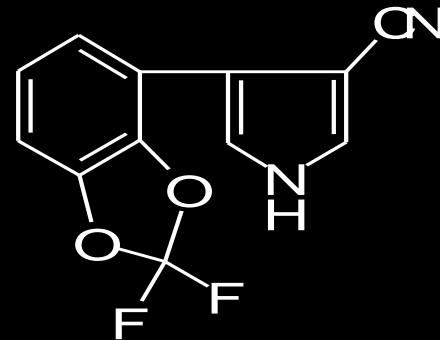
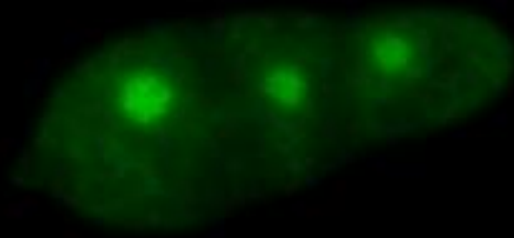


SIGNALING NETWORKS IN FUNGAL PATHOGENS

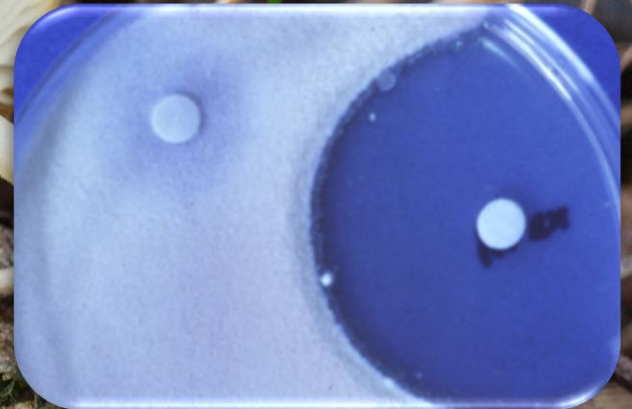
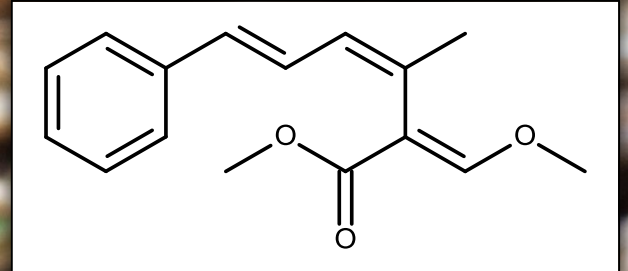
OSMOREGULATION AS FUNGICIDE TARGET



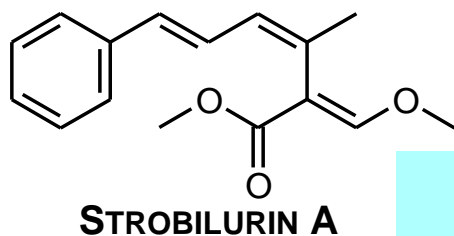
STROBILURUS TENACELLUS

PINECONE CAP

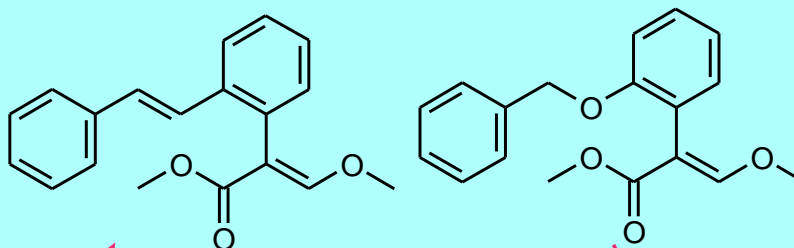
BITTERER KIEFERN-ZAPFENRÜBLING



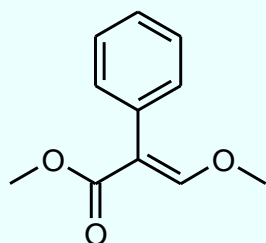
FUNGAL SECONDARY METABOLITES AS LEAD STRUCTURES FOR AGRO-CHEMICALS



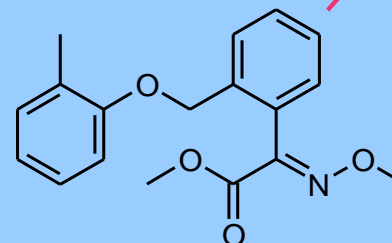
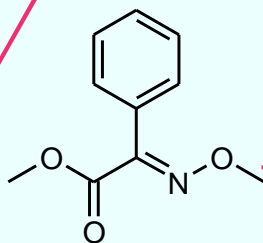
STRONG ANTIFUNGAL ACTIVITY, NON-TOXIC FOR MICE;
MoA: INHIBITION OF ELECTRON TRANSPORT IN THE RESPIRATORY CHAIN (BC1).



