPB: Purple phototrophic bacteria

Table S6. Mass balance calculations of PWW treatment under steady state conditions inPBR-AB and PBR-PPB in Stage III.

Parameter	PWW	PBR-AB	PBR-PPB
TOC (mg/L)	564	246	156
TOC (mg/d) in		412.3	412.3
TOC (mg/d) out		142.1	93.5
% TOC RE		66	77
IC (mg/L)	62	117	122
IC (mg/d) in		45.3	45.3
IC (mg/d) out		67.6	73.1
% IC RE		-49	-61
TN (mg/L)	176	118	110
TN (mg/d) in		128.7	128.7
TN (mg/d) out		68.1	65.9
% TN RE		47	49
Nitrite (mg/L)		< 0.5	< 0.5
Nitrate (mg/L)		< 0.5	< 0.5
TP (mg/L)	5.74	1.22	1.22
TP (mg/d) in		4.20	4.20
TP (mg/d) out		0.70	0.73
% TP RE		83	83
Zinc (mg/L)	0.69	0.08	0.06
Zn (mg/d) in		0.50	0.50
Zn (mg/d) out		0.05	0.04
% Zn RE		91	93
TSS (mg/L)	265	638	553
Productivity (g/m2/d)		18	17
C %		47	51
C (mg/d) biomass		173	169
% Carbon recovered		70	58
N %		6.55	8.74
N (mg/d) biomass		24.13	28.97
% Nitrogen recovered		40	46
P%		0.48	0.64
P (mg/d) biomass		1.77	2.12
% Phosphorus recovered		51	61

Table S7. Relative abundance (%) of the taxonomic composition for algal-bacterial (PBR-AB) and purple phototrophic bacteria (PBR-PPB) photobioreactors.

Dhadaaaa	Clear	Orden	E	Gamma]	PBR-A	B (%)			PBR-P	PB (%)	
Phylum	Class	Order	Family	Genus	Inoc.	SI	SII	SIII	Inoc.	SI	SII	SIII
Acidobacteria	Acidobacteriia	Solibacterales	Solibacteraceae (Subgroup 3)	Paludibaculum						0.5	2.5	1.5
Actinobacteria	Coriobacteriia	OPB41	uncultured Actinobacteridae bacterium	uncultured Actinobacteridae bacterium		1.3	3.4					
		Anaerolineales	Anaerolineaceae	Anaerolinea			0.7	1.5			1.7	0.9
		Anderonneales	Anaerolineaceae	uncultured		11.6	10.6	1.6			2.1	4.5
Chloroflexi	Anaerolineae	SBR1031	A4b	uncultured gamma proteobacterium		1.6						
		Chloroflexales	Chloroflexaceae	Chloronema				4.3				
	Chloroflexia		Chloroflexaceae					1.8				
Cyanobacteria	Oxyphotobacteria	obacteria Chloroplast		Tetradesmus obliquus		2						
5	~ 1		_		26.9	23.7	3.1	13.2				
Epsilonbacteraeota	Campylobacteria	Campylobacterales	Arcobacteraceae	Arcobacter		18.8	8.2	10.7		46.6	19.9	19.8
Firmicutes	BRH-c20a	uncultured bacterium mle1-9	uncultured bacterium mle1-9	uncultured bacterium mle1-9				1.6				1.4
			Christensenellaceae	Christensenellaceae R-7 group	1.0	0.8	6.2	3	1.5	0.8	7.2	3.5
Firmicutes	Clostridia	Clostridiales	Family XI	Sedimentibacter				14.7			1.2	7.8
			Family XIII	Anaerovorax		0.9	13.5	3.6			2	3.1
				uncultured							1.2	6.6

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Dhadaaa	Class	Orden	Family	Commo	I	PBR-A	B (%)		PBR-PPB (%)			
Phylum	Class	Order	Family	Genus	Inoc.	SI	SII	SIII	Inoc.	SI	SII	SIII
		Clostridiale		Fastidiosipila		1	8.3	5.1	2.1	4.5	11.3	9.3
				Ruminococcaceae UCG-013							1.6	
			Ruminococcaceae	Ruminococcaceae UCG-014			1				1.7	
Firmicutes	Clostridia			[Eubacterium] coprostanoligenes group		1.2	1.1	0.5				
						0.7	1.2	1.1			1.1	2.2
		DTU014	uncultured Selenomonadales bacterium	uncultured Selenomonadales bacterium			6.9	14.2			2.1	12.8
Patescibacteria	Parcubacteria	Candidatus Kaiserbacteria	Candidatus Kaiserbacteria bacterium RIFOXYB1_FULL_ 46_14	Candidatus Kaiserbacteria bacterium RIFOXYB1_FULL _46_14				1			4.1	
	Saccharimonadia	Saccharimonadales	Saccharimonadaceae								1.4	
	WS6 (Dojkabacteria)	uncultured bacterium	uncultured bacterium	uncultured bacterium		5.5	0.6			4.5	1	

