

Delft University of Technology

### Drivers of bioaggregation from flocs to biofilms and granular sludge

Aqeel, Hussain; Weissbrodt, David G.; Cerruti, Marta; Wolfaardt, Gideon M.; Wilén, Britt Marie; Liss, Steven N.

DOI 10.1039/c9ew00450e

**Publication date** 2019 **Document Version** 

Final published version

Published in Environmental Science: Water Research and Technology

#### Citation (APA)

Ageel, H., Weissbrodt, D. G., Cerruti, M., Wolfaardt, G. M., Wilén, B. M., & Liss, S. N. (2019). Drivers of bioaggregation from flocs to biofilms and granular sludge. Environmental Science: Water Research and Technology, 5(12), 2072-2089. https://doi.org/10.1039/c9ew00450e

#### Important note

To cite this publication, please use the final published version (if applicable). Please check the document version above.

Copyright

Other than for strictly personal use, it is not permitted to download, forward or distribute the text or part of it, without the consent of the author(s) and/or copyright holder(s), unless the work is under an open content license such as Creative Commons.

**Takedown policy** Please contact us and provide details if you believe this document breaches copyrights. We will remove access to the work immediately and investigate your claim.



Featuring work from Prof. Steven N Liss, Chemistry and Biology, Faculty of Science, and Vice-President Research and Innovation, Ryerson University, Toronto, Canada.

Drivers of bioaggregation from flocs to biofilms and granular sludge

Microbial aggregation reflect dynamic processes; and, structures that form are not necessarily discrete, but fall along a continuum. This review focuses on the drivers affecting the formation and stability of flocs, granules, and biofilms, and conditions that support hybrid and integrated systems for biological wastewater treatment.

## As featured in:



See Steven N. Liss *et al., Environ. Sci.: Water Res. Technol.,* 2019, **5**, 2072.





Registered charity number: 207890

# Environmental Science Water Research & Technology



**View Article Online** 

# **CRITICAL REVIEW**



Cite this: Environ. Sci.: Water Res. Technol., 2019, 5, 2072

Received 30th May 2019, Accepted 22nd September 2019

DOI: 10.1039/c9ew00450e

rsc.li/es-water

#### Water impact

# Drivers of bioaggregation from flocs to biofilms and granular sludge

Hussain Aqeel, <sup>®</sup> a David G. Weissbrodt, <sup>®</sup> Marta Cerruti, <sup>®</sup> Gideon M. Wolfaardt, <sup>®</sup> d Britt-Marie Wilén <sup>®</sup> and Steven N. Liss <sup>®</sup>

Microorganisms in natural and engineered environments interact with surfaces and form aggregates consisting of cells and an extracellular matrix. The design of the process and appropriate operational conditions drive the formation of these biofilms, flocs, and granular structures. The application of granular sludge technologies for nutrient removal is relatively new. Although research and practice benefit from several decades of investigation of biofilm and anaerobic granular sludge systems, a thorough understanding of factors affecting granulation is only beginning to emerge from bench, pilot, and full-scale investigations. Challenges intrinsic to maintaining granular and biofilm structures include management of resistance to substrate transport, establishment of targeted microbial niches, role of extracellular polymeric substances, and impacts of toxic compounds, among others. There is increasing recognition of the potential value of hybrid process configurations that optimize interactions between flocs, granules, and/or biofilm features for process enhancement and robustness. While these structures appear distinct, it is not uncommon to find a mixture of these structures present in a single system and dynamics leading to a transition from one structure to another. The transitions are dependent on changes in the microbial community and properties of the extracellular matrix. This review focuses on the drivers affecting formation and stability of flocs, biofilms, and granules and conditions that support integrated technologies for biological wastewater treatment.

The future of water/wastewater infrastructure must innovatively evolve in response to pressures (*e.g.* climate change, water quality deterioration, energy/ resource costs, and urbanization) on legacy systems, and inadequate or ageing infrastructure, as well as opportunities created by the circular economy. Fixed-film and granule-based processes are increasingly important in wastewater treatment; therefore conditions that drive microbial communities that form these structures and contribute to their stability are key to successful implementation and operations.

## 1. Introduction

Environmental biotechnology processes such as those designed for biological nutrient removal (BNR) from wastewater rely on the proliferation and metabolic activity of specialist microorganisms present in the seeding incoming wastewater to form a microbial community engineered under deterministic and stochastic conditions. In such processes, microorganisms seldom function as single cells dispersed in suspension (such

E-mail: D.G.Weissbrodt@tudelft.nl, M.Cerruti@tudelft.nl

E-mail: britt-marie.wilen@chalmers.se

as commonly conceptualized in industrial biotechnology) but form aggregates of various architectures from suspended activated sludge flocs to biofilms attached on fixed or mobile carrier materials to advanced forms of mobile aggregates of biofilms, called "granules". Bioaggregation is the predominant mode of growth in nature as it protects the microorganisms from predation, environmental stresses, enhances access to nutrients and facilitates microbial syntrophy.<sup>1</sup> The key to mixed-culture processes is to harness this natural propensity for process design, stability, and resiliency. The microbial composition and activity play an important role in the removal of organic matter and nutrients from the wastewater, with bioaggregation facilitating overall process efficiency through improved separation of treated water and biomass retention.<sup>2</sup> Therefore, the aim is to create conditions that favor the development of compact and stable bioaggregates that readily separate from the bulk aqueous phase without leaving excessive concentrations of suspended solids in the effluent. Bioaggregation is a complex process that is influenced by

<sup>&</sup>lt;sup>a</sup> Queen's University, Kingston, Ontario, Canada. E-mail: 9ha11@queensu.ca <sup>b</sup> Delft University of Technology, Delft, Netherlands.

<sup>&</sup>lt;sup>c</sup> Department of Chemistry and Biology, Office of the Vice-President, Research and Innovation, Ryerson University, Toronto, Ontario, Canada.

E-mail: steven.liss@queensu.ca, steven.liss@ryerson.ca

<sup>&</sup>lt;sup>d</sup> Stellenbosch University, Stellenbosch, South Africa. E-mail: gmw@sun.ac.za, gwolfaar@ryerson.ca

<sup>&</sup>lt;sup>e</sup> Chalmers University of Technology, Sweden.

several factors. The mechanisms involved in the formation of suspended bioaggregates into flocs, biofilms and granules are essentially the same, where environmental factors such as feeding regime and hydrodynamic conditions are decisive for the structure to form.<sup>3</sup> Biological aggregation phenomena should, therefore, be considered in such systems as a continuum from flocs to biofilms and granules (Fig. 1). A thorough understanding of biological, chemical, and physical mechanisms that govern microbial immobilization and structural dynamics is necessary to manage bioaggregation.<sup>2</sup>

fellow

Oueen's Canada.

research

dynamics

formed the

The activated sludge process for treatment of wastewater relies on the ability of bacteria to form flocs which can be separated from the treated water. Activated sludge flocs are suspended aggregates of bacterial cells assembled in a matrix of extracellular polymeric substances (EPS).<sup>1</sup> Secretion of EPS promotes bacterial adhesion and embed the different floc components. Bacteria which lack the ability to aggregate are washed out from the secondary settlers (clarifiers) which causes poor effluent quality and inefficient recycling of return sludge to the activated sludge process. The settling and stability properties depend on the floc



**Hussain Ageel** 

workshop proposed and held at the "IWA Biofilms: Granular Sludge Conference 2018".

of



David Weissbrodt

Prof. David Weissbrodt is Assistant Professor at Delft University of Technology, Netherlands. He leads the Weissbrodt Group for Environmental Life Science Engineering in the Environmental Biotechnology Section of the Department of Biotechnology. He benefits from a multi-disciplinary skillset envirobiochemical in engineering developed from HES-SO Valais/Wallis to EPFL, ETH Zürich, and Eawag in Switzerland prior to international research as

Wolfaardt

Environmental Microbiologist at

the Departments of Chemistry

University and Microbiology at

Stellenbosch. He held a Canada

Research Chair in Biofilms and

Interfaces and ERWAT Chair in

Water Management. He is an

adjunct professor at Applied

and

Toronto, and was a visiting

professor at the Thayer School

University

at

Biology

is

an

Ryerson

Chemical

of

Swiss NSF Fellow at TU Delft and Centre for Microbial Communities at Aalborg University, Denmark. His expertise targets systems microbiology integration into bioprocess design. He teaches in Life Science & Technology and Civil Engineering & Geoscience, and received one of the Best Teacher Awards of the Years 2017/18 and 2018/19 of the Environmental Engineering track. He organized and chaired the "IWA Biofilms: Granular Sludge Conference 2018" with Merle de Kreuk in Delft.

Gideon

and



Marta Cerruti

Marta Cerruti is а PhD candidate at TU Delft, the Netherlands. the In Life Environmental Science Engineering group (Environmental Biotechnology, Biotechnology department) she is investigating the metabolic versatility and physiology of purple non-sulfur bacteria for water treatment and bioproducts valorization. Previously, she obtained her MSc in Plant Biotechnology at University of

Hussain Aqeel is a post-doctoral

the

and applied microbiology. His

Ph.D. research work, on the role

and matrix in the formation and

stability of microbial structures,

basis

microbial

School

biotechnology,

community

of the

extracellular

Studies

University, Kingston,

He is working on

projects related

of

at

to

at

Environmental

environmental

Torino, Italy, with a master project on Tetrasphaera japonica as a possible model for the metabolism of polyphosphateaccumulating organisms at EPFL, Switzerland.



Gideon Wolfaardt

of Engineering at Dartmouth College, NH. Current projects are funded by EU H2020, Global Challenges Research Fund, East Rand Water Care Company, Water Research Commission and First Rand Bank, with collaborators from Germany, UK, Austria, Italy, Spain, South Africa, Canada, Mozambique.

Chemistry

Engineering,



**Fig. 1** Schematic representation of the dynamics in biofilm, flocs and granular structures. In a hybrid system, biofilms serve as a nursery where dispersed planktonic cells can attach to the substratum to initiate biofilm floc or to each other to initiate floc formation; and, dispersed planktonic biofilms can associate with flocs and develop into granular sludge or granules. Similarly, flocs and granules can disintegrate or change structure (*e.g.* filamentous outgrowth) or transition to intermediary forms (*e.g.* granular sludge).

structure, size, and intra-particle forces holding the floc components together. Most bacteria grow in colonies which contribute to the formation of compact and strong aggregates.<sup>4</sup> The filamentous bacteria often grow in flocs for which they may

contribute to floc strength by providing backbones, but are definitely linked to poorer settling properties.<sup>5</sup> Enhancement of aggregation and controlling settling properties of biomass is less of a consideration for biomass retention in some BNR systems



Britt-Marie Wilén

Britt-Marie Wilén has а background as chemical engineer and is Professor at the department of Architecture and Civil Engineering at Chalmers University of Technology and head of division of Water Environment Technology. She is leading a research group on wastewater process technology specializing in biological processes such as activated sludge flocculation, biofilm and granular sludge formation,

nutrient removal, bioelectrochemical systems, microbial community analysis and removal of micropollutants.



Steven Liss

Steven Liss' research interests span areas of environmental biotechnology, applied microbial microbiology, and structures and processes in engineered and environmental systems. He is a professor in Chemistry Biology at and Ryerson University, Toronto, and is the Vice-President Research and Innovation. He has held faculty appointments at the University of Guelph and at Queen's University, Kingston, in

Environmental Studies and Chemical Engineering, where he is Professor Emeritus. He holds a visiting professorship at Stellenbosch University, South Africa; and, has had appointments at the Advanced Environmental Biotechnology Centre, Nanyang Technological University, Singapore, and in Environmental Science and Engineering, Tongji University, Shanghai, China. and lagoons with extended aeration and/or long hydraulic retention times. These systems can include quiescent zones that allow for settling that occurs naturally over longer periods; whereas, structural properties and settling behavior are important in other systems including sequencing batch reactors (SBR), to form BNR granules.

