Structure and Function of Stony Coral Intraskeletal Polysaccharides

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1 Supporting Information

ABSTRACT: Polysaccharides represent a main weight fraction of the intraskeletal organic matrix of corals, but their structure, as well as their function in the calcification process, has been poorly investigated. This communication shows by a combination of techniques (nuclear magnetic resonance, Fourier transform infrared, and monosaccharide composition) that their key component is a 1→3 β-D glucuronic acid polymer and evidences its influence in vitro in the calcification process.

1. INTRODUCTION

The key component of biominerals is the organic matrix (OM), which is involved in the control of nucleation, growth, form, and function of the mineral phase. In stony corals, OM is synthesized by cells of the calicoblastic epithelium and then secreted into the adjacent subepithelial space in which calcification takes place. The OM is formed by a mixture of macromolecules (proteins, polysaccharides, and lipids) that share some common chemical features through diverse coral genus. The structural and functional relationship between OM and mineral phases in scleractinian coral skeleton only recently has started to be understood at the molecular level, for what regards the proteic moieties. These proteins, as many others associated with biominerals, are characterized by the presence of high mol % (around 40 mol %) of acidic residues.

Polysaccharides represent a relevant mass fraction of the OM in corals, but their function in the crystallization process is not known. In mollusks, chitin has been commonly observed to build an insoluble three-dimensional scaffold in which the acid proteins adsorb, establishing a confined mineralizing space. This polysaccharide is not common in corals OM, and its presence has been reported only in some species. In coccolithopores, three families of polysaccharides interact directly with the CaCO₃. Indeed, it was demonstrated that polysaccharides replace the water molecules that form hydrogen bonds with the surface of the growing crystals. That polysaccharides shield cation and anion trafficking to the crystals’ surfaces inhibiting their growth. Usually, OM polysaccharides (pscOMs) bear sulfate groups that cooperate with the carboxylates one to induce CaCO₃ nucleation. For instance, the investigation on the effects of carbohydrate additives during the early stages of CaCO₃ precipitation revealed a variety of interaction between these additives and CaCO₃ species, such as the complexation of Ca²⁺, the influence on the stability of solute prenucleation clusters, and the inhibition or promotion of nucleation.

Here, it is reported as the characterization of a fraction of pscOM from coral skeletons and the study of its influence on inorganic mineral phase.

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in vitro CaCO₃ precipitation. *Astroides calycularis* (Aca), *Balanophyllia europaea* (Beu), and *Stylophora pistillata* (Spi), which are representatives of diverse trophic strategies and growth forms.¹¹

### 2. RESULTS AND DISCUSSION

pscOM was obtained by the mild alkaline digestion at room temperature of the whole intraskeletal OM (wOM, that is, soluble and insoluble fractions). Alkaline hydrolysis leads to the removal of saccharides which are O-linked to Ser or Thr and to the random breaking of peptide bonds in proteins, producing single amino acids or small peptides. Alkaline hydrolysis can also destroy monosaccharides and remove polysaccharide critical function groups (e.g., acetyl).²⁷ This choice was carried out because protease activity by PNGase F (cleaving N-linked oligosaccharides at the N-glycosidic bond) and by other proteolytic enzymes having a broad specificity (see the Supporting Information), which was expected to hydrolyze the protein regions, failed. This is because their activity is restricted in the vicinity of densely glycosylated regions of the protein, presumably because of steric hindrance.²⁸

The pscOM represents about 8 wt % of the wOM, with a great variability among different (*n* = 3) alkaline digestion experiments, more specifically: 7 ± 4 wt % for Aca, 8 ± 4 wt % for Beu, and 11 ± 6 wt % for Spi.

A first characterization of pscOM was performed by FTIR spectroscopy. In Figure 1, the most representative spectra of pscOM for each species (small differences in absorption profiles were recorded among samples from the same species) are shown. A description of the assigned absorption bands is listed in Table S1.²⁹,³⁰ The relative intensities of these bands changed according to the pscOM source. The spectra from all of the species showed that the most intense bands were those because of polysaccharides (1200–1000 cm⁻¹), namely the vibration modes associated with ring vibration overlapped with stretching vibrations of carbonyl group and the glycosidic bond vibration.

