Review

Characterisation of Phosphate Accumulating Organisms and Techniques for Polyphosphate Detection: A Review

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Abstract: Phosphate minerals have long been used for the production of phosphorus-based chemicals used in many economic sectors. However, these resources are not renewable and the natural phosphate stocks are decreasing. In this context, the research of new phosphate sources has become necessary. Many types of wastes contain non-negligible phosphate concentrations, such as wastewater. In wastewater treatment plants, phosphorus is eliminated by physicochemical and/or biological techniques. In this latter case, a specific microbiota, phosphate accumulating organisms (PAOs), accumulates phosphate as polyphosphate. This molecule can be considered as an alternative phosphate source, and is directly extracted from wastewater generated by human activities. This review focuses on the techniques which can be applied to enrich and try to isolate these PAOs, and to detect the presence of polyphosphate in microbial cells.

Keywords: PAO; polyphosphate; detection; single cell technologies

1. Introduction

Phosphorus is an essential element required for food production. It has been estimated that about 19.7 MT of phosphorus were extracted from phosphate rocks in 2000. About 15.3 MT were used to produce fertilizers out of this total production. The global growing population induces a progressive increase of fertilizers’ requirement, which means an increasing need for phosphate (P). In this context, the establishment of circular economy applicable to P resources appears obvious [1]. Europe does not have significant P mines and must import ores from the other continents. The total P-reserves have been estimated at 67,000 Tg, and Morocco owns 75% of these known deposits. The remaining percentage mainly belongs to China and the USA. This geopolitical context can generate unexpected increases of fertilizers’ prices, which occurred in 2008 [2]. It is, therefore, necessary to find new P sources. Some wastes contain recoverable P, such as wastewater. A growing attention has been paid to the research of new techniques to remove P from it. More specifically, enhanced biological phosphorus removal (EBPR) is currently applied in wastewater treatment plants (WWTPs). Sewage sludge resulting from wastewater treatment contains significant quantities of P. This is due to the presence of phosphate
accumulating organisms (PAOs) [3]. PAOs have been studied in many research papers. Under specific conditions, these microorganisms can store P as polyphosphate (Poly-P). This ability has been found in bacteria, microalgae, and fungi [3–5].

This article focuses on bacterial PAOs and investigates the difficulties encountered in isolation attempts. It presents an inventory of analytical techniques which can be used to detect the presence of PAOs and Poly-P.

2. Techniques for the Isolation of PAOs

The methods used to detect and identify the PAOs are mainly based on the detection of Poly-P and polyhydroxyalkanoate (PHA). However, the detection of Poly-P itself is much more reliable. Prior to the detection of the polymers which are specific of PAOs (mainly Poly-P), an enrichment step of microorganisms is required. The original samples are mostly collected in WWTPs, in which the metabolism of PAOs is used to remove free phosphate from wastewater by accumulating it as Poly-P [6]. However, it is important to note that the maximal efficiency of PAOs is far from being reached in common EBPR plants.

2.1. Isolation of PAOs

Poly-P is the main form of phosphate accumulation. However, other forms of phosphate storage can be encountered. These components were investigated in a previous review [7]. The P storage compounds can be mineral or organic and depend on the type of microorganism. Low-solubility orthophosphate compounds are the simplest forms of P-accumulation. Some archaea can precipitate MgPO$_4$·OH·4H$_2$O (such as Halobacterium salinarium and Halorubrum distributum). These strains concentrate P from aqueous solution during their growth phase. Brevibacteria can also accumulate P as intracellular NH$_4$MgPO$_4$·6H$_2$O. Finally, cyanobacteria accumulate P in their sheaths, combined with calcium. The second type of P storage compounds is organic. Teichoic acid is synthesized by bacteria and is composed of polyol and glycosyl polyol residues bound by phosphodiester bonds. It has been suggested that this molecule is a phosphate reserve. The yeast Kuraishia capsulata can also store phosphate in extracellular phosphomannan [7].

