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Corresponding Author:	Andrew J. Collins The Forsyth Institute Cambridge, Massachusetts UNITED STATES
Corresponding Author's Institution:	The Forsyth Institute
Corresponding Author E-Mail:	andrewjcollins85@gmail.com
Order of Authors:	Andrew J. Collins
	Dr. Pallavi P. Murugkar
	Dr. Floyd E. Dewhirst
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1 TITLE:

Establishing Stable Binary Cultures of Symbiotic Saccharibacteria from the Oral Cavity

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AUTHORS AND AFFILIATIONS:

5 Andrew J. Collins^{†1,2}, Pallavi P. Murugkar^{†1,2}, Floyd E. Dewhirst^{1,2}

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¹The Forsyth Institute, Cambridge MA, USA

²Harvard School of Dental Medicine, Boston, MA, USA

8 9

10 †These authors contributed equally.

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12 Email addresses of co-authors:

13 Andrew J. Collins (acollins@covaris.com)

14 Pallavi P. Murugkar

(murugkar@stanford.edu)

15

16 Email address of corresponding author:

17 Floyd E. Dewhirst (fdewhirst@forsyth.org)

18 19

KEYWORDS:

Unculturable bacteria, Candidate phyla, Saccharibacteria, oral cavity, binary cultures

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SUMMARY:

We demonstrate a method for isolating difficult-to-grow members of the novel bacterial phylum, Saccharibacteria, by filtering dental plaque and co-culturing with host bacteria.

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ABSTRACT:

Many bacterial species cannot be cultured in the laboratory using standard methods, posing a significant barrier to studying the majority of microbial diversity on earth. Novel approaches are required to culture these uncultured bacteria so that investigators can effectively study their physiology and lifestyle using the powerful tools available in the laboratory. The Candidate Phyla Radiation (CPR) is one of the largest groups of uncultivated bacteria, comprising ~15% of the living diversity on earth. The first isolate of this group was a member of the Saccharibacteria phylum, 'Nanosynbacter lyticus' strain TM7x. TM7x is an unusually small bacterium that lives as a symbiont in direct contact with a bacterial host, Schaalia odontolytica, strain XH001. Taking advantage of the unusually small cell size and its lifestyle as a symbiotic organism, we developed a protocol to rapidly culture Saccharibacteria from dental plaque. This protocol will show how to filter a suspension of dental plaque through a 0.2 µm filter, then concentrate the collected Saccharibacteria cells and infect a culture of host organisms. The resulting coculture can be passaged as any normal bacterial culture and infection is confirmed by PCR. The resulting binary culture can be maintained in the laboratory and used for future experiments. While contamination is a possibility, the binary culture can be purified by either further filtering and reinfection of host, or by plating the binary culture and screening for infected colonies. We hope this protocol can be expanded to other sample types and environments, leading to the cultivation of many more species in the CPR.

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INTRODUCTION:

Culturing novel species of bacteria and bringing them into the laboratory allows for powerful experiments to better understand their physiology and broader interactions within their microbial community. While there are culture-free methods of interrogating these questions (e.g., "meta-omics"), the complex interactions of diverse microbial populations make it difficult to tease apart single variables and reach meaningful conclusions. While culturing bacteria has many benefits, there are many potential barriers to isolating a bacterium and growing it in pure culture. Potential specific growth requirements include pH, oxygen tension, vitamins, growth factors, signaling molecules or even direct cell contact to elicit growth¹. However, it is believed that specific auxotrophies are the primary deterrent to culturing new species of bacteria. Standard media formulations lack many nutrients required by uncultivated bacteria, such as specific vitamins or carbons sources. These missing molecules can be key to the physiology of the uncultured bacteria and are usually provided by either another organism in the microbial community or a host organism. For example, complex carbohydrates such as mucins can be provided by animal hosts. Adding these to media has allowed cultivation of several bacteria from animal guts, including Akkermansia muciniphila and Mucinivorans hirudinis^{2–4}. Many pathogenic bacteria have evolved the ability to use iron bound to hemin in animal cells, including the oral pathogen Porphyromonas gingivalis⁵. In the laboratory, the growth of Porphyromonas and other organisms, can be stimulated by the addition of hemin⁶.

