

# Precipitation of carbonates by bacteria isolated from wastewater samples collected in a conventional wastewater treatment plant

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**Abstract** This research studied the precipitation of calcium carbonate by populations of bacteria from domestic wastewater cultivated in both natural and artificial solid culture media. The only carbonate-forming bacteria detected appeared in an artificial medium added with calcium acetate. Precipitation occurred three days after inoculation, and the percentage was slightly higher than 65 %.

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Our results showed that nine major carbonate-forming colony types were the dominant heterotrophic plateable bacteria growing aerobically in artificial media added with calcium acetate. According to their taxonomic affiliations (based on partial sequencing of the 16S-rRNA), the nine strains belonged to the following nine genera of Gram-negative and Gram-positive bacteria: *Caulobacter*, *Blastomonas*, *Roseobacter*, *Staphylococcus*, *Bacillus*, *Gemmatimonas*, *Saccharopolyspora*, *Microthrix*, and *Sphingomonas*. All of these strains formed calcium carbonate, precipitated as calcite and vaterite in different proportions and shapes (spheres, hemispheres, dumbbells, and pseudopolyhedral forms). The results of this study suggest that in real domestic wastewater, the precipitation of carbonates through bacterial action could not take place in situ because the concentrations of calcium did not create the optimal circumstances for biomineralization. However, in the artificial media, it was possible to induce this process by adding calcium ions.

**Keywords** Calcite · Calcium carbonate · Domestic wastewater · Vaterite

## Introduction

Calcium carbonate precipitation is a common phenomenon found in different environments, such as sea water, freshwater, industrial wastewaters, and soil (Delgado et al. 2008; Ehrlich 2002; Hammes and Verstraete 2002; Hammes et al. 2003). It is a chemical process that requires sufficient calcium concentration and carbonate ions so that the ion activity product exceeds the solubility constant. It also depends on the presence of nucleation sites.



Microorganisms, mainly bacteria, contribute to the precipitation of a wide variety of minerals (Dove et al. 2003; Ehrlich 2002). Numerous laboratory studies have demonstrated the bacterial precipitation of carbonates (Rivadeneira et al. 1999; Rivadeneira et al. 2000, 2006a, b; Sánchez-Román et al. 2007) and have proposed different mechanisms for the bacterially mediated precipitation of carbonate minerals (Ehrlich 2002; Rivadeneira et al. 1998; Rivadeneira et al. 2010; van Lith et al. 2003). One of the mechanisms most often proposed is the production of ammonium and  $\text{CO}_2$  by microorganisms (which increases the pH) in the presence of calcium and magnesium-produced calcium or calcium and/or magnesium carbonates. This is an induced precipitation mechanism (Lowenstam and Weiner 1989), where the metabolic activity alters the physicochemical parameters of the habitats of these microorganisms, allowing the precipitation of minerals. Other authors have reported that the bacteria can serve as a nucleus for mineral precipitation upon adsorbing  $\text{Ca}^{2+}$ ,  $\text{Mg}^{2+}$  and other metallic cations onto the cell surface, and that the matrix of extracellular polymeric secretions affects mineral precipitation (Beveridge and Fyfe 1985; Rivadeneira et al. 2006a; van Lith et al. 2003). In this context, the role of bacterial surfaces in the nucleation of certain minerals (such as some types of carbonate) has been widely discussed. However, in many cases, the exact role of the bacteria in such nucleation remains unclear.

Different bacterial species have previously been detected and assumed to be associated with carbonate precipitation in diverse environments including bioreactor systems for industrial wastewater treatments (Hammes et al. 2003; van Lith et al. 2003). Species-specific carbonate precipitation has been hypothesized by several authors (Hammes and Verstraete 2002) though the exact mechanisms of precipitation and the way that this process works within the microbial ecology of the precipitating organism remains unresolved.

The activated sludge process is a wastewater treatment method in which the carbonaceous organic matter of wastewater provides an energy source for the production of new cells for a mixed population of microorganisms in an aquatic aerobic environment. The microbes convert carbon into cell tissue and oxidized end products that include carbon dioxide and water. The majority of microorganisms in activated sludge are bacteria. For the most part, they are heterotrophic, and thus require organic compounds for their supply of carbon and energy. In contrast, autotrophic bacteria, which use inorganic compounds for cell growth, occur in proportion to concentrations of carbon and nitrogen. Important genera of heterotrophic bacteria include *Achromobacter*, *Sphingomonas*, *Alcaligenes*, *Arthrobacter*, *Citromonas*, *Flavobacterium*, *Pseudomonas*, and *Zoogloea*

(Jenkins et al. 1993). Previous studies have shown that microbial populations influence the precipitation of different minerals in wastewater treatment systems (Dove et al. 2003; Hammes et al. 2003). For example, the interest in dedicated reactors for struvite precipitation is currently growing because of the extensive operational problems caused by struvite accumulation in wastewater treatment plants (Straful et al. 2001).

In this paper, we study carbonate precipitation by bacteria isolated from domestic wastewater collected from the municipal wastewater treatment plant of the city of Granada (Puente de los Vados, Granada, Spain) in culture media made from domestic wastewater (natural media) and in artificial culture media (conventional media for carbonate precipitation). The main goal was to demonstrate which culture conditions influence carbonate formation caused by bacteria isolated from real domestic wastewater. This paper also discusses the relation of these microorganisms to biomineralization in culture media.

