

Nanoparticles applied to membrane bioreactors: potential impact on reactor performance and microbial communities

Hong Cheng ^a and Pei-Ying Hong ^{*a}

^a King Abdullah University of Science and Technology (KAUST), Water Desalination and Reuse Center (WDRC), Division of Biological and Environmental Science & Engineering (BESE), Thuwal 23955-6900, Saudi Arabia

* Corresponding author:

Pei-Ying Hong

Email: peiyong.hong@kaust.edu.sa

Phone: +966-12-8082218

1 **Abstract**

2

3 Both aerobic and anaerobic membrane bioreactors (MBRs) are able to remove emerging
4 contaminants from wastewater at high efficiencies. However, the main bottleneck of
5 this technology is membrane biofouling. Coating heavy metal nanoparticles on the
6 surface of membrane has been proposed as an effective antifouling strategy.

7 Nevertheless, metal nanoparticles can potentially result in detrimental impact on the
8 overall functionality of the MBRs. This review aims to understand how nanoparticles
9 impact MBRs. To achieve this aim, the review starts off by illustrating the antibacterial
10 mechanisms of nanoparticles. The review then critically evaluates past studies that
11 illustrate the antibacterial effect of nanoparticles against pure bacterial cultures and
12 biofilm-associated populations. Finally, the review discusses if the presence of
13 nanoparticles would affect the overall performance of aerobic and anaerobic biological
14 processes. Specifically, the impact of heavy metal nanoparticles on nitrogen and
15 phosphorus removal process of aerobic MBRs were discussed. The effect on anaerobic
16 MBR, which is comprised of hydrolysis, acidogenesis, acetogenesis and methanogenesis
17 stages, was also reviewed.

18

19

20

21 **Keywords:** Heavy metals, nanoparticles, antibacterial, functional genes

22

23

24

25 **Water impact**

26 Nanoparticles have been widely applied to membrane modification for its excellent
27 antibacterial and antifouling effects. Nevertheless, the unintended leaching of
28 nanoparticles would be potentially detrimental to the biological activity in MBR,
29 resulting in inadequately treated effluent. This review collates past studies and critically
30 discuss the antibacterial mechanisms of nanoparticles and its potential impacts on
31 aerobic and anaerobic biological processes.

32 **1. Introduction**

33 Over the last century, continued industrialization and population growth have resulted
34 in a wastewater stream that is becoming increasingly complex with time. The presence
35 of emerging contaminants (ECs) such as pharmaceuticals, illicit drugs, personal care
36 products, antibiotic-resistant pathogens and antibiotic resistance genes in untreated
37 wastewaters have been widely reported by many studies ¹⁻⁴. Heavy metals in influent of
38 wastewater treatment plant located in Thuwal, Saudi Arabia was measured, and the
39 result revealed heavy metal concentration to range from several ppb to more than
40 20000 ppb in the untreated wastewater ⁵. Many of these ECs can detrimentally impact
41 the natural ecosystems if left unremoved from the wastewaters. To illustrate, silver
42 nanoparticles and silver ions can result in a significant decrease in marine biofilm
43 volume and biomass, and perturb the major bacterial groups in a marine ecosystem ⁶.
44 Furthermore, as more countries pursue water reuse to mitigate water scarcity, the
45 presence of ECs in the reclaimed water can impose public health concerns.

46 To remove ECs from wastewaters, processes including physicochemical treatment (e.g.
47 activated carbon absorption), biological treatment (e.g. activated sludge) and advanced
48 treatments (e.g. ultra-violet (UV) photolysis, and membrane filtration) ⁷ can be used.
49 However, each process has its strengths and limitations. To exemplify, although
50 activated carbon adsorb hydrophobic compounds and non-polar compounds effectively
51 by up to 1 log, it is less effective against hydrophilic and polar compounds ⁸⁻¹⁰. Activated
52 sludge can remove a wide range of emerging contaminants by absorption and
53 biodegradation, but is unable to remove recalcitrant compounds like polar ECs
54 efficiently ¹¹. UV photolysis requires high UV intensity and contact time to remove ECs ¹²,
55 ¹³, which makes it less cost-effective.

56 In contrast, membrane bioreactor (MBR) is able to remove ECs from wastewater at high
57 log removal rates, usually reducing the concentrations by more than 1-2 log compared
58 to the influent stream ¹⁴. MBR systems couple microfiltration or ultrafiltration
59 membranes to activated sludge and are able to achieve an efficient solid-liquid

60 separation. The result is a permeate stream of higher quality than that obtained from
61 activated sludge tanks. Furthermore, MBR is typically operated with a longer sludge
62 retention time and has higher sludge concentration which may improve the removal
63 efficiency of ECs through adsorption and biodegradation. Despite MBR having these
64 advantages, the main bottleneck of this technology is membrane biofouling.

65 Biofouling is the unwanted deposition of materials, particles, and bacteria on the
66 membrane surface during the filtration process ¹⁵. Soluble microbial products (SMP),
67 bacteria and other colloidal particles first attach onto the membrane through adhesive
68 forces. Nutrients and other trace metals which are highly abundant in wastewater
69 streams then serve as substrates for the immobilized bacteria, resulting in the growth of
70 bacteria and contributing to the biofilm formation. The inherent resistance associated
71 with the biofilm layer would decrease water flux, increase energy consumption and
72 operational costs, and shorten the lifetime of membranes ¹⁶.

73 Considering the problems caused by biofouling, devising antifouling strategies has
74 become a widely researched topic. Examples of antifouling strategies include physical
75 membrane maintenance (e.g. backwash, chemical wash, sonication) ¹⁷, chemical
76 modification (e.g. nanoparticles coating, organic functional groups modification) ¹⁸ and
77 biocidal control (e.g. quorum quenching, phage-based decomposition) ^{19, 20}. Among
78 them, coating heavy metal nanoparticles on the surface of membrane has been widely
79 applied to inhibit fouling layer. It is thought that antibacterial effects associated with
80 heavy metals and nanoparticles ^{21, 22} would inhibit the initial attachment of bacteria to
81 membranes. Although nanoparticles have been widely applied to control biofouling, the
82 mechanism behind how nanoparticles decrease biofouling is relatively less understood
83 at the molecular level. Furthermore, it should be brought to attention that heavy metal
84 nanoparticles are themselves, a form of ECs. While coating membranes with heavy
85 metal nanoparticles to create antifouling membranes can potentially delay or eradicate
86 membrane biofouling, it may also result in unintended leaching of these ECs into the
87 biological processes. The presence of heavy metal nanoparticles in the biological system

88 can impede biological activity and in turn cause a potential detrimental impact on the
89 overall treatment efficiency.

90 This review therefore aims to understand how nanoparticles can potentially impact
91 MBRs. To achieve this aim, the review starts off by illustrating how nanoparticles induce
92 antibacterial effect on microorganisms. Past studies that illustrate the antibacterial
93 effect of nanoparticles against pure bacterial cultures and biofilm-associated
94 populations were collated and critically assessed. Finally, this review evaluates if the
95 presence of nanoparticles in wastewater treatment systems would affect the overall
96 performance of aerobic and anaerobic biological processes. This is done by assessing the
97 gene expression profiles of keystone bacterial or archaeal populations that are
98 contributing to the overall functionality of those biological processes upon exposure to
99 heavy metal nanoparticles.

100

101 **2. Antimicrobial mechanisms of nanoparticles**

102 There are four main mechanisms to account for the antimicrobial effects associated
103 with nanoparticles. First, the production of reactive oxygen species (ROS) induces
104 oxidative stress on microorganisms. Second, metal ion toxicity directly kills or affect
105 microbial cell growth. Third, nanoparticles accumulate on cells' surfaces and induce
106 toxicity. Lastly, internalization of nanoparticles within the cells induce cell death ²³⁻²⁷ (Fig.
107 1).

108 *2.1 ROS induced damage*

109 The toxicity of nanoparticles is frequently attributed to oxidative stress induced by ROS
110 such as hydroxyl radicals, hydrogen peroxide, superoxide ions, and singlet oxygen ²⁶⁻³³.
111 Different nanoparticles produce ROS under different conditions. Some metal oxide
112 nanoparticles (i.e., ZnO and TiO₂ nanoparticles) produce ROS upon UV or visible light
113 illumination ³⁴. However, for silver nanoparticles, it can spontaneously generate ROS
114 without illumination ³⁵.

115

116 The molecular mechanisms of ROS damage has been widely studied. ROS could induce
117 lipid peroxidation, which is a highly deleterious reaction. Furthermore, ROS can also
118 oxidize membrane-bound proteins, and then alter the integrity of cellular components.
119 Intracellular ROS in cytoplasm could lead to a series of damage to bacteria ³⁶. For
120 example, superoxide and hydrogen peroxide (H₂O₂) produced by nanoparticles can
121 oxidize 4Fe-4S clusters and other enzymes within the cytoplasm. In addition, hydrogen
122 peroxide can directly react with cysteine residues while hydroxyl radicals can damage
123 any biological molecules, including DNA, in a diffusion limited manner. These types of
124 damage collectively result in metabolic defects and eventually cell death ³⁶.

125 In order to protect themselves from oxidative stress induced by ROS, bacteria are
126 equipped with an array of antioxidant defenses that range from small, oxidant-
127 scavenging molecules to self-regulating gene networks ³⁷. ROS-induced damage would
128 be regulated by a cascade of antioxidants, including superoxide dismutases, catalases,
129 glutathione, peroxidase and some non-enzymatic antioxidants (i.e., α -tocopherol and
130 ascorbic acid) ³⁸. A triple mutant *sodABC E. coli* strain could not produce superoxide
131 dismutases, and was very sensitive to silver nanoparticles ³⁹, highlighting the role of this
132 enzyme in protecting against ROS induced by silver nanoparticles. Similarly, Xie and co-
133 authors showed that oxidative stress genes (i.e., *katA* coding catalase and *ahpC* coding
134 alkyl hydroperoxide reductase) were upregulated by 52- and 7-fold upon exposure to
135 ZnO nanoparticles ⁴⁰. These studies highlight that one of the main mechanisms behind
136 antimicrobial effect of heavy metal nanoparticles is through the production of ROS.

