Long-Distance Electron Transport in

Multicellular Freshwater Cable Bacteria

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Abstract

Filamentous multicellular cable bacteria perform centimeter-scale electron transport in a process that couples oxidation of an electron donor (sulfide) in deeper sediment to the reduction of an electron acceptor (oxygen or nitrate) near the surface. While this electric metabolism is prevalent in both marine and freshwater sediments, detailed electronic measurements of the conductivity previously focused on the marine cable bacteria (*Candidatus* Electrothrix), rather than freshwater cable bacteria, which form a separate genus (*Candidatus* Electronema) and contribute essential geochemical roles in freshwater sediments. Here, we characterize the electron transport characteristics of *Ca.* Electronema cable bacteria from Southern California freshwater sediments. Current-voltage measurements of intact cable filaments bridging interdigitated electrodes confirmed their persistent conductivity under a controlled atmosphere and the variable sensitivity of this conduction to air exposure. Electrostatic and conductive atomic force microscopies mapped out the characteristics of the cell envelope's nanofiber network, implicating it as the conductive pathway in a manner consistent with previous findings in marine cable bacteria. Four-probe measurements of microelectrodes addressing intact cables demonstrated nanoampere currents up to 200 μm lengths at modest driving voltages, allowing us to quantify the nanofiber conductivity at 0.1 S/cm for freshwater cable bacteria filaments under our measurement conditions. Such a high conductivity can support the remarkable sulfide-to-oxygen electrical currents mediated by cable bacteria in sediments. These measurements expand the knowledgebase of long-distance electron transport to the freshwater niche while shedding light on underlying conductive network of cable bacteria.

Significance

Cable bacteria are multicellular filaments composed of up-to-thousands of end-to-end cells and are found worldwide in both marine and freshwater sediments. Remarkably, these cells gain energy from a long-distance electron transport process that carries electrons generated by sulfide oxidation in deeper sediment layers to drive oxygen reduction near the sediment-water interface. This electric metabolism requires an unusually high electronic conductivity, previously thought impossible in natural biological materials. However, the underlying mechanism(s) remain poorly understood, and previous characterization of the conductivity largely focused on marine cable bacteria. Here, we characterize and quantify the electronic conductivity of freshwater cable bacteria from sediments in Southern California, with emphasis on the role of the conductive periplasmic fiber network in routing electron transport along cables.

Introduction

Electron transfer reactions are fundamental to biological energy conversion. Respiratory organisms, for instance, gain energy by routing electron flow from an electron donor to a terminal electron acceptor through the network of reduction-oxidation (redox) cofactors that constitute the cellular electron transport chain (1). While the length scales of biological electron transfer events were long thought to be limited to nanometer distances and confined within individual cells, recent observations of fast long-distance extracellular and intercellular electron transport in microbial communities have upended this view (2). In fact, with the discovery of micron-scale electron transport via cytochrome nanowires from metal-reducing bacteria (3, 4), conductive bacterial biofilms (5, 6), inter-species electron

transfer in microbial consortia (7), and multicellular cable bacteria (8), the length scales of

microbial electron transport observations have jumped from the nanometer to centimeter

scale during the last 15 years.

The novel multicellular filamentous cable bacteria (CB), which belong to the

Desulfobulbaceae family of Deltaproteobacteria (8), are a particularly interesting system

for understanding the limits of biological electron transport given that they are the only

known living structures capable of macroscopic (up to centimeter-scale) electron

conduction. This macroscopic electron transport allows CB to gain energy from coupling

sulfide oxidation by cells that dwell in the deeper sulfidic zone to oxygen reduction by cells

near the surface sediment in both marine and freshwater sediments (8-11). The energy

harnessed from this process allows CB to form a dense matrix (up to hundreds of meters

per cm^2) that composes a majority of the biomass in sediment when blooming (12-16).

While the molecular pathway of the conductive network responsible for centimeter-scale

electron transport along the cables is not yet understood, the genomic basis and components

allowing CB to carry out the two spatially-separated half-reactions are just beginning to

come into focus. CB appear to oxidize sulfide by reversing the canonical sulfate reduction

pathway, and are capable of sulfur disproportionation, while oxygen reduction is

hypothesized to be driven by periplasmic cytochromes without energy conservation due to

the absence of terminal oxidases (17, 18). The electrogenic sulfur oxidation and its

coupling to distant redox events by CB significantly affects sulfur cycling and other

essential geochemical processes, including nitrogen and metal cycling in global sediments

(18, 19).

But what is the conductive pathway responsible for long-distance electron transport along

an entire CB filament? The answer appears to lie in the unusual cell envelope of CB, where

up-to-thousands of cells share a common outer membrane and parallel longitudinal ridges

run along the surface of the entire filament (8, 20). These ridges reflect the presence of a

network of long nanofibers (33 – 67 nm diameter) along the entire length of each cable in

the periplasmic space between the common outer membrane and the cellular inner

membranes (20, 21). Initial electrostatic force microscopy of these ridges showed striking

electrostatic contrast (8), implicating the periplasmic nanofibers as the current carrying

structures in marine CB filaments. More recently, the conductivity of intact air-dried

marine CB filament and chemically extracted sheaths containing the nanofiber network

was measured directly by various electronic techniques (22-24). Tens of nanoampere

currents could be observed along both intact cable filaments and extracted nanofiber

sheaths spanning hundreds of microns between microelectrodes in N_2/vacuum

environments, but the conductance declined rapidly upon exposure to ambient air (22).