## Materials and methods

### Domestic wastewater samples

Wastewater samples were collected from the wastewater treatment plant of the city of Granada (Puente de los Vados, Granada), operated by EMASAGRA S.A. This wastewater was taken from the primary settling tank of the treatment plant. The average composition of wastewater was determined by standard methods (Clesceri et al. 2001) and was found to be the following: chemical oxygen demand (COD), 450 mg/l; biological oxygen demand at 5 days (BOD<sub>5</sub>), 300 mg/l;  $\text{NO}_3^-$ , 5 mg/l;  $\text{NH}_4^+$ , 75 mg/l;  $\text{Ca}^{2+}$ , 15 mg/l; total suspended solids (TSS), 0.01 g/l; volatile suspended solids (VSS), 0.78 g/l. The samples were collected in sterile bottles (1 L), shipped to the laboratory, and refrigerated (4 °C) until their analysis in the laboratory.

### Culture media

The culture media used in this study were wastewater media (WWM) and artificial media added with calcium acetate (MC). The WWM medium was composed of domestic wastewater with the addition of 18 g/l of Bacto-Agar. The pH of the medium was adjusted to 7.2 with 0.1 M KOH. The MC medium was composed of 10.0 g/l yeast extract, 5.0 g/l protease peptone, 1.0 g/l glucose, and 4.0 g/l calcium acetate. To obtain a solid medium, 18 g/l Bacto-Agar was added, and the pH was adjusted to 7.2 with 0.1 M KOH. The culture media were autoclaved at 112 °C for 20 min.

## Microorganisms

The experiments were performed with bacterial strains isolated from urban wastewater samples. Aliquots (0.1 ml) of the wastewater samples were serially diluted and spread on plates containing WWM and MC solid media. Plates were aerobically incubated at 25 °C for 30 days and checked periodically for the presence of crystals using optical microscopy. The percentage of crystal-forming and non-crystal-forming colonies was counted. Isolated representatives of the dominant colony morphologies with crystal-forming capacity were selected and purified, by restreaking them twice on trypticase soy agar (TSA). All the experiments were carried out in triplicate.

## Genetic identification of the isolated bacterial strains

In this study, all of the strains isolated (with crystal-forming capacity) were identified by analyzing the partial sequence of the gene encoding 16S rRNA. Primers fD1 and rD1 (Weisburg et al. 1991) were synthesized by Sigma Genosis (UK) and used to amplify nearly the full length of the 16S rRNA gene. Fresh cultured colonies of each strain were lysed by the addition of 20 µl of a mixture of NaOH (0.05 M)–SDS (0.25 %, w/v), which was then boiled for 15 min. The lysates were adjusted to 200 µl with sterile water and centrifuged at 2500g for 5 min in a table-top centrifuge. The cleared lysates (4 µl) were used as a template for amplification. PCR was carried out by adding the following to the lysates: 1xPCR Gold buffer (Applied Biosystems, Germany); 1.5 mM MgCl<sub>2</sub> (Applied Biosystems, Germany); 200 µM dNTPs (Roche Molecular Biochemicals, Germany); 20 pmol of each primer; and 1 U of Ampli-Tag Gold polymerase (Applied Biosystems, Germany). The final volume of the reaction tubes was adjusted to 50 µl. Reactions were run in a Perkin Elmer GeneAmp PCR system 2400 (Perkin Elmer, Norwalk, CT, USA). The temperature profile was the one previously described by Vinuesa et al. (1998), except for the extension of the initial denaturation step to 7 min, as required, using the Quiaex II kit (Qiagen, Germany). The nucleotide sequence of the purified bands was determined by the dideoxy chain terminator method, using the ABI-PRISM Big Dye Terminator Cycle Sequencing Ready Reaction kit (Perkin Elmer, Norwalk, CT, USA) and automated sequencer ABI-PRISM 3100 Avant Genetic Analyzer (Applied Biosystems, Germany). The sequenced fragment analyzed corresponded to the first 650 bp of the 16S-rRNA gene, comprising hypervariable regions V1, V2, and V3 (Neefs et al. 1990).

DNA sequences were analyzed using the biocomputing tools provided on-line by the European Bioinformatics Institute (<http://www.ebi.ac.uk>). The BLASTn (Altschul et al. 1997) program was used for preliminary sequence

similarity analysis, and the ClustalX v.1.8 software (Jeanmougin et al. 1998) was used for sequence alignment.

## Study of mineral formation

All the nine strains (major colony types) isolated from the wastewater samples with crystal-forming capacity were surface-inoculated onto WWM and MC solid media, incubated aerobically at 25 °C, and periodically examined with an optical microscope for the presence of carbonate crystals up to 30 days after inoculation. The experiments were carried out in triplicate and were repeated three times. A control consisting of uninoculated culture media and media inoculated with autoclaved bacterial cells were included in all experiments.