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Recently, many breakthroughs in culturing novel isolates of bacteria have come through coculturing, using a "feeder" organism to provide specific factors to uncultured bacteria necessary for their growth. An elegant study by Vartoukian and colleagues showed that siderophores, iron binding molecules produced by bacteria, stimulated the growth of several novel oral isolates. Pyoverdines, a type of siderophore produced by pseudomonad species, were shown to significantly facilitate the growth of a novel *Prevotella* species⁷. In the same study, the first oral isolate for the phylum *Chloroflexi* was cultivated, also using *F. nucleatum* as a helper for providing some as yet unknown compound⁷. More recently, a bacterium from the genus *Ruminococcaceae* was isolated using *Bacteroides fragilis* as helper organism⁸. It was later shown that gamma aminobutyric acid (GABA), an inhibitory neurotransmitter, was required for growth on laboratory media. Using feeder organisms has proved to be a key strategy to mimicking specific microenvironments where uncultured bacteria grow, being more efficient than continuously reformulating growth media with different additives in varying concentrations.

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One of the largest groups of uncultured bacteria are in the "Candidate Phyla Radiation" (CPR), a monophyletic group of several candidate bacterial phyla^{9,10}. As of this writing, only members of the Saccharibacteria phylum within the CPR have been successfully cultured in the laboratory. The first isolate, 'Nanosynbacter lyticus' strain TM7x, was isolated using the antibiotic streptomycin, which had been predicted to enrich for the uncultured TM7^{11,12}. A key discovery of this work was that the new isolate grew as a parasite growing in direct contact with a bacterial host, *Schaalia odontolytica*, and microscopy showed these parasites were ultrasmall bacteria.

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Using these clues, we devised a method to quickly establish binary cocultures of Saccharibacteria

with their partners by filtering dental plaque and other oral samples through a 0.2 µm filter, collecting cells in the filtrate by centrifugation and using them to infect cultures of candidate host bacteria. This method has the advantage of avoiding enrichment cultures, which can be overwhelmed with fast-growing organisms. It also avoids the use of antibiotics, which could stop the growth of either the targeted Saccharibacteria species or their hosts. Using the method demonstrated here, we have successfully cultured 32 isolates from the Saccharibacteria phylum.

PROTOCOL:

When developing this protocol, IRB approval was sought and approved (#14-10) and informed consent was obtained from all subjects.

1. Preparation

102 1.1. When working with human subjects, obtain necessary IRB approval and informed consent.

1.2. Start cultures of host bacteria with enough time to grow to early stationary phase. For example, inoculate 2-5 mL of tryptic soy broth (TSBY) with 0.1% yeast extract added with *Arachnia propionica* and incubate at 37 °C for 24 h.

1.3. Assemble filter holders with track-etched 0.2 µm filter membrane. Wrap the assembly in foil and sterilize by autoclaving.

112 1.4. Sterilize centrifuge tubes and cap assemblies.

2. Obtain sample of oral bacteria

NOTE: While many oral samples contain Saccharibacteria (e.g., saliva, swabs of tonsils, tongue scrapings) dental plaque routinely is the most successful.

2.1. Take a plaque scraping using a sterile paper point, Gracey curette, or, if self-sampling, use a sterile toothpick or pipette tip. Transfer plaque to a suitable buffer, such as maximal recovery diluent (MRD, 0.85% NaCl, 0.1% peptone) or PBS. If not processed immediately, keep the sample on ice until ready to proceed.

124 2.2. Vigorously resuspend the dental plaque in MRD buffer using a combination of vortexing and
 125 pipetting with a small pipet tip.

