

Pyranoanthocyanins Interfering with the Quorum Sensing of *Pseudomonas aeruginosa* and *Staphylococcus aureus*

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Supplementary Materials

Table S1: Antimicrobial resistance pattern of the multidrug-resistant isolates used in this study.

| Isolate | Antimicrobial resistance pattern |
|---------|--|
| Pa3 | FEP, GEN, CIP |
| PA004 | CIP, GEN, IPM, PTZ, TOB |
| SA007* | CIP, CLI, ERI, FOX, GEN, LEV, MOX, OXA |
| SA011* | CIP, CLI, ERI, LEV, MOX, OXA |

CIP: ciprofloxacin; CLI: clindamycin; ERI: erythromycin; FEP: ceftazidime; FOX: cefotaxime; GEN: gentamicin; IPM: imipenem; LEV: levofloxacin; MOX: moxifloxacin; OXA: oxacillin; PTZ: Piperacillin/tazobactam; TOB: Tobramycin. * These isolates are methicillin-resistant *S. aureus* (MRSA).

Table S2: Primers used in *P. aeruginosa* RT-qPCR analysis.

| Gene | Sequence (5' → 3') | Reference |
|-------------|---------------------------------|-----------|
| <i>rpoS</i> | Forward: CTCCCCGGGCAACTCCAAAAG | [1] |
| | Reverse: CGATCATCCGCTTCCGACCAAG | |
| <i>pqsA</i> | Forward: GTTCTGGTTCCTACCTGCC | [1] |
| | Reverse: CAGCAGGATCTGGTTGTCGT | |
| <i>pqsE</i> | Forward: GGTGAAGGAGGGATCAGCC | [1] |
| | Reverse: AGTGGTCGTAGTGCTTGTGG | |
| <i>pqsR</i> | Forward: GATGCCCTGGCGACGATCAA | [1] |
| | Reverse: CACTGGTTGAAGCGGGAGAT | |
| <i>lasI</i> | Forward: CAGAACGACATCCAGACGCT | [1] |
| | Reverse: TCGATGCCGATCTCAGGTG | |
| <i>lasR</i> | Forward: AGATCCTGTTGGCCTGTTG | |

| | |
|-------------|-------------------------------|
| | Reverse: GGGTAGTTGCCGACGATGAA |
| <i>rhlI</i> | Forward: CAGTCGACCATCCGCAAAC |
| | Reverse: GACGTCCCTGAGCAGGTAGG |
| <i>rhlR</i> | Forward: GTTGCGTAGCGAGATGCAG |
| | Reverse: GGCGTAGTAATCGAAGCCCC |

Table S3: Primers used in *S. aureus* RT-qPCR analysis.

| Gene | Sequence (5' -> 3') | Reference |
|---------------|---|-----------|
| 16S | Forward: CCATAAAGTTGTTCTCAGTT | |
| rRNA | Reverse: CATGTCGATCTACGATTACT | |
| <i>agrA</i> | Forward: ACGTGGCAGTAATTCACTGTATGTT Reverse: GGCAATGAGTCTGTGAGATTTGT | |
| <i>sarA</i> | Forward: GCTGTATTGACATACATCAGCGAAA Reverse: CGTTGTTGCTTCAGTGATTCGT | |
| <i>RNAIII</i> | Forward: GAATTGTTCACTGTGTCGATAATCCATT Reverse: GAAGGAGTGATTCAATGGCACAAAGATAT | [2] |
| <i>ica</i> | Forward: TCGCACTCTTATTGATAGTCGCTACGAG Reverse: TGGGACAAGAACTACTGCTCGTTAAT | |
| <i>hla</i> | Forward: ATGGCTCTATGAAAGCAGCAGA Reverse: AAGGTGAAAACCCTGAAGA | |

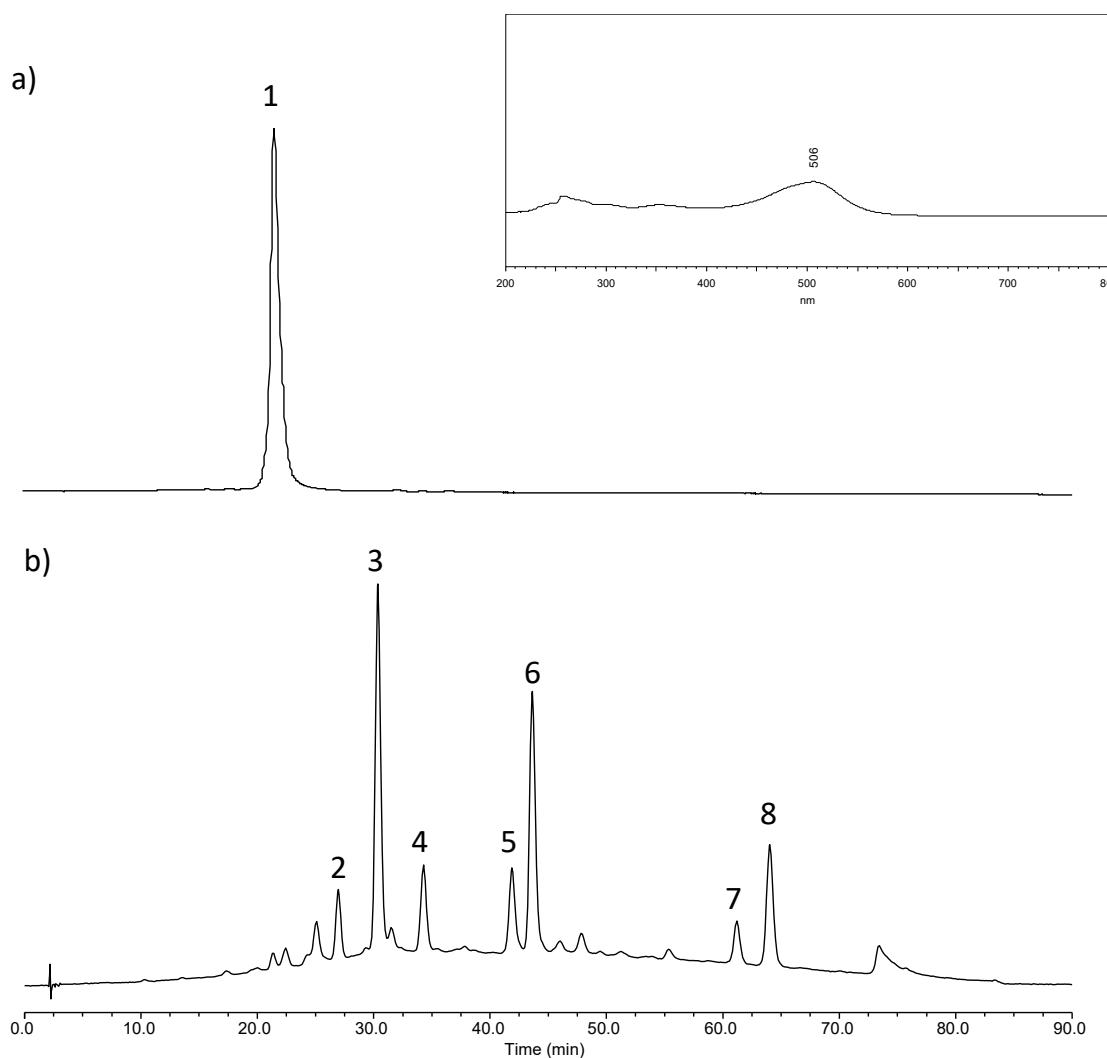


Figure S1. Chromatographic profile of a) 1: carboxypyranocyanidin-3-O-glucoside (carboxypyCy-3-glc), UV-vis spectra at the top right and b) red wine carboxypyrananthocyanins, at 520 nm. 2: Carboxypyranopeonidin-3-glucoside, 3: Carboxypyranomalvidin-3-glucoside; 4: Carboxypyranomalvidin-3-(6''-acetylglucoside); 5: Carboxypyranopeonidin-3-(6''-coumaroylglucoside); 6: Carboxypyranomalvidin-3-(6''-coumaroylglucoside); 7: Peonidin-3-(6''-coumaroylglucoside); 8: Malvidin-3-(6''-coumaroylglucoside).

References

1. Birmes, F.S.; Säring, R.; Hauke, M.C.; Ritzmann, N.H.; Drees, S.L.; Daniel, J.; Treffon, J.; Liebau, E.; Kahl, B.C.; Fetzner, S. Interference with *Pseudomonas aeruginosa* quorum sensing and virulence by the mycobacterial pseudomonas quinolone signal dioxygenase AQDC in combination with the N-Acylhomoserine lactone lactonase QSDA. *Infect Immun* **2019** *87*(10). <https://doi.org/10.1128/IAI.00278-19>
2. Chen, Y.; Liu, T.; Wang, K.; Hou, C.; Cai, S.; Huang, Y.; Du, Z.; Huang, H.; Kong, J.; Chen, Y. (2016). Baicalein inhibits *Staphylococcus aureus* biofilm formation and the quorum sensing system in vitro. *PLoS One* **2016**, *11*(4):e0153468. <https://doi.org/10.1371/journal.pone.0153468>