In particular, at 1078 and 1037 cm⁻¹, ether bonds and C–C single bond vibration modes were present. Weak absorption bands relative to methylene and methyl group vibration mode and to absorption bands at 1634 and 1649 cm⁻¹ could be associated with the presence of both N-acetyl and uronic acid residua. The FTIR spectrum of pscOM from Spi showed the presence of absorption bands at 1453 and 1412 cm⁻¹ that were probably because of carboxylic groups. The pscOM from Beu showed additional weak bands at 1255 and 1235 cm⁻¹ that can be associated with the S=O stretching, suggesting the presence of sulfate groups.

The pscOM was further characterized by nuclear magnetic resonance (NMR) spectroscopy. The ¹H NMR pscOM spectrum of Spi was the most detailed one (Figure S2), whereas the other samples showed, despite the same purification process, large amount of broad signals suggesting the presence of different molecular components. The spectrum of Spi shows a predominant set of well-resolved signals coming into the typical chemical shift region of polysaccharides together with minor signals because of other minority components, as suggested by the FTIR spectrum. Mono- and bidimensional NMR techniques were used for the structure identification. Through 2D correlation spectroscopy (COSY) and total correlation spectroscopy (TOCSY), it was possible to identify the anomeric signal and four secondary alcohol signals belonging to the ring proton, whereas any –CH₂– or primary alcohol signals were detected. The investigation was prosecuted by heteronuclear ¹H–¹³C COSY [heteronuclear single-quantum coherence (HSQC)] to verify the number of carbon atoms and to obtain the chemical shift values of their signals (Figure 2). The experiment confirmed the absence of diprotonated carbons suggesting the presence of a uronic acid residue. Using an NMR spectrometer equipped with cryoprobe, it was possible to perform a heteronuclear multiple bond correlation experiment (HMBC). The result of Spi NMR studies confirmed the working hypothesis of the presence of a uronic acid, identifying the three and two bond distances carbon–proton correlation, that is, the correlations of carboxyl in C6 with the protons of carbon 4 and 5 of the sugar ring. The assigned chemical shift values are reported in Table S2.

The proton and carbon chemical shift values of C1, C3, and C6 as well as the anomeric proton coupling constant (*J*₁₋₂) indicate the presence of a β 1–3 glucuronic acid copolymer. The observed chemical shifts are in agreement with the literature data reporting a synthetic model obtained by oxidation of (1→3) β-curdan, in spite of the different spectrum calibration that cause a systematic 3 ppm shift in signals.²⁸

Furthermore, experiments evidenced the presence of an N-acetyl residue (¹H: 2.02 ppm; ¹³C: 25 and 177 ppm) likely linked to unidentified saccharidic minor component(s), whose detectable signals appear in the proton spectrum between 4.6 and 3.8 ppm and having different anomeric signals at 5.52, 5.08, and 5.03 ppm.

The above data were supplemented by the analysis of constituent monosaccharides. In view of the presence of a polyglucuronate polymer, the Taylor and Conrad method was applied.³³ Table 1 summarizes the results. The monosaccharide compositional pattern of pscOM from Aca, Beu, or Spi was similar; all extracts were extremely rich in glucuronic acid, which represented about 80 wt % of the total monosaccharides in each of these fractions. Arabinose, galactose, mannose, and xylose were also detected.
crystals were generated by the steps given by the combination of Aca3. Image (Figure 3; Aca3) showed that the curved surfaces of the agglomerates (up to randomly aggregated in polycrystalline units that joined in big presence of pscOM from Aca, crystals of calcite appeared edge-free (Figure 3; Aca2). The higher magnification image (Figure 3; Aca3) showed that the curved surfaces of the crystals were generated by the steps given by the combination of \{10.4\} and \{hk.l\} faces. Among the latter, the presence of small \{00.1\} faces was identified (arrow in Figure 3; Aca2). The surfaces of the \{10.4\} faces showed particles of about 200 nm. In the presence of pscOM from Beu, calcitic spherulites having a diameter up to 180 \(\mu m\) formed (Figure 3; Beu1). Their flat surface regions, associated with the \{10.4\} faces, were surrounded by rough regions, associated with \{hk.l\} faces. The latter was shown on the surface platelike particles of about 500 nm (Figure 3; Beu2−3). In the presence of pscOM from Spi, the rhombohedral calcite crystals clustered in aggregates having a diameter up to 150 \(\mu m\) (Figure 3; Spi1). In these aggregates, calcite crystals penetrated and preserved most of the \{10.4\} faces. They also exposed \{hk.l\} faces that smoothed the crystalline edges (Figure 3; Spi1). The \{10.4\} faces were very flat, whereas crystallites of about 100 nm were observed on the \{hk.l\} faces.