2.3. Add resuspended plaque sample to an additional 9 mL of MRD buffer.

3. Prepare filtrate from oral sample

3.1. Using aseptic technique, unwrap a sterile filter assembly. Untwist a quarter turn and retighten the filter holder to be sure threads are properly engaged and the apparatus is closed

- properly. Using a syringe, wash the membrane by passing 10 mL of MRD buffer through the apparatus. Improper assembly is revealed at this step by fluid leaking out of the filter holder. If leakage occurs, obtain another sterile filter assembly and repeat the wash step.
- 3.2. Apply the sample to the washed filter. Remove the plunger from a syringe and attach it to
 the filter apparatus. Pour the dispersed dental sample, now in 10 mL of MRD buffer, into a syringe
 and load it onto the filter.
- 3.3. Place a sterile centrifuge tube beneath the filter apparatus to catch the filtrate, then apply
 light pressure to the plunger to push the sample through the filter.
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- 144 3.4. Repeat the procedure with another 10 mL of MRD buffer to wash the membrane, collecting the flow-through in the same tube as the filtered sample. Aseptically cap the tube.

4. Concentrate Saccharibacteria cells by centrifugation

- 4.1. Make an orientation mark on the tube and cap and place the tube in a high-speed centrifuge with the mark on the upper side. The pellet formed from centrifugation is usually invisible. The marking will help determine where the pellet of Saccharibacteria cells is located when centrifugation is done and the tube removed from the centrifuge.
- 4.2. Centrifuge the samples at $60,000 \times g$ for 1 h at 4 °C.

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- NOTE: This force and time is sufficient to pellet all Saccharibacteria cells. However, Saccharibacteria cells can be at least partially pelleted by centrifuging for as little as 20 min at $20,000 \times q$.
- 4.3. Carefully remove the tubes from centrifuge. Pour out the liquid from the tube, keeping the
 pellet on the upper side of the tube.
- 163 4.4. Resuspend the usually invisible pellet in 1-2 mL of MRD buffer by vigorous vortexing.

5. Infect host cultures with Saccharibacteria-enriched filtrate

- 5.1. Prepare culture tubes by aliquoting 2 mL of appropriate growth media (e.g., TSBY, BHI, etc.)
 into tubes.
- 5.2. Add 200 μL of overnight culture of host organisms to each tube. Add 100-200 μL of
 resuspended, filtered sample to each tube.
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- 5.3. Incubate combined samples as appropriate for the host organism (e.g., 37 °C, in an aerobic atmosphere for *A. propionica*).
- 176 5.4. Passage the cells every two to three days by transferring 200 μL of binary culture to 2 mL of

fresh growth medium in a new tube. If it appears that the passaged cultures show no growth (i.e., the turbidity/optical density of the culture doesn't increase after passage into fresh media) the Saccharibacteria could be overwhelming or killing all the host organism. To remedy this, add 200 µL of uninfected host culture when passaging the cells. Repeat for at least 5 passages.

6. Confirm infection by PCR

6.1. After 5 passages, confirm infection with PCR. Five passages will ensure than any non-growing Saccharibacteria cells have been depleted beyond the limit of detection using 25 cycles of PCR.

6.2. Prepare a PCR mastermix. A suggested recipe using is as follows, for each tube needed, prepare 12.5 μ L 2x PCR buffer, 0.75 μ L 10 μ M Forward primer (580F¹² - AYT GGG CGT AAA GAG TTG C), 0.75 μ L 10 μ M Reverse primer (1177R¹³ – GAC CTG ACA TCA TCC CCT CCT TCC), 1 μ L 25 mM MgCl₂ and 9 μ L water. Mix by vortexing.

6.3. Aliquot 24 μ L of mastermix into a 0.2 mL PCR tube. Add 1 μ L of Saccharibacteria-infected culture to the PCR reaction. Place the tube in a thermocycler and perform the following protocol: Initial denaturation 95 °C for 5 min, 25 cycles of 95 °C for 30 s, 60 °C for 30 s, 72 °C for 30 s, final elongation at 72 °C for 2 min, then final hold at 4 °C.

6.4. Load and run the PCR products in a 1% agarose gel. A band of ~600 bases will indicate the presence of Saccharibacteria.

7. Check for purity and remove contaminating organisms

7.1. Confirm purity of positive cultures by plating the coculture of Saccharibacteria on nutrient agar sufficient for the growth of the host organism. Perform a 10-fold serial dilution of the culture in sterile buffer (e.g., MRD or PBS) and spread 100 μ L on an agar plate. Perform dilutions such that there will be approximately 20-200 colonies growing on the agar plate.