Flocs and granules can be viewed as suspended biofilms, without carrier material.<sup>6,7</sup> The landscape of the microbial population within a community of an attached biofilm, or suspended biofilm (floc or granule) depends on different internal gradients of nutritional and environmental condition. The size of a biofilm, floc and granule influence the diffusion of dissolved oxygen and nutrients and complexity of microbial community.<sup>8,9</sup> The flocs and biofilms are integrated structures. However, the physical, chemical and microbial structures of biofilms and flocs may vary and complement each other for BNR in a hybrid system with biofilms and flocs growing in the same bioreactor.<sup>10</sup>

The granular sludge process has premium BNR capacity in a more compact system compared to conventional activated sludge processes.<sup>11</sup> Nevertheless, the demonstrated instability of granular sludge has initially limited its widespread largescale application.<sup>12,13</sup> Despite the discovery of so-called "aerobic" granules nearly two-decades ago for the aerobic removal of COD<sup>14</sup> there is little information on microbial community interactions and dynamics in relation to the operating conditions of the treatment process.<sup>15</sup> The application of anaerobic selectors do make the difference in stabilizing the granular biofilm architectures by selecting for slow-growing organisms.15-17 The advent of advanced molecular and microscopy techniques allow for in-depth elucidation of microbial community composition and architectures of granules<sup>18,19</sup> and could, therefore, pave the way for new strategies to improve water treatment. The use of multivariate analyses in numerical ecology enables to screen for multi-scale correlations,<sup>18,20,21</sup> while in-depth examination of parameter effects<sup>16</sup> is needed toward process design, understanding, and control. Furthermore, there are inconsistent and conflicting reports on factors affecting granulation including dissolved oxygen and shear rate.<sup>22-25</sup> Unlike anaerobic granules, a wider diversity of aerobic (O<sub>2</sub> as terminal electron acceptor), anoxic (nitrite or nitrate as e-acceptor) and anaerobic (no terminal eacceptor)† microorganisms can contribute to granulation in

BNR systems, each of which may have different ecological requirements to establish across redox and substrate gradients within these aggregates (Table 1).

Recent analytical developments associated with the use of molecular methods and advanced microscopy have further increased our understanding of microbial interactions and structural features of the EPS found in granular biofilm structures.<sup>15,26</sup> In this review, we examine the drivers of the formation and stability of the microbial structures that are functional components of BNR in wastewater treatment. Flocs, granules and biofilms, distinct entities, in one respect, are relatively pleomorphic in another and these structures can be viewed as dynamic shifting from one structure to another (Fig. 1). It is not uncommon to find that systems designed and operated for one type is in fact a hybrid system. The structural characteristics of these bioaggregates are dependent on microbial community dynamics, activity, and the extracellular matrix. The tendency of mature bioaggregates to disperse demand implementation of strategies (such as hybrid systems) to enhance performance and increased resiliency and stability.

# 2. Formation of flocs, biofilms and granules in BNR systems

Granulation typically arises from microbial flocs with the same initial aggregation of cells involving a number of mechanisms supporting bioflocculation. For example, initial cell-cell contact (attractive forces between cells causing them to aggregate), formation of microbial aggregates embedded in EPS onto which cells can attach and multiply,27-29 followed by the hydrodynamic shaping of the aggregate into spherically shaped aggregates.23 The management of the microbial resource via engineering of operational conditions can therefore drive granule formation and stability.15 Granules have stratified structures like mature biofilms, therefore granules are also called suspended biofilms.<sup>15</sup> The attached biofilm formation can be broadly described as a stepwise process, which includes three phases: (a) an initial reversible phase (biofilm initiation), (b) an irreversible attachment phase (biofilm growth and maturation), and (c) a dispersion phase to start a new biofilm.9 The dispersed cells reattach and regrow to form new biofilms that result in overall stable biofilm systems.

Three basic strategies involving a feast/famine regime, shear force, and short settling time have been proposed to stimulate BNR granule formation (Fig. 2).

Nonetheless, the essence of granulation originates at the microbial level. The invention of granular sludge development was based on the observation that slow-growing microorganisms are capable to convert easily degradable organic substrates under feast conditions to form complex storage polymers such as polyhydroxyalkanoates (PHA), which they can subsequently utilize when easily degradable organic substrates are less available during famine conditions.<sup>30</sup> The continuous research and gradual optimization of operational

<sup>†</sup> In the wastewater engineer's lexicon, redox conditions are considered as "aerobic" in the presence of dissolved oxygen as terminal electron acceptor *via* aeration, as "anoxic" in the presence of nitrite and nitrate as e-acceptors, and "anaerobic" in the absence of terminal e-acceptors in the bulk. This is in relative mix up with microbiology terms of "aerobic" respiration (O<sub>2</sub> as terminal e-acceptors), "anaerobic" respiration (inorganics like nitrite and nitrate as terminal e-acceptors), and fermentative conditions. One typical illustration of this apparent wording opposition resides in the anaerobic ammonium oxidation (anammox) process that has been characterized by microbiologists (therefore "anaerobic") whilst engineers should call it "anoxic" ammonium oxidation according to their systematics (presence of nitrite as terminal e-acceptor). The engineer's lexicon that is applied in the wastewater sector is primarily used in this article.

Table 1 Predominant bacterial genera found in BNR granules

Genus	Bacterial class	Metabolic characteristic	Type of wastewater	Ref.
Rhodanobacter	γ-Proteobacteria	Complete DEN	Synthetic	Aqeel <i>et al.</i> <sup>26</sup>
Klebsiella	γ-Proteobacteria	Aerobic DEN	Synthetic	Taheri <i>et al.</i> <sup>141</sup>
Thauera	β-Proteobacteria	Aerobic DEN	Piggery wastewater	Zhao <i>et al.</i> <sup>142</sup>
Zooglea	β-Proteobacteria	DEN	Synthetic	Zhao <i>et al.</i> <sup>143</sup>
Competibacter	γ-Proteobacteria	GAO	Synthetic	Weissbrodt et al.15
Accumulibacter	β-Proteobacteria	PAO	Synthetic	Weissbrodt et al.15
Candidatus Jettenia	Planctomycetes	Anammox	Synthetic	Quan <i>et al.</i> <sup>144</sup>
Brocadiacea	Planctomycetes	Anammox	Full-scale wastewater	Gonzalez-Gil et al. <sup>100</sup>

DEN = denitrifying, PAO phosphorous accumulating bacteria, GAO glycogen accumulating bacteria.



Fig. 2 Microenvironmental conditions in BNR granules. The microenvironment in oxic and anoxic zones of BNR granules helps the growth of specific bacteria (*e.g.* autotrophs and heterotrophs) to support simultaneous nutrient removal processes. There are several pathways for biological nutrient removal where bacteria compete for nutrients.

conditions such as anaerobic feeding (feast) and oxic/anoxic react time during famine conditions facilitate slow-growing bacteria and discourage fast-growing bacteria.<sup>24</sup> This can notably be achieved by designing efficient anaerobic selectors that select for preferential storage of organics by slow-growing bacteria such as polyphosphate-accumulating organisms (PAOs) and glycogen accumulating organisms (GAOs) prior to switching on aeration.

The use of shear force for BNR granules was adapted based on observations of anaerobic granulation in up-flow anaerobic sludge bioreactors (UASB), where the up-flow velocity of wastewater and gases (produced during fermentation) and the downward settling of the biomass supported anaerobic granule formation.<sup>31</sup> The high shear force results in erosion of surface particles, physically break filamentous outgrowth and facilitates mass transfer to the deep layers of the granules and biofilms. The erosion of

surface particles results in short mean cell residence times of fast-growing heterotrophic bacteria located in the outer layers.<sup>32–35</sup> Slow-growing organisms such as autotrophic nitrifying and anammox bacteria, PAOs, and GAOs form strong microcolonies that remain intact at a high shear rate.<sup>15,36</sup>

The nitrifying bacteria form dense microcolonies that are selected when a short settling time for granulation is applied. The growth of these dense microcolonies results in granulation that in turn can support the growth of denitrifying bacteria in the core of the granules. Additionally, a model that proposes the clumping of different microcolonies with a filamentous backbone or adhesive bacteria to form large aggregate and support growth of denitrifying bacteria facilitates granulation.<sup>37,38</sup> Weissbrodt *et al.*<sup>38</sup> reported on the role of heterotrophic *Zoogloea* in the formation of granules that has adhesive properties to support granule formation. It was observed that after initial

granulation, which was dependent on fast-growing bacteria, the slow-growing PAOs and GAOs out-compete the fast-growing heterotrophic bacteria. Therefore, a gradual shortening of the settling time (15 to 3 minutes) was adopted for granule maturation. Nevertheless, the presence of filamentous bacteria and of *Zoogloea*-like organisms are mainly a sign of a deficient anaerobic selector. The design of an efficient anaerobic selector is the key for a stable process.<sup>39,40</sup>

# 3. Extracellular matrix maintaining the cohesion of bioaggregates

The complex arrangement of EPS creates a distinct environment supporting both the diffusion of nutrients and retention of metabolites and a highly selective environment for the microbial community.<sup>41</sup> EPS are important constituents of flocs, granules, and biofilms contributing to both structural and functional features of as well protecting cells from harsh environmental conditions.<sup>6,22</sup> The proteins and polysaccharides (major components of EPS) form attractive and repulsive interactions within EPS supporting aggregation of cells and the formation of a compact structure.<sup>42,43</sup> In spite of decades of research, the knowledge about its production, composition, and function is limited. Hence, contradictory results can be found in the literature regarding the importance of EPS in granule formation, partly due to differences in extraction and analytical methods.<sup>44</sup> The protein to polysaccharide ratio is higher in granules as compared to flocs;<sup>26,45</sup> whereas the protein to polysaccharide ratio is higher in flocs as compared to biofilms, cultivated in an integrated fixed-film activated sludge system.<sup>10</sup> The protein content of the EPS contributes to a more hydrophobic surface and a more compact and tightly packed granular structure, compared to flocs.<sup>46</sup> In contrast, the polysaccharide content supports hydration and facilitates retention of nutrients and metabolites through interactions of the granules.22,47 charged surface and sorption in Exopolysaccharides or glycosides have been found to be an important gelling agent in BNR granules, more adhesive than EPS extracted from activated sludge.48,49

Transmission electron microscopy observations have revealed cell to cell interaction with the help of thin fibers in biofilms flocs.27 environmental and Later, the immunohistochemistry studies have shown that these fibers were the amyloid adhesins that are abundant in the environmental biofilms and flocs.<sup>36,50</sup> **Bioinformatics** analyses indicate that amyloid adhesins are widespread in bacteria related to phyla Proteobacteria, Bacteroidetes, Firmicutes and Thermodesulfobacteria.51 Whole genome microarray analyses of flocs have shown that the expression of pili and lectins is upregulated under phosphorus limiting conditions that can also affect microbial cohesion.52 Venter et al.53 applied "whole-genome shotgun sequencing" to microbial population collected from the Sargasso Sea that revealed about 300 flagellar genes.

The extracellular adhesins (e.g., flagella, pili, lectins and amyloid adhesins) are abundant in environmental, multispecies microbial communities of biofilms and flocs. However, the specific role of the extracellular adhesins in the environmental biofilms, flocs and granules is not well understood. Most of the information about the potential significance of extracellular adhesins is inferred from the pure culture and co-culture biofilm studies. The extracellular protein forms three types of specific interactions in the extracellular matrix: (a) protein to protein interaction (amyloid adhesins), (b) protein to polysaccharide interaction (lectins), and relatively less understood (c) protein to extracellular DNA interaction (type IV pili binds specifically to the extracellular DNA, where DNase treatment can inhibit development of Pseudomonas<sup>54</sup> and Acidovorax the biofilms<sup>55</sup>).