B. SCHWALGE, DISSERTATION 1986

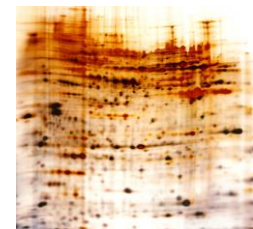
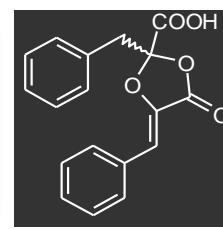
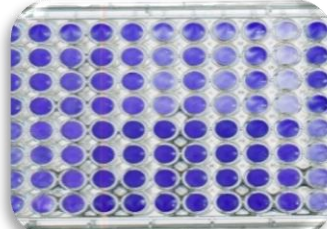
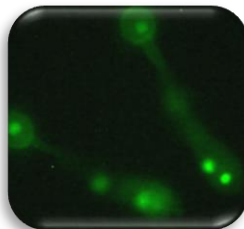
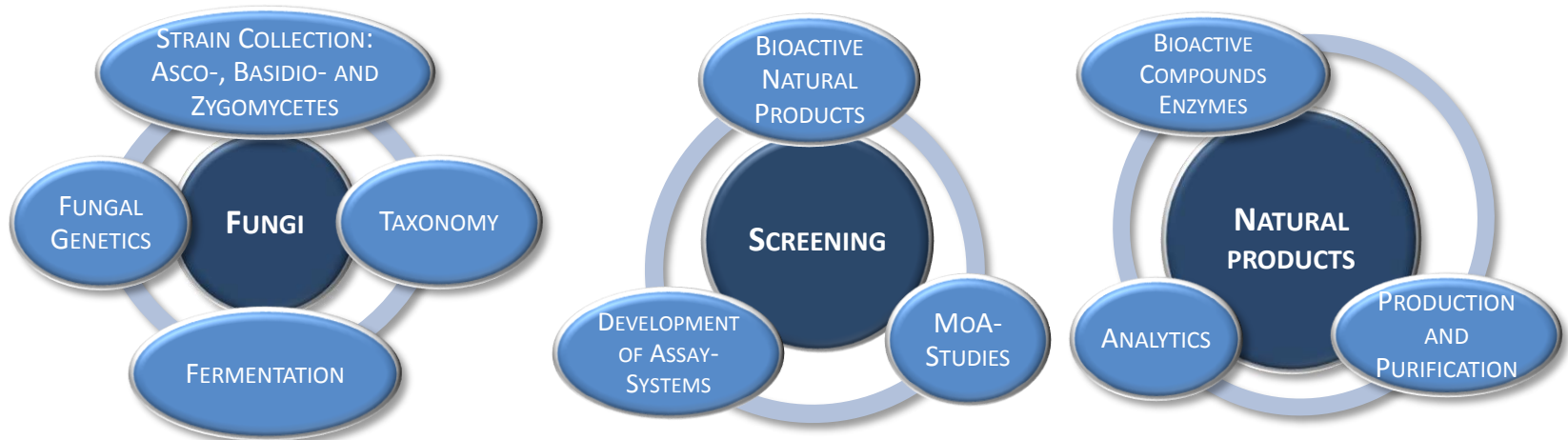


G. SCHRAMM, DISSERTATION 1980



**KRESOXIM-
METHYL
(BASF)**

BIOTECHNOLOGY AT THE IBWF gGMBH



OUTLINE

- **PART I: INTRODUCTION**

- SIGNALING AND OSMOREGULATION
- PHYTOPATHOGENIC FUNGUS *MAGNAPORTHE ORYZAE*

- **PART II: FUNGAL GENETICS**

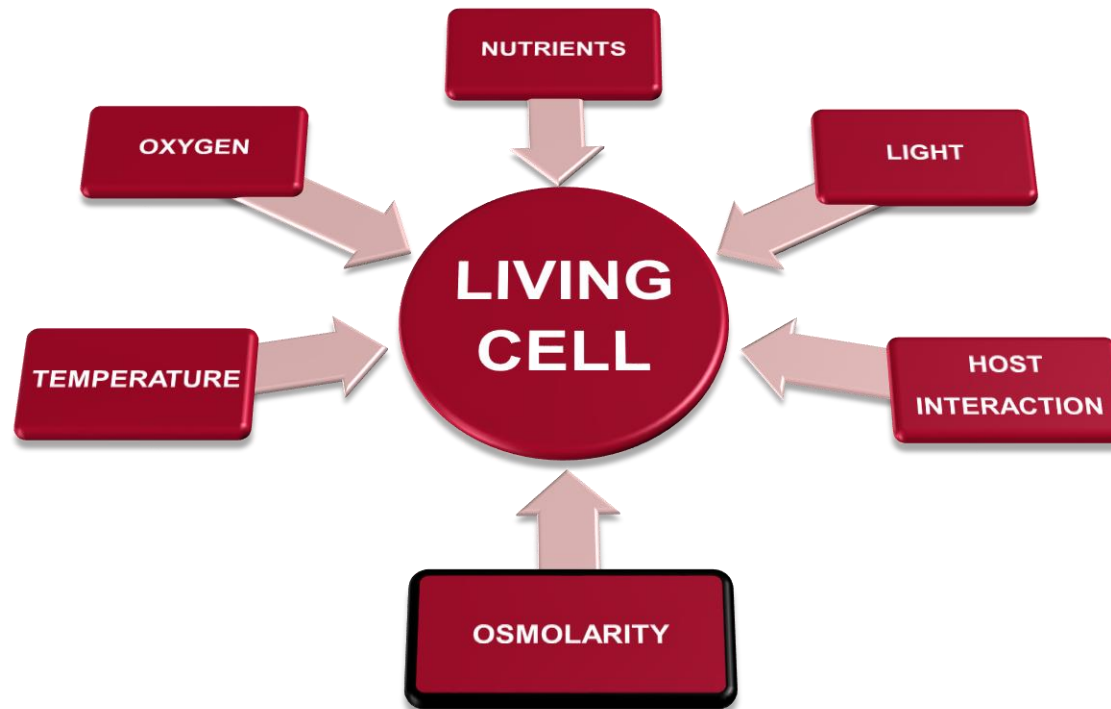
- “LOSS OF FUNCTION”-MUTANTS
- BIOLOGICAL CHARACTERIZATION OF MUTANT STRAINS
- GFP-TAGGING FOR PATHWAY VISUALIZATION