Conductive atomic force microscopy (C-AFM) with stiff tips that scrape outer cell surface

layers revealed that the conductive path indeed correlates with the network of individual

periplasmic nanofibers (23). For the marine CB nanofibers, conductivities spanning the 10^{-2}-10^1 S/cm range, with the upper end reaching as high as 79 S/cm, have been estimated

(22); these values represent a record high for any biological structure. Furthermore,

temperature-dependent electronic measurements suggest a thermally activated Arrhenius-

type charge transport process with a low activation energy, suggesting that the CB

conductive properties are comparable to organic semiconductor materials (24). The

composition of the CB conductive nanofibers remains unknown, but recent evidence points

to yet-unknown core proteins rich in sulfur-ligated nickel cofactors that mediate electron

transport through an enigmatic physical mechanism, and that the conductivity is

diminished upon oxidation of the Ni/S groups (25).

On the basis of 16S rRNA gene sequencing, the CB clade consists of two novel candidate

genera: the mostly marine *Candidatus* Electrothrix and mostly freshwater *Candidatus*

Electronema (26). In addition, a more distant group of CB has been identified in ground

water environments (17, 27). However, the abovementioned electronic characterization of

CB conduction largely focused on marine CB, with the exception of two-probe

measurements as part of a recent comparative study of diverse CB that included one

freshwater CB enrichment culture (21). Freshwater sediments can contain CB populations

at densities comparable to their marine counterparts, and their activity strongly influences

the sediment geochemistry (11). Indeed, freshwater CB play an essential role in a cryptic

sulfur cycle by stimulating sulfate reduction (28), and have been harnessed to significantly

reduce methane emissions from rice-planted soil (29). From a materials perspective,

investigations of freshwater CB may also uncover conductive biomaterials suited for

applications in less saline environments that support higher electric fields than marine

environments (28, 30). Here, we report on the recovery, phylogenetic analysis, imaging,

and electronic characterization of *Ca.* Electronema CB from Southern California

freshwater sediments using atomic force and multi-probe transport techniques.

Materials and Methods

122 **Sediment sampling and incubation.** Surface sediment and pond water was collected from

123 two sampling sites. The first site, the Ballona Freshwater Marsh pond (33°58'15"N,

118°25'51"W), is a restored wetland that was previously filled and farmed for decades and is located 18 miles from downtown Los Angeles. The pond is seasonal (accessible ~ December to early June during the wet season). The second sampling site is Mungi Lake (34°6'6.192"N, 118°0'44.2224"W), located in the Los Angeles River Basin in the city of Arcadia, California. The sediment samples were collected from shallow water because the dissolved oxygen (DO) typically declines to zero during the summer months at depths greater than 5 meters (31). In the wintertime, less water stratification results in a higher DO concentration that can support cable bacteria growth in the lake bottom sediment. Mungi Lake samples were accessible year-round despite the lake shrinking during the dry season.

The top ~5 cm of sediment from both sites was sampled approximately every 3 months. Upon returning to the laboratory, the sediment was homogenized, sieved through a 5 mm-pore-size sieve, and carefully filled into glass jars, avoiding air bubbles in the sediment. The sediment jars were then incubated in a water bath at 15 °C, submerged in circulating, air-saturated water collected from the respective sites.

Cable bacteria filaments pickup. Filamentous bacteria were detectable after 2 weeks of incubation; occasionally, the filament density was high enough to be directly observed in sediment cracks (Figure 1). Over the incubation period, small sediment cores were taken from the sediment jars using plastic straws (8 mm diameter, 80 mm length) and transferred into small petri dishes filled with DI water. Long filaments were hand-picked under a dissection microscope using glass hooks made from capillary tubes, then washed successively at least three times in MilliQ water droplets. This extraction of cable bacteria was performed in a glove bag under an N_2 atmosphere.

16S rRNA gene sequence-based phylogenetic identification. Under a dissection microscope, three long filaments from each sampling site were washed and transferred into a sterile PCR-tube. Cells were lysed by freeze-thaw three times as described in a previous study (13). The lysates were directly used for PCR using the primer pair 8F and DSBB/1297R (32). PCR was performed according to manufacturer's instructions for the Q5® Hot Start High-Fidelity 2X Master Mix (New England Biolabs, USA). After 30 s at 98 °C for initial denaturation, PCR was performed with 35 cycles, consisting of 10 s at 98 °C, 30 s at 65 °C, and 45 s at 72 °C, followed by a final step of 2 min at 72 °C. The PCR amplicons were mixed and cloned using a Zero Blunt™ TOPO™ PCR Cloning Kit. Clones with a correct insertion were commercially sequenced (GENEWIZ, USA) using the vector primers M13F and M13R. Sequences were assembled in Sequencher 5.4, and their quality was checked manually. Alignment was performed in the ARB software package (arb-6.0.6) (33) using aligned sequences from SILVA database SSU 132 for reference (34), and manually refined subsequently. The 16S rRNA phylogenetic trees were generated by ARB. Trees that were built by the ARB neighbor joining algorithm and the maximum likelihood algorithm (RaxML) with a 50% positional conservation filter, with 1000 bootstrap replicates in all trees, were compared to make sure the trees had no conflicting branching. Representative sequences from this study were deposited in GenBank with accession numbers (OR244071 - OR244134).

Transmission electron microscopy (TEM). Bundles of filaments were washed thoroughly, deposited in small droplets of MilliQ water, and the CB filament-containing water droplets were then transferred and air-dried on PELCO® hexagonal Carbon-Formvar coated copper grids. The filament samples were unstained. High vacuum TEM was

performed by using an FEI Talos™ F200C (Thermo Fisher, USA) Electron Microscope at

80 kV, and the images were captured by a CCD camera.

Electrostatic Force Microscopy (EFM). EFM measurements were performed using an

Oxford Instruments Asylum Research Cypher ES atomic force microscope (AFM).