7.2. Incubate the culture in appropriate growth conditions and observe for contaminating organisms.

NOTE: Axenic Saccharibacteria colonies have never been observed, so only the host organism is observed growing on these plates. More than one colony type usually indicates contamination. Should contamination occur, the culture should be re-filtered to remove the unwanted contaminants and re-inoculated onto fresh host.

7.3. Occasionally a double infection occurs, where two Saccharibacteria species infect the same host. To separate the Saccharibacteria species, resuspend individual colonies in 20 μ L sterile PBS and use 1 μ L of suspension in a PCR reaction to detect colonies with Saccharibacteria infections.

7.4. Transfer the suspensions that gave a positive result to growth medium to start a binary culture. The success rate will depend on the titer of each Saccharibacteria species in the culture.

221 It may be necessary to screen more than 50 colonies to find an infected one for propagation.

8. Storage of cultures

8.1. Grow a binary culture of sufficient volume (10-50 mL) overnight.

8.2. Pellet cells by centrifugation $(4,000 \times q \text{ for } 10 \text{ min})$.

NOTE: High-speed centrifugation is not necessary here as Saccharibacteria cells will be attached to the larger, heavier cells of the host and will pellet with them.

8.3. Resuspend cells in cryoprotectant. Growth media supplemented with either 5% DMSO or 20% glycerol is usually sufficient. Ensure that the host organism is compatible with cryoprotectant media (e.g., Growth of *A. propionica* is inhibited by glycerol and frozen stocks may not revive). Test the viability of an uninfected host culture with cryoprotectant to ensure stocks can survive freezing.

8.4. Resuspend cell pellet in the same volume of culture (10-50 mL). Aliquot 0.5 mL into labeled cryovials. Freeze cell cultures at -80 °C.

REPRESENTATIVE RESULTS:

PCR for detection of Saccharibacteria may appear negative (i.e., no product seen) in initial infection cultures due to a low number of Saccharibacteria symbionts. However, after a few passages a strong PCR product should appear showing that a stable infection has occurred (**Figure 1A**). Conversely, some infections will initially appear positive by PCR, but diminish to undetectable after 1-4 passages (not shown). This indicates that a large amount of Saccharibacteria cells were present in the initial filtrate but were diluted out by passage and none were able to enter into a stable symbiosis with the host culture. Testing several host species with the same Saccharibacteria filtrate from the oral cavity will usually have a low success rate (**Figure 1B**) as the symbiont-host interaction is very specific. A researcher may also test the same host with filtrates from several different subjects. If a good host organism (such as *Arachnia propionica* or *Schaalia odontolytica*) is used, a success rate of 50% can be expected.

Testing by PCR is crucial. Experienced researchers have attempted to confirm infection by microscopy, only to report false positives. The Saccharibacteria cells are small and difficult to distinguish between vesicles or irregular cell envelope bulges. A PCR signal stable through several passages is key to confirm a successful infection.

As the infected cultures grow, normal turbid growth should appear. As the symbiosis establishes itself, infected cultures will appear less turbid than cultures of uninfected host organisms (**Figure 2A**). In some cases, infected cultures may appear to stop growing completely and not become turbid at all. This may be due to an infection where the Saccharibacteria cells are overwhelming the host organism. Adding "fresh" (uninfected) host to these cultures should provide a sufficient population of host to support the continued growth of the Saccharibacteria parasites.

Overgrowth, or excessively turbid cultures, may indicate contamination by either a laboratory contaminant or another small oral bacterium that was able to pass through the 0.2 μ m filter. Campylobacter and Capnocytophaga spp. are typical oral contaminants of these experiments. This can be confirmed by plating the culture and looking for colonies that are atypical of the host organism followed by 16S rRNA sequencing. If contamination is seen, filtering these cultures through a 0.2 μ m filter is usually sufficient to remove the contaminants. Saccharibacteria cells in the filtrate can be concentrated by centrifugation and used to re-infect a pure host culture.

