# Co-digestion strategies to enhance microalgae anaerobic digestion: A

# review

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#### Abstract

Microalgae biorefineries for the production of biofuels and high-value products have captured the attention of academia and industry. Implementing an anaerobic digestion step can enhance resource recovery from microalgae and microalgae residues. Anaerobic codigestion, the simultaneous digestion of two or more substrates, is an opportunity to overcome the low biodegradability and the risk of ammonia inhibition associated with microalgae and microalgae residues mono-digestion. Besides, microalgae can also be used as co-substrate in biogas plants, with the aim of increasing the organic loading rate while providing alkalinity, macro- and micronutrients. Sewage sludge is the most researched cosubstrate for microalgae since microalgae photobioreactors can be used for secondary, tertiary and anaerobic digestion supernatant treatment in wastewater treatment plants. However, microalgae and microalgae residues have been successfully co-digested with a wide variety of wastes, including crops, energy crops, paper waste, animal manure, vinasse, olive mill waste, and fat, oil and grease. Lipid-spent microalgae and glycerol co-digestion has also been largely researched due to the growing interest on microalgal-derived biodiesel. Most studies have assessed the impact of co-digestion on the methane yield and process kinetics through biochemical methane potential (BMP) tests. However, BMP test is not the most suitable method to assess the impact of co-digestion on other important factors such as supernatant nutrient content, digestate dewaterability, biosolids quality, and H<sub>2</sub>S concentration in the biogas. Overall, more lab-scale and pilot-scale continuous experiments are needed to get a holistic understanding of microalgal anaerobic co-digestion.

**Keywords:** biogas; anaerobic co-digestion; biorefinery; microalgal biomass; cyanobacteria; microalgae residues

#### 1. Introduction

The development of integrated microalgae-based facilities, so-called microalgae biorefineries, has attracted a substantial amount of attention from both academia and industry [1-3]. Microalgae biorefineries combine the production of biofuels (e.g. biodiesel, bioethanol, biogas) and high-value products (e.g. pigments, proteins, omega-3). Thus, microalgae biorefineries go one-step beyond the "third-generation biofuels" concept, which only aims at the production of liquid biofuels from microalgae. In biorefineries, microalgae cultivation costs can be reduced by using wastewater streams as nutrient source; achieving the dual goal of wastewater treatment and high-value chemicals production [4-6]. However, biosecurity regulations may restrict the application of microalgae products when using wastewater streams as source of nutrients [7-9].

Anaerobic digestion (AD), a biological process that transforms organic matter into renewable biogas energy, has been stated as the most suitable technology to maximize resource recovery from microalgae [10-13]. This technology is particularly suitable to treat the large amount of microalgae residues produced from the extraction of metabolites and reduce costs associated with microalgae residues treatment and disposal [2, 14-16]. Additional benefits of treating microalgae or microalgae residues via anaerobic digestion are the mobilization of nutrients (N and P) and the availability of CO<sub>2</sub>, which can be recycled for microalgae cultivation [17-19]. However, microalgae AD is generally limited by its low methane potential (degradation extent) and conversion rate (degradation speed) as well as the risk of ammonia nitrogen inhibition. Pre-treatment methods, with or without co-products recovery, have been applied to disrupt microalgae cell wall, make their intracellular content more available and improve microalgae anaerobic biodegradability (extent and rate). Microalgae pre-treatments without co-products recovery have been reported to increase microalgae methane yield up to 100% [20, 21]. However, in most applications the increase in methane yield do not compensate the pre-treatment implementation and operational costs [22]. Indeed, the co-production of high-value chemicals and biogas has been identified as a more conceivable scenario than the production of biogas alone [3, 12, 23, 24]. Microalgae residues after co-product extraction have shown an increased anaerobic biodegradability when compared to raw microalgae, since the extraction step acts as a pre-treatment. For example, Ramos-Suarez and Carreras [14] observed an increase on Scenedesmus sp. methane yield from 140 to 272 and 212 mLCH<sub>4</sub>/gVS after the extraction of proteins and lipids respectively, while Parimi et al. [25] reported a methane yield increase from 181 to 254 mLCH<sub>4</sub>/gVS for protein-spent Spirulina platensis. Even if upstream processing increases microalgae's anaerobic biodegradability, microalgae and microalgae residues are generally characterized by low methane yields (150-300 mLCH<sub>4</sub>/gVS) and degradation rates (0.05-0.25 day<sup>-1</sup>) when compared to common anaerobic digestion substrates, such as sewage sludge (200-350 mLCH4/gVS, 0.20-0.40 day<sup>-1</sup>), animal manure (200-400 mLCH4/gVS, 0.10-0.30 day<sup>-1</sup>) and food waste (400-550 mLCH<sub>4</sub>/gVS, 0.30-0.70 day<sup>-1</sup>) [14, 26-35].

Another key limitation for microalgae and microalgae residues anaerobic digestion is the risk of ammonia nitrogen inhibition, typically associated with microalgae's low carbon-tonitrogen (C/N) ratio. Ammonia nitrogen is a potential inhibitor of the AD process that is released during the degradation of nitrogenous organic matter (e.g. proteins, amino acids, urea and nucleic acids) [36-38]. Indeed, microalgae biodegradability can be improved by different strategies, such as cultivating a different microalgae strain, tuning cultivation conditions, and using pre-treatments [22, 39, 40]. However, a high protein content and a low C/N ratio is common across all microalgae species. The risk of ammonia inhibition limits the maximum organic loading rate (OLR) at which microalgae digesters can be operated. An OLR around 2 gVS/(L<sub>r</sub>·day) has been observed as OLR threshold prior evidence of process inhibition [16, 41-44]. This low OLR threshold is a critical constraint for microalgae and microalgae residues anaerobic digestion feasibility, requiring (i) longer hydraulic retention times (HRT), i.e. larger digester volume or (ii) lower influent organic matter concentration. Either way, the resulting low volumetric methane yields (LCH<sub>4</sub>/(L<sub>r</sub>·day)) compromises the economic feasibility of microalgae AD.

Anaerobic co-digestion (AcoD), the simultaneous digestion of two or more substrates, is a well-established option to overcome the drawbacks of mono-digestion and improve the economic feasibility of biogas plants. The latter is a result of the higher methane production and the treatment of several wastes in a single facility [45-48]. Microalgae and microalgae residues have been co-digested with a large variety of co-substrates, such as sewage sludge, animal manure, food waste, crops, glycerol, paper waste, and fat, oil and grease (FOG). Although the improvement of the methane production is mainly a consequence of the increased OLR rather than synergisms, ideal co-substrates for microalgae are highly biodegradable carbon-rich substrates, which boost methane production without increasing the nitrogen load [47]. Additionally, microalgae can be used as co-substrate in existing biogas plants. For instance, Schwede [49] explored the possibility of substituting pig

manure by microalgae as source of alkalinity and macro- and micronutrients in corn silage anaerobic digestion.

Multiple microalgae anaerobic co-digestion mixtures and scenarios have been researched. Fig. 1 illustrates the most researched microalgae co-digestion scenarios in the literature, including:

- High-value products biorefinery (Fig. 1A): high-value products are extracted from microalgae and microalgae residues are co-digested with an external co-substrate [14, 24, 25, 50]. In such scenario, CO<sub>2</sub> from biogas combustion can be recycled for microalgae cultivation, while biosecurity regulations may restrict the use of the AD supernatant (liquid fraction after digestate solid/liquid separation) for microalgae cultivation.
- Biodiesel biorefinery (Fig. 1B): lipid-spent microalgae is co-digested with glycerol, by-product of lipids transesterification for biodiesel production [14, 16, 51, 52]. Anaerobic digestion supernatant and CO<sub>2</sub> from biogas combustion are recycled for microalgae cultivation.
- Secondary treatment in wastewater treatment plants (Fig. 1C): a high rate algal pond (HRAP) is used for municipal wastewater treatment (instead of waste activated sludge (WAS)) and harvested microalgae are co-digested with primary sludge [20, 53-58].
- Tertiary treatment in wastewater treatment plants (Fig. 1D): a photobioreactor is installed after the activated sludge unit to improve the quality of the final effluent and harvested microalgae are co-digested with sewage sludge [12, 59, 60].

- Anaerobic digestion supernatant treatment (Fig. 1E): a HRAP is used to remove nutrients from the anaerobic digestion supernatant and harvested microalgae are used as co-substrate. This approach have been studied to decrease the nutrient content of the return stream in wastewater treatment plants [6, 59, 61-64] and to improve the effluent quality of animal manure anaerobic digesters [9, 15, 50, 65-67].
- Microalgae as co-substrate in an existing biogas plants (Fig. 1E): microalgae cultivated outside the biogas plant [49, 68-71] or collected from microalgae blooms [44, 72, 73] are added as co-substrate to an existing anaerobic digester.

The aim of this publication is to present a comprehensive and critical review about the recent achievements and perspectives of microalgae (including cyanobacteria) anaerobic co-digestion. The following sections discuss the anaerobic co-digestion of microalgae and microalgae residues with sewage sludge, animal manure, and a wide variety of agriindustrial wastes. Literature results are summarized in tables, where the methane yield improvement was calculated by comparing the experimental methane yield of the mixture with its theoretical methane yield. The review also identifies several knowledge gaps that warrant further investigation.

## 2. Co-digestion of microalgae and sewage sludge

Sewage sludge (mix of primary and waste activated sludge) is the most researched cosubstrate for microalgae. The cultivation of microalgae in wastewater treatment plants (WWTP) has been used as an alternative to the conventional activated sludge reactor (Fig.

1C), to polish the final effluent (Fig 1D) and to treat the anaerobic digestion supernatant (Fig. 1E) [7, 12, 56, 57, 64]. Additionally, microalgae ponds are a well-known technology for wastewater treatment [5, 74], which eases the adoption of microalgae cultivation systems in WWTP.