Substrates were affixed and electrically connected to sample discs. The CB were extracted

from sediment using fine glass hooks, washed several times in DI water, and placed on the

prepared substrates to dry. The sample discs were wired to the grounding pin of the AFM

upon loading. Si probes with a Ti/Ir (5/20) coating, a nominal resonant frequency of 75

kHz (58-97 kHz range), a nominal spring constant of 2.8 N/m (1.4-5.8 N/m range), and a

tip radius of 28±10 nm were used for all EFM measurements. High resolution EFM scans

were performed following initial 'wide view' tapping AFM imaging to locate the CB

filaments. For each EFM scan, a first pass scans the conductive probe in standard tapping

without an applied tip bias, followed by a second 'nap' pass that retraces the first pass

topography at a constant height above the sample. EFM was performed at different nap

pass heights and tip voltage biases to arrive at optimal conditions and evaluate the voltage

dependence. Electrostatic forces between the tip and sample shift the resonance frequency

of the probe, resulting in a phase change of the cantilever, which was quantified for pixels

representing specific features (e.g. nanofibers). The phase differences for each image (e.g.

specific tip voltage bias) were averaged and their standard deviation taken.

Current-Voltage (I-V) measurements on Inter-Digitated Array (IDA)

Microelectrodes. A 3 µm gap-size Au IDA (IDA Model#012129, ALS Co., Ltd, Japan)

and a Pine® WaveDriver 200 Bipotentiostat/Galvanostat were used for electrical

measurements. The IDA consists of two individually addressable arrays of interdigitated

gold microelectrodes (length, 2 mm; width, 3 µm; thickness: 100 nm). Copper wires were

affixed to these two electrodes with silver paint (Product #: 16035, TED PELLA, Inc.) and

covered with an epoxy layer at the contact points. The wires were then connected to the

potentiostat, which swept the potential from -0.2 V to 0.2 V at a 10 µV/s sweep rate. All

measurements were performed in an anaerobic chamber (95% N_2 and 5% H_2) (Bactron 300,

Sheldon Manufacturing, Inc. Cornelius, OR). Control measurements prior to

measurements of CB included open circuit (without water or cable bacteria), I-V

measurements of an IDA on which a water droplet has been deposited (no cable bacteria),

and I-V measurements made after the MilliQ water droplets were dried on the IDA (no

cable bacteria). After these controls, CB filaments were washed and deposited in a small

droplet of MilliQ water on the IDA surface. The CB filament-containing water droplet was

then left to dry prior to the I-V measurement.

Conductive Atomic Force Microscopy. Conductive atomic force microscopy (C-AFM)

was performed using an Oxford Instruments Asylum Research Cypher ES AFM. CB

filaments were placed and dried across the insulating gaps between Au (Model#012129, 3

µm gap) or ITO (Model#012128, 5 µm gap) IDA microelectrodes from ALS Co., Ltd,

Japan. For C-AFM current-mapping, ultra-sharp conductive single crystal diamond probes

(AD-2.8-SS, Adama Innovations, Ltd), with < 5 nm tip radius, were used to image Mungi

Lake CB in contact mode under N_2 perfusion. C-AFM current mapping was performed

with a 0.1 Hz scan rate, 300 nN force, and a constant 5 V DC voltage bias applied between

the probe and IDA. The probes and imaging parameters were selected to minimize

disturbance to the sample while simultaneously eroding the insulating outer membrane

enough to electrically probe the underlying periplasmic fibers. Prior to C-AFM, high

resolution tapping mode imaging was performed to spatially correlate the topography of

the CB nanofibers to conductivity.

Point current-voltage (I-V) measurements were also performed to measure conduction

along Ballona Marsh CB between the IDA microelectrodes and C-AFM tip (Si Probe

ASYELEC.01-R2, Oxford Instruments, with a conductive Ti/Ir (5/20) coating and 28±10

nm tip radius) probing CB spots above the insulating quartz IDA gap under ambient

conditions. At each point, the IDA voltage was swept for 5 consecutive cycles in the -10–

10 V range with a 0.1 cycle/second scan rate, while the C-AFM probe was grounded and

held in contact with the CB surface at a 450-510 nN force. All point-IV curves shown,

including the insulating quartz background curve, are the average values taken from the 5

voltage cycles.

Four-probe Device Fabrication. Four-probe (4P) microelectrode devices were designed

inhouse and fabricated by the University of California San Diego Nano3 cleanroom

foundry service. Each device consisted of a series of parallel electrode bands. The bands

were 3.5 mm long, 100 μm wide, and each one was connected on alternating sides to a 1.5

mm by 5 mm contact pad. The electrode band separation distances were variable, including

multiple 20 μm and 200 μm gaps to assess different CB lengths. The fabrication process is

briefly outlined as follows. 100 mm diameter and 0.5 mm thick SiO_2/Si wafers (Wafer

model: 1583, University Wafer, Inc.) were used as substrates. Solvent cleaned wafers were

coated with photoresist (PR) and a laser writer was used to project repeats of the device

pattern onto the PR with UV light. E-beam evaporation was used to deposit a 5 nm Ti

adhesion layer followed by a 100 nm Au layer onto the wafers. Solvents were used to

remove the excess Au/Ti layer and PR, leaving just the electrode patterns. As a final step,

the wafers were coated in PR and each device was then diced into an 18 mm x 18 mm chip.

Four-probe Electrical Measurements. Four-probe (4P) measurements were made using

an Agilent 4156C Precision Semiconductor Parameter Analyzer along with a Signatone

1160 series probe station. Thin copper wires (Product #: 1227, Train Control Systems)

were electrically connected to the 4P contact pads with silver paint (Product #: 16035, TED

PELLA, Inc.) and alligator clips attached to tungsten probes were used to connect the 4P

device to the analyzer. Mungi Lake CB filaments were placed and dried across the

electrodes of our devices. To deal with the high degree of variability between individual

filaments, they were placed one at a time and I-V measurements performed after each

deposition until current (from a conductive filament) was observed. To make the 4P

measurement, a set of four electrodes on the device, with either a 20 μm or a 200 μm inner

probe separation distance, were each wired to a source monitor unit within the analyzer.