137 2.2 Metal ion toxicity

138 Heavy metals dissolve to release metal ions. When present in the ionic state, they
139 induce toxicity due to their affinity for cellular components and biomolecules, and the
140 subsequent formation of metal–biomolecule complexes. Metal ions can bind or block
141 the functional groups in a biological molecule. It can also displace the essential metals in
142 enzymes, and bind to the cellular thiol pool or participate in deleterious chemical
143 reactions within the cell. Ultimately, metal ions damage proteins, DNA and biological

144 membranes, and interfere in enzyme function and cellular processes ⁴¹. In addition,
145 toxicity of metal ions is normally accompanied by ROS-mediated toxicity. Bondarenko
146 and co-authors showed that Cu ions released from CuO nanoparticles was the key factor
147 in triggering the ROS and DNA damage ⁴². Ag nanoparticles can easily destroy the
148 membrane, pass through the microbial body, then convert to silver ions in cytoplasm to
149 damage the intracellular structure ⁴³. In one study, 19 different types of nanoparticles
150 were examined and it was determined that ions released from nanoparticles were
151 cytotoxic ⁴⁴. In another study, Zheng et al observed similar inhibitory effects on
152 microbial denitrification by either Zn ions or ZnO nanoparticles, hence indicating that
153 the toxicity of ZnO nanoparticlse was due to Zn ions release ⁴⁵.

154

155 However, discerning the mode by which dissolved ions and nanoparticles lead to
156 microbial cell death is difficult to prove experimentally and only few studies focused on
157 this. To illustrate, Neal et al. compared the effects of exposing sublethal concentrations
158 of ZnO nanoparticles and Zn ions to *Cupriavidus necator*, and results indicated
159 nanoparticles mainly affected membrane-associated proteins. In contrast, zinc ions
160 exposure affected metabolic processes, likely because Zn ions can act as important co-
161 factor for a number of enzymes including DNA and RNA polymerases ⁴⁶. Similar result
162 was observed when *Daphnia magna* was exposed to Ag nanoparticles and ions ⁴⁷. Ag
163 nanoparticles could disrupt the major biological processes including protein metabolism
164 and signal transduction, while the Ag ions disrupted developmental processes. It is likely
165 that both metal ions and nanoparticles affect different metabolic pathways due to
166 varying affinities towards certain biomolecules, but synergistically complement each
167 other to result in cell death.

168 *2.3 Nanoparticles accumulation on membrane surface*

169 The interaction between microbial cell wall and nanoparticles may lead to toxicity. It
170 was reported that nanoparticles accumulate on cell wall by complexing with
171 lipopolysaccharide and lipoteichoic acid through hydrogen bonding and ligand exchange.
172 This interaction can subsequently change protein structure ⁴⁸. Positively charged CeS₂

173 nanoparticles are electrostatically attached to the negatively charged bacterial
174 membranes at neutral pH and display strong electrostatic attraction toward bacterial
175 outer membrane ²⁶. Alternatively, the electrostatic interactions are attributed to
176 negatively charged phospholipids in cell membrane absorbing to positively charged
177 nanoparticles ³⁰.

178 The toxicity of nanoparticles accumulation was reviewed by Djuriscic et al. ²³. After
179 nanoparticles attach onto the cell surface, nanoparticles facilitate redox reactions and
180 result in oxidative stress. Hence the toxicity of nanoparticles accumulation can also be
181 attributed to ROS generation. In addition, Ivask et al. pointed out that nanoparticles
182 attachment may cause mechanical damage, such as change in cell morphology and
183 deformation of membranes, disorganization or leakage of intracellular structures ³⁸.

184

185 *2.4 Internalization of nanoparticles*

186 Normally, microorganisms have rigid cell wall structures which would inhibit the
187 nanoparticles from entering into the cell. However, nanoparticles can still be
188 internalized by non-specific diffusion ⁴⁹. It was observed that ZnO nanoparticles below
189 10 nm and Ag nanoparticles below 16 nm in diameter could enter into *E. coli* by non-
190 specific diffusion ³⁸. For those nanoparticles larger in diameter, it is likely that non-
191 specific diffusion through compromised membranes would play a more favorable role.
192 To exemplify, previous injuries induced by ROS, metal ions and nanoparticles
193 accumulation would possibly increase permeability of cell wall. Liu et al. found that ZnO
194 nanoparticles could lead to destruction of lipids and proteins on membrane of *E. coli*,
195 which then increase the amount of nanoparticles entering the cells ⁵⁰.

196

197 **3. Antibacterial effect of heavy metal nanoparticles against pure cultures**

198 Based on the known mechanisms by which heavy metal nanoparticles induce cell death,
199 many studies have hypothesized that gram-negative and gram-positive bacteria would
200 exhibit different susceptibilities towards nanoparticles (Table 1). Unlike gram-positive

201 bacteria, gram-negative bacteria has a thinner peptidoglycan layer ⁵¹. Besides, there is a
202 layer of negatively-charged lipopolysaccharide (i.e., O-polysaccharide, core
203 polysaccharide, lipid A, and proteins) at outer membrane layer of cell wall of gram-
204 negative bacteria ^{23, 51}, which may enhance the interaction between bacteria and
205 nanoparticles. These differences of gram-positive and gram-negative bacteria can
206 possibly result in different susceptibility to nanoparticles.

207 To evaluate this, Kim et al. tested the antibacterial effects of silver nanoparticles with
208 less than 20 nm diameter against yeast, *E. coli* and *Staphylococcus aureus*. The authors
209 showed that the minimum inhibitory concentration (MIC) of silver nanoparticles to *E.*
210 *coli* was only 3.3 nM to 6.6 nM, while higher MIC of > 33 nM would be required for *S.*
211 *aureus* ²¹. Contrasting results were however obtained by Azam and co-authors, who
212 evaluated the antibacterial effects of zinc oxide, copper oxide on both gram-positive (i.e.,
213 *Escherichia coli* and *Pseudomonas aeruginosa*) and gram-negative (i.e., *Bacillus subtilis*
214 and *S. aureus*) bacteria. Results showed that gram-negative *E. coli* had a 24% and 28%
215 smaller inhibition zone size than *B. subtilis*. Similarly, gram-negative *P. aeruginosa* had a
216 16% and 33% smaller inhibition zone size than *S. aureus* in the case of ZnO nanoparticles
217 and CuO nanoparticles, respectively ²². Nevertheless, these antibacterial effects were
218 only evaluated by disk diffusion method and the inhibition zones can be affected by
219 particles sizes and diffusion rate of nanoparticles or the dissolved ions in agar plate.
220 Another study conducted by Suresh and his co-authors suggest that gram-negative
221 bacteria (i.e., *E. coli* and *Shewanella oneidensis*) were more resistant to silver
222 nanoparticles than gram-positive bacterium (i.e., *B. subtilis*). This is done based on
223 several evaluation tests, including disk diffusion, minimum inhibitory concentrations,
224 live/dead assays, and atomic force microscopy ⁵².

225 Bacteria could resist stress from environment by regulating associated genes. The
226 reasons to account for these contradictory findings can be due to how different
227 bacterial species upregulate genes to cope against the antimicrobial effects. For
228 examples, nanoparticles can induce *S. aureus* to produce catalase, which could

229 neutralize the hydrogen peroxide and further decrease the toxicity of nanoparticles ⁵³. It
230 should be noted that the coping mechanisms of bacteria is complex - even different
231 bacterial strains in same species have different coping mechanisms under same
232 environmental stress. To illustrate, Al-Jassim and her co-authors exposed two different *E.*
233 *coli* strains to solar irradiation, and RNA-sequencing results indicated that these two
234 strains had different expression and regulation of genes ⁵⁴. It is therefore likely that
235 susceptibility towards nanoparticles is not simply governed by whether a bacterium is
236 gram-negative or gram-positive, but instead, also attributed by other factors (e.g.,
237 genomic contents and the ability to upregulate genes related to oxidative stress
238 response and repair mechanisms and so on).

239

240 **4. Antibacterial effect of heavy metal nanoparticles against biofilm**

241 Even though nanoparticles have demonstrated antibacterial activity against pure
242 cultures of gram-positive and gram-negative bacteria, biofilm attached on membranes
243 are comprised of complex microbial populations and extracellular polymeric substance
244 (EPS), and therefore make it hard to predict the effect on biofilm by heavy metal
245 nanoparticles. Generally, biofilm-associated bacteria should exhibit higher tolerance to
246 heavy metal NPs compared to planktonic cells. To illustrate, it was reported that 1 mg
247 Ag/L could inhibit most planktonic bacteria isolated from wastewater biofilm within 1 h.
248 In contrast, 24 h of treatment on biofilm with 200 mg Ag/L was needed to achieve
249 significant reduction of biofilm-associated bacteria ⁵⁵. In the case of anaerobic
250 membrane biofilm, same results have been observed. Reyes et al. indicated that the
251 growth rates and yields of *P. denitrificans* in biofilms exposed to Cu nanoparticles were
252 40-fold to 50-fold less inhibited than those in planktonic cells ⁵⁶.

253

254 In recent years, membranes coated with nanoparticles have been applied to control
255 biofouling in MBRs. For example, palladium nanoparticles on polysulfone membrane
256 was used to minimize bacteria number and EPS content in bacterial monoculture (i.e., *P.*

257 *aeruginosa*) and to also minimize EPS content in complex biofilm (i.e., sludge) ¹⁸. In
258 another study, polyvinylidene fluoride membrane coated with MnO₂ nanoparticles
259 could greatly reduce presence of bacteria and organic matter attached on the
260 membrane, resulting in a significantly thinner cake layer ⁵⁷. Similarly, Maximous et al.
261 immobilized ZrO₂ nanoparticles on polyethylsulfone membrane and then evaluated its
262 antifouling effect. They successfully demonstrated that 5% weight fraction of ZrO₂ with
263 polyethersulfone membrane has the lowest fouling rate ⁵⁸. Ag nanoparticles entrapped
264 on polysulfone membrane can delay pore clogging of membrane and decrease EPS
265 content in biofilm maxtrix, indicating an improvement in the antifouling effect ⁵⁹. The
266 schematic representation of antifouling mechanism is shown in Fig. 2. However, these
267 studies evaluated the potential antifouling effects by aerobic sludge, and it remains
268 unknown whether the nanoparticles would achieve the same effect on anaerobic sludge.