Another way to purify a contaminated culture is by picking infected colonies from plating an infected culture. Colonies of infected hosts can sometimes be identified by irregular colony shape compared to uninfected colonies (Figure 2B-D). These irregular colonies are dependent on the titer of Saccharibacteria in the binary culture and a smaller proportion of colonies will appear irregular if the titer is low. This can make it easy to identify infected colonies, which can be picked and used to start a pure binary culture. If no irregular colonies are seen on plating from a Saccharibacteria infected culture, it is possible that the symbiont is at a low titer or does not cause irregularly shaped colonies. In such a case, PCR screening colonies with a normal appearance can show infection by Saccharibacteria, but at a low rate (2-10% of all colonies).

FIGURE AND TABLE LEGENDS:

Figure 1: Typical PCR results of Saccharibacteria infections of host cultures. (**A**) A PCR product indicating presence of Saccharibacteria may not appear with the initial infection but can appear and become stronger in subsequent passages as the coculture establishes itself. (**B**) Most hosts infected with Saccharibacteria-enriched filtrate will not support their growth due to the specificity of the symbiosis. In this example, only *A. propionica* was successfully infected.

Figure 2: Growth characteristics of Saccharibacteria infected cultures. (A) Growth curves of infected and uninfected cultures of *A. propionica* showing infected culture will not grow to the same density as the uninfected control and appear less turbid. (B-D) Plating of cocultures can produce irregular colonies, caused by Saccharibateria infections. Irregular colonies will decrease in proportion relative to the titer of Saccharibacteria in the coculture. (B) Host with a high level of Saccharibacteria. (C) 10-fold diluted Saccharibacteria (D) Uninfected host culture. White arrows indicate examples of irregular colonies. Scale bar= 1 cm.

DISCUSSION:

Our method of filtering plaque and applying it to pure cultures of host organisms is largely based on earlier observations on the first cultured Saccharibacteria, 'Nanosynbacter lyticus' strain TM7x^{11,14,15}. Given the small cell size, we deduced that they could be separated from dental plaque using a filter and concentrated with centrifugation. Second, as these organisms live as parasites, providing these cells pure cultures of hosts would allow them to enter into a symbiosis and grow as binary cultures.

One advantage of this method is that it does not require an enrichment culture or selective pressure. 'Nanosynbacter lyticus' strain TM7x was cultured from an enrichment culture using

streptomycin as a selective agent, which sequencing had suggested would be effective at enriching for Saccharibacteria. Fortuitously, the host for 'Nanosynbacter lyticus' Schaalia odontolytica, is known to be resistant to streptomycin¹⁶. Using antibiotics as a selective agent could also prevent the host organism from growing, which in turn would preclude the growth of Saccharibacteria.

A larger issue of using enrichment cultures is that fast-growing organisms will quickly displace organisms of interest. In the oral cavity, for example, *Streptococcus* species can grow quickly and, if sugar is present in the growth medium, produce enough acid to acidify the medium, further selecting against organisms of interest. By avoiding an enrichment culture and selective antibiotics, our method provides a general approach that could be applied to a wider range of Saccharibacteria and potential hosts without the complications of these other methods.

There are some obstacles to the method presented here. First, this method assumes that Saccharibacteria live in a binary culture. We have not tested combinations of trinary or ternary cultures to gauge their effectiveness, but there are likely Saccharibacteria that require growth factors that a single host organism cannot supply. Testing the vast combinations of oral bacteria that could support the growth of Saccharibacteria would be a daunting task. Second, the method assumes all Saccharibacteria are small enough to pass through a 0.2 µm filter. It could be that other Saccharibacteria are larger than believed and the filter is selecting against these organisms. A filter with a larger pore size could be used, but this runs the risk of allowing more unwanted oral bacteria into the infected co-culture. Lastly, it is very difficult to find host species outside of those that have already been published. Thus far, the only successful hosts are species from the genera *Actinomyces, Schaalia, Arachnia* and *Cellulosimicrobium,* all members of the phylum Actinobacteria^{15, 17, 18}. However, these hosts only support the growth of specific Saccharibacteria. To culture more species of Saccharibacteria, many more hosts must be explored.

