

有孔虫：環境と生物進化の語り部

Foraminifera: Story teller of both Earth's environments and biotic evolution

*北里 洋¹

*Hiroshi Kitazato¹

1.国立大学法人東京海洋大学

1.Tokyo University of Marine Sciences and Technology (TUMSAT)

原生生物有孔虫類は、海洋生態系の重要な一員である。生元素循環とくに炭素循環、窒素循環に深く関わっている。また、真核生物進化の初期に現れ、現在に至るまで栄えている。有孔虫はさまざまな殻を持っている事から化石記録が豊富であり、海洋環境、生物進化双方の情報を読み出す事が出来る。しかし、生物としての有孔虫の実態は十分に理解されているとは言えない。講演では、生物としての有孔虫に焦点を当て、そのユニークな生態と生物像を紹介する。

キーワード：有孔虫、地球史、生物進化

Keywords: Foraminifera, Earth's history, biotic evolution

有孔虫のチャンバー形成時のpH変化

The variable ambient pH during chamber formation process of benthic foraminifera

*豊福 高志¹

*Takashi Toyofuku¹

1. 独立行政法人海洋研究開発機構

1. Japan Agency for Marine-Earth Science and Technology (JAMSTEC)

Some of major studies recently showed that calcifying marine organisms respond differently to ocean acidification (OA). Prediction of OA impact is getting tangled by this variability. Moreover, this complicates modelling future ecological cycles. Carbon usage is key to understanding calcification and hence understanding impact of OA. Using fluorescent techniques to visualize pH gradients, this shows foraminifera actively pump protons to promote passive CO₂ uptake. This appearance the basis of calcification and effectively support carbon uptake independent from seawater pH. The resulting fundamentally new calcification model has major implications for understanding past changes in atmospheric CO₂ as well for predicting future CO₂.

キーワード：有孔虫、石灰化

Keywords: Foraminifera, Calcification

Living (stained) benthic foraminifera from the Mozambique Channel (eastern Africa):
Exploring deep-sea biodiversity and biogeochemistry of unicellular fossilizing meiofauna

*Christophe Fontanier¹

1.Ifremer, 2.University of Bordeaux, EPOC, 3.JAMSTEC

Fontanier C.^a, Garnier E.^{a,b}, Brandily C.^c, Toyofuku T.^d, Saburo S.^d, Dennielou B.^a, Bichon S.^b, Gayet N.^c, Eugene T.^d, Rovere M.^a, Grémare A.^b, Deflandre B.^b

^aIfremer, Laboratoire Environnements Sédimentaires, Centre de Brest, Technopôle de Brest-Iroise, BP 70, F-29280 Plouzané, France.

^bUniversité de Bordeaux, CNRS, Environnements et Paléo-environnements Océaniques et Continentaux, UMR 5805, F-33400 Talence, France.

^cIfremer, Laboratoire Environnements Profonds, Centre de Brest, Technopôle de Brest-Iroise, BP 70, F-29280 Plouzané, France.

^dJAMSTEC Japan Agency for Marine Earth Science and Technology, 2-15 Natsushima-cho, Yokosuka F-237-0061, Japan.

^dIfremer, Service Cartographie, Traitement de Données et Instrumentation, Centre de Brest, Technopôle de Brest-Iroise, BP 70, F-29280 Plouzané, France.

