

1 **SUPPLEMENTAL MATERIAL FOR: Single-Cell Analysis Reveals that the**
2 **Enterococcal Sex Pheromone Response Results in Expression of Full-length**
3 **Conjugation Operon Transcripts in All Induced Cells**

4
5 Rebecca J. B. Erickson,^a Arpan A. Bandyopadhyay,^b Aaron M. T. Barnes,^{a,c} Sofie A.
6 O'Brien,^b Wei-Shou Hu,^b Gary M. Dunny^a

7 ^aDepartment of Microbiology and Immunology, University of Minnesota, Minneapolis,
8 Minnesota, USA, ^bDepartment of Chemical Engineering and Materials Science,
9 University of Minnesota, Minneapolis, Minnesota, USA, ^cDepartment of Laboratory
10 Medicine and Pathology, University of Minnesota, Minneapolis, Minnesota, USA

11

12 **SUPPLEMENTAL TEXT**

13 **Not all induced cells appear to transfer by fluorescent reporter analysis of**
14 **plasmid induction and transfer.** Fluorescent reporters were designed to examine
15 functional conjugation by induced cells at the single cell level (1). Microscopic tracking
16 of the fluorescent expression of individual cells after mixing of donors and recipients
17 embedded in agar pads allowed observation of donor induction and in some cases,
18 formation of new transconjugants cells via transfer of the plasmid from an induced
19 donor to a recipient cell as shown in the main paper (Fig. 2). Figure S1 shows a much
20 larger field that was imaged as a merge of red, blue, and green fluorescence after 330
21 minutes of incubation of the embedded mating mixture. It is readily apparent that the
22 vast majority of the donor cells are surrounded by recipients, with a significant fraction
23 of this population appearing to be induced (green or yellow cells). However, many

24 uninduced cells (red) surrounded by recipients remained at the end of the experiment.
25 This may be due to relatively low levels of **C** production in the embedded recipient cells,
26 Quantification of induction and conjugation on larger populations of mixed donor
27 and recipient cells by flow cytometry (Fig. S4) also suggested that only a fraction of
28 induced donors transferred the plasmid, even when recipients were in the majority (1).
29 The fluorescent reporter is fused to early Q_{Op}, so it is possible that some of the cells that
30 failed to transfer did not express all of the required conjugation proteins because they
31 lacked full length Q_{Op} transcripts. However, this is unlikely, as we tested that the pCF10
32 derivative containing fluorescent reporters showed very similar transfer frequencies to
33 those of wild type donors in standard liquid or plate matings.

34

35

36 **SUPPLEMENTAL METHODS**

37 **Fluorescent protein reporter strain construction and Microscopic analysis**
38 **were described in the main paper).**

39 **Flow cytometric analysis of donor induction and conjugation in liquid**
40 **matings.** Overnight cultures of *E. faecalis* donors (OG1RF+pCF10_tdTomato_iGFP)
41 and recipients (OG1RF_CFP) were diluted 1:10 in fresh medium and grown to
42 exponential growth phase without antibiotics for 1h at 37 °C. Donors were induced with
43 10 ng/ml of the **C** pheromone followed by 1h incubation at 37°C. The donors and
44 recipients were then mixed in 1:10 ratio and mating was carried out at 37 °C for 2h. The
45 mating suspensions were fixed using 2% (w/v) PFA for 10 mins at 4 °C. The bacterial
46 clumps were broken up by EDTA treatment (100 mM), followed by Proteinase K

47 treatment (50 µg/mL, 10 mins at 55 °C). The cell suspension was further sonicated for
48 10 secs at 20% amplitude. The mating suspensions were then analyzed by flow
49 cytometry using a Fortessa X-20 flow cytometer. Expression of CFP, GFP, and
50 tdTomato was measured for 100,000 - 500,000 cells using the 405, 488, and 561 nm
51 laser excitation lines respectively. The cells were gated and analyzed using FlowJo v8
52 (Tree Star, Ashland, OR).