However, in most microorganisms, inorganic Poly-P is the main P reserve, composed of three to several hundred phosphate units. This molecule is a polymeric reserve which also plays a role in the regulation of enzymatic activities, the expression of many genes and the stress adaptation process [7]. Many studies report trials of isolation of PAOs. Acinetobacter was isolated from an EBPR plant, but subsequent studies using culture-independent techniques revealed that this bacterium had a small, but significant, effect in the process. Microlunatus phosphovorus was also isolated and proposed as a PAO, but did not show their characteristics (presence of PHA granules). Molecular techniques allowed to identify Candidatus Accumulibacter phosphatis (Rhodocyclus-related bacteria) as an important subclass involved in P removal at the lab scale, but it was never isolated in pure culture. This subclass is considered as the model showing all the characteristics of a typical bacterial PAO. It is important to note the diversity of this subclass, including two types (I and II), and seven clades in Type II (IIA, IIB, IIC, IID, IIE, IIF and IIG). A recent paper reported the existence of two new clades (IIH and III) and concluded that the configuration of WWTPs has a direct impact on the structure of PAO communities [8]. However, the attempts of isolation of true PAOs have not been successful so far [9].

Some screening techniques have been proposed. Morohoshi et al. [10] developed a method based on the detection of PAOs by the apparition of a blue color due to the hydrolysis of 5-bromo-4-chloro-3-indolyolphosphate on agar plates. Chaudhry and Nautiyal [11] also developed a visual technique based on the use of Toluidine blue-O dye in agar plates containing sodium citrate as the main carbon source. It is important to note that it has not been possible to isolate pure microorganisms seen as true PAOs so far, although many efforts have been made in this respect. This is due to the symbioses in microbial consortia, broken by the isolation techniques.
The isolation attempts have tested many enrichment media containing acetate and citrate as carbon sources [6,12–14]. Next, phosphate uptake assays must be achieved in order to confirm the P-accumulation ability of the strain isolated. At this level, the mechanisms of P-accumulation as Poly-P (aerobic/anoxic conditions) and P-release (anaerobic conditions) in bacteria have been described well. However, it must be kept in mind that many parameters affect the capacity of P-accumulation, such as time of anaerobic and aerobic/anoxic phases, pH, temperature, volatile fatty acid composition, \(\text{PO}_4^{3-}\), \(\text{SO}_4^{2-}\), \(\text{NH}_4^+\), \(\text{K}^+\), \(\text{Na}^+\), \(\text{Mg}^{2+}\), \(\text{Ca}^{2+}\), and other metal ion concentrations [11,12,15].

These mechanisms have a direct impact on isolation strategies. Firstly, an anaerobic step is necessary to remove Poly-P before undertaking phosphate uptake assays [6]. Secondly, the detection of Poly-P can be performed only after having grown the bacterial strains in aerobic conditions. Anaerobic phases can be considered as stress conditions for PAOs. They take up the carbon sources and store them to cope with a potentially long-term absence of oxygen (Figure 1). In wastewater, acetate is the main carbon source and is converted into Acetyl-CoA. This reaction requires energy provided by the hydrolysis of ATP, leading to ADP and the release of P in the medium. Acetyl-CoA is next metabolized into \(\beta\)-hydroxyalkanoates (PHAs), which are carbon storage compounds [9].

![Figure 1. Anaerobic metabolism of phosphorus in phosphate accumulating bacteria.](image)

In aerobic conditions, this trend is reversed. PHAs are used for cell growth and the reconstitution of the Poly-P reserves (Figure 2). They are degraded, which leads to the synthesis of Acetyl-CoA, processed through the TCA cycle. The TCA cycle produces energy from oxidation and carbon for new cell growth. A part of this energy is used to take up soluble P from the environment and to incorporate it into Poly-P. Another part of carbon and energy is used to regenerate glycogen [16].

The case of phosphate accumulating microalgae and mycetes has much less been investigated. However, it has been determined that microalgae accumulate P under stress conditions in specific organites: the acidocalcisomes [17].

The techniques applied to detect Poly-P in microorganisms are diversified and were reviewed in several articles [18–23]. Here, we propose a global overview of the methods which can be applied to the detection of PAOs and Poly-P.
2.2. Poly-P Detection and Identification of PAOs

The methods used to detect Poly-P in microbial cells are diversified. The technique used depends on the information required (detection, quantification, degree of polymerization (DP)). They can focus on the PAOs or the Poly-P molecule itself. The methods applicable are summarized in Table 1.

