

Publication

A liquid chromatography-coupled tandem mass spectrometry method for quantitation of cyclic di-guanosine monophosphate

JournalArticle (Originalarbeit in einer wissenschaftlichen Zeitschrift)**ID** 491053**Author(s)** Spangler, Christian; Böhm, Alex; Jenal, Urs; Seifert, Roland; Kaever, Volkhard**Author(s) at UniBasel** [Jenal, Urs](#) ;**Year** 2010**Title** A liquid chromatography-coupled tandem mass spectrometry method for quantitation of cyclic di-guanosine monophosphate**Journal** Journal of Microbiological Methods**Volume** 81**Number** 3**Pages / Article-Number** 226-31**Keywords** HPLC-MS/MS, Quantitation, c-di-GMP, Di-guanylate cyclase

Cyclic di-guanosine monophosphate (c-di-GMP) represents an important ubiquitous second messenger in bacteria. It controls the transition between a sessile and a motile lifestyle of bacteria and, hence, affects the formation of biofilms which are highly resistant to antimicrobial treatment. c-di-GMP is synthesized by di-guanylate cyclases (DGCs) and degraded by specific phosphodiesterases (PDEs), two highly abundant protein families in bacteria. We have established a robust and highly sensitive high performance liquid chromatography-coupled tandem mass spectrometry (HPLC-MS/MS) based method for the quantitation of c-di-GMP and investigated various method performance parameters such as limit of detection (LOD), lower limit of quantitation (LLOQ), linearity, accuracy, recovery and analyte stability. As a proof of principle we used this method to accurately measure the activity of the prototype DGC PleD* from *Caulobacter crescentus* in vitro. In addition the methodology was successfully applied to determine in vivo levels of c-di-GMP in bacterial extracts of *E. coli* at different stages of bacterial growth. This demonstrates that our method is suitable for the sensitive and specific quantitation of c-di-GMP in bacterial cell extracts.

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