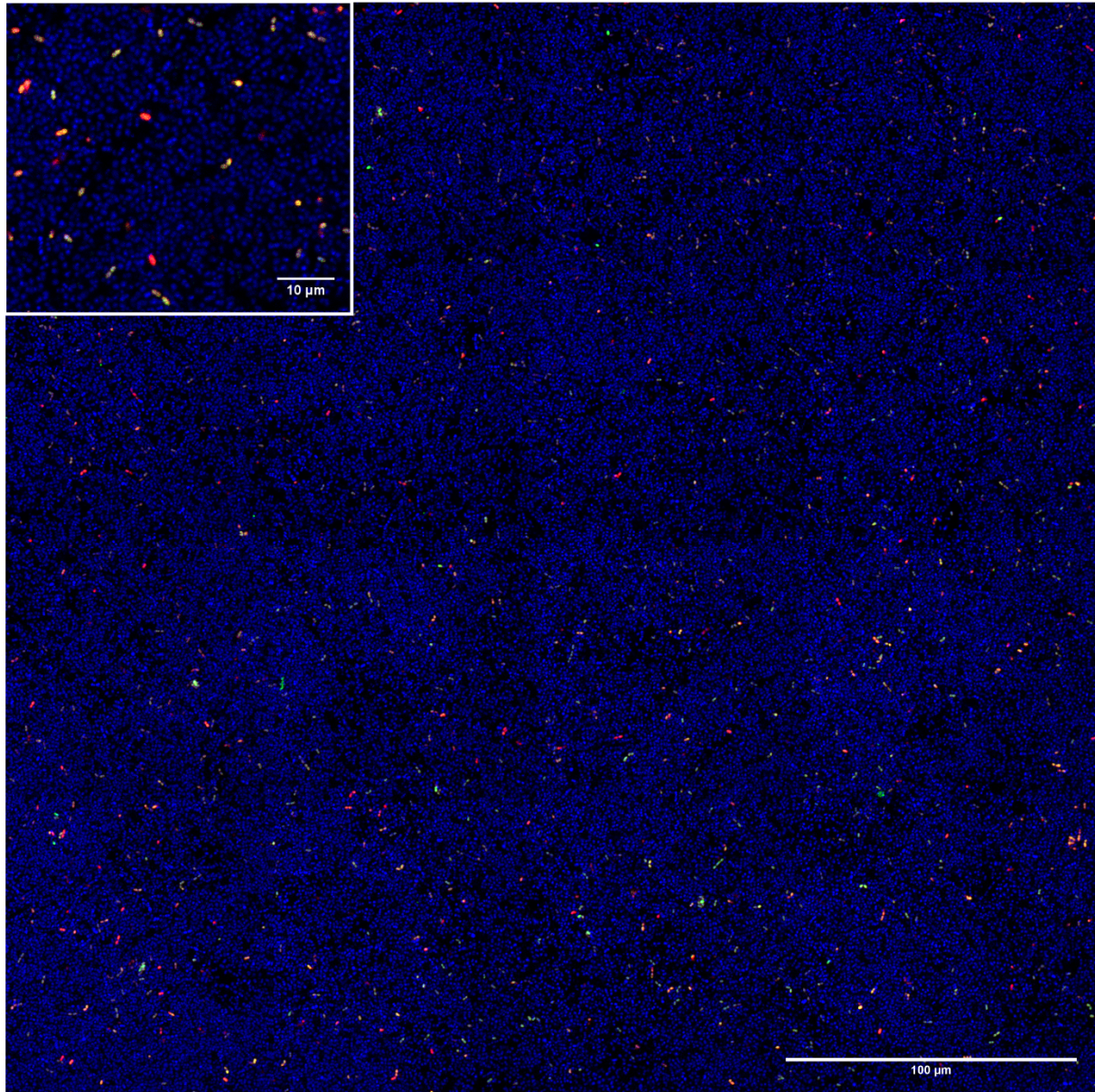
53

54

55 **Supplemental figures / legends**

56 **FIGURE S1: Visualization of induction and conjugation via fluorescent reporter**
57 **protein expression.**

58



59

60

61 A lawn of *E. faecalis* cells on the surface of the agarose pad. This image is composed of
62 16 tiled image stacks stitched together (total area: 375μm × 375μm); a small portion of
63 this image is also shown in Fig. 2. Recipients (blue; Hoechst 33342) and uninduced

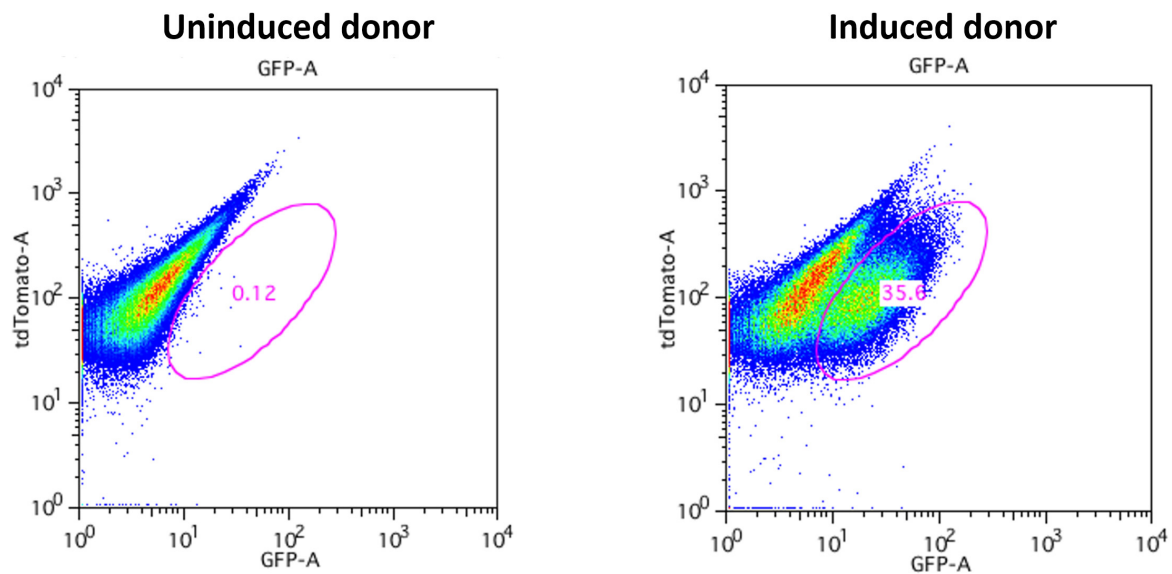
64 donors (red; constitutive tdTomato expression) were mixed (D:R=1:100) and
65 sandwiched between an agarose pad and coverslip-bottomed Petri dish. Donors
66 express GFP (green) when induced by **C**. The entire field was tracked every 30 mins for
67 6 hours using time-lapse laser-scanning confocal microscopy; t=330 min shown here.
68 Note the relative paucity of induced donors relative to uninduced donor cells even
69 though virtually all of the plasmid-carrying cells (td Tomato positive and/or GFP-positive;
70 red, green or yellow in this image) are in close proximity to numerous recipients (Blue).

71

72

73 **FIGURE S2: GFP expression increases ~300-fold after pheromone induction.**
74 Uninduced or induced (10 ng/ml **C** for 60 minutes) exponential phase donor cultures
75 were treated to disperse large clumps, and subjected to flow cytometry as described in
76 the supplemental methods. Nearly all cells express robust red fluorescence, indicative
77 of carriage of pCF10. A. The image demonstrates the lack of green fluorescent cells in
78 uninduced donor populations. B. The image demonstrates the fidelity of the GFP tag as
79 a reporter of induction.