Amyloid adhesins are beta-sheet rich, fibrous proteins, which are insoluble and resistant to the proteolytic enzyme activity of proteases.<sup>56</sup> Amyloid fibers of neighboring bacteria cluster together and twist to form rope-like structures. The clustering and rope formation of amyloid adhesins at interbacterial surfaces pulls the neighboring bacteria close to each other. The amyloid-adhesin-expressing bacteria are tightly packed, 20-fold stiffer and relatively more hydrophobic compared to non-amyloid adhesin producing bacteria.<sup>57,58</sup> Amyloid adhesins mediate cell-cell interactions for microbial aggregation in flocculation and biofilm formation.<sup>59</sup> The inhibition of amyloid adhesin synthesis results in a thin biofilm, due to the loss of cell-cell interaction.<sup>57</sup>

Lectins, sugar-binding proteins (that are highly specific for their target polysaccharide) are produced by many microorganisms that facilitate cell-cell and cell-surface adhesions. Usually, these lectins are small binding sites present on the tip of the cellular appendages, which make these appendages adhesive for biofilm formation and development.<sup>60</sup> For example, the type 1 fimbrin D-mannose specific adhesin (FimH) present on the tip of type I pili binds specifically to mannose sugars.<sup>61</sup> FimH is very adaptive to the environmental stress and there are multiple wild type FimH mutants.<sup>62</sup> The binding force of the FimH to the target polysaccharide is positively correlated with shear stress. The FimH has a polysaccharide-recognizing domain, which can bind to mono-mannose. The FimH possess a second polysaccharide-binding domain, which cannot bind to the second polysaccharide in hydrostatic conditions unless there is a shear force. The shear stress in hydrodynamic conditions stretches the inter-domain space of FimH to bind the trimannose.61,63

Cellular appendages (flagella and type IV pili) help motile bacteria to move towards the favorable conditions (a) in an aquatic system for biofilm initiation, (b) within biofilm or (c) detach from the biofilm. In the (a) case, flagella (chemotactic swimming) and type IV pili (twitching or gliding) mediate initial bacterial attachment and spread of biofilm on a substrate. Flagellated bacteria can resist the hydrodynamic conditions (where non-flagellated bacteria are washed away)

to anchor on a substrate for biofilm initiation.<sup>64</sup> In the (b) case, type IV pili enable bacteria to move vertically within biofilms for favorable conditions. Motile bacteria (with flagella and type IV pili) are present in the outer layer of biofilm, which is relatively rich in the nutrient source; whereas nonmotile bacteria are predominant in the core of the biofilms.<sup>65,66</sup> In the (c) case, bacterial attachment downregulates the flagellar synthesis and induces synthesis of exopolysaccharides and amyloid adhesins.<sup>67</sup> In contrast, flagellar synthesis is induced prior to biofilm sloughing. The motile bacteria (with flagella and type IV pili) mediate sloughing when conditions are not favorable in a biofilm. Additionally, it has been observed in Bacillus subtilis biofilms that p-amino acids are secreted during the late stage of biofilm development, before biofilm sloughing. D-Amino acids trigger the disassembly of amyloid fibers, which result in the disintegration of the biofilm.68

A two-layer model with respect to the arrangement of EPS as a tightly bound layer at the core and a loosely bound layer at the outer region of bioaggregates appears to be an important feature of formation and stability of granules and biofilms and their structure that requires greater attention.<sup>42,69,70</sup> Basuvaraj *et al.*<sup>69</sup> described the formation of a granular sludge composed of a predominantly tightly bound protein-rich EPS and a smaller proportion of loosely bound EPS. The loosely bound layer was more pronounced in the seeding activated sludge floc structure and was richer in polysaccharides. The autotrophic nitrifying biofilms and flocs show that these are predominantly composed of tightly bound EPS.<sup>71</sup>

EPS in flocs and granules may be hydrolyzed to form soluble microbial products (SMP).<sup>72,73</sup> The EPS hydrolysis for SMP production has been largely attributed to heterotrophs; whereas, autotrophs have been reported to account for <10% of the SMP, with 5% associated with ammonia oxidizing bacteria and 3% related to nitrite oxidizing bacteria.<sup>73</sup> Ni *et al.*<sup>74</sup> have reported for granular sludge that soluble microbial products generated can be used as an organic substrate by denitrifying bacteria. Therefore, the growth of autotrophic nitrifying bacteria in the outer layer helps to conserve the nutrient resources, by minimizing the loss of SMP to the aquatic column. Whereas, the SMP secreted by the heterotrophic bacteria to budget the nutrient resources.

# 4. Microbial community architectures and selection phenomena in bioaggregates

Granules and biofilms are organized multicellular structures composed of mixed microbial populations, which cooperate for the optimum harvesting of resources from the surrounding environment and competitive advantage for the capture of limited resources (Fig. 3).<sup>24,75,76</sup> The cooperation in microbial community include: (a) production of EPS that

can be hydrolyzed by other bacteria and synthesis of complex storage polymers that serve as nutrient source for some bacteria; (b) production of metabolic byproducts that can be used by other bacteria, *e.g.*, nitrite and nitrate resulting from the metabolism of nitrifying bacteria are used as electron acceptors by denitrifying bacteria. The effective cooperation between bacteria present in the outer oxic zone and deep anaerobic layer of the granules results in structural and functional stability of the granular sludge.

The spherical architecture of BNR granules with outer aerobic layer and core with anoxic/anaerobic is a widely accepted conceptual model of the granular structure, typically implemented in mathematical models. The stratified structure and the ability to support simultaneous nitrification and denitrification in granules are owing to the thickness of large size of granules (diameters  $> 200 \ \mu m$ ) where the penetration of dissolved oxygen is restricted to the outer 50-100  $\mu$ m.<sup>77,78</sup> However, there are few contradicting observations, where the distinction between communities across the depth is absent.<sup>79</sup> Barr et al.<sup>79</sup> hypothesized that granules with stratified structure form by the growth of single microcolony; whereas, granules with homogeneous microbial community form by aggregation of microcolonies. Weissbrodt et al.15 highlighted with CLSM different types of internal architectures of granules (i) with smooth continuous matrices when dominated by fast-growing biofilm heterotrophs like Zoogloea spp., (ii) with conglomerates of microcolonies when slow-growing organisms like PAOs, GAOs, and nitrifiers populate across the granule crosssection, and (iii) unfavorably overgrown by filamentous bacteria when the organic matter was not adequately removed in an (anaerobic) selector. In contrast to Barr et al.,<sup>79</sup> Weissbrodt et al.<sup>15</sup> stated that the growth physiology of the predominant microorganisms do impact on the resulting internal structure of granules. This statement is valid across flocs, granules, and biofilms, and meets with previous modelling reports of Alpkvist et al.<sup>80</sup> Gonzalez-Gil and Holliger<sup>78</sup> recreated the three-dimensional image of the granules that revealed a "cavernlike" granular structure. The "cavernlike" channels in the granules improve the diffusion of nutrients and deviate the microbial growth, compared to the widely accepted model of a spherical granular structure. Additionally, the denitrifying bacteria could be observed in the outer layer because some bacteria are capable of aerobic denitrification.81

#### 4.1. Microbial niches of the oxic zone of bioaggregates

The operational conditions for wastewater treatment are engineered to drive the selection of cooperative microbial communities that maintain granular sludge with optimum size and activity. For example, in the oxic zones of these granules, the metabolism and biomass assimilation of the nitrite-oxidizing bacteria (NOB) that convert nitrite to nitrate are dependent on ammonia oxidizing bacteria (AOB) that convert ammonium to nitrite. Furthermore, since NOB such



**Fig. 3** Conditions selecting for BNR granule formation and retention (a) within sequencing batch reactors the settling phase facilitates selection of the larger, compact and fast settling granules and eradicate the relatively slow settling microbial aggregates. (b) Shear forces erode the fast-growing bacteria from the outer layer resulting in its low cell residence time, whereas within the core slow-growing cells are retained (until cell lysis due to aging) and out-compete fast-growing bacteria. Shear forces induce PS secretion resulting in a PS rich outer layer, which facilitates granule formation (c) "Feast famine" conditions facilitate the development of slow-growing bacteria and conversion of readily degrading organic matter to storage polymers including PHA. Gradients and stratification arise with respect to the diffusion of external nutrients and secretion of metabolic byproducts.

as *Nitrobacter* and *Nitrospira* compete for available nitrite resources in wastewater, the operational and nutritional conditions can favor the growth of one bacterial group over another. *Nitrospira* outcompetes *Nitrobacter* at microaerophilic and low nitrite concentrations.<sup>82</sup> Conversely, *Nitrobacter* outcompetes *Nitrospira* at higher nitrite and oxygen concentration.

The dependence of NOB on AOB is reflected by a low NOB/ AOB ratio (0.5) in conventional activated sludge flocs. However, in granules (where *Nitrobacter* predominates) a higher NOB/AOB ratio (3) has been reported. The ping-pong theory and nitrite loop theory have been used to explain the abundance of NOB in granules.<sup>79,83</sup> According to the pingpong theory: in the oxic region, NOB convert nitrite to nitrate autotrophically and then metabolize nitrates heterotrophically in the anoxic region to produce nitric oxide. The NOB (which typically grow autotrophically are also observed in deeper anoxic region) grow mixotrophically to outnumber the AOB. NOB are capable to use organic substrate present in the feed during the feast phase and convert these to storage polymers such as PHA.<sup>79</sup> During famine phase, NOB uses organic matter either stored as PHA or released by cell lysis. The heterotrophic growth of NOB (which is independent from the availability of nitrite) results in an increase in the NOB/AOB ratio in granular biomass.<sup>79</sup> Alternatively, a nitrite loop may develop whereby the nitrite is oxidized by NOB to nitrate, followed by reduction back to nitrite by denitrifying bacteria. The accumulated nitrite is now available again to be used by NOB.<sup>79,83</sup> The recent insights in *Nitrospira*-like organisms able a complete ammonium oxidation (comammox) into nitrate.<sup>84–86</sup> can further explain the predominance of NOB at the expense of AOB in granules.