Current was swept between the outer two electrodes from -1 nA to 1 nA in step sizes of

100 pA. The voltage difference between the inner two electrodes (V_{inner}) was then recorded

at each current step. Auto range voltage sensitivity with a voltage compliance of 5 V and

an integration time of 1 PLC were used. A 2 second delay was used between each V_{inner}

measurement to eliminate capacitive current contributions. Current sweeps were performed

three times for the two inner probe gap sizes used. Average V_{inner} values for each current

step were obtained from the three sweeps along with an error bar indicating the standard

deviation. All 4P measurements were performed under an N_2 environment.

The single periplasmic fiber conductivity for the CB filaments measured at each inner

electrode gap size was calculated from the formula $\sigma_{fiber} = \frac{L}{A\,N\,R_{cable}}$, where L is the

known spacing between the two inner electrodes, R_{cable} is the CB filament resistance, N is the average number of periplasmic fibers per cable bacteria filament, and A is the average cross-sectional area of a single periplasmic fiber. An average $N = 35$ fibers per CB was determined from TEM imaging. A was calculated from $\pi \left(\frac{d}{2}\right)^2$, with the periplasmic fibers assumed to be cylinders with average diameter $d = 50$ nm estimated from TEM imaging. R_{cable} was obtained as the inverse slope of best fit lines for the first CB current vs V_{inner} scan. Since current in each CB is carried by multiple periplasmic fibers, the $N\,R_{cable}$ factor, which represents the resistance of a single fiber, is based on a simplistic equivalent circuit model of a series of N identical parallel fiber resistors. Open circuit 2-probe controls (for each pair of electrodes on the 4P device) were performed prior to placing CB on the devices.

Results

1. Freshwater Cable Bacteria from Southern California Sediments

1.1 Identification of cable bacteria sampling sites. We identified two freshwater sediment sampling sites for recovery and subsequent measurements of cable bacteria (CB) in the Los Angeles area. The first site, Ballona Freshwater Marsh (Figure 1A), is a shallow seasonal freshwater pond which is heavily vegetated by aquatic plants that supply sufficient organic matter to the pond bottom sediment in the wet season (December to early June), and is dried out from approximately late June to December every year. The sediment profile from the site was brown-orange-colored at the surface and dark in color with a sulfide-smell below the surface, indicating a sharp oxic to sulfidic transition in the sediment. The second sampling site is Mungi Lake (Figure 1A), which is primarily fed by washes from

surrounding areas in Eastern Los Angeles and therefore is precipitation dependent. Mungi

Lake has a much thinner and more densely packed sediment layer, compared to the Ballona

Freshwater sediment, as the lake was converted from a gravel pit, with almost no aquatic

plants and therefore no roots in the bottom surface sediment.

1.2 Southern California cable bacteria morphology: Thin silk-like filaments were

directly observed by eye within cracks of sediment at both sampling sites. Following

laboratory incubation in circulating water to enrich for CB (*Materials and Methods*), a high

density of CB filaments was observed (Figure 1B). When observed with optical

microscopy, the filament diameters varied from 0.7 µm to 3 µm, with thinner CB filaments

usually observed within Mungi Lake sediments. Other than the diameter, there were no

obvious morphological difference between the CB from the two sites. Occasionally, darker

cells with slightly larger diameters than neighboring cells could be observed at seemingly

random locations along the cable filaments (Figure 1C).

Transmission electron microscopy (TEM) imaging revealed that the freshwater CB display

the characteristic previously observed parallel longitudinal periplasmic nanofibers that run

along the along the entire CB filaments (Figure 2A). The estimated width of the fibers

ranged from 49-57 nm, which is comparable to those of the marine cables (ca. 50 nm)

(20). TEM imaging of CB recovered from the sediments occasionally revealed degraded

cells along the filaments where only an 'empty cage' of the parallel nanofibers remained,

appearing as darker material reflecting a higher electron-dense composition (Figure 2B)

and likely corresponding to the enlarged cells in Figure 1. Imaging of the freshwater CB

also revealed small intracellular electron-dense globules, consistent with polyphosphate

granules commonly observed in marine cables but not in freshwater CB previously (11,

18).

1.3 16S rRNA gene phylogeny of Southern California freshwater cable filaments and

their associated bacteria. Three Ballona sediment CB filaments and three Mungi Lake

sediment CB filaments were analyzed for 16S rRNA gene-based phylogenetic

identification (Figure 3). All the Ballona sequences were within the family

Desulfobulbaceae, and the majority (24/27 total Ballona sequences) grouped together with

the freshwater CB sequence of *Ca.* Electronema palustris (96.7% - 97.3% similarity) (26).

The three representative sequences slightly differ from each other indicating coexistence

of different species in Ballona sediment. The few non-CB sequences all affiliated to

sulfate-reducing bacteria from freshwater sediments (35, 36), with one sequence closely

related to polycyclic aromatic hydrocarbon (PAH) degrading bacteria from a novel genus

Desulfoprunum benzoelyticum (37).