On the one hand, the integration of microalgae cultivation as tertiary treatment and supernatant treatment aims to improve nutrients removal (N and P) from wastewater, while generating an additional co-substrate for sewage sludge. The cultivation of microalgae on anaerobic digestion supernatant is of special interest since it has the potential to (i) reduce the nutrient load of the return stream to the head of the plant, which represents up to 20% of the WWTP nutrient load, (ii) mitigate greenhouse gases emissions by using CO<sub>2</sub> from biogas or biogas combustion for microalgae growth, and (iii) generate significant amounts of microalgae as onsite co-substrate, which lowers the uncertainty about co-substrate availability and seasonality [47, 75]. Nonetheless, the supernatant may need to be pretreated and/or diluted to reduce the presence of microalgae growth inhibitory compounds and improve light transmittance [7, 59, 76]. On the other hand, microalgae-based WWTPs, where HRAPs are used as secondary treatment, stand as a low-energy wastewater treatment system for regions with sufficient surface area and solar radiation [5, 56, 58]. In HRAP, microalgae grow in symbiosis with heterotrophic bacteria responsible of organic matter degradation; thus, the harvested biomass consists of a mix community of microalgae, bacteria and protozoa [77]. In this scenario, microalgae from the HRAP are co-digested with primary sludge from the primary settler.

Microalgae and sewage sludge co-digestion is not a new concept, since the first published study dates from 1983, when Samson and LeDuy [78] co-digested *Spirulina maxima* with three different wastes, including sewage sludge. However, the number of papers dealing with this topic has grown exponentially over the last years alongside the growing interest on microalgal-derived biofuels. Most of these studies have been carried out using batch assays, so-called biochemical methane potential (BMP) tests, at mesophilic conditions (Table 1). Nevertheless, several studies have researched the performance of this mixture in lab-scale continuous systems, such as continuous stirred tank reactors (CSTR) (Table 2). The main differences between those studies are related to the microalgae strain, sewage sludge composition (primary and/or waste activated sludge) and the composition of the co-digestion mixture.

Most of the BMP-based studies analyzed a wide range of proportions between both cosubstrates. Beltrán et al. [79], Garoma and Nguyen [63], Lee et al. [80], Mahdy et al. [20], and Neumann et al. [81] tested the co-digestion of different microalgae species and WAS at 25, 50 and 75 %. The same mixture range was tested by Mahdy et al. [20] and Solé-Bundó et al. [53, 55] for primary sludge, by Caporgno et al. [82], Du et al. [83], and Olsson et al. [62] for sewage sludge, and by Lu and Zhang [84] for septic sludge. Exploring a wide range of proportions between microalgae and sludge is needed since the amount and characteristics of both substrates would vary through the year depending on the wastewater temperature and composition, as well as on the treatment plant design and operational conditions [55, 57, 58]. In this regard, Passos et al. [56], who explored the feasibility of a microalgae-based wastewater treatment plant (similar to Fig. 1C), calculated that the proportion between microalgal biomass and primary sludge would be around 30/70% in winter and 60/40% in summer (VS-basis). Peng and Colosi [12], who performed a life cycle assessment on the implementation of a microalgae pond as tertiary treatment (similar to Fig. 1D), estimated that proportion between microalgae and sewage sludge would vary between 5/95% and 20/80% (VSS-basis). Therefore, mixtures where microalgae proportion represent less than 50% of the mixture may represent better WWTP scenarios. Peng and Colosi [12], Olsson et al. [62], Wang et al. [85], and Yuan et al. [59] studied mixtures with microalgae proportion below 50% (Table 1). Finally, Wagner et al. [86] studied the possibility of using bacterial biomass from an enhanced biological phosphorus removal system (similar to WAS) as bioflocculant for microalgae harvesting and subsequent anaerobic co-digestion. According to the authors, using 10 grams of bacterial biomass per gram of microalgae reduced the polymer dosing by 40%.

BMP tests results show that the methane yield obtained from microalgae and sludge mixtures is proportional to the amount of microalgae and sludge. However, some authors have reported synergies (increased methane yield compared to the proportional one) of up to 25% [79, 85, 86]. Thorin et al. [48] noted that in most cases the improved methane yield could not be substantiated if the methane yield uncertainty was considered. Additionally, it should be considered that in full-scale plants minor methane yield improvements due to synergisms would be masked by natural variations of the substrates load, composition and biodegradability.

Although microalgae and sludge co-digestion research has primarily focused on the methane yield, the feasibility of the process is also linked to the kinetics of the limiting step [87, 88]. As a highly particulate substrates, microalgae and sewage sludge anaerobic

digestion is limited by the hydrolysis rate. The reported first-order constant rates for microalgae range between 0.03 and 0.24 day<sup>-1</sup> (average of 0.12 day<sup>-1</sup>); which is at the lower end of the first-order constant rates reported for sewage sludge [28, 89]. With the exception of Wagner et al. [86], publications comparing the degradation kinetics of microalgae and sewage sludge mono-digestion and co-digestion observed a 20 - 50% improvement of the degradation kinetics under co-digestion conditions [34, 55, 79-81]. An improvement in degradation kinetics has also been reported when microalgae was co-digested with other substrates [14, 35, 90, 91]. The reasons behind the kinetic improvement under co-digestion conditions remain unexplored and warrants further research, since it opens the possibility to reduce treatment time or, if treatment time is maintained, improve waste stabilization. However, it should be noted that in BMP tests the apparent degradation kinetics are influenced by the inoculum capabilities [92, 93]. In this regard, Beltran et al. [79], Lee et al. [80], Olsson et al. [34], Solé-Bundó et al. [55], and Wagner et al. [86] used digested sewage sludge as inoculum, while Neumann et al. [81] used granular biomass from a UASB reactor. Digested sewage sludge is a suitable inoculum for this mixture. Moreover, it is the recommended inoculum by Holliger et al. [94] and Raposo et al. [30] when a fully adapted inoculum is not available. However, the correlation between the degradation kinetics observed in BMP tests and continuous reactors is a topic of current research and discussion within the anaerobic community.

Despite the higher methane production, the implementation of anaerobic co-digestion in a WWTP has a direct impact on other key factors, such as the supernatant nutrient content, digestate dewaterability, biosolids quality and biogas composition (e.g.  $H_2S$ ); all of them directly affecting the treatment costs [95-97]. The impact of a co-substrate on digestate

dewaterability, biosolids stability and amount of residual solids to be handled (nonbiodegradable organic matter) are of particular importance since they affect the volume of biosolids to be transported outside the WWTP as well as the digestate management opportunities [34, 53, 59, 98].

Regarding digestate dewaterability, Yuan et al. [59] reported that co-digesting 5 and 15% of Spirulina platensis with WAS improved the digestate dewaterability when compared to WAS alone. Nonetheless, in the same study, the digestate dewaterability worsened when 5 and 15% of Chlorella sp. were co-digested with WAS [59]. Conversely, Wang et al. [85] reported that the anaerobic co-digestion of Chlorella sp. and WAS improved digestate dewaterability at low Chlorella sp. proportions (4 and 11% weigh-basis), but worsened it at a higher proportion (41% Chlorella sp.). However, these results should be interpreted carefully since the dewaterability was measured on digestates obtained from BMP tests. In a BMP test, the properties of the digestate are mostly controlled by the inoculum properties rather than by substrates properties [42, 50]. In continuous lab-scale digesters, Solé-Bundó et al. [53, 55] showed that the dewaterability from the digester treating a mix of primary sludge and pre-treated microalgae (75/25% VS-basis) was better than the dewaterability from the digester treating pre-treated microalgae and the digester treating non-pre-treated microalgae. Olsson et al. [34] also reported an improvement in digestate dewaterability when microalgae was added to sewage sludge (37/63% VS-basis). All previous studies evaluated digestate dewaterability by determining the capillarity suction time (CST), likely due to its simplicity and affordability. However, the CST is a proxy parameter for dewaterability since it does not resemble the actual dewatering process and it fails to predict the solids concentration of the dewatered cake [99]. Future research should

complement CST with other dewaterability methods, such as thermo-gravimetric [100], filtration-centrifugation [101] and rheology analysis [99, 102, 103].

Finally, the circular economy paradigm and the cradle-to-cradle concept require the development of auto-regenerative production systems, where waste products are converted into useful materials [95]. Therefore, beyond biogas production, AD plants need to find sustainable management and disposal solutions for the biosolids [104, 105]. Agricultural reuse is regarded as the best option to recycle the nutrients contained in the digestate [47, 106]. However, this can only be done when the digestate quality fulfils the legal quality requirements. Solé-Bundó et al. [53], who assessed the digestate quality (i.e. concentration of nutrients, heavy metals, pathogens, phytotoxicity and organic matter stability), concluded that the digestate from primary sludge and microalgae co-digestion was more suitable for agricultural reuse than the digestate from microalgae grown on flue gas as  $CO_2$  source increased the heavy metal content in the digestate, making it unsuitable to be used as fertilizer. The authors related the higher heavy metal content in microalgae to the uptake of heavy metals from the flue gas and recommended to carefully assess the source of  $CO_2$ .

#### 3. Co-digestion of microalgae and agri-industrial wastes

Agri-industries supply products to the food and fodder markets as well as a wide range of processing industries. Nonetheless, the production and processing of these products result in the generation of large amount of wastes [107]. AD stands out as a suitable technology to reduce the environmental impact of agri-industrial wastes and increase the energy self-

sufficiency of these industries. However, agri-industrial wastes are characterized by a high C/N ratio, which can affect AD performance due to poor alkalinity and deficit of macroand micronutrients [49, 108-110]. Co-digesting microalgae with agri-industrial wastes has been suggested as an option to overcome these limitations [47, 111]. Additionally, microalgae can be cultivated using marginal soil in regions where other suitable cosubstrates are not available [49, 81]. Conversely, agri-industrial wastes can be used as cosubstrates in microalgae digesters to increase the digester OLR and methane yield without increasing (or even diluting) the nitrogen concentration.

Microalgae have been co-digested with a wide range of agri-industrial wastes, including crops (e.g. corn silage, corn stover, wheat straw), energy crops (e.g. switchgrass, *Pennisetum*), waste paper/sludge, olive mill waste, FOG, and glycerol. Most of the studies focused on improving AD performance by balancing the C/N ratio since agri-industrial wastes present relatively high C/N ratios (>45), while microalgae present relatively low C/N ratios (< 12). Table 3 and Table 4 summarize the studies co-digesting microalgae and agri-industrial waste in BMP tests and in continuous lab-scale reactors respectively.