For mineralogical and morphological analysis, precipitates were removed by cutting out pieces of the media, which were placed in boiling water to dissolve the agar. The sediments were re-suspended and washed in distilled water to free them of impurities. In this treatment, calcium carbonate dissolution was not significant, and the morphology of crystals was not altered, as observed by optical microscopy both before and after their recovery. The washed carbonate crystals were finally air-dried at 37 °C.

## XRD study

The precipitates obtained after 30 days of incubation were examined by powder X-ray diffraction (PXRD) using a Philips PW 1710/00 diffractometer, with a graphite monochromator automatic slit, CuK $\alpha$  radiation, and on-line connection with microcomputer. Data were collected for a 0.4 s integration time in 0.02 °C 2 $\theta$  steps at 40 kV and 40 mA, in a 2 $\theta$  interval between 3 and 80 °C. Data were processed using the XPPowder program for a qualitative and quantitative determination of the mineral composition (Martín 2004). The crystalline mosaic size on hkl reciprocal vectors was obtained from full width at half of the maximum intensity (FWHM) after instrumental broadening and  $K_{\alpha 2}$  corrections.

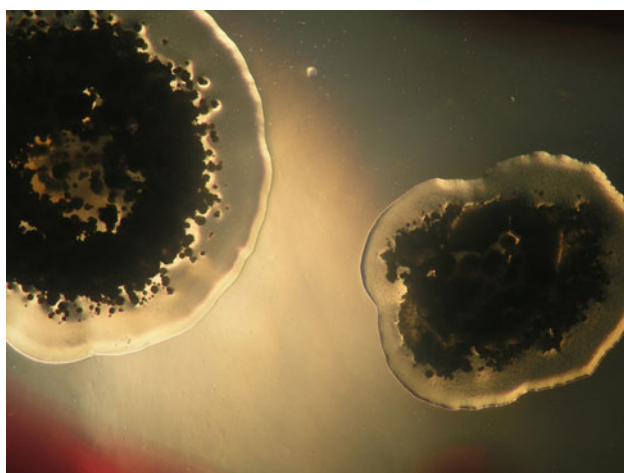
## SEM study

Secondary electron micrographs of bacterial precipitates were made with gold-coated samples using a Zeiss DMS SEM (LEO Electron Microscopy, Oberkochen, Germany), operated at an acceleration voltage of 20 kV to examine the micromorphology of the crystals. Some selected samples were coated with carbon for energy dispersive X-ray (EDX) microanalysis. High-resolution secondary electron images were prepared with a field emission scanning electron microscopy (FESEM) LEO 1525, under 2–3 kV on carbon-coated samples.



## Results and discussion

The formation of calcium carbonate was only observed in artificial media (MC). No precipitation was detected in natural media (WWM), uninoculated control media, or media inoculated with a high concentration of dead bacteria and thus without metabolic activity. The number of bacteria (CFU) per ml of urban wastewater recounts in WWM medium was  $5.5 \times 10^5$ , whereas in MC medium, it was  $2.3 \times 10^6$ . The percentage of carbonate-forming bacteria in the MC medium was slightly higher than 65 %. Precipitation took place rapidly, and crystal formation began 3 days after inoculation. After 10 days, the crystals had significantly increased in quantity and were of a large size. Figure 1 shows colonies with precipitates of carbonates.



**Fig. 1** Colonies with precipitates of carbonates

Nine major colony types were detected as the dominant heterotrophic plateable bacteria growing aerobically in MC media. Nine different colonies of identical morphology were randomly selected from plates, re-isolated, and tested for phenotypical characters. Identical results were obtained. The taxonomic affiliations of the nine strains, based on partial sequencing of the 16S-rRNA gene (V1 to V3 hypervariable regions, ca. 650 nt) are shown in Table 1. The strains fell into eight different genera of Gram-negative and Gram-positive bacteria. Sequence comparison with databases demonstrated the affiliation of strain WW1 to *Caulobacter* sp. (95.8 % identity), WW2 to *Blastomonas* sp. (99.7 % identity), WW3 to *Roseobacter* sp. (89.4 % identity), WW4 to *Staphylococcus epidermidis* (100 % identity), WW5 to *Bacillus cereus* (98.8 % identity), WW6 to *Gemmatimonas aurantica* (89.4 % identity), WW7 to *Saccharopolyspora* sp. (82.3 % identity), WW8 to *Microthrix parvicella* (96.9 % identity) and WW9 to *Sphingomonas* sp. (99.4 % identity).