This study was done in the framework of the PAMELA project ("Passive Margin Exploration Laboratories") funded by TOTAL and Ifremer. Live (Rose-Bengal stained) deep-sea foraminiferal faunas have been studied at four stations between 530–3200-m depth in the Mozambique Channel (eastern Africa) to understand how complex environmental conditions (e.g., organic matter, oxygenation) control (1) their ecological structure (i.e., diversity, standing stocks, and microhabitats) and (2) their geochemical signatures (stable isotopes, and trace elements). Two upper-slope stations, located at 530- and 780-m depth off Madagascar, are bathed by well-oxygenated bottom waters. They are characterized by fine sediments enriched in highly degraded organic matter (low amino-acid bio-availability and reduced chlorophyllic freshness). Mineralization of organic compounds results in relatively moderate oxygen penetration depth (i.e., 15 and 30 mm) in sediment. Interestingly, foraminiferal simple diversity (S) is exceptionally high at both sites. The higher standing stocks are observed in the 780-m deep station, where peculiar sedimentary facies of organic matter focusing are recorded (OC >2.0% DW). Redox conditions and sedimentary organic matter control the composition and the vertical distribution (i.e. microhabitat) of benthic faunas at both upper-slope sites. *Bolivina alata*, *Bulimina marginata*, *Haplophragmoides bradyi* and *Nouria compressa* are relevant bio-indicators of enhanced burial of organic matter prevailing at the 780-m deep station (i.e., eutrophic settings), whereas *Uvigerina hispida* and *Uvigerina semiornata* are dominant at the 530-m deep station (i.e., relatively mesotrophic settings). Two other stations are located on well-ventilated terraces from the deep-sea canyons of Tsiribihina and Zambezi (>3000-m depth). They are characterized by carbonate ooze, which is depleted in degraded organic matter and, where oxygen penetration depth is relatively deep (i.e., > 80 mm). Because of food scarcity, foraminiferal simple diversity (S) and standing stocks are relatively low, and agglutinated and organic-walled taxa dominate foraminiferal faunas. *Hospitella fulva*, a foraminifera belonging to Allogromiida, occupy very deep infaunal microhabitat, what disrupts the classical scheme of microhabitat patterns in oligotrophic settings.

Keywords: Mozambique Channel, Foraminifera, Diversity, Microhabitat, Sedimentary Organic Matter, Stable Isotopes and Trace Elements

小規模データを有効活用した有孔虫石灰化プロセス動力学の再構築

Reconstruction of foraminiferal calcification process by effective use of restricted data

*山本 美希¹、豊福 高志¹、白石 史人²、de Nooijer Lennart³、阪口 秀¹*Miki Y. Matsuo¹, Takashi Toyofuku¹, Fumito Shiraishi², Lennart Jan de Nooijer³, Hide Sakaguchi¹

1.海洋研究開発機構数理科学先端技術研究分野、2.広島大学大学院理学研究科地球惑星システム学専攻、3.オランダ王立海洋研究所

1.Mathematical Science and Advanced Technology, Japan Agency for Marine-Earth Science and Technology, 2.Department of Earth and Planetary Systems Science, Graduate School of Science, Hiroshima Univ., 3.Royal Netherlands Institute for Sea Research

有孔虫の石灰化プロセスの動力学の詳細を理解することは、古環境を再構築する手段としての有孔虫の役割をより明確に確立するために必要である。しかしながら石灰化過程における周辺カルシウムの挙動やそれと関わる殻表面成長など、重要なプロセスをリアルタイムで詳細に観測できていない。本公演では、このような石灰化過程における重要なプロセスを、カルシウムプローブなどから得られる少ないカルシウム挙動情報から予測・再構築する情報処理手法を提案し、提案手法を実際に適用した結果について報告する。

アコヤガイ貝殻稜柱層カルサイト結晶内の有機物ネットワークと結晶欠陥の関係

Relation between the organic network and the crystal defects in the calcite of the prismatic layer of *Pinctada fucata*

*近都 浩之¹、奥村 大河²、根岸 瑠美³、小暮 敏博²、作田 庄平¹、鈴木 道生¹

*Hiroyuki Kintsu¹, Taiga Okumura², Lumi Negishi³, Toshihiro Kogure², Shohei Sakuda¹, Michio Suzuki¹

1.東京大学大学院農学生命科学研究科、2.東京大学大学院理学系研究科、3.東京大学分子細胞生物学研究所
1.Graduate School of Agricultural and Life Sciences, The University of Tokyo, 2.Graduate School of Science, The University of Tokyo, 3.Institute of Molecular and Cellular Biosciences, The University of Tokyo