Fig. 2 illustrates the calculated improvement of the methane yield depending on the C/N ratio for the studies co-digesting microalgae and agri-industrial wastes. The methane yield improvement was calculated by comparing the experimental methane yield of the mixture with its theoretical methane yield. The latter was calculated by the product summation of each substrate methane yield in mono-digestion and their proportion in the mixture [79, 112]. Positive values (>10%) indicate synergism (i.e. the mixture produces more methane than expected), while negative values (<10%) indicate antagonism (i.e. the mixture

produces less methane than expected). Values between -10% and 10% were considered neutral (neither synergistic nor antagonistic) in order to account for the uncertainty around measured methane yields and the propagation of multifarious analytical errors. Most studies target mixtures with C/N ratios ranging between 15 and 30, which falls into the reported optimum range for optimum AD performance [16, 108, 111]. However, both neutral and synergistic responses are observed within this C/N range. Given the variability of improvement in methane yields for a certain C/N ratio, it is clear that optimizing co-digestion mixtures based on the C/N ratio is an oversimplification. The C/N ratio is a proxy for macronutrients availability, ammoniacal nitrogen concentration and system alkalinity. However, it does not consider other important factors, such as substrate biodegradability, secondary risk of inhibition and micronutrients. Thus, the long legacy of using the C/N ratio as key factor to explain the synergisms and antagonisms occurring during anaerobic co-digestion has caused an overlook of the actual mechanisms behind such phenomena.

Regarding the impact of the C/N on methane yield, Solé-Bundó et al. [91] observed that adding NH<sub>4</sub>Cl to wheat straw did not increase its methane yield in BMP tests. One could argue that the impact of adding NH<sub>4</sub>Cl was masked by the inoculum which in BMP testing is typically the main source of macro- and micronutrients, alkalinity and microorganisms [92, 94]. However, a similar result was obtained by Yen and Brune [41] in a research study devoted to co-digest microalgae and waste paper in continuous reactors. Yen and Brune [41] observed that adding NH<sub>4</sub>Cl to decrease the waste paper C/N ratio from 2000 to 21.5 was not enough to explain the synergism occurring during microalgae and waste paper co-digestion. The authors hypothesized that microalgae improved waste paper anaerobic digestion by balancing the C/N ratio and providing a range of essential micronutrients.

Herrmann et al. [42], who co-digested *Spirulina platensis* with three different carbon-rich substrates (i.e. barley straw, beet silage, and brown seaweed) each in a separate CSTR, also noted that the C/N ratio is not the only parameter to consider when optimizing co-digestion mixtures. Besides the digester treating only *Spirulina platensis*, the other three CSTRs were fed with the co-digestion mixture that provided a C/N ratio of 25 (i.e. 15% barley straw, 45% beet silage and 55% brown seaweed on a VS-basis). Hermann et al. [42] reported that the reactor digesting *Spirulina platensis* was inhibited (substantial decrease of the methane yield) when the OLR was increased from 1 to 2 gVS/(L<sub>r</sub>·day), whereas the CSTRs co-digesting barley straw, beet silage and 5 gVS/(L<sub>r</sub>·day) respectively. As the maximum OLR for stable AD operation increased together with the co-substrate proportion, Hermann et al. [42] concluded that the difference in performance was linked to the occurrence of ammonia inhibition rather than the C/N ratio itself.

Synergisms associated with microalgae anaerobic co-digestion have also been linked to other parameters more difficult to quantify and monitor than the C/N ratio, ammonia concentration, and alkalinity. For instance, Schwede et al. [49] claimed that the micronutrients (i.e. Co, Mo, Ni, Na) supplemented by *Nannochloropsis salina* were one of the key factors preventing digestion failure when the OLR was increased from 2 up to 4.7 gVS/( $L_r$ ·day). Indeed, micronutrients (e.g. Co, Mo, Fe, Ni and Se) are well-known cofactors in numerous enzymatic reactions involved in the biochemistry of methane formation [109, 113]. Yen and Brune [41] results may also indicate that the observed increase on cellulase activity (enzyme that catalyzes cellulose hydrolysis) was partly related to the supplementation of micronutrients by microalgae. However, Zhong et al. [44] did not

observe an improvement of cellulose activity when *Microcystis* sp. was co-digested with corn straw, as cellulase activity decreased as the corn straw proportion in the mixture decreased. The role of micronutrients and enzymes activity on anaerobic (co-)digestion performance warrants further research.

Although most studies have emphasized possible synergisms between substrates, more attention should be given to inhibition/antagonism phenomena occurring during anaerobic co-digestion, since they are clear indicators of constraints associated with the co-digestion of a particular co-substrate. In practice, co-substrate selection and dose are primarily controlled by the availability and occurrence of secondary inhibition phenomena (e.g. salinity, heavy metals, ammoniacal nitrogen, volatile fatty acids (VFA), long chain fatty acid (LCFA), biogas H<sub>2</sub>S concentration) [45, 47, 96, 97, 114-118]. For instance, the addition of microalgae into a digester could increase the heavy metals concentration in the digestion media, which may not only impact the AD performance but also the possibility of reusing the digestate on land [14, 34, 53]. In the same way, the addition of microalgae grown on brackish and brine water can increase the concentration of Na<sup>+</sup> and other cations (e.g.  $Ca^{2+}$ ,  $K^+$  and  $Mg^{2+}$ ) in the digestion media, all of them well-known inhibitors of the AD process [116, 119]. Na<sup>+</sup> and K<sup>+</sup> concentrations may also be increased when crude glycerol, by-product of biodiesel production, is used as co-substrate in a microalgae digester; although the main limitation when using crude glycerol as co-substrate is linked to the accumulation of propionate [98]. Similarly, the risk of LCFA inhibition limits the dose of FOG as co-substrate [47, 115, 120]. Finally, it is worth highlighting that antagonisms occurring during co-digestion are more difficult to detect and quantify than synergisms. This is because (i) the impact of inhibitors and intermediate metabolites in BMP testing is diluted, and (ii) long operation time and a certain co-substrate loading rate may be required prior an inhibitor reaches its inhibitory concentrations.

### 4. Co-digestion of microalgae and animal manure

The life cycle assessment and energy analysis carried out by Wang et al. [67] and Zhang et al. [15] showed that treating animal manure anaerobic digestion supernatant with microalgae ponds is an opportunity to reduce the environmental impacts associated with manure management (e.g. eutrophication, global warming) and increase bioenergy production through co-digestion. The configuration analyzed in both studies (similar to Fig. 1E) was found environmentally and energy superior to direct land application and manure anaerobic mono-digestion [15, 67]. Nonetheless, Zhang et al. [15] observed that the profitability of this scheme was highly dependent on the sale price of nutrient credits.

Animal manure (i.e. pig, cattle, and poultry) and microalgae co-digestion has received less attention than microalgae co-digestion with sewage sludge (Section 2) or agri-industrial waste (Section 3). This is likely due to the relatively low C/N ratio of both substrates, which increases the risk of ammonia inhibition. However, the possibility of recovering nutrients, improving the effluent quality, and producing an onsite co-substrate through microalgae cultivation makes manure and microalgae co-digestion worth investigating. Even more when Mahdy et al. [66], who co-digested *Chlorella vulgaris* and cattle manure, showed that anaerobic biomass could be acclimated to tolerate free ammonia and total ammonical nitrogen (TAN) concentrations up to 650 mgNH<sub>3</sub>-N/L and 3.8 gTAN/L respectively.

Most of the animal manure and microalgae co-digestion research has been carried out in BMP tests, with pig manure being the most studied (Table 5). The BMP test is a suitable analytical method to understand the interaction between substrates occurring during codigestion. However, a BMP test is not the most suitable method to assess the impact of inhibitors (e.g. free ammonia), since they are masked by the inoculum [42, 50]. Regarding substrates interaction, Astals et al. [50], Gonzalez-Fernandez et al. [65], Tsapekos et al. [71] and Wang et al. [67] observed that co-digesting microalgae with pig manure increased microalgae anaerobic biodegradability to different extent. An improvement of the methane yield (compared to the proportional one) was also obtained by Mahdy et al. [66] and Prajapati et al. [121] when co-digesting microalgae and cattle manure, and by Li et al. [9] and Menenses-Reves et al. [122] when co-digesting microalgae and poultry manure (Table 5). Mahdy et al. [66] and Prajapati et al. [121] attributed the synergistic effect to the improved C/N ratio, while Li et al. [9] attributed it to the N/P ratio. Although the C/N ratio is the most reported parameter to explain the synergies occurring during anaerobic codigestion, synergism could not always be linked to the C/N ratio [50, 65, 70, 122]. In this regard, Astals et al [50] hypothesized that synergism was due to the addition of specific microbes from pig manure, since other factors previously used to explain co-digestion synergisms (e.g. macro- and micronutrients, C/N ratio, ammonia inhibition, alkalinity) were unlikely to occur under the trialed experimental conditions. The impact of incoming microbes (microbes arriving with the substrate) on anaerobic digestion microbial community and performance is a topic that warrants further research.

Due to BMP tests limitations, continuous experiments are required to better understand the benefits and constraints of co-digesting microalgae and animal manure. However, few

studies have reported the operation of continuous anaerobic digesters co-treating microalgae and animal manure (Table 5). Under mesophilic conditions, Wang et al. [67] co-digested Chlorella sp. and pig manure (10/90% VS-basis) at a HRT of 21 days and an OLR around 1.4 gVS/( $L_r \cdot day$ ), Mahdy et al. [66] co-digested *Chlorella vulgaris* and cattle manure (80/20% VS-basis) at a HRT of 23 days and an OLR of 2.1 gVS/(Lr day), and Menenses-Reyes [70] co-digested lipid-spend Chlorella vulgaris with glycerol and poultry litter (30/3/67% dry-basis) at a HRT of 30 days and an OLR of 0.7 gVS/( $L_r$ ·day). Despite the differences on manure type and OLR, which resulted in quite different pH, TAN, and NH<sub>3</sub> concentrations, these three studies showed that co-digesting Chlorella with manure is technically feasible under mesophilic conditions. Tsapekos et al. [71] is the only study codigesting microalgae (Nannochloropsis limnetica) and animal manure under thermophilic conditions. Tsapekos et al. [71] showed that adding microalgae to a pig manure digester (40/60% VS-basis) increased the digester methane yield and reduced the concentration of VFAs. Menenses-Reyes et al. [70] also reported a lower VFA concentration when microalgae (30/70% dry-basis) or microalgae and glycerol (30/3/67% dry-basis) where codigested with poultry litter. The lower VFA concentration under co-digestion conditions indicates that adding microalgae to an animal manure digester can lead to a more stable process.