- **PART III: *IN VIVO* TEST SYSTEM**

- TARGET BASED *IN VIVO* TEST SYSTEM FOR THE IDENTIFICATION OF INHIBITORS
- VALIDATION OF INHIBITOR ACTIVITY

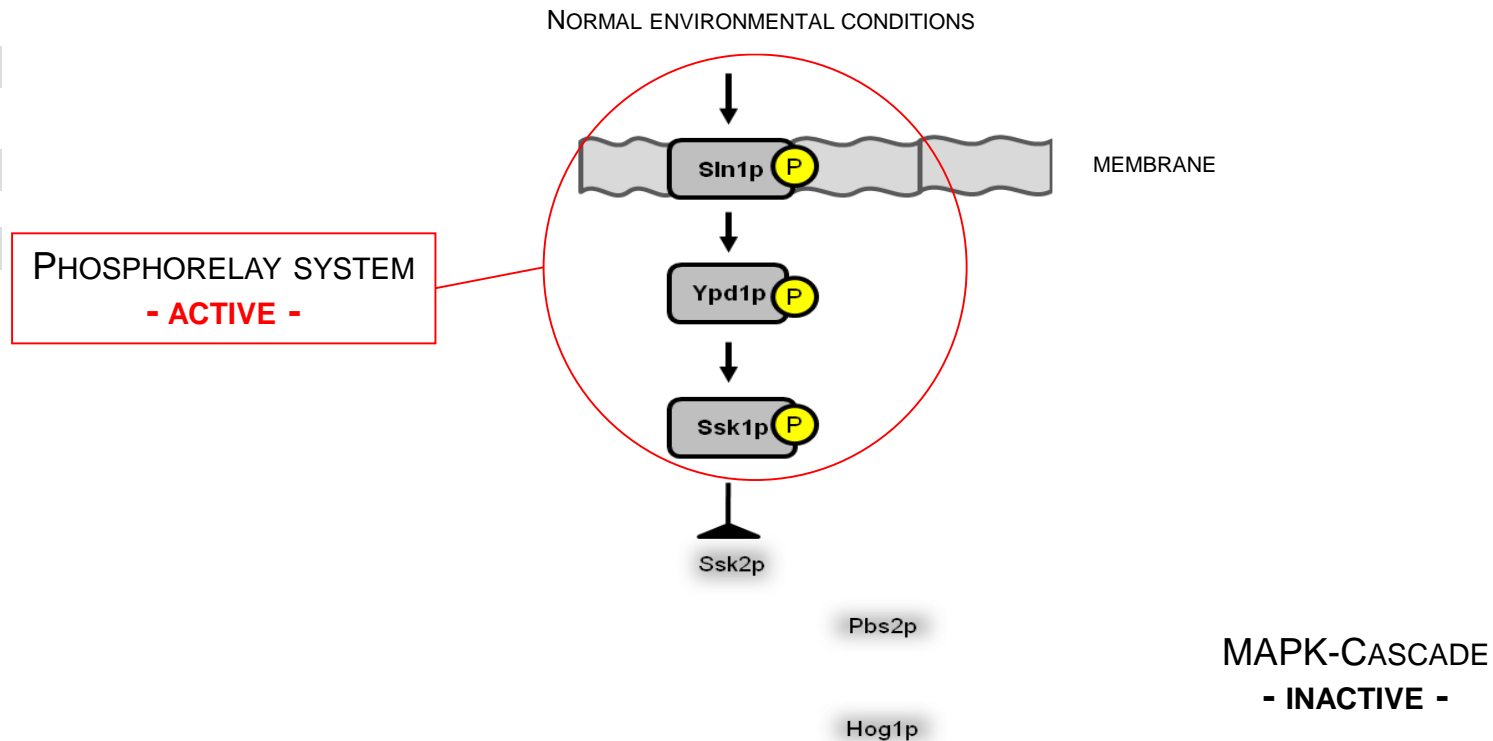
- **CONCLUSION AND DISCUSSION**

MICROORGANISMS AND THEIR ENVIRONMENT



MICROORGANISMS HAVE TO ADAPT RAPIDLY TO CHANGING ENVIRONMENTAL CONDITIONS!