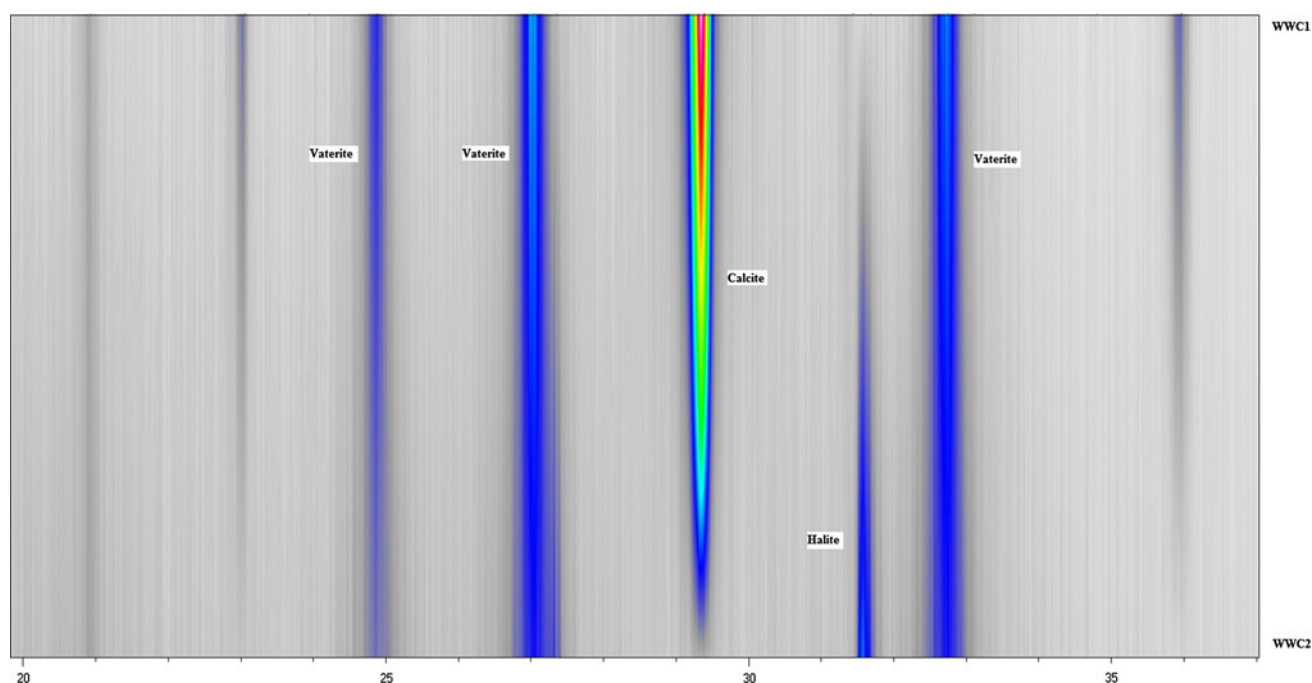
The results of the mineralogical analysis with XRD (Fig. 2; Table 2) showed that all of the strains formed calcium carbonate, precipitated as calcite and vaterite in different proportions, depending on the strain. Precipitation of small amounts of calcium phosphates and different percentages of amorphous crystals were also observed in all cases (Fig. 1).

WW7 and WW8 strains affiliated as Actinobacteria precipitated minerals with a low percentage of calcite and a high percentage of vaterite of lower crystallinity than the precipitated mineral produced by bacterial strains affiliated to other taxonomic classes. These results were also verified by measuring the crystal size of all the samples (Table 2).

**Table 1** Identification of strains with crystal-forming capacity isolated from urban waste waters

Strain	Closest taxonomy affiliation (class/family)	Overlap (nt)	Most similar organisms	Access nc.	Percent identity
WW1	$\alpha$ -proteobacteria/Caulobacteraceae	120	<i>Caulobacter</i> sp. 5142	AY97383	95.8
			<i>Brevundimonas</i> sp. K2/98-FUNDUS	AJ313427	90.2
WW2	$\alpha$ -proteobacteria/Sphingomonadaceae	123	<i>Blastomonas</i> sp.	AB242676	99.7
			<i>Sphingomonas</i> sp.	AJ812013	98.1
WW3	$\alpha$ -proteobacteria/	123	<i>Roseobacter</i> sp.	AY136130	89.4
			<i>Sinorickettsia chlamys</i>	AY174894	86.1
WW4	Bacilli/Staphylococcae	159	<i>Staphylococcus epidermidis</i>	AF270147	100
WW5	Bacilli/Bacillaceae	141	<i>Bacillus cereus</i>	AY138274	98.8
			<i>Bacillus anthracis</i>	AY138291	81.4
WW6	Gammatimonadetes	159	<i>Gemmatimonas aurantiaca</i>	AB072735	89.4
			Bacterium Ellin 5290	AY234641	86.3
WW7	Actinobacteria/Pseudonocarficaceae	141	<i>Saccharopolyspora</i> sp.	H7131491	82.3
WW8	Actinobacteria	124	<i>Microthrix parvicella</i>	X93044	96.4
			<i>Collinsella</i> sp.	AB064936	82.7
WW9	$\alpha$ -proteobacteria/Sphingomonadaceae	123	<i>Sphingomonas</i> sp.	AJ812013	99.4
			<i>Blastomonas</i> sp.	AB242676	91.2





**Fig. 2** Map drawn from XRD patterns. Colors in the map indicate the changing intensity of the diffracted X-rays as a function of  $2\theta$ , with warmer colors for progressively higher intensities. Residues of halite of the culture media are also present in the patterns

**Table 2** Quantitative analysis of precipitates and size of crystal

Bacteria	Vaterite		Calcite		Ca phosphates + amorphous % Crystalline components
	% Crystalline components	Crystal size (nm)	% Crystalline components	Crystal size (nm)	
Actinomycetes	$72.5 \pm 4.25$	$20 \pm 1$	$8.3 \pm 0.8$	Very low, Shortly content	$19.2 \pm 1.7$
Other bacteria	$42.2 \pm 4.3$	$29 \pm 4.6$	$52.8 \pm 8.9$	$50 \pm 6.5$	$5 \pm 4.5$

In general, it was observed that the calcite showed higher crystallinity than the vaterite. In contrast, the vaterites precipitated by Actinobacteria showed a lower crystallinity than the vaterites precipitated by the other microorganisms studied. Moreover, in the minerals precipitated by Actinobacteria, a high percentage of amorphous forms was also observed.

