

Peer reviewed publications of Vittorio Venturi as of May 2014

1. Zhang, J., L. Du, F. Liu, F. Xu, B. Hu, V. Venturi, and G. Qian, *Involvement of both PKS and NRPS in antibacterial activity in Lysobacter enzymogenes OH11*. FEMS Microbiol Lett, 2014. **in press**.
2. Qian, G., F. Xu, V. Venturi, L. Du, and F. Liu, *Roles of a Solo LuxR in the Biological Control Agent Lysobacter enzymogenes Strain OH11*. Phytopathology, 2014. **104**(3): p. 224-31.
3. Patel, H.K., P. Ferrante, S. Covaceuszach, D. Lamba, M. Scortichini, and V. Venturi, *The Kiwifruit Emerging Pathogen Pseudomonas syringae pv. actinidiae Does Not Produce AHLs but Possesses Three LuxR Solos*. PLoS One, 2014. **9**(1): p. e87862.
4. Passos da Silva, D., M.P. Castaneda-Ojeda, C. Moretti, R. Buonauro, C. Ramos, and V. Venturi, *Bacterial multispecies studies and microbiome analysis of a plant disease*. Microbiology, 2014. **160**(Pt 3): p. 556-66.
5. Licciardello, G., R.W. Jackson, P. Bella, C.P. Strano, A.F. Catara, D.L. Arnold, V. Venturi, M.W. Silby, and V. Catara, *Draft genome sequence of Pseudomonas corrugata, a phytopathogenic bacterium with potential industrial applications*. J Biotechnol, 2014. **175**: p. 65-6.
6. Venturi, V. and C. Fuqua, *Chemical signaling between plants and plant-pathogenic bacteria*. Annu Rev Phytopathol, 2013. **51**: p. 17-37.
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9. Patel, H.K., Z.R. Suarez-Moreno, G. Degrassi, S. Subramoni, J.F. Gonzalez, and V. Venturi, *Bacterial LuxR solos have evolved to respond to different molecules including signals from plants*. Front Plant Sci, 2013. **4**: p. 447.
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12. Gonzalez, J.F. and V. Venturi, *A novel widespread interkingdom signaling circuit*. Trends Plant Sci, 2013. **18**(3): p. 167-74.
13. Gonzalez, J.F., M.P. Myers, and V. Venturi, *The inter-kingdom solo OryR regulator of Xanthomonas oryzae is important for motility*. Mol Plant Pathol, 2013. **14**(3): p. 211-21.
14. Covaceuszach, S., G. Degrassi, V. Venturi, and D. Lamba, *Structural Insights into a Novel Interkingdom Signaling Circuit by Cartography of the Ligand-Binding*

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