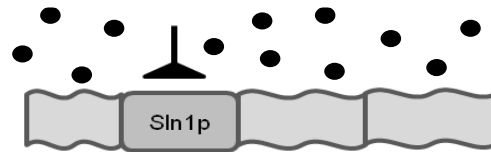
HIGH OSMOLARITY GLYCEROL (HOG) PATHWAY IN YEAST



➔ NORMAL ENVIRONMENTAL CONDITIONS: REGULATORY SYSTEM **INACTIVE**

HIGH OSMOLARITY GLYCEROL (HOG) PATHWAY IN YEAST

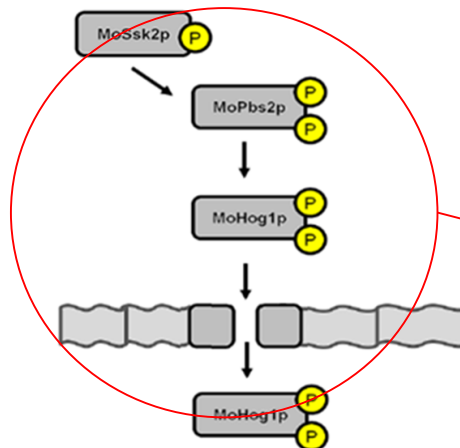
HIGH OSMOLARITY (E.G. SALT STRESS)



Ypd1p

PHOSPHORELAY SYSTEM
- INACTIVE -

Ssk1p

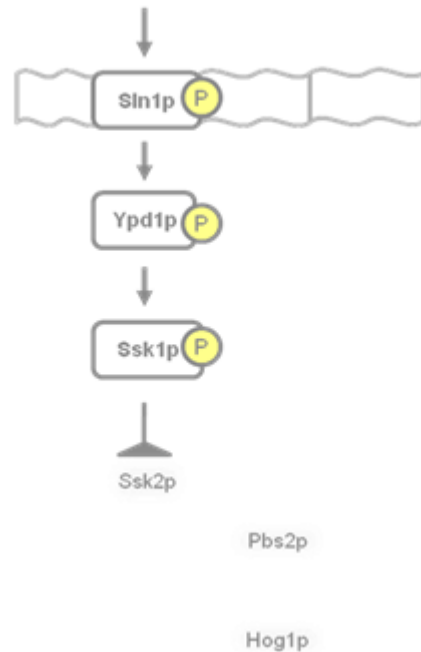


MAPK-CASCADE
- ACTIVE -

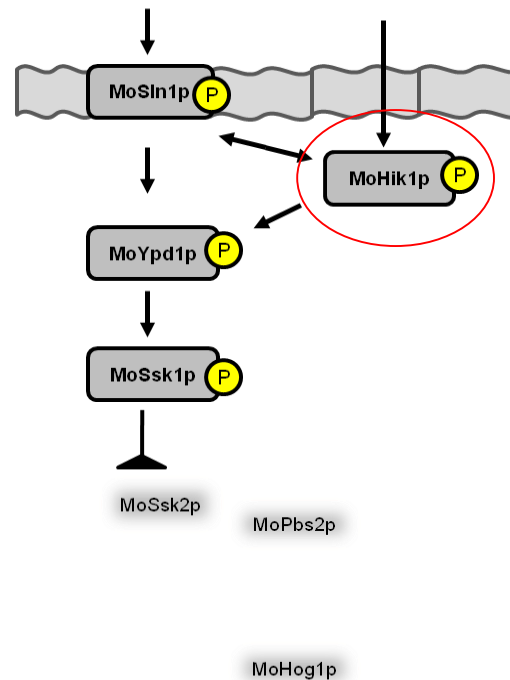
→ HIGH OSMOLARITY: REGULATORY SYSTEM **ACTIVE** (E.G. GLYCEROL PRODUCTION)