Biominerals are biogenic mineralized tissues containing not only inorganic components but also a small amount of organic matrices that play an important role in formation of biominerals. Molluscan shells, which are typical biominerals consisting of calcium carbonate and organic matrices, have various kind of microstructures. The shell of a pearl oyster, *Pinctada fucata*, has two different layers. The outer layer of the shell is a prismatic layer consisting of calcite crystals. Each prism, which is surrounded by the organic framework, is composed of some single crystals of calcite. The single crystal has the small distortion of crystal orientation because this crystal has some small-angle grain boundaries inside of the crystal which cause the minim lattice distortion or defect and divide the crystal into subgrain units of a few hundred nanometers. Such minim lattice distortion and defects may increase the toughness of the shell by inhibiting the cleavage and fracture of the crystals in the prismatic layer. The small-angle grain boundaries were observed by TEM along with the localized organic matrices like networks, indicating that the organic networks may cause the small-angle grain boundaries. TEM observations showed that the organic networks are a few dozen nanometers in thickness and divide the crystal into subgrain units of a few hundred nanometers. However, what the organic networks consist of or how they cause the small angle grain boundaries has not been reported yet.

To reveal the components of the organic networks and formation mechanism of the small angle grain boundaries in the calcite crystal of the prismatic layer, we extracted the organic network from the prismatic layer and tried to identify the components. The IR spectrum of acetic acid-insoluble materials from the prisms revealed that the major component of the organic network was chitin. LC-MS/MS analysis of acetic acid-insoluble and SDS/DTT-soluble fractions showed that the chitinolytic enzymes such as chitinase and chitobiase were involved in the intracrystalline organic matrices of the prismatic layer. These results suggested that the chitinase and chitobiase regulate the formation of chin fibers that interact with the calcium carbonate to make the small-angle grain boundaries. To understand the function of the chitinolytic enzymes in the prismatic layer, calcium carbonate was precipitated in the chitin hydrogel after being treated with the commercially available enzyme. The calcite crystals precipitated in the chitin hydrogel appeared to contained larger crystal defects as the chitinolytic enzyme concentration increase and crystal defects similar to those in the prismatic layer were observed as to some extent concentration. In addition to observation, a variance of lattice spacing was calculated from peak broadening in powder X-ray diffraction and compared among the these calcite crystals. As a result, a variance lattice spacing tended to increase depending on chitinolytic enzyme concentration, implying that chitinolytic enzyme decrease the thickness of the chitin fiber to increase the interaction between chitin fibers and calcium carbonate. Such interaction is probably important to produce the small-angle grain boundaries in the calcite crystal and strengthens the toughness of shell.

キーワード：バイオミネラル、貝殻、キチナーゼ

Keywords: biomineral, shell, chitinase

有孔虫の殻における水平および垂直なテクスチャー

Lateral and Vertical Textures in Shells of Calcitic and Aragonitic Foraminifers

*中島 希実子¹、鈴木 萌菜美¹、長井 裕季子²、緒明 佑哉¹、豊福 高志²、今井 宏明¹

*Kimiko Nakajima¹, Monami Suzuki¹, Yukiko Nagai², Yuya Oaki¹, Takashi Toyofuku², Hiroaki Imai¹

1.慶應義塾大学工学部応用化学科、2.独立行政法人海洋研究開発機構

1.Department of Applied Chemistry, Keio University, Yokohama, Japan, 2.Japan Agency for Marine-Earth Science and Technology, Yokosuka, Japan

Although shells of various foraminifers were reported to be composed of unique layered structures, details of the crystallographic structure have not been clarified sufficiently. Here, we investigated the shells of foraminifers, calcitic *Ammonia beccarii* and aragonitic *Hoeglundina elegans*, using scanning electron microscopy (SEM), transmission electron microscopy (TEM), and selected area electron diffraction (SAED). Despite the difference in their crystal polymorph and shapes, they exhibited similar textures in micro- and nano-scales. The shells were revealed to consist of single-crystalline micrometric domains with their c axes perpendicular to the shell surface from SAED patterns of platy samples 100 nm thick prepared using a focused ion beam (FIB) technique. Cross-sectional SEM and TEM observations showed the presence of lateral organic layers ~10-20 nm thick with ~200-400 nm intervals in the single-crystalline domains. Whereas a micrometric layered structure has previously been reported as a periodic growth pattern, we discovered finer lateral textures. On the other hand, a columnar structure ~100-400 nm wide derived from vertical textures was observed on the cross-sections after mild etching of the shells in a diluted acetic acid solution. We also found the vertical textures which originated from strains of the crystal lattice as a difference of contrast in the TEM images of the FIB-cut samples. The columns were produced through selective dissolution of the strained area. Therefore, the foraminifer shells of the two species were deduced to consist of micrometric single-crystalline domains having lateral and vertical textures with organic thin layers and strains, respectively, regardless of the difference of polymorph.