269

270 Furthermore, most of the studies only evaluate the efficacy of heavy metals by
271 measuring parameters related to the phenotypic characteristics of the biofilm and did
272 not study the effect on biofilm formation at the molecular level. In the subsequent
273 sections, we introduce genes and enzymes involved in biofilm formation so as to provide
274 a basis for future studies to determine if these would be affected by nanoparticles.

275

276 *4.1 Initial surface attachment*

277 Biofilm formation relies on bacteria to first attach on a surface. Previous studies indicate
278 that outer surface appendages (e.g. flagella, pili, curli) and outer membrane adhesion
279 proteins can contribute to the initial biofilm attachment process ⁶⁰. To exemplify, the
280 conjugation pilus acts as an adhesion factor for cell-cell and cell-surface interaction. It is
281 widely accepted that initial bacterial adhesion is a key part of the biofilm development
282 process ⁶¹. However, in a high transmembrane pressure system like an MBR, the initial
283 attachment is also contributed by filtration pressure and forces, and the subsequent
284 adsorption of bacteria onto the inorganic or organic constituents in a cake layer present
285 on the membrane.

286

287 4.2 Microcolony formation

288 After initial attachment, bacteria would form highly structured cell cluster, which is also
289 called microcolony⁶². Two processes, namely bacterial motility and clonal growth
290 contribute to microcolony formation. Bacterial motility are dependent on flagella, pili
291 structures of bacteria, and chemotaxis. For example, type IV pilus of *Pseudomonas*
292 *aeruginosa* is important for biofilm formation⁶³. Type IV pilus could mediate twitching
293 motility for the bacteria to move from one location to another. Similar to type IV pili,
294 CupA fimbriae, encoded by three *cup* clusters, can also mediate cell-surface interactions
295⁶⁴.

296 In addition, EPS, the predominant “glue” in a biofilm matrix, is required for microcolony
297 formation. Bacteria secrete exopolysaccharides. Using *P. aeruginosa* as an example, this
298 bacterium could produce three kinds of polymers (i.e., alginate, Pel, and Psl), which are
299 correspondingly regulated by *alg*, *pel* and *psl* genes⁶⁵. *Staphylococcus aureus* and
300 *Staphylococcus epidermidis* could also produce polysaccharides intercellular adhesion
301 (PIA) which is encoded by PIA synthesis gene (*icaADBC* operon)⁶³. It is likely that such
302 production of polysaccharides are ubiquitous in most microorganisms. To illustrate, *E.*
303 *coli* has the *pga* gene, which shares high sequence similarity with *ica* genes, and is found
304 to be related to polymer production that facilitates biofilm formation⁶⁶. *hms* genes,
305 which shared high sequence homology to *ica* genes, have also been identified in *Yersinia*
306 *pestis*⁶⁷.

307

308 4.3 Biofilm maturation

309 Once the bacteria attach and microcolony is initiated, the usual metabolic processes
310 occur among the microorganisms since an MBR treating municipal wastewater is a
311 nutrient-rich ecosystem. Bacterial cells are not substrate-limited and would reproduce
312 to a high cell density. Microorganisms then perform cell-to-cell communication (i.e.,
313 quorum sensing). Gram-positive and gram-negative bacteria utilize quorum sensing
314 communication circuits to regulate or coordinate certain traits and activities.

315

316 In *P. aeruginosa* and many other gram-negative bacteria, N-acyl homoserine lactones
317 (AHLs) are produced. There are two AHLs signaling systems, *lac* and *rhl*, that are
318 relatively well-studied. The *lac* system contains the signal synthase LasI and associated
319 signal receptor LasR. Similarly, in *rhl* system, bacteria could synthesize signals by
320 synthase RhlI and regulate associated genes expression by receptor RhlR⁶⁸. It was
321 reported that biosilver nanoparticles could reduce the levels of LasIR-RhIR and reduce
322 AHLs production⁶⁹. However, QS involves other signal molecules like autoinducer-2 (AI-
323 2) and autoinducer peptides (AIP). Less is known about how heavy metal nanoparticules
324 would affect cell-to-cell signaling via these other molecules.

325

326 **5. Impacts induced by nanoparticles on performance of aerobic and anaerobic** 327 **biological processes.**

328 MBR has been widely applied to treat wastewater. Aerobic MBR couples a membrane-
329 based filtration process with activated sludge to achieve nitrogen, phosphorus and COD
330 removal⁷⁰. Similarly, anaerobic MBR couples a membrane-based filtration process with
331 anaerobic digester to convert organic carbon to biogas. Although coating heavy metal
332 nanoparticles on polymer-based organic membrane could mitigate biofouling, one has
333 to consider the potential impact on biological processes if the nanoparticles were to
334 leach into the system. It should be taken into account that the nanoparticles are
335 classified as one of the new emerging contaminants, and it would pose a challenge to
336 the overall functionality of MBR². Here, we will briefly introduce how the nanoparticles
337 affect the biological nitrogen removal, biological phosphorus removal and anaerobic
338 fermentation process.

339 *5.1 Nitrogen removal*

340 Untreated wastewater contains excessive nitrogen compounds which would cause
341 adverse effects (e.g. eutrophication) if directly discharged into environmental waters.

342 The nitrogen compounds include ammonium (NH_4^+), nitrite (NO_2^-), nitrate (NO_3^-) and
343 other dissolved organic nitrogen. In biological aerobic processes, ammonium is oxidized
344 to hydroxylamine by ammonia-oxidizing bacteria via ammonia monooxygenase (*amo*),
345 and subsequently to nitrite (hydroxylamine oxidoreductase, *hao*) and nitrate (nitrite
346 oxidoreductase, *nxr*) by nitrifiers. The nitrate is then reduced to nitrite (nitrate
347 reductase, *nap* and *nar*), nitric oxide (nitrite reductase, *nirK* and *nirS*) and nitrous oxide
348 (nitric oxide reductases, *qNor* and *cNor*) and finally nitrogen gas (nitrous oxide reductase,
349 *nos*) via denitrifiers ⁷¹.

350 In the past years, many studies have reported the effects of nanoparticles on nitrogen
351 removal. To illustrate, it was reported that alumina nanoparticles would affect the
352 nitrogen removal after both short-term and long-term exposure ⁷². The study found that
353 50 mg/L alumina nanoparticles would affect nitrification and denitrification during
354 short-term exposure. In contrast, long-term exposure only led to decrease in
355 denitrifying bacteria and denitrifying enzymatic activities (i.e., NAP, NAR and NIK). These
356 decreases then resulted in a deterioration of total nitrogen removal from 80.4% to
357 62.5%. Zheng et al. spiked TiO_2 nanoparticles into anaerobic-anoxic sequencing batch
358 reactor, and found that the total nitrogen removal also decreased from 80.3% to 24.4%
359 after 70 days of exposure. Further examination revealed TiO_2 nanoparticles decrease
360 the relative abundance of nitrifying bacteria, especially ammonia-oxidizing bacteria, and
361 simultaneously inhibited the activities of ammonia monooxygenase and nitrite
362 oxidoreductase ⁷³. Similarly, zinc oxide and copper nanoparticles also affected the
363 denitrification process and led to nitrous oxide accumulation via inhibitory effect on the
364 catalytic activities of key denitrifying enzymes after long-term exposure ^{45, 74}.
365 Nevertheless, not all nanoparticles showed inhibitory effects to nitrogen removal. For
366 instance, 200 $\mu\text{g/L}$ gold nanoparticles reported no influence on ammonia removal and
367 *amo* gene abundance in a sequencing batch reactor, even though such nanoparticles
368 changed microbial community in reactor ⁷⁵. It is likely that the types and concentrations
369 of nanoparticles affect the outcome on nitrogen cycle.

370 5.2 Phosphorus removal

371 Phosphorus can usually be removed from municipal wastewater by enhanced biological
372 phosphorus removal (EBPR) method. This process is operated in an anaerobic/aerobic
373 condition via the enrichment of phosphorus accumulating organisms (PAOs). In general,
374 PAOs are capable of storing organic compounds, such as short chain fatty acids (SCFA)
375 and glycogen as internal storage compounds (i.e., polyhydroxyalkanoates (PHA)) by
376 utilizing energy obtained from hydrolysis of the intracellular stored polyphosphates
377 (poly-P). Ortho-phosphate is released into wastewater under anaerobic condition.
378 However, in aerobic condition, PAOs are able to rapidly take in excess phosphorus and
379 store them as intracellular poly-P by using stored PHAs as the energy source. The net
380 phosphorus removal can be achieved by wasting excess sludge with high phosphorus
381 content because the aerobic phosphorus uptake is greater than the anaerobic
382 phosphorus release.

383 Similar to nitrogen removal, nanoparticles have been reported to affect phosphorus
384 removal. Hu et al. evaluated the effects of ZnO nanoparticles on phosphorus removal
385 based on one anaerobic-aerobic sequencing batch reactor, and results indicated long-
386 term exposure to 6 mg/L ZnO nanoparticles would significantly deteriorate phosphorus
387 removal via inhibitory effects on *Accumulibacter* and *Competibacter*⁷⁶. Zheng and his
388 co-authors spiked 50 mg/L ZnO nanoparticles into anaerobic-anoxic sequencing batch
389 reactor. The phosphorus concentration in effluent increased from non-detectable to
390 16.5 mg/L, indicating a decrease in phosphorus removal efficiency. The authors
391 attributed the deterioration to inhibitory effects on PAOs, exopolyphosphatase (i.e.,
392 PPX, required to hydrolyze polyphosphates) and polyphosphate kinase (i.e., PPK,
393 required to convert orthophosphates to intracellular poly-P)⁷⁷. Nevertheless, not all
394 nanoparticles cause deterioration in phosphorus removal. To illustrate, Chen and his co-
395 authors spiked alumina nanoparticles into anaerobic-anoxic sequencing batch reactor
396 and found that long-term exposure did not affect the phosphorus removal⁷². Similarly,
397 50 mg/L TiO₂ nanoparticles was spiked in the same kind of reactor, and no inhibition on

398 biological phosphorus removal was observed after 70 days of exposure. Furthermore,
399 the PPX and PPK activities were also not affected ⁷³. Hence, similar to the nitrogen cycle,
400 it is likely that the types and concentrations of nanoparticles would impact phosphorus
401 removal in a different manner.