SIGNAL PERCEPTION IN FILAMENTOUS FUNGI IS DIFFERENT TO YEAST

SACCHAROMYCES CEREVISIAE

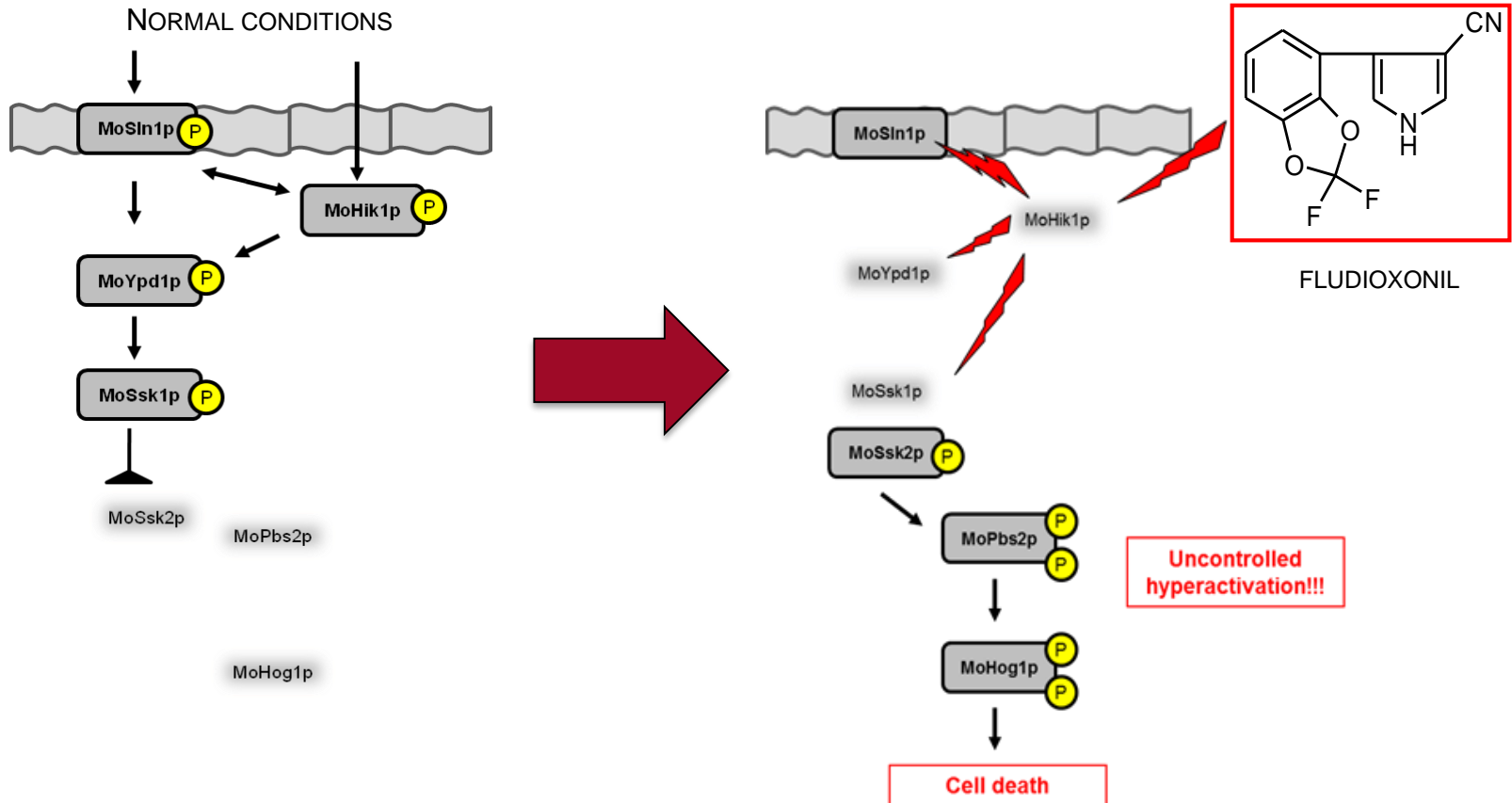


FILAMENTOUS FUNGI



HOG SIGNALLING IN FILAMENTOUS FUNGI COMPRISES MORE ELEMENTS, REFINING BUT ALSO COMPLICATING THE REGULATION SYSTEM!

FLUDIOXONIL HYPERACTIVATES THE PATHWAY

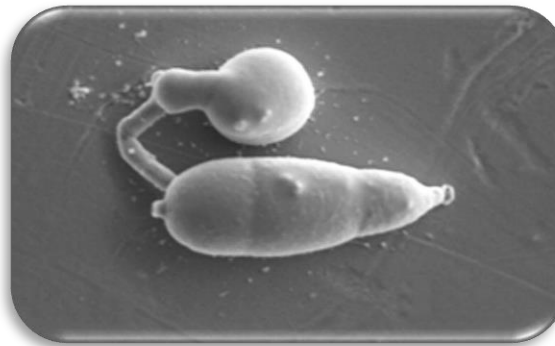


FLUDIOXONIL APPEARS TO INACTIVATE THE PHOSPHORELAY SYSTEM BY INTERACTION WITH MOHIK1!

MAGNAPORTHE ORYZAE – RICE BLAST DISEASE



BIOSPRCTRUM ASIA, 2013



SKAMNIOTI/GURR, 2007



AGRICULTURAL RISK MANAGEMENT IN DEVELOPMENT, 2012

- HEMIBIOTROPHIC ASCOMYCETE, FACULTATIVE PATHOGEN
- GOOD TO HANDLE IN LABORATORY
- GENOME SEQUENCED SINCE 2002
- GENETIC MANIPULATION POSSIBLE (E.G. *AGROBACTERIUM TUMEFACIENS*)

PART II:

FUNGAL GENETICS

MOLECULAR TOOLBOX FOR THE MANIPULATION OF FUNGAL GENOMES

OBTAIN SEQUENCE OF TARGET GENE

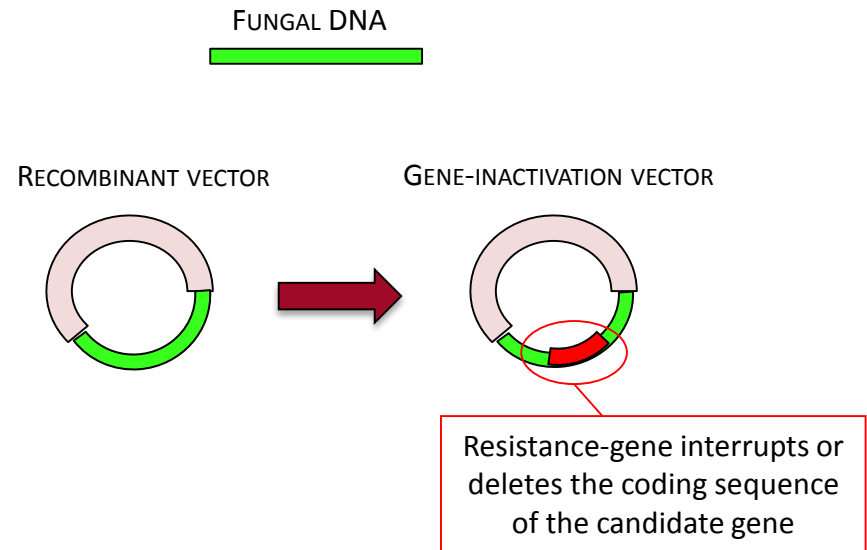
ISOLATE FUNGAL GDNA, AMPLIFICATION BY MEANS OF PCR, RUN AGAROSE GELS, PURIFY DNA, ...