Dissister	Class	Orden	E a su llas	Gamma	PBR-AB	(%)			PBR-PPB (%)			
Phylum	Class	Order	Family	Genus	Inoc.	SI	SII	SIII	Inoc.	SI	SII	SIII
		Caulobacterales	Caulobacteraceae	Caulobacter	3.7							
			Beijerinckiaceae	Methylocystis			1.9			0.7	9	3.1
			Rhizobiaceae			1.3				10.2	1	
		Rhizobiales		Rhodoplanes			18.7	1.5		9.2	11.4	7.5
	steria		Xanthobacteraceae	Rhodopseudomonas					81.7	1.7		
	Alphaproteobacteria	Rhodobacterales Rhodospirillales	Rhodobacteraceae	Paracoccus			1					
Proteobacteria				Rhodobacter				6.3				
						7.2		1.5		4.6	1	
	Alp		Magnetospirillaceae	Magnetospirillum				1.6				1.6
			Sphingomonadaceae	Blastomonas	49.7							
		Sphingomonadales		Novosphingobium	6.3	10.6			1	4.2	2.3	
				Sphingomonas	7.4				1			
Synergistetes	Synergistia	Synergistales	Synergistaceae	Aminobacterium					5.3			
				Others	5	11.8	13.5	11.1	7.3	12.6	13.5	14.4
				Number of cells	34903	18964	13683	28884	44503	29644	19355	14765

_____ **(** 198 **)**_____



Conclusions and future work

Several photobioreactor configurations were evaluated during wastewater treatment under indoor and outdoor conditions, along with several photosynthetic microorganisms. In addition, state-of-the-art operational strategies such as CO₂ supplementation, biomass settling and recycling, and wastewater dilution, were tested to boost pollutant biodegradation and biomass productivity during wastewater treatment. Overall, the results herein obtained confirmed the potential of photosynthetic processes for domestic and livestock wastewater treatment as a cost-effective and environmental friendly technology platforms.

Firstly, a novel anoxic-aerobic algal-bacterial photobioreactor with biomass settling and recycling was evaluated using real domestic wastewater. This study was conducted to improve processes performance by overcoming the main limitation present in domestic wastewater as a result of its low C/N/P ratio (inorganic carbon). The results obtained in **Chapter 1** demonstrated that algal-bacterial symbiosis supported an efficient biodegradation of TOC and TN when carbon dioxide from biogas was supplemented. In this context, biogas supplementation avoided the need for pH control, and enhanced TOC and TN removals by promoting the activity of nitrifying bacteria up to complete ammonium oxidation. On the other hand, a stable and high biomass concentration was achieved by recirculation of the settled biomass, which also supported an effluent TSS concentration complying with EU regulations. Likewise, biomass settling and recycling also contibuted to the enrichment of dominant monoalgal cultures with good settling properties.

The long-term dominance of a specific microalga or cyanobacterium under indoor conditions treating piggery wastewater was evaluated in four open photobioreactors. Despite the optimal environmental conditions of light, temperature and pH (controlled via CO₂ addition) prevailing throughout the experiment, the results obtained in **Chapter 2** showed a difficulty to maintain a monoalgal culture in all photobioreactors. Interestingly, the highest biomass concentrations were recorded in the control photobioreactor at the end of the experiment as a result of the acclimation of native species. *Chlorella sp.* was dominant in R3 and R4, which highlighted the high tolerance of this microalga to organic pollution. Finally, an efficient PWW treatment occurred regardless of the microalgae species inoculated, with stripping identified as the main mechanism responsible for carbon and nitrogen removal.