Morphospecies, cryptic species or biological species...? Searching for the evolutionary significance of diversity in foraminifera

*Agnes Katharina Weiner¹, Masashi Tsuchiya², Takashi Toyofuku², Hiroshi Kitazato²

1.JSPS/JAMSTEC, 2.JAMSTEC

Many groups of foraminifera are characterized by the formation of elaborate shells. These shells provide detailed morphological features that can be used for species classification. Since the majority of works on foraminifera focuses on their fossil record and their application as proxies in micropaleontological studies, a comprehensive morphotaxonomy has been established, describing tens of thousands of extant and extinct morphospecies. On the other hand, genetic analyses of the group revealed an even higher diversity on the molecular level, hidden within the traditional morphospecies. These cryptic species are usually marked by large genetic distances, differentiated biogeographic distribution patterns and ecological adaptations, implying that cryptic species rather than morphospecies represent the level of biological species.

Biological species are defined as reproductively isolated groups of organisms and their identification forms the basis for all studies on the biodiversity, biogeography and ecology of any group of organisms. Yet so far in foraminifera the evolutionary significance of both the morphological as well as genetic diversity and their relation to reproductive isolation remain uncertain, even though this knowledge is indispensable to improve the application of foraminifera as paleoceanographic proxies. E.g. the amount of genetic variation that represents species level divergence instead of intra-population variability is not yet known. In addition, it cannot be objectively stated which morphological features represent species-level differences and which are the result of environmental adaptations. Thus, in order to achieve an objective identification of the level of biological species in foraminifera, culturing experiments are needed to observe reproductive isolation.

We apply a single cell approach to survey the extent of cryptic diversity within foraminifera morphospecies and to examine their distribution and ecological adaptations. In addition, we conduct morphometric analysis in order to establish a connection between morphological and genetic diversity. Furthermore, we carry out mating experiments in order to determine the degree of genetic divergence that corresponds to reproductive isolation and thus represents the level of biological species in foraminifera. In addition, we will try to investigate the existence of different mating types, in order to elucidate the influence of the mating system on the evolution and diversification of the group.

Keywords: foraminifera, genetics, morphology

軟体動物 *Lymnaea stagnalis* の貝殻プロテオーム解析

Proteomic analysis of shell matrix proteins in the pond snail *Lymnaea stagnalis*:
identification of potentially functional proteins

*石川 彰人¹、清水 啓介²、磯和 幸延¹、紀藤 圭治³、藤江 学⁴、佐藤 矩行⁴、遠藤 一佳⁵

*Akito Ishikawa¹, Keisuke Shimizu², Yukinobu Isowa¹, Keiji Kito³, Manabu Fujie⁴, Nori Satoh⁴,
Kazuyoshi Endo⁵

1. 東京大学大学院理学系研究科地球惑星科学専攻、2. 海洋研究開発機構、3. 明治大学、4. 沖縄科学技術大学院
大学、5. 東京大学大学院理学系研究科

1. University of Tokyo, Graduate School of Science, Department of Earth and Planetary Science,
2. JAMSTEC, 3. Meiji Univ., 4. OIST, 5. Univ. of Tokyo, Graduate School of Science