Table 1. Inventory of techniques applied to the detection of PAOs and Poly-P.

| Technique                                                                 | Information                                                                 |
|---------------------------------------------------------------------------|----------------------------------------------------------------------------|
| Light and fluorescence microscopy coupled with specific staining (LEM)     | Presence/absence of Poly-P                                                  |
| Flow cytometry (FC)                                                       | Location and quantification of Poly-P granules, cell sorting               |
| FISH analysis (FISH)                                                      | Detection of PAOs                                                          |
| Extraction procedures and phosphate quantification (EXT)                   | Quantification of Poly-P                                                    |
| Polyacrylamide gel electrophoresis (PAGE)                                 | Detection of Poly-P, determination of DP                                    |
| Electron microscopy (EM)                                                  | Presence/absence of Poly-P, location and composition of Poly-P granules    |
| X-ray analysis (X-RAY)                                                    | Composition of Poly-P granules, possible quantification                    |
| Nuclear Magnetic Resonance Spectroscopy (NMRS)                            | Detection of Poly-P, study of Poly-P structure                             |
| RAMAN microscopy (RAM)                                                    | Detection and quantification of Poly-P                                      |
| Enzyme assays (EA)                                                        | Detection and quantification of Poly-P                                      |
| Cryoelectron tomography and spectroscopic imaging (CTSI)                  | Detection of Poly-P, study of Poly-P structure                             |
| Mass spectrometry (MS)                                                    | Detection of Poly-P, study of Poly-P structure                             |
| Proteic affinity (PA)                                                     | Detection of Poly-P, location of Poly-P granules                           |
| “Omics techniques” (OMICS)                                               | Study of PAOs in complex communities                                       |

2.2.1. Staining Techniques—Light and Epifluorescence Microscopy (LEM)

The most common techniques consist in dying Poly-P with chemicals and observing the Poly-P granules with a light or fluorescence microscope. These qualitative methods were the first used
and are still being applied, owing to their simplicity. Loeffler’s methylene blue staining is based on the affinity of anionic Poly-P with methylene blue. The bonds between the dying molecules and Poly-P lead to a specific pink-violet color. Toluidine blue can be used instead of methylene blue because its metachromatic properties are identical. Neisser staining, the second common method, also uses methylene blue combined with a counterstaining solution (Bismark brown or Chrysoidin Y). This protocol improves the detection efficiency by increasing the contrast between Poly-P and the other cell components. Neisser staining protocol leads to a purple color for Poly-P, while the other cellular components are yellowish-brown. Regardless of what technique is used, methylene blue requires an acidic pH to improve the linkages with Poly-P [19]. Neutral red is a pH-sensitive dye, as well. It is used as a probe which detects acidic cellular compartments, such as the ones which contain Poly-P (under the form of “volutin” granules). The neutral form of neutral red is membrane-permeable, while the protonated form is not. Consequently, the acidic compartments tend to accumulate neutral red and can be observed through fluorescence microscopy. The accumulation can be detected on the basis of the absorbance shift: neutral red shows a $\lambda_{\text{max}}$ of 450 nm at pH 8.1 while its protonated form shows a $\lambda_{\text{max}}$ of 535 nm at pH 5.3 [24]. The case of 4′,6-diamidino-2-phenylindole (DAPI), another dying molecule, has been reported in many research articles. This dye stains Poly-P granules with a yellow fluorescence when used at high concentrations (see the part “Flow cytometry”). However, DAPI is not specific: it also stains other polymeric ions, such as DNA and lipids [20]. DAPI-DNA fluorescence is blue, but both DAPI-Poly-P and DAPI-lipids show a yellow fluorescence. The fluorescence of DAPI-lipids is shorter in time than the one of DAPI-Poly-P [25]. High DAPI concentrations induce a high fluorescence background, and lower DAPI concentrations coupled with a longer incubation time can be applied [19]. Due to its lack of specificity, the detection of Poly-P with DAPI can be performed after having checked that the reserves of PHA have been totally consumed. This storage polymer can be detected with Sudan black and Nile blue [19]. Other marginal dyes were mentioned in some articles, such as tetracycline (fluorescence based on the linkage with Poly-P) and Fura-2 (fluorescence depending on the competition of Mn$^{2+}$ affinity with Fura-2 and Poly-P) [26,27].