#### 4.2. Microbial niches of the anoxic zone of bioaggregates

Heterotrophic bacteria typically prefer dissolved oxygen as an electron acceptor; however, denitrifying heterotrophs can use nitrate and nitrite as an electron acceptor under anoxic conditions. In conventional wastewater treatment (floccular sludge and biofilm-based systems) denitrification is usually performed in a separate tank under anoxic conditions. In the granular sludge that removes nutrients simultaneously,

In BNR granules, non-denitrifying heterotrophs can outcompete denitrifying bacteria in the presence of limited organic substrate and extensive aeration.87 Therefore, strategies to ensure optimum granulation involve conditions (e.g., feast-famine) to discourage the growth of heterotrophs in the oxic zone. When this strategy does not efficiently inhibit the heterotrophic growth, usually higher organic substrate in the wastewater combined with improved biomass transport with shear force is required to support denitrifying heterotrophic growth in the anoxic zone. Denitrifying bacteria, which are able to utilize storage polymers (e.g., PHA) as an organic substrate can outcompete growing heterotrophs and facilitate fast granule formation.<sup>17,30</sup> Furthermore, in the anoxic zone, PAOs and GAOs are also capable to use storage polymers (PHA) and thus compete with denitrifying microorganisms for the limited organic substrate.<sup>88,89</sup> However, PAOs (which are desired for enhanced biological phosphorous removal -EBPR) and GAOs (which are typically considered undesired in EBPR systems) are both capable of denitrification.<sup>15,90-92</sup> The competition between the microbial community for limited PHA can be manipulated by optimizing the operational conditions according to the requirement of the system.<sup>93</sup>

Conditions where substrates (external or stored) are limited, denitrifying bacteria prefer nitrate as an electron acceptor. Denitrifying bacteria reduce nitrite and nitrate sequentially, to nitric oxide, nitrous oxide, and dinitrogen gas, with the help of reductase enzymes. Reduction of nitrate is preferred over nitrite reduction (by nitrate reductase and nitrite reductase, respectively) during denitrification because nitrate can accept relatively more electrons than nitrite, which results in accumulation of nitrite at low organic loading, in anammox process granules. The accumulated nitrite thus favors the coexistence of anammox and denitrifying bacteria in granules.<sup>74</sup> Conversely, nitrous oxide reduction to nitrogen gas is halted at low organic loading to nitrogen loading ratios, to budget the organic loading. Consequently, nitrous oxide (N<sub>2</sub>O, a greenhouse gas) emission is undesired in simultaneous nitrification and denitrification in granules.<sup>94</sup> Additionally, the activity of nitrous oxide reductase is inhibited at low dissolved oxygen concentrations, resulting in the unfavorable emission of N<sub>2</sub>O.<sup>95</sup>

The microbial ecology of denitrifying bacteria varies with operational and nutritional conditions such as solid retention time, temperature, pH, carbon source, chemical oxygen demand and nitrogen ratios.<sup>96</sup> For example, most denitrifying bacteria require a pH of 6.5–7.5, for nitrogen removal. However, *Rhodanobacter* related to the class *Gammaproteobacteria* (complete denitrifying bacteria, which can reduce nitrate and nitrite to nitrogen gas) are capable of denitrification at a broader range of pH 4–8.<sup>97,98</sup> *Rhodanobacter* were predominant in granules where pH was not maintained during the react cycle (pH ranged 7–4) in sequencing batch reactor (SBR).<sup>26</sup>

The microbial community composition in the anoxic zone depends upon the concentration and composition of nutrients in the influent, activity of bacteria in the outer layer (that depends on operational conditions including temperature, pH, dissolved oxygen concentration) and architecture of nutrient transport channels in the bioaggregates. The anammox and denitrifying bacteria can coexist in the anoxic zone of the granules.<sup>99,100</sup> The growth rate of anammox bacteria is lower compared to their competitors: AOB and NOB in the outer layer and denitrifying bacteria in the inner layer. To support the growth of anammox bacteria: (1) the growth of AOB must be limited and NOB inhibited, for surplus ammonia and nitrite for anammox activity. The activity of AOB and NOB populations can be suppressed by several mechanisms, in bioreactors with simultaneous nitrification and denitrification activity. For example, by decreasing the temperature from 29 °C to 15 °C (ref. 101) and intermittent and low aeration rate.<sup>102</sup> Ma et al.<sup>102</sup> showed that the dissolved oxygen concentration during aeration ranged between  $0.08-0.25 \text{ mg L}^{-1}$ , which was enough to support ammonium oxidation. Aeration was interrupted when nitrite start to accumulate, to discourage nitrite oxidation and thereby facilitate anammox bacteria;<sup>102</sup> (2) the growth of denitrifying bacteria is discouraged by feeding high strength nitrogenous wastewater to the system, with relatively low organic substrate  $(300 \text{ mg L}^{-1})^{74}$  and a low carbon to nitrogen ration of two.

#### 4.3. Microbial niches of the anaerobic zone of bioaggregates

Most of the environmental bacteria are opportunist that grow when conditions are favorable. With increase in the size of the granule or biofilm, anaerobic bacteria proliferate in the core of these bioaggregates.<sup>103</sup> In the anaerobic zone a number of bacteria related to different functional activities have been observed, including bacterial groups with anammox, denitrifying, hydrolyzing, and fermentative activities.<sup>18,19,99</sup> Hydrolyzing bacteria degrade the complex storage polymers, EPS, and dead cells to produce organic substrate to support the growth of nutrient-removing bacteria (denitrifiers, PAO and GAO).<sup>18</sup> As size of the granules further increases, mass transport of nutrients in and waste-products out of the core is reduced that suppress the growth of bacterial subpopulations in the inner parts of the bioaggregates. A core of dead cells has been observed at a depth of 900  $\mu m.^{31}$  However, the threshold size for mass transport may vary depending upon local operation condition and/or due to clogging of the mass transport channels by excessive EPS.<sup>104</sup> The clogging of mass transport channels reduces the metabolic activity of nutrient removing bacteria. The growth of fermentative bacteria in the core is favored when organic substrate starts to accumulate with hydrolytic activity.<sup>105</sup> Fermentative bacteria create acidic conditions that further inhibit the growth of competitive bacteria and promote cell lysis that results in granule disintegration. In large size granules with clogged channels,<sup>104</sup> hydrolysis of dead cells and EPS complemented with

fermentation (that produce fermentative gases) creates a hollow core.<sup>106</sup> The granules with hollow core loose density, float on the surface and are washed out of the system.<sup>107</sup> Therefore, maintaining an optimal granule size is imperative for the stability of the system.

#### 4.4. Microbial community dynamics and granule instability

The dominant populations in BNR granular sludge as revealed by various studies are summarized in Table 1. It has been suggested that operational or nutritional conditions that do not favor the growth of these dominant bacteria, result in changes in the relative abundance of unfavorable microbial populations, and consequently in granule instability.<sup>26,108</sup> Zou et al.<sup>108</sup> reported that changes in nitrogen concentration in influent wastewater resulted in an imbalance in nitrogen removal and phosphorous accumulating organisms. This shift in the microbial community structure resulted in functional instability and subsequent disintegration of the BNR granules. Isanta et al.<sup>109</sup> examined the sensitivity of the anammox process and the microbial community of granular sludge to temperature changes (temperature increased from 35-46 °C for eight days). An imbalance in the microbial community and functional instability of anammox processes was observed following this temperature change for eight days. The functional stability of the bioreactor recovered after 70 days of operation. In these two recent studies, the microbial diversity was further reduced in the recovered bioreactor following the transient instability and recovery period.<sup>26,109</sup>

# 5. Instability and control of floc, biofilms, and granular structures

Activated sludge floc instability can be caused by fluctuations in chemical composition and physical properties of incoming wastewater and dynamics in operational environmental conditions, such as temperature which affects microbial activity as well as the structural and functional properties of the EPS; pH and ionic composition which determine surface charge; divalent cations such as Ca<sup>2+</sup> which contribute to the cation-bridging of EPS; and various toxins.<sup>1</sup> These parameters have a similar influence on biofilms and granules.<sup>3,43</sup>

Biofilm sloughing can be triggered by several physical, chemical or biological stressors that can be induced due to changes in environmental or nutritional conditions. The microcolonies growing in a biofilm have varied responses to nutritional and environmental stress.<sup>110</sup> Biofilm dispersion increases with an increase in surface area loading rate and hydraulic retention time in moving bed bioreactors treating wastewater.<sup>111,112</sup> In membrane aerated biofilm reactors, higher chemical oxygen demand favours biofilm stability; whereas higher oxygen partial pressure results in loss of EPS at the bottom of biofilm that triggers biofilm detachment.<sup>113</sup> The biomass in BNR system undergoes consistent hydraulic shear force. Biofilms and granules that are formed at higher shear force form dense, compact and stable biofilms and

granules. The higher shear force improves EPS production and efficiency of mass transfer to improve the structural and functional stability of the biofilms and granules.<sup>23</sup>

Bacteria can relocate within biofilms towards a more favorable microenvironment or detach from a biofilm to search for better environmental and nutritional resources.<sup>114</sup> A subpopulation of bacteria may lyse when conditions are not favorable for bacteria in deeper layers of the biofilms. The organic matter released due to cell lysis serves as nutrition for the nearby surviving bacteria.<sup>115</sup> An increase in carbon source up-regulates the flagellum expression and down-regulates the adhesive extracellular pili.<sup>116</sup> The switch in the expression of extracellular adhesins results in sloughing, which leaves a hollow biofilm.<sup>110</sup> Overall, these phenomena demonstrate that biofilms are organized structures with the ability to reorganize with changes in environmental or nutritional conditions.<sup>32,117</sup>

In granular sludge, filamentous outgrowth, granule disintegration, presence of gas bubbles, and higher polysaccharide content in the extracellular matrix, can lead to poor settleability. Shorter settling time helps in the removal of these compromised structures; however, if instability events become dominant in the system, it results in biomass washout and system failure.<sup>26,118</sup> The application of short settling time is also difficult at full scale. A good management of hydraulic regimes during fill/draw phases is essential to maintain fastersettling aggregates in the reactor while washing-out slowsettling flocs.40,119 Liu and Liu120 reviewed and listed several operational and nutritional conditions causing granule instability, including long solid retention time and low dissolved oxygen concentration. Interestingly granules have cultivated successfully under these been operating conditions.<sup>119</sup> Additionally, filaments are usually considered to cause granule instability and biomass washout.<sup>26,118</sup> Conversely, it has been observed that filaments play a role in granule formation. The filaments serve as the backbone and immobilize the bacteria community to form large flocs, which can be converted into granular sludge by hydrodynamic shear forces. Furthermore, filaments can bridge the large flocs and enmesh them to form granules.7,26,121,122 Nevertheless, the presence of filaments will mainly always remain a latent issue. A good anaerobic selector should specifically suppress the growth of filamentous organisms, protecting granules from a filamentous bulking outbreak.15

# 5.1. Practical considerations to manage structural and functional characteristics

Key issues in the development and scaling of biofilm- and granule-based wastewater treatment processes are presented in Table 2. Anaerobic feeding of a readily degradable organic substrate facilitates the conversion of organic substrates to complex storage polymers, to support slow-growing PAOs and GAOs. If the organic substrate is fed under oxic conditions, the substrate is consumed by fast-growing heterotrophic bacteria present in the outer layer which results in starvation

Table 2	Key issues in t	the development an	d scaling of biofiln	n and granular	based wastewater	treatment processes <sup>a</sup>
	<i>,</i>		5	2		

Formation and stability	Scaling and full-scale operations	Fundamental questions offering new insights
i) Feeding conditions & pulse vs. slow feed	i) Controlling hydrodynamics and mixing	i) How to better integrate or design BNR granular, and biofilm systems to support anammox processes
ii) Composition and variability of industrial wastewater	ii) Selective sludge removal and design criteria for height to diameter ratios	ii) How to selectively remove specific groups of bacteria ( <i>e.g.</i> heterotrophs to enhance partial nitration and
iii) Nutriant loading and stability	moving from lab to full-scale	anammox (PN/A)) from granules
( <i>e.g.</i> high strength organics, N & P)	solids concentrations	hybrid systems of flocs-films, floc-granules and/or film-granules better path to stability, treatment, and operational efficiency outcomes?
iv) Sludge volume loading	iv) Applicability of current biofilm models for granular systems	iv) Increased understanding of the role of EPS in the formation and stability of granular sludge
v) Effect of metals		v) Resolving microbial interactions and functions related to formation, function and stability of films and granules

<sup>*a*</sup> Summary of key issues identified by wastewater engineering researchers and practitioners in industry, covering expertise in microbiology, civil and environmental engineering, and chemical engineering, in attendance at the *Drivers of Granular Sludge and Biofilm Stability Workshop* (March 18, 2018), "IWA Biofilms: Granular Sludge Conference 2018", IWA Biofilms Specialist Group, Delft University of Technology, Delft, Netherlands, March 18–21, 2018.

and lysis of slow-growing bacteria present in the core.<sup>123</sup> Polymeric substrates which are slowly degradable, and which are not consumed under anaerobic conditions in the core, contribute to conditions that enable fast-growing filamentous outgrowth at the surface of the granules where oxic conditions exist. The presence of polymeric organic substrates in wastewater is one of the limiting factors for stability of granular sludge in full-scale wastewater treatment plants.<sup>124,125</sup> Wagner *et al.*<sup>126</sup> proposed to treat wastewater with high polymeric substrates content by operating SBRs with longer anaerobic conditions for effective hydrolysis and to improve the granule stability. For optimum nutrient removal and structural stability of granules, a balance between aeration and non-aeration periods during the operating cycles of the reactor is therefore required.

Another example of the need to understand the relationships of these microbial systems is the formation of a relatively less dense and "fluffy" granular structure in the presence of a higher EPS polysaccharide content. While these structures retain many of the features of a functional granule including size, they settle slowly in the bioreactor (like flocs)<sup>127,128</sup> and are thus no longer granules by virtue of their settleability characteristics. If it is desirable to retain these granular structures in the reactor, an adjustment upward in settling time is required. Washing out slow-settling aggregates by gradually decreasing the settling time from 15 minutes to 3 minutes has been efficient to form dense granules from "fluffy" structures at lab scale.<sup>15,39</sup>

Simultaneous nitrogen removal is possible because BNR granules retain metabolic byproducts, which can be used by other bacteria,<sup>24</sup> however, it also results in retention of metabolites, which may be toxic to the microbial community.<sup>26</sup> A few studies have reported the presence of bacteria capable of producing toxic metabolites or enzymes in BNR granules, which can lead to the disintegration of the granular structure. Adav *et al.*<sup>129</sup> identified bacteria, which produce protease enzymes that were responsible for a decline

in EPS protein content resulting in granule disintegration. The adhesive protein content can also decline due to loss of specific bacterial populations which contribute to the secretion of protein in EPS, resulting in granule instability.<sup>26,130</sup> *Janthinobacterium* (capable of producing an antibiotic pigment called violacein) have been observed in BNR granules that may contribute to bacterial cell lysis resulting in granule instability.<sup>26,131</sup> Incubation of BNR granules with enzymes degrading the  $\alpha$  (1–4) glucans and proteins and exposure to shear stress resulted in aggregate instability.<sup>132</sup> Little is, however, known about the biosynthesis of these molecules, or their concentrations in full scale operations.

Overgrowth of a microbe that is producing less common non-toxic products of metabolism may contribute to microbial community dynamics that lead to granule instability. Aqeel *et al.*<sup>26</sup> reported on the outgrowth of *Auxenochlorella* in BNR granules that can accumulate lipid and chitin. Successional changes in the community resulted in an initial decline of *Gammaproteobacteria* followed by the growth of the bacteria *Janthinobacterium* and *Chitinophaga*, which can use these byproducts as nutrient substrates. In this instance, *Chitinophaga* was antagonistic towards *Auxenochlorella* resulting in the recovery of the granular structure and return to a granular microbial community dominated by the denitrifying genus *Rhodanobacter.*<sup>26,133</sup>

Most of the information on granulation is based on intensive research on laboratory scale SBRs, which are operated under well-controlled operational and nutritional conditions. Few reports have been published that describe structural and functional characteristics of granules in fullscale plants.<sup>119,134,135</sup> Pronk *et al.*<sup>119</sup> reported the detailed operational conditions for granulation in full-scale wastewater treatment, where feast-famine conditions and fast settling times were applied. The full-scale SBR was seeded with surplus biomass from an existing granular bioreactor that was operated with a long solid retention time (20–38 days) for selection of slow-growing bacteria. The full-scale SBR was fed anaerobically for 60 minutes, aerated for 300 minutes during the react phase, settled for 30 minutes, and the treated water was drained in 60 minutes, during dry weather conditions. During a period of rain the number of SBR cycles was increased to allow for a constant daily organic loading of the plant: the reactor was anaerobically fed for 90 minutes, aerated for 60 minutes during the react phase, and settled for 30 minutes. The length of anaerobically feeding and react phase were optimized during seasonal variations according to the fluctuation in environmental and nutritional conditions. In full-scale SBR, the stable granule formation was achieved after five months of startup period that was maintained by optimizing the operational conditions.

Current efforts further granulation in existing flowthrough installations to retrofit and intensify them while keeping the original plant design and components. The key remains in the design of a procedure to select for fast-settling aggregate while separating slower-settling flocs.<sup>136–138</sup> While process engineering considerations are under elaboration world-wide, the underlying microbiology still remains a central element of the puzzle. Several full-scale plants operated for full BNR with well-operating anaerobic selectors have been observed to granulate "spontaneously". A good conjunction of process engineering and microbial community engineering should be kept in mind in order to manage the continuum from flocs to granules.

# 5.2. Hybrid systems to optimize granule structure-function relationship

In SBRs, the larger and dense granules settle to the bottom of the tank and relatively smaller and less dense structures are drained from the system. This results in a gradual development of granules with a large diameter. The large diameter limits the transport of nutrients in the granules which ultimately lead to cell lysis of the slow-growing bacteria in the anaerobic core and granule disintegration.<sup>123,139</sup> The large granules are also more prone to channel blockage which limits the mass transfer of nutrients even more. The blocked channels also result in entrapment of gas bubbles and lowering of granule density. Eventually, these structures float to the surface and are washed out of the system.<sup>107</sup>

Hybrid biomass with a wider range of bioaggregate sizes is suggested to support the stability of granular sludge. The hybrid biomass may reduce the overall settleability compared to uniform larger granules. However, hybrid biomass has advantages that the disintegration of larger granules has minimal effect on the overall settleability of the granular sludge; where the disintegrated structures can reform into granules. Pronk *et al.*<sup>119</sup> reported stable full-scale granular sludge system with granules ranging from 0.2 mm to larger than 1 mm, with 20% flocs smaller than 0.2 mm in size. Zhu *et al.*<sup>140</sup> suggested selective biomass removal to achieve hybrid granular sludge that improved the stability of the system. In addition to removal of floccular sludge (to keep an SRT of 2.7 days for flocs) from of the settled biomass, the aged larger granules were also removed (SRT of 9.9 days for aged granules) from the bottom of the tank, to promote a hybrid biomass in the system.<sup>140</sup> However, the selective removal of biomass is possible in bioreactors with higher height to diameter ratios (h/d), where the efficient separation of biomass is possible during settling time. Higher aeration rate is used to minimize the development of large size granules where h/d is lower, but only with limited success.

Granulation in integrated fixed-film activated sludge (a hybrid system) represents an interesting development to maintain hybrid biomass and minimize the development of large size granules. Granule formation is relatively reproducible, and the resulting granules are relatively stable in bioreactors operated under controlled conditions, as compared to full-scale wastewater treatment plants. The selective pressure applied for granule sludge production results in a decrease in overall microbial diversity. The dominant microbial population is usually distinct from one system to the other (Table 1) due to unique nutritional and environmental conditions for most wastewater systems. The loss of microbial diversity may contribute to granule instability due to the inability to adapt to changes in operational nutritional/environmental or conditions. Whereas, full-scale wastewater treatment plants are operated under diverse environmental and nutritional conditions, some of which is due to seasonal changes (winter) and precipitation events (rain). The seasonal variation also affects the efficiency of conventional full-scale wastewater treatment. In contrast, full-scale integrated fixed-film activated sludge (IFAS) systems show year-round stable BNR due to the presence of both attached and suspended biomass.<sup>10</sup> The biomass in hybrid reactors shows bioaggregates with a wider range in size and density. The wider range of this type of biomass has diverse redox properties. The wider size range of the hybrid biomass in IFAS systems conserves a more diverse microbial community, where different microbial groups are dominant in attached and suspended biomass.<sup>10</sup> This result in improved adaptability for structural and functional stability during a change in environmental and nutritional conditions. In an experiment where the attached and suspended biomasses were separated to understand the capability of hybrid biomass in BNR, it has been observed that the contribution from the attached and suspended biomass is not additive. Instead, it has been synergistic to improve the overall BNR in IFAS system. Additionally, it has been observed that biofilms in IFAS systems tend to slough and readily associate with the suspended biomass. The sloughed biomass forms the stable granular flocs that improve the overall settling properties of the biomass.

## 6. Conclusion

Microbial aggregation phenomena should be considered on a continuum from flocs to biofilms and granules. Biofilm and

granule formation start with initial bioaggregation that grow in size and when it reaches a threshold limit, biofilms and granules disperse to produce floccular biomass and start a new cycle of biofilm and granule formation. The microbial aggregates are organized structures that can be engineered to achieve a target microbial niche for the desired function. Biofilm sloughing results in partial loss of biomass and transient decline in biological nutrient removal; whereas when granules disperse (in fast settling operations) it may result in a major biomass loss and system failure. Granular sludge management should, therefore, include a strategy that prepares for dispersion events, by keeping (a) a heterogeneous size range of granules that compensate for dispersion of a fraction of the granules and simultaneously favor regranulation of dispersed biomass; and (b) a hybrid microbial aggregates (flocs, granules, and biofilms) in the system. The hybrid biomass ensures microbial structures with diverse redox condition and diverse microbial population for enhanced performance and resilience.

# Conflicts of interest

There are no conflicts of interest to declare.

# Acknowledgements

SNL and HA acknowledge funding from Queen's University and Natural Sciences and Engineering Research Council of Canada (NSERC) Collaborative Research and Training Experience (CREATE) program. DGW benefitted from a fellowship from the Swiss National Science Foundation (grant no. 151977) for his work on granular sludge microbiomes. The review follows a workshop organized by the authors and held in Delft, Netherlands, March 2018, in conjunction with the "TWA Biofilms: Granular Sludge Conference 2018" of the Biofilms Specialist Group of the International Water Association.

# References

- 1 A. Suresh, E. Grygolowicz-Pawlak, S. Pathak, L. S. Poh, M. bin Abdul Majid, D. Dominiak, T. V. Bugge, X. Gao and W. J. Ng, Understanding and optimization of the flocculation process in biological wastewater treatment processes: A review, *Chemosphere*, 2018, 210, 401–416.
- 2 M. K. Winkler, C. Meunier, O. Henriet, J. Mahillon, M. E. Suárez-Ojeda, G. Del Moro, M. De Sanctis, C. Di Iaconi and D. G. Weissbrodt, An integrative review of granular sludge for the biological removal of nutrients and recalcitrant organic matter from wastewater, *Chem. Eng. J.*, 2018, 336, 489–502.
- 3 B. M. Wilén, R. Liébana, F. Persson, O. Modin and M. Hermansson, The mechanisms of granulation of activated sludge in wastewater treatment, its optimization, and impact on effluent quality, *Appl. Microbiol. Biotechnol.*, 2018, **102**, 5005–5020.
- 4 M. Klausen, T. R. Thomsen, J. L. Nielsen, L. H. Mikkelsen and P. H. Nielsen, Variations in microcolony strength of

probe-defined bacteria in activated sludge flocs, *FEMS Microbiol. Ecol.*, 2004, **50**, 123–132.