The relevance of studying the structure of pscOM and their influence on the deposition of calcium carbonate in organisms are first given by their high relative mass content with respect to other macromolecules (i.e., proteins and lipids) into many calcified tissues.1−4 This is particularly true for stony corals from the Mediterranean Sea, where their content was much higher than that of proteins.5 However, also other studies on tropical coral species provided a similar outcome.11,13

The pscOM investigated in this study represents only a fraction of the whole content of polysaccharides associated with the wOM. This is because the quite harsh treatment used for its extraction provoked the degradation and loss of sugars (not quantifiable). For that reason, its low mass content (about 8 wt %) in the wOM is not in conflict with the 52 wt % of monosaccharides detected in the soluble OM from A. calycularis.34 On the other side, the remaining material was still polymeric (MW > 3500 Da, according to dialysis) and of reduced complexity, making easier its structural characterization. Most importantly, its relevance for in vitro calcification studies is proved by the capability to affect the precipitation of calcium carbonate.

Table 1. Relative Molar Percentages of the Monosaccharide Detected in the pscOM from S. pistillata (Spi), B. europea (Beu), and A. calycularis (Aca)

| samples | Ara | Xyl | Man | Gal | Glc |
|---------|-----|-----|-----|-----|-----|
| Spi     | 6.5 | 5.0 | 6.0 | 2.0 | 80.5|
| Beu     | 4.1 | 3.8 | 7.1 | 7.0 | 78.0|
| Aca     | 3.0 | 4.1 | 7.2 | 5.3 | 80.4|

“Percentage of peak area relative to total peak areas, determined by GC−MS. Ara = arabinose; Man = mannose; Xyl = xylose; Gal = galactose; and Glc = glucose from glucuronic acid reduction (see Section 4.3.1, Monosaccharide Analysis).
The monosaccharidic distribution of the pscOM from Aca, Beu, and Spi is similar (Table 1) and different from what was reported for other species of coral;35,36 for example, in A. millepora, a content of about 70 wt % of arabinose and of about 3.9 wt % of glucuronic acid was detected, but again, a diverse extraction procedure was used. The corals monosaccharidic composition is also diverse with most of sugar moieties associated with mollusks, where carbohydrates are quantitatively minor in comparison to proteins.37

In the case of Spi, the NMR investigations detected the presence of 1→3 β-D glucuronic acid polymers. To our knowledge, this is the first identification of a polysaccharidic sequence in the OM of corals.

This signature is in line with what was observed in coccolith-associated polysaccharides.9,15 One of this polysaccharide, referred to as PS2, contains β-linked D-glucuronic acid sequences and has the role to accumulate calcium ions in the coccolith formation process.38 A coccolith contamination of our sequences and has the role to accumulate calcium ions in the associated polysaccharides.18,19 One of this polysaccharide, sequence in the OM of corals.

The presence of an N-acetyl residue, linked to unidentified saccharide minor component(s), may indicate the presence of small amount of chitinlike structures, as already reported in old studies,16,17 which were resistant to the chemical treatment.

The function of polysaccharide on coral calcification is completely unknown, different from what was recently reported for proteins6−12 and/or lipids.18

Here, the influence of pscOM on CaCO₃ precipitation was studied by comparing the results of experiments performed in the presence of the pscOM, the wOM, and in the absence of additives. Without additives, only rhombohedral calcite crystals precipitated, whereas in the presence of pscOM, the morphologies observed were different from those imaged in the presence of wOM (Figure S5), according to their diverse composition (Figure 3).