2.2.2. Flow Cytometry (FC)

Flow cytometry has been cited in many research articles relating to the analysis of Poly-P, mainly with DAPI. The staining reaction is exactly the same as in fluorescence microscopy, but flow cytometry brings much more data [28]. DAPI stains the cells with Poly-P contents higher than 400 µmol/g (dry weight) under a concentration of at least 18 µM (5 to 50 µg/mL). It is considered that a lower concentration in DAPI (0.24 to 5 µM) leads to a blue fluorescence related to bacterial DNA only, while the fluorescence due to Poly-P is yellow and induced by higher DAPI concentrations [27]. Flow cytometry allows the measuring and analysis of multiple physical characteristics, and the particles of interest can be separated from another to make further analyses through cell sorting [28]. Flow cytometry has already been used to detect Poly-P in bacterial cells but also in mammal cells [29]. This technique can be used to locate the Poly-P granules inside the living cells. Many research articles cited DAPI in the topic of Poly-P research [27–31]. Tetracycline can also be used to detect Poly-P through flow cytometry. This antibiotic and its derivatives are usually used in medicine as a labelling molecule for the assessment of calcium deposition. The association of diamagnetic divalent cations (Mg$^{2+}$, Ca$^{2+}$) with tetracycline leads to a green fluorescence (emission wavelength of 515 nm), with an excitation wavelength of 390 nm at a neutral pH [27]. This protocol was applied in several research articles [27,32,33].

2.2.3. Fluorescence in Situ Hybridization (FISH) Analysis

Fluorescence in situ hybridization (FISH) is rather linked to the microorganisms themselves and is based on the use of labelled probes which can detect DNA or RNA molecules in situ. This technique is applicable to the identification of microorganisms, the study of gene expression and the study of ecophysiology, which is particularly interesting in the case of PAOs [34]. However,
the technique suffers from disadvantages due to its specificity: the sequence of the target molecules and their abundance are required, and the probes must also penetrate the cells with a sufficient efficiency [28]. Albertsen et al. [35] used quantitative FISH with 32 oligonucleotide probes to investigate the composition of the bacterial communities in Danish EBPR plants. FISH analysis allows targeting of specific bacterial clades. Such probes were used to detect the presence of *Accumulibacter*, a well-known PAO [36,37]. It can also be used to determine the location of PAOs in complex samples [38]. FISH analysis may be combined with other techniques such as in the study of Liu et al. [39], which used DAPI staining, electron microscopy, and X-ray analysis as complementary techniques. The labelled cells can be observed through microscopy techniques [40], but flow cytometry is also applicable. This combination was used in the study of Kim et al. [41] to separate four *Accumulibacter* clades with Fluorescence-Activated Cell Sorting (FACS). Another possible combination is FISH/micro-autoradiography (MAR-FISH), which used labelled substrates (³H, ¹⁴C) to investigate the assimilation of nutrient sources according to the conditions used [42].

2.2.4. Extraction Procedures and Polyphosphate Quantification (EXT)

Many articles report extraction procedures of Poly-P followed by its quantification. This procedure is applied to microorganisms the ability to accumulate Poly-P of which has been confirmed. Several problems must be considered: inorganic Poly-P is usually combined with other macromolecules, such as nucleic acids and proteins; the Poly-P solubility depends on the chain length and the presence of cations; the cellular breakdown required to extract Poly-P may lead to undesired modifications, such as changes in chain length, concentrations, and linkage to inorganic molecules. The extraction procedures were reviewed before [20]. The extraction of Poly-P from living cells and the subsequent phosphate (resulting from Poly-P hydrolysis) measurement can be made through different techniques. An extraction by NaOH followed by an acidic hydrolysis with HCl was performed by Aravind et al. [12], before phosphate measurement by the method of molybdate reagent and ascorbic acid. Another research article reported the extraction of different Poly-P fractions (acid-soluble with HClO₄, salt-soluble with NaClO₄ and HClO₄, and alkali-soluble with NaOH) from *Saccharomyces cerevisiae*, before the digestion by HClO₄ [43,44]. Poly-P can also be extracted by the method of alkaline hypochlorite [11]. Jimenez-Nuñez et al. [29] used a digestion technique using RNase and DNase followed by an extraction with phenol-chloroform. Next, exopolyphosphatase was used to assess the Poly-P concentration. The extraction with boiling water was also tested, but this technique leads to a partial hydrolysis of Poly-P, which is not desired until the true hydrolysis step [20]. The analysis of phosphate resulting from the hydrolysis step can also be achieved by ion chromatography and inductively coupled plasma (ICP) atomic emission spectrometry (AES) [22,23].

