

Conjugation Protocol Optimised for *Roseburia inulinivorans* and *Eubacterium rectale*

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[Abstract] *Roseburia* and *Eubacterium* species of the human gut microbiota play an important role in the maintenance of human health, partly by producing butyrate, the main energy source of our colonic epithelial cells. However, our knowledge of the biochemistry and physiology of these bacteria has been limited by a lack of genetic manipulation techniques. Conjugative transposons previously introduced into *Roseburia* species could not be easily modified, greatly limiting their applicability as genetic modification platforms. Modular plasmid shuttle vectors have previously been developed for *Clostridium* species, which share a taxonomic order with *Roseburia* and *Eubacterium*, raising the possibility that these vectors could be used in these organisms. Here, we describe an optimized conjugation protocol enabling the transfer of autonomously replicating plasmids from an *E. coli* donor strain into *Roseburia inulinivorans* and *Eubacterium rectale*. The modular nature of the plasmids and their ability to be maintained in the recipient bacterium by autonomous replication makes them ideal for investigating heterologous gene expression, and as a platform for other genetic tools including antisense RNA silencing or mobile group II interon gene disruption strategies.

Keywords: Conjugation, Gut microbiota, Lachnospiraceae, Gene transfer, Shuttle vector

[Background] *Roseburia* and *Eubacterium* species are among the most abundant bacteria in the human gut microbiota (Zhernakova *et al.*, 2016), impacting human health by utilising dietary and host derived polysaccharides (Scott *et al.*, 2006 and 2011; Cockburn *et al.*, 2015; Sheridan *et al.*, 2016) and producing the health promoting metabolite butyrate as a fermentation end product (Duncan *et al.*, 2002 and 2006). Additionally, these species are capable of modulating host immunity via flagella (Neville *et al.*, 2013). The lack of genetic modification techniques for these organisms has prevented a more complete understanding of the complex interactions between these bacteria and their human host.

Previously, conjugative transposons were successfully transferred into *Roseburia inulinivorans* from *Eubacterium cellulosolvens* and *Clostridium cf. saccharolyticum* (Scott *et al.*, 2008). These large, novel mobile genetic elements could not be easily modified and thus were a suboptimal platform for detailed genetic modification. This work did however illustrate that conjugative mating was possible between Lachnospiraceae bacteria including *Roseburia* species. The development of easily modified conjugative plasmids for clostridial species (Purdy *et al.*, 2002; Heap *et al.*, 2009) raised the possibility that these techniques could be adapted for *Roseburia* and *Eubacterium* species.

The detailed protocol presented here is based on procedures established in Sheridan *et al.* (2019). In this work, the different conjugative plasmids developed for use in *Clostridium* species (Heap *et al.*, 2009) were tested for transferability into the *Roseburia* and *Eubacterium rectale* species. Plasmid pMTL83151

was successfully transferred into two strains of *E. rectale*, while pMTL83151 and pMTL82151 were transferred into *Roseburia inulinivorans* A2-194. Transfer frequencies of 10^{-6} - 10^{-8} per potential recipient were obtained. These frequencies are similar to those observed when suicide vectors were introduced into other Gram-positive bacteria (Williams *et al.*, 1990; Aquino de Muro and Priest, 2000). The ability to add exogenous DNA to bacterial species opens up opportunities for genetically manipulation, including knockout mutagenesis. Alternatively, these plasmids could be modified as expression vectors for mobile group II interon gene disruption strategies, as has been demonstrated in several clostridial species (Heap *et al.*, 2007). Additionally, Plasmid pMTL83151 was shown to be a suitable vector for heterologous gene expression (Sheridan *et al.*, 2019), producing an enzymatically active *Streptococcus* glycoside hydrolase in both species and thus proving the utility of this technique in studying researcher-selected functional gains in these important bacteria. The protocol below is a stepwise guide to introducing foreign DNA to these bacteria.