The majority of Mungi Lake sequences affiliated within *Deltaproteobacteria* (26/35

clones). Three CB clones are closely related (> 99% similarity) to uncultured sequences

(accession number FQ658891 and FQ658831) from a PAH-contaminated wetland

collecting highway road runoffs (38). These Mungi Lake CB sequences share only 96.5%

similarity to the *Ca.* Electronema nielsenii and are even further distantly related to the

sequences of Ballona CB and *Ca.* Electronema palustris. Most of the non-CB Mungi Lake

clones (18/26 clones) grouped together to sequences identified from an altiplanic cold lake

in South America (39). Also in this group was a cluster (representative sequence:

AJ389622, similarity 98.5%) resembling uncultured sulfate-reducers from a meromictic

lake in Switzerland (40). Other Mungi Lake sequences affiliated to uncultured *Geobacter*

clones (EF192881) from a temperate artificial lake which receives mining wastes, and a polychlorinated-dioxin-dechlorinating community that can also perform oxidative degradation of the dechlorinated products, enriched from a polluted river sediment enrichment (41). Outside the *Deltaproteobacteria*, there were few sequences belonging to uncultured *Chlorobi* group (99.15% identity). One Mungi Lake clone was identified as *Ignavibacteria* (98.9% similarity to CP053446), which have been consistently found as the dominant flanking community in freshwater anammox bioreactors, presumably reducing nitrate to nitrite and degrading peptides, amino acids and EPS produced by anammox bacteria (42). One Mungi Lake clone shares 99.2% similarity to the uncultured sub group 1 of *Acidobacteria* from an Australian pasture soil (2-4 cm) (43); other Mungi Lake sequences include one that loosely related to a potential *Hydrogendentes* bacterium (44) and 4 clones that reside deep in the branch.

2. Electronic Characterization of Southern California freshwater cable bacteria

2.1 Electrostatic Force Microscopy. Electrostatic force microscopy (EFM) was used to visualize the electric field variations on Ballona CB filaments using a dual-pass measurement approach. Topographic (Figure 4A) and phase (Figure 4B) tapping mode images of CB were generated during the first pass, revealing in high resolution the surface ridge pattern along two consecutive CB cells. During the second pass, performed with a tip voltage bias at a set height (50 nm) above the retraced CB topography, the phase image (Figure 4C) captured the electric force gradient distribution as a result of the change in the resonant frequency of the probe arising from the CB-tip electrostatic interaction. Consistent with a previous report (8), the ridges and associated nanofibers exhibited conspicuously stronger electric forces relative to inter-ridge regions (Figure 4C), hinting at the unique

electronic properties of the underlying nanofibers. The voltage dependence of the CB-tip

interaction was also investigated, revealing that the phase difference between the elevated

ridges and inter-ridge regions is proportional to the tip voltage squared (Figure 4D).

2.2 Current-voltage measurements of freshwater cable bacteria across interdigitated

array microelectrodes. For an initial assessment of freshwater CB conduction, current-

voltage (I-V) measurements were performed along Ballona CB filaments bridging

interdigitated array (IDA) microelectrodes in the oxygen-free environment of an anaerobic

chamber (95% N_2, 5% H_2). In contrast to controls (open circuit IDA and deposited drops

lacking CB), which showed no detectable current, multiple devices with deposited CB

showed linear up-to-μA currents along intact filaments in response to voltage in the -0.2-

0.2 V range (Figure 5).

In contrast to previous observations of marine CB, which exhibited a rapid and massive

decline in conductance upon exposure to air (22), our observations of conductance in at

least some freshwater CB filaments hinted at a higher degree of robustness upon oxygen

exposure. For example, an IDA device with 2 CB filaments showed a modest slow increase

in conductance during repeated I-V measurements over 3 hours in the oxygen-free

atmosphere (Figure 5), ultimately reaching $\pm$ 3 μA in response to $\pm$ 0.2 V. Upon removal

from the anaerobic chamber and exposure to air, the conductance immediately declined to

80% of the previous value, and slowly decayed until it ultimately stabilized at 63% of the

oxygen-free value over 2.7 hours. When the IDA was moved back to the anaerobic chamber

for further I-V measurements, the conductance again increased slowly until it stabilized

after 2.4 hours (Figure 5). Interestingly, this behavior was filament-dependent; in another

IDA that started with significantly smaller conductance under N_2 flow conditions (nA

current values for 0.8 V), exposure to ambient air caused immediate dramatic decline

within minutes (Figure S1).

2.3 Conductive atomic force microscopy: point current-voltage spectra. The

abovementioned conductance measurements probe transport longitudinally along multiple

CB sections bridging IDA microelectrodes. We also performed point I-V conductance

measurements through CB filaments between the underlying IDA and conductive AFM

(C-AFM) tips; these measurements simultaneously probe longitudinal transport locally

along specific CB sections and transversely across specific cells to the C-AFM tip above

the filaments (Figure 6).

Tapping mode AFM images were first taken without voltage bias to visualize CB filaments

laid on top of IDA microelectrodes and bridging 3-5 μm sections of the insulating quartz

substrates (Figure 6A-C). No current response between the IDA and C-AFM tip was

detected in response to applied voltage when the tip then probed the bare quartz surface

near CBs. However, when the C-AFM tip probed points on the top surface of CB over

quartz gaps, we detected currents up to tens of pA for voltages (-10-10 V) between the IDA

and tip. These currents reflect a continuous path for electron transport from the underlying

IDAs into cells, longitudinally along the conductive network of CB, and transversely into

the overlying C-AFM tip. Interestingly, when such point I-V measurements were

performed on CB exposed to air, the observed conductivity did not decay immediately, but

slowly decreased over several hours, consistent with the IDA measurements reported above.

