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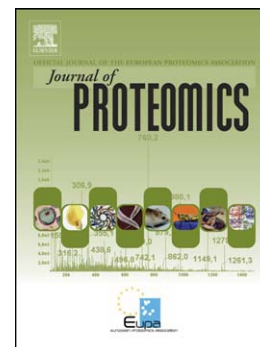
Quantitative proteomics reveals the mechanism and consequence of gliotoxin-mediated dysregulation of the methionine cycle in *Aspergillus niger*

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Mass spectrometry, *Aspergillus fumigatus*, methyltransferase, epigenetics, glycoside hydrolases, biofuels.

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