Materials and Reagents

1. Pipette tips
2. Cuvettes (Bio-Rad, catalog number: 1652083)
3. Petri dishes (Greinerbio-one, catalog number: 633180)
4. Glass Pasteur pipettes (Fisher Scientific, catalog number: FB50261)
5. Nylon membranes (Roche, catalog number: 11417240001)
6. X-ray film (Fujifilm) (Fisher Scientific, catalog number: 12735325)
7. 50 ml conical centrifuge tubes (Corning, catalog number: 10038980)
8. Strains (Table 1)
9. Plasmids (Table 1)
Note: Modular plasmids can be obtained from CHAINbiotools (<http://clostron.com/pMTL80000.php>).
10. Primers (Table 2)
11. PBS tablets (Sigma, catalog number: P4417)
12. Chloramphenicol (Sigma-Aldrich, catalog number: C0378) stock solution 10 µg/ml, stored at -20 °C
13. HindIII restriction endonuclease (NEB, catalog number: R0104S)
14. Wizard genomic DNA Purification kit (Promega, catalog number: A1120)
15. DIG High Prime DNA Labelling and Detection Starter Kit II (Roche, catalog number: 11585614910)
16. PCR reagents (Taq Polymerase kit Biotline, catalog number: BIO-21040 and dNTP's Promega, catalog number: U1240)
17. Dipotassium phosphate, K_2HPO_4 (Fisher Scientific, catalog number: P/5240/53)
18. Potassium dihydrogen phosphate, KH_2PO_4 (Fisher Scientific, catalog number: P/4800/60)
19. Ammonium sulfate, $(NH_4)_2SO_4$ (Fisher Scientific, catalog number: A/6480/53)

20. Sodium chloride, NaCl (Fisher Scientific, catalog number: S/3160/53)
21. Magnesium sulfate, MgSO₄ (Sigma-Aldrich, catalog number: M7506)
22. Calcium chloride, CaCl₂ (Sigma-Aldrich, catalog number: C1016)
23. Acetic acid (Fisher Scientific, catalog number: A/10400/PB17)
24. Propionic acid (Sigma-Aldrich, catalog number: P1386)
25. n-Valeric acid (Sigma-Aldrich, catalog number: V9759)
26. Iso-Valeric acid (Sigma-Aldrich, catalog number: I7128)
27. Iso-Butyric acid (Sigma-Aldrich, catalog number: I1754)
28. Biotin (Sigma-Aldrich, catalog number: B4501)
29. Cobalamin (Sigma-Aldrich, catalog number: V2876)
30. p-Aminobenzoic acid (Sigma-Aldrich, catalog number: A9878)
31. Folic acid (Sigma-Aldrich, catalog number: F7876)
32. Pyridoxamine (Sigma-Aldrich, catalog number: P9755)
33. Potassium hydroxide, KOH (Sigma-Aldrich, catalog number: P5958)
34. Ethanol 95% (Fisher Scientific, catalog number: E/0650DF/17)
35. Haemin (Sigma-Aldrich, catalog number: H5533)
36. Bacto tryptone (BD Diagnostics Systems, catalog number: 211705)
37. Yeast Extract (BD Diagnostics Systems, catalog number: 212750)
38. Potassium chloride, KCl (Sigma-Aldrich, catalog number: P3911)
39. Magnesium chloride hexahydrate, MgCl₂·6H₂O (Sigma-Aldrich, catalog number: M9272)
40. Magnesium sulfate heptahydrate, MgSO₄·7H₂O (Sigma-Aldrich, catalog number: 230391)
41. Bacto casitone (BD Diagnostics Systems, catalog number: 225930)
42. Sodium bicarbonate, NaHCO₃ (Sigma-Aldrich, catalog number: S576)
43. Glucose (Fisher Scientific, catalog number: G/0500/53)
44. Soluble starch (Sigma-Aldrich, catalog number: S2004)
45. Cellobiose (Sigma-Aldrich, catalog number: C7252)
46. Resazurine (Sigma-Aldrich, catalog number: R2127)
47. L-cysteine (Sigma-Aldrich, catalog number: C1276)
48. Thiamin (Sigma-Aldrich, catalog number: T1270)
49. Riboflavin (Sigma-Aldrich, catalog number: R9504)
50. Gas mix 10% Carbon Dioxide, 10% Hydrogen balance Nitrogen (Anaerobic) Cylinder (BOC, catalog number: 290564-L)
51. Agar (Oxoid, catalog number: LP0011)
52. Resazurin solution (see Recipes)
53. Anaerobic phosphate buffered saline (PBS) (see Recipes)
54. Mineral solution 1 (see Recipes)
55. Mineral solution 2 (see Recipes)
56. Short chain fatty acid solution (see Recipes)
57. Vitamin solution 1 (see Recipes)

58. Vitamin solution 2 (see Recipes)
59. Haemin solution (see Recipes)
60. SOC (see Recipes)
61. LB (see Recipes)
62. LA (see Recipes)
63. YCFAGSC and AMM (see Recipes)

Equipment

1. Pipettes Gilson P1000 (Gilson, catalog number: F123602)
2. Pipettes Gilson P200 (Gilson, catalog number: F123601)
3. Pipettes Gilson P20 (Gilson, catalog number: F123600)
4. Pipettes Gilson P2 (Gilson, catalog number: F144801)
5. Concept Plus Anaerobic Workstation, Ruskinn Technology
6. Hungate tubes (Sciquip, catalog number: 2047-00125)
7. Hungate lids butyl rubber septa (Sciquip, catalog numbers: 2047-11600 and 2047-16000)
8. Wheaton bottles (Merck, catalog number: 33110-U)
9. CO₂ hooks (made in-house) and CO₂ piped gas supply
10. Shaking Incubator for *E. coli* growth (Sanyo Orbital incubator)
11. Static Incubator for anaerobic bacteria (Sciquip incu-160S)
12. Gene pulser (Bio-Rad, model: 1652078)
13. Platform rocker (Stuart, model: STR6)
14. UV-linker, Bio-Rad GS Gene linker UV chamber (UVP CL-1000 Ultraviolet Crosslinker)
15. Hybridizer (UVP laboratory products HB 1000 hybridizer)
16. Blot transfer pump (Hybaid Vacu-aid blot processing pump, DA7C.VAC/T)
17. Centrifuge (Jouan MR1822)
18. Transilluminator (UVtec BXT-20.L)

Procedure

- A. Electroporation of plasmids into *E. coli* CA434
 1. Pre-chill 1 mm electroporation cuvette in refrigerator overnight.
 2. Dilute 1 µl of plasmid solution (10 µg/ml) in 4 µl of dH₂O (final concentration 2 µg/ml).
 3. Thaw 55 µl electrocompetent *E. coli* CA434 (prepared in house following standard protocols, <https://www.protocols.io/view/Making-your-own-electrocompetent-cells-imsv6m>) on ice and add the plasmid solution, mixing gently.
 4. Transfer the mixture to the cuvette and place in Gene Pulser.
 5. Electroporate at 1.8 kV, 200 Ohms, 25 µF.
 6. Transfer into 1 ml SOC medium, pre-warmed to 37 °C.

7. Incubate for 1 h at 37 °C, 200 RPM.
8. Dilute in sterile dH₂O in ten-fold dilutions (10¹, 10² and 10³).
9. Spread 50 µl of these dilutions onto LA plates (see Recipes) supplemented with chloramphenicol 10 µg/ml (Cm10) and incubate at 37 °C for 24 h or until colonies appear.
10. Pick individual colonies into LB Cm10 and incubate at 37 °C, 200 RPM for 24 h.

B. Anaerobic culturing of recipient bacterium

1. Prepare anaerobic media YCFAGSC and AMM in broth as 7.5 ml aliquots in Hungate tubes, sealed with butyl rubber septa and 2% agar solutions divided into 100 ml aliquots in Wheaton bottles, with all dispensing and inoculating carried out under CO₂ using the Hungate technique as described by Bryant, 1972 (Figure 1).



Figure 1. Culturing of strictly anaerobic bacteria using the Hungate methodology (growth in tubes) and anaerobic workstation (growth on Petri plates)

2. Incubate anaerobic YCFAGSC liquid cultures at 37 °C, without agitation in Hungate tubes.
3. Pour anaerobic agar media into Petri plate (20 ml per plate) within anaerobic workstation (gas mix CO₂:N₂:H₂, ratio 7:11:2) approximately 24 h before use, enabling agar to adjust to anaerobic atmosphere.
4. Incubate anaerobic agar cultures at 37 °C in an anaerobic workstation.

C. Mating of donor *E. coli* CA434 and recipient bacterium

1. Prepare 7.5 ml overnight culture of recipient. Donor, recipients and plasmids are described in Table 1.

Table 1. Strains and plasmids

Strain/plasmid	Relative characteristics	Source/Reference
Strains		
<i>Eubacterium rectale</i> A1-86	Butyrate producing, strict anaerobes of the	(Barcenilla <i>et al.</i> , 2000)
<i>Eubacterium rectale</i> T1-815	<i>Lachnospiraceae</i> family.	(Barcenilla <i>et al.</i> , 2000)
<i>Roseburia inulinivorans</i> A2-194		(Duncan <i>et al.</i> , 2006)
<i>Eubacterium rectale</i> EAM3	<i>E. rectale</i> A1-86 harbouring pMTL83151	(Sheridan <i>et al.</i> , 2019)
<i>Eubacterium rectale</i> ETM3	<i>E. rectale</i> T1-815 harbouring pMTL83151	(Sheridan <i>et al.</i> , 2019)
<i>Eubacterium rectale</i> ETBglu	<i>E. rectale</i> T1-815 harbouring pMTL3 β -glu	(Sheridan <i>et al.</i> , 2019)
<i>Roseburia inulinivorans</i> RAM2	<i>R. inulinivorans</i> A2-194 harbouring pMTL82151	(Sheridan <i>et al.</i> , 2019)
<i>Roseburia inulinivorans</i> RAM3	<i>R. inulinivorans</i> A2-194 harbouring pMTL83151	(Sheridan <i>et al.</i> , 2019)
<i>Escherichia coli</i> CA434	Conjugative donor. Genotype: HB101 (<i>thi-1</i> hsdS20 (r-B, m-B) <i>supE44</i> <i>recAB</i> <i>ara-14</i> <i>leuB5proA2</i> <i>lacY1</i> <i>galK</i> <i>rpsL20</i> (str ^R) <i>xyl-5</i> <i>mtl-1</i>) carrying R701 (Tra+, Mob+ conjugative plasmid)	(Williams <i>et al.</i> , 1990)
Plasmids*		
pMTL82151	ColE1, <i>catP</i> , <i>traJ</i> , MCS and pBP1	(Heap <i>et al.</i> , 2009)
pMTL83151	ColE1, <i>catP</i> , <i>traJ</i> , MCS and pCB102	(Heap <i>et al.</i> , 2009)

*pBP1, pCB102, pCD6 and pIM13 (replicons of these plasmids). ColE1 (Gram-negative replicon), *catP* (chloramphenicol resistance gene), *traJ* (origin of transfer) and MCS (multiple cloning site).

- Inoculate donor *E. coli* CA434 in 40 ml of LB Cm10 in 50 ml conical centrifuge tubes and incubate at 37 °C, 200 RPM overnight.
- Centrifuge donor overnight culture at 1,200 × *g* for 10 min.
- Decant supernatant and resuspend pellet in 20 ml of anaerobic PBS.
- Centrifuge donor again at 1,200 × *g* for 10min.
- Decant supernatant in anaerobic cabinet.
- Transfer overnight culture of recipient bacterium (grown as described in Steps B1 and B2) to anaerobic cabinet.
- Resuspend donor pellet in 1 ml of recipient overnight culture in anaerobic cabinet.
- Spot 100 μ l of this solution on the centre of AMM agar plate and incubate in the anaerobic cabinet at 37 °C for 48 h.
- Scrape mating mix off centre of AMM plate with sterile loop and suspend in 500 μ l of anaerobic PBS.
- Spread 50 μ l of this solution on to the YCFAGSC plates supplemented with either 5 or 7.5 μ g/ml chloramphenicol and incubate anaerobically at 37 °C until colonies appear (usually 48 to 96 h).

12. Restreak single colonies on fresh YCFAGSC plates supplemented with either 5 or 7.5 µg/ml chloramphenicol and incubate anaerobically at 37 °C until single colonies appear. This may take 3 days incubation.
13. Diagrammatic representation of the optimised conjugation protocol shown in Figure 2.

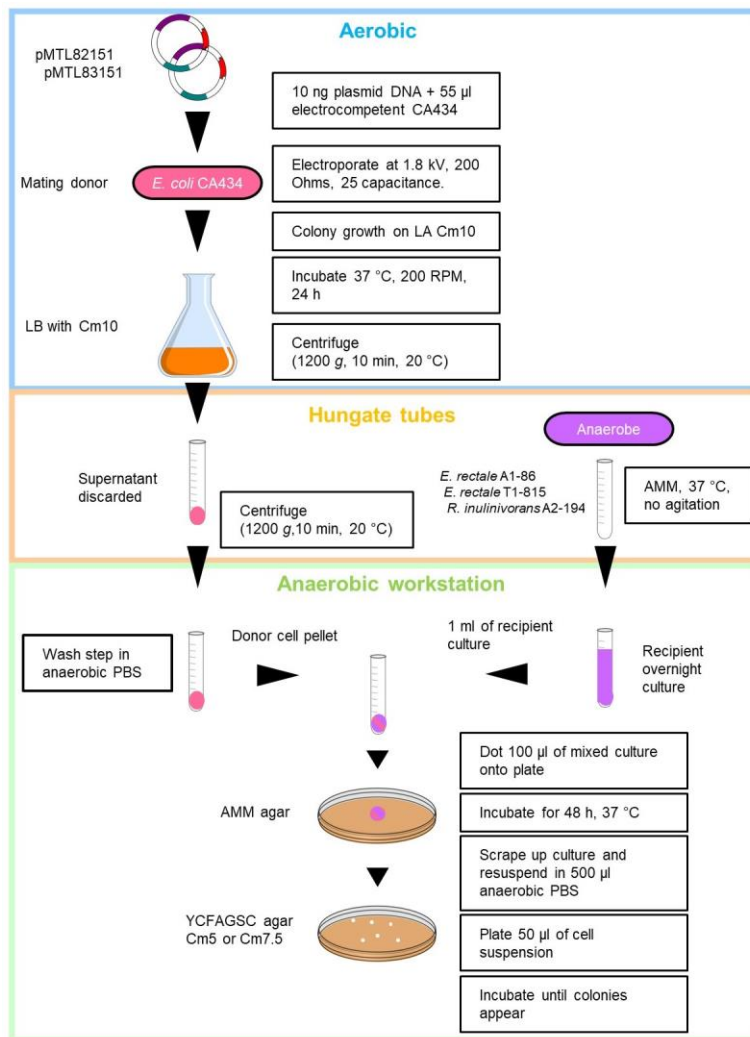


Figure 2. Detailed diagrammatic representation of the optimised conjugation protocol for *Roseburia inulinivorans* and *Eubacterium rectale*. Supplemented with 5 or 7.5 µg/ml chloramphenicol (Cm5 or Cm7.5).

D. Verification of putative transconjugants

It is good practice to perform various tests to confirm the validity of transconjugants. Some simple tests can readily eliminate bacteria that are not transconjugants, and can save time and expense.

1. Incubate putative transconjugants aerobically on AMM agar at 37 °C. The recipient bacterium is incapable of aerobic growth while the donor *E. coli* bacterium will grow.
2. Gram-stain using standard procedures. Coccus species are common contaminants in gut microbiology and are easily differentiated from the rod-like *Roseburia* and *Eubacterium* by

Gram-staining.

3. PCR amplify the 16S rRNA gene directly from colonies or liquid culture using the universal bacterial 16S rRNA gene primers FD1 and RP2 (Table 2) (annealing temp. 52 °C), generating an amplicon of 1,495 bp. PCR amplification conditions are provided in Table 3.

Table 2. Primers

Primer	Sequence (5'-3')	Target	Reference
PS#MTL-for	TATCTATGATACCGTGGTCAAC	pMTL80000 series plasmids	(Sheridan <i>et al.</i> , 2019)
PS#MTL-rev	CTGCTGAAGCCAGTTACC	pMTL80000 series plasmids	(Sheridan <i>et al.</i> , 2019)
FD1	AGAGTTTGATCCTGGCTCAG	Full 16S rRNA gene	(Wood <i>et al.</i> , 1998)
RP2	ACGGCTACCTTGTTACGACTT	Full 16S rRNA gene	(Wood <i>et al.</i> , 1998)
519R	GWATTACCGCGGCKGCTG	16S rRNA gene (universal)	(Turner <i>et al.</i> , 1999)
926F	ACTCAAAGGAATTGACGG	16S rRNA gene (universal)	(Muyzer <i>et al.</i> , 1995)

Nucleotide code: Guanine (G), adenine (A), thymine (T), cytosine (C), adenine or thymine (W) and guanine or thymine (K)

Table 3. PCR protocol

	Temperature	Time
Initial denaturation	94 °C	5 min
25 cycles	95 °C	30 s
	Depending on primers	30 s
	72 °C	2 min
Final Extension	72 °C	8 min
Hold	4 °C	

4. Sanger sequence the resulting amplicon with the primers 519R and 926F (Table 2).
5. BLASTn query each sequence against the NCBI 16S rRNA gene database to confirm identity.
6. Confirm the presence of the plasmid in putative transconjugants by amplifying a nucleotide sequence common to all of the modular plasmids but absent in the recipient's chromosome. The primers PS#MTL-for and PS#MTL-rev (Table 2) (annealing temp 60 °C) amplify a 514 bp region incorporating sections of *catP* gene and ColE1 Gram-negative replicon.
7. Expected results and comments are described in Table 4.

Table 4. Verification procedures for putative transconjugants

	Confirmation	Expected result	Comment
Gram-staining	Absence of contamination	Pink or purple bacilli	Common contaminants, such as coccus species are easily distinguished from <i>Roseburia</i> and <i>Eubacterium</i> by Gram staining.
16S rRNA gene sequencing	Absence of contamination	Match to recipient DNA	
Plasmid-specific PCR	Presence of plasmid	514 bp amplicon	Recipient chromosomes lack the amplified region of DNA
Anaerobic incubation	Absence of donor	No aerobic growth	Recipient anaerobes are incapable of aerobic growth
Southern blotting	Autonomous replication	Single band of same size as linear plasmid	Chromosome insertion would change the size of the probe hybridising band

E. Verification of autonomous plasmid replication by Southern blotting

1. Extract DNA from transconjugants using the Wizard Genomic DNA Purification kit, following the manufacturer's instructions.
2. Digest 1 µg of extracted DNA with HindIII for 3 h at 37 °C to produce restriction fragments of various sizes.

Note: HindIII only cuts the plasmid once, resulting in a single linear fragment, whereas restriction of the genomic DNA results in fragments of various sizes.

3. Separate restriction fragments by size by gel electrophoresis (0.8% agarose, TBE).
4. Photograph gel image with transilluminator to facilitate size inference in the final blot.
5. Gently rock gel in depurination solution (0.25 M HCl) for 7 min and rinse in ddH₂O (double distilled water).
6. Gently rock the gel in denaturation solution (0.5 M NaOH, 1.5 M NaCl) for 30 min three times and rinse in ddH₂O.
7. Gently rock the gel in neutralisation solution (0.5 M Tris-HCl [pH 7.4], 3 M NaCl) for 30 min three times and rinse in ddH₂O.
8. Transfer DNA from gel to nylon membrane using the Hybaid blotter (1.5 h) and UV-crosslinked.
9. Create probe by digesting pMTL83151 with HindIII and ApaLI. This produces two fragments (~3,000 bp and ~1,500 bp), the smaller of which is specific to a region common to all of the plasmids, but not present in the recipient chromosome. Gel purify the smaller fragment and use as probe template DNA.
10. Perform Southern blotting with DIG High Prime DNA Labelling and Detection Starter Kit II, following the manufacturer's instructions.
11. Visualise hybridizing bands by exposing membrane to x-ray film.

This protocol enables the establishment and verification of autonomously replicating plasmids transferred into *Roseburia inulinivorans* and *Eubacterium rectale* and will provide an essential tool in investigating the biochemistry and physiology of these important organisms by genetic manipulation.

Recipes

1. Resazurin solution
 - Add 100 mg of powdered resazurin to 100 ml ddH₂O
2. Anaerobic phosphate buffered saline (PBS)
 - a. Dissolve PBS tablets in 1 L of ddH₂O and add 1 ml resazurin solution
 - b. Place solution in boiling waterbath for 15 min
 - c. Bubble solution with 100% CO₂ until liquid turns from purple to clear
 - d. Dispense in 100 ml aliquots in Wheaton bottles
3. Mineral solution 1
 - K₂HPO₄ 3.0 g
 - ddH₂O to 1 L
 - Store at 4 °C
4. Mineral solution 2
 - KH₂PO₄ 3.0 g
 - (NH₄)₂SO₄ 6.0 g
 - NaCl 6.0 g
 - MgSO₄ 0.6 g
 - CaCl₂ 20.6 g
 - ddH₂O to 1 L
 - Store at 4 °C
5. Short chain fatty acid solution
 - Acetic acid 17 ml
 - Propionic acid 6 ml
 - n-Valeric acid 1 ml
 - Iso-Valeric acid 1 ml
 - Iso-Butyric acid 1 ml
 - Store at 4 °C
6. Vitamin solution 1
 - Biotin 1 mg
 - Cobalamin 1 mg
 - p-Aminobenzoic acid 3 mg
 - Folic acid 5 mg
 - Pyridoxamine 15 mg
 - ddH₂O to 100 ml

- Store at -20 °C
7. Vitamin solution 2
 - Thiamin 5.0 mg
 - Riboflavin 5.0 mg
 - ddH₂O to 100 ml
 8. Haemin solution
 - KOH 0.28 g
 - Ethanol 95% 25 ml
 - Haemin 100 mg
 - ddH₂O to 100 ml
 - Store at 4 °C
 9. SOC
 - Bacto tryptone 2 g
 - Yeast Extract 0.5 g
 - NaCl 200 µl of 5 M
 - KCl 250 µl of 1 M
 - ddH₂O to 100 ml
 - a. Stir to dissolve, autoclave, then cool to room temperature
 - b. Add 1 ml of filter sterile 2 M Mg stock solution (1 M MgCl₂·6H₂O and 1 M MgSO₄·7H₂O) to give a final conc of 20 mM
 - c. Before use add 20 µl of 1 M sterile glucose per 1 ml of SOC
 10. LB
 - Bacto tryptone 1 g
 - Yeast extract 0.5 g
 - NaCl 1 g
 - Deionized H₂O to 100 ml
 11. LA
 - Bacto tryptone 1 g
 - Yeast extract 0.5 g
 - NaCl 1 g
 - Agar 1.5 g
 - Deionized H₂O to 100 ml
 12. YCFAGSC and AMM
 - Bacto casitone 10.0 g
 - Yeast extract 2.5 g
 - NaHCO₃ 4.0g
 - Glucose 2.0 g
 - Soluble starch 2.0 g
 - Cellobiose 2.0 g

Mineral solution 1 150.0 ml
 Mineral solution 2 150.0 ml
 Haemin solution 10.0 ml
 Vitamin solution 1 (before autoclaving) 1.0 ml
 Vitamin solution 2 (after autoclaving) 1.0 ml
 Resazurine 1.0 ml
 L-cysteine 1.0 g
 Acetic acid (0.7 ml) to make AMM or short chain fatty acid solution (3.1 ml) to make YCFAGSC
 Add 20 g of agar to these recipes to make AMM or YCFAGSC agar
 Distilled water up to 1 L

Acknowledgments

The Rowett Institute (University of Aberdeen) receives financial support from the Scottish Government Rural and Environmental Sciences and Analytical Services (RESAS). The protocol is derived from work published in “Heterologous gene expression in the human gut bacteria *Eubacterium rectale* and *Roseburia inulinivorans* by means of conjugative plasmids” *Anaerobe* 59: 131-140 (2019) (Sheridan *et al.*, 2019).

Competing interests

The authors state that there are no competing interests

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