2.4 Conductive atomic force microscopy: mapping the conductive network of cable

bacteria. In addition to the point I-V measurements described above, C-AFM was used in

contact scanning mode to image the electron transport path along CB hanging off

microelectrodes under N_2 perfusion conditions (Figure 7). A constant voltage (5 V) was

applied along the freshwater CB filaments between the microelectrode and an ultra-sharp

(< 5 nm tip radius) conductive single crystal diamond C-AFM tip. Following a previous

report on marine CB (23), a force set point was selected to gently scrape off the insulating

outer membrane of the freshwater CB while allowing access to image the conductive

network underneath. Figure 7 displays a current map, stitched together from 3 consecutive

C-AFM images, of two CB filaments lying side-by-side and extending approximately 30

μm off a microelectrode. The imaging revealed parallel conductive paths consistent with

the periplasmic nanofibers as the charge carriers in CB. Indeed, when compared with

topographic imaging (Figure 7), the conductive paths coincided with the CB cell surface

ridge pattern.

2.5 Four-probe transport measurements. To quantify the intrinsic conductivity of

freshwater CB and their periplasmic nanofibers, we measured electronic transport along

Mungi Lake CB deposited on custom fabricated four-probe (4P) microelectrode devices.

In contrast to the two-probe IDA and C-AFM measurements described above, which

include contributions from the CB-electrode contact resistances, the 4P measurements

excluded contact resistance by routing current between two external microelectrodes and

separately measuring the voltage drop across two inner microelectrodes. The resulting I-V

plots (Figure 8) are shown for two separate inner probe separations, with the inverse slopes

yielding whole CB resistances of 47 MΩ and 240 MΩ for the 20 μm and 200 μm lengths,

respectively. By modeling the CB periplasmic nanofibers as current-carrying simple

cylindrical wires forming a simplified circuit of parallel resistors, and taking into account

each nanofiber's estimated diameter (50 nm) and number of fibers per CB (~35), we arrive

at a single nanofiber conductivity on the order of 0.1 S/cm, from both gap sizes.

Discussion

The appearance in many gene sequencing studies and increasing field observations suggest

a cosmopolitan global distribution for cable bacteria (CB) in both marine and freshwater

sediments (11, 45). Here, we identified two sampling sites (Figure 1) in the Los Angeles

basin area, which served as a source for downstream phylogeny, microscopy, scanning

probe, and electronic studies of freshwater CB. While CB have not been isolated into pure

culture (46), the freshwater CB from both Ballona wetlands sediment and Mungi Lake

sediment could be reliably enriched in laboratory incubations to provide a continuous

supply for studies (Figure 1). Based on 16S rRNA gene sequence phylogeny (Figure 3),

CB from the heavily vegetated Ballona sediments are closely related to *Ca.* Electronema

palustris (26), which has been previously observed to associate with the roots of aquatic

plants (47). On the other hand, CB from the non-vegetated Mungi Lake sediments are more

closely related to those from PAH-contaminated sediment collecting water runoffs (38).

The 16S rRNA gene analysis also revealed additional non-CB clones from both Ballona

and Mungi Lake samples, many of which are highly similar to those from stratified and

oligotrophic freshwater ecosystems (34). Although the relationship and functions of these

bacteria are unclear, they likely represent cells firmly attached to CB filaments, as all the

filaments were thoroughly washed before DNA extraction. The communities from both

sites have representative lineages related to organic contaminant degradation, which

corelate to the environmental background of the two sites. In Mungi Lake, for example,

various contaminants (chlordane, DDT, lead, dieldrin, polycyclic aromatic hydrocarbon,

and polychlorinated biphenyls) have been identified previously (31).

Morphologically, our freshwater CB filaments exhibited the expected characteristics of

previously studied CB (8, 20), including the cell surface ridge pattern and underlying

periplasmic nanofiber network (Figures 2-4). The latter could be observed in transmission

electron microscopy (TEM) of freshwater CB both within intact cells and, occasionally, as

the remaining cage scaffold from naturally degraded cells (Figure 2). The typical single

nanofiber width encountered in our freshwater CB was 53 ± 8 nm, which is consistent with

the previously reported nanofiber thickness from marine CB (20) and recent

characterization of a single strain freshwater CB enrichment culture (33 nm) (21). These

results suggest that, while general morphological features are shared amongst diverse CB

strains, the number and size of the ridges (and possibly their internal components) may

depend on strain and environmental condition. However, similarly to previous

measurements of marine CB (8), electrostatic force microscopy (EFM) captured stronger

electric forces when scanning the cell surface above the ridges (Figure 4), again indicating

the unique electric potential and charge distribution from the underlying conductive

periplasmic nanofibers. By scanning the long-range electrostatic interaction between CB

and tip at a fixed retrace height above the sample, we observed the expected voltage

squared dependence of the EFM phase contrast (48), thereby unambiguously assigning

EFM phase contrast to the electric force gradient rather than a topographic contribution.

Conductive atomic force microscopy (C-AFM) with an ultra-sharp diamond tip that

disrupts the insulating outer cell layer (Figure 7) offered an even more direct confirmation

of the conductivity of underlying periplasmic nanofibers, by observing conductive paths

between the microelectrode and scanning tip that correlate with the CB ridge pattern.

Furthermore, C-AFM point I-V spectra (Figure 6) confirmed a direct path for charge

transport originating from the microelectrode below intact CB, along the internal

periplasmic nanofiber, and into the C-AFM tip above the sample, consistent with the

previously observed interconnectivity of the nanofiber network between cells (23). The

latter measurements also illustrate a high degree of variability (Figure 6) in the conduction

signal across different cables; it remains unclear whether this variability reflects

heterogeneities intrinsic to different CB filaments or stems from culturing and/or sample

preparation effects (including possible oxidative damage). Taken collectively, however,

the EFM and C-AFM observations, performed here for the first time on freshwater CB, are

consistent with previous analogous measurements on marine CB (8, 22, 23).

While their two-probe nature includes a contact resistance contribution, we found

interdigitated array (IDA) microelectrodes to offer a convenient assessment of CB

conduction properties (Figure 5) ahead of more sophisticated AFM and four-probe based

techniques (Figures 6-8). The IDA platforms, which are commercially available, have been

previously used to investigate conductivity in bacterial biofilms (5, 6), and result in high

current signals (μA currents in Figure 5) as a result of assessing transport along multiple

CB sections spanning the IDA inter-electrode gaps. Interestingly, these measurements hint

that freshwater CB conductivity may have a somewhat higher degree of tolerance to

oxygen exposure than previously reported for marine CB. Instead of an instantaneous and

dramatic decline in conduction upon exposure to air (22), some of our freshwater CB

samples exhibited a slow (hours) and only partial decline of the conduction upon exposure

to air, which could be partially restored by re-insertion into an anaerobic chamber (Figure

5). While these observations showed a high degree of variability and therefore require a

more detailed investigation, it is interesting to consider the possibility that the oxidative

decline (or other damaging processes), thought to be a consequence of oxidation of Ni

cofactors involved in electron transport (25), may not affect all sections of the cm long CB

filaments simultaneously; under these conditions, IDA measurements, which probe

multiple micrometer-scale electrode-crossing CB regions (e.g. 372 crossings in Figure 5

inset) may offer an advantage over techniques addressing entire CBs or specific CB regions.

It is also interesting to consider an alternative possibility that the conductive properties of

freshwater CB maybe intrinsically more oxygen-resistant than marine CB. In general, salt

content affects the solubility of oxygen, which dissolves to 25% high levels in fresh water

than saltwater. Indeed, previous CB studies verified 20% - 25% higher oxygen

concentration at the surface of freshwater sediment (310 – 350 µM) (11, 28), compared to

marine sediments (200 – 280 µM) (8-10, 13, 14, 49-52). Our results motivate a systematic

investigation to correlate the oxygen availability in CB environments to both their

conductivity and tolerance of electron transport to oxygen exposures.

When taking contact resistances of intact freshwater CB into account with four-probe

measurements (Figure 8), we estimated the single nanofiber conductivity to be in the 10^{-1}

S/cm range. This figure falls within the 10^{-2}-10^{1} S/cm range recently reported for marine

CB (22, 24). While these measurements are performed under *ex situ* conditions, they

motivate the question of whether the measured conductivities are sufficient to support the

in situ activity of CB in sediments. From previous measurements of the CB filaments

density in sediment incubations ($\sim 10^{8}$ - 10^{9} m^{-2}) and the electron current densities linking

electrogenic sulfur oxidation to oxygen reduction (74 - 96 mA m^{-2}) (10, 13, 53, 54), one

arrives at a per cable current around 170 pA, which amounts to per nanofiber current of 5

pA given the average number of nanofibers seen in our cables. Assuming a driving voltage

up to 1 V, representing the redox potential difference between the anodic sulfide oxidation

and cathodic oxygen reduction half-reactions, for a typical nanofiber cross sectional area

and length (e.g. several mm), these conditions requires a minimum nanofiber conductivity

on the order of 0.1 S/cm. Remarkably, this heuristic calculation therefore sets a limit on

conductivity that is consistent with the electronic measurements. It is also interesting to

consider what the observed conductivities ((22-24) and this study) may teach us about the

underlying physical electron transport mechanism in CB. It is common to interpret such

extended transport in molecular systems in one of two limits: hopping between charge

localizing sites such as redox factors (e.g. the hypothesized Ni/S groups of CB), or coherent

band-like transport familiar from periodic systems including metals or semiconductors. In

this context, the thermally activated nature of CB transport, and the estimate of the room

temperature electron mobility at ~ 0.1 cm^2/Vs (24) are consistent with hopping transport,

as the latter figure approaches but is still below the generally accepted minimum mobility

($\sim$1 cm^2/Vs) for band transport (55, 56). Still, future electronic transport and structural

studies (e.g. to identify the precise molecular makeup of the nanofibers) are needed to

resolve the underlying mechanism, particularly if conduction operates in a more complex

regime than band theory or hopping, where electron transport may be delocalized

transiently or permanently over several sites along the nanofibers (57, 58).

Conclusion

In summary, this study presents a detailed electronic characterization of *Candidatus*

Electronema cable bacteria from Southern California freshwater sediments. The cell

envelope characteristics of the freshwater cable bacteria, including the characteristic

periplasmic nanofiber network, are consistent with previous observations of the globally

distributed cable bacteria. Conductive atomic force and electrostatic force microscopies

confirmed the role of the nanofiber network as the conductive conduit of freshwater cable

bacteria, as previously observed in marine cable bacteria. In addition, current-voltage

measurements confirmed the conductivity of whole cable bacterial filaments, while hinting

that the conductivity of the freshwater filaments may be less sensitive to oxygen exposure

than previously observed. The conductivity of the periplasmic nanofibers from four-probe

electronic transport measurements was estimated at 0.1 S/cm, a value high enough to

support the electron current densities linking sulfide oxidation to oxygen reduction through

the electric metabolism of cable bacteria in sediment environments. While electronic

characterization of this long-distance electron transport previously focused on marine

cables, our measurements shed light on the conductivity of freshwater cable bacteria, and

highlight the potential applications of cable bacteria as bioelectronic conduits with both

environmental and future biotechnological relevance.

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Figures and Tables

Figure 1. (A) Sampling sites for cable bacteria from the Ballona Freshwater Marsh and Mungi Lake, California, USA. **(B)** Enrichment in laboratory incubations resulted in a high density of long cable bacteria filaments, some of which could be observed directly in cracked sediments. Scale bar: 35 mm. **(C)** Optical microscopy revealed the characteristic end-to-end multicellular morphology of the cable bacteria filaments, revealing occasional darker cells (black arrows) with slightly larger diameters, consistent with 'empty cage' cells scaffolded by the nanofiber network of cable bacteria (see Figure 2). Scale bar: 5 µm.