Shape and morphology were different for particles formed in the presence of pscOM from the diverse species. In them, the observed morphology was generated by different crystal aggregation and the relative extension of {10.4} and a-specific {hk.l} stepping faces. However, in the presence of pscOM from Aca specific {00.1} faces were observed. This face was observed in the calcite crystal grown in the presence of the polysaccharidic component of the OM extracted from the skeleton of the mollusk Atrina rigida and associated with the interaction with polysaccharidic moieties.39 In addition, the observation of diverse kind of nanoparticles (or crystal units) on the surface of the growing crystals agree with the recent observations that scleractinia coral skeleton forms through the aggregation of spherical particles of amorphous calcium carbonate.6−12

These differences in the texture of the calcite crystals occur despite the similarity of the pscOM composition and the common presence, directly detected only for Spi, of 1→3 β-D glucuronic acid polymers. This observation is intriguing also considering that these scleractinia corals belong to diverse families, Beu and Aca to Dendrophylliidae, whereas Spi to Pocilloporidae.

In agreement with what was observed in the precipitation experiment of calcite, glucuronic acid concentrates and binds calcium ions38,40 and inhibits the nucleation process of calcium carbonate.22 Interestingly, the capability of the glucuronic acid moieties to interact with calcium ions40 and calcite crystals22 is dependent on the way they are bonded to a polypeptide core. This feature also changes the assembly of calcite crystals from...
compensated units to spherulites, with a concomitant reduction of the size of monocrystals. This knowledge may justify the diverse calcite crystal shapes and textures observed in the presence of pscOM from different coral species.

3. CONCLUSIONS

In conclusion, in this study, for the first time, it is observed that 1→3 β-D glucuronic acid polymers are in the carbohydrate fraction of the wOM from corals. These polysaccharidic moieties can act as calcite morphology modifiers in a species-specific manner. These discoveries shed a new light on the function of polysaccharide in the molecular processes which govern coral calcification.

4. MATERIALS AND METHODS

4.1. OM Extraction. The samples of B. europaea and A. calycularis from Palinuro (Italian coast, North-Western Mediterranean Sea) and S. pistillata from the Gulf of Eilat (Red Sea, Israel) were randomly collected by scuba diving at 6, 9, and 10 m depths, respectively. Coral skeletons were cleaned and ground, and wOM was extracted through decalcification using a 0.1 M acetic acid solution, as previously reported. wOM was weighed and analyzed by FTIR spectroscopy using a Bruker ALPHA FTIR spectrometer equipped with a single-reflection diamond-attenuated total reflection accessory working in the range of wavenumbers 4000–400 cm⁻¹ at a resolution of 4 cm⁻¹.

4.2. Polysaccharide Purification. Proteolytic digestion: 50 mg of wOM were dispersed in sodium phosphate buffer (100 mM, pH 7.8, 5 mL). Pronase (protease type XIV from Streptomyces griseus, Sigma-Aldrich; 10 pg) was added and the solution was left for 24 h at 37 °C. Trypsin (Sigma-Aldrich; 50 μg) was added and left for an additional 18 h at 37 °C. Then, the dispersion was extensively dialyzed against deionized water and finally lyophilized. PNGase F treatment: 50 mg of wOM were dispersed in 5 mL of a denaturing pH 7.5, 0.05 M phosphate buffer containing 0.5 wt % sodium dodecyl sulfate and 1 wt % mercaptoethanol by heating at 100 °C for 2 min. PNGase F (20 units; Sigma-Aldrich) was added and incubated for 24 h at 37 °C. The dispersion was extensively dialyzed against deionized water and finally lyophilized. Alkaline digestion: 50 mg of wOM were dispersed in 5 mL of a 0.1 M NaOH solution at room temperature and the resultant solution was stirred for 4 days at room temperature. Then, the dispersion was extensively dialyzed against deionized water and finally lyophilized and weighed.

The product of wOM enzymatic or chemical treatment was first characterized by FTIR spectroscopy. Only the alkaline digestion provoked a reduction of the protein content, estimated by the intensity of FTIR absorption of the amide I and amide II proteic bands.