2.2.5. Polyacrylamide Gel Electrophoresis (PAGE)

Gel electrophoresis can provide data about the DP of Poly-P and requires a Poly-P extraction procedure. The samples must be treated to obtain a soluble form of Poly-P. Cation-exchange resins can be used to reach these conditions, and the migration of standards allows to assess the size of Poly-P molecules. A 6%–30% polyacrylamide gel with 7 M urea was used in several research articles [29,43,44]. Next, the gels can be stained for visualization (Toluidine blue) [44]. Another publication reports the use of PAGE added with DAPI, which led to a yellow fluorescence, while nucleic acids and glycosaminoglycans fluoresced blue [45].

2.2.6. Electron Microscopy (EM)

Electron microscopy is a powerful technique which allows to visualize and locate the Poly-P granules inside cells. The technique used alone cannot quantify the concentrations in Poly-P, but confirms the presence of Poly-P granules. Scanning electron microscopy (SEM) is usually used with other techniques, such as staining protocols [22,23]. Scanning transmission electron microscopy (STEM) can be used to detect high-density granules, such as Poly-P [46]. Electron microscopes are usually
equipped with X-ray detectors, which can confirm the elementary composition of the material and the composition of Poly-P granules [43]. Another article reports a complete analysis of phosphate granules in fossil bacteria through several microscopy techniques (SEM, STEM, and Scanning Transmission X-ray Microscopy) [47]. Such a combination was also used in another work which studied the accumulation of Poly-P in Comamonadaceae through SEM combined with energy dispersive X-ray spectroscopy [48]. A metallization can be applied to the samples to improve the performance of electron microscopy, with osmium tetroxide as in a previous study [13]. Another article investigated the structure of precipitated phosphate minerals (struvite) produced by phosphate forming bacteria through SEM and field-emission scanning electron microscopy (FESEM) [49].

2.2.7. X-Ray Analysis Techniques (X-RAY)

X-ray analysis provides information about the composition of the samples and is usually combined with electron microscopes. Energy-dispersive X-ray microanalysis was previously used to assess the composition of Poly-P granules observed through electron microscopy (abbreviated by EM-EDX) [46]. Such combination was used in many studies [43,47,48]. However, the treatments applied before the observation stage must be limited to prevent P losses from the cells [20]. X-ray analyses highlighted the effect of redox conditions on different types of Poly-P: the polymers associated with Mg and K are more easily metabolized than the ones associated with Ca, which means that this latter type of phosphate storage is not influenced by the presence or absence of aerobic conditions [20]. Another useful combined technique is X-ray fluorescence spectromicroscopy, a combination of X-ray spectroscopy and X-ray microscopy. This method was found to be able to elucidate the spatial concentration and speciation of elements at submicron scale in minimally-prepared particulate samples including individual cells. It can analyze thicker samples than electron microscopy and is less restrictive during the preparation procedure [22,23]. The technique developed by Rivadeneyra et al. [49] used an X-ray diffractometer to study struvite crystals produced in culture media. This device allowed to determine the qualitative and quantitative composition of the salts produced by bacteria [49].

2.2.8. Nuclear Magnetic Resonance Spectroscopy (NMRS)

This non-invasive and non-destructive technique requires labelled substrates ($^{31}$P) and is applied to PAOs to study their physiology. The labelled substrates are included in Poly-P granules, which allows assessing the substrate degradation, and the synthesis of end-products and metabolites in only one sample [28]. The method can be applied to liquid (culture media) and solid samples and provides an excellent overview of the molecules containing phosphorus (Poly-P but also orthophosphate, pyrophosphate, phosphate mono- and di-esters, phosphonates). However, the interference caused by the heterogeneous physical and chemical properties of the samples and the association of P containing molecules with metallic ions (iron, manganese) must be highlighted. Another point is the similarity of linkage in Poly-P and other molecules containing phosphoanhydride bonds, such as nucleotides. Consequently, NMR usually provides results of the distribution of chemical linkages detected in the sample considered without providing data about the real Poly-P concentration [23]. NMR can also be coupled with bioreactors to follow up the concentrations in P containing metabolites [21]. A recent research article investigated the P transfers under aerobic and anaerobic conditions through NMR spectroscopy and studied the extracellular polymeric substances, focusing on Poly-P [50]. This work showed that NMR can be used as a relevant quantitative technique applicable to the study of PAOs.