402 *5.3 Anaerobic fermentation*

403 Anaerobic membrane bioreactor (anMBR) couples a membrane-based filtration process
404 with anaerobic fermentation, which has lower sludge production rates than activated
405 sludge processes, and can achieve high COD removal. Anaerobic fermentation not only
406 eliminates the need for aeration, but also generates methane as an energy source ⁷⁸.

407 Anaerobic fermentation is mainly divided into four major steps, including hydrolysis,
408 acidogenesis, acetogenesis, and methanogenesis ⁷⁹. Representative keystone bacterial
409 and archaeal species of each step are shown in Fig. 3. In this section, we briefly
410 introduce related genes involved in anaerobic fermentation and methanogenesis which
411 would possibly be affected by nanoparticles.

412 *5.3.1 Hydrolysis*

413 During the hydrolysis process, a consortia of bacteria break down the complex
414 compounds (e.g. carbohydrates, protein, lipids) into smaller fractions (e.g. fatty acids,
415 amino acids, polysaccharides etc.). The microorganisms involved in hydrolysis include
416 *Clostridium thermocellum*, *Bifidobacterium longum*, *Clostridium cellulolicum*,
417 *Bacteroides thetaiotaomicron*, *Enterococcus faecalis*, *Bacteroides capillosus* ⁸⁰. There are
418 four main classifications of hydrolases that would contribute to this step, namely lipases
419 (EC 3.1.1), phosphatases (EC 3.1.3), glucosidases (EC 3.2.1) and proteases (EC 3.4).

420 Hydrolysis is typically defined as the rate limiting step of the whole anaerobic
421 fermentation since it is the first step of the whole process. This means increasing
422 hydrolytic rate could improve the overall anaerobic fermentation rate. To illustrate, Yu
423 and his coauthors enriched endogenous amylase and then added these enzymes into
424 anaerobic digester. It was observed endogenous amylase could increase volatile fatty

425 acids (VFAs) concentration in supernatant after 7 h treatment, and could increase biogas
426 production after 11 days of anaerobic digestion compared to the control ⁸¹.

427 In presence of nanoparticles, deleterious effect on the overall anaerobic fermentation
428 process were observed because nanoparticles affected the hydrolytic rates. To illustrate,
429 Mu and his co-authors showed that ZnO nanoparticle decreased the bovine serum
430 albumin (BSA) and dextran degradation efficiencies by 37.5% and 12.3%, respectively.
431 This suggests that nanoparticles can inhibit hydrolysis of protein and polysaccharide ⁸²,
432 both of which are important rate limiting steps in anaerobic fermentation.

433

434 5.3.2 Acidogenesis

435 Hydrolytic and acidogenic microorganisms are about ten times faster growing than
436 methanogens ⁸³. Fermentative acidogenic microorganisms (e.g. *Slackia*
437 *heliotrinireducens*, *Candidatus Cloacamonas acidaminovorans*, *Clostridium kluyveri*,
438 *Clostridium acetibutylicum*, *Clostridium perfringens*, *Clostridium saccharolyticum*,
439 *Caldanaerobacter subterraneus*, *Finegoldia magna*, *Enterococcus faecium*, *Lactobacillus*
440 *helveticus*) could further convert the hydrolyzed products into VFAs (e.g. formic acid,
441 acetic acid, propionic acid, butyric acid) ⁸⁰. Most of the VFAs are formed in the
442 metabolism of glucose via the glycolytic Embden-Meyerhof-Parnas (EMP) pathway ⁸³.
443 Hydrolyzed glucose is decomposed to produce pyruvic acid. The pyruvic acid
444 fermentation then produces a number of VFAs. The production of VFAs by anaerobic
445 bacteria is mediated by several kinds of enzymes. To illustrate, butyrate synthesis
446 requires crotonase while alcohol synthesis requires alcohol dehydrogenase.
447 Methylmalonyl-CoA mutase catalyzes the isomerization between methylmalonyl-CoA
448 and succinyl-CoA, and are involved in propionate synthesis ⁸⁴. Certain nanoparticles
449 were shown to inhibit acidogenesis process. It was reported that 150 mg/L ZnO
450 nanoparticles could inhibit 40-80% acidogenesis efficiency of waste active sludge⁸⁵. In
451 addition, Cu nanoparticles added into fermentation system could inhibit hydrolysis and
452 acidification and then result in VFAs reduction ⁸⁶.

453

454 5.3.3 Acetogenesis

455 In a well-functioning anaerobic fermentation process, VFAs, especially long chain VFAs
456 do not accumulate but are instead converted into acetic acid, carbon dioxide and
457 hydrogen by the acetogenic microorganisms (e.g. *Carboxydotherrmus hydrogenoformans*,
458 *Morella thermoacetica*, *Pelotomaculum thermopropionicum*). Acetogenic
459 microorganisms are strict anaerobes since they utilize the acetyl coenzyme A pathway
460 that contains enzymes sensitive to oxygen. It was reported that phospho-transacetylase,
461 acetate kinase (AK) and acetyl-CoA synthase were involved in the acetogenesis step ⁸⁴.
462 For example, acetate kinase (AK) is responsible for the transformation of
463 monosaccharides and amino acids into acetate. In addition, homoacetogens (a
464 subpopulation of acetogens) convert hydrogen and carbon dioxide to acetate ⁸⁷.
465 Homoacetogens rely on pyruvate ferredoxin oxidoreductase, CO dehydrogenase/acetyl-
466 CoA synthase, tetrahydrofolate-dependent enzymes, methyltransferase (e.g. MeTr, AcsE)
467 and corrinoid iron sulfur protein (CFeSP, AcsCD), all of which are enzymes involved in
468 Wood-Ljungdahl pathway, to achieve this conversion.

469

470 It was reported that both Ag NPs and MgO NPs at the dosage of 500 mg/g total
471 suspended solids could inhibit acetogenesis by significantly decreasing the AK activity ⁸⁸.
472 In contrast, certain concentrations of NiO and CoO nanoparticles (i.e., 1.5 mg/L and 1.0
473 mg/L, respectively) could enhance the acetic acid and butyric acid concentration, and
474 finally increase the hydrogen yield by 1.51 and 1.67 fold respectively. This is achieved
475 through enhancing electron flow efficiency and enzyme activities in both acidogenesis
476 and acetogenesis processes ⁸⁹.

477

478

479 4.3.4 Methanogenesis

480 In methanogenesis, *Methanosarcina barkeri*, *Methanosarcina acetivorans*,

481 *Methanoculleus marisnigri*, *Methanoregula boonei*, *Methanosphaerula palustris*,
482 *Methanospirillum hungatei*, *Methanoplanus petrolearius*, *Methanocorpusculum*
483 *labreanum* were reported to be involved in anaerobic digestion to produce methane ⁸⁰.

484 The two distinct methanogenic pathways are from H₂/CO₂ to methane
485 (hydrogenotrophic pathway) and from acetate to methane (aceticlastic
486 methanogenesis). Hydrogenotrophic methanogenesis converts CO₂ and H₂ to methane.
487 The H₂ and formate are the major electron donors. According to the KEGG-Orthology,
488 there are seven genes involved in this process, including formate dehydrogenase (*fdh*,
489 EC: 1.2.1.2), formylmethanofuran dehydrogenase (*fwdA/fmdA*, EC: 1.2.99.5),
490 formylmethanofuran-tetrahydromethanopterin formyltransferase (*ftr*, EC: 2.3.1.101),
491 methenyltetrahydromethanopterin cyclohydrolase (*mch*, EC: 3.5.4.27), oxidoreductases
492 (*mtd*, EC: 1.5.99.9), coenzyme F420 hydrogenase (*frhB*, EC: 1.12.98.1) and coenzyme
493 F420-dependent N5,N10-methenyltetrahydromethanopterin reductase (*mer*, EC:
494 1.5.99.11).

495 In aceticlastic methanogenesis, acetate is directly converted to methane. The enzymes
496 involved include acetate kinase (*ackA*, EC: 2.7.2.1), phosphate acetyl-transferase (*pta*,
497 EC: 2.3.1.8), acetyl-CoA decarbonylase/synthase complex (ACDS; composed of CdhA1,
498 CdhB, CdhD, CdhE and CdhC), and acetyl-CoA synthetase (*acs*, EC: 6.2.1.1). Besides, the
499 two modes of methanogenesis share the same enzymes and genes (i.e.,
500 Tetrahydromethanopterin S-methyltransferase (*mtr*, EC: 2.1.1.86), methyl coenzyme M
501 reductase (*mcr*, EC: 2.8.4.1) and reductase heterodisulfide reductase (*Hdr*, EC: 1.8.98.1)),
502 which are required to obtain methane as final product ⁹⁰.

503 Despite knowing the genes and pathways involved in anaerobic fermentation process,
504 studies that reported the effects of heavy metal nanoparticles on anaerobic MBR
505 performance mainly rely on end point measurements of biogas and VFAs production but
506 not on gene expressions. To illustrate, Luna-delRisco and the co-authors added 15 mg/L
507 CuO nanoparticles and 120 mg/L ZnO nanoparticles to anaerobic digester of cattle
508 manure for 14 days, and found that methane production was severely affected. The

509 authors further estimated that the half effective concentration (i.e., EC₅₀) for methane
510 was 10.7 mg Cu/L and 57.4 mg Zn/L⁹¹. Mishra et al. indicated optimal concentration of
511 NiO and CoO nanoparticles (i.e., 1.5 mg/L and 1.0 mg/L, respectively) could enhance
512 hydrogen yield in anaerobic digestion of palm oil mill effluent, suggesting that methane
513 production is also affected which would result in accumulation of hydrogen gas. The
514 author attributed this to be due to nanoparticles enhancing catalytic effect of
515 hydrogenases enzymes, but did not verify the effect on enzymes related to
516 methanogenesis⁸⁹.

517 In another long-term exposure study, different concentrations of ZnO nanoparticles
518 were spiked into anaerobic digester, and it was determined that 30 mg of ZnO and 150
519 mg of ZnO per gram of total suspended solids could inhibit about 18.3% and 75.1% of
520 methane production, respectively, when compared to the control. It was further shown
521 that the release of Zn ions from Zn nanoparticles may inhibit hydrolysis and
522 methanogenesis steps of anaerobic digestion by decreasing the activities of protease
523 and coenzyme F₄₂₀ associated with archaeal methanogens⁸² (Table 2).

524 Besides affecting activities of archaeal methanogens, exposure to heavy metal
525 nanoparticles could also result in changes in the abundance of specific microbial groups.
526 To exemplify, it was reported that when landfill solids were exposed to 10 mg Ag per kg
527 of solids at an exposure time of 250 days, biogas production was reduced. These
528 observations correlated with a decrease in the total copy numbers of 16S rRNA gene of
529 methanogens from $1.97 \pm 0.21 \times 10^7$ to $5.79 \pm 2.83 \times 10^5$ ⁹². However, this study
530 determined abundance of methanogens by 16S rRNA genes, which may not relate to
531 absolute abundance since some methanogens may contain > 1 copy of 16S rRNA gene
532 per cell⁹³. The decrease in methanogen-related 16S rRNA gene copies may be due to
533 changes in specific methanogenic community in response to the nanoparticles.

534 Despite the earlier discussed studies, there still remain some contradictory observations
535 on whether heavy metal nanoparticles would indeed affect methanogenesis. To
536 exemplify, Garcia et al. reported that the Ag nanoparticles can lead to about 33%-50%

537 decrease in biogas production, while Au nanoparticles showed only slight or no
538 inhibition to biogas production after 50 days exposure ⁹⁴. The impact of CeO₂-NPs on the
539 acidogenesis and methanogenesis of anaerobic granule sludge and anaerobic flocculent
540 sludge was also studied ⁹⁵. High concentrations of CeO₂ nanoparticles (i.e., 150 mg CeO₂-
541 NPs/g-VSS) could inhibit 19% VFAs production in flocculent sludge and 35% VFAs
542 production in granular sludge, but showed no inhibition to methane production. The
543 toxicity of ceria nanoparticles to anaerobic sludge was attributed mainly to direct
544 contact and interaction between nanoparticles with cell membrane. This finding
545 indicated that the acidogenesis was more sensitive to CeO₂ nanoparticles than
546 methanogenesis. Since acidogenesis is mediated by bacterial populations while
547 methanogenesis is mainly contributed by archaeal populations, it is inferred that
548 archaea is less susceptible to nanoparticles than bacteria when the main toxicity
549 mechanism is nanoparticle-cell interaction, possibly due to the fact that archaea has a
550 cell wall structure different from that of bacteria ⁵¹. More studies detailing how heavy
551 metal nanoparticles would impact anaerobic fermentation and methane production
552 were listed in Table 2.

553 It should be noted that not all nanoparticles result in negative effects to anaerobic
554 fermentation. This is especially so for Fe and other heavy metals introduced at sub-
555 lethal dosage. For example, Fe nanoparticles can enhance methane production by
556 enhancing immobilization of other metals, hence decreasing the potential toxicity of
557 solubilized heavy metals on anaerobic fermentation process ⁹⁶. Alternatively, Fe acts as
558 an electron donor to reduce oxidation/reduction potential (ORP) and as a buffer to
559 control pH through reduction of protons to hydrogen gas, which would facilitate
560 methanogenic activity. As such, the overall effect of heavy metal nanoparticles on MBR
561 process would be dependent on the type of metal and the concentration exposed to the
562 microbial consortium ⁹⁷.

563

564 **5. Conclusion**

565 Heavy metal nanoparticles exhibit high antibacterial effects through production of
566 reactive oxygen species (ROS), metal ion dissolution, accumulation on cell surfaces and
567 internalization within cells. Therefore, nanoparticles can be coated on membranes
568 surface to achieve antifouling effect in MBR. Many studies have applied the
569 nanoparticles-coated membranes to lab-scale MBR, and demonstrated promising results
570 in terms of high antibacterial effect and reduction in transmembrane pressures.
571 However, nanoparticles from such coated membranes can leach into the system and
572 impact the overall performance of MBR (e.g. biological nitrogen removal, biological
573 phosphorus removal, and fermentation). A review on related papers revealed
574 contradictory conclusions on how nanoparticles would affect these biological processes,
575 especially anaerobic fermentation and/or anaerobic MBR. The contradictory outcomes
576 reported by the various studies are possibly explained by the differences in heavy
577 metals tested, concentrations of heavy metal nanoparticles spiked into the system
578 and/or differences in the microbial consortium. Therefore, depending on the intended
579 purpose of using heavy metal nanoparticles in MBR processes, for example if the
580 purpose is for inhibiting membrane biofouling process while not detrimentally
581 affecting the overall functionality of the reactor system, one should first determine the
582 lethal dosage for that particular heavy metal nanoparticles and for that particular
583 microbial consortium. Subsequently, it would be beneficial to understand the microbial
584 consortium and map out the gene pathways affected by the nanoparticles so as to
585 ensure that undesirable and unintended impact on the overall functionality of the
586 reactor systems are minimized.

587

588 **Conflicts of interest**

589 There are no conflicts to declare.

590

591

592

593 **Acknowledgments**

594 This study is supported by KAUST Center Applied Research Funding URF/1/1971-32-

595 awarded to P.-Y. Hong.

Table 1. List of studies demonstrating antibacterial effect of heavy metal nanoparticles.

Type of heavy metal	Form	Test methods	Bacteria	Outcome	References
Ag	Nanoparticles (NPs)	Cell density on Muller Hinton agar plates in the presence/absence of the NPs	<i>Escherichia coli</i> (gram -ve) <i>Staphylococcus aureus</i> (gram +ve)	MIC of <i>E. coli</i> > 3.3 nM MIC of <i>S. aureus</i> > 33 nm	21
ZnO CuO Fe ₂ O ₃	Metal oxides NPs	Well-diffusion method on agar to observe for zone of inhibition; broth-dilution method and viable bacterial cell counts	<i>E. coli</i> <i>Pseudomonas aeruginosa</i> (gram -ve) <i>S. aureus</i> <i>Bacillus subtilis</i> (gram +ve)	Antibacterial activity (in ug/mL) and in decreasing order of efficacy: ZnO > CuO > Fe ₂ O ₃ ; Based on zone of inhibition, observed higher tolerance against tested NPs for gram-ve vs. gram+ve	22
CoO	Metal oxides thin nanosheets	Broth cultivation and measurement of optical density at 600 nm to generate growth curves	<i>E. coli</i> <i>P. aeruginosa</i> <i>Micrococcus luteus</i> (gram +ve) <i>S. aureus</i>	More effective against gram +ve (MIC > 50 ug/mL) than gram -ve (MIC > 100 ug/mL)	98
ZnO TiO ₂ CuO Cu ₂ O Ag	NPs	Well-diffusion method on agar to observe for zone of inhibition; activity and cationic assays (ATP, K+, H+)	<i>E. coli</i> <i>Enterococcus hirae</i> (gram +ve) grown anaerobically	Only Ag NPs showed antibacterial effect, most likely by affecting the proton and potassium ion fluxes	99
CuO	NPs	Broth cultivation and measurement of optical density at 600 nm to generate growth curves; Reactive oxygen species measurement; DNA damage and cell morphology characterization	<i>E. coli</i> <i>Proteus vulgaris</i> <i>M. luteus</i> <i>B. subtilis</i>	Sheet shaped CuO NPs were more effective on gram +ve; spherical shaped CuO NPs were more active on gram -ve	100
ZnO	NPs	Diffusion method on agar to observe for zone of inhibition; broth-dilution method and viable bacterial cell counts	<i>Salmonella enterica serotype Typhi</i> (gram -ve)	Antibacterial effect observed, activity increased by reducing size of NPs or by increasing the molarity of NPs	101
ZnO	NPs	Growth inhibition measured using colony counts and optical density at 600 nm; Reactive oxygen species measurement; DNA fragmentation, and membrane permeability	<i>Vibrio cholera</i> strains O395 and N16961 (gram -ve)	Antibacterial effect observed, ROS generation and induction of DNA damage are accounted as main antibacterial mechanisms; variation in response depending on the strain type	102
Fe ₂ O ₃	NPs; functionalized with -NH ₃ ³⁺	Broth dilution method; cell viability assay using Live/Dead BacLight kit; crystal violet staining on biofilm	<i>B. subtilis</i>	Biofilm formation affected without detrimental impact on cell viability	103
Al ₂ O ₃	NPs	Culture-based viability test; Reactive oxygen	Alga (<i>Pseudokirchneriella</i>	CuO and ZnO exhibit the highest toxicity (<1	104

Co ₃ O ₄	species measurement	<i>subcapitata</i> ,	mg/L)
CuO		Bacteria	
Fe ₃ O ₄		(<i>V. fischeri</i> , <i>E. coli</i> , <i>S. aureus</i>),	
MgO		Protozoa (<i>Tetrahymena</i>	
Mn ₃ O ₄		<i>thermophila</i>)	
Sb ₂ O ₃			
SiO ₂			
ZnO			
TiO ₂			
WO ₃			
Pd			

Table 2. List of studies detailing the effects of heavy metal nanoparticles on anaerobic fermentation

ZnO TiO ₂	NPs	Cell viability; proteomics and biochemical assays	<i>B. subtilis</i>	130 µg/mL TiO ₂ and 170 µg/mL ZnO stressed the cells and reduced viability by 50%; proteins related to central metabolism were affected by ZnO but not by TiO ₂	105
Ag	NPs immobilized in PVDF	Well-diffusion method on agar to observe for zone of inhibition; Filtration based colony count and SEM observation.	<i>E. coli</i>	0.5%-1.5% Ag showed clear halo zones and inhibited bacteria. Few bacteria grew on the membrane surface.	106
Cu	NPs stabilized on acrylic polymer	Broth cultivation and measurement of optical density at 600 nm to generate growth curves, Atomic absorption spectroscopy applied to quantify copper ions release.	<i>Chlamydomonas</i> sp. strain CD1 Red. <i>Synechocystis</i> sp. PCC 6803, <i>Phaeodactylum tricornutum</i> CCMP 1327	All Cu NPs modified membrane exhibit good antibacterial activity.	107
Cu	NPs load on thin-film composite polyamide membrane	Viable cell attachment experiment: Cell viability measured using colony counts; SEM observed cells attached.	<i>E. coli</i> <i>P. aeruginosa</i> <i>S. aureus</i>	1 h contact decreased about 87% ± 0.2% <i>E. coli</i> , 96% ± 3% <i>P. aeruginosa</i> and 79.5% ± 13% <i>S. aureus</i> live cells attached on membrane.	108

Nanoparticles	NPs diameter (nm)	Dosage	Reactor type	Exposure time (days)	Impact on anaerobic digestion	Reference
Ag	29	40 mg/L	Anaerobic digestion	14	No negative effects in biogas and methane production. Dominant methanogens remained largely unchanged.	109
Ag	21	1 /10 mg Ag/kg TS	Landfill digestion	>250	High concentration Ag NPs reduce biogas production rate, accumulate volatile fatty acid (VFAs), decrease methanogens number.	92
Ag/Au	30/10	N.A	Anaerobic digestion	50	Ag NPs decreased biogas production	94
CeO ₂	<25	5-150 mg/g VSS	Anaerobic granule/flocculent sludge	6	150 mg/g VSS CeO ₂ NPs inhibited acidification process, resulting in less short-chain fatty acid through physical penetration and membrane reduction. However, there was no observed inhibition on methanogenesis. CeO ₂ NPs has less influence on flocculent than granular sludge.	95
Cu	20-40	50 mg/L	Sequencing batch reactors followed with anaerobic digestion	>121	Releasing NPs through wastewater improved sludge solubilization, showed no effect on VFAs production. Directly adding NPs to anaerobic digestion inhibited hydrolysis and acidification, resulting in VFAs decreasing.	86
CuO	37	1.4 mg/L	Upflow Anaerobic Sludge Blanket	107	Short-term exposure caused minor inhibition to methanogenesis, while extended exposure caused severe toxicity and reduced 85% of the acetoclastic methanogenesis. >50% reduction in methane production.	110
Fe/ Fe ₃ O ₄	50/20	N.A.	Anaerobic digestion	12	Both NPs improved biogas production.	96
Fe	119	1.25 g/L	Anaerobic batch reactor	10	Nanoscale zero valent iron increased 28% in methane production via directly decreasing reactor's oxidation/reduction potential.	97
NiO / CoO	14 / 16.8	0-3 mg/L	Anaerobic digestion	4	1.5 mg/L NiO and 1.0 mg/L CoO NPs enhanced VFAs accumulation and hydrogen yield, also increase the COD removal.	89
ZnO	<100	0.32 / 34.5 mg Zn/L	Upflow Anaerobic Sludge Blanket	165	High ZnO NPs concentration resulted in rapid deterioration of methane production and VFAs removal; Deterioration more gradual and partial in presence of low ZnO NPs concentration; acetoclastic methanogens severely inhibited	111
ZnO	140	1-150 mg/g TSS	Anaerobic digestion	>150	High concentration ZnO NPs inhibited methane production due to Zn ²⁺ inhibiting hydrolysis and methanogenesis.	82

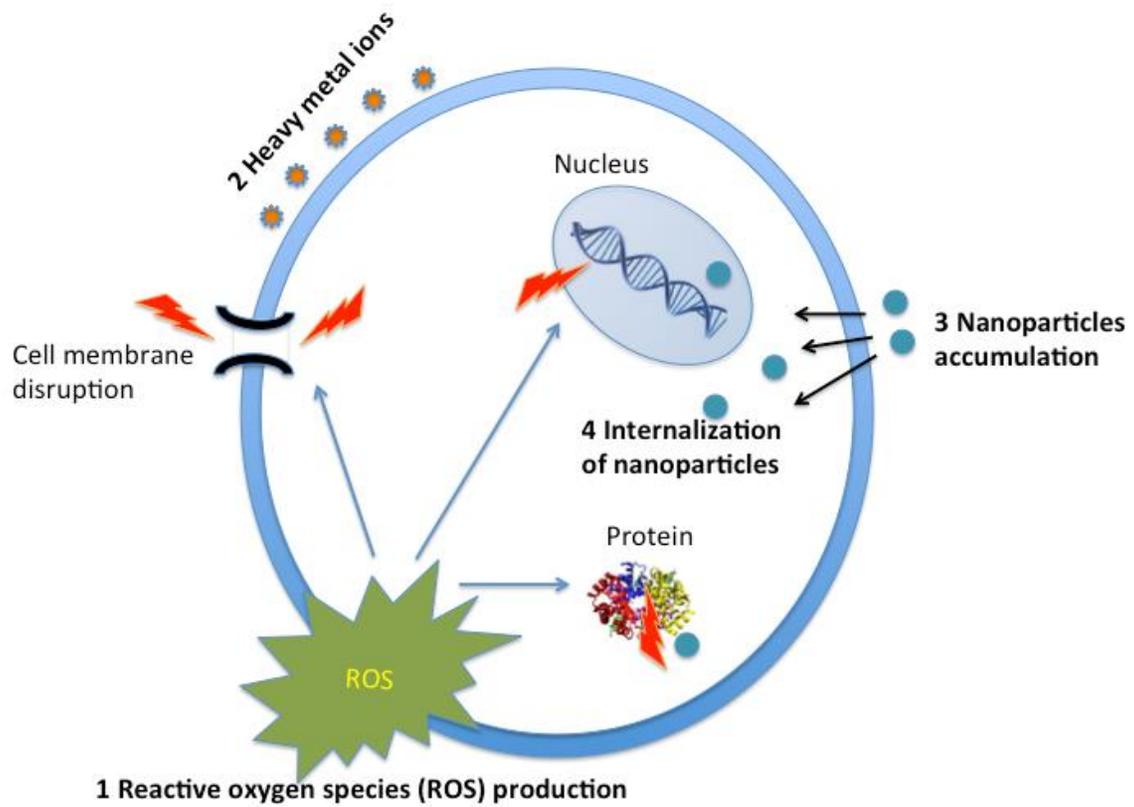


Fig. 1 Schematic representation of antibacterial effects induced by heavy metal nanoparticles.

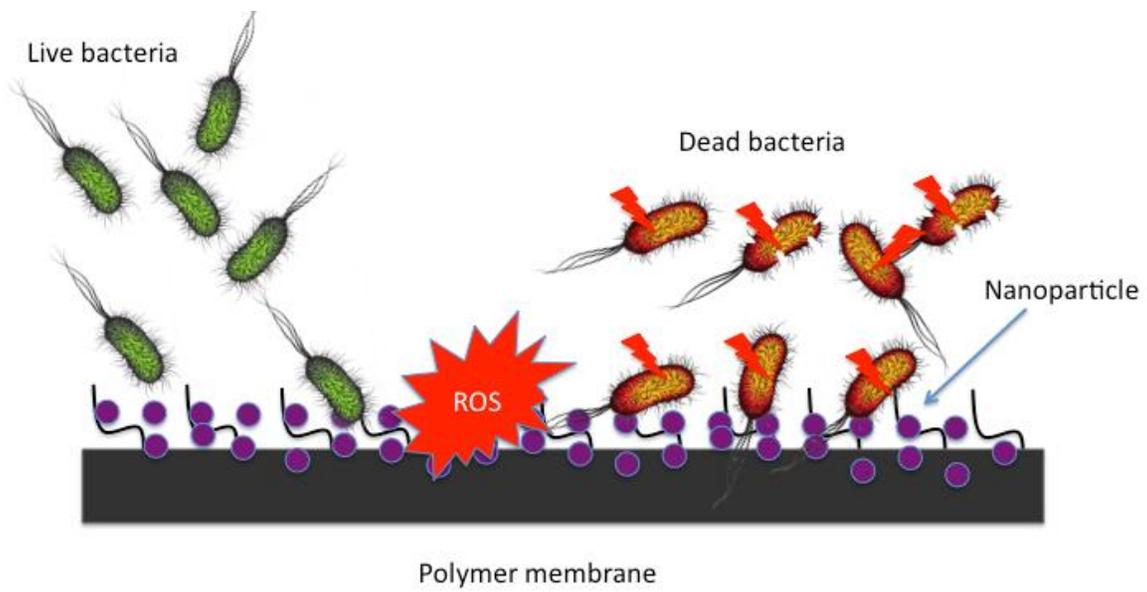


Fig. 2 Schematic representation of antifouling mechanism by heavy metal nanoparticles.

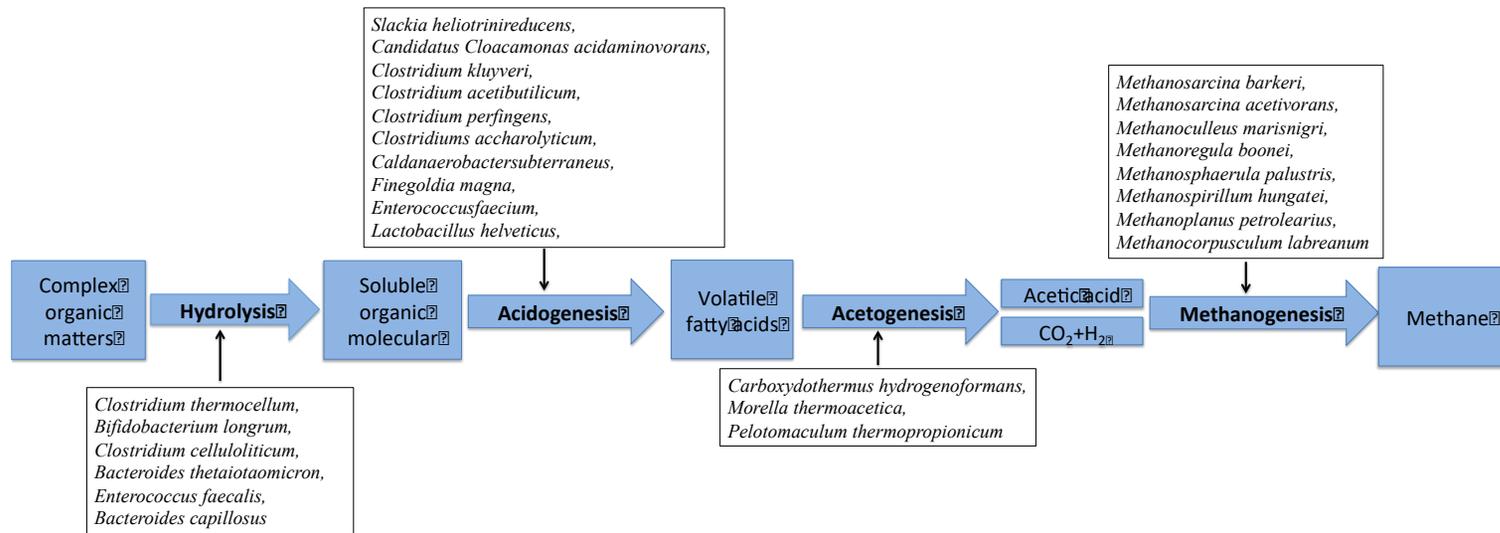


Fig. 3 Schematic representation of anaerobic fermentation process and the corresponding keystone microbial populations.

Reference:

1. S. Sauve and M. Desrosiers, *Chemistry Central Journal*, 2014, **8**, 15.
2. B. P. Colman, B. Espinasse, C. J. Richardson, C. W. Matson, G. V. Lowry, D. E. Hunt, M. R. Wiesner and E. S. Bernhardt, *Environmental science & technology*, 2014, **48**, 5229-5236.
3. A. Pruden, R. Pei, H. Storteboom and K. H. Carlson, *Environmental Science & Technology*, 2006, **40**, 7445-7450.
4. B. Petrie, R. Barden and B. Kasprzyk-Hordern, *Water Research*, 2015, **72**, 3-27.
5. M. R. Jumat, M. F. Haroon, N. Al-Jassim, H. Cheng and P.-Y. Hong, *Water*, 2018, **10**, 436.
6. J. Fabrega, R. Zhang, J. C. Renshaw, W.-T. Liu and J. R. Lead, *Chemosphere*, 2011, **85**, 961-966.
7. N. Bolong, A. Ismail, M. R. Salim and T. Matsuura, *Desalination*, 2009, **239**, 229-246.
8. G. Ying, R. Kookana and T. Waite, 2004.
9. A. Katsigiannis, C. Noutsopoulos, J. Mantziaras and M. Gioldasi, *Chemical Engineering Journal*, 2015, **280**, 49-57.
10. L. Kovalova, D. R. Knappe, K. Lehnberg, C. Kazner and J. Hollender, *Environmental Science and Pollution Research*, 2013, **20**, 3607-3615.
11. M. Petrovic, S. Gonzalez and D. Barcelo, *TrAC Trends in Analytical Chemistry*, 2003, **22**, 685-696.
12. C. W. McKinney and A. Pruden, *Environmental science & technology*, 2012, **46**, 13393-13400.
13. C. Adams, Y. Wang, K. Loftin and M. Meyer, *Journal of environmental engineering*, 2002, **128**, 253-260.
14. M. B. Ahmed, J. L. Zhou, H. H. Ngo, W. Guo, N. S. Thomaidis and J. Xu, *Journal of hazardous materials*, 2017, **323**, 274-298.
15. P. Le-Clech, *Applied microbiology and biotechnology*, 2010, **88**, 1253-1260.
16. P. Xu, J. E. Drewes, T.-U. Kim, C. Bellona and G. Amy, *Journal of Membrane Science*, 2006, **279**, 165-175.
17. A. Lim and R. Bai, *Journal of membrane science*, 2003, **216**, 279-290.
18. H. Cheng, Y. Xie, L. F. Villalobos, L. Song, K.-V. Peinemann, S. Nunes and P.-Y. Hong, *Scientific reports*, 2016, **6**.
19. J.-H. Kim, D.-C. Choi, K.-M. Yeon, S.-R. Kim and C.-H. Lee, *Environmental science & technology*, 2011, **45**, 1601-1607.
20. M. Simoes, L. C. Simoes and M. J. Vieira, *LWT-Food Science and Technology*, 2010, **43**, 573-583.

21. J. S. Kim, E. Kuk, K. N. Yu, J.-H. Kim, S. J. Park, H. J. Lee, S. H. Kim, Y. K. Park, Y. H. Park and C.-Y. Hwang, *Nanomedicine: Nanotechnology, Biology and Medicine*, 2007, **3**, 95-101.
22. A. Azam, A. S. Ahmed, M. Oves, M. S. Khan, S. S. Habib and A. Memic, *International journal of nanomedicine*, 2012, **7**, 6003.
23. A. B. Djurusic, Y. H. Leung, A. Ng, X. Y. Xu, P. K. Lee and N. Degger, *Small*, 2015, **11**, 26-44.
24. P. AshaRani, G. Low Kah Mun, M. P. Hande and S. Valiyaveetil, *ACS nano*, 2008, **3**, 279-290.
25. M. Mortimer, K. Kasemets and A. Kahru, *Toxicology*, 2010, **269**, 182-189.
26. A. Thill, O. Zeyons, O. Spalla, F. Chauvat, J. Rose, M. Auffan and A. M. Flank, *Environmental science & technology*, 2006, **40**, 6151-6156.
27. V. Sharma, D. Anderson and A. Dhawan, *Apoptosis*, 2012, **17**, 852-870.
28. M. I. Khan, A. Mohammad, G. Patil, S. Naqvi, L. Chauhan and I. Ahmad, *Biomaterials*, 2012, **33**, 1477-1488.
29. P. Chairuangkitti, S. Lawanprasert, S. Roytrakul, S. Aueviriyavit, D. Phummiratch, K. Kulthong, P. Chanvorachote and R. Maniratanachote, *Toxicology in vitro*, 2013, **27**, 330-338.
30. A. Chompoosor, K. Saha, P. S. Ghosh, D. J. Macarthy, O. R. Miranda, Z. J. Zhu, K. F. Arcaro and V. M. Rotello, *Small*, 2010, **6**, 2246-2249.
31. M. Ahamed, M. J. Akhtar, M. A. Siddiqui, J. Ahmad, J. Musarrat, A. A. Al-Khedhairi, M. S. AlSalhi and S. A. Alrokayan, *Toxicology*, 2011, **283**, 101-108.
32. N. Tran, A. Mir, D. Mallik, A. Sinha, S. Nayar and T. J. Webster, *International journal of nanomedicine*, 2010, **5**, 277.
33. W. Zhang, Y. Li, J. Niu and Y. Chen, *Langmuir*, 2013, **29**, 4647-4651.
34. Y. H. Leung, X. Xu, A. P. Ma, F. Liu, A. M. Ng, Z. Shen, L. A. Gethings, M. Y. Guo, A. B. Djurusic and P. K. Lee, *Scientific reports*, 2016, **6**.
35. O. Choi and Z. Hu, *Environmental science & technology*, 2008, **42**, 4583-4588.
36. J. M. Slauch, *Molecular microbiology*, 2011, **80**, 580-583.
37. J. J. Harrison, V. Tremaroli, M. A. Stan, C. S. Chan, C. Vacchi Suzzi, B. J. Heyne, M. R. Parsek, H. Ceri and R. J. Turner, *Environmental microbiology*, 2009, **11**, 2491-2509.
38. A. Ivask, K. Juganson, O. Bondarenko, M. Mortimer, V. Aruoja, K. Kasemets, I. Blinova, M. Heinlaan, V. Slaveykova and A. Kahru, *Nanotoxicology*, 2014, **8**, 57-71.
39. A. Ivask, O. Bondarenko, N. Jephthina and A. Kahru, *Analytical and bioanalytical chemistry*, 2010, **398**, 701-716.
40. Y. Xie, Y. He, P. L. Irwin, T. Jin and X. Shi, *Applied and environmental microbiology*, 2011, **77**, 2325-2331.
41. J. L. Hobman and L. C. Crossman, *Journal of medical microbiology*, 2015, **64**, 471-497.
42. O. Bondarenko, A. Ivask, A. Kakinen and A. Kahru, *Environmental pollution*, 2012, **169**, 81-89.