CLONING & MOLECULAR MANIPULATION

PCR-PRODUCT LIGATION IN VECTOR, RESTRICTION ANALYSIS, INTERRUPT OR REPLACE CODING SEQ, SELECTION OF BACTERIAL TRANSFORMANTS, ...

GENERATION OF FUNGAL MUTANT STRAINS (LOSS-OF FUNCTION MUTANTS)

AGROBACTERIUM TUMEFACIENS MEDIATED TRANSFORMATION, SELECTION OF TRANSFORMANTS, SOUTHERN ANALYSIS, PHENOTYPIC CHARACTERIZATION, ...



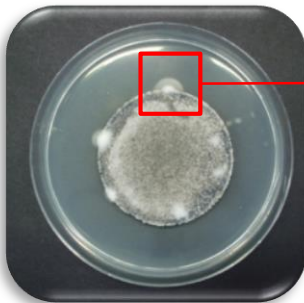
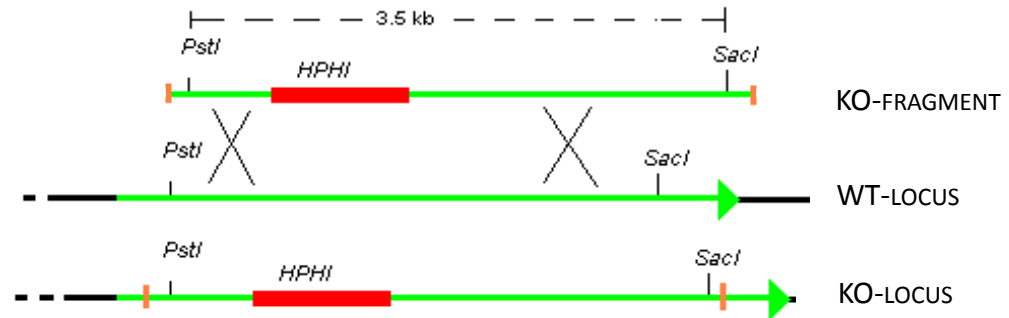
INTERNATIONAL SOCIETY FOR
MICROBIOL ECOLOGY



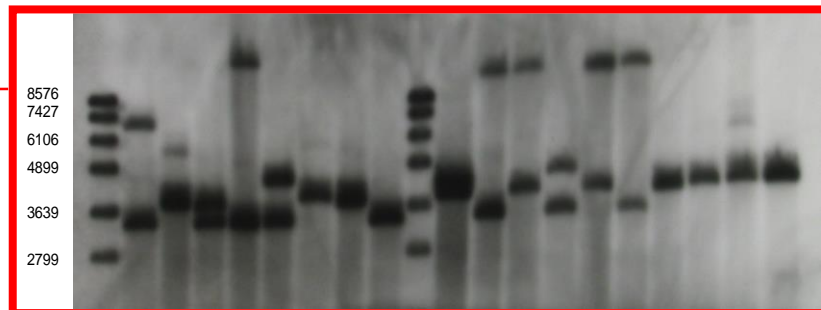
MALCOLM STOREY, 2010,
WWW.BIOIMAGES.ORG.UK