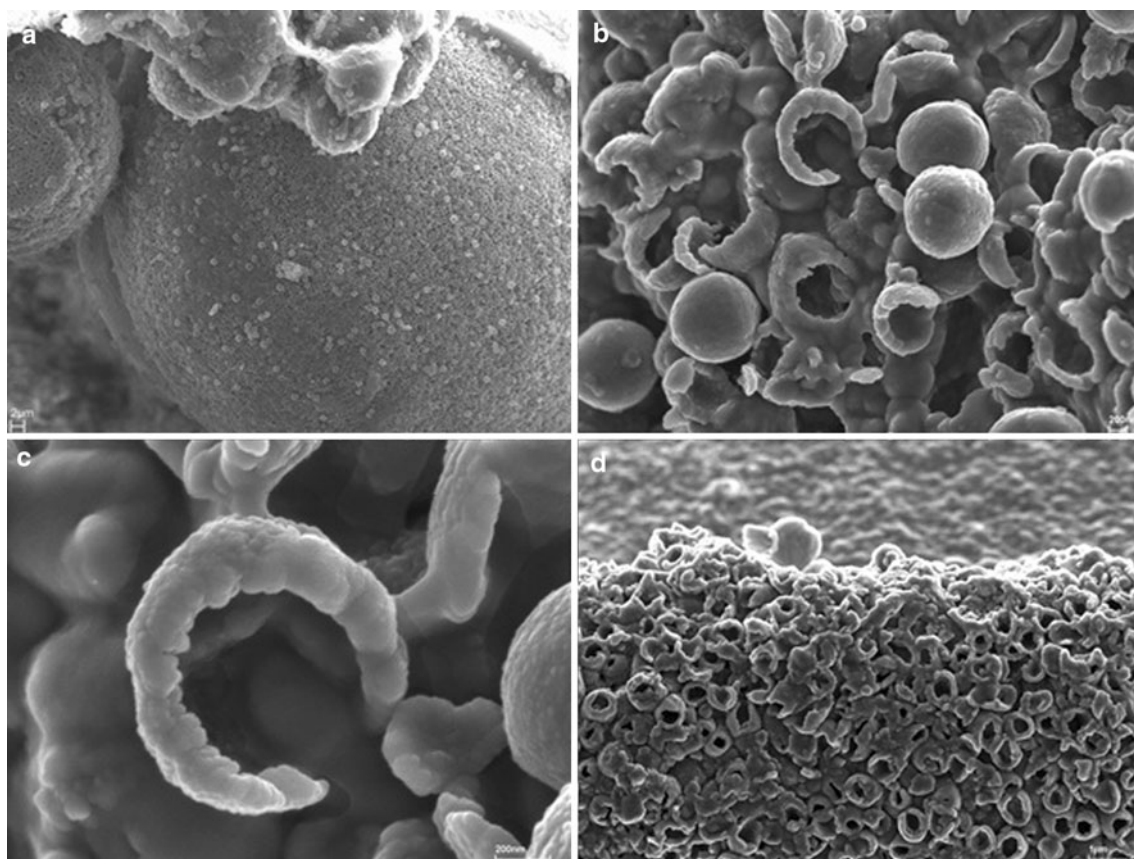
A variety of shapes were observed by SEM (Figs. 3, 4). Of these, the most important were spheres, hemispheres, dumbbells, and pseudopolyhedral forms, which appeared either in isolation or in groups. However, spherulitic forms were also predominant. Certain differences were detected in the spherulites produced by Actinobacteria strains and the spherulites formed by the rest of the microorganisms. More specifically, the spherulites formed by non-Actinobacteria microorganisms had rough surfaces with small holes, and high porosity (Fig. 3a). Many holes were of

bacteria-like size and shape, and in many cases, mineralized bacteria were clearly evident (Fig. 3b, c, d). However, most of the bioliths generated by Actinobacteria were formed by cell aggregation strains. Other spherulites did not show any cell mark, but rather were frequently covered by mineralized filaments (Fig. 4). The microscopical observation of the plates showed that the majority of spherulites were not only in the bacterial mass but also near the colonies. The EDX analysis (Fig. 5) confirmed the DRX results and showed that spherulites formed by Actinobacteria had a greater concentration of phosphates than the spherulites precipitated by other bacteria. The analysis also showed that the calcified filament in some Actinobacteria spherulites contained a greater proportion of calcium phosphate than the rest of the spherulites.