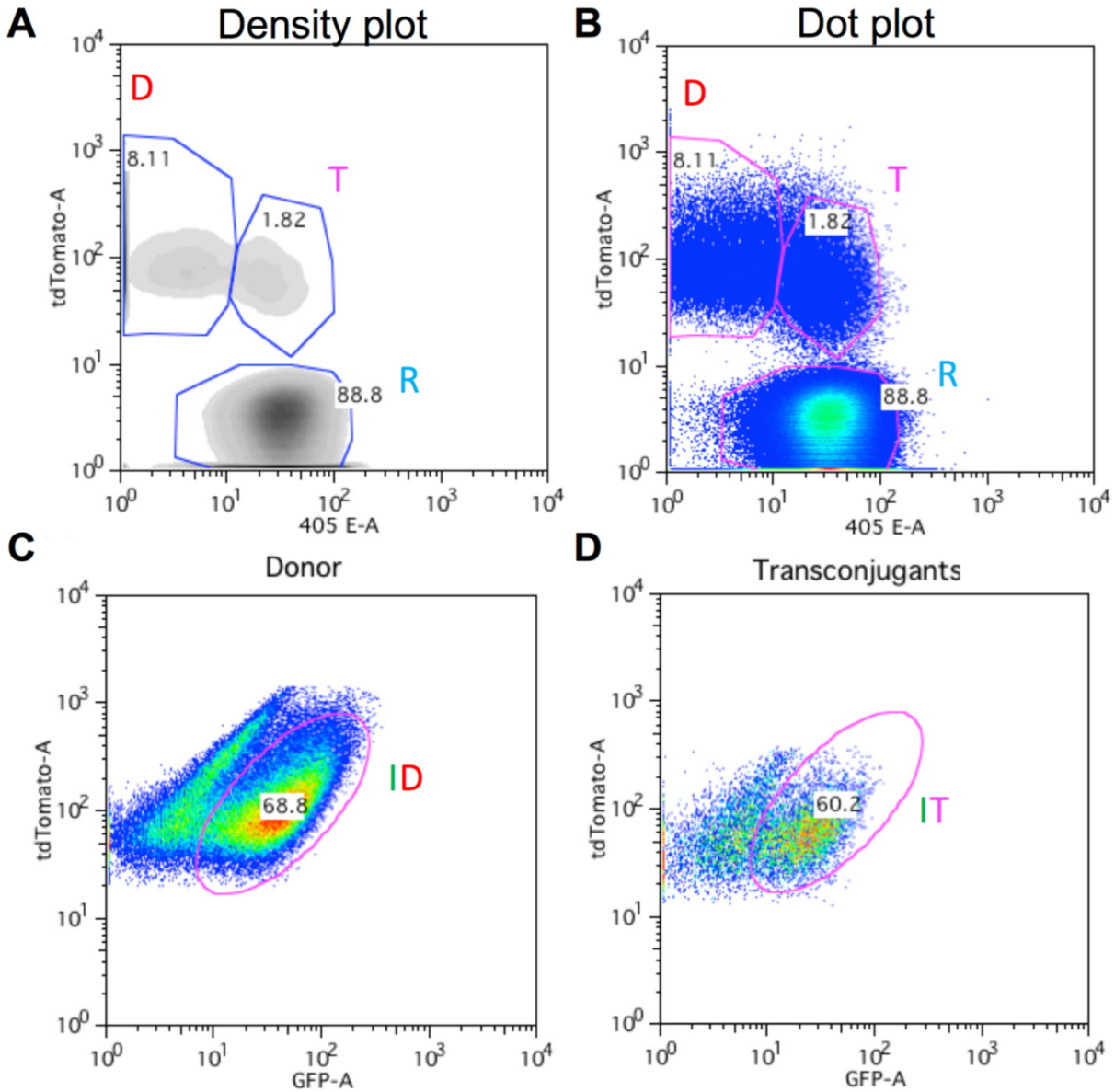
80



81

82 **FIGURE S3: Results of flow cytometry-based analysis of fluorescent protein**
83 **reporters of induction and conjugation.**

84



85

86

87 (A) Two-parameter fluorescence density plot. (B), (C), and (D) Two-parameter
88 fluorescence dot plots. (A-B) Three distinct populations can be identified from
89 differential fluorescence patterns in the density and dot plots, D: donors, R: recipients,
90 and T: transconjugants. The donor: recipient ratio is approximately 1:10. Although over
91 60% of donors were observed to be induced (C) and recipients were in excess,

92 transconjugants only make up about 1.8% of the total population (A) and (B). (D) About
 93 60% of transconjugants were also observed to be induced.

