a result of lower wind stress since the LIA. The absence of the "Suess effect" (the lowering of Δ^{14} C values from the late 19th century as fossil fuel burning released ¹⁴C-free CO₂) in these records is interpreted as a long-term change in ocean circulation within the southwestern Pacific since the late 1800s (*38*). Rising Δ^{14} C values since the late 1800s (*38*). Rising Δ^{14} C values since the 1880s indicate a greater influx of younger surface waters, and this reduced oceanic "reservoir effect" overrides the anticipated Suess effect (Fig. 3F). We interpret this shift as a weakening of the wind-driven SEC since the late 19th century, which reduced the transport of upwelled equatorial ¹⁴C-depleted water to the GBR corals.

Our SSTA reconstructions suggest that a stronger latitudinal temperature gradient, relative to the present, may have prevailed during the LIA. General circulation model experiments (9) indicate that the global climate responds to an exaggerated latitudinal temperature difference by intensifying the largescale atmospheric dynamics of the Hadley circulation cells in order to maintain thermal balance. During the LIA, if warm tropical Pacific SSTs encouraged evaporation (as implied by the coral δ^{18} O records) and a stronger Hadley circulation dried the subtropics, then global average water vapor should have increased. A critical question is where that moisture was transported. Generally, snow cover increases with a strong latitudinal temperature gradient, in part because of cooler high-latitude conditions, but also as a result of a greater atmospheric connection between the tropics and extratropics (10). Our results imply that the tropical oceans may have played an important role in driving the LIA glacial expansion during the repeated advances between 1600 and 1860 (1). Cooling and abrupt freshening of the tropical southwestern Pacific coincided with the weakening of atmospheric circulation at the end of the LIA, when glaciers worldwide began to retreat.

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- 16. The composite reconstructions are the average of all records, after normalization relative to the longest continuous record (HAV-01B). The δ¹⁸O composite is based on eight cores; the Sr/Ca and U/Ca composites are constructed from seven (PAN08B excluded). Error bounds were calculated using 95% confidence intervals for each 5-year period. Periods of higher U/Ca variability coincide with die-offs in two of the cores in 1782–1785 and 1817. Data are available at www. ngdr.noaa.gov/paleo/pubs/hendy2002.
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Microbial Activity at Gigapascal Pressures

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We observed physiological and metabolic activity of *Shewanella oneidensis* strain MR1 and *Escherichia coli* strain MG1655 at pressures of 68 to 1680 megapascals (MPa) in diamond anvil cells. We measured biological formate oxidation at high pressures (68 to 1060 MPa). At pressures of 1200 to 1600 MPa, living bacteria resided in fluid inclusions in ice-VI crystals and continued to be viable upon subsequent release to ambient pressures (0.1 MPa). Evidence of microbial viability and activity at these extreme pressures expands by an order of magnitude the range of conditions representing the habitable zone in the solar system.

Microbial communities adapt to a wide range of pressures, temperatures, salinities, pH, and oxidation states. Although the chemical and physical conditions in these extreme environments are reasonably well constrained, the consequence of these physical parameters on the physiology of microbial communities is not well understood. Significant attention has been focused on the effects of high and low temper-

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*To whom correspondence should be addressed. Email: sharma@gl.ciw.edu (A.S.); j.scott@gl.ciw.edu (J.H.S.) ature on physiology (1, 2). There is some evidence that elevated pressure may also manifest interesting effects on cellular physiology (3, 4). For example, recent studies report that elevated pressure may lead to enzyme inactivation, compromise cell-membrane integrity, and suppress protein interactions with various substrates (3-6). Whereas the cumulative impact of these pressure-induced effects on microbial metabolism and physiology is an inhibition in growth rate and cellular division in microorganisms (7-9), exactly how these factors affect intact cells is not well understood (4, 10).

Numerous high-pressure studies have been conducted on biological systems; however, these have been either on individual biomolecules or "indirect" measurements of cellular growth. In most instances, biological techniques for high-pressure studies (>200 MPa or >2 kbar) do not allow for in situ measurements and observations. Typical experimental designs for high-pressure studies extract samples for subsequent analysis at ambient pressure (4, 11). It would be useful to directly examine the effects of pressure on microbial viability to understand physiological responses and cellular biochemistry (10). Here, we adapted diamond anvil cells (12) to explore the effects of high pressure on microbial life.

We used the rate of microbial formate oxidation as a probe of metabolic viability. The utilization of formate by microorganisms is a fundamental metabolic process in anaerobic environments (13) that can be easily monitored via molecular spectroscopy. We monitored in situ microbial formate oxidation for Shewanella oneidensis MR1 (12) to pressures of 1060 MPa, at room temperature (25°C), using Raman spectroscopy (12) in diamond anvil cells (Fig. 1). The Raman data revealed rapid oxidation of formate determined from the reduction in the peak height of C-H vibration mode (14) (Fig. 1). Microscopic observations to pressures of 1060 MPa confirmed continued cell viability (intact and motile) over a duration of more than 30 days. As anticipated, formate oxidation rates were significantly reduced upon the addition of sodium cyanide (0.2 M). Cyanide acts as an inhibitor of respiration by acting in competition as an alternate electron acceptor. This result indicates that the oxidation of formate is coupled to biological respiration. Experiments with heat-inactivated MR1 cells, inactivated at 50°C for ~ 1 hour until lysis was observed, did not exhibit any detectable chemical changes (Fig. 2). Both cell-free and dead cell controls indicate



that formate oxidation was exclusively a result of biological activity. The decrease in the formate oxidation rates with increasing pressure in living cells, as indicated from the Raman data (Fig. 2), may simply reflect a decrease in the cell numbers (15).