Figure 2. (A) Transmission electron microscopy of Ballona Freshwater Marsh cable

bacteria showing the characteristic cell surface ridge pattern. Scale bar: 1 µm. **(B)** The

underlying network of parallel periplasmic nanofibers can be seen clearly in occasional

degraded cells as the electron-dense scaffold of "empty cages". Small intracellular dark

globules (white arrows) are consistent with polyphosphate granules commonly found in

marine cable filaments. Scale bar: 2.5 µm.

Figure 3. Phylogenetic tree based on 16S rRNA gene sequences of the Ballona Freshwater

Marsh and Mungi Lake cable bacteria, other cable bacterial Candidatus taxa, and

representatives of isolated members from the family *Desulfobulbaceae*. Scale bar

represents 10% estimated sequence divergence. Bootstrap numbers (1000 re-samplings) >

70 are listed.

Figure 4. (A) Tapping mode atomic force microscopy of freshwater cable bacteria near the

junction between two cells. Inset is a cross-sectional line scan showing cell surface ridge

pattern. **(B)** Atomic force phase image corresponding to the topographical scan and

showing the ridge pattern in high contrast. **(C)** Electrostatic force microscopy image with

a 6 V tip bias, collected from the retrace scan at 50 nm fixed height above the cable surface.

The contrast in this image stems from the electric force rather than topography. **(D)** The

voltage dependence of the electrostatic interaction (phase shift, $\Delta\phi$, of the ridges with

higher electric force relative to inter-ridge regions) between the conductive tip and the same

cable bacterial filament from (C).

Figure 5. Current-voltage (I-V) measurements of Ballona Freshwater Marsh cable bacteria on gold interdigitated array (IDA) microelectrodes (3 µm gap between the fingers of the two electrode bands). For this measurement, two cable filaments were deposited on the IDA (inset). The I-V measurements (black) show a modest increase in conductance over time (3 hours) during consecutive measurements in an air-free environment (95% N_2 and 5% H_2). Upon air exposure (red) the conductance slowly declined over time (2.7 hours), and conductance recovered partially after re-insertion into an air-free environment (green). Arrow direction denotes measurements over time for a particular condition.

Figure 6. Representative atomic force microscopy amplitude images of Ballona freshwater

cable bacteria filaments spanning the insulating quartz gap between yellow false colored

gold **(A and B)** and indium tin oxide **(C)** microelectrodes. Colored crosses mark the

conductive tip location in each sample where point current-voltage (I-V) spectra were

collected to measure electron transport between the microelectrodes and tip. Scale bars:

2.5 μm. **(D)** I-V responses from the three representative cables. Line color corresponds to

the color of the corresponding cross mark in (A-C).

Figure 7. (A) A spatial montage of three consecutive tapping atomic force microscopy

amplitude images of two parallel Mungi Lake freshwater cable bacteria in contact with,

and extending away from, an electrode (false colored yellow). Scale bar: 3 μm. **(B)**

Conductive atomic force current map images of the same sections shown in (A) with 5 V

bias between the underlying electrode and conductive tip. The observed current signals

largely correlate with the corresponding cell surface ridge pattern in (A).

Figure 8. Four-probe current-voltage measurements of Mungi Lake freshwater cable

bacteria filaments, shown for two different separations between the two inner probes.

Inset: representative image of cable bacteria filaments across a four-probe device. Each

data point is the average of three data points, and the error bars represent the standard

deviation.

Figure S1. An example of sensitivity of conduction in cable bacteria to air exposure.

Consistent current is monitored overnight for a Ballona freshwater sediment cable bacteria

filament on an IDA under N_2 flow conditions, with an 800 mV difference between the two

microelectrode bands. The current immediately declined upon interruption of N_2 flow and

exposure to air.

A

Ballona Freshwater Marsh

Mungi Lake

Los Angeles

20 km

B

C

Uncultured contaminants-degrading Deltaproteobacteria

Ballona_1 (14 clones)
Ballona_2 (3 clones)
Ballona_6 (7 clones)
Candidatus Electronema palustris (KP728463)

***Candidatus* Electronema VI**
(Freshwater CB)

Mungi_24 (3 clones)
Candidatus Electronema nielsenii (KP728462)

***Candidatus* Electronema V**
(Freshwater CB)

Candidatus Electrothrix japonica (KR912349)
Candidatus Electrothrix communis (KR912344)
Candidatus Electrothrix aarhusiensis (KR912338)
Candidatus Electrothrix marina (KR912340)

***Candidatus* Electrothrix**
(Marine CB)

Desulfobulbus propionicus (AY548789)
Desulfobulbus propionicus DSM 2032 (CP002364)
Desulfobulbus elongatus (X95180)
Desulfobulbus alkaliphilus (HM750216)
Desulfobulbus rhabdoformis (U12253)
Desulfobulbus rhabdoformis (AB546247)
Desulfobulbus japonicus (AB110550)
Desulfobulbus japonicus (AB110549)
Desulfobulbus mediterraneus (AF354663)
Desulfobulbus mediterraneus (AJ866934)

Desulfobulbus

Desulfurivibrio alkaliphilus AHT 2 (EF422413)
Desulfurivibrio alkaliphilus AHT 2 (CP001940)
Delta proteobacterium MLMS−1 (AY459365)

Desulfurivibrio

Desulfobulbaceae

Geobacter
Uncultured Deltaproteobacteria
Desulfovibrio desulfuricans subsp. *desulfuricans* (AF192153)
Desulfovibrio marinus (DQ365924)

0.10

A Quartz background / Au

B Current (pA)