2.2.9. RAMAN Spectromicroscopy (RAM)

RAMAN spectromicroscopy requires a simple preparation protocol of samples and may be used with complementary molecular techniques [22,23]. Near-infrared reflectance spectroscopy was first found to be able to provide precious data about carbon, carbonate, N, and P in sediments [51]. RAMAN spectromicroscopy is a related technique which allowed to identify and quantify Poly-P and other storage polymers involved in phosphate accumulation in individual bacterial cells. It also
identified orthophosphate, trimers, hexamers, and other oligomers on the basis of the specificity of spectra [23]. RAMAN spectroscopy was also found to be a reliable technique applicable to the study of microbiota composition [52]. More specifically, a research article reports the successful use of RAMAN spectromicroscopy to investigate the dynamics of intracellular Poly-P in PAOs in EBPR process [53]. Further research was applied to Poly-P, PHA, and glycogen, and developed a reliable technique to assess the intracellular distribution and dynamics of these polymers in different metabolic stages of the EBPR process [54]. Another research paper described RAMAN spectromicroscopy as a powerful tool to identify bacteria as PAOs or GAOs on the basis of their intracellular polymers [23]. Indeed, GAOs and PAOs are known to compete to consume carbon sources, and their metabolic properties relating to phosphorus are completely different. GAOs do not participate in the P-removal process [15].

2.2.10. Enzyme Assays (EA)

The metabolism of Poly-P depends on the activity of different types of enzymes. It is synthesized in bacterial cells by Poly-P kinases (PPK1 and PPK2) and degraded by exopolyphosphatase (PPX). A first technique to detect Poly-P in samples consists in using a yeast exopolyphosphatase on the extract and measuring the resulting phosphate concentration. Another enzyme assay is based on the action of PPK, which converts ADP to ATP in the presence of Poly-P [21]. After the enzymatic hydrolysis, the reaction products can be analyzed by thin-layer chromatography or gel electrophoresis [46]. Other techniques based on the activity of Poly-P-glucokinase and Poly-P AMP phosphotransferase were also reported [20]. Such enzyme assays require a good quality of the extract supposed to contain Poly-P.

2.2.11. Cryoelectron Tomography and Spectroscopic Imaging (CTSI)

This technique was used in the study of Comolli et al. [55] to investigate polymeric structures in Caulobacter crescentus and Deinococcus grandis. Here, the study of subcellular structure and subcellular bodies in whole bacteria was undertaken. This combination led to the identification of P-rich and carbon-rich bodies in Caulobacter crescentus, and provided structural information without altering the granules. P-rich granules were also detected in Deinococcus grandis.

2.2.12. Mass Spectrometry (MS)

Electrospray ionization mass spectrometry (ESI-MS) was tested on orthophosphate, pyrophosphate, tripolyphosphate, and tetrapolyphosphate [21]. This technique shows a high selectivity and does not require pre-separation steps. However, this method was not really investigated on samples extracted from EBPR experiments or pure cultures of PAOs.

2.2.13. Proteic Affinity (PA)

Saito et al. [56] developed a proteic affinity technique which leads to the detection of Poly-P. The principle is based on the use of a recombinant Poly-P binding domain (PPBD) of an exopolyphosphatase from Escherichia coli. The PPBD showed a high affinity for long-chain Poly-P and a low affinity for short-chain Poly-P. The technique was found to be able to locate Poly-P by an immunocytochemical method.

2.2.14. “Omics” Techniques (OMICS)

Omics techniques have been widely used in many studies over the past decade. Here, we present some applications of those data-rich methods to the study of PAOs. Metagenomics is mainly based on the analysis of 16S rRNA extracted from complex microbiotas, and is perfectly applicable to the study of the strains involved in the EBPR process [57]. This technique was combined with quantitative FISH in a previous work and was applied to the bacterial community involved in EBPR process [35]. Metatranscriptomics, based on the analysis of mRNA sequences, has also been used to investigate
the EBPR process. Such study was successfully applied to the research of Poly-P kinase sequences and focused on *Accumulibacter* [58]. Finally, metaproteomics provides data about the expression of proteins in complex samples, but the results obtained by this method must be considered carefully. Genes are regulated at the transcriptional, translational, and post-translational levels, and another point comes from the short half-life of gene products. Consequently, metaproteomics’ data should not be considered alone [57]. The study of Wilmes et al. [59] reported a metaproteomic analysis of bacterial consortia according to the efficiency of the EBPR process. A successful EBPR process led to the detection of a majority of *Accumulibacter*, while the failure of the process led to a majority of *α*-Proteobacteria [58,59]. The study of Wexler et al. [60] used radiolabelled metaproteomics to investigate the effect of anaerobic and aerobic conditions on the EBPR process. Another combined study was based on proteogenomics and assessed the role of *Accumulibacter* in the EBPR process, taking the other co-existing strains into account. The metaproteome was found to be very close in both aerobic and anaerobic conditions, and denitrification, fatty acid cycling and the glyoxylate bypass were revealed to have an important role in the EBPR process [59].