#### 5. Anaerobic co-digestion of microalgae residues

The pre-treatment of microalgae has been largely researched since microalgae low anaerobic biodegradability (extent and rate) is one of the major bottlenecks of microalgae anaerobic digestion [22, 125, 126]. Microalgae pre-treatment prior to its anaerobic codigestion has also been used to improve microalgae biodegradability [20, 49, 55, 66, 90, 91,

123, 127]. However, in many cases, the pre-treatment energy/economic balance is negative, i.e. the pre-treatment requires more resources than what is recovered from the additional methane production [22]. Additionally, the impact of the pre-treatment on AD performance is less evident when microalgae are co-digested than when microalgae are mono-digested [55, 91]. Consequently, the incentive to pre-treat microalgae prior to its co-digestion is low. Microalgae pre-treatment (out of the scope of this literature review) have been extensively reviewed in Jankowska et al. [126] and Passos et al. [22].

A more suitable approach may be to pre-treat microalgae to recover high-value compounds (e.g. lipids, proteins, antioxidants, pigments) and treat microalgae residues through anaerobic digestion [23, 128]. Interestingly, several authors have reported that the methane yield of microalgae residues is between 20 and 100% higher than the methane yield of raw microalgae [14, 20, 121, 129-131]. However, as highlighted by Astals et al. [50], the recovery of high-value products will reduce the amount of microalgae diverted to AD and, consequently, methane yields cannot be used to directly compare the amount of methane that will be produced in each scenario. Finally, a factor that is not always taken into account is that microalgae pre-treatment also increases microalgae hydrolysis rate, which further contributes to improve the methane yield of a continuous AD system. The anaerobic co-digestion of microalgae residues after lipid and/or protein extraction with a range of co-substrates is discussed in the following subsections and summarized in Table 6 and Table 7.

# 5.1. Co-digestion of lipid-spent microalgae with glycerol and other co-substrates

 The anaerobic co-digestion of lipid-spent microalgae and glycerol (by-product of biodiesel production) has been investigated by several researchers [14, 16, 51, 52, 81]. The integration of biodiesel production from microalgal lipids and the anaerobic co-digestion of by-products is a biorefinery approach that aims to make the process more economically feasible by (i) maximizing the energy recovery from microalgae, (ii) reducing the amount of residues to be managed, and (iii) reusing the nutrients released during the AD and the  $CO_2$  from biogas combustion for microalgae cultivation (Fig. 1B).

Ehimen et al. [51], who produced biodiesel from *Chlorella* sp. (oil fraction of 0.27 TSbasis) using both conventional (via solvent extraction) and insitu transesterification, calculated a maximum yield of 0.028 g of glycerol per g (dry) of *Chlorella* sp. or 0.038 g of glycerol per g (dry) of lipid-spent *Chlorella* sp. The co-digestion BMP tests carried out using this relative quantity showed that glycerol addition increased the methane yield by 4% and 7% when co-digested with insitu and conventional lipid-spent microalgae respectively. These values are in agreement with the values obtained when the experimental methane yield of the insitu (270 mLCH<sub>4</sub>/gTS) and conventional (220 mLCH<sub>4</sub>/gTS) lipidspent microalgae are combined with the glycerol theoretical methane yield (426 mLCH<sub>4</sub>/gTS). Interestingly, the combination of the glycerol maximum yield (0.038 g of glycerol per g (dry) of lipid-spent *Chlorella* sp.) and a hypothesized volatile-to-total solids (VS/TS) ratio of 0.8 for the lipid-spent *Chlorella* sp. shows that the addition of glycerol would only represent a ~5% increase of the digester OLR (VS-basis). In a subsequent study, Ehimen et al. [16] evaluated the feasibility of co-digesting lipid-spent *Chlorella* sp. with glycerol in continuous digesters under several treatment conditions (i.e. HRT, OLR, C/N ratio and temperature). The addition of glycerol to increase the C/N ratio from 5.4 (mono-digestion) up to 12.4 improved the methane yield from 190 to 300 mLCH<sub>4</sub>/gVS. However, when the glycerol dose was further increased to reach a C/N of 24.2, there was a reduction of the methane yield linked to the accumulation of VFA. It is worth highlighting that the amount of glycerol needed to increase the C/N ratio from 5.4 to 12.4 is much higher than the glycerol generated from microalgal lipids transesterification, with the literature average being 0.03 g of glycerol per g (dry) of lipid-spent microalgae.

The results obtained by Ramos-Suarez and Carreras [14], who co-digested lipid-spent *Scenedesmus* sp. with crude glycerol, showed the same trend as Ehimen et al. [16, 51]. On the one hand, the methane yield of the mixture with the relative proportion between lipid-spent microalgae and glycerol (0.0235 g of glycerol per g (VS) of lipid-spent *Scenedesmus* sp.) did not show a significant difference compared to the methane yield of the lipid-spent microalgae. This is likely due to the small amount of glycerol in the mixture. On the other hand, larger amounts of glycerol (11% VS-basis) were able to increase the methane yield; but when the glycerol concentration was further increased (29% VS-basis) the test showed clear signs of inhibition. From Ehimen et al. [16] and Ramos-Suarez and Carreras [14] results, it can be concluded that a lipid-spent microalgae digester is capable of accepting all crude glycerol produced during the biodiesel production and still has capacity to accept other suitable co-substrates. This organic and volumetric loading spare capacity could be used to digest other waste and further improve the biorefinery economic feasibility.

Besides glycerol, lipid-spent microalgae have been co-digested with other wastes, such as FOG [120], waste activated sludge [81], food waste leachate [132], pig manure [50],

poultry litter [70, 122], and cellulose [131]. Most of these studies have been carried out using BMP tests, however, results suggest that the co-digestion of lipid-spent microalgae with other co-substrates, particularly carbon-rich wastes, is not antagonistic. Therefore, the co-substrate loading rate and subsequent methane production improvement will depend on the (i) AD plant capacity, (ii) co-substrate availability and biodegradability, (iii) secondary inhibitors, and (iv) the impact of the co-substrates on supernatant and digestate quality. However, as previously discussed, most of these factors can only by reliably evaluated in continuous experiments. Park and Li [120], who operated a continuous digesters codigesting lipid-spent Nannochloropsis salina and FOG, observed that the addition of FOG allowed to increase the OLR from 2 to 3 gVS/( $L_r$ ·day) whereas the control reactor (microalgae residues only) was inhibited when the same OLR change occurred; likely due to ammonia inhibition. The co-digester was inhibited when the OLR was subsequently increased to 4 gVS/( $L_r$ ·day); likely due to LCFA inhibition. Park and Li [120] results showed a clear synergy between Nannochloropsis salina and FOG since microalgae provided alkalinity and nutrients while FOG boosted the methane production and diluted ammonia concentration. However, Park and Li [120] results also showed that there was a risk associated with the addition of a co-substrate, particularly when a certain threshold is surpassed. The benefits and constraints of using FOG as co-substrate have already been discussed by Long et al. [115] and Mata-Alvarez et al. [47].

### 5.2. Co-digestion of protein-spent microalgae

The anaerobic co-digestion of protein-spent microalgae has received less attention than the co-digestion of lipid-spent microalgae. This is likely due to (i) the past few years' interest on the production of microalgal-derived biodiesel [11, 17], and (ii) the lower production

costs and higher nutritional value obtained when the whole microalgal biomass is used as feed source [8, 133]. However, protein hydrolyzates have several applications in the food and drink industry (e.g. sport drinks) and the fermentation industry [14, 133]. Additionally, the extraction of proteins would reduce the risk of ammonia inhibition associated with microalgae anaerobic digestion.

To the best of our knowledge, only Ramos-Suarez and Carreras [14] and Astals et al. [50] have studied the anaerobic co-digestion of protein-spent microalgae. Ramos-Suarez and Carreras [14] co-digested protein-spent microalgae with paper sludge and *Opuntia maxima*, while Astals et al. [50] co-digested protein-spent microalgae with pig manure. Although both studies used *Scenedesmus* sp., the method used to release the protein was different since Ramos-Suarez and Carreras [14] used an enzymatic pre-treatment and Astals et al. [50] used free nitrous acid (chemical pre-treatment). Both studies observed that the extraction of protein significantly increased microalgae's methane yield from 140 to 273 mLCH<sub>4</sub>/gVS [14] and from 163 to 222 mLCH<sub>4</sub>/gVS [50]. However, Astals et al. [50] also showed that protein extraction reduced by 54% the amount microalgae diverted to anaerobic digestion, while lipid extraction only reduced it by 14%. The amount of proteins and lipids in microalgae varies depending on the microalgae strain and cultivation conditions, however, microalgae are typically characterized by a larger proportion of proteins than lipids [18, 133]. Consequently, the need to implement anaerobic co-digestion.