Matrix proteins have important roles in molluscan shell formation, and their amino acid sequences have been characterized for some species. However, the mechanisms of shell formation have barely been clarified. In order to setup a platform for a systematic functional analysis of shell matrix proteins, we performed a combined transcriptome and proteomic analysis of the shell matrix proteins for the pond snail *Lymnaea stagnalis*. We found a total of 203 shell matrix proteins from the shell matrix of *L. stagnalis*. A total of 161 amino acid sequences of them showed sequence similarities to known proteins, including four paralogs of dermatopontin, which was previously reported from the shell matrix of *L. stagnalis*, when searched against public databases, while the remaining 42 showed no similarity to the proteins in the current databases. Next, in order to discriminate 'functional' shell matrix proteins from those that were accidentally entombed in the shells, we compared the levels of expression of these shell matrix proteins between the right side and the left side of the mantle, which makes the shell, underlying assumption being that genuine functional shell matrix protein genes would be more strongly expressed in the right hand side of the mantle in the dextral shells, while there would be no such differential expression pattern for the proteins which were accidentally trapped within the shells. Actin is the most abundant shell matrix protein found in the shell of *L. stagnalis*, but the expression patterns of the actin gene showed no difference between right and left of the mantle. On the other hand, the second most abundant shell matrix protein called Pif-like protein, which is an acidic shell matrix protein identified also in *Pinctada fucata* and *Crassostrea gigas*, showed that its gene is more strongly expressed in the right hand side than the left hand side of the mantle. Our results suggest that Pif-like protein is a functional shell matrix protein, while actin is a protein occluded in the shell accidentally. Finally, we searched for conserved domains of these amino acid sequences. We found various domains and classified them into six categories. Those are extracellular protein, enzyme, cation-interaction protein, polysaccharide interaction protein, proteinase inhibitor, and others. Conserved domains allow us to estimate possible functions of novel shell matrix proteins. Characterization of those novel proteins, and functional analyses of all those proteins identified in this study for this 'model organism' will help understand the mechanisms of biomineralization as well as the evolutionary processes of shell formation in molluscs.

キーワード：トランスクリプトーム解析、プロテオーム解析、生体鉱物、貝殻形成、基質タンパク質

Keywords: Transcriptome, Proteomic analysis, Biomineralization, Shell formation, Matrix protein

琉球列島完新世サンゴ礁コアにおけるマイクロビアライト（礁性微生物岩層）の発見

Reefal microbial crusts found in a Holocene reef sediment core, off Okinawa Island, the Ryukyu Archipelago

*藤田 和彦¹、佐々木 徹¹、知念 正昭¹

*Kazuhiko Fujita¹, Tohru Sasaki¹, Masaaki Chinen¹

1. 琉球大学理学部物質地球科学科

1. Department of Physics and Earth Sciences, University of the Ryukyus

Reefal microbialites are fine-grained, non-skeletal crusts, locally common in late Quaternary coral reef deposits. They are interpreted as microbial carbonates produced by heterotrophic bacterial communities. The carbonate crusts are typically up to 10 cm thick and form the late stage of reef developments, covering on framework skeletons. Major questions still remains unanswered concerning their uneven distributions in space and time, formation process and controlling factors. Here we first report fine-grained, non-skeletal carbonate crusts in a Holocene reef core drilled at Naha New Port area, off west of Okinawa Island, the Ryukyu Archipelago, which are similar in texture and fabric to reefal microbialites reported from deglacial reef deposits from other coral reef regions. The carbonate crusts are up to several centimeters in thickness, and mainly developed in a particular stratigraphic horizon (depth: 5~6 mbsl, age: ca. 7 ka), from which the crust thickness decreased downward and upward. Based on their external surfaces and internal sections, these crusts are classified mostly as a digitate type (thrombolite) and partly as a weakly layered type (stromatolite). Surface elemental analysis showed that crusts are composed mainly of Ca and Mg (Mg calcite). Microscopic observations clearly showed a biological succession from in situ bioeroded corals, overlain by coralline algae and encrusting foraminifers, finally to fine-grained crusts. Silt-sized grains were mainly made of peloids, with subordinate bioclastic and siliciclastic grains. SEM-EDX analysis recognized four different elemental spectrum patterns in the carbonate crusts: sulfur (S)-rich pattern (interpreted as trapped skeletal Mg calcite grains), poor S pattern (precipitated Mg calcite crystals with Mg/Ca ratio of ~0.14), Si and Al common pattern (precipitated Mg calcite crystals with trapped siliciclastic grains or clay minerals), and no Mg pattern (trapped coral aragonite grains). These petrographic and geochemical features are very similar to reefal microbialites found in deglacial reef deposits from other coral reef regions. Some differences in thickness and elemental compositions are possibly related to environmental settings (a volcanic hinterland vs. a mixed carbonate-siliciclastic hinterland).