Table 2 summarizes the specificities of each technique described above.

Table 2. Specificities of techniques applied to the detection of PAOs and Poly-P.

| Technique | Investment | Advantages | Disadvantages |
|-----------|------------|------------|---------------|
| LFM-MB    | low        | Simplicity, rapidity<br>Direct visualization of Poly-P granules | Not adapted to visualize small granules<br>Non-quantitative technique<br>Requirement of a preparation protocol and an acidic pH<br>Destructive technique |
| LFM-NR    | low        | Simplicity, rapidity<br>Direct visualization of Poly-P granules | Technique targeting acidic vacuoles and not Poly-P itself<br>Non-quantitative technique<br>Requirement of a preparation protocol and an acidic pH<br>Destructive technique |
| LFM-DAPI  | low        | Possibility to visualize polyhydroxyalkanoate<br>Possibility of combination with other techniques (FC) | Expensive staining reagent<br>Non-quantitative and destructive technique<br>No specificity to Poly-P<br>Importance of the noise with high DAPI concentrations |
| FC        | high       | Possibility to visualize polyhydroxyalkanoate<br>Possibility to sort the cells in a complex sample<br>Possibility to study the phenotypic heterogeneity<br>Possibility to locate Poly-P inside the cells<br>Possibility to combine different staining techniques<br>Possibility to combine with FISH-A<br>Possibility to adapt the technique to animal cells | Same disadvantages as LFM-DAPI<br>Obligation to filtrate the samples<br>Destructive technique |
| FISH-A    | low        | Possibility to combine with FC and FISH-A<br>Possibility to study specific groups of PAOs<br>Possibility to study gene expression<br>Possibility to study complex microbiotas | Complexity of sample preparation<br>Efficiency depending on the penetration of the probes inside the cells<br>No specificity to Poly-P<br>Destructive technique |
| EXT       | low        | Possibility to measure phosphate resulting from the hydrolysis by diversified techniques<br>Possibility to characterize different Poly-P fractions<br>Quantitative technique | Efficiency depending on the association of Poly-P with other molecules<br>Effect of cations and chain length on the analysis<br>The extraction can lead to undesired modifications of Poly-P<br>Destructive technique |
| PAGE      | low        | Possibility to measure the size and the DP of Poly-P<br>Possibility to combine with DAPI staining<br>Semi-quantitative technique | The technique requires a step of Poly-P extraction<br>Poly-P must be treated to be soluble<br>Destructive technique |
### Table 2. Cont.

| Technique | Investment | Advantages                                                                 | Disadvantages                                                                 |
|-----------|------------|----------------------------------------------------------------------------|------------------------------------------------------------------------------|
| EM        | high       | Possibility to locate Poly-P inside the cells                             | Complexity of sample preparation                                           |
|           |            | Possibility to combine the technique with staining protocols and X-RAY    | Non-quantitative and destructive technique                                   |
|           |            | Possibility to observe Poly-P granules inferior to 100 nm                | SEM may be not adapted to observe internal Poly-P granules                    |
| X-RAY     | high       | Powerful combination with EM                                              | Possible loss of Poly-P during the preparation protocol                      |
|           |            | Quantitative analysis                                                     | Destructive technique                                                        |
|           |            | Possibility to combine with X-ray fluorescence spectromicroscopy leading to the study of the repartition of Poly-P inside the cells |
| NMRS      | high       | Global assessment of phosphorus metabolism inside the cells              | Tagged substrates required                                                   |
|           |            | Applicability to liquid and solid samples                                | Frequency of interferences due to the medium                                 |
|           |            | Possibility to use the method as a follow-up technique of bioreactors     | No specificity to Poly-P but measurement of phosphodiester linkages           |
|           |            | Nondestructive technique                                                 |                                                                              |
| RAM       | high       | Simplicity of preparation protocol                                        | Weakness of RAMAN signal which requires an adapted device                   |
|           |            | Quantitative detection of Poly-P and other storage polymers              | Interferences due to impurities and fluorescence                             |
|           |            | Applicability to the study of microbial consortia                         |                                                                              |
|           |            | Distinction of PAOs and GAOs                                              |                                                                              |
| EA        | low        | Quantitative technique                                                    | Complexity of preparation of samples to avoid an inactivation of enzymes due to impurities |
|           |            | Possibility to assess the metabolism of Poly-P extracts                  | Complexity of enzymatic reaction, less simple than with simple chemical reagents |
| CTSI      | high       | Possibility to measure the diameter of Poly-P granules with a high accuracy | Non-quantitative and visual technique                                         |
|           |            | Possibility to observe many cellular structures                          |                                                                              |
|           |            | Nondestructive technique                                                 |                                                                              |
| MS        | high       | Possibility to characterize different Poly-P fractions                   | Destructive technique                                                        |
|           |            | Possible combination with other techniques (FISH-A)                      | The technique requires standard samples                                       |
|           |            | Simple sample preparation protocol                                        |                                                                              |
| PA        | low        | Quantitative technique                                                    | Complexity of sample preparation                                             |
|           |            | Possibility to locate Poly-P                                              | The technique requires antibodies specific to Poly-P                          |
|           |            | High resolution when the technique is combined with EM                    |                                                                              |
| OMICS     | high       | Global overview of microbial consortia                                    | Not adapted to measure the Poly-P content                                   |
|           |            | Possible combination with other techniques (FISH-A)                      | Requirement of up-to-date databases                                          |
|           |            | Possibility to assess the evolution of complex consortia in time and according to specific conditions |                                                                              |