#### 6. Conclusions

Anaerobic co-digestion is an opportunity to overcome the drawbacks of microalgae monodigestion and boost the methane production of microalgae and microalgae residues biogas plants. Microalgae can also be used as co-substrate that, besides increasing the digester organic loading rate, can represent a valuable source of alkalinity, macro- and micronutrients. Microalgae have been co-digested with a large variety of co-substrates, such as sewage sludge, animal manure, food waste, crops, glycerol, paper waste, and fat, oil and grease. Most studies have focused on the impact of co-digestion on the methane yield, while less attention has been paid to other important factors, such as supernatant nutrient content, digestate dewaterability, biosolids quality and H<sub>2</sub>S concentration in the biogas. Sewage sludge is the most researched co-substrate since the cultivation of microalgae in wastewater treatment plants could be used as secondary, tertiary and anaerobic digestion supernatant treatment. For microalgae biogas plants, highly biodegradable carbon-rich wastes are the preferred co-substrates since they can increase the digester organic loading rate and methane yield without increasing the nitrogen concentration. Several studies optimized the co-substrate dose by balancing the C/N ratio, however, positive interactions occurring during anaerobic co-digestion could not always be linked to the C/N ratio indicating that other factors have to be considered. Overall, more lab and pilotscale continuous experiments are needed to get a holistic understanding of microalgal anaerobic co-digestion.

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**Fig. 1.** Most researched scenarios for microalgae anaerobic co-digestion: (A) high-value products biorefinery, (B) biodiesel biorefinery, (C) HRAP as secondary treatment in a wastewater treatment plant, (D) photobioreactor as tertiary treatment in a wastewater treatment plant, (E) HRAP as anaerobic digestion supernatant treatment, and (F) microalgae as co-substrate in an existing biogas plant.



**Fig. 2.** Methane yield improvement vs. C/N ratio in studies co-digesting microalgae and agriindustrial wastes

Chlorella sp.WAS41.Chlorella sp. & Scenedesmus sp.Sewage sludge37.Isochrysis galbanaSewage sludge25.Isochrysis galbanaSewage sludge25.Selenastrum capricornutumSewage sludge50.Chlorella vulgarisPrimary sludge50.5Chlorella vulgarisPrimary sludge50.5Lipid-spent Botryococcus brauniiWAS75.2Chlorella sp.WAS21.Chlorella sp.WAS21.Chlorella sp.WAS21.	41:59 (TS) 37:63 (TS) 12:88 (TS) 25:75 (VS) 50:50 (VS) 75:25 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	37 37 33 33 35 35 35 35 35 35	468 <sup>b</sup> 408 378 413 <sup>b</sup> 392 <sup>b</sup> 107 <sup>c</sup> 283 <sup>c</sup>	23 n.d. - 8	Wang et al. [85]
Chlorella sp. & Scenedesmus sp.       Sewage sludge       37         Isochrysis galbana       Sewage sludge       25         Isochrysis galbana       Sewage sludge       50         Selenastrum capricornutum       Sewage sludge       50         Chlorella vulgaris       WAS       75:2         Chlorella vulgaris       Primary sludge       50:5         Lipid-spent Botryococcus braunii       WAS       75:2         Micractinium sp.       WAS       21:         Micractinium sp.       WAS       21:         Othorella sp.       WAS       21:	37:63 (TS) 12:88 (TS) 25:75 (VS) 50:50 (VS) 75:25 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	37 55 33 35 35 35 35	408 378 392 <sup>b</sup> 107 <sup>c</sup> 283 <sup>c</sup>	n.d. n.d. - 8	
12:         Isochrysis galbana       Sewage sludge       25:         Selenastrum capricornutum       Sewage sludge       50:         Chlorella vulgaris       WAS       75:2         Chlorella vulgaris<(pretreated)	12:88 (TS) 25:75 (VS) 50:50 (VS) 75:25 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	55 33 35 35 35 35 35 35 35 35 35 35 35 3	378 413 <sup>b</sup> 392 <sup>b</sup> 107 <sup>c</sup> 283 <sup>c</sup> 293 <sup>c</sup>	n.d. - 8	Olsson et al. [62]
Isochrysis galbanaSewage sludge25Selenastrum capricornutumSewage sludge50Chlorella vulgarisWAS75:2Chlorella vulgarisPrimary sludge50:5Chlorella vulgarisPrimary sludge50:5Lipid-spent Botryococcus brauniiWAS75:2Micractinium sp.WAS21:Chlorella sp.WAS21:	25:75 (VS) 50:50 (VS) 75:25 (COD) 50:50 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	33 35 35 35 35 35 35 35 35 35 35 35 35 3	413 <sup>b</sup> 392 <sup>b</sup> 107 <sup>c</sup> 283 <sup>c</sup> 203 <sup>c</sup>	- 8	
Selenastrum capricornutumSewage sludge50Chlorella vulgarisWAS75:2Chlorella vulgarisPrimary sludge50:5Chlorella vulgarisPrimary sludge50:5Lipid-spent Botryococcus brauniiWAS75:2Micractinium sp.WAS21:Chlorella sp.WAS21:	50:50 (VS) 75:25 (COD) 50:50 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	33 35 35 35	392 <sup>b</sup> 107° 283° 203°		Caporgno et al. [82]
Chlorella vulgarisWAS75:2Chlorella vulgaris (pretreated)Primary sludge50:5Lipid-spent Botryococcus brauniiWAS75:2Micractinium sp.WAS21:Chlorella sp.WAS21:	75:25 (COD) 50:50 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	35 35 35	107° 283° 203°	6	Caporgno et al. [82]
Chlorella vulgaris (pretreated)Primary sludge50:575:275:2Lipid-spent Botryococcus brauniiWAS75:2Micractinium sp.WAS21:Chlorella sp.WAS21:	50:50 (COD) 75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	35 35 35	283° 203°	9	Mahdy et al. [20]
75:2         Lipid-spent Botryococcus braunii       WAS       75:         Micractinium sp.       WAS       21:         Chlorella sp.       WAS       21:	75:25 (COD) 75:25 (VS) 21:79 (VS) 21:79 (VS)	35 35	703°	16	Mahdy et al. [20]
Lipid-spent Botryococcus brauniiWAS75.Micractinium sp.WAS21:Chlorella sp.WAS21:	75:25 (VS) 21:79 (VS) 21:79 (VS)	35	6/1	13	
Micractinium sp. WAS 21: Chlorella sp. WAS 21:	21:79 (VS) 21:79 (VS)		393	7	Neumann et al. [81]
Chlorella sp. WAS 21:	21:79 (VS)	37	236	0	Wang and Park [6]
		37	253	5	
Chlorella sorokiniana WAS 23:	25:75 (VS)	37	442	26	Beltran et al. [79]
Scenedesmus quadricauda WAS 49:	49:51 (VS)	35	222	12	Garoma and Nguyen [63]
Microalgal biomass <sup>d</sup> Primary sludge 92.5	92.5:7.5 (VS)	35	$168^{b}$	n.d.	Hlavinek et al. [54]
Chlorella sp. Septic sludge 50:	50:50 (VS)	35	547	84	Lu and Zhang [84]
Chlorella sorokiniana & WAS <sup>e</sup> 9.5 Scenedesmus sp.	9:91 (VS)	37	560	28	Wagner et al. [86]
WAS <sup>f</sup> 9:	9:91 (VS)	37	400	11	
Scenedesmus sp. (pretreated) WAS (pretreated) 20:	20:80 (VS)	37	187	- 2	Arias et al. [60]
Spirulina platensis Sewage sludge 67:	67:37 (VS)	35	343	15	Du et al. [83]
Scenedesmus sp. & Chlorella sp. Sewage sludge 40:	40:60 (VS)	35	237	2	Olsson et al. [34]
Microalgal biomass <sup>d</sup> Primary sludge 25:	25:75 (VS)	37	291	- 5	Solé-Bundó et al. [55]
Microalgal biomass <sup>d</sup> (pretreated) Primary sludge 25:	25:75 (VS)	37	339	9	

Table 1. Summary of sewage sludge and microalgae co-digestion studies in BMP tests

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Microalgae	Co-substrate	Mixture ratio	AD operation	T (0°)	Working volume (L)	OLR (gVS/(Lr·day))	HRT (days)	Methane yield (LCH4/gVS)	Reference
Spirulina maxima	Sewage sludge	50:50 (VS)	Continuous	35	1.5	3.9	20	0.36	Samson and LeDuy [78]
Scenedesmus sp.	Sewage sludge	25:75 (VS)	Semi- continuous <sup>a</sup>	35	0.45	$0.5^{b}$	15	0.39°	Peng and Colosi [12]
		50:50 (VS)						$0.33^{\circ}$	
Microalgal biomass <sup>d</sup>	Sewage sludge	25:75 (VS)	Semi- continuous <sup>a</sup>	35	0.45	0.5 <sup>b</sup>	15	0.51°	Peng and Colosi [12]
		50:50 (VS)						$0.44^{\circ}$	
Chlorella sp.	Sewage sludge	4-15:96-85 (v/v)	Continuous	55	10	n.d.	28	0.40	Hidaka et al. [64]
Scenedesmus sp. & Chlorella sp.	Sewage sludge	40:60 (VS)	Semi- continuous	37	5	2.4	15	0.20	Olsson et al. [34]
						3.5	10	0.17	
Microalgal biomass <sup>d</sup>	Primary sludge	25:75 (VS)	Semi- continuous	35	1.5	1.2	20	0.46	Solé-Bundó et al. [55]
<sup>a</sup> Feed every 48 hours; <sup>b</sup>	Expressed as gVSS/(Lr.0	day); <sup>c</sup> Expressed as LC.	H4/kgVSS; <sup>d</sup> Micro	valgae colle	scted from a WWTI	<sup>5</sup> ; n.d. stands for not defi	ned		

Table 2. Summary of sewage sludge and microalgae co-digestion in continuous lab-scale reactors