We hope the method presented here will aid future research of the Saccharibacteria and other CPR organisms. Metagenomic sequencing suggests that these organisms also have small genomes and are suspected to be symbionts or rely on the local microbial community to supply metabolites and other factors critical to their survival¹⁹. A similar filtering strategy could be used to isolate these organisms, provided they are small enough and that their host organisms(s) can be cultured. The methods described here are a first step to bringing the powerful tools of laboratory culture to this large and diverse group of bacteria.

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DISCLOSURES:

354 The authors have nothing to disclose.

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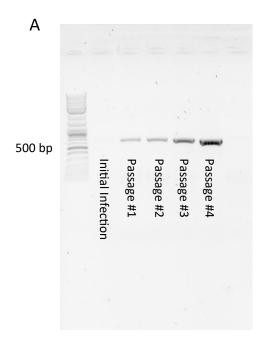
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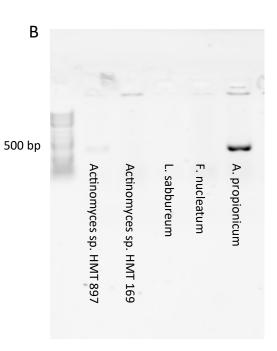
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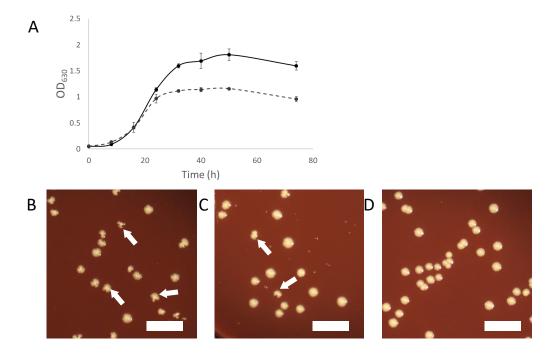
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Name of Material/ Equipment	Company	Catalog Number	Comments/Description
Agarose	Fisher Scientific	BP160-100	
Alphaimager	Cell Biosciences	FluorChem HD2	Or equivalent UV gel imaging system
Aluminum foil	Fisher Scientific	01-213-101	
Brain Heart Infusion Broth (dehydrated			
powder)	Becton-Dickinson	211059	Or other growth media suitable for ta
Centrifuge Rotor 70-Ti	Beckman Coulter	337922	2
Cryovials	Fisher Scientific	12-567-500	
DMSO	Fisher Scientific	BP231-100	
Electrophoresis Power Supply	Bio-Rad	1645052	2
Electrophoresis Rig	Bio-Rad	1704467	7
Filter Forceps	Millipore Sigma	XX6200006P	Not essential, helps ensure filters are
Glycerol	Fisher Scientific	G33-500	
GoTaq Green Mastermix	Promega	M7122	
Mastercycler Pro Thermocycler	Eppendorf	950040025	Or equivalent thermocycler for PCR
MgCl2 solution 25mM	Promega	A3513	
Molecular Biology grade water	Fisher Scientific	BP2819100	
O2 Control InVitro Glove Box	Coy Laoratories	031615	If needed for microaerobic organisms
Optima L-100 XP High Speed Centrifuge	Beckman Coulter	8043-30-1124	
P-10 micro pipette	Gilson	F144802	
P-1000 micro pipette	Gilson	F123601G	
P-2 micro pipette	Gilson	F144801	
P-20 micro pipette	Gilson	F123600	
P-200 micro pipette	Gilson	F123602G	
PBS	Fisher Scientific	BP399500	
PCR tubes 0.2 mL	Fisher Scientific	14-230-205	
Peptone	Fisher Scientific	BP1420-500	
Pipette tips - 10 μL	Fisher Scientific	02-717-157	
Pipette tips - 1000 μL	Fisher Scientific	02-717-166	
Pipette tips - 20 μL	Fisher Scientific	02-717-161	
Pipette tips - 200 μL	Fisher Scientific	02-717-165	
Polycarbonate filters - 47mm, 0.2 μm pore size	e Millipore	GTTP04700	