HOMOLOGOUS RECOMBINATION AND ANALYSIS OF MUTANTS

HOMOLOGOUS RECOMBINATION OF
THE T-DNA INTO THE FUNGAL GENOME

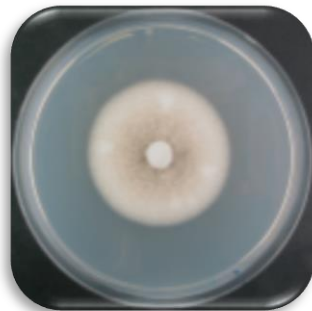


SELECTION OF TRANSFORMANTS

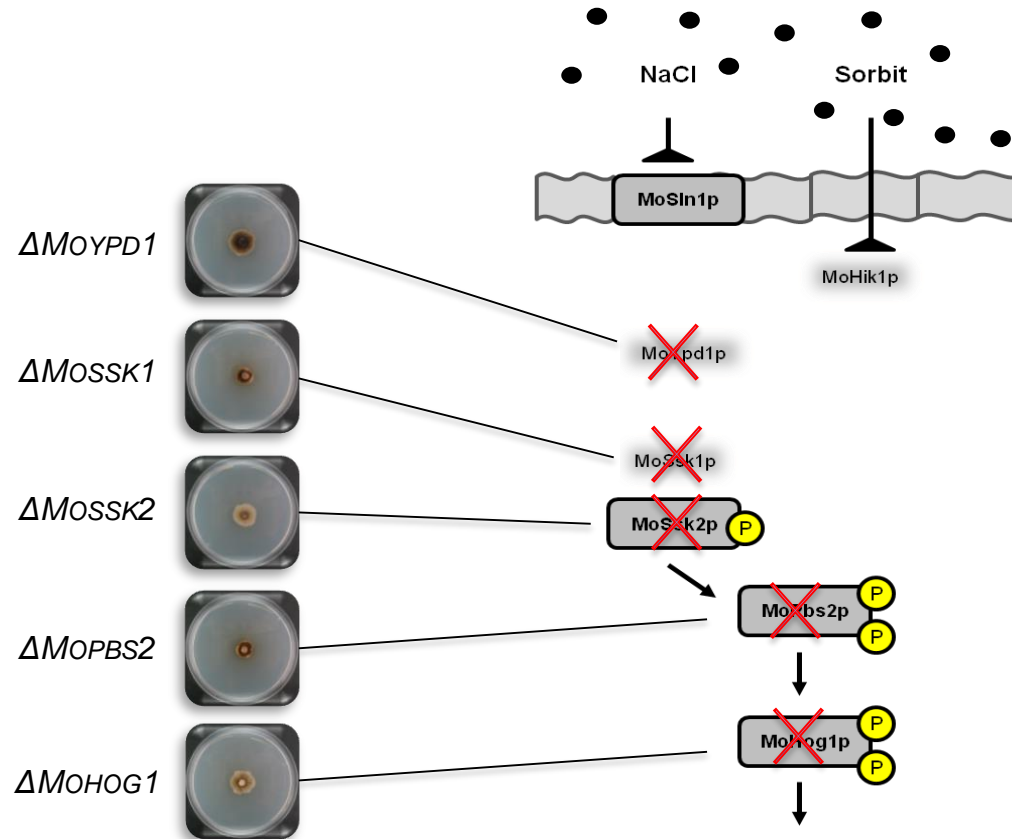


SOUTHERN ANALYSIS OF TRANSFORMANTS

„LOSS OF FUNCTION“-MUTANTS: PHENOTYPE CHARACTERIZATION

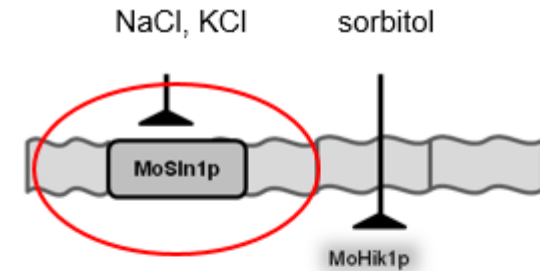
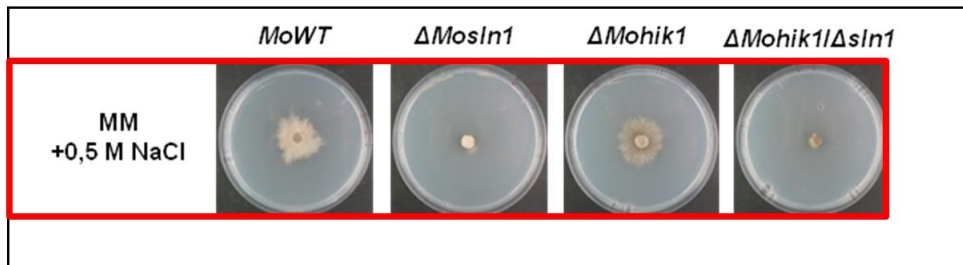


WILDTYPE STRAIN



THE MUTANTS ARE **UNABLE TO ADAPT TO OSMOTIC STRESS** (I.E. NaCl, SORBITOL)

MoSLN1p: A PUTATIVE SALT SENSOR



MoYpd1p

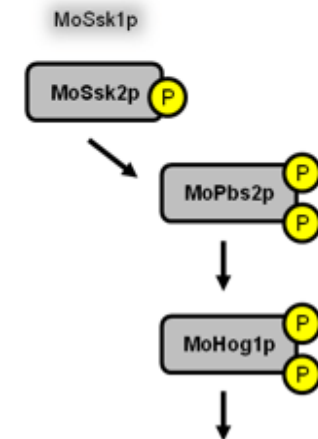
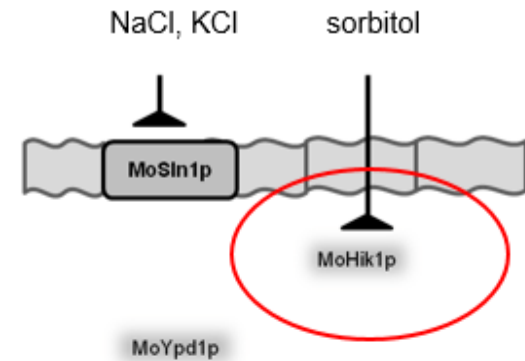
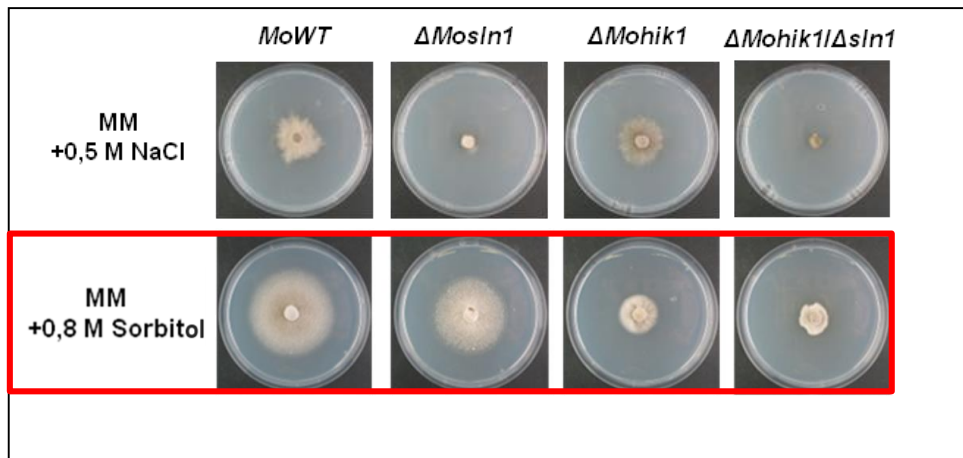
MoSsk1p