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Microalgae	Co-substrate	Mixture ratio	T (°C)	Methane yield (mLCH4/gVS)	Improvement <sup>a</sup> (%)	Reference
Taihu blue algae	Corn straw	n.d.	35	325	62	Zhong et al. [72]
Spirulina platensis	Switchgrass	33:67 (VS)	35	198	- 2	El-Mashad [68]
		33:67 (VS)	50	236	ŝ	
Nannochloropsis salina	Corn silage	14:86 (v/v)	37	660°	15	Schwede et al. [49]
	Corn cob	25:75 (v/v)		$610^{\circ}$	18	
Dunaliella salina	Olive mill solid waste	50:50 (VS)	35	285	48	Fernández-Rodríguez et al. [111]
Scenedesmus sp.	Opuntia maxima	25:75 (VS)	37	234	65	Ramos-Suárez et al. [110]
Scenedesmus sp.	Opuntia maxima	10:90 (VS)	37	166	L	
Spirulina platensis	Pretreated switchgrass	50:50 (TS)	50	354	10	El-Mashad [117]
Botryococcus braunii	Glycerol	90:10 (VS)	37	430	6	Neumann et al. [81]
Chlorella sp. & Scenedesmus sp.	Sida hermaphrodita	40:60 (VS)	35	352	56	Dębowski et al. [32]
Chlorella sp. & Monoraphidium sp.	Wheat straw	50:50 (VS)	37	289	9	Solé-Bundó et al. [91]
Scenedesmus acuminatus & Scenedesmus quadricauda	Cheese whey (deproteinated)	17:83 (VS)	35	302	6 -	Carminati et al. [35]
	Cellulose	16:84 (VS)		272	- 5	
<i>Chlorella</i> sp. <sup>d</sup>	Coffee husks <sup>d</sup>	14:86 (VS)	35	196	87	Passos et al. [90]
$^{\rm a}$ The methane yield improvement is calculated as as mLCH4/gTS; $^{\rm c}$ Expressed as mLbiogas/gVS; $^{\rm d}$	the product summation of each sub pretreated at 120 °C; n.d. stands for	ostrate methane yield, control of the state	onsidering the	co-substrates proportion	and their experimenta	.1 methane yield in mono-digestion; <sup>b</sup> Expressed

Table 3. Summary of agri-industrial wastes and microalgae co-digestion studies in BMP tests

Namochloropsis adinaCorn slage $14:86$ (v/v) $2$ $n.d.$ $1.0-1.5^{a}$ SchwedeNamochloropsis adina (pretreated)Corn slage $14:86$ (v/v) $2$ $n.d.$ $1.5-1.8^{a}$ SchwedeScenedesmus sp.Corn slage $14:86$ (v/v) $2$ $n.d.$ $1.5-1.8^{a}$ SchwedeScenedesmus sp.Opuntia maxima $25:75$ (VS) $2$ $30$ $0.21$ Ramos-StScenedesmus sp.Opuntia maxima $25:75$ (VS) $2$ $4$ $1.6$ $0.20$ Scenedesmus sp.Opuntia maxima $25:75$ (VS) $2$ $4$ $0.20$ Ramos-StScenedesmus sp.Opuntia maxima $25:75$ (VS) $2$ $4$ $0.20$ Ramos-StScenedesmus sp.Opuntia maxima $25:75$ (VS) $2$ $4$ $0.20$ $0.20$ Nonoraphidium sp. & Monoraphidium sp.Wheat straw $50:50$ (VS) $1$ $2$ $0.21$ Solé-BunScenedesmusSociedesmus sp.Totreated $50:50$ (VS) $1$ $2$ $0.21$ Solé-BunSigeoconium sp. and Scenedesmus sp.Papaya waste $50:50$ (VS) $1$ $2$ $0.24$ $50:6$ -BunScenedesmus acuminatus & ScenedesmusCohese whey $17:83$ (VS) $1,5^{b}$ $30$ $0.22$ CarminatiCohese whey $17:83$ (VS) $15^{b}$ $20$ $0.22$ CarminatiScenedesmusCohese whey $17:83$ (VS) $15^{b}$ $0.22$ Carminati	Microalgae	Co-substrate	Mixture ratio	OLR (gVS/(Lr·day))	HRT (days)	Methane yield (LCH4/gVS)	Reference
Namochloropsis salina (pretreated)Corn silage $14:86 (v'v)$ 2n.d. $1.5-1.8^{a}$ SchwedeScenedesmus sp. $  -$	Nannochloropsis salina	Corn silage	14:86 (v/v)	2	n.d.	$1.0-1.5^{a}$	Schwede et al. [49]
Scenedesmus sp.       4       n.d.       1.8-2.0 <sup>6</sup> Scenedesmus sp.       Opuntia maxima       25:75 (VS)       2       30       0.21       Ramos-St         Scenedesmus sp.       0       1       5       10       0.29       Ramos-St         Scenedesmus sp.       0       25:75 (VS)       2       40       0.29       Ramos-St         Scenedesmus sp.       0       0       2       40       0.32       Ramos-St         Chiorella sp. & Monoraphidium sp.       0       0       2       40       0.31       2         Monoraphidium sp.       Wheat straw       50:50 (VS)       1       20       0.21       Solé-Bun         Sigeoclonium sp. and Scenedesmus sp.       Papaya waste       50:50 (VS)       1       20       0.24       Solé-Bun         Sigeoclonium sp. and Scenedesmus sp.       Cheese whey       17:83 (VS)       1       30       0.25       Caminati         Genedesmus acuminatus & Scenedesmus sp.       Cheese whey       17:83 (VS)       1       0       0.20       0.29         Granded       1       31       0.55       0.29       0.24       Solé-Bun	Nannochloropsis salina (pretreated)	Corn silage	14:86 (v/v)	7	n.d.	$1.5 - 1.8^{a}$	Schwede et al. [49]
Scenedesmus sp.Cpuntia maxima $25:75$ (VS) $2$ $30$ $0.21$ Ramos-StRamos-St $4$ $15$ $0.29$ $6$ $10$ $0.29$ Scenedesmus sp.Cpuntia maxima $25:75$ (VS) $2$ $40$ $0.32$ Ramos-StScenedesmus sp.Cpuntia maxima $25:75$ (VS) $2$ $40$ $0.32$ Ramos-StScenedesmus sp.Cpuntia maxima $25:75$ (VS) $2$ $40$ $0.32$ Ramos-StScenedesmus sp.Chlorella sp. & Monoraphidium sp.Wheat straw $50:50$ (VS) $1$ $20$ $0.21$ Solé-BunMonoraphidium sp.Wineat straw $50:50$ (VS) $1$ $20$ $0.21$ Solé-BunSigeoclonium sp. and Scenedesmus sp.Papaya waste $50:50$ (WW) $1$ $20$ $0.24$ Solé-BunSigeoclonium sp. and ScenedesmusCheese wheey $17:83$ (VS) $1.5^b$ $30$ $0.22$ CaminatiGenedesmus acuminatus & ScenedesmusCheese wheey $15:84$ (VS) $15^b$ $30$ $0.22$ CaminatiColubics $16:84$ (VS) $15^b$ $15^b$ $20$ $0.12$ $0.12$				4	n.d.	$1.8-2.0^{a}$	
4 $15$ $0.29$ Scenedesmus sp. $0$ $0.20$ $0.20$ Scenedesmus sp. $0$ $0.32$ $8$ amos-St $6$ $10$ $2$ $4$ $20$ $0.31$ $7$ $12$ $0.31$ $7$ $12$ $0.31$ $6$ $10$ $12$ $0.31$ $7$ $12$ $0.28$ $6$ $10$ $12$ $0.21$ $0.21$ $0.21$ $505$ -Bun $6$ $11$ $20$ $11$ $20$ $0.21$ $505$ -Bun $6$ $11$ $20$ $11$ $20$ $0.21$ $505$ -Bun $6$ $11$ $11$ $20$ $0.21$ $0.24$ $505$ -Bun $6$ $11$ $11$ $20$ $12$ $0.24$ $505$ -Bun $50$ $50$ $50$ $50$ $50$ $50$ $50$	Scenedesmus sp.	Opuntia maxima	25:75 (VS)	7	30	0.21	Ramos-Suárez et al. [110]
Scenedesmus sp. $6$ $10$ $0.20$ Scenedesmus sp. $0$ puntia maxima $25:75$ (VS) $2$ $40$ $0.32$ Ramos-St $4$ $20$ $0.30$ $3$ $7$ $12$ $0.31$ $7$ $7$ $12$ $0.31$ $7$ $12$ $0.28$ $7$ $7$ $12$ $0.28$ $0.31$ $7$ $7$ $12$ $0.28$ $0.31$ $7$ $7$ $12$ $0.28$ $0.31$ $7$ $7$ $12$ $0.28$ $0.31$ $7$ $7$ $12$ $0.28$ $0.28$ $7$ $7$ $12$ $0.28$ $0.28$ $7$ $7$ $12$ $0.28$ $0.28$ $8$ choraphidium sp. & Chlorella sp. (pretreated) $50:50$ (VS) $1$ $20$ $0.24$ $50:-Bun5igeolonium sp. and Scenedesmus sp.Papaya waste50:50 (VS)1200.25Ca-Barci8 cenedesmus acuminatus & Scenedesmus(deproteinated)17:83 (VS)1.5^{b}300.22Carmination and and and and and and and and and an$				4	15	0.29	
Scenedesmus sp.Chante and sp. $25:75$ (VS) $2$ $40$ $0.32$ Ramos-St. $6$ $7$ $15$ $0.30$ $7$ $15$ $0.31$ $7$ $7$ $12$ $0.31$ $7$ $12$ $0.28$ $7$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $11$ $20$ $0.21$ $806$ -Bund $7$ $11$ $20$ $0.24$ $806$ -Bund $8$ figeoclonium sp. & Chlorella sp. (pretreated) $50:50$ (VS) $1$ $20$ $0.24$ $806$ -Bund $8$ figeoclonium sp. and Scenedesmus sp.Papaya waste $50:50$ (WN) $1$ $20$ $0.24$ $806$ -Bund $8$ figeoclonium sp. and Scenedesmus sp.Papaya waste $50:50$ (WN) $1$ $31$ $0.55$ $Cea-Barci8 figeoclonium sp. and Scenedesmus(deproteinated)17:83 (VS)1.5^b300.22Carminati6 dudricauda6.84 (VS)15^b300.22Carminati$				9	10	0.20	
$4$ $20$ $0.30$ $5$ $15$ $0.31$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.28$ $7$ $12$ $0.21$ $8$ Chlorella sp. (pretreated) $50:50$ (VS) $1$ $7$ $20$ $0.24$ $50!6$ -Bun $5igeoclonium sp. and Scenedesmus sp.Papaya waste50:50 (w/w)15igeoclonium sp. and Scenedesmus20:50 (w/w)1200.55Cea-Barci5ieendesmus acuminatus & Scenedesmus(deproteinated)17:83 (VS)1.5^b300.22Carminati6-Bunce16-sec whey17:83 (VS)1.5^b300.22Carminati71010^{2}10^{2}10^{2}10^{2}10^{2}$	Scenedesmus sp.	Opuntia maxima	25:75 (VS)	7	40	0.32	Ramos-Suárez et al. [110]
5       15       0.31         7       12       0.28         7       12       0.28         7       12       0.21         8       Monoraphidium sp. & Chlorella sp. (pretreated)       50:50 (VS)       1       20       0.21       Solé-Bund         8       Monoraphidium sp. & Chlorella sp. (pretreated)       Wheat straw       50:50 (VS)       1       20       0.24       Solé-Bund         8       Stigeoclonium sp. and Scenedesmus sp.       Papaya waste       50:50 (w/w)       1       31       0.55       Cea-Barci         8       Scenedesmus acuminatus & Scenedesmus       Cheese whey       17:83 (VS)       1.5 <sup>b</sup> 30       0.22       Carminati         9       uadricauda       1       5 <sup>b</sup> 30       0.22       Carminati				4	20	0.30	
7       12       0.28         Chlorella sp. & Monoraphidium sp.       Wheat straw       50:50 (VS)       1       20       0.21       Solé-Bunc         Monoraphidium sp. & Chlorella sp. (pretreated)       Wheat straw       50:50 (VS)       1       20       0.24       Solé-Bunc         Stigeoclonium sp. & Chlorella sp. (pretreated)       Wheat straw       50:50 (VS)       1       20       0.24       Solé-Bunc         Stigeoclonium sp. and Scenedesmus sp.       Papaya waste       50:50 (w/w)       1       31       0.55       Cea-Barci         Scenedesmus acuminatus & Scenedesmus       Checse whey       17:83 (VS)       1.5 <sup>b</sup> 30       0.22       Carminati         quadricauda       totalintee       16:84 (VS)       1.5 <sup>b</sup> 30       0.22       Carminati				5	15	0.31	
Chlorella sp. & Monoraphidium sp.Wheat straw50:50 (VS)1200.21Solé-BundMonoraphidium sp. & Chlorella sp. (pretreated)Wheat straw50:50 (VS)1200.24Solé-BundMonoraphidium sp. & Chlorella sp. (pretreated)Wreat straw50:50 (WN)1200.24Solé-BundStigeoclonium sp. and Scenedesmus sp.Papaya waste50:50 (WN)1310.55Cea-BarciScenedesmus acuminatus & ScenedesmusCheese whey17:83 (VS)1.5b300.22CarminatiquadricaudaCollubee16:84 (VS)1.5b300.12Carminati				7	12	0.28	
Monoraphidium sp. & Chlorella sp. (pretreated)Wheat straw (pretreated)50:50 (VS)1200.24Solé-BuncStigeoclonium sp. and Scenedesmus sp.Papaya waste50:50 (w/w)1310.55Cea-BarciScenedesmus acuminatus & ScenedesmusCheese whey17:83 (VS)1.5b300.22CarminatiquadricaudaCellulose16:84 (VS)1.5b300.12Carminati	Chlorella sp. & Monoraphidium sp.	Wheat straw	50:50 (VS)	1	20	0.21	Solé-Bundó et al. [91]
Stigeoclonium sp. and Scenedesmus sp.Papaya waste50:50 (w/w)1310.55Cea-BarciScenedesmus acuminatus & ScenedesmusCheese whey17:83 (VS)1.5b300.22Carminatiquadricauda(deproteinated)17:83 (VS)1.5b300.22CarminatiGellinbee16:84 (VS)1.5b300.12Carminati	Monoraphidium sp. & Chlorella sp. (pretreated)	Wheat straw (pretreated)	50:50 (VS)	1	20	0.24	Solé-Bundó et al. [91]
Scenedesmus acuminatus & Scenedesmus Cheese whey 17:83 (VS) 1.5 <sup>b</sup> 30 0.22 Carminati quadricauda (deproteinated) 17:84 (VS) 1.5 <sup>b</sup> 30 0.12 Carminati	Stigeoclonium sp. and Scenedesmus sp.	Papaya waste	50:50 (w/w)	1	31	0.55	Cea-Barcia et al. [118]
$Cellificee = 16.84 \text{ (VS)} = 15^{\text{b}} = 30 = 0.12$	Scenedesmus acuminatus & Scenedesmus quadricauda	Cheese whey (deproteinated)	17:83 (VS)	1.5 <sup>b</sup>	30	0.22	Carminati et al. [35]
		Cellulose	16:84 (VS)	$1.5^{b}$	30	0.12	

