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Review

The Occurrence and Role of *Tetrasphaera* in Enhanced Biological Phosphorus Removal Systems

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Abstract: The application of enhanced biological phosphorus removal (EBPR) in wastewater treatment plants (WWTPs) has commonly been utilized worldwide. However, the optimum efficiency has not been realized over the past decades, prompting many studies and publications. The limitations, especially comprehension of the abundance and actual potential of polyphosphate-accumulating organisms (PAOs), are not fully understood. Recently identified putative PAOs, *Tetrasphaera*, present a vast metabolic versatility compared to *Candidatus* Accumulibacter. The characterisation of *Tetrasphaera* unique abilities to utilize various carbon substrates, volatile fatty acids production and consistent high abundance, presents potential boosts towards the process efficiency improvement. This paper provides the existing knowledge on the physiology, morphology and genetic description of PAOs with a special attention to the current state of research on *Tetrasphaera* and its potential. In addition, process conditions and their influence on the microbial activities in EBPR systems are discussed.

Keywords: phosphorus removal; *Tetrasphaera*; denitrification; enhanced biological phosphorus removal

1. Introduction

Enhanced biological phosphorus removal (EBPR) has emerged as the most powerful phosphorus (P) removal process during municipal and industrial wastewater treatment. For a long time, the EBPR has been considered as one of the most complex processes involved in the metabolic activity of activated sludge systems and has shown promise in terms of the cost, reliability and sustainability [1]. Recent years have brought many research contributions to expand knowledge and improve the process efficiency based on the recognition of the pathways and microorganisms involved.

The EBPR in biological nutrient removal (BNR) systems is mainly carried out by a group of microorganisms known as polyphosphate-accumulating organisms (PAOs). Conventionally, P removal via PAO activity is achieved by triggering anaerobic–aerobic conditions, which considerably increase operational costs related to energy consumption by aerators. The focus has recently been on P removal by denitrifying PAOs (DPAOs) under anaerobic–anoxic conditions to reduce the costs. The DPAOs are capable of using alternative electron sources (nitrate or nitrite) to metabolize intracellular organic compounds under anoxic conditions, and P uptake and denitrification is performed simultaneously [2–5].

Due to the growing interest in the implementation of P removal under anaerobic-anoxic conditions, a special attention has been paid to the microorganisms responsible for that process. Representatives of the genus *Tetrasphaera* are among the recently confirmed putative denitrifying PAO attracting attention of the scientific community. Members of *Tetrasphaera* are able to perform either denitrification or aerobic respiration, depending on the local environmental conditions [6]. All currently characterized *Tetrasphaera* isolates have proven the capability of reducing nitrate only to nitrite, whereas some members revealed the ability to reduce nitric acid to nitrous oxide. Moreover, the *Tetrasphaera* group is capable of carrying out the complete physiological EBPR process compared to other known PAOs, whose activity is more dependent on interspecies relationships [7].

The key aspects of *Tetrasphaera* have been studied intensively, including their classification and taxonomy, development of methods for detection [8], abundance in wastewater treatment plants (WWTPs) [9,10], functions in EBPR and biochemistry [6,11,12]. The ubiquitous occurrence of *Tetrasphaera* in diverse ecological niches, the use of various carbon sources and the ability to produce volatile fatty acids (VFAs) show their extraordinary metabolic potential [4,13].

Interactions and competition between DPAOs and other functional microbial groups in the anaerobic-aerobic cycle enable P removal optimization in activated sludge systems [14–16]. In addition, the interest in DPAO ecophysiology, in particular in the context of *Tetrasphaera* activity related to nitrous oxide emissions, was significant [3]. An emerging approach to enhance the full-scale EBPR is optimization by the application of mathematical modelling. The conventional models are thought to favor *Ca.* Accumulibacter over other PAOs, such as *Ca.* Halomonas phosphatis, *Tessaracoccus*, as well *Tetrasphaera* [8]. To ensure the highest prediction accuracy of the model, it is strongly recommended to extend and develop currently available models for multiple PAO groups, differentiated in terms of the growth rate and physiology.

This study aims to consolidate the existing knowledge on the role of *Tetrasphaera* by reviewing their physiological and metabolic characteristics, the occurrence and abundance in WWTPs and factors influencing their growth. Additionally, it highlights the knowledge gaps and research challenges in the field of EBPR microbiology as well as presents scientific approaches to overcome these limitations, including the meta-analysis and models. Furthermore, this study investigates the abundance of *Tetrasphaera* and their response to the local process conditions, such as dissolved oxygen (DO) concentration, pH, temperature and influent characteristics.

2. Historical Perspective of Microorganisms Involved in EBPR

The first observations of EBPR date back to the mid-20th century and the findings obtained in laboratory scale experiments [17] and to a minor extent at full scale plants [18]. The principles of EBPR were formulated by Barnard [19,20], whose experiments clarified the need for anaerobic contact between activated sludge and influent wastewater prior to aerobic treatment to accomplish P removal. Subsequently, Barnard [21] used the term Phoredox to represent any process with an anaerobic/aerobic sequence to promote the EBPR technology concept (Figure 1a).

The performance and start-up process of the first full scale Phoredox system, which was launched in 1973, was briefly reported by Levin et al. [22]. In parallel, an alternative side-stream P removal technology, called PhoStrip, [1]. was developed based on the separation of the enriched side stream liquor treated with lime (Figure 1b).

While the nature of P removal was initially considered as chemical, Fuhs and Chen [23] found *Acinetobacter* as the primary microorganisms responsible for EBPR. These organisms responded to VFA in the influent wastewater under anaerobic conditions by releasing stored phosphate. Bacteria affiliated to *Acinetobacter* were considered as the key PAO responsible for the EBPR, mainly due to the limitations of the cultivation techniques applied for the microbial characterization at that time [11]. Significant advances in the microbial research have been achieved over the years and novel bacterial groups involved in the P metabolism were identified. Those microorganisms were able to store P in their cells in the form of energy-rich polyphosphates, resulting in the P content as high as 20 to 30 percent by dry weight [1]. The anaerobic zone free of nitrate and DO was found to favour the PAOs activity over other heterotrophs. In the following years, Betaproteobacteria were determined as the dominant bacterial group in the P metabolism [24], as well as the presence of *Rhodocyclus*-related bacteria was observed and linked with EBPR [25,26]. Subsequently, modern microbial tools without the cultivation step, established *Ca.* Accumulibacter as the most important member of PAOs, with the share ranging from 0.6 to 33.1% [27,28]. Insights into the biochemical characteristics of the *Ca.* Accumulibacter were applied to propose mathematical models of the EBPR within Activated Sludge Model [29]. Basic metabolic

models are based on the assumption that PAOs exhibit intracellular phosphorus and energy storage in the form of poly-P and polyhydroxyalkanoates (PHA), respectively [30]. The stored PHA provide energy for the growth of PAOs when exposed to anoxic condition due to the capability of simultaneous denitrification and P uptake.
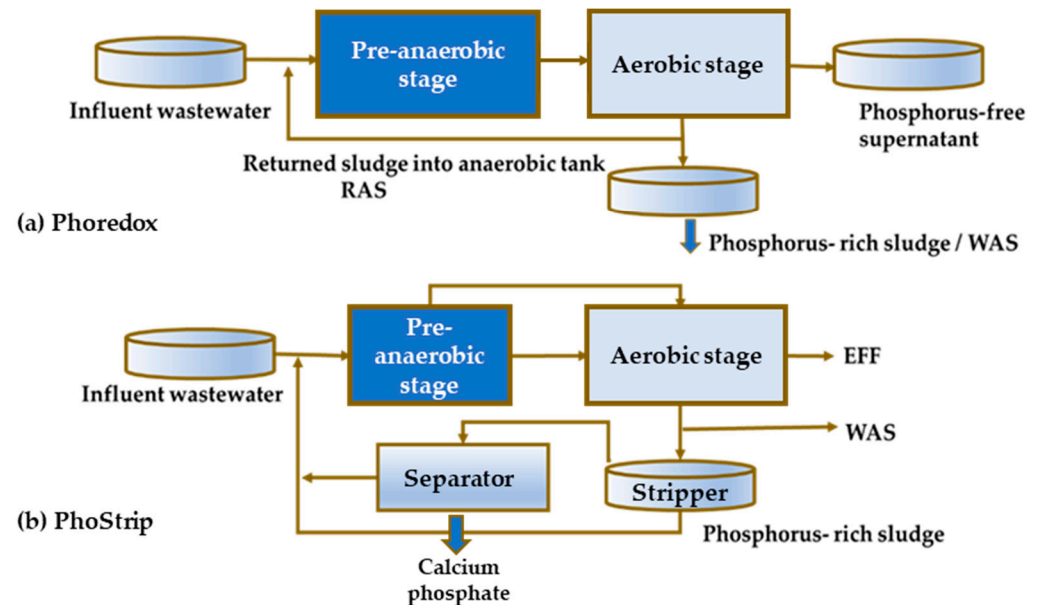
Figure 1. A typical configuration for P removal in the mainstream the mainstream EBPR (Phoredox process) (**a**), sidestream EBPR (PhoStrip process) (**b**) (abbreviations: EFF—effluent, RAS—recirculated activated sludge, WAS—waste activated sludge).

The novel microbial tools comprise deoxyribonucleic acid (DNA) and ribonucleic acid (RNA) polymorphism analyses (e.g., 16S rRNA high-throughput gene sequencing, metagenomics, fluorescence in situ hybridization (FISH) and their modification) as well as flow cytometry and Raman spectroscopy. Along with the development of those tool, the contribution of particular bacterial groups in EBPR could be revised greatly [4,28,31].

The next advance in understanding the microbiology of EBPR was the discovery of new PAO and capability of simultaneous P and nitrogen (N) removal under anoxic conditions. Genus *Tetrasphaera* is the most recently confirmed putative PAO. From the first characterization of the isolated *Tetrasphaera* strain [32,33], it was found that representatives of this genera show a large and often predominant number in full-scale WWTPs [7,34]. Moreover, versatile metabolic capabilities of *Tetrasphaera* have been recognised, including the capability of fermenting glucose and amino acids to produce VFA in the anaerobic zone, thereby enhancing a pool of the available substrates for EBPR. The main difference with respect to the typical PAO is that *Tetrasphaera* are capable of storing other (than PHA) intracellular compounds and use nitrate, but not nitrite, in addition to DO as an electron acceptor [1]. The characterization of *Tetrasphaera* and their assignment to PAO drew attention to this group, especially in terms of the competition and interaction with *Ca.* Accumulibacter [3,35].

In the recent years, several new genera have been proposed as potential PAOs, including *Dechloromonas* or *Candidatus* Microthrix, but only members of the betaproteobacterial genera *Ca.* Accumulibacter [7] and the actinobacterial genus *Tetrasphaera* were consistently found in high abundances in full-scale EBPR plants [34]. For instance, it was proven that approximately 24–70% of total P removed in Danish WWTPs was directly attributed to *Ca.* Accumulibacter and *Tetrasphaera* [7]. The relative abundances of other PAO within activated sludge are usually significantly lower. In the study by Seviour and McIlroy [36], the relative abundance of *Acinetobacter* reached 1.2%, whereas the relative abundances below 1% were reported for *Dechloromonas*, another newly characterized PAO group [37].

3. Morphology, Physiology and Phylogeny of *Tetrasphaera*

Tetrasphaera is a bacterial genus that belongs to the *Intrasporangiaceae* family within the *Actinomycetia* class and initially contained eight proposed species [38]. Zhang and Kinyua [8] listed the following representatives of the genus *Tetrasphaera*: *Tetrasphaera japonica*, *Tetrasphaera australiensis*, *Tetrasphaera elongata*, *Tetrasphaera jenkinsii*, *Tetrasphaera vanveenii*, *Tetrasphaera veronensis*, *Tetrasphaera duodecadis* and *Tetrasphaera remsis*. Based on the distinct morphological, biochemical characteristics (the capability of PHA storage) and probes applied for detection via FISH), particular *Tetrasphaera* representatives were divided into three clades: clade I including *T. elongata* and *T. duodecadis;* clade II *including T. jenkinsii*, *T. australiensis*, *T. veronensis* and the filamentous *Candidatus* Nostocoida limicola, clade III containing uncultured clones [39]. However, the phylogenetic classification of the genus *Tetrasphaera* has not been definitively clarified and is subject to continuous revision. For instance, important updates to representatives of clade III were provided by Singleton et al. [40], who postulated separation of the two novel genera *Ca.* Phosphoribacter and *Ca.* Lutibacillus from the genus *Tetrasphaera*. Similarly, the whole genome sequence analyses by Nouioui et al. [41] revealed the need to reclassify *T. duodecadis, T. remsis,* and *T. elongata* into *Phycicoccus duodecadis, Knoellia remsis* and *Phycicoccus elongatus*, respectively.

First important insights into the morphology and biochemical characteristics of *Tetrasphaera* have been provided by Maszenan et al. [33]. Representatives of *Tetrasphaera* were characterized as an aerobic, Gram-positive cocci, mostly clustered in tetrads and less often in pairs. Due to the lack of flagella, *Tetrasphaera* were considered as a non-motile. Despite P removal activity, members of *T. japonica* and *T. australiensis* did not reflect the ability to store PHA granules, which suggested the role of other intracellular compounds in their EBPR metabolism.

Further cultivation studies by Hanada et al. [32] on biomass from a EBPR system led to the characterization of new species T. elongata., which showed a versatile morphology (oval to rod-shaped) and capability to metabolize wide groups of complex organic compounds, including sugars, alcohols and organic acids. By a positive result of Neisser staining, intracellular polyphosphate granules occurrence has been confirmed and provide evidence of the characterized isolate contribution to P metabolism. Further studies focused on pure cultures of T. elongata have demonstrated archetypical PAO characteristics [8].

A consensus is yet to be built on generally accepted biochemical transformation models for EBPR by *Tetrasphaera*, in particular recognition of the storage compounds involved in EBPR. To date, the capability of intracellular PHA storage has been identified in several *Tetrasphaera* species, including *T. japonica T. jenkinsii T. vanveenii T. veronensis* [14]. However, the ability of *Tetrasphaera* species for anaerobic P release, aerobic/anoxic uptake patterns, and accumulation of intracellular poly-P granules remain inconclusive [12].

Important updates into the characterization of the *Tetrasphaera* metabolism have been by Close et al. [42], who conducted their study on an enriched *Tetrasphaera* culture. In that study, more complex nature of the intracellular compounds cycling within EBPR has been highlighted, where PHA storage was accompanied with the complex amino acids' metabolism, mainly related to aspartic and glutamic acid accumulation within the cell. Moreover, the authors identified that polyhydroxyvalerate (PHV), rather than polyhydroxybutyrate (PHB) (typical for PAO) seems to be more specific feature of *Tetrasphaera* metabolism within EBPR. Despite PHA and intracellular amino acids, members of Tetrasphaera have shown the ability to store glycogen [11,12]. In contrast to the other PAOs, *Tetrasphaera* synthesize glycogen and release phosphate under anaerobic conditions, then metabolize glycogen as an energy source to produce polyphosphate after transition to the aerobic/anoxic environment [2]. Those metabolic features makes this bacterial group versatile in terms of ecophysiology [38].

As postulated by Barnard et al. [2], *Tetrasphaera* exhibits an alternative P depletion mechanism compared to *Ca.* Accumulibacter and other "traditional" PAOs with a potentially more effective P removal efficiency. There is scarce evidence concerning the ability of *Tetrasphaera* to perform denitrification within EBPR. The existing studies emphasize the

fact that *Tetrasphaera* are less capable of removing P under anoxic conditions compared to *Ca*. Accumulibacter [6]. On the other hand, *Tetrasphaera* showed a high performance of N removal (>80%), indicating the ability of *Tetrasphaera* to reduce NO_3-N to NO_2-N similar to their counterpart DPAOs. In the studies by Marques et al. [6], a strong denitrifying activity by *Tetrasphaera* was observed in a mixed culture, where *Tetrasphaera* contributed to 60% of the general bacterial population. *Tetrasphaera* had the capability of denitrification, even though the specific P-uptake rate under anoxic conditions was insignificant compared to *Ca*. Accumulibacter [14].

3.1. Carbon Sources

Currently, the research interest has focused on the identification of preferable carbon sources that can be metabolized by *Tetrasphaera* to release P. The most frequently tested substrates were acetate, glucose, glutamate, glycine and lactate [14]. Moreover, *Tetrasphaera* exhibit the ability to assimilate a wider range of carbons sources, including amino acids, sugars, higher VFAs under anaerobic conditions [11,38,43]. Other processes have been demonstrated to produce an energy source for aerobic P uptake by *Tetrasphaera*, including fermentation of amino acids and sugars, the subsequent storage of either amino acids or glycogen anaerobically, and the use of internally stored substrates [11,12,39]. For instance, laboratory experiments with an enriched culture of *Tetrasphaera*, fed with casein hydrolysate as the sole carbon source, showed that *Tetrasphaera* can metabolize amino acids and were the main microorganisms responsible for aerobic P removal [39]. Moreover, P uptake and release has been detected in addition to the typical substrates (acetate and glucose) with other organics, including formic acid, propionate, butyric acid, pyruvate, lactate, ethanol, glucose, oleic acid, aspartic acid, glutamic acid, leucine, glycine, thymidine and mixed amino acids [43]. In some cases, *Tetrasphaera* isolates revealed different preferences for carbon sources in achieving P release. For instance, in the studies by Nguyen et al. [12], the application of glucose and glutamic acid was performed under anaerobic condition with *Tetrasphaera* isolates, despite substrate utilization no P release was observed. Species *T. australiensis*, *T. japonica* and *T. elongate* responded positively to acetate, whereas propionate in addition to acetate favored *T. jenkinsii*, *T. vanveenii* and *T. veronensis* [33]. In the studies by Marques et al., [39], glucose, aspartate, glutamate and glycine were explored with an enriched *Tetrasphaera* culture. The results from fluorescence in situ fluorescence in situ hybridization with microradiography FISH-MAR showed that *Tetrasphaera* could perform P-release anaerobically with each of those carbon sources used solely.

In terms of the intracellular compounds, glycogen has been considered an important energy storage compound in *Tetrasphaera* [11]. Kong et al. [43] and Nguyen et al. [38] demonstrated that mixed cultures predominated by *Tetrasphaera* were able to consume glucose anaerobically to promote P uptake aerobically. Other experiments with a pure culture of *T. elongate* showed the typical PAO phenotype with glucose as a preferable carbon source [11]. Moreover, glycogen production in the anaerobic phase was also observed in another study not only with glucose, but also glutamate and aspartate supplied as an external carbon source [39].

3.2. Metabolic Models of EBPR

In contrary to *Ca*. Accumulibacter, *Tetrasphaera* are able to synthesize and store anaerobically wider range of the extracellular compounds. Thus, metabolic pathways of P removal by *Tetrasphaera* differ from the metabolic models established for the typical PAOs with acetate as the main substrate. Moreover, due to the capability to decompose complex organic compounds, *Tetrasphaera* form substrate dependencies with other PAOs, including *Ca*. Accumulibacter.

Initially, EBPR has not been linked with denitrification, e.g., in the Activated Sludge Model No 2 (ASM2) [29]. Using the theory of Mino et al. [44], Smolders et al. [45] developed the anaerobic metabolic model of PAO fed with acetate as a single carbon source. According to that theory, PAO transported acetate across the cell membrane and convert it into acetyl-

CoA with the process energy of cleaving poly-P and releasing phosphate from the cell. The parameter represents the ATP required for the transport of 1 C-mmol acetate across the cell membrane [45,46]. Acetyl-CoA was found to be linearly dependent on pH. The origin of the reducing power (i.e., nicotinamide adenine dinucleotide (NADH)), required for PHA synthesis, has been debated by many authors, with Mino et al. [44] supporting its origin from internal glycolysis.

However, Vlekke et al. [47] had earlier demonstrated the capacity of DPAO which led to an update of the ASM2 to the ASM2d [29]. The characterization of the *Tetrasphaera* metabolism, especially in terms of the capability of complex carbon compounds utilization, provided further insights into the potential pathways within EBPR (Figure 2).
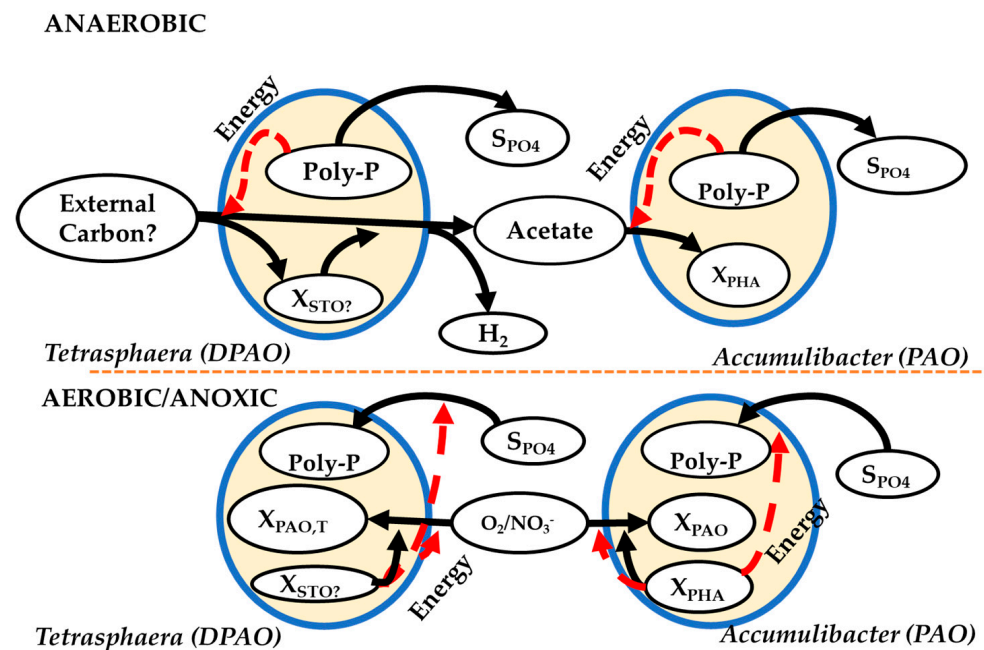
Figure 2. Comparison of the metabolic pathways and biochemical transformation of P between *Tetrasphaera* and *Ca.* Accumulibacter in full-scale EBPR. S_{PO4}—external phosphate, X_{PHA}—internal polyhydroxyalkanoates, X_{PAO}—biomass of the PAO, $X_{STO?}$—not yet characterized internal carbon storage material, $X_{PAO, T}$—biomass of *Tetrasphaera*. Adapted from Makinia and Zaborowska [48].

During the anaerobic phase, complex carbon compounds, such as, glucose, are taken up and either stored as glycogen or fermented to acetate. The energy required for glycogen synthesis is supplied by fermentation and/or polyphosphate decomposition to orthophosphate. In the aerobic/anoxic phase, the stored glycogen is degraded, supplying energy for the growth and enhanced P uptake, followed by replenishing the polyphosphate storage.

Various experimental studies have highlighted metabolic models regarding P removal by *Tetrasphaera* [11,12,39]. The majority of those models have been based on the applied carbon sources and microbial cultures (mixed or pure culture) [49]. Kristiansen et al. [11] proposed a model describing the metabolism of *T. elongata* as a representative in EBPR with glucose as the substrate. Under anaerobic conditions, *Tetrasphaera* take up glucose using poly-P as an energy source and glucose can be stored as glycogen. Under the subsequent aerobic conditions, the stored glycogen can be used for the growth and replenishing supplies of poly-P. A model of Nguyen et al. [12] incorporated glycine as a carbon source with no glycogen. The intracellular glycine was accumulated under anaerobic conditions along with small amounts of glutamine, serine, and alanine. These intracellular metabolites could subsequently be used to support the aerobic P uptake. Moreover, *Tetrasphaera* share some key metabolic pathways with *Ca.* Accumulibacter, such as tricarboxylic acid cycle (TCA) and poly-P degradation/synthesis. Overall, representatives of *Tetrasphaera* are extremely versatile, capable of surviving in highly dynamic environments and highly

abundant in WWTPs. Therefore, a reliable generic model is still missing, and a consensus has not yet been built on general metabolic models for *Tetrasphaera* within EBPR.

Metabolic modelling and prediction of bacterial activity can provide useful information for the process optimization and design purposes [27]. For instance, modelling studies by Oehmen et al., [50] enabled to distinguish two subgroups of PAOs, PAO I able to denitrify from nitrate to N_2 gas and PAO II performing denitrification with nitrite, as the preferable N source. In addition, the authors found that despite anaerobic kinetic parameters for all PAO (PAO I and PAO II) and GAO subgroups are constant, the actual process rates were strongly dependent on the activity of each specific bacterial group.

With regard to *Tetrasphaera*, research on their metabolism is still ongoing, particularly on storage products in the anaerobic phase [49]. *Tetrasphaera* models are considered directly relevant to the models of the EBPR incorporated in ASM and metabolic models [51]. However, an update of the currently available models is necessary as the knowledge of *Tetrasphaera* biochemical properties and interactions with other bacterial functional groups is continuously increasing.

4. Occurrence of *Tetrasphaera* in EBPR Systems

Tetrasphaera abundance has been consistent and demonstrated their predominant role in various studies. Full-scale EBPR systems had higher abundances of *Tetrasphaera* representatives than *Ca.* Accumulibacter in many previous studies, where their maximum contribution was estimated at 30% of the total biomass [12,34,38]. Stokholm-Bjerregaard et al. [34] detected *Tetrasphaera* in large amounts, i.e., up to 35% of the bacterial population, significantly outcompeting *Ca.* Accumulibacter. Moreover, using 16S rRNA amplicon sequencing and quantitative FISH, Herbst et al. [52] found that *Tetrasphaera* the most abundant genus in a Danish WWTP, accounting for 30% of the activated sludge community. These findings were confirmed by a survey of 32 full-scale EBPR plants in 12 countries, where higher abundances of *Tetrasphaera* were reported in most cases [3] by 16S rRNA high-throughput gene sequencing. In that study, the *Tetrasphaera* abundance in EBPR systems was in the range from 1.3% to 11.9%. According to the recent global survey over the bacterial community structure in EBPR systems across 12 countries from 5 continents (MIDAS project), the highest average abundance of *Tetrasphaera* was indicated in terms of both DPAOs subpopulation and a general bacterial community. The average *Tetrasphaera* abundance constituted 4.60%, and prevailed other DPAOs from *Dechloromonas* (2.84%), *Ca.* Accumulibacter (1.19%), *Ca.* Microthrix 0.85% and *Halomonas* at 0.01% (Figure 3). The predominance of *Tetrasphaera* in EBPR systems was confirmed also with the application of other than next generation sequencing techniques. For instance, the use of Raman spectroscopy technology for in situ intracellular compound quantification, detected a higher abundance of *Tetrasphaera* than *Ca.* Accumulibacter in full-scale WWTPs in Denmark [7]. Singleton et al. [40] showed that representatives of the former *Tetrasphaera* clade III, in particular the newly established genus *Ca.* Phosphoribacter, were the dominant PAOs in EPBR systems in Denmark.

However, as suggested by Close et al. [42], the overall abundance and contribution of *Tetrasphaera* within the total bacterial community is not unequivocally related to the potentially obtained P removal rates. The authors suggested that in case of this issue, the composition of *Tetrasphaera* clades played more important role.

An increased scientific interest in *Tetrasphaera* has been observed in terms of the number of publications with the key words "*Tetrasphaera* in wastewater treatment plants" and "PAO in wastewater treatment plants" based on the Scopus database between 1999 to 2021 (Figure 4).
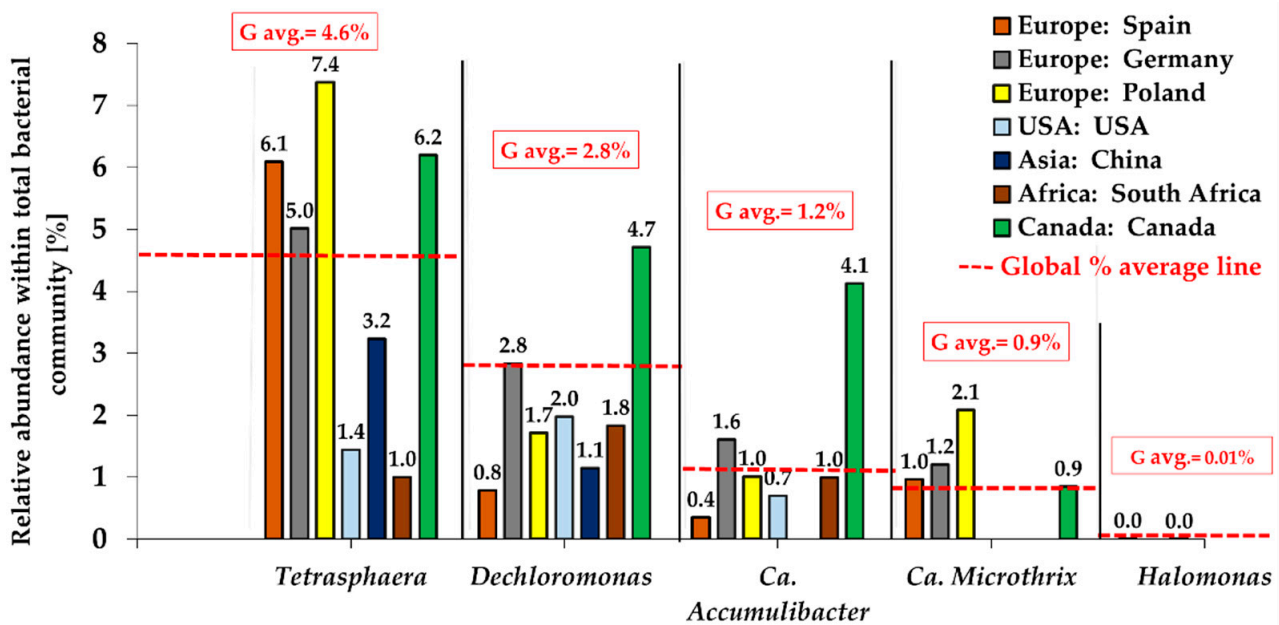
Figure 3. Overview of the global abundance of the denitrifying PAOs within the general microbial community in EBPR systems (based on data from [4,34]).
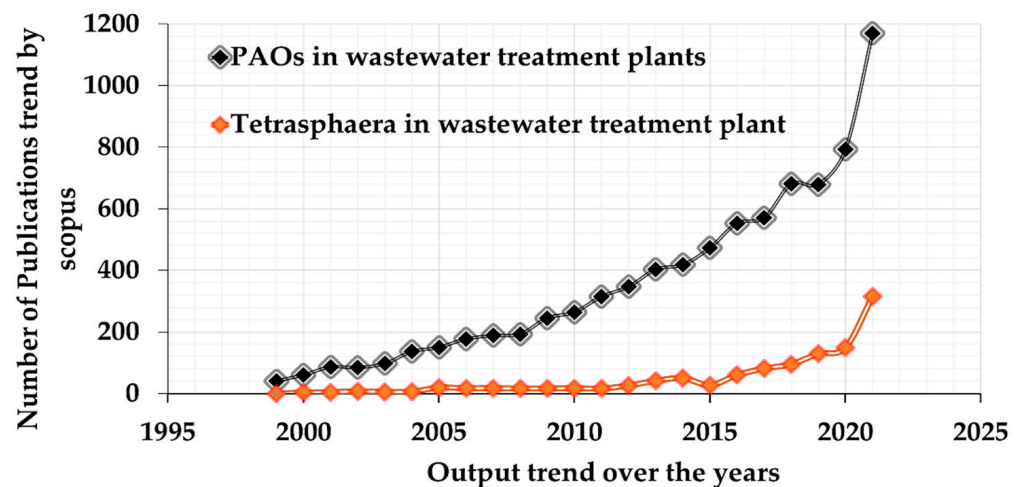

Figure 4. Comparison of PAO and *Tetrasphaera* mentions in publications related to EBPR. Summary of the research trends over the past two decades against the number of scientific papers with the mention of PAOs and *Tetrasphaera*.

5. Implications of *Tetrasphaera* on EBPR Configuration and Operation

The identification of PAOs/DPAOs as well their relationship is a crucial step in optimizing P removal efficiency in WWTPs [14]. Maximizing the P removal fraction achieved in anoxic conditions can significantly reduce the operational costs of EBPR systems. Currently, several different process configurations are available, in which both P and N removal are combined [27].

The studies by Meinhold et al. [53] highlighted two different groups of PAO, including aerobic PAOs (APAOs) and denitrifying PAOs (DPAOs). The APAOs can use only DO as an electron acceptor, whereas DPAO can use either DO or nitrate as an electron acceptor. Oehmen et al. [50] postulated that DPAOs have different denitrification capabilities and can be classified based on their reduction abilities towards nitrate or nitrite. Type I Accumulibacter (DPAO I) showed the ability to reduce nitrate, whereas Type II Accumulibacter (DPAO II) had the ability to reduce nitrite only (Table 1). The representatives

of *Tetrasphaera* reflect differential preferences for electron donors and acceptors. While genera *T. australiensis, T. japonica, T. elongata* show similar metabolic properties to DPAO I with acetate as the key carbon source, whereas the aerobic pathway and preference of propionate are specific features for of *T. jenkinsii, T. vanveenii* and *T. veronensis* [14].

Table 1. Summary of preferable electron acceptors by the main functional bacterial groups involved in EBPR.

Bacterial Functional Group	Electron Acceptors
APAO	DO
DPAO I	$NO_3^-\rightarrow NO_2^-\rightarrow N_2O\rightarrow N_2$
DPAO II *	$NO_2^-\rightarrow N_2O\rightarrow N_2$
GAO	$NO_3^-\rightarrow NO_2^-\rightarrow N_2O\rightarrow N_2$
T. australiensis T. japonica T. elongata	$NO_3^-\rightarrow NO_2^-\rightarrow N_2O\rightarrow N_2$
T. jenkinsii, T. vanveenii, T. veronensis	DO

Note(s): * Reduction abilities towards nitrate or nitrite of the EBPR microbes.

According to Mino et al. [54], the denitrification ability of PAO is essential for design of activated sludge system configurations. Two stage anaerobic-anoxic configurations have been considered with the focus of DPAOs activity with EBPR, and their potential role in the anoxic P uptake and available carbon source for denitrification. The main idea of the modern EBPR systems, is to ensure more space to the anoxic zone as compared to the aerobic zone. However, even with these modifications, the DPAOs contributions to denitrification and the total P uptake usually vary in the range of 0–25% and 0–62%, respectively.

This suggests that despite DPAOs capability to grow under both anoxic and aerobic conditions, there is still a lower efficiency in the utilization of stored intracellular compounds, such as PHA or glycogen. It is apparent that DPAOs may have lower performance than APAO in terms of the usage of readily biodegradable COD [48].

The presence of other microorganisms, such as glycogen accumulating organisms (GAO) in EBPR systems, is an additional, important issue that can affect the EBPR performance [55,56]. With relative abundance of 36.0–42.6% [57,58]. *Defluviicoccus* and 15.3–24.9% of *Candidatus* Competibacter, are considered as the most dominant GAOs in WWTPs. The predominance of *Defluviicoccus* is strongly associated with a higher relative ratio of propionate over acetate in the feed. Propionate uptake rates by these bacteria are comparable to *Candidatus* Accumulibacter and much higher than *Candidatus* Competibacter [59]. Notably, it remains unclear which factors govern the changes in the known GAO community composition as well competition with PAOs, including solids retention time (SRT) [11,60], C/P ratios [61] and available carbon sources [62], which have been evidenced to be associated with GAO shifts. The challenge of identifying PAO and GAO metabolisms is merely based on the functional genes associated with phenotypes characteristics, such as the ability of cycling of phosphorus, polyhydroxyalkanoates (PHA) and/or glycogen, as intracellular storage materials within bacterial cells [13,63]. On the other hand, investigations by Tu and Schuler [64] led to the conclusion that the role of GAOs in full scale EBPR systems had been overestimated. Most of the previous microbial characterizations had been conducted in laboratory or pilot scale, where a common practice was application of the enormous concentrations of acetate, which favoured the growth of GAO. Similar suppositions have been formulated by Nielsen et al. [4]. The occurrence of GAOs was related to the excess of available carbon, which does not normally occur in typical EBPR system.

Configurations of the EBPR Systems

In general, EBPR systems can be categorized as main- and side stream configurations. The common feature of conventional side-stream configurations was sole treatment of return sludge anaerobically combined with chemical precipitation, whereas in mainstream configurations all mixed liquors flow through a sequence of anaerobic, anoxic and/or aerobic conditions for P removal [1].

Modern mainstream EBPR systems by design, are meant to avail conditions that sustain the parallel processes of N and P removal. Achieving optimal operating conditions for the biological processes, such as phosphorus release and uptake, nitrification and denitrification, requires consideration of specific environmental conditions within anaerobic, aerobic and anoxic zones. Due to the limited resources and stricter operational regulations, continuous modernization and revaluation of the BNR configurations has been observed [65].

A successful EBPR process is dependent on the presence of readily biodegradable organic carbon and phosphorus, anaerobic zone prior to aerobic zone and sufficient amount of nutrients since it relies on growth and selection of PAOs which are capable of storing orthophosphate in excess.

Earlier configurations achieved P removal with >90% efficiency [19,20,66]. By changing the sequential steps in the specific configuration, a low P removal efficiency was achieved, leading to an understanding of necessary conditions of EBPR [2]. The availability of COD favours heterotrophic activities (PAO, denitrifiers) under anaerobic/anoxic conditions, whereas nitrification takes place in the aerobic zone. P removal depends on several conditions that are essential for microbial metabolism of PAO [67]. However, simultaneous N and P removal are not straightforward as the addition of the anaerobic zone, which favours the PAO growth only [68]. In most systems, nitrification and denitrification may cause detrimental impact on EBPR due to the presence of nitrite and nitrate in the external recycle stream which enters the anaerobic zone, leading to a process failure. The presence of electron acceptors, such as nitrate and nitrite, under anaerobic conditions potentially sparks heterotrophic denitrifying bacteria growth and outcompete PAO [69]. Moreover, as indicated by Conidi et al. [70], the most common cause of instability in EBPR systems is the underestimated size of anaerobic zones, often less than 10% by mass of solids, whereas 15–25% is recommended for a stable operation.

Currently, the main direction towards increasing EBPR stability is the implementation of the novel sidestream EBPR (S2EBPR) configurations. With respect to the existing bioreactor configurations, providing a side stream fermentation zone (S2EBPR reactor) of recirculated activated sludge (RAS) or mixed liquor fermentation have emerged as a perspective solution to solve this issue. The main potential advantages offered by these solutions are increased anaerobic mass fraction and potential of the selective GAO suppression [71].

Due to P metabolism combined with the capability of fermenting a wide range of organic compounds and denitrification with simultaneous P uptake, *Tetrasphaera* are considered the key players of the S2EBPR systems. However, it should be noted that currently available data from the S2EBPR operations are limited, basically due to a narrow range of the operational conditions studied to date. As mentioned by Dold and Conidi [72], more extensive data are highly required to develop models specifically dedicated for S2EBPR systems.

The most typical configurations of the novel S2EBPR systems are summarized in Figure 5. A more comprehensive description of the mainstream systems can be found elsewhere (e.g., [48]), whereas examples of S2EBPR configurations have been presented by Dold and Conidi [72] or Gu et al. [73].
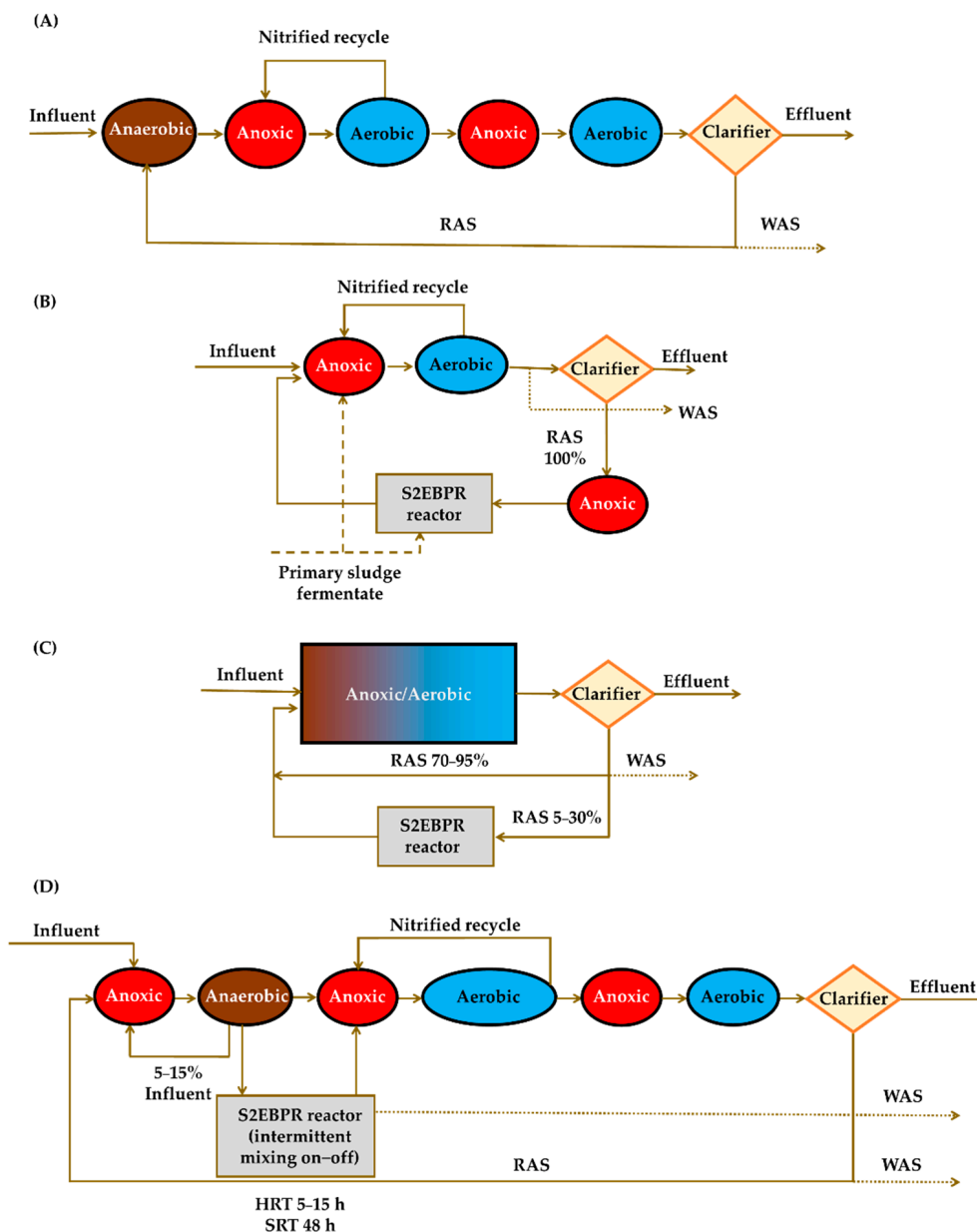
Figure 5. Examples of the mainstream EBPR system configurations ((**A**)—Bardenpho) and sidestream application ((**B**)—sidestream RAS plus carbon [SSRC], (**C**)—sidestream RAS [SSR], (**D**)—Sidestream Mixed Liquor Suspended Solids [SSM]). RAS—recirculated activated sludge, WAS—waste activated sludge, HRT—hydraulic retention time, SRT—sludge retention time.

6. Factors Affecting the Occurrence and Activity of *Tetrasphaera* and Other DPAOs in EBPR Systems

The recognition of dependencies between activity and occurrence of *Tetrasphaera* in relation to the changes of the environmental and operational conditions (pH, temperature, influent wastewater characteristics and DO or nitrate presence), as well interactions with other functional bacterial groups (especially GAO) are crucial for the development and optimization of EBPR [60,74–76]. However, in practice, it is not possible to achieve stable and fully controlled operational parameters, such as temperature and pH [77]. In general, other factors, such as the availability of the readily biodegradable organic compounds, P load, cation (specifically K^+ and M^{2+}) concentration, pH, and the food-to-microorganism ratio should be considered [16].

6.1. Temperature

Temperature is known to affect the operational efficiency of EBPR systems [78]. A study on psychrophilic nature of PAOs indicated their functionality at the temperatures around 20 °C or lower [4]. The increasing temperatures between 5 and 24 °C, improve the process efficiency. At elevated temperatures, the mesophilic GAOs seem to outcompete PAOs in terms of the carbon sources uptake, so the process efficiency becomes problematic at tropical temperatures of (25–32 °C) [4,27,31,57,77].

Other studies also draw attention to the deterioration of the efficiency of the EBPR process at high temperatures (above 20 °C) [51,79–81]. This indicates that the threshold temperature for tropical EBPR failure ranges from 35 to 40 °C, and above a significant deterioration in the PAO activity is observed. In addition, Panswad et al. [79] found that under elevated temperature, the competition between GAOs and PAOs is believed to enhance in terms of uptake of the available VFAs under anaerobic conditions. There is a substantial effect of temperature on lowering the relative abundance of PAOs with the increasing operating temperature from 20 to 35 °C. According to Whang and Park [81] and Lopez -Vazquez et al., [51], GAOs had faster VFA uptake rates at higher temperatures, leading to their excessive proliferation in the system. *Ca.* Accumulibacter and Tetrasphaera reflect higher abundances in the systems operated below 20 °C [27,51,82], whereas *Ca.* Competibacter become more numerous at higher temperatures. However, studies by Qiu et al. [83] and Wang et al. [84] showed the possibility of achieving highly efficient EBPR at the temperatures ranging from 28 to 32 °C.

The effect of temperature on the occurrence of *Tetrasphaera* was studied by Liu et al. [14] with the correlation analysis of experimental data from over 60 references. The results of the analysis revealed that the highest abundances of *Tetrasphaera* were mainly recorded in the temperature range from 10 to 20 °C. In contrast, in the countries with the warm climate and temperatures ranging from 20 to 30 °C, such as Australia or Spain, the abundance of *Tetrasphaera* was significantly lower. This finding is not fully consistent with the characteristics of pure *Tetrasphaera* cultures, where the temperature of 25 °C was considered suboptimal [32,33]. The lowered occurrence of *Tetrasphaera* at elevated temperatures, should therefore be explained by outcompetition from other microbial groups, in particular GAOs.

6.2. Influent Wastewater Characteristics

Influent wastewater characteristics have the potential to affect the abundance of *Tetrasphaera* and alter the metabolic pathways. The availability of preferred carbon sources and electron acceptors are essential to promote P-removal physiology of *Tetrasphaera*. As pointed out by Kong et al. [43], industrial WWTPs reflect usually higher abundances of *Tetrasphaera* compared to the systems treating mainly domestic wastewater. Mielczarek et al. [85] studied the correlation of abundance of *Tetrasphaera* and wastewater characteristics, process design, and operation, with the data of over 3 years from 28 Danish WWTPs. Their results showed weak correlations with the increased amount of industrial wastewater in the influent. As reported by Lopez-Vazquez et al., [51], the diversity of carbon compounds in the influent (the ratios of acetate to propionate of 75–25% and 50–50%) provided more favorable conditions for PAO than GAO, despite the elevated temperature (30 °C) in contrary to the experimental trials when single carbon source was used.

The capability to metabolize and internally store many carbon compounds provides *Tetrasphaera* with enormous adaptability to the dynamic conditions of substrate availability in WWTPs [52]. Such physiological plasticity may gain *Tetrasphaera* advantage over *Ca.* Accumulibacter, which is highly dependent on the acetate availability under anaerobic conditions [56].

6.3. pH

It is considered that, pH set point and its control are critical for enhancing typical PAO and *Tetrasphaera* activity in P removal [86,87].

Experimental results showed that P removal linearly increased with initial pH increasing from 6.6 to 7.8, but slightly decreased when initial pH increased from 7.8 to 8.2. Initial optimal pH of 7.8 favored P removal offering approximately 1.7 times compared to the pH at 6.6. Additionally, the modeling studies have showed that the biomass cultured at initial pH 7.8 contained elevated abundances of PAOs [88]. A recent study by Kang et al. [89] showed that the significant effect of pH control at a pH greater than or equal to 7.5 improved the P removal efficiency from 90.8% to 99.6%, whereas a pH below 7.0 ensured only about 63.1%.

The results of technological studies cover expectations from the characteristics of cultivable *Tetrasphaera* strains, for which the pH range for growth is 6.0–9.0 with the optimum at 7.0 [32]. Filipe et al. [89] observed the improved phosphate removal in the studied EBPR system when the pH was allowed to increase to a maximum of 7.5.

Another critical issue related to the pH influence is the competition between PAOs and GAOs, where pH is considered to affect anaerobic uptake kinetics of organic carbon. Several studies confirmed that pH in the range of 7–8.5 leads to an increased abundance of PAOs, whereas GAOs were inhibited or their proliferation rate was reduced under those conditions [27,51,56]. There is now strong evidence that the stability of EBPR systems can be improved by increasing the pH in the anaerobic zone. This allows for creating conditions under which PAOs are able to uptake acetate faster than GAOs, which leads to the positive shift in the composition of activated sludge biomass.

On the other hand, researchers seek clarity on how alkaline conditions can inhibit GAO proliferation and induce PAO activity [10]. The main explanation of this observation is that metabolism of the bacterial cells and enzymes activity is highly dependent on pH, thus under unfavourable pH more energy is needed for the substrate uptake and P release. It is postulated that carbon source uptake and phosphorus release/uptake are provided by proton motive force (PMF), strongly dependent on the extracellular pH and related to cation release [45,90].

In the study of Schuler and Jenkins [56], it was found that pH significantly affects the anaerobic phase of EBPR performed by *Ca. Accumulibacter*. However, different correlations were observed between pH and acetate uptake rates as well as between pH and anaerobic P release rates. For acetate uptake rates, a continuous positive correlation was found with the increasing pH, but the increase in acetate uptake slowed down above pH 7.2. On the contrary, in the case of P release rate, a plateau was gained at pH 7.2 and above. This suggests that at elevated pH, the energetic cost of acetate transport could not be compensated by phosphate lysis. Based on the behaviour of cations, correlated with pH during anaerobic carbon uptake, Saunders [91] postulated a secondary transport model as the dominant mechanism of EBPR by PAOs. In that model, acetate was co-transported with cations (typically H^+), which started to accumulate inside the bacterial cell and decrease PMF, thus slowing down the P release rate.

A recent study by Belka [92] showed that similar patterns were observed for pH-dependent anaerobic P release by *Tetrasphaera* as in previous studies with *Ca. Accumulibacter*. However, the anaerobic carbon uptake by *Tetrasphaera* was not correlated with pH and did not slow down at pH higher than 7.2. Due to more versatile metabolic properties (i.e., capability of accumulation of free organic carbon solutes and anaerobic amino acids uptake), *Tetrasphaera* could gain an important advantage over *Ca. Accumulibacter* under highly dynamic feeding and pH conditions.

