

Imaging mass spectrometry and genome mining reveal highly antifungal virulence factor of mushroom soft rot pathogen.

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Abstract

Caught in the act: imaging mass spectrometry of a button mushroom infected with the soft rot pathogen *Janthinobacterium agaricidamnosum* in conjunction with genome mining revealed jagaricin as a highly antifungal virulence factor that is not produced under standard cultivation conditions. The structure of jagaricin was rigorously elucidated by a combination of physicochemical analyses, chemical derivatization, and bioinformatics.

Beteiligte Forschungseinheiten

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