The influence of microorganisms in mineral precipitation has been recognized for a wide variety of minerals. In







**Fig. 3** SEM and FESEM images of carbonate bioliths formed by WW4 strain (*Staphylococcus epidermidis*). **a** Spherulites and dumbbells. **b, c** Close-up of the boxed area in **a**, showing a dense aggregate of mineralized bacterial cells composed of Ca-carbonate. Note the

abundance of calcium-carbonate nanoparticles delimiting the bacterial-cell contours. **d** Edge of a hemispherical biolith (hemispheres) which shows its formation by accumulation of calcified cocci

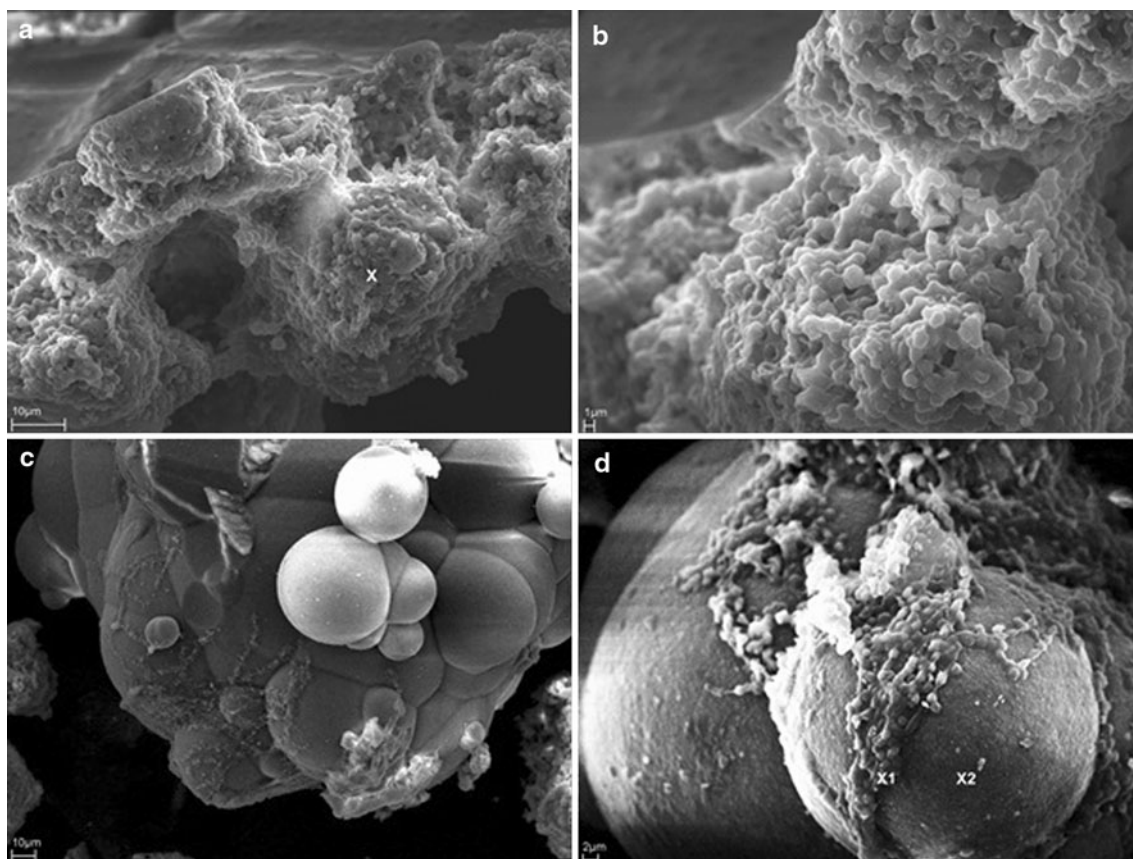
fact, there have been various research studies on microbial involvement in carbonate precipitation in natural environments (Ehrlich 2002; Parraga et al. 2004). Generally speaking, calcium carbonate biomineralization is not necessarily linked to any particular group of microorganisms, and has been extensively documented in a variety of natural environments.

A culture-dependent approach was used to analyze the widely diverse bacteria in the urban wastewater of our study. The results showed a complex cultivable community mainly integrated by members of the genus *Caulobacter*, *Blastomonas*, *Roseobacter*, *Staphylococcus*, *Bacillus*, *Gemmatimonas*, *Saccharopolyspora*, and *Collinsella*. This technique has been extensively applied to the study of a variety of different bacterial ecosystems, including wastewaters, sludges, and biofilms. However, recently, the introduction of molecular biology techniques based on the in situ detection of nucleic acids has provided important information about microorganisms in their natural habitats. In this context, denaturing gradient gel electrophoresis (DGGE/TGGE) yields extensive data concerning the complexity and behavior of microbial communities

(Muyzer 1999). Most of these techniques have been used in recent studies on the ecology of biological processes in wastewaters (Wagner et al. 2002). We found that the community isolated in our standard domestic wastewater using culture-dependent methods was analogous to what has been described by other authors using culture-independent techniques, such as DGGE and TGGE (Cortes-Lorenzo et al. 2006; Molina-Muñoz et al. 2007).

Numerous laboratory studies and observations in natural environments have demonstrated the bacterial precipitation of calcium carbonate (Rivadeneyra et al. 1999, 2000, 2006b; Sánchez-Román et al. 2007). One of the mechanisms of calcium carbonate precipitation most often proposed is that microorganisms produce  $\text{NH}_4^+$  by metabolizing nitrogenated organic substrates, which increase the pH. In the presence of calcium ions, the  $\text{CO}_2$  also produced by the bacterial activity precipitates at a basic pH in the form of calcite, aragonite, vaterite, etc. Consequently, the  $\text{CO}_2$  produced by the microbial community during the biodegradation of the organic matter is totally or partially used for the formation of calcium carbonate.





**Fig. 4** SEM images of carbonate bioliths formed by actinobacteria strains WW7 (*Saccharopolyspora sp*) and WW8 (*Microthrix parvicella*) (a) and (b) Spherulites in formation, showing a dense aggregate of mineralized bacterial cells composed principally of Ca-carbonate

(see EDX spectrum b). c, d spherulites group with smooth surface, where they are not observed traces of bacteria. Some spherulites are covered by mineralized filaments EDX spectrum indicates that they are composed of Ca and P

To date, most systematic studies on bacterial calcium carbonate biomineralization have been carried out in natural aquatic environments (many of which are marine environments) or in artificial culture media using different microorganisms previously isolated from aquatic or soil habitats (Delgado et al. 2008; Parraga et al. 2004; Riva-deneyra et al. 2006a, b). Nevertheless, there have been no publications that have reported the bioprecipitation of calcium carbonate associated with a microbial community in domestic wastewaters.

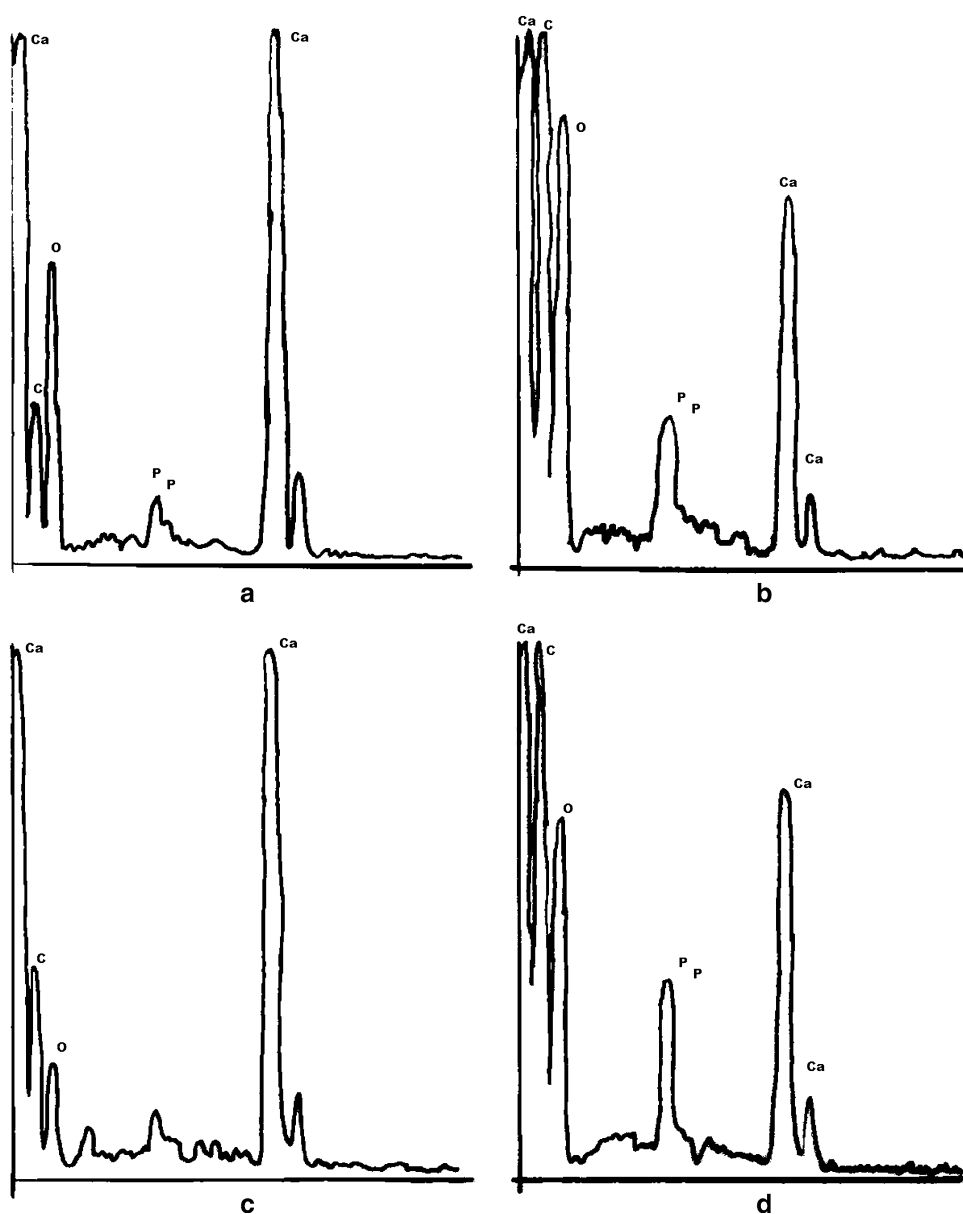
Our study did not detect the formation of calcium carbonate when bacterial colonies were grown in a natural culture medium containing urban wastewater as a source of nutrients (WWM medium). In the case of the WWM medium, all of the bacterial strains grew there very well, and formed colonies in 48 h though no crystal formation was observed after the 30-day incubation period. In contrast, 65 % of the colonies detected in the artificial culture media, containing yeast extract, protease peptone, glucose, and calcium acetate, formed calcium carbonate after an incubation period of 3 days. Previous results have shown that calcium and metabolizable organic substrate