キーワード：微生物岩、琉球列島、完新世

Keywords: microbialite, Ryukyu Archipelago, Holocene

Syn depositional formation of calcareous nodules on muddy sea floor: elucidating depositional history by C, O, S isotope characterization

Syn depositional formation of calcareous nodules on muddy sea floor: elucidating depositional history by C, O, S isotope characterization

*長谷川 卓¹、媚山 陽介²

*Takashi Hasegawa¹, Yosuke Kobiyama²

1.金沢大学自然システム学系、2.金沢大学理工研究域自然システム学専攻

1.Department of Earth Sciences, Kanazawa University, 2.Division of Natural System, Graduate School of Natural Science and Technology, Kanazawa University

Calcareous nodules are commonly observed in mudstone sequences, however, paleodepositional condition including paleothermometry have never been discussed based on its chemistry without exceptional case such as methane seepage. It partly owes to their complicated depositional history. We characterized dark grey muddy carbonate nodules collected from Cretaceous strata in Hokkaido, Japan with carbon, oxygen, sulfide sulfur and sulfate sulfur isotopes, and demonstrated the syn depositional formation of some nodules. It is notable that initial formation of carbonate "precursor" for all muddy nodules studied here were syn depositional near oxic/anoxic boundary at shallower depth of muddy sea floor.

Some outcrop observations demonstrate calcareous nodules can be formed just below the sea bottom (JBSB). Structure suggesting consolidation JBSB includes burrows that eject calcareous precursor material from nodule. JBSB origin of the nodules consolidated associated with anaerobic oxidation of methane (identified by carbon isotope ($d^{13}C$) values) with sulfate reduction are also demonstrated with oxygen isotope ($d^{18}O$) and sulfur isotope ($d^{34}S$) values. Such nodules show the exactly same $d^{18}O$ values with that of benthic foraminifers. A bivalve fossil found on one of the methane seep nodules preserved aragonite of the shell and yielded close $d^{18}O$ paleotemperature with that of host nodule.

The cross-plot of the $d^{13}C$ and $d^{18}O$ data can emerge "upper limit line" of $d^{18}O$ values showing syn depositional formation. On the other hand, nodules from the Oyubari area characterized with $d^{34}S$ (sulfate) lower than -18 permil indicating JBSB formation had conflicting $d^{18}O$ values. It can be ascribed to the difference of burial depth between sediments of the Haboro and Oyubari areas. Even if precursor of the nodules initially crystallized JBSB, strong compaction during burial would have caused permeation of pore water into the incomplete nodules. Carbon dioxide or bicarbonate ions derived from decomposed organic matter would have caused recrystallization of calcite with $d^{18}O$ as low as -10 permil in the nodule.

キーワード：石灰質ノジュール、酸素同位体比、硫黄同位体比、炭素同位体比

Keywords: calcareous nodule, oxygen isotope, sulfur isotope, carbon isotope

A novel approach in utilising marine methane seep derived bivalve shells for developing local ΔR corrections in deep sea environments

*Kazuhiro Yagasaki¹, Juichiro Ashi¹, Yusuke Yokoyama¹, Yosuke Miyairi¹, Shin'ichi Kuramoto²

1.Atmosphere and Ocean Research Institute, The University of Tokyo, 2.Center for Deep Earth Exploration, JAMSTEC

Shells and corals are commonly used as proxies for sea surface temperature reconstructions in paleoclimatology. These species utilise the dissolved inorganic carbon (DIC) in seawater making it suitable for analysing ocean circulation through the measurement of radiocarbon ^{14}C . It is also possible to estimate the age of the shell through analysing the growth increments within the shells. Yet calibration models in deeper marine environment remains complicated due to the multiple variables that must be considered which include problems such as carbon sinks, ocean current flux and dead carbon effect from seafloor vents.

This study proposes a novel approach in ^{14}C dating shells found in a deeper marine environment (600m) located near methane vents off Tokai using a Single-Stage Accelerated Mass Spectrometer (AMS), to develop a suitable calibration model with a greater goal of observing local marine environmental changes in the deep sea. This method may allow bivalves living in deeper marine environments to be used as future tools for measuring samples located in complicated environments such as deep hydrothermal vents and cold seep vents, possibly leading to assessing local shallow fault movements occurrences in the past.