MoSsk2p **P**MoPbs2p **P**
PMoHog1p **P**
P

Stress response

 **MoSLN1p** APPEARS TO BE ESSENTIAL FOR **SALT** SENSING!

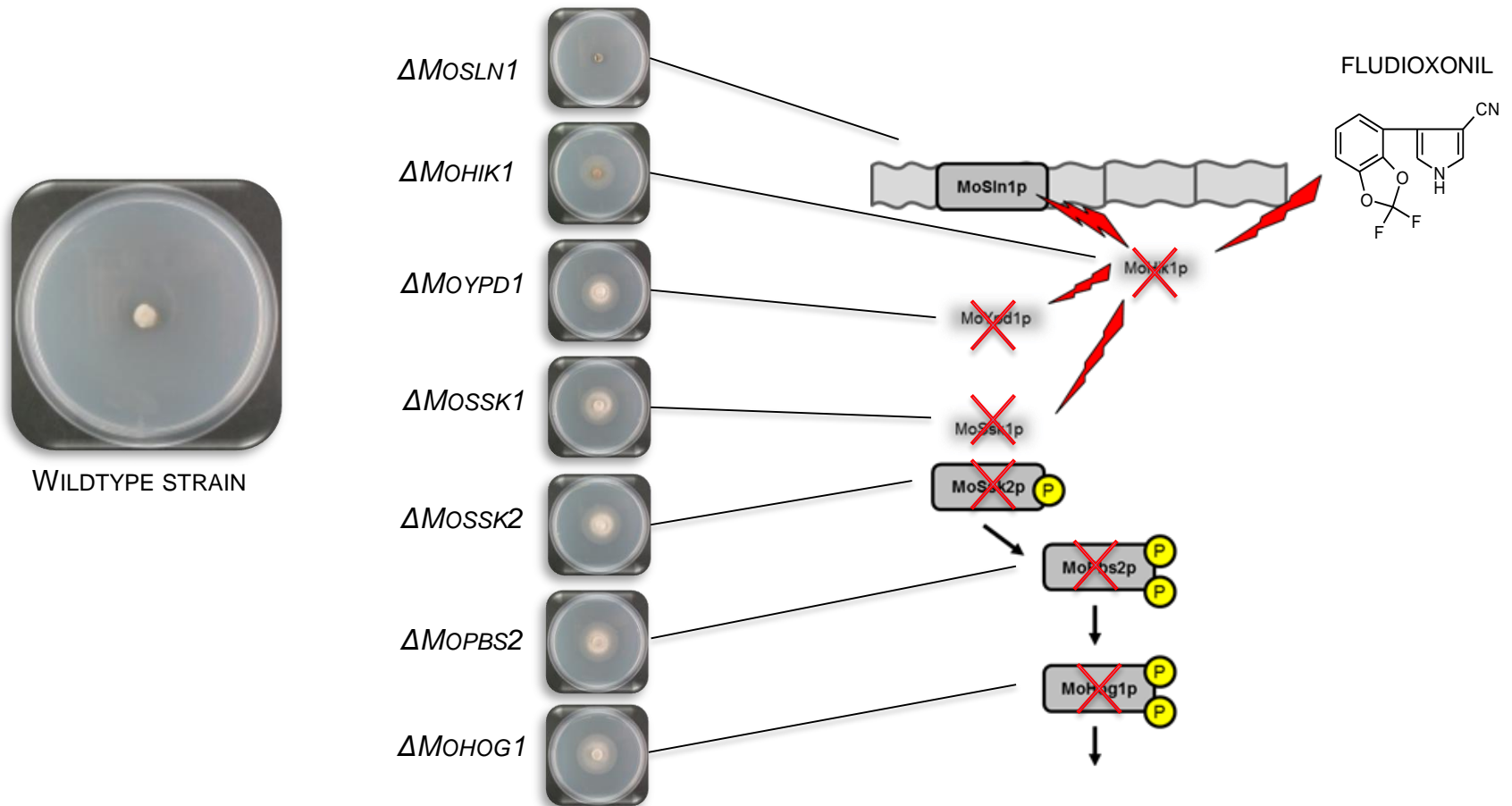
MoHik1p: A PUTATIVE SUGAR SENSOR



Stress response

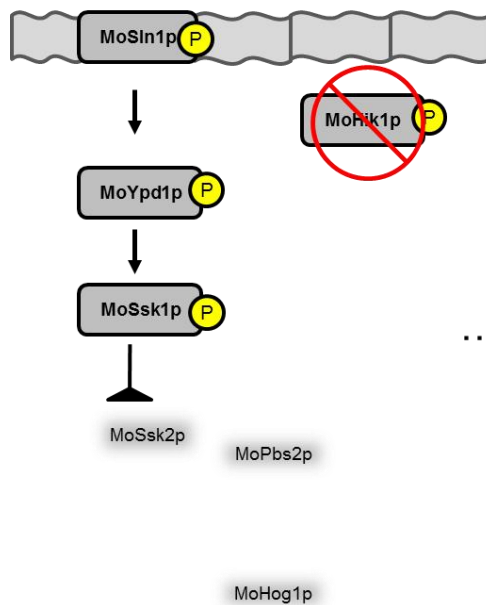
MoHik1p APPEARS TO BE ESSENTIAL FOR **SUGAR** SENSING!

PHENOTYPE OF THE „LOSS OF FUNCTION“-MUTANTS - FUNGICIDE RