Figure 1

A. Microscopic image of E. faecalis – GFP

B. 3D rendering of E. faecalis – GFP

C. Graph showing counts of E. faecalis – GFP with error bars, indicating 27.8 ± 2.2%

D. Graph showing fold change of bacterial growth over time (min), comparing E. faecalis and E. faecalis + MCs

E. Bar graph showing growth inhibition (%) for different isolates and infections

Legend: blood isolate (E1), endocarditis (E3), blood isolate (E5), urinary tract infection (E8), endocarditis (E18), aortic valve infection (E19), endocarditis (E21), endocarditis (E26)
Figure 2
Figure 4
Figure 5
Figure 6

A

B

C

D

E

F

G

E. faecalis
E. faecalis + MCs
E. faecalis + cromolyn
E. faecalis + MCs + cromolyn

Fold change of bacterial growth

Fold change of bacterial growth

- - + + + +
- + - - +
- + - - +

E. faecalis
MCs
nuclease
cromolyn

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Figure 7

A

Fold change in bacterial growth

Bacteria + 10µg/ml LL-37
Bacteria + 1µg/ml LL-37
Bacteria + 0.1µg/ml LL-37

B

C

Scale bars: 1 µm