43. J. R. Morones, J. L. Elechiguerra, A. Camacho, K. Holt, J. B. Kouri, J. T. Ramirez and M. J. Yacaman, *Nanotechnology*, 2005, **16**, 2346.
44. M. Horie, K. Fujita, H. Kato, S. Endoh, K. Nishio, L. K. Komaba, A. Nakamura, A. Miyauchi, S. Kinugasa and Y. Hagihara, *Metallomics*, 2012, **4**, 350-360.
45. X. Zheng, Y. Su, Y. Chen, R. Wan, K. Liu, M. Li and D. Yin, *Environmental science & technology*, 2014, **48**, 13800-13807.
46. A. L. Neal, N. Kabengi, A. Grider and P. M. Bertsch, *Nanotoxicology*, 2012, **6**, 371-380.
47. H. C. Poynton, J. M. Lazorchak, C. A. Impellitteri, B. J. Blalock, K. Rogers, H. J. Allen, A. Loguinov, J. L. Heckman and S. Govindasmaw, *Environmental science & technology*, 2012, **46**, 6288-6296.
48. W. Jiang, K. Yang, R. W. Vachet and B. Xing, *Langmuir*, 2010, **26**, 18071-18077.
49. A. Kumar, A. K. Pandey, S. S. Singh, R. Shanker and A. Dhawan, *Chemosphere*, 2011, **83**, 1124-1132.
50. Y. Liu, L. He, A. Mustapha, H. Li, Z. Hu and M. Lin, *Journal of applied microbiology*, 2009, **107**, 1193-1201.
51. M. T. Madigan, J. M. Martinko and J. Parker, *Brock biology of microorganisms*, Prentice hall Upper Saddle River, NJ, 1997.
52. A. K. Suresh, D. A. Pelletier, W. Wang, J.-W. Moon, B. Gu, N. P. Mortensen, D. P. Allison, D. C. Joy, T. J. Phelps and M. J. Doktycz, *Environmental science & technology*, 2010, **44**, 5210-5215.
53. S. Makhluף, R. Dror, Y. Nitzan, Y. Abramovich, R. Jelinek and A. Gedanken, *Advanced Functional Materials*, 2005, **15**, 1708-1715.
54. N. Al-Jassim, D. Mantilla-Calderon, T. Wang and P.-Y. Hong, *Environmental Science & Technology*, 2017, **51**, 3649-3659.
55. Z. Sheng and Y. Liu, *Water research*, 2011, **45**, 6039-6050.
56. V. C. Reyes, S. O. Opot and S. Mahendra, *Environmental toxicology and chemistry*, 2015, **34**, 887-897.
57. W. Yu, M. Brown and N. J. Graham, *Scientific reports*, 2016, **6**, 30144.
58. N. Maximous, G. Nakhla, W. Wan and K. Wong, *Journal of Membrane Science*, 2010, **352**, 222-230.
59. D. Y. Koseoglu-Imer, B. Kose, M. Altinbas and I. Koyuncu, *Journal of membrane science*, 2013, **428**, 620-628.
60. Q. Ma and T. K. Wood, *Environmental microbiology*, 2009, **11**, 2735-2746.
61. O. Habimana, A. Semiao and E. Casey, *Journal of Membrane Science*, 2014, **454**, 82-96.
62. H. Koo, J. Xiao, M. Klein and J. Jeon, *Journal of bacteriology*, 2010, **192**, 3024-3032.
63. N. R. Stanley and B. A. Lazazzera, *Molecular microbiology*, 2004, **52**, 917-924.
64. S. S. Branda, A. Vik, L. Friedman and R. Kolter, *Trends in microbiology*, 2005, **13**, 20-26.
65. M. J. Franklin, D. E. Nivens, J. T. Weadge and P. L. Howell, *Frontiers in microbiology*, 2011, **2**.

66. X. Wang, J. F. Preston and T. Romeo, *Journal of bacteriology*, 2004, **186**, 2724-2734.
67. C. Darby, J. W. Hsu, N. Ghori and S. Falkow, *Nature*, 2002, **417**, 243-244.
68. M. Schuster and E. P. Greenberg, *International journal of medical microbiology*, 2006, **296**, 73-81.
69. B. R. Singh, B. N. Singh, A. Singh, W. Khan, A. H. Naqvi and H. B. Singh, *Scientific reports*, 2015, **5**.
70. S. Yang, F. Yang, Z. Fu, T. Wang and R. Lei, *Journal of Hazardous Materials*, 2010, **175**, 551-557.
71. J. Pang, M. Matsuda, M. Kuroda, D. Inoue, K. Sei, K. Nishida and M. Ike, *Frontiers of Environmental Science & Engineering*, 2016, **10**, 7.
72. Y. Chen, Y. Su, X. Zheng, H. Chen and H. Yang, *Water research*, 2012, **46**, 4379-4386.
73. X. Zheng, Y. Chen and R. Wu, *Environmental science & technology*, 2011, **45**, 7284-7290.
74. Y. Chen, D. Wang, X. Zhu, X. Zheng and L. Feng, *Environmental science & technology*, 2012, **46**, 12452-12458.
75. J. W. Metch, N. D. Burrows, C. J. Murphy, A. Pruden and P. J. Vikesland, *Nature nanotechnology*, 2018, **1**.
76. Z. Hu, X. Lu, P. Sun, Z. Hu, R. Wang, C. Lou and J. Han, *Bioresource technology*, 2017, **225**, 279-285.
77. X. Zheng, R. Wu and Y. Chen, *Environmental science & technology*, 2011, **45**, 2826-2832.
78. H. Cheng and P.-Y. Hong, *Environmental science & technology*, 2017, **51**, 12200-12209.
79. O. M. Amaya, M. T. C. Barragan and F. J. A. Tapia, in *Biomass Now-Sustainable Growth and Use*, Intech, 2013.
80. R. Wirth, E. Kovacs, G. Maroti, Z. Bagi, G. Rakhely and K. L. Kovacs, *Biotechnology for biofuels*, 2012, **5**, 41.
81. S. Yu, G. Zhang, J. Li, Z. Zhao and X. Kang, *Bioresource technology*, 2013, **146**, 758-761.
82. H. Mu and Y. Chen, *Water research*, 2011, **45**, 5612-5620.
83. P. M. Christy, L. Gopinath and D. Divya, *Renewable and Sustainable Energy Reviews*, 2014, **34**, 167-173.
84. M. Zakrzewski, A. Goesmann, S. Jaenicke, S. Junemann, F. Eikmeyer, R. Szczepanowski, W. A. Al-Soud, S. Sorensen, A. Puhler and A. Schluer, *Journal of biotechnology*, 2012, **158**, 248-258.
85. H. Mu, Y. Chen and N. Xiao, *Bioresource technology*, 2011, **102**, 10305-10311.
86. H. Chen, Y. Chen, X. Zheng, X. Li and J. Luo, *Water research*, 2014, **63**, 125-134.
87. S. W. Ragsdale, *Annals of the New York Academy of Sciences*, 2008, **1125**, 129-136.
88. T. Wang, D. Zhang, L. Dai, Y. Chen and X. Dai, *Scientific reports*, 2016, **6**.
89. P. Mishra, S. Thakur, D. M. Mahapatra, Z. Ab Wahid, H. Liu and L. Singh, *International Journal of Hydrogen Energy*, 2018.

90. A. Li, J. Yan, J. Yu, L. Zhang, L. Ren, S. Li, S. Wu, X. Liu, X. Wang and Y. n. Chu, *Biotechnology for biofuels*, 2013, **6**, 3.
91. M. Luna-delRisco, K. Orupold and H.-C. Dubourguier, *Journal of hazardous materials*, 2011, **189**, 603-608.
92. Y. Yang, M. Xu, J. D. Wall and Z. Hu, *Waste management*, 2012, **32**, 816-825.
93. S. F. Stoddard, B. J. Smith, R. Hein, B. R. Roller and T. M. Schmidt, *Nucleic acids research*, 2014, **43**, D593-D598.
94. A. Garcia, L. a. Delgado, J. A. Tora , E. Casals, E. Gonzalez, V. Puntès, X. Font, J. Carrera and A. Sanchez, *Journal of hazardous materials*, 2012, **199**, 64-72.
95. J. Ma, X. Quan, X. Si and Y. Wu, *Bioresource technology*, 2013, **149**, 346-352.
96. F. Suanon, Q. Sun, D. Mama, J. Li, B. Dimon and C.-P. Yu, *Water research*, 2016, **88**, 897-903.
97. A. W. Carpenter, S. N. Laughton and M. R. Wiesner, *Environmental engineering science*, 2015, **32**, 647-655.
98. S. T. Khan, R. Wahab, J. Ahmad, A. A. Al-Khedhairi, M. A. Siddiqui, Q. Saquib, B. A. Ali and J. Musarrat, *Korean Chemical Engineering Research*, 2015, **53**, 565-569.
99. Z. Vardanyan, V. Gevorkyan, M. Ananyan, H. Vardapetyan and A. Trchounian, *Journal of nanobiotechnology*, 2015, **13**, 69.
100. D. Laha, A. Pramanik, A. Laskar, M. Jana, P. Pramanik and P. Karmakar, *Materials Research Bulletin*, 2014, **59**, 185-191.
101. R. Meraat, A. A. Ziabari, K. Issazadeh, N. Shadan and K. M. Jalali, *Acta Metallurgica Sinica (English Letters)*, 2016, **29**, 601-608.
102. S. Sarwar, S. Chakraborti, S. Bera, I. A. Sheikh, K. M. Hoque and P. Chakrabarti, *Nanomedicine: Nanotechnology, Biology and Medicine*, 2016, **12**, 1499-1509.
103. D. Ranmadugala, A. Ebrahiminezhad, M. Manley-Harris, Y. Ghasemi and A. Berenjian, *Process Biochemistry*, 2017, **62**, 231-240.
104. V. Aruoja, S. Pokhrel, M. Sihtmae, M. Mortimer, L. Madler and A. Kahru, *Environmental Science: Nano*, 2015, **2**, 630-644.
105. S. Luche, E. Eymard-Vernain, H. I. n. Diemer, A. Van Dorsselaer, T. Rabilloud and C. c. Lelong, *Journal of proteomics*, 2016, **135**, 170-180.
106. X. Li, R. Pang, J. Li, X. Sun, J. Shen, W. Han and L. Wang, *Desalination*, 2013, **324**, 48-56.
107. K. C. Anyaogu, A. V. Fedorov and D. C. Neckers, *Langmuir*, 2008, **24**, 4340-4346.
108. M. Ben-Sasson, K. R. Zodrow, Q. Genggeng, Y. Kang, E. P. Giannelis and M. Elimelech, *Environmental science & technology*, 2013, **48**, 384-393.
109. Y. Yang, Q. Chen, J. D. Wall and Z. Hu, *Water research*, 2012, **46**, 1176-1184.
110. L. Otero-Gonzalez, J. A. Field and R. Sierra-Alvarez, *Water research*, 2014, **58**, 160-168.
111. L. Otero-Gonzalez, J. A. Field and R. Sierra-Alvarez, *Journal of environmental management*, 2014, **135**, 110-117.