Sclerochronological analysis of the shells proved difficult since growth increments of deep sea shells are not determined according to simple changes in temperature such is the case with most surface marine shells which often reveal daily, fortnightly or annual bands during seasonal temperature extremes. The radiocarbon ^{14}C age of these bivalve shell measurements ranged between three age groups of 1396 ± 36 - 1448 ± 34 , 1912 ± 31 - 1938 ± 35 and 5975 ± 34 . The ^{14}C age of shells that were alive upon collection and the dissolved inorganic carbon (DIC) in seawater show little difference (around 100 ^{14}C age) indicating that shells are not heavily affected by the dead carbon effect from cold seeps that is of biogenic or thermogenic origin, which can make the age of the shells to become considerably older than their actual age. The novel calibration model used was therefore based on the seawater DIC collected above the *Calyptogena* spp. colony site (1133 ± 31), resulting in the dead shells to be clustered around 1900 Cal AD. This age group proves to be interesting as the Ansei-Tokai earthquake (M 8.4) in 1854 is extremely close to the bivalve colony site. Based on the theory that the bivalve shells formed sometime after the venting of methane fluids, it may support the validity for applying such novel calibrations methods in complicated deep sea environments. Using geological data obtained using visual analysis and sub-seafloor structural analysis that show multiple shallow faults and chaotic sediment structure below the colony site, the *Calyptogena* spp. shells have a strong connection to the coseismic faulting activity and could show potential for radiocarbon dating to be applied on marine samples providing the necessary calibration tools are available.

Keywords: Radiocarbon dating, Active fault, Cold seep, Bivalves, Methane

大型底生有孔虫の高マグネシウム方解石の殻における水温プロキシとしての酸素同位体比および Mg/Ca比

Oxygen isotope and Mg/Ca ratio of high magnesium calcite of benthic foraminifera as a proxy for water temperature

*前田 歩¹、藤田 和彦²、鈴木 淳³、堀川 恵司⁴、吉村 寿紘¹、川幡 穂高¹

*Ayumi Maeda¹, Kazuhiko Fujita², Atsushi Suzuki³, Keiji Horikawa⁴, Toshihiro Yoshimura¹, Hodaka Kawahata¹

1.東京大学 大気海洋研究所、2.琉球大学 理学部物質地球科学科、3.独立行政法人産業技術総合研究所地質情報研究部門、4.富山大学 大学院理工学研究部

1.Atmosphere and Ocean Research Institute, The University of Tokyo, 2.Department of Physics and Earth sciences, University of the Ryukyus, 3.Geological Survey of Japan National Institute of Advanced Industrial Science and Technology, 4.Graduate School of Engineer and Science, University of Toyama

Although metal/Ca or oxygen isotopes ($\delta^{18}\text{O}$) of organic precipitated calcium carbonate (calcite and aragonite) of marine species have been used to estimate paleotemperature, the relationships between components of large benthic foraminiferal shells (high-magnesium calcite) and seawater temperature has yet to be established. We investigated the possibility as a proxy for seawater temperature through culture experiments of three species of reef-dwelling large benthic foraminifera in a laboratory. Three species found commonly in the western Pacific were selected including perforate species *Calcarina gaudichaudii*, and imperforate species *Amphisorus kudakajimensis*. All of them are host to algal symbionts. They grew sufficiently during culture experiments and showed a maximum in terms of shell length and weight at 27°C and 29°C while they were significantly small at 30°C. Mg/Ca ratios of three species in similar range showed high correlation to water temperature, suggesting Mg/Ca ratios as a precise proxy for paleo-temperature in shallow-reef environment. In terms of $\delta^{18}\text{O}$, *C. gaudichaudii* showed strong correlation versus temperature while $\delta^{18}\text{O}$ of *A. kudakajimensis* showed less significant correlation possibly caused by poor growth of unhealthy individuals. The trends in the temperature and oxygen isotope ratios were similar for both species, suggesting the potential of oxygen isotope ratios in the tests of reef-dwelling foraminifera as a paleo-thermometer. Species-specific calibration may be necessary for the use of $\delta^{18}\text{O}$ of reef-dwelling large benthic foraminifera as a proxy of paleo-temperature.

キーワード：高マグネシウム方解石、水温プロキシ、大型底生有孔虫

Keywords: high magnesium calcite, proxy for water temperature, large benthic foraminifera