### 3. Conclusions

Analytical techniques focusing on the isolation or the detection of Poly-P are quite diversified. However, there is evidence that the culture of pure PAOs requires more investigation, and the enrichment of phosphate-accumulating consortia is an alternative to this constraint. This method requires a perfect understanding of the effect of environmental conditions on PAOs in order to establish phosphate accumulation-oriented consortia. This field is still lacking knowledge, although the general metabolism has been described in many publications. Flow cytometry is also a powerful technique applicable to the concentration of PAOs. Screening the genes responsible for phosphate accumulation is another solution which could lead to the engineering of phosphate-accumulating strains, more efficient than the wild-types. In these conditions, the accumulation of phosphate should be feasible without requiring specific conditions, such as aeration rate or the presence of specific components. Then, the recovery of phosphate from wastewater or other types of liquid effluents will be much easier.
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Abbreviations

The following abbreviations are used in this manuscript:

| Abbreviation | Description |
|--------------|-------------|
| ADP          | Adenosine Diphosphate |
| ATP          | Adenosine Triphosphate |
| CTSI         | Cryoelectron Tomography and Spectroscopic Imaging |
| DAPI         | 4',6-Diamidino-2-Phenylindole |
| DP           | Degree of Polymerisation |
| EA           | Enzyme Assays |
| EBPR         | Enhanced Biological Phosphorus Removal |
| EM           | Electron Microscopy |
| EM-EDX       | Electron Microscopy—Energy-dispersive X-ray |
| ESI-MS       | Electrospray Ionisation Mass Spectrometry |
| EXT          | Extraction procedures |
| FACS         | Fluorescence-Activated Cell Sorting |
| FC           | Flow Cytometry |
| FESEM        | Field-Emission Scanning Electron Microscopy |
| FISH         | Fluorescence in Situ Hybridization |
| GAO          | Glycogen Accumulating Organism |
| ICP-AES      | Inductively Coupled Plasma—Atomic Emission Spectrometry |
| LFM          | Light and Epifluorescence Microscopy |
| MAR          | Micro-autoradiography |
| MS           | Mass Spectrometry |
| MT           | Megaton |
| NMRS         | Nuclear Magnetic Resonance Spectroscopy |
| OMICS        | “Omic” techniques |
| P            | Phosphate |
| PA           | Proteic Affinity |
| PAGE         | Polyacrylamide Gel Electrophoresis |
| PAO          | Phosphate Accumulating Organism |
| PHA          | Polyhydroxyalkanoate |
| Poly-P       | Polyphosphate |
| PPBD         | Polyphosphate Binding Domain |
| PPK          | Polyphosphate kinase |
| PPX          | Exopolys phosphatase |
| RAM          | Raman spectromicroscopy |
| SEM          | Scanning Electron Microscopy |
| STEM         | Scanning Transmission Electron Microscopy |
| TCA          | Tricarboxylic acid |
| Tg           | Teragram |
| WWTP         | Wastewater Treatment Plant |
| X-RAY        | X-ray analysis techniques |

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