6.4. Presence of DO and Nitrate

The presence of nitrate and DO in the anaerobic zone are considered inhibitors of the PAO activity. When the anaerobic zone is free of nitrate and DO, the PAO activity is favoured over any other rapidly growing heterotrophs that may use DO or nitrate as electron acceptors. The DO concentration of DO has been found to strongly affect EBPR performance and PAO dominance. Izadi et al. [10] achieved efficient P-removal at low DO concentrations (in the range of 0.5 or 0.8 mg/L), which was attributed to the favoured growth of PAOs compared to

GAOs. The presence of nitrate and/or DO reduces the amount of VFAs available for the PAO activity, while hampering P removal [93]. At low DO levels, PAOs have an advantage over *Ca.* Competibacter due to a higher DO affinity. Excessive aeration may induce instable P-removal as a result of GAO competition. Chen et al. [94] found that maintaining the DO level at 0.5 mgO$_2$/L promoted a higher efficiency EBPR.

Regardless of the dynamic conditions associated with the transition between anaerobic, anoxic and aerobic phases, *Tetrasphaera* representatives were found to be the dominant genus in most of the European WWTPs [34]. Herbst et al. [52] provided important insights into the characteristics of metabolic traits, which favored *Tetrasphaera* over most other functional bacterial groups under such dynamic conditions. By the use of label-free quantitative proteomics and nuclear magnetic resonance (NMR), the physiology of *T. longate* str. LP2 isolate was verified under dynamic shifts between anaerobic and aerobic conditions. Unlike the reference bacterial strain of *Escherichia coli*, *Tetrasphaera* reflected a stable proteome profile during transitions, which suggested that specific metabolic pathways for anaerobic processes remained induced under aerobic conditions. This metabolic property ensures a physiological advantage of *Tetrasphaera* over more specialized microbial groups. Moreover, it is suggested that by enlarging the anaerobic zone in EBPR systems, with a reduced oxidation-reduction potential (ORP), *Tetrasphaera* may gain a completive advantage over other heterotrophic bacteria [14].

7. Conclusions

The implementation of P-removal technologies based on application of the anaerobic–anoxic conditions has been recognized as the perspective approach for the energy efficient wastewater treatment. In such systems, the development of *Tetrasphaera* population and control of its activity may become a critical step. On the contrary, EBPR systems dominated by *Ca.* Accumulibacter, which are strongly dependent on acetate availability, are considered more susceptible to failure under dynamic conditions in WWTPs.

The development of *Tetrasphaera*-based EPBR systems is strongly determined by understanding the operational conditions, available carbon source and the presence of electron acceptors in the anaerobic zone. In addition, interactions with other functional bacterial groups (especially GAOs and other PAOs) have to be understood for further EBPR development and optimization.

The availability of complex carbon sources in the influent, moderate temperature (10–20 °C), elevated pH (>7.5) and increased sizes of the anaerobic zones are currently recognized as the possible main factors favouring *Tetrasphaera* abundance over typical PAOs (*Ca.* Accumulibacter). The conclusiveness of operational conditions such as pH and temperature specifically to promote the growth of Tetrasphaera over Accumulibacter is not definite at present. The known advantages exhibited by *Tetrasphaera* therefore, are attributable to versatile metabolism in line with their denitrification, fermentation and polyphosphate accumulation ability. Due to these versatile metabolic properties, *Tetrasphaera* are considered the key microorganisms for the novel S2EBPR systems.

However, there are still issues that need to be resolved in the future, including a phylogeny of *Tetrasphaera* genus, more complex characterization of metabolic traits of individual *Tetrasphaera* species, as well as kinetic studies based on *Tetrasphaera*-enriched cultures from full-scale systems.

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References

1. Metcalf and Eddy Inc. *Wastewater Engineering Treatment and Reuse*, 5th ed.; McGraw-Hill: New York, NY, USA, 2014.
2. Barnard, J.L.; Dunlap, P.; Steichen, M. Rethinking the Mechanisms of Biological Phosphorus Removal. *Water Environ. Res.* **2017**, *89*, 2043–2054. [CrossRef]
3. Nielsen, P.H.; Saunders, A.M.; Hansen, A.A.; Larsen, P.; Nielsen, J.L. Microbial communities involved in enhanced biological phosphorus removal from wastewater—A model system in environmental biotechnology. *Curr. Opin. Biotechnol.* **2012**, *23*, 452–459. [CrossRef] [PubMed]
4. Nielsen, P.H.; McIlroy, S.J.; Albertsen, M.; Nierychlo, M. Re-evaluating the microbiology of the enhanced biological phosphorus removal process. *Curr. Opin. Biotechnol.* **2019**, *57*, 111–118. [CrossRef] [PubMed]
5. Rubio-Rincón, F.; Lopez-Vazquez, C.; Welles, L.; van Loosdrecht, M.; Brdjanovic, D. Cooperation between Candidatus Competibacter and Candidatus Accumulibacter clade I, in denitrification and phosphate removal processes. *Water Res.* **2017**, *120*, 156–164. [CrossRef]
6. Marques, R.; Ribera-Guardia, A.; Santos, J.; Carvalho, G.; Reis, M.A.; Pijuan, M.; Oehmen, A. Denitrifying capabilities of Tetrasphaera and their contribution towards nitrous oxide production in enhanced biological phosphorus removal processes. *Water Res.* **2018**, *137*, 262–272. [CrossRef]
7. Fernando, E.Y.; McIlroy, S.J.; Nierychlo, M.; Herbst, F.-A.; Petriglieri, F.; Schmid, M.C.; Wagner, M.; Nielsen, J.L.; Nielsen, P.H. Resolving the individual contribution of key microbial populations to enhanced biological phosphorus removal with Raman–FISH. *ISME J.* **2019**, *13*, 1933–1946. [CrossRef] [PubMed]
8. Zhang, Y.; Kinyua, M.N. Identification and classification of the Tetrasphaera genus in enhanced biological phosphorus removal process: A review. *Rev. Environ. Sci. Technol.* **2020**, *19*, 699–715. [CrossRef]
9. Cydzik-Kwiatkowska, A.; Zielińska, M. Bacterial communities in full-scale wastewater treatment systems. *World J. Microbiol. Biotechnol.* **2016**, *32*, 66. [CrossRef] [PubMed]
10. Izadi, P.; Eldyasti, A. Understanding microbial shift of Enhanced Biological Phosphorus Removal process (EBPR) under different Dissolved Oxygen (DO) concentrations and Hydraulic Retention Time (HRTs). *Biochem. Eng. J.* **2021**, *166*, 107833. [CrossRef]
11. Kristiansen, R.; Nguyen, H.T.T.; Saunders, A.M.; Nielsen, J.L.; Wimmer, R.; Le, V.Q.; McIlroy, S.J.; Petrovski, S.; Seviour, R.J.; Calteau, A.; et al. A metabolic model for members of the genus Tetrasphaera involved in enhanced biological phosphorus removal. *ISME J.* **2013**, *7*, 543–554. [CrossRef] [PubMed]
12. Nguyen, H.T.T.; Kristiansen, R.; Vestergaard, M.; Wimmer, R.; Nielsen, P.H. Intracellular Accumulation of Glycine in Polyphosphate-Accumulating Organisms in Activated Sludge, a Novel Storage Mechanism under Dynamic Anaerobic-Aerobic Conditions. *Appl. Environ. Microbiol.* **2015**, *81*, 4809–4818. [CrossRef]
13. Roy, S.; Guanglei, Q.; Zuniga-Montanez, R.; Williams, R.B.; Wuertz, S. Recent advances in understanding the ecophysiology of enhanced biological phosphorus removal. *Curr. Opin. Biotechnol.* **2021**, *67*, 166–174. [CrossRef]
14. Liu, R.; Hao, X.; Chen, Q.; Li, J. Research advances of Tetrasphaera in enhanced biological phosphorus removal: A review. *Water Res.* **2019**, *166*, 115003. [CrossRef] [PubMed]
15. Welles, L.; Tian, W.; Saad, S.; Abbas, B.; Lopez-Vazquez, C.; Hooijmans, C.; van Loosdrecht, M.; Brdjanovic, D. Accumulibacter clades Type I and II performing kinetically different glycogen-accumulating organisms metabolisms for anaerobic substrate uptake. *Water Res.* **2015**, *83*, 354–366. [CrossRef]
16. Wisniewski, K.; Kowalski, M.; Makinia, J. Modeling nitrous oxide production by a denitrifying-enhanced biologically phosphorus removing (EBPR) activated sludge in the presence of different carbon sources and electron acceptors. *Water Res.* **2018**, *142*, 55–64. [CrossRef] [PubMed]
17. Levin, G.V.; Shapiro, J. Metabolic Uptake of Phosphorus by Wastewater Organisms. *Water Pollut. Control. Fed.* **1965**, *37*, 800–821.
18. Srinath, E.G.; Sastry, C.A.; Pillai, S.C. Rapid removal of phosphorus from sewage by activated sludge. *Experientia* **1959**, *15*, 339–340. [CrossRef]
19. Barnard, J.L. Cut P and N without chemicals. *Water Wastes Eng. Part 1* **1974**, *11*, 33–36.
20. Barnard, J.L. Cut P and N without chemicals. *Water Wastes Eng. Part 2* **1974**, *11*, 41–43.
21. Barnard, J.L. Nutrient removal in biological systems. *Water Pollut. Control* **1975**, *74*, 143–154.
22. Levin, G.V.; Topol, G.J.; Tarnay, A.G. Operation of full-scale biological phosphorus removal plant. *Water Pollut. Control Fed.* **1975**, *47*, 577–590.
23. Fuhs, G.W.; Chen, M. Microbiological basis of phosphate removal in the activated sludge process for the treatment of wastewater. *Microb. Ecol.* **1975**, *2*, 119–138. [CrossRef] [PubMed]
24. Wagner, M.; Loy, A.; Nogueira, R.; Purkhold, U.; Valeeva, A.V.; Daims, H. Microbial community composition and function in wastewater treatment plants. *Antonie Leeuwenhoek* **2002**, *81*, 665–680. [CrossRef]
25. Bond, P.L.; Hugenholtz, P.; Keller, J.; Blackall, L.L. Bacterial community structures of phosphate-removing and non-phosphate-removing activated sludges from sequencing batch reactors. *Appl. Environ. Microbiol.* **1995**, *61*, 1910–1916. [CrossRef]