Table 4. Summary of agri-industrial wastes and microalgae co-digestion in mesophilic continuous lab-scale reactors

Chlorella wlgaris & Scenedesnus obliquusFig manueBMP50:50 (VS)3522015Gonzalez-Ferr ToSeenedesnus obliquusCattle manueBMP50:50 (VS)3729270Prajapati et al.Scenedesnus sp.Pig manueBMP15:85 (VS)372969Scenedesnus sp.Pig manueBMP50:50 (VS)3726011Scenedesnus sp.Pig manueBMP6:94 (VS)35348111Wang et al. [6Chlorella sp.Pig manueBMP6:94 (VS)35348111Wang et al. [6Chlorella sp.Pig manueBMP6:94 (VS)35348111Wang et al. [6Chlorella sp.Chicken manueBMP20:80 (VS)3531Li et al. [9]Chlorella vulgarisChicken manueBMP20:80 (VS)35331000hChlorella vulgarisCattle manueBMP30:50 (VS)35331000hChlorella vulgarisCattle manueBMP33:67 (VS)35331010Mahdy et al. [9]Chlorella vulgarisPig manueBMP30:50 (VS)3532331010Mahdy et al. [9]Chlorella vulgarisPig manueBMP33:67 (VS)351310Mahdy et al. [9]Scenedesnus sp. &Pig manueBMP50:50 (VS)351310MaChlorella sp.Pig manueBMP <th>Microalgae</th> <th>Co-substrate</th> <th>AD operation</th> <th>Mixture ratio</th> <th>T (°C)</th> <th>Methane yield (mLCH4/gVS)</th> <th>Improvement<sup>a</sup> (%)</th> <th>Reference</th>	Microalgae	Co-substrate	AD operation	Mixture ratio	T (°C)	Methane yield (mLCH4/gVS)	Improvement <sup>a</sup> (%)	Reference
Chroacoccus sp.Cattle manureBMP $50:50$ (VS) $36$ $292$ $70$ Prajapati et al.Scenedesmus sp.Pig manureBMP $15:85$ (VS) $37$ $300$ $0$ $0$ Astals et al. [5Scenedesmus sp.Pig manureBMP $15:85$ (VS) $37$ $296$ $9$ $111$ Ghorella sp.Pig manureBMP $6:94$ (VS) $35$ $348$ $111$ Wang et al. [6Chlorella sp.Pig manureBMP $6:94$ (VS) $35$ $348$ $111$ Wang et al. [6Chlorella ulgarisChlorella ulgarisContinuous $10:90$ (VS) $35$ $239$ $31$ $Li et al. [9]$ Chlorella ulgarisCattle manureBMP $80:20$ (VS) $35$ $431$ $10$ Mahdy et al. [7]Chlorella vulgarisCattle manureBMP $80:20$ (VS) $37$ $351$ $n.d.$ $Cheng et al. [7]Chlorella vulgarisCattle manureBMP80:20 (VS)373511000^{5}Chlorella vulgarisPig manureBMP50:50 (VS)37351n.d.Cheng et al. [7]Scenedesmus sp. &Pig manureBMP50:50 (VS)355101n.d.Cheng et al. [7]Scenedesmus sp. &Pig manureBMP50:50 (VS)25:31326n.d.101n.d.Scenedesmus sp. &Pig manureBMP50:50 (VS)55101n.d.101n.d.Scenedesmus sp. Π$	Chlorella vulgaris & Scenedesmus obliquus	Pig manure	BMP	50:50 (COD)	35	220	15	Gonzalez-Fernandez et al. [65]
Scenedesmus sp.Pig manureBMP $15:85 (VS)$ $37$ $300$ $0$ Astals et al. [5 $30:70 (VS)$ $37$ $296$ $9$ $30:70 (VS)$ $37$ $266$ $11$ $Chlorella sp.$ Pig manureBMP $6:94 (VS)$ $35$ $348$ $11$ Wang et al. [6 $Chlorella sp.$ Pig manureBMP $6:94 (VS)$ $35$ $348$ $11$ Wang et al. [6 $Chlorella vulgarisChicken manureBMP20:80 (VS)351900^{9}Chlorella vulgarisCattle manureBMP20:80 (VS)35239311.64  al. [9]Chlorella vulgarisCattle manureBMP80:20 (VS)3543110Mahdy et al. [10]Chlorella vulgarisPig manureBMP33:67 (VS)37351n.d.20.80  colds)Scillatoria tenuisPig manureBMP30:20  (VS)35191n.d.1.64  colds)Scenedesmus sp. &Pig manureBMP30:50  (VS)35191n.d.2.66  colds)Scenedesmus sp. &Pig manureBMP50:50  (VS)55191n.d.2.66  colds)Scenedesmus sp. &Pig manureBMP50:50  (VS)53191n.d.2.66  colds)Scenedesmus sp. &Pig manureBMP50:50  (VS)53191n.d.2.66  colds)Scenedesmus sp. &Pig manureBMP$	Chroococcus sp.	Cattle manure	BMP	50:50 (VS)	36	292	70	Prajapati et al. [121]
$30:70$ (VS) $37$ $296$ $9$ $Chlorella$ sp.Pig manureBMP $6:94$ (VS) $37$ $260$ $11$ $Chlorella$ sp.Pig manureBMP $6:94$ (VS) $35$ $348$ $11$ Wang et al. [6] $Chlorella$ ulgarisContinuous $10:90$ (VS) $35$ $190$ $0^b$ $11$ $Chlorella ulgarisChicken manureBMP20:80 (VS)3523931Li et al. [9]Chlorella vulgarisCattle manureBMP80:20 (VS)352393110Mahdy et al. [6]Chlorella vulgarisCattle manureBMP80:20 (VS)3543110Mahdy et al. [7]Chlorella vulgarisPig manureBMP33:67 (VS)37351n.d.Cheng et al. [1]Scendesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.200Scendesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.n.d.Scendesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.n.d.Scendesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.n.d.Nanochloropsis linneticaPig manureBMP40:60 (VS)5335512n.d.Nanochloropsis linneticaPig manureBMP40:60 (VS)53355121.d.$	Scenedesmus sp.	Pig manure	BMP	15:85 (VS)	37	300	0	Astals et al. [50]
Chlorella sp.Fig manureBMP $6:94$ (VS) $37$ $260$ $11$ Chlorella sp.Pig manureBMP $6:94$ (VS) $35$ $348$ $11$ Wang et al. [6Chlorella 1067Continuous $10:90$ (VS) $35$ $348$ $11$ Wang et al. [9]Chlorella vulgarisChicken manureBMP $20:80$ (VS) $35$ $239$ $31$ Li et al. [9]Chlorella vulgarisCattle manureBMP $80:20$ (VS) $55$ $431$ $10$ Mahdy et al. [10]Chlorella vulgarisCattle manureBMP $80:20$ (VS) $37$ $351$ $n.d.$ $Meho et al. [10]Coscillatoria tenuisPig manureBMP33:67 (VS)35191n.d.Meho et al. [10]Scenedesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.Meho et al. [10]Scenedesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.Meho et al. [10]Namochloropsis limmeticaPig manureBMP40:60 (VS)535312n.d.Meho et al. [10]Namochloropsis limmeticaPig manureBMP40:60 (VS)5353121.d.Meho et al. [10]Namochloropsis limmeticaPig manureBMP40:60 (VS)5353121.d.Meho et al. [10]$				30:70 (VS)	37	296	6	
Chlorella sp.Pig manureBMP $6:94$ (VS) $35$ $348$ $11$ Wang et al. [6]Chlorella 1067Cnicken manureBMP $20:80$ (VS) $35$ $239$ $0^{b}$ Chlorella vulgarisCattle manureBMP $20:80$ (VS) $35$ $239$ $31$ Li et al. [9]Chlorella vulgarisCattle manureBMP $20:80$ (VS) $35$ $431$ $10$ Mahdy et al. [6]Chlorella vulgarisCattle manureBMP $80:20$ (VS) $37$ $351$ $n.d.$ Coscillatoria tenuisPig manureBMP $33:67$ (VS) $35$ $191$ $n.d.$ <i>Oscillatoria tenuis</i> Pig manureBMP $50:50$ (v:v) $25-31$ $326$ $n.d.$ <i>Scenedesmus</i> sp. &Pig manureBMP $50:50$ (v:v) $25-31$ $326$ $n.d.$ <i>Namochloropsis limnetica</i> Pig manureBMP $40:60$ (VS) $53$ $355$ $n.d.$				50:50 (VS)	37	260	11	