- 5 W. Burger, K. Krysiak-Baltyn, P. J. Scales, G. J. Martin, A. D. Stickland and S. L. Gras, The influence of protruding filamentous bacteria on floc stability and solid-liquid separation in the activated sludge process, *Water Res.*, 2017, 123, 578–585.
- 6 S. N. Liss, Microbial flocs suspended biofilms, *The Encyclopaedia of Environ. Microbiol.*, 2002, vol. 4, p. 2000.
- 7 J. J. Beun, A. Hendriks, M. C. M. van Loosdrecht, E. Morgenroth, P. A. Wilderer and J. J. Heijnen, Aerobic granulation in a sequencing batch reactor, *Water Res.*, 1999, 33, 2283–2290.
- 8 Y. Han, J. Liu, X. Guo and L. Li, Micro-environment characteristics and microbial communities in activated sludge flocs of different particle size, *Bioresour. Technol.*, 2012, 124, 252–258.
- 9 P. Stoodley, K. Sauer, D. G. Davies and J. W. Costerton, Biofilms as complex differentiated communities, *Annu. Rev. Microbiol.*, 2002, 56, 187–209.
- 10 B. Mahendran, L. Lishman and S. N. Liss, Structural, physicochemical and microbial properties of flocs and biofilms in integrated fixed-film activated sludge (IFFAS) systems, *Water Res.*, 2012, **46**, 5085–5101.
- 11 S. Bengtsson, M. de Blois, B. M. Wilén and D. Gustavsson, Treatment of municipal wastewater with aerobic granular sludge, *Crit. Rev. Environ. Sci. Technol.*, 2018, 48, 119–166.
- 12 A. J. Li, T. Zhang and X. Y. Li, Fate of aerobic bacterial granules with fungal contamination under different organic loading conditions, *Chemosphere*, 2010, 78, 500–509.
- 13 R. D. Franca, H. M. Pinheiro, M. C. M. van Loosdrecht and N. D. Lourenço, Stability of aerobic granules during long-term bioreactor operation, *Biotechnol. Adv.*, 2017, 36, 228–246.
- 14 E. Morgenroth, T. Sherden, M. C. M. van Loosdrecht, J. J. Heijnen and P. A. Wilderer, Aerobic granular sludge in a sequencing batch reactor, *Water Res.*, 1997, 31, 3191–3194.
- 15 D. G. Weissbrodt, T. R. Neu, U. Kuhlicke, Y. Rappaz and C. Holliger, Assessment of bacterial and structural dynamics in aerobic granular biofilms, *Front. Microbiol.*, 2013, 4, 175.
- 16 L. B. Guimarães, J. Wagner, T. R. Akaboci, G. C. Daudt, P. H. Nielsen, M. C. Van Loosdrecht, D. G. Weissbrodt and R. H. da Costa, Elucidating performance failures in use of granular sludge for nutrient removal from domestic wastewater in a warm coastal climate region, *Environ. Technol.*, 2018, 30, 1–6.
- 17 M. K. De Kreuk and M. C. M. van Loosdrecht, Selection of slow growing organisms as a means for improving aerobic granular sludge stability, *Water Sci. Technol.*, 2004, 49, 9–17.
- 18 D. G. Weissbrodt, N. Shani and C. Holliger, Linking bacterial population dynamics and nutrient removal in the granular sludge biofilm ecosystem engineered for wastewater treatment, *FEMS Microbiol. Ecol.*, 2014, 88, 579–595.
- 19 E. Szabó, R. Liébana, M. Hermansson, O. Modin, F. Persson and B. M. Wilén, Microbial population dynamics

and ecosystem functions of anoxic/aerobic granular sludge in sequencing batch reactors operated at different organic loading rates, *Front. Microbiol.*, 2017, 8, 770.

- 20 F. L. de los Reyes, J. E. Weaver and L. Wang, A methodological framework for linking bioreactor function to microbial communities and environmental conditions, *Curr. Opin. Biotechnol.*, 2015, 33, 112–118.
- 21 D. G. Weissbrodt, G. S. Schneiter, J. M. Fürbringer and C. Holliger, Identification of trigger factors selecting for polyphosphate-and glycogen-accumulating organisms in aerobic granular sludge sequencing batch reactors, *Water Res.*, 2013, 47, 7006–7018.
- 22 J. H. Tay, Q. S. Liu and Y. Liu, The effects of shear force on the formation, structure and metabolism of aerobic granules, *Appl. Microbiol. Biotechnol.*, 2001, 57, 227–233.
- 23 Y. Liu and J. H. Tay, The essential role of hydrodynamic shear force in the formation of biofilm and granular sludge, *Water Res.*, 2002, **36**, 1653–1665.
- 24 M. K. De Kreuk, J. J. Heijnen and M. C. M. van Loosdrecht, Simultaneous COD, nitrogen, and phosphate removal by aerobic granular sludge, *Biotechnol. Bioeng.*, 2005, **90**, 761–769.
- 25 H. Zhang, F. Dong, T. Jiang, Y. Wei, T. Wang and F. Yang, Aerobic granulation with low strength wastewater at low aeration rate in A/O/A SBR reactor, *Enzyme Microb. Technol.*, 2011, 49, 215–222.
- 26 H. Aqeel, M. Basuvaraj, M. Hall, J. D. Neufeld and S. N. Liss, Microbial dynamics and properties of aerobic granules developed in a laboratory-scale sequencing batch reactor with an intermediate filamentous bulking stage, *Appl. Microbiol. Biotechnol.*, 2016, 1, 447–460.
- 27 S. N. Liss, I. G. Droppo, D. T. Flannigan and G. G. Leppard, Floc architecture in wastewater and natural riverine systems, *Environ. Sci. Technol.*, 1996, **30**, 680–686.
- 28 B. Q. Liao, D. G. Allen, I. G. Droppo, G. Leppard and S. N. Liss, Surface properties of sludge and their role in bioflocculation and settleability, *Water Res.*, 2001, 35, 339–350.
- 29 Y. V. Nancharaiah and G. K. Reddy, Aerobic granular sludge technology: mechanisms of granulation and biotechnological applications, *Bioresour. Technol.*, 2018, 247, 1128–1143.
- 30 M. C. M. Van Loosdrecht, M. A. Pot and J. J. Heijnen, Importance of bacterial storage polymers in bioprocesses, *Water Sci. Technol.*, 1997, 35, 41–47.
- 31 J. H. Tay, V. Ivanov, S. Pan and S. L. Tay, Specific layers in aerobically grown microbial granules, *Lett. Appl. Microbiol.*, 2002, 34, 254–257.
- 32 C. Picioreanu, M. C. M. van Loosdrecht and J. J. Heijnen, Two-dimensional model of biofilm detachment caused by internal stress from liquid flow, *Biotechnol. Bioeng.*, 2001, 72, 205–218.
- 33 E. Morgenroth and P. A. Wilderer, Influence of detachment mechanisms on competition in biofilms, *Water Res.*, 2000, 34, 417–426.
- 34 S. S. Adav, D. J. Lee and J. Y. Lai, Effects of aeration intensity on formation of phenol-fed aerobic granules and

extracellular polymeric substances, *Appl. Microbiol. Biotechnol.*, 2007, 77, 175–182.

- 35 M. K. Winkler, R. Kleerebezem, W. O. Khunjar, B. de Bruin and M. C. M. van Loosdrecht, Evaluating the solid retention time of bacteria in flocculent and granular sludge, *Water Res.*, 2012, 46, 4973–4980.
- 36 P. Larsen, J. L. Nielsen, D. Otzen and P. H. Nielsen, Amyloid-like adhesins produced by floc-forming and filamentous bacteria in activated sludge, *Appl. Environ. Microbiol.*, 2008, 74, 1517–1526.
- 37 M. K. Winkler, J. P. Bassin, R. Kleerebezem, D. Y. Sorokin and M. C. M. van Loosdrecht, Unravelling the reasons for disproportion in the ratio of AOB and NOB in aerobic granular sludge, *Appl. Microbiol. Biotechnol.*, 2012, 94, 1657–1666.
- 38 D. G. Weissbrodt, S. Lochmatter, S. Ebrahimi, P. Rossi, J. Maillard and C. Holliger, Bacterial selection during the formation of early-stage aerobic granules in wastewater treatment systems operated under wash-out dynamics, *Front. Microbiol.*, 2012, 3, 332.
- 39 S. Lochmatter and C. Holliger, Optimization of operation conditions for the startup of aerobic granular sludge reactors biologically removing carbon, nitrogen, and phosphorous, *Water Res.*, 2014, **59**, 58–70.
- 40 D. G. Weissbrodt, C. Holliger and E. Morgenroth, Modeling hydraulic transport and anaerobic uptake by PAOs and GAOs during wastewater feeding in EBPR granular sludge reactors, *Biotechnol. Bioeng.*, 2017, 114, 1688–1702.
- 41 H. C. Flemming, T. R. Neu and D. J. Wozniak, The EPS matrix: The house of biofilm cells, *J. Bacteriol.*, 2007, 189, 7945–7947.
- 42 G. P. Sheng, H. Q. Yu and X. Y. Li, Extracellular polymeric substances (EPS) of microbial aggregates in biological wastewater treatment systems: a review, *Biotechnol. Adv.*, 2010, 28, 882–894.
- 43 H. Flemming and J. Wingender, The biofilm matrix, *Nat. Rev. Microbiol.*, 2010, 8, 623–633.
- 44 T. Seviour, N. Derlon, M. S. Dueholm, H. C. Flemming, E. Girbal-Neuhauser, H. Horn, S. Kjelleberg, M. C. M. van Loosdrecht, T. Lotti, R. Nerenberg, T. R. Neu, E. Paul, H. Yu and Y. Lin, Extracellular polymeric substances of biofilms: suffering from an identity crisis, *Water Res.*, 2018, 151, 1–7.
- 45 B. S. McSwain, R. L. Irvine, M. Hausner and P. A. Wilderer, Composition and distribution of extracellular polymeric substances in aerobic flocs and granular sludge, *Appl. Environ. Microbiol.*, 2005, 71, 1051–1057.
- 46 K. Hori and S. Matsumoto, Bacterial adhesion: From mechanism to control, *Biochem. Eng. J.*, 2010, 48, 424–434.
- 47 F. Ahimou, M. J. Semmens, G. Haugstad and P. J. Novak, Effect of protein, polysaccharide, and oxygen concentration profiles on biofilm cohesiveness, *Appl. Environ. Microbiol.*, 2007, 73, 2905–2910.
- 48 T. Seviour, M. Pijuan, T. Nicholson, J. Keller and Z. Yuan, Gel-forming exopolysaccharides explain basic differences between structures of aerobic sludge granules and floccular sludges, *Water Res.*, 2009, 43, 4469–4478.