concentrations are influential factors in the biomineralization of calcium carbonate (Delgado et al. 2008). Our data suggest that the calcium concentration in the urban wastewaters was not sufficient to produce the precipitation of calcium carbonate under our experimental conditions. However, in artificial culture media amended with high concentrations of metabolizable organic matter and significant amounts of calcium, the microbial populations in the urban wastewaters were able to create the optimal conditions for the formation of carbonates.

The most significant finding of this research study is that bacterial populations in domestic wastewater can precipitate calcium carbonate in the form of calcite and vaterite in different proportions, depending on the strain. Moreover, the precipitation of small amounts of calcium phosphates and different percentages of amorphous crystals can also be produced in all cases. However, this precipitation capacity was only induced when the microorganisms were cultivated in artificial laboratory media and never when the microorganisms were cultivated in natural media derived from the wastewater. This suggests that in actual domestic wastewater, the precipitation of carbonates through



**Fig. 5** EDX spectra of carbonate bioliths **(a)** carbonate bioliths formed by WW4 strain (See location in Fig. 3a). Bioliths formed by actinobacteria strains: **b** bioliths in formation (See location in Fig. 4a), **c** spherulites group with smooth surface (See location in Fig. 4d) and **d** mineralized filaments covering some spherulites (See location in Fig. 4d)



bacterial action could not take place in situ. Obviously, the chemical composition of the wastewater can affect the result of the process. If the wastewater had a high concentration of calcium, then the biomineralization of calcium carbonate could be bacterially induced.

SEM was used to detect a wide range of morphological features and shapes (i.e. spheres, hemispheres, dumbbells, and pseudopolyhedral forms), either in isolation or in groups (Figs. 3, 4). All of them are typical of bacterial carbonate precipitation. The dependence of the crystal morphology produced by the tested strains on the artificial culture media was difficult to establish since the morphology was quite heterogeneous. Electron microscopy was used to verify that the calcium carbonate bioliths produced by the bacteria showed a significant quantity of

cell marks inside the mass as well as on the surface. This fact confirms that they are formed by the accumulation of calcified organisms. These findings have been previously reported by other authors (Parraga et al. 2004; Rivadeneyra et al. 1998, 2004, 2006a). The observation with FESEM shows mineralized cells defined by rounded cells to Ca-carbonate nanoparticles. Similar nanoparticles have also been found in other bacterial Ca–Mg carbonate precipitates, called “nanoglobules” (Aloisi et al. 2006; Sánchez-Román et al. 2008). Sánchez-Navas et al. (2009) interpreted them as the nanocrystalline building units that form bacterially precipitated carbonate mesocrystals.

In the case of actinomycetes, these cell marks are not always present. The microscopical observation of the plates shows that the majority of spherulites are not only in the





bacterial mass but also nearby the colonies. This result indicates that these spherulites were not formed by the aggregation of bacteria, and in those cases, Actinobacteria only contributed to this formation by changing the media as a consequence of their metabolic activity.

The samples of carbonates also showed precipitation of calcium phosphate and small amounts of amorphous minerals. Rivadeneyra et al. (2010) described the precipitation of carbonates in *C. marismortui*, beginning with the early formation of amorphous calcium phosphate nanoglobules. During the formation and maturation of the spherulites, these nanoglobules were transformed into amorphous calcium carbonate, which gradually reached higher crystallinity to finally become aragonite. Based on this, in our opinion, the poorly crystalline calcium phosphates and small amounts of amorphous calcium phosphate observed in our precipitates (WWC1, WWC2 and WWC3) may correspond to stages previous to the precipitation of calcium carbonate. The spherulites produced by actinomycetes often comprise calcified filaments on their surface. The microanalysis (Fig. 5) showed a higher percentage of calcium phosphates, a fact which leads us to believe that these filaments may be at an earlier stage of the crystallization process, and confirms that the calcium phosphate may be the precursor of the carbonate. Our findings support the existence of this process, and also indicate that this process is not unique to *C. marismortui*, but may be a more widespread type of precipitation.

In conclusion, all the results of this study, which is limited to cultivable bacteria, confirm that bacterial activity has an importance role in the formation of calcium carbonate studied in vitro, and that consequently, this is a biomineralization process. The formation of calcium carbonate by microorganisms isolated from domestic wastewater is mainly calcite and vaterite in artificial media, which contains a high concentration of calcium and metabolizable organic matter. However, for the study of calcium carbonate precipitation by bacteria from domestic wastewaters, it is advisable not to use media derived from the wastewater itself since environmental conditions do not create the optimal circumstances for the precipitation of minerals.

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