There are studies from Antarctic ice that point to it as a viable ecosystem even when the availability of liquid water is limited due to solidification (16-18). We have extended these experiments to pressures where ice-VI becomes stable and explored how the pressure-sensitive bacteria (4, 19), Shewanella MR1 and Escherichia coli, respond to this change of state, in a nutrient-rich medium (20). Optical observations at pressures ranging from 40 to 1200 MPa, lower than ice-VI formation (21), detect a gradual decrease in cell motility, an increase in cellular adhesion to surfaces, decrease in cell numbers and reduction in metabolic activity (Fig. 2). Furthermore, methylene blue, when added as an electron acceptor to test for respiration at elevated pressures, remained reduced throughout the experiment, indicating that bacterial respiration continued at elevated pressures (22). At ~1250 MPa, the formation of ice-VI was instantaneous. Initially, mosaics of ice-VI crystals were observed separated by thin "veins" of organic-rich fluid containing the bacteria (Fig. 3). However, over time the ice melted partially along the interstitial veins and developed numerous organic-rich fluid inclusions containing microbial clusters (Fig. 3) [Web movie 1(12)]. These textural changes in the ice, observed to pressures of 1600 MPa, might either be the consequence of the phase separation associated with the formation of ice in any organic-rich medium, or the result of the microorganisms protecting ice-sensitive cellular components (23, 24). Upon subsequent lowering of pressure to melt the ice, cell survival and viability at the higher pressures were determined by direct cell counts (Fig. 2) on both E. coli [ranging from 10^6 cells (at 0 hours) to 10^4 cells (after 30 hours)] and Shewanella MR1 [ranging



Fig. 1. Raman spectra of the formic-biological system (A) shown with the vibration peaks of formic and diamond anvils used in this study. The outlined boxed region is shown at higher resolution (B) to quantify the successive decrease in the peak intensity of the C-H stretch of formic acid at pressures of 68, 142, and 324 MPa. The equivalent formate concentrations (C), corresponding to each peak height change, are based on comparisons with a known calibration curve. All experiments were performed at 25° C, with diamond anvil cells (32) with gold-lined sample chambers (12). Pressures were estimated using Raman shifts in quartz used as an internal calibrant (14).

Fig. 2. Shown are formate oxidation rates for experiments at high pressures for the first 48 hours. Whereas biological formate oxidation rates exhibit an inversely exponential relationship with pressure, the cyanide-inhibited system shows a lower formate oxidation rate.

from 3×10^6 cells (at 0 hours) to 10^4 cells (after 30 hours) (20)] (Fig. 3). Distinct changes in cellular morphology were also observed, including a significant increase in cell size and polar invaginations, both clear indications of arrested cellular division (25, 26). Upon a subsequent increase of pressure to 1680 MPa, the veins of organic-rich fluids developed a "stringy" texture as well as an increase in fluorescence background in Raman spectra. This could be an indication of cellular lysis, because no live cells were observed upon return to ambient pressure.

Adaptability of *Shewanella* MR1 as well as *E. coli* in these experiments indicates that microorganisms can continue to metabolize substrate at pressures far beyond those previously reported in the literature (3, 4, 10). Although an evolutionary component to the adaptation of microbial communities to temperature and salinity is well known (27), whether there might

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be any evolutionary component for pressure adaptation is still in question. Shewanella MR1 belongs to a genus that contains a number of piezophiles; however, E. coli clearly does not. Despite this, there is evidence that exposure of E. coli to pressures up to 800 MPa selects a population of cells less sensitive to pressure inactivation (19). Furthermore, it is well known that the increase in pressure tolerance is also associated with heat tolerance (19). Our observations on pressure-sensitive bacteria may indicate that piezo-tolerance of these bacteria could be a protective mechanism in response to stress (4, 28, 29). Whether this is a short-term or an evolutionary adaptation remains unanswered.

These results imply that pressure may not be a significant impediment to life. The maximum pressure explored in this work is equivalent to a depth of ~ 50 km below Earth's crust, or ~ 160 km in a hypothetical ocean. The pressures encountered at the depths of

Escherichia coli





Fig. 3. Microbial activity and viability in ice-VI. Upon ice nucleation (0 hours, 1400 MPa), organic fluid veins (*v*) filled with bacteria (*Shewanella* MR1 on the left, *E. coli* stained with methylene blue on the right) appear. After \sim 1 hour, textural changes in the ice occur, defined by the formation of organic-rich inclusions (*i*) containing motile bacteria [Web movie 2 (*12*)]. Viable (methylene blue in solution remains colorless indicating respiration) and countable bacteria (*b*) were observed upon subsequent lowering of pressure [Web movie 3 (*12*)].

thick ice caps and deep crustal subsurface may not be a limiting factor for the existence of life. This suggests that deep (water/ice) layers of Europa, Callisto, or Ganymede, subduction zones on Earth, and the polar ice caps of Mars might provide viable settings for life unhindered by the high pressures (30, 31).

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