



Corrigendum: TaqMan qPCR for Quantification of *Clonostachys rosea* Used as a Biological Control Agent Against *Fusarium graminearum*

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Gimeno A, Sohlberg E, Pakula T, Limnell J, Keller B, Laitila A and Vogelgsang S (2019) Corrigendum: TaqMan qPCR for Quantification of Clonostachys rosea Used as a Biological Control Agent Against Fusarium graminearum. Front. Microbiol. 10:1839. doi: 10.3389/fmicb.2019.01839 Alejandro Gimeno^{1,2}, Elina Sohlberg³, Tiina Pakula³, Jenni Limnell³, Beat Keller², Arja Laitila^{3*} and Susanne Vogelgsang^{1*}

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A Corrigendum on

TaqMan qPCR for Quantification of *Clonostachys rosea* Used as a Biological Control Agent Against *Fusarium graminearum*

by Gimeno, A., Sohlberg, E., Pakula, T., Limnell, J., Keller, B., Laitila, A., et al. (2019). Front. Microbiol. 10:1627. doi: 10.3389/fmicb.2019.01627

In the original article, there was a mistake in **Table 2** as published. The primer sequence (5'-3') of *VTTact*—reverse contained a false nucleobase. The corrected **Table 2** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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TABLE 2 | Primer and hydrolysis probe sequences (5'-3') for the specificamplification of *Clonostachys rosea* with TaqMan qPCR.

Primer/probe	Sequence (5' to 3')
VTTact-forward	GGCCAGAGATTGTGTTGATGA
VTTact-reverse	ACAGGTTAGGCTCAATGCTC
VTTact probe	GAGGCTGGCAAGAGAGGTCAGTCAC