26. Bond, P.L.; Keller, J.; Blackall, L.L. Characterisation of enhanced biological phosphorus removal activated sludges with dissimilar phosphorus removal performances. *Water Sci. Technol.* **1998**, *37*, 567–571. [CrossRef]

27. Oehmen, A.; Lemos, P.C.; Carvalho, G.; Yuan, Z.; Keller, J.; Blackall, L.L.; Reis, M.A. Advances in enhanced biological phosphorus removal: From micro to macro scale. *Water Res.* **2007**, *41*, 2271–2300. [CrossRef] [PubMed]

28. McIlroy, S.J.; Saunders, A.; Albertsen, M.; Nierychlo, M.; McIlroy, B.; Hansen, A.A.; Karst, S.M.; Nielsen, J.L.; Nielsen, P.H. MiDAS: The field guide to the microbes of activated sludge. *Database* **2015**, *2015*, bav062. [CrossRef]

29. Henze, M.; Gujer, W.; Mino, T.; van Loosdrecht, M. *Activated Sludge Models, ASM1, ASM2, ASM2d and ASM3. Scientific and Technical Report (Volume 5)*; IWA Publishing: London, UK, 2000. [CrossRef]

30. Jenkins, D.; Wanner, J. *Activated Sludge—100 Years and Counting*; IWA Publishing: Glasgow, UK, 2014. [CrossRef]

31. Bertanza, G.; Menoni, L.; Capoferri, G.U.; Pedrazzani, R. Promoting biological phosphorus removal in a full scale pre-denitrification wastewater treatment plant. *J. Environ. Manag.* **2020**, *254*, 109803. [CrossRef]

32. Hanada, S.; Liu, W.-T.; Shintani, T.; Kamagata, Y.; Nakamura, K. Tetrasphaera elongata sp. nov., a polyphosphate-accumulating bacterium isolated from activated sludge. *Int. J. Syst. Evol. Microbiol.* **2002**, *52*, 883–887. [CrossRef]

33. Maszenan, A.; Seviour, R.; Patel, B.; Schumann, P.; Burghardt, J.; Tokiwa, Y.; Stratton, H. Three isolates of novel polyphosphate-accumulating gram-positive cocci, obtained from activated sludge, belong to a new genus, *Tetrasphaera* gen. nov., and description of two new species, *Tetrasphaera japonica* sp. nov. and *Tetrasphaera australiensis* sp. nov. *Int. J. Syst. Evol. Microbiol.* **2000**, *50*, 593–603. [CrossRef]

34. Stokholm-Bjerregaard, M.; McIlroy, S.J.; Nierychlo, M.; Karst, S.M.; Albertsen, M.; Nielsen, P.H. A Critical Assessment of the Microorganisms Proposed to be Important to Enhanced Biological Phosphorus Removal in Full-Scale Wastewater Treatment Systems. *Front. Microbiol.* **2017**, *8*, 718. [CrossRef]

35. Seviour, R.J.; Mino, T.; Onuki, M. The microbiology of biological phosphorus removal in activated sludge systems. *FEMS Microbiol. Rev.* **2003**, *27*, 99–127. [CrossRef]

36. Seviour, R.J.; McIlroy, S.J. The microbiology of phosphorus removal in activated sludge processes-the current state of play. *J. Microbiol.* **2008**, *46*, 115–124. [CrossRef] [PubMed]

37. Petriglieri, F.; Singleton, C.; Peces, M.; Petersen, J.F.; Nierychlo, M.; Nielsen, P.H. "Candidatus Dechloromonas phosphatis" and "Candidatus Dechloromonas phosphovora", two novel polyphosphate accumulating organisms abundant in wastewater treatment systems. *BioRxiv* **2020**. [CrossRef]

38. Nguyen, H.T.T.; Le-Quy, V.; Hansen, A.A.; Nielsen, J.L.; Nielsen, P.H. High diversity and abundance of putative polyphosphate-accumulating Tetrasphaera-related bacteria in activated sludge systems. *FEMS Microbiol. Ecol.* **2011**, *76*, 256–267. [CrossRef]

39. Marques, R.; Santos, J.; Nguyen, H.; Carvalho, G.; Noronha, J.; Nielsen, P.H.; Reis, M.A.; Oehmen, A. Metabolism and ecological niche of Tetrasphaera and Ca. Accumulibacter in enhanced biological phosphorus removal. *Water Res.* **2017**, *122*, 159–171. [CrossRef] [PubMed]

40. Singleton, C.M.; Petriglieri, F.; Wasmund, K.; Nierychlo, M.; Kondrotaite, Z.; Petersen, J.F.; Peces, M.; Dueholm, M.S.; Wagner, M.; Nielsen, P.H. The novel genus, 'Candidatus Phosphoribacter', previously identified as Tetrasphaera, is the dominant polyphosphate accumulating lineage in EBPR wastewater treatment plants worldwide. *ISME J.* **2022**, *16*, 1605–1616. [CrossRef]

41. Nouioui, I.; Carro, L.; García-López, M.; Meier-Kolthoff, J.P.; Woyke, T.; Kyrpides, N.C.; Pukall, R.; Klenk, H.-P.; Goodfellow, M.; Göker, M. Genome-Based Taxonomic Classification of the Phylum Actinobacteria. *Front. Microbiol.* **2018**, *9*, 2007. [CrossRef] [PubMed]

42. Close, K.; Marques, R.; Carvalho, V.C.; Freitas, E.B.; Reis, M.A.; Carvalho, G.; Oehmen, A. The storage compounds associated with Tetrasphaera PAO metabolism and the relationship between diversity and P removal. *Water Res.* **2021**, *204*, 117621. [CrossRef]

43. Kong, Y.; Nielsen, J.L.; Nielsen, P.H. Identity and Ecophysiology of Uncultured Actinobacterial Polyphosphate-Accumulating Organisms in Full-Scale Enhanced Biological Phosphorus Removal Plants. *Appl. Environ. Microbiol.* **2005**, *71*, 4076–4085. [CrossRef]

44. Mino, T.; Arun, V.; Tsuzuki, Y.; Matsuo, T. Effect of phosphorus accumulation on acetate metabolism in the biological phosphorus removal process. In *Biological Phosphate Removal from Wastewaters, Proceedings of an IAWPRC Specialized Conference held in Rome, Italy, 28–30 September 1987*; Ramadori, R., Ed.; Elsevier: Amsterdam, The Netherlands, 1987; pp. 27–38. [CrossRef]

45. Smolders, G.J.F.; van der Meij, J.; van Loosdrecht, M.C.M.; Heijnen, J.J. Model of the anaerobic metabolism of the biological phosphorus removal process: Stoichiometry and pH influence. *Biotechnol. Bioeng.* **1994**, *43*, 461–470. [CrossRef] [PubMed]

46. Filipe, C.D.M.; Daigger, G.T.; Grady, C.P.L. A metabolic model for acetate uptake under anaerobic conditions by glycogen accumulating organisms: Stoichiometry, kinetics, and the effect of pH. *Biotechnol. Bioeng.* **2001**, *76*, 17–31. [CrossRef]

47. Vlekke, G.; Comeau, Y.; Oldham, W. Biological phosphate removal from wastewater with oxygen or nitrate in sequencing batch reactors. *Environ. Technol. Lett.* **1988**, *9*, 791–796. [CrossRef]

48. Mąkinia, J.; Zaborowska, E. *Mathematical Modelling and Computer Simulation of Activated Sludge Systems*; International Water Association Publishing: London, UK, 2020. [CrossRef]

49. Izadi, P.; Eldyasti, A. A review of biochemical diversity and metabolic modeling of EBPR process under specific environmental conditions and carbon source availability. *J. Environ. Manag.* **2021**, *288*, 112362. [CrossRef]

50. Oehmen, A.; Carvalho, G.; Lopez-Vazquez, C.; van Loosdrecht, M.; Reis, M. Incorporating microbial ecology into the metabolic modelling of polyphosphate accumulating organisms and glycogen accumulating organisms. *Water Res.* **2010**, *44*, 4992–5004. [CrossRef]

51. Lopez-Vazquez, C.M.; Oehmen, A.; Hooijmans, C.M.; Brdjanovic, D.; Gijzen, H.J.; Yuan, Z.; van Loosdrecht, M.C. Modeling the PAO–GAO competition: Effects of carbon source, pH and temperature. *Water Res.* **2009**, *43*, 450–462. [CrossRef]

52. Herbst, F.-A.; Dueholm, M.S.; Wimmer, R.; Nielsen, P.H. The Proteome of Tetrasphaera elongata is adapted to Changing Conditions in Wastewater Treatment Plants. *Proteomes* **2019**, *7*, 16. [CrossRef]

53. Meinhold, J.; Filipe, C.D.; Daigger, G.T.; Isaacs, S.H. Characterization of the denitrifying fraction of phosphate accumulating organisms in biological phosphate removal. *Water Sci. Technol.* **1999**, *39*, 31–42. [CrossRef]

54. Mino, T.; van Loosdrecht, M.; Heijnen, J. Microbiology and biochemistry of the enhanced biological phosphate removal process. *Water Res.* **1998**, *32*, 3193–3207. [CrossRef]

55. Meng, Q.; Zeng, W.; Wang, B.; Fan, Z.; Peng, Y. New insights in the competition of polyphosphate-accumulating organisms and glycogen-accumulating organisms under glycogen accumulating metabolism with trace Poly-P using flow cytometry. *Chem. Eng. J.* **2020**, *385*, 123915. [CrossRef]

56. Schuler, A.J.; Jenkins, D. Enhanced biological phosphorus removal from wastewater by biomass with different phosphorus contents, Part I: Experimental results and comparison with metabolic models. *Water Environ. Res.* **2003**, *75*, 485–498. [CrossRef]

57. Winkler, M.-K.; Bassin, J.; Kleerebezem, R.; de Bruin, L.; Brand, T.V.D.; van Loosdrecht, M. Selective sludge removal in a segregated aerobic granular biomass system as a strategy to control PAO–GAO competition at high temperatures. *Water Res.* **2011**, *45*, 3291–3299. [CrossRef]

58. Wong, M.-T.; Tan, F.M.; Ng, W.J.; Liu, W.-T. Identification and occurrence of tetrad-forming Alphaproteobacteria in anaerobic–aerobic activated sludge processes. *Microbiology* **2004**, *150*, 3741–3748. [CrossRef]

59. Carvalheira, M.; Oehmen, A.; Carvalho, G.; Reis, M.A. The effect of substrate competition on the metabolism of polyphosphate accumulating organisms (PAOs). *Water Res.* **2014**, *64*, 149–159. [CrossRef]

