

Table S1. Comparison of inhibition activities of lutein (Lut) and 3'-dehydrolutein (DHLut) towards PAO1

| PAO1 Virulence | % of inhibition | |
|-----------------------------------|-----------------|----------------|
| | Lut at 22 µM | DHLut at 22 µM |
| • Biofilm formation | 56 ± 2 | 57 ± 2 |
| • Motilities | | |
| • Swarming | 40 ± 9 | 36 ± 3 |
| • Twitching | 34 ± 5 | 37 ± 5 |
| • Virulence factors production | | |
| • Pyocyanin | 67 ± 4 | 66 ± 6 |
| • Elastase B | 19 ± 2 | 8 ± 3 |
| • Rhamnolipids | 62 ± 5 | 64 ± 5 |
| • QS-related genes expression | | |
| • <i>lasB</i> | 60 ± 4 | 61 ± 3 |
| • <i>rhlA</i> | 62 ± 2 | 64 ± 4 |
| • <i>rhII</i> | 60 ± 3 | 50 ± 7 |
| • <i>rhIIR</i> | 64 ± 3 | 68 ± 8 |
| • <i>lasI</i> | 56 ± 3 | 53 ± 5 |
| • <i>lasR</i> | 60 ± 5 | 61 ± 6 |
| • <i>vfr</i> | 16 ± 3 | 12 ± 2 |
| • <i>gacA</i> | 2 ± 3 | 5 ± 6 |
| • QS-independent genes expression | | |
| • <i>aceA</i> | 1 ± 3 | 2 ± 4 |

Value in bold were considered significant compared to DMSO condition

Commenté [PD1]: NON : tester !!!!

Table S2: *Pseudomonas aeruginosa* strains and plasmids used in this study

| Strains or plasmids | Relevant characteristics | References |
|------------------------------|---|---------------------------------|
| Strains | | |
| <i>P. aeruginosa</i> PAO1 | Wild-type (strain PAO0001; http://www.pseudomonas.med.ecu.edu/) | |
| <i>P. aeruginosa</i> ΔPA1432 | <i>P. aeruginosa</i> transposon mutant ID11174; <i>lasI</i> ::IS <i>lacZ</i> /hah; Tet ^R | (Jacobs et al., 2003) |
| <i>P. aeruginosa</i> ΔPA3476 | <i>P. aeruginosa</i> transposon mutant ID32454; <i>rhlI</i> ::IS <i>phoA</i> /hah; Tet ^R | (Jacobs et al., 2003) |
| Plasmids | | |
| pLP170 | Broad-host-range <i>lacZ</i> transcriptional fusion vector containing an RNase III splice sequence positioned between the multiple cloning site and <i>lacZ</i> ; Cb ^r | (Pesci et al., 1997) |
| pPCS1001 | pLP170-derivative containing P _{lasR} - <i>lacZ</i> transcriptional fusion | (Pesci et al., 1997) |
| pLPR1 | pLP170-derivative containing P _{rhlR} - <i>lacZ</i> transcriptional fusion | (Van Delden and Iglewski, 1998) |
| pPCS1002 | pLP170-derivative containing P _{rhlR} - <i>lacZ</i> transcriptional fusion | (Pesci et al., 1997) |
| pLP170_gacA | pLP170- derivative containing P _{gacA} - <i>lacZ</i> transcriptional fusion | (Rasamiravaka et al., 2015) |
| pLP170_vfr | pLP170- derivative containing P _{vfr} - <i>lacZ</i> transcriptional fusion | (Rasamiravaka et al., 2015) |
| pQF50 | Broad-host-range promoter-less <i>lacZ</i> transcriptional fusion vector; Cb ^r | (Ishida et al., 2007) |
| pø01 | pQF50-derivative containing P _{lasB} - <i>lacZ</i> transcriptional fusion | (Ishida et al., 2007) |
| pø02 | pQF50-derivative containing P _{rhlA} - <i>lacZ</i> transcriptional fusion | (Ishida et al., 2007) |
| pø03 | pQF50-derivative containing P _{lasI} - <i>lacZ</i> transcriptional fusion | (Ishida et al., 2007) |
| pTB4124 | pQF50-derivative containing P _{aceA} - <i>lacZ</i> transcriptional fusion | (Kretzschmar et al., 2008) |

Tcr,tetracycline resistance; Cbr carbenicillin resistance