- 49 T. Seviour, Z. Yuan, M. C. M. van Loosdrecht and Y. Lin, Aerobic sludge granulation: a tale of two polysaccharides?, *Water Res.*, 2012, **46**, 4803–4813.
- 50 P. Larsen, J. L. Nielsen, M. S. Dueholm, R. Wetzel, D. Otzen and P. H. Nielsen, Amyloid adhesins are abundant in natural biofilms, *Environ. Microbiol.*, 2007, **9**, 3077–3090.
- 51 M. S. Dueholm, M. Albertsen, D. Otzen and P. H. Nielsen, Curli functional amyloid systems are phylogenetically widespread and display large diversity in operon and protein structure, *PLoS One*, 2012, 12, 51274.
- 52 J. R. Liu, C. T. Liu, E. A. Edwards and S. N. Liss, Effect of phosphorus limitation on microbial floc structure and gene expression in activated sludge, *Water Sci. Technol.*, 2006, 54, 247–255.
- 53 J. C. Venter, K. Remington, J. F. Heidelberg, A. L. Halpern, D. Rusch, J. A. Eisen, D. Wu, I. Paulsen, K. E. Nelson, W. Nelson and D. E. Fouts, Environmental genome shotgun sequencing of the Sargasso Sea, *Science*, 2004, 304, 66–74.
- 54 E. J. Van Schaik, C. L. Giltner, G. F. Audette, D. W. Keizer, D. L. Bautista, C. M. Slupsky, B. D. Sykes and R. T. Irvin, DNA binding: a novel function of Pseudomonas aeruginosa type IV pili, *J. Bacteriol.*, 2005, 187, 1455–1464.
- 55 B. D. Heijstra, F. B. Pichler, Q. Liang, R. G. Blaza and S. J. Turner, Extracellular DNA and Type IV pili mediate surface attachment by Acidovorax temperans, *Antonie van Leeuwenhoek*, 2009, **95**, 343–349.
- 56 E. A. Epstein and M. R. Chapman, Polymerizing the fibre between bacteria and host cells: the biogenesis of functional amyloid fibres, *Cell. Microbiol.*, 2008, **10**, 1413–1420.
- 57 C. Prigent-Combaret, G. Prensier, T. T. Le Thi, O. Vidal, P. Lejeune and C. Dorel, Developmental pathway for biofilm formation in curli-producing Escherichia coli strains: role of flagella, curli and colanic acid, *Environ. Microbiol.*, 2000, 2, 450–464.
- 58 G. Zeng, B. S. Vad, M. S. Dueholm, G. Christiansen, M. Nilsson, T. Tolker-Nielsen, P. H. Nielsen, R. L. Meyer and D. E. Otzen, Functional bacterial amyloid increases Pseudomonas biofilm hydrophobicity and stiffness, *Front. Microbiol.*, 2015, 6, 1099.
- 59 M. S. Dueholm, D. Otzen and P. H. Nielsen, Evolutionary Insight into the Functional Amyloids of the Pseudomonads, *PLoS One*, 2013, 8, e76630.
- 60 C. Park and J. T. Novak, Characterization of lectins and bacterial adhesins in activated sludge flocs, *Water Environ. Res.*, 2009, **81**, 755–764.
- 61 T. K. Lindhorst, K. Bruegge, A. Fuchs and O. Sperling, A bivalent glycopeptide to target two putative carbohydrate binding sites on FimH, *Beilstein J. Org. Chem.*, 2010, 6, 801–809.
- 62 M. A. Schembri, G. Christiansen and P. Klemm, FimHmediated autoaggregation of Escherichia coli, *Mol. Microbiol.*, 2001, 41, 1419–1430.
- 63 W. E. Thomas, E. Trintchina, M. Forero, V. Vogel and E. V. Sokurenko, Bacterial adhesion to target cells enhanced by shear force, *Cell*, 2002, **109**, 913–923.

- 64 J. W. McClaine and R. M. Ford, Characterizing the adhesion of motile and nonmotile *Escherichia coli* to a glass surface using a parallel-plate flow chamber, *Biotechnol. Bioeng.*, 2002, 78, 179–189.
- 65 M. L. Gibiansky, J. C. Conrad, F. Jin, V. D. Gordon, D. A. Motto, M. A. Mathewson, W. G. Stopka, D. C. Zelasko, J. D. Shrout and G. C. L. Wong, Bacteria use type IV pili to walk upright and detach from surfaces, *Science*, 2010, 330, 197.
- 66 J. C. Conrad, M. L. Gibiansky, F. Jin, V. D. Gordon, D. A. Motto, M. A. Mathewson, W. G. Stopka, D. C. Zelasko, J. D. Shrout and G. C. L. Wong, Flagella and Pili-Mediated Near-Surface Single-Cell Motility Mechanisms, *Biophys. J.*, 2011, 100, 1608–1616.
- 67 D. O. Serra, A. M. Richter, G. Klauck, F. Mika and R. Hengge, Microanatomy at cellular resolution and spatial order of physiological differentiation in a bacterial biofilm, *mBio*, 2013, 4, 103–113.
- 68 I. Kolodkin-Gal, D. Romero, S. Cao, J. Clardy, R. Kolter and R. Losick, D-amino acids trigger biofilm disassembly, *Science*, 2010, 328, 627–629.
- 69 M. Basuvaraj, J. Fein and S. N. Liss, Protein and polysaccharide content of tightly and loosely bound extracellular polymeric substances and the development of a granular activated sludge floc, *Water Res.*, 2015, **82**, 104–114.
- 70 B. Tang, C. Yu, L. Bin, Y. Zhao, X. Feng, S. Huang, F. Fu, J. Ding, C. Chen, P. Li and Q. Chen, Essential factors of an integrated moving bed biofilm reactor-membrane bioreactor: adhesion characteristics and microbial community of the biofilm, *Bioresour. Technol.*, 2016, 211, 574–583.
- 71 Z. Liang, W. Li, S. Yang and P. Du, Extraction and structural characteristics of extracellular polymeric substances (EPS), pellets in autotrophic nitrifying biofilm and activated sludge, *Chemosphere*, 2010, **81**, 626–632.
- 72 B. J. Ni, W. M. Xie, Y. P. Chen, F. Fang, S. Y. Liu, T. T. Ren and Y. C. Tian, Heterotrophs grown on the soluble microbial products (SMP) released by autotrophs are responsible for the nitrogen loss in nitrifying granular sludge, *Biotechnol. Bioeng.*, 2011, 108, 2844–2852.
- 73 W. M. Xie, B. J. Ni, T. Seviour, G. P. Sheng and H. Q. Yu, Characterization of autotrophic and heterotrophic soluble microbial product (SMP) fractions from activated sludge, *Water Res.*, 2012, 46, 6210–6217.
- 74 S. Q. Ni, J. Y. Ni, D. L. Hu and S. Sung, Effect of organic matter on the performance of granular anammox process, *Bioresour. Technol.*, 2012, **110**, 701–705.
- 75 J. B. Xavier, M. K. De Kreuk, C. Picioreanu and M. C. M. van Loosdrecht, Multi-scale individual-based model of microbial and bioconversion dynamics in aerobic granular sludge, *Environ. Sci. Technol.*, 2007, 41, 6410–6417.
- 76 Y. Kagawa, J. Tahata, N. Kishida, S. Matsumoto, C. Picioreanu, M. C. M. van Loosdrecht and S. Tsuneda, Modeling the nutrient removal process in aerobic granular sludge system by coupling the reactor-and granule-scale models, *Biotechnol. Bioeng.*, 2015, 112, 53–64.

- 77 R. L. Meyer, R. J. Zeng, V. Giugliano and L. L. Blackall, Challenges for simultaneous nitrification, denitrification, and phosphorus removal in microbial aggregates: mass transfer limitation and nitrous oxide production, *FEMS Microbiol. Ecol.*, 2005, 52, 329–338.
- 78 G. Gonzalez-Gil and C. Holliger, Aerobic granules: microbial landscape and architecture, stages, and practical implications, *Appl. Environ. Microbiol.*, 2014, 80, 3433–3441.
- 79 J. J. Barr, A. E. Cook and P. L. Bond, Granule formation mechanisms within an aerobic wastewater system for phosphorus removal, *Appl. Environ. Microbiol.*, 2010, 76, 7588–7597.
- 80 E. Alpkvist, C. Picioreanu, M. C. van Loosdrecht and A. Heyden, Three-dimensional biofilm model with individual cells and continuum EPS matrix, *Biotechnol. Bioeng.*, 2006, 94, 961–979.
- 81 L. A. Robertson and J. G. Kuenen, Aerobic denitrification: a controversy revived, *Arch. Microbiol.*, 1984, 139, 351–354.
- 82 A. Schramm, D. De Beer, A. Gieseke and R. Amann, Microenvironments and distribution of nitrifying bacteria in a membrane-bound biofilm, *Environ. Microbiol.*, 2000, 2, 680–686.
- 83 M. K. Winkler, Q. H. Le and E. I. Volcke, Influence of partial denitrification and mixotrophic growth of NOB on microbial distribution in aerobic granular sludge, *Environ. Sci. Technol.*, 2015, 49, 11003–11010.
- 84 E. Costa, J. Pérez and J. U. Kreft, Why is metabolic labour divided in nitrification?, *Trends Microbiol.*, 2006, 14, 213–219.
- 85 M. A. Van Kessel, D. R. Speth, M. Albertsen, P. H. Nielsen, H. J. den Camp, B. Kartal, M. S. Jetten and S. Lücker, Complete nitrification by a single microorganism, *Nature*, 2015, 528, 555.
- 86 H. Daims, E. V. Lebedeva, P. Pjevac, P. Han, C. Herbold, M. Albertsen, N. Jehmlich, M. Palatinszky, J. Vierheilig, A. Bulaev and R. H. Kirkegaard, Complete nitrification by Nitrospira bacteria, *Nature*, 2015, 528, 504.
- 87 M. S. I. Mozumder, C. Picioreanu, M. C. M. van Loosdrecht and E. I. Volcke, Effect of heterotrophic growth on autotrophic nitrogen removal in a granular sludge reactor, *Environ. Technol.*, 2014, 35, 1027–1037.
- 88 G. J. F. Smolders, J. Van der Meij, M. C. M. van Loosdrecht and J. J. Heijnen, A structured metabolic model for anaerobic and aerobic stoichiometry and kinetics of the biological phosphorus removal process, *Biotechnol. Bioeng.*, 1995, 47, 277–287.
- 89 C. M. Lopez-Vazquez, C. M. Hooijmans, D. Brdjanovic, H. J. Gijzen and M. C. M. van Loosdrecht, Temperature effects on glycogen accumulating organisms, *Water Res.*, 2009, 43, 2852–2864.
- 90 Z. Bin, X. Bin, Q. Zhigang, C. Zhiqiang, L. Junwen, G. Taishi and W. Jingfeng, Denitrifying capability and community dynamics of glycogen accumulating organisms during sludge granulation in an anaerobic-aerobic sequencing batch reactor, *Sci. Rep.*, 2015, 5, 12904.
- 91 T. Kuba, E. Murnleitner, M. C. Van Loosdrecht and J. J. Heijnen, A metabolic model for biological phosphorus

removal by denitrifying organisms, *Biotechnol. Bioeng.*, 1996, 52, 685–695.

- 92 A. B. Lanham, A. Oehmen, G. Carvalho, A. M. Saunders, P. H. Nielsen and M. A. Reis, Denitrification activity of polyphosphate accumulating organisms (PAOs) in full-scale wastewater treatment plants, *Water Sci. Technol.*, 2018, 78, 2449–2458.
- 93 C. M. Lopez-Vazquez, A. Oehmen, C. M. Hooijmans, D. Brdjanovic, H. J. Gijzen, Z. Yuan and M. C. M. van Loosdrecht, Modeling the PAO–GAO competition: effects of carbon source, pH and temperature, *Water Res.*, 2009, 43, 450–462.
- 94 M. J. Kampschreur, H. Temmink, R. Kleerebezem, M. S. Jetten and M. C. M. van Loosdrecht, Nitrous oxide emission during wastewater treatment, *Water Res.*, 2009, 43, 4093–4103.
- 95 X. Quan, M. Zhang, P. G. Lawlor, Z. Yang and X. Zhan, Nitrous oxide emission and nutrient removal in aerobic granular sludge sequencing batch reactors, *Water Res.*, 2012, 46, 4981–4990.
- 96 H. Lu, K. Chandran and D. Stensel, Microbial ecology of denitrification in biological wastewater treatment, *Water Res.*, 2014, 64, 237–254.
- 97 S. J. Green, O. Prakash, P. Jasrotia, W. A. Overholt, E. Cardenas, D. Hubbard and J. E. Kostka, Denitrifying bacteria from the genus *Rhodanobacter* dominate bacterial communities in the highly contaminated subsurface of a nuclear legacy waste site, *Appl. Environ. Microbiol.*, 2012, 78, 1039–1047.
- 98 O. Prakash, S. J. Green, P. Jasrotia, W. A. Overholt, A. Canion, D. B. Watson and J. E. Kostka, *Rhodanobacter denitrificans* sp. nov., isolated from nitrate-rich zones of a contaminated aquifer, *Int. J. Syst. Evol. Microbiol.*, 2012, 62, 2457–2462.
- 99 D. R. Speth, S. Guerrero-Cruz, B. E. Dutilh and M. S. Jetten, Genome-based microbial ecology of anammox granules in a full-scale wastewater treatment system, *Nat. Commun.*, 2016, 7, 11172.
- 100 G. Gonzalez-Gil, R. Sougrat, A. R. Behzad, P. N. Lens and P. E. Saikaly, Microbial community composition and ultrastructure of granules from a full-scale Anammox reactor, *Microb. Ecol.*, 2015, **70**, 118–131.
- 101 H. De Clippeleir, S. E. Vlaeminck, F. De Wilde, K. Daeninck, M. Mosquera, P. Boeckx, W. Verstraete and N. Boon, One-stage partial nitritation/anammox at 15 C on pretreated sewage: feasibility demonstration at lab-scale, *Appl. Microbiol. Biotechnol.*, 2013, 97, 10199–10210.
- 102 B. Ma, B. P. Bao, Y. Wei, G. Zhu, Z. Yuan and Y. Peng, Suppressing nitrite-oxidizing bacteria growth to achieve nitrogen removal from domestic wastewater via anammox using intermittent aeration with low dissolved oxygen, *Sci. Rep.*, 2015, 5, 13048.
- 103 Y. Lv, C. Wan, D. J. Lee, X. Liu and J. H. Tay, Microbial communities of aerobic granules: granulation mechanisms, *Bioresour. Technol.*, 2004, 169, 344–351.
- 104 S. F. Corsino, M. Capodici, M. Torregrossa and G. Viviani, Fate of aerobic granular sludge in the long-term: the role of

EPSs on the clogging of granular sludge porosity, *J. Environ. Manage.*, 2016, 183, 541–550.