60. Onnis-Hayden, A.; Majed, N.; Li, Y.; Rahman, S.M.; Drury, D.; Risso, L.; Gu, A.Z. Impact of solid residence time (SRT) on functionally relevant microbial populations and performance in full-scale enhanced biological phosphorus removal (EBPR) systems. *Water Environ. Res.* **2020**, *92*, 389–402. [CrossRef] [PubMed]

61. Majed, N.; Gu, A.Z. Phenotypic dynamics in polyphosphate and glycogen accumulating organisms in response to varying influent C/P ratios in EBPR systems. *Sci. Total Environ.* **2020**, *743*, 140603. [CrossRef] [PubMed]

62. Shen, N.; Zhou, Y. Enhanced biological phosphorus removal with different carbon sources. *Appl. Microbiol. Biotechnol.* **2016**, *100*, 4735–4745. [CrossRef]

63. Oyserman, B.; Noguera, D.R.; del Rio, T.G.; Tringe, S.G.; McMahon, K.D. Metatranscriptomic insights on gene expression and regulatory controls in Candidatus Accumulibacter phosphatis. *ISME J.* **2016**, *10*, 810–822. [CrossRef] [PubMed]

64. Tu, Y.; Schuler, A.J. Low Acetate Concentrations Favor Polyphosphate-Accumulating Organisms over Glycogen-Accumulating Organisms in Enhanced Biological Phosphorus Removal from Wastewater. *Environ. Sci. Technol.* **2013**, *47*, 3816–3824. [CrossRef]

65. Izadi, P.; Eldyasti, A. Design, operation and technology configurations for enhanced biological phosphorus removal (EBPR) process: A review. *Rev. Environ. Sci. Technol.* **2020**, *19*, 561–593. [CrossRef]

66. Barnard, J.L. A review of biological phosphorus removal in the activated sludge process. *Water SA* **1976**, *2*, 136–144.

67. Alasino, N.; Mussati, M.C.; Scenna, N.; Aguirre, P. Combined nitrogen and phosphorus removal. Model-based process optimization. *Comput. Aided Chem. Eng.* **2008**, *25*, 163–168. [CrossRef]

68. Guerrero, J.; Guisasola, A.; Baeza, J.A. A novel control strategy for efficient biological phosphorus removal with carbon-limited wastewaters. *Water Sci. Technol.* **2014**, *70*, 691–697. [CrossRef] [PubMed]

69. Guerrero, J.; Flores-Alsina, X.; Guisasola, A.; Baeza, J.A.; Gernaey, K.V. Effect of nitrite, limited reactive settler and plant design configuration on the predicted performance of simultaneous C/N/P removal WWTPs. *Bioresour. Technol.* **2013**, *136*, 680–688. [CrossRef]

70. Conidi, D.; Andalib, M.; Andres, C.; Bye, C.; Umble, A.; Dold, P. Modeling quaternary ammonium compound inhibition of biological nutrient removal activated sludge. *Water Sci. Technol.* **2018**, *79*, 41–50. [CrossRef]

71. Onnis-Hayden, A.; Srinivasan, V.; Tooker, N.B.; Li, G.; Wang, D.; Barnard, J.L.; Bott, C.; Dombrowski, P.; Schauer, P.; Menniti, A.; et al. Survey of full-scale sidestream enhanced biological phosphorus removal (S2EBPR) systems and comparison with conventional EBPRs in North America: Process stability, kinetics, and microbial populations. *Water Environ. Res.* **2020**, *92*, 403–417. [CrossRef]

72. Dold, P.; Conidi, D. Achieving Enhanced Biological P Removal: Have we forgotten how to design a bioP plant? In Proceedings of the 92nd Annual Water Environment Federation Technical Exhibition and Conference [CD-ROM], Chicago, IL, USA, 21–25 September 2019; pp. 1452–1466.

73. Gu, A.Z.; Tooker, N.; Onnis-Hayden, A.; Wang, D.; Srinivasan, V.; Li, G.; Takács, I.; Vargas, E. *Optimization and Design of a Side-Stream EBPR Process as a Sustainable Approach for Achieving Stable and Efficient Phosphorus Removal*; Project No. U1R13/4869; WRF: Alexandria, VA, USA, 2019.

74. Ferrera, I.; Sánchez, O. Insights into microbial diversity in wastewater treatment systems: How far have we come? *Biotechnol. Adv.* **2016**, *34*, 790–802. [CrossRef]

75. Guo, G.; Wu, D.; Hao, T.; Mackey, H.R.; Wei, L.; Chen, G. Denitrifying sulfur conversion-associated EBPR: The effect of pH on anaerobic metabolism and performance. *Water Res.* **2017**, *123*, 687–695. [CrossRef]

76. Mulkerrins, D.; Dobson, A.; Colleran, E. Parameters affecting biological phosphate removal from wastewaters. *Environ. Int.* **2004**, *30*, 249–259. [CrossRef]

77. Gebremariam, S.Y.; Beutel, M.W.; Christian, D.; Hess, T.F. Research Advances and Challenges in the Microbiology of Enhanced Biological Phosphorus Removal—A Critical Review. *Water Environ. Res.* **2011**, *83*, 195–219. [CrossRef]

78. Jiang, L.; Wang, M.; Wang, Y.; Liu, F.; Qin, M.; Zhang, Y.; Zhou, W. The condition optimization and mechanism of aerobic phosphorus removal by marine bacterium Shewanella sp. *Chem. Eng. J.* **2018**, *345*, 611–620. [CrossRef]

79. Panswad, T.; Doungchai, A.; Anotai, J. Temperature effect on microbial community of enhanced biological phosphorus removal system. *Water Res.* **2003**, *37*, 409–415. [CrossRef]

80. Weissbrodt, D.G.; Neu, T.R.; Kuhlicke, U.; Rappaz, Y.; Holliger, C. Assessment of bacterial and structural dynamics in aerobic granular biofilms. *Front. Microbiol.* **2013**, *4*, 175. [CrossRef]

81. Whang, L.-M.; Park, J.K. Competition between Polyphosphate- and Glycogen-Accumulating Organisms in Enhanced-Biological-Phosphorus-Removal Systems: Effect of Temperature and Sludge Age. *Water Environ. Res.* **2006**, *78*, 4–11. [CrossRef] [PubMed]

82. Poh, P.K.; Ong, Y.H.; Arumugam, K.; Nittami, T.; Yeoh, H.K.; Bessarab, I.; William, R.; Chua, A.S.M. Tropical-based EBPR process: The long-term stability, microbial community and its response towards temperature stress. *Water Environ. Res.* **2021**, *93*, 2598–2608. [CrossRef] [PubMed]

83. Qiu, G.; Zuniga-Montanez, R.; Law, Y.; Thi, S.S.; Nguyen, T.Q.N.; Eganathan, K.; Liu, X.; Nielsen, P.H.; Williams, R.B.; Wuertz, S. Polyphosphate-accumulating organisms in full-scale tropical wastewater treatment plants use diverse carbon sources. *Water Res.* **2019**, *149*, 496–510. [CrossRef]

84. Wang, L.; Shen, N.; Oehmen, A.; Zhou, Y. The impact of temperature on the metabolism of volatile fatty acids by polyphosphate accumulating organisms (PAOs). *Environ. Res.* **2020**, *188*, 109729. [CrossRef] [PubMed]

85. Mielczarek, A.T.; Nguyen, H.T.; Nielsen, J.L.; Nielsen, P.H. Population dynamics of bacteria involved in enhanced biological phosphorus removal in Danish wastewater treatment plants. *Water Res.* **2013**, *47*, 1529–1544. [CrossRef]

86. Liau, K.F.; Shoji, T.; Ong, Y.H.; Chua, A.S.M.; Yeoh, H.K.; Ho, P.Y. Kinetic and stoichiometric characterization for efficient enhanced biological phosphorus removal (EBPR) process at high temperatures. *Bioprocess Biosyst. Eng.* **2015**, *38*, 729–737. [CrossRef]

87. Xu, Y.; Hu, H.; Liu, J.; Luo, J.; Qian, G.; Wang, A. pH dependent phosphorus release from waste activated sludge: Contributions of phosphorus speciation. *Chem. Eng. J.* **2015**, *267*, 260–265. [CrossRef]

88. Wang, D.; Tooker, N.B.; Srinivasan, V.; Li, G.; Fernandez, L.A.; Schauer, P.; Menniti, A.; Maher, C.; Bott, C.B.; Dombrowski, P.; et al. Side-stream enhanced biological phosphorus removal (S2EBPR) process improves system performance—A full-scale comparative study. *Water Res.* **2019**, *167*, 115109. [CrossRef] [PubMed]

89. Kang, A.J.; Munz, G.; Yuan, Q. Influence of pH control on material characteristics, bacterial community composition and BNR performance of mature aerobic granules. *Process Saf. Environ. Prot.* **2019**, *124*, 158–166. [CrossRef]

90. Filipe, C.D.; Daigger, G.T.; Grady, C.L. pH as a key factor in the competition between glycogen-accumulating organisms and phosphorus-accumulating organisms. *Water Environ. Res.* **2001**, *73*, 223–232. [CrossRef] [PubMed]

91. Saunders, A.M.; Mabbett, A.N.; McEwan, A.G.; Blackall, L.L. Proton motive force generation from stored polymers for the uptake of acetate under anaerobic conditions. *FEMS Microbiol. Lett.* **2007**, *274*, 245–251. [CrossRef] [PubMed]

92. Belka, D. The Effect of pH on Organic Carbon Uptake and Biological Phosphorus Removal by Tetrasphaera Polyphosphate Accumulating Organisms. 2021. Available online: https://digitalrepository.unm.edu/ce_etds/259 (accessed on 8 October 2022).

93. Zuthi, M.; Guo, W.; Ngo, H.; Nghiem, L.; Hai, F. Enhanced biological phosphorus removal and its modeling for the activated sludge and membrane bioreactor processes. *Bioresour. Technol.* **2013**, *139*, 363–374. [CrossRef]

94. Chen, H.-B.; Wang, D.-B.; Li, X.-M.; Yang, Q.; Luo, K.; Zeng, G.-M. Temperature influence on biological phosphorus removal induced by aerobic/extended-idle regime. *Environ. Sci. Pollut. Res.* **2014**, *21*, 6034–6043. [CrossRef]