# Two-dimensional proteome reference maps for the human pathogenic filamentous fungus *Aspergillus fumigatus*.

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#### **ILRS Authors**

Daniela Albrecht

## **Projects**

Integration of transcriptome and proteome data from human-pathogenic fungi Details

### **Abstract**

The filamentous fungus *Aspergillus fumigatus* has become the most important airborne fungal pathogen causing life-threatening infections in immunosuppressed patients. We established a 2-D reference map for *A. fumigatus*. Using MALDI-TOF-MS/MS, we identified 381 spots representing 334 proteins. Proteins involved in cellular metabolism, protein synthesis, transport processes and cell cycle were most abundant. Furthermore, we established a protocol for the isolation of mitochondria of A. fumigatus and developed a mitochondrial proteome reference map. 147 proteins represented by 234 spots were identified.

## **Identifier**

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