- 105 M. Pronk, B. Abba, R. Kleerebezem and M. C. Van Loosdrecht, Effect of sludge age on methanogenic and glycogen accumulating organisms in an aerobic granular sludge process fed with methanol and acetate, *Microb. Biotechnol.*, 2015, 8, 853–864.
- 106 N. Kosaric, R. Blaszczyk, L. Orphan and J. Valladarfs, The characteristics of granules from upflow anaerobic sludge blanket reactors, *Water Res.*, 1990, 24, 1473–1477.
- 107 H. F. Lu, P. Zheng, Q. X. Ji, H. T. Zhang, J. Y. Ji, L. Wang and J. W. Chen, The structure, density and settlability of anammox granular sludge in high-rate reactors, *Bioresour. Technol.*, 2012, 123, 312–317.
- 108 J. Zou, Y. Li, L. Zhang, R. Wang and J. Sun, Understanding the impact of influent nitrogen concentration on granule size and microbial community in a granule-based enhanced biological phosphorus removal system, *Bioresour. Technol.*, 2015, 177, 209–216.
- 109 E. Isanta, T. Bezerra, I. Fernández, M. E. Suárez-Ojeda, J. Pérez and J. Carrera, Microbial community shifts on an anammox reactor after a temperature shock using 454pyrosequencing analysis, *Bioresour. Technol.*, 2015, 181, 207–213.
- 110 B. Purevdorj-Gage, W. Costerton and P. Stoodley, Phenotypic differentiation and seeding dispersal in nonmucoid and mucoid Pseudomonas aeruginosa biofilms, *Microbiology*, 2005, **151**, 1569–1576.
- 111 M. S. Karizmeh, R. Delatolla and R. M. Narbaitz, Investigation of settleability of biologically produced solids and biofilm morphology in moving bed bioreactors (MBBRs), *Bioprocess Biosyst. Eng.*, 2014, 37, 1839–1848.
- 112 D. Forrest, R. Delatolla and K. Kennedy, Carrier effects on tertiary nitrifying moving bed biofilm reactor: an examination of performance, biofilm and biologically produced solids, *Environ. Technol.*, 2016, 37, 662–671.
- 113 I. X. Zhu, D. G. Allen and S. N. Liss, Effect of oxygen partial pressure and chemical oxygen demand loading on the biofilm properties in membrane-aerated bioreactors, *Water Environ. Res.*, 2009, **81**, 2892–2897.
- 114 M. Klausen, A. Aaes-Jørgensen, S. Molin and T. Tolker-Nielsen, Involvement of bacterial migration in the development of complex multicellular structures in Pseudomonas aeruginosa biofilms, *Mol. Microbiol.*, 2003, 50, 61–68.
- 115 D. McDougald, S. A. Rice, N. Barraud, P. D. Steinberg and S. Kjelleberg, Should we stay or should we go: mechanisms and ecological consequences for biofilm dispersal, *Nat. Rev. Microbiol.*, 2012, **10**, 39.
- 116 K. Sauer, M. Cullen, A. Rickard, L. Zeef, D. Davies and P. Gilbert, Characterization of nutrient-induced dispersion in *Pseudomonas aeruginosa* PAO1 biofilm, *J. Bacteriol.*, 2004, **186**, 7312–7326.
- 117 M. C. M. Van Loosdrecht, J. J. Heijnen, H. Eberl, J. Kreft and C. Picioreanu, Mathematical modelling of biofilm structures, *Antonie van Leeuwenhoek*, 2002, **81**, 245–256.

- 118 S. S. Adav, D. J. Lee, K. Y. Show and J. H. Tay, Aerobic granular sludge: recent advances, *Biotechnol. Adv.*, 2008, 26, 411-423.
- 119 M. Pronk, M. K. de Kreuk, B. de Bruin, P. Kamminga, R. Kleerebezem and M. C. M. van Loosdrecht, Full scale performance of the aerobic granular sludge process for sewage treatment, *Water Res.*, 2015, 84, 207–217.
- 120 Y. Liu and Q. S. Liu, Causes and control of filamentous growth in aerobic granular sludge sequencing batch reactors, *Biotechnol. Adv.*, 2006, 24, 115–127.
- 121 J. Williams and F. de los Reyes, Microbial community structure of activated sludge during aerobic granulation in an annular gap bioreactor, *Water Sci. Technol.*, 2006, 54, 139–146.
- 122 Y. M. Zheng, H. Q. Yu, S. J. Liu and X. Z. Liu, Formation and instability of aerobic granules under high organic loading conditions, *Chemosphere*, 2006, 63, 1791–1800.
- 123 M. Pronk, B. Abbas, S. H. K. Al-zuhairy, R. Kraan, R. Kleerebezem and M. C. M. Van Loosdrecht, Effect and behaviour of different substrates in relation to the formation of aerobic granular sludge, *Appl. Microbiol. Biotechnol.*, 2015, **99**, 5257–5268.
- 124 M. K. De Kreuk, N. Kishida, S. Tsuneda and M. C. M. van Loosdrecht, Behavior of polymeric substrates in an aerobic granular sludge system, *Water Res.*, 2010, 44, 5929–5938.
- 125 M. Figueroa, A. V. del Río, J. L. Campos, R. Méndez and A. Mosquera-Corral, Filamentous bacteria existence in aerobic granular reactors, *Bioprocess Biosyst. Eng.*, 2015, 38, 841–851.
- 126 J. Wagner, D. G. Weissbrodt, V. Manguin, R. H. R. da Costa, E. Morgenroth and N. Derlon, Effect of particulate organic substrate on aerobic granulation and operating conditions of sequencing batch reactors, *Water Res.*, 2015, 85, 158–166.
- 127 N. Schwarzenbeck, J. M. Borges and P. A. Wilderer, Treatment of dairy effluents in an aerobic granular sludge sequencing batch reactor, *Appl. Microbiol. Biotechnol.*, 2005, 66, 711–718.
- 128 A. N. Anuar, Z. Ujang, M. V. Loosdrecht and M. D. Kreuk, Settling behaviour of aerobic granular sludge, *Water Sci. Technol.*, 2007, **56**, 55–64.
- 129 S. S. Adav, D. J. Lee and J. Y. Lai, Proteolytic activity in stored aerobic granular sludge and structural integrity, *Bioresour. Technol.*, 2009, **100**, 68–73.
- 130 S. S. Adav, D. J. Lee and J. Y. Lai, Potential cause of aerobic granular sludge breakdown at high organic loading rates, *Appl. Microbiol. Biotechnol.*, 2010, 85, 1601–1610.
- 131 F. Pantanella, F. Berlutti, C. Passariello, S. Sarli, C. Morea and S. Schippa, Violacein and biofilm production in *Janthinobacterium lividum*, *J. Appl. Microbiol.*, 2007, **102**, 992–999.
- 132 C. Caudan, A. Filali, M. Spérandio and E. Girbal-Neuhauser, Multiple EPS interactions involved in the cohesion and structure of aerobic granules, *Chemosphere*, 2014, **117**, 262–270.
- 133 V. Sangkhobol and V. B. D. Skerman, *Chitinophaga*, a new genus of chitinolytic myxobacteria, *Int. J. Syst. Bacteriol.*, 1981, 31, 285–293.

- 134 A. Giesen, L. M. M. de Bruin, R. P. Niermans and H. F. van der Roest, Advancements in the application of aerobic granular biomass technology for sustainable treatment of wastewater, *Water Pract. Tech.*, 2013, **8**, 47–54.
- 135 J. Li, L. B. Ding, A. Cai, G. X. Huang and H. Horn, Aerobic sludge granulation in a full-scale sequencing batch reactor, *BioMed Res. Int.*, 2014, 2014, 1–12.
- 136 N. Kishida, A. Kono, Y. Yamashita and S. Tsuneda, Formation of aerobic granular sludge in a continuous-flow reactorcontrol strategy for the selection of well-settling granular sludge, *J. Water Environ. Nanotechnol.*, 2010, 8, 251–258.
- 137 T. R. Kent, C. B. Bott and Z. W. Wang, State of the art of aerobic granulation in continuous flow bioreactors, *Biotechnol. Adv.*, 2018, 36, 1139–1366.
- 138 C. Cofré, J. L. Campos, D. Valenzuela-Heredia, J. P. Pavissich, N. Camus, M. Belmonte, A. Pedrouso, P. Carrera, A. Mosquera-Corral and A. V. del Río, Novel system configuration with activated sludge like-geometry to develop aerobic granular biomass under continuous flow, *Bioresour. Technol.*, 2018, 267, 778–781.
- 139 K. Y. Show, D. J. Lee and J. H. Tay, Aerobic granulation: advances and challenges, *Appl. Biochem. Biotechnol.*, 2012, **167**, 1622–1640.

- 140 L. Zhu, Y. Yu, X. Dai, X. Xu and H. Qi, Optimization of selective sludge discharge mode for enhancing the stability of aerobic granular sludge process, *Chem. Eng. J.*, 2013, 217, 442–446.
- 141 E. Taheri, M. H. Khiadani, M. M. Amin, M. Nikaeen and A. Hassanzadeh, Treatment of saline wastewater by a sequencing batch reactor with emphasis on aerobic granule formation, *Bioresour. Technol.*, 2012, **111**, 21–26.
- 142 Y. Zhao, J. Huang, H. Zhao and H. Yang, Microbial community and N removal of aerobic granular sludge at high COD and N loading rates, *Bioresour. Technol.*, 2013, 143, 439–446.
- 143 X. Zhao, Z. Chen, X. Wang, J. Li, J. Shen and H. Xu, Remediation of pharmaceuticals and personal care products using an aerobic granular sludge sequencing bioreactor and microbial community profiling using Solexa sequencing technology analysis, *Bioresour. Technol.*, 2015, **179**, 104–112.
- 144 Z. X. Quan, S. K. Rhee, J. E. Zuo, Y. Yang, J. W. Bae, J. R. Park, S. T. Lee and Y. H. Park, Diversity of ammoniumoxidizing bacteria in a granular sludge anaerobic ammonium oxidizing (anammox) reactor, *Environ. Microbiol.*, 2008, **10**, 3130–3139.