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For publication

**Profiling of secondary metabolite gene clusters regulated by LaeA in
Aspergillus niger FGSC A1279 based on genome sequencing and
transcriptome analysis**

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Abstract

The global regulator LaeA controls the production of many fungal secondary metabolites, possibly via chromatin remodeling. Here we aimed to survey the secondary metabolite profile regulated by LaeA in *Aspergillus niger* FGSC A1279 by genome sequencing and comparative transcriptomics between the *laeA* deletion

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