A comparative study (Chapter 3) was carried to evaluate microalgae evolution and process performance under indoor and outdoor conditions during the treatment of PWW. Four open photobioreactors were operated to elucidate the representativeness of the results obtained indoor under artificial irradiation. A comparable TOC removal performance was recorded regardless of the photobioreactor location. Higher TN removal efficiencies were recorded at lowest dilutions (×20) both indoors and outdoors. Stripping was the main mechanism responsible for TOC and TN removal during the treatment of 10× PWW. Chlorella vulgaris became dominant in all photobioreactors regardless of the environmental conditions and PWW dilution, which confirmed the high pollutiontolerance of this microalga species. The highest microalgae concentrations were recorded during the treatment of 20× PWW regardless of the environmental conditions likely due to the lower toxicity of this effluent. On the other hand, the molecular analysis of the bacterial populations revealed a high dominance of the Proteobacteria phylum in all photobioreactors, and the key influence of temperature and irradiation on the final bacterial population structure. The results herein obtained demonstrated for the first time that neither pollutant removal nor the structure of microalgae and bacterial communities under indoor conditions can be directly extrapolated to outdoors photobioreactors.

Finally, the potential of a microalgal-bacterial consortium and purple phototrophic bacteria was comparatively evaluated during the continuous treatment of PWW under indoor conditions in two open photobioreactors. In addition, the influence of PWW dilution on the biodegradation performance of these photosynthetic consortia was assessed batchwise (Chapter 4). Overall, the results revealed a similar treatment performance of both photosynthetic microorganisms in terms of carbon, nutrients and zinc removal. PBR-PPB exhibited a slightly better capacity to remove organic matter (which was not observed during batch PWW treatment), while a superior carbon and nutrient recovery was recorded in PBR-AB. Assimilation was the main mechanism responsible for carbon removal (except in PBR-PPB at high HRT), while stripping governed N removal in both PBRs. Phosphorus assimilation and Zn biosorption were likely the main mechanisms responsible for the removal of these inorganic pollutants based on the moderate pH values prevailing in both PBRs. Higher biomass productivities were recorded in PBR-AB mediated by microalgae CO₂ recovery. The stepwise decrease in HRT, rather than the type of illumination used, caused significant changes in the structure of microalgae and bacterial population. The batch biodegradation tests conducted at multiple PWW dilutions (5, 10 and 15%) confirmed the slightly better performance of algal-bacterial systems regardless of PWW dilution.

Based on the outcomes and limitations found in this study, further research on photosynthetic biodegradation should focus on:

1. Use of pollution tolerant microorganisms in order to increase pollutant removal and enhance biomass production based on the findings observed in **Chapter 2**.

2. The scale-up of photobioreactors under outdoor conditions, which will provide more representative information about pollutant removal performance and biomass production based on the results obtained in **Chapter 1 and 3**.

3. Evaluation of alternative photobioreactors configuration in order to avoid high evaporation rates and their negative influence on the quality of the treated effluents based on the observations in **Chapter 2 and 3**.

4. Optimization of the cultivation conditions of PPB in order to identify the full potential of these microorganisms for PWW treatment based on the results recorded in **Chapter 4**.

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About the author

Biography

Dimas Alberto García Guzmán (Masaya, Nicaragua, 1973)

Dimas A. García studied Chemistry at Universidad Nacional Autónoma de Nicaragua UNAN-Managua (1994-1999). In 2010, Dimas was awarded with a scholarship to study a Master's Degree in Water Science at Centro para la Investigación en Recursos Acuáticos de Nicaragua (CIRA/UNAN-Managua). His Master's thesis was entitled "Zonificación Hídrica de la microcuenca del río La Trinidad, en la subcuenca del Río Viejo, Estelí, Nicaragua". In July 2014, Dimas was awarded with a scholarship from the **EURICA** project (**Erasmus Mundus Action 2, Strand 1, Lot 15, Grant Agreement number 2013-2587**) of the European Union to study a doctorate in Chemical Engineering and Environmental Technology at Valladolid University, Spain.

In September 2014, Dimas joined the Biological Gas Treatment and Microalgae Technology Research Group headed by Professor Raúl Muñoz Torre within the Environmental Technology Group of the Department of Chemical Engineering and Environmental Technology. His PhD thesis was focused on the development of innovative biotechnological processes for the treatment of domestic and agroindustrial wastewater using photobioreactors (September 2014 - September 2018).

Dimas A. García has been working at CIRA/UNAN-Managua since July 2000 as a lecturer on Limnology and Hydrogeochemestry. In addition, Dimas has an extensive experience in monitoring water resources such as rivers and lakes, and industrial effluents and wastewater treatment plants. Dimas has been an internal auditor in analytical quality management at CIRA/UNAN-Managua.

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