4.3. Polysaccharide Characterization. 4.3.1. Monosaccharide Analysis. The polysaccharide fraction from Spi, Beu, or Aca was carboxyl-reduced by the carbodiimide method, using two successive cycles to obtain reduced CH₂OH from COOH groups of uronic acid units. NaBH₄ was used as the reducing agent. Monosaccharide components of the carboxyl-reduced polysaccharide and their ratios were determined, following hydrolysis with 1 M trifluoroacetic acid for 16 h at 100 °C. The acid was then evaporated, and the residue was dissolved in water. The hydrolyzate was converted to alditol acetates by treatment with NaBH₃O₃ followed by acetylation with acetic anhydride—pyridine (1:1 v/v) at 100 °C for 30 min, and the resulting alditol acetates were extracted with chloroform. The alditol acetate analysis was carried out using a Varian gas chromatograph, model CP-3800, and mass spectrometer, model Saturn 4000R, with He as the carrier gas (1 mL/min). A capillary column (30 m × 0.25 mm i.d.) of DB-1, held at 100 °C during injection for 1 min, then programmed at 5 °C/min to 220 °C, and held at 280 °C for 18 min was used for the quantitative analysis. The alditol acetates were identified by their retention times and typical electron impact breakdown profiles compared with standards.

4.3.2. NMR Characterization. The product of the alkaline digestion was characterized by NMR spectroscopy. NMR spectra were recorded at 25 °C using Bruker AVANCE III spectrometers (600 and 500 MHz) equipped with high-sensitivity 5 mm TCI cryoprobes. After exchanging 2 mg of the sample thrice in 2H₂O, samples were dissolved in 0.6 mL of a 3 mM EDTA d₂ ²H₂O (99.996%) solution and placed in 5 mm NMR tubes. Proton spectra were recorded with presaturation of the residual water signal with a recycle delay of 6 s and 32 scans. Bidimensional DQF (double-quantum-filter)—COSY and 2D (two-dimensional)—TOCSY spectra were acquired using 16 scans per series of 2048 × 256 data points with zero-filling in F1 (4096 × 1024), and a shifted (π/3) squared cosine function was applied prior to Fourier transformation. HSQC and HMBC (heteronuclear multiple bond correlation) spectra were obtained in phase-sensitivity-enhanced pure-absorption mode with decoupling in the acquisition period. Scans (40 and 96) were collected for each free-induction decay. The matrix size of 1024 × 320 data points was zero-filled to 4096 × 1024 by the application of a squared cosine function prior to Fourier transformation.

4.4. Calcium Carbonate Crystallization Experiments. Calcium carbonate (CaCO₃) in vitro crystallization experiments were carried out by vapor diffusion in a desiccator containing (NH₄)₂CO₃ (Carlo Erba) and anhydrous CaCl₂ (Sigma-Aldrich). Microplates for cellular culture (Microplate 24 well with lid, Iwaki), containing a round glass coverslip in each well, were used as previously reported. CaCl₂ solutions (750 μL of 10 mM; CaCl₂ 2H₂O, Sigma-Aldrich) were poured in each well. wOM (0.2 mg) or pscOM (0.2 mg) were added in some well. This concentration (0.27 mg/mL) ensured the optimal precipitation yield having a change of the particle shape and morphology. Microplates were located in the desiccator for 4 days at room temperature. The obtained crystals were washed with distilled water and dried. CaCO₃ crystallization experiments were replicated at least five times.

Characterization of CaCO₃ precipitates was carried out by FTIR spectroscopy and microscopic observations. FTIR spectra of samples in KBr disks were collected at room temperature by using a Bruker ALPHA FTIR spectrometer working in the range of 4000–400 cm⁻¹ at a resolution of 4 cm⁻¹. Optical microscopic observations of CaCO₃ precipitates were carried out using a Leica microscope equipped with a digital camera. SEM observations were conducted using a Hitachi FEG 6400 scanning electron microscope after coating the samples with gold.

ASSOCIATED CONTENT

Supporting Information
The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acsomega.7b02053.
Characterization data; 1H NMR spectra; FTIR spectra; and SEM images (PDF)

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Notes
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