94

95 **Supplemental Tables**

96

97 **Supplemental Table S1: HCR probe sequences**

98

HCR probes	Sequence (portion of probe homologous to transcripts of interest)	Amplifier code
Short QL-0 Short QL-1	GCATTGAATTATTCCACAAGCGGGTCATTTTTAGAAAAATGAGCGTGCTTG CACGTTGTTTGCACGGCTCTTACGAGTAGTTCCAGTACAAACAATGCGTG	B4
prgA-1 prgA-5 prgA-16 prgA-21 prgA-23	TCAGGTGTTTTTGGTTGCGCTTGTTCCTGCTGCCCTGACTTCATTTCCCC TCCACGACTTTTTTAGCTTCGTCTACAACCTGCTTGTGGTCAGTCACTGC GATTTTCATGCTTGCCACTTCGCCGTTTTGGTATTGACTACTATCCTCCAG CTTTCAGCACTTTAGCAGACGTTTGTTCAAATCGCTTGTGCTTCCGCAACG CATAGGCTTCTTGTGCTTGAACCACTGCCGCTTCTGCTTTGTTAACTCG	B1
prgB-3 prgB-5 prgB-8 prgB-16 prgB-22	TTCGCTTCATTTTCAGCTGGTGCAACTTCTGTTGGCTGCCCTAGAGGTTT TTGAACGCATTGTTGGCCCCAAAATCAGAAAACAAAATCCGCATGGCCACC CCCCATTGCTAAAACAACAGAGCCGTCCGCGTTGGTTTTTCATCATTTGCTG TACGCTAGAATAAAGGCTTGTGGGTCTTTGGCAGAAAATCGTCACCGTGCC TGGCTCTACTGTTGCTTGCAGGTGTTTTTGGTGTGGTGGTACAATCACTG	B2D1
prgJ-3 prgJ-6 prgJ-17 prgJ-21 prgJ-27	GACGGTAGTTATCTAAGTGATCTCCCTGATACGCCAATAAAAACGTCTTCC GTCCGTGGCGCAAAAAATAGCTGAACAAAGACAATCGCTCTAAGCCAGTC CGCCATTCTTTACCAGAACCAGACGTTCCGAAAATAAGCCCACTTGGTG CGGTCAACAAGTCCCGCATCCACGTCCGTATAGCTTTTTTAGTAGCGATTC TTGCCCCGCTTTACTGTCTAATAACGTCGTAGGGTTTTGCGTGATACCTG	B3
pcfC-7 pcfC-10 pcfC-15 pcfC-26 pcfC-27	GTCCAATCTCTGGCAATATCCCACCGTCTGGGTCTGTTGTGATAAAAGCTG TCGTCCCCGGTTCGTTTTTCCACTTCTTTTGTGCTTCAGTGACAGCTTC TCATGGTCAAAAACAGAGTAACGTGCAGCAGCTATCCCTAGAACGCTGGC AAACCAGACACTCGTCTCCTCCAATAAGCGCAACTTCACTCCTGTCAATC TCATCAGTCGGTCTGTGCTAGTTCATCTGCTCGGTTCATGCTGAAAAGC	B4
pcfG-4 pcfG-18 pcfG-21 pcfG-24 pcfG-29	TCTGATTTTGGATCGAATGTGTACCCGATTTTTTCGTTGAATTGCTGCTTG CCTTATACCCTCCTACAAAGAGCTTTCCAGAATTGCCATAGCCATAATCG TCACAGGTTCTCTTTTTCAGTGGTGTGTTGTCCATTATACAGTCGTAATTGT GCTCTAACATCTTCTCGTCAAGAGTCTCTAAAGTTTTTTCTGCTTCAAGA AGTTTTGGGCTTAATGTCGGTTTTGCTTCTCTGGCTCTTTTTTCTGTACGGC	B5

99

100 **Supplemental Table S2: HCR amplifier and fluorophore details**
 101

Transcript label	Amplifiers : Fluorophore
Short QL	B4H1, B4H2 : Alexa Fluor (AF) 647
prgA	B1H1, B1H2 : AF 488
prgB	B2H1, B2H2 : AF 488
prgB	B2H1, B2H2 : AF 647
prgJ	B3H1, B3H2 : AF 488
pcfC	B4H1, B4H2 : AF 488
pcfC	B4H1, B4H2 : AF 647
pcfG	B5H1, B5H2 : AF 488

102

103

104 **Supplemental Table S3: Flow cytometry settings (A) and compensation matrix (B)**
 105 **used for analysis shown in Fig. 3.**

A.

Flow cytometry parameter	Function	Voltage
FSC-A	Cell size	444
SSC-A	Cell granularity	253
Alexa Fluor 488	HCR-labeled transcripts	450
Alexa Fluor 647	WGA lectin (cell envelope)	135

106

B.

	Alexa Fluor 488	Alexa Fluor 647
Alexa Fluor 488	100.00	0.00
Alexa Fluor 647	8.59	100.00

107

108 **SUPPLEMENTAL REFERENCES**

- 109 1. Bandyopadhyay AA. 2018. Systems analysis of pheromone signaling and
 110 antibiotic resistance transfer in *Enterococcus faecalis*. Ph.D. thesis. University of
 111 Minnesota. <http://hdl.handle.net/11299/197623>