Chlorella 1067Continuous $10:90$ (VS) $35$ $190$ $0^b$ Chlorella 1067Chicken manureBMP $20:80$ (VS) $35$ $239$ $31$ Li et al. [9]Chlorella vulgarisCattle manureBMP $80:20$ (VS) $55$ $431$ $10$ Mahdy et al. [1Chlorella vulgarisCattle manureBMP $80:20$ (VS) $55$ $431$ $10$ Mahdy et al. [1ContinuousBMP $30:20$ (VS) $37$ $351$ $1.0$ $n.d.$ Oscillatoria tenuisPig manureBMP $33:67$ (VS) $35$ $191$ $n.d.$ Cheng et al. [1Scenedesmus sp. & Chlorella sp.Pig manureBMP $50:50$ (v:v) $25-31$ $326$ $n.d.$ Panyaping et aNamochloropsis linneticaPig manureBMP $40:60$ (VS) $53$ $355$ $12$ $n.d.$	Chlorella sp.	Pig manure	BMP	6:94 (VS)	35	348	11	Wang et al. [67]
Chlorella 1067Chicken manure $BMP$ $20:80$ (VS) $35$ $239$ $31$ $Li et al. [9]$ $Chlorella$ vulgarisCattle manure $BMP$ $80:20$ (VS) $55$ $431$ $10$ $Mahdy et al. [1]$ $Chlorella vulgarisCattle manureBMP80:20 (VS)3735110mdh et al. [1]Oscillatoria tenuisPig manureBMP33:67 (VS)37351n.d.Cheng et al. [1]Oscillatoria tenuisPig manureBMP50:50 (v:v)25-31326n.d.Panyaping et a 2000 (v:v)Scenedesmus sp. &Scenedesmus sp. &Pig manurePig manureBMP50:50 (v:v)25-31326n.d.Pinder al. [1]Namochloropsis linneticaPig manureBMP40:60 (VS)5353121Tapekos et al. [1]$			Continuous	10:90 (VS)	35	190	$0^{ m p}$	
Chlorella vulgaris (pretreated)Cattle manureBMP80:20 (VS)5543110Mahdy et al. [1(pretreated)Continuous80:20 (VS)37351n.d.Cheng et al. [1Oscillatoria tenuisPig manureBMP33:67 (VS)35191n.d.Cheng et al. [1Scenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et al. [1Scenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et al. [1Scenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et al. [1Nanochloropsis linneticaPig manureBMP40:60 (VS)5335512Tsapekos et al. [1	Chlorella 1067	Chicken manure	BMP	20:80 (VS)	35	239	31	Li et al. [9]
Continuous80:20 (VS)37351n.d.Oscillatoria tenuisPig manureBMP33:67 (VS)35191n.d.Scenedesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aScenedesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aScenedesmus sp. &Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aScenedesmus sp. &Pig manureBMP50:50 (v:v)25-31325n.d.Nanochlorella sp. (pretreated)(pretreated)BMP40:60 (VS)5335512Tsapekos et al	Chlorella vulgaris (pretreated)	Cattle manure	BMP	80:20 (VS)	55	431	10	Mahdy et al. [66]
Oscillatoria tenuisPig manureBMP33:67 (VS)35191n.d.Cheng et al. [1Scenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aScenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aScenedesmus sp. & Chlorella sp. (pretreated)Pig manure292n.d.Panyaping et aScenedesmus sp. & 			Continuous	80:20 (VS)	37	351	n.d.	
Scenedesmus sp. & Chlorella sp.Pig manureBMP50:50 (v:v)25-31326n.d.Panyaping et aChlorella sp.Pig manureScenedesmus sp. & Chlorella sp. (pretreated)Pig manure292n.d.Nannochloropsis linneticaPig manureBMP40:60 (VS)5335512Tsapekos et al	Oscillatoria tenuis	Pig manure	BMP	33:67 (VS)	35	191	n.d.	Cheng et al. [123]
Scenedesmus sp. & Pig manure 292 n.d. Chlorella sp. (pretreated) (pretreated) 355 12 Tsapekos et al Nannochloropsis limnetica Pig manure BMP 40:60 (VS) 53 355 12 Tsapekos et al	Scenedesmus sp. & Chlorella sp.	Pig manure	BMP	50:50 (v:v)	25-31	326	n.d.	Panyaping et al. [124]
Nannochloropsis limnetica Pig manure BMP 40:60 (VS) 53 355 12 Tsapekos et al	Scenedesmus sp. & Chlorella sp. (pretreated)	Pig manure (pretreated)				292	n.d.	
	Nannochloropsis limnetica	Pig manure	BMP	40:60 (VS)	53	355	12	Tsapekos et al. [71]
Continuous 216 n.d.			Continuous			216	n.d.	

Table 5. Summary of animal manure and microalgae co-digestion studies in batch and continuous conditions

Microalgae	Co-substrate	Mixture ratio	T (°C)	Methane yield (mLCH4/gVS)	Improvement <sup>a</sup> (%)	Reference
Lipid-spend microalgae:						
Chlorella sp.	Glycerol	67:3 (v/v)	37	$267^{\mathrm{b}}$	4-7	Ehimen et al. [51]
Scenedesmus sp.	Glycerol	89:11 (VS)	37	255	n.d.	Ramos-Suárez and Carreras [14]
Nannochloropsis gaditana	Cellulose	50:50 (VS)	37	268	n.d.	Barontini et al. [131]
Chlorella vulgaris	Chicken litter & glycerol	30:67:3 (TS)	37	131	$16^{\circ}$	Meneses-Reyes et al. [122]
Protein-spend microalgae:						
Scenedesmus sp.	Paper sludge	74:26 (VS)	37	173	35	Ramos-Suárez and Carreras [14]
Scenedesmus sp.	Pig manure	15:85 (VS)	37	319	5	Astals et al. [50]
		30:70 (VS)	37	302	4	
$^{\rm a}$ The methane yield improvement is calculated as th as mLCH4/gTS; $^{\rm c}$ Result compared to mono-digestio	he product summation of each sub on of chicken litter; n.d. stands for	strate methane yield, co not defined	insidering the	co-substrates proportion	and their experimenta	l methane yield in mono-digestion; <sup>b</sup> Expressed

Table 6. Summary of microalgae residues co-digestion studies in BMP tests

Microalgae	Co-substrate	Mixture ratio	OLR (gVS/(Lr·day))	HRT (days)	Methane yield (LCH4/gVS)	Reference
Lipid-spent Nannochloropsis salina	FOG	33:67 (VS)	2	40	0.45	Park and Li [120]
			4	20	0.12	
			9	13	fail	
Lipid-spent Nannochloropsis salina	FOG	50:50 (VS)	7	40	0.40	Park and Li [120]
			4	20	0.54	
Lipid-spent Nannochloropsis salina	FOG	67:33 (VS)	7	40	0.38	Park and Li [120]
			4	20	0.28	
Lipid-spent Chlorella vulgaris	Chicken litter & glycerol	30:67:3 (VS)	0.7	30	0.27	Meneses-Reyes et al. [70]

Table 7. Summary of microalgae residues co-digestion studies in mesophilic continuous lab-scale reactors