Screw-cap conical centrifuge tubes 15 mL	Falcon	352096 Or other tube suitable for bacterial cu
Sodium chloride	Fisher Scientific	BP358-1
Swin-Lok Filter - 47mm	Whatman	4200400
SYBR Safe DNA Gel stain	ThermoFisher Scientific	S33102
Syringes - 20 mL	Fisher Scientific	14-955-460
TAE Buffer (50x) concentrate	Fisher Scientific	P1332500
Thickwall Polycarbonate 25 x 89 mm (26.3mL		
capacity) centrifuge tubes with caps	Beckman Coulter Northeast Laboratory	355618
Tryptic Soy Blood Agar Plates	Services	P1100 Or other agar plate sufficient for grow
Tryptic Soy Broth (dehydrated powder)	Becton-Dickinson	211825 Or other growth media suitable for ta
Vinyl Anaerobic Chamber	Coy Laboratories	032714 If needed for anaerobic organisms
Vortex mixer	Scientific Industries	SI-0236
Yeast Extract	Fisher Scientific	BP1422-500

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February 25, 2021

Dear Sir:

We are pleased to see our manuscript was received so well by our peerreviewers. Attached is the revised document and figures that should correct the issues brought to our attention. Our responses to specific issues are in red below.

- 1. Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammar issues. E.g., Line 30: check the term CPR, etc. We have gone through and made some changes to grammar and added some clarifying language. All initialisms should be used correctly.
- 2. Use appropriate symbols for units. E.g., use " μ m" instead of "micron" (line 36, 86, 285 etc.), " μ L" instead of "uL" (line 203, Table of Materials). Add a single space between the quantity and its unit. E.g. "37 oC" instead of "37oC" (line 103, 156, 181etc.), "-80 oC" instead of "-80C" (line 255), "3 s" instead of "3s" (line 205), "500 bp" instead of "500bp" (figure 1 labels), "1 cm" instead of "1cm" (figure 2 legend : scale bar), and in the Table of Materials.

Changes have been made accordingly.

- 3. Please adjust the numbering of the Protocol to follow the JoVE Instructions for Authors. For example, 1 should be followed by 1.1 and then 1.1.1 and 1.1.2 if necessary. Do not start with step "0" or substep "0" (e.g. 5.0). Changes have been made accordingly.
- 4. Please include an ethics statement before the numbered protocol steps, indicating that the protocol follows the guidelines of your institution's human research ethics committee.

Language describing institutional review and using human subjects has been added.

- 5. Lines 124, 128,135,141, 168 etc.: Specify which buffer is used. We have clarified which buffers used at all points.
- 6. Line 186-178: How is this checked? Language clarifying that researchers should look for turbidity over time was added.
- 7. Line 250-251: How is this checked? Language suggesting that researchers verify the host organism can be revived from cryoprotectant before making frozen stocks has been added.
- 8. JoVE cannot publish manuscripts containing commercial language. This includes trademark symbols (™), registered symbols (®), and company names before an instrument or reagent. Please remove all commercial language from your manuscript and use generic terms instead. All commercial products should be sufficiently referenced in the Table of Materials. E.g. Promega GoTag

Commercial language has been removed from the manuscript.

9. Please highlight up to 3 pages of the Protocol (including headings and spacing) that identifies the essential steps of the protocol for the video, i.e., the steps that should be visualized to tell the most cohesive story of the Protocol. Remember that non-highlighted Protocol steps will remain in the manuscript, and therefore will still be available to the reader.

A section has been highlighted.

10. Please ensure that the references appear as the following: [Lastname, F.I., LastName, F.I., LastName, F.I. Article Title. Source. Volume (Issue), FirstPage – LastPage (YEAR).] For more than 6 authors, list only the first author then et al. Please do not abbreviate the journal names (lines 402, 406).

References should be in the right format and no journals are abbreviated.

11. Please sort the Materials Table alphabetically by the name of the material. Materials are now alphabetized.

We hope this satisfies the issues raised with our manuscript's first draft and look forward to moving forward with the publication process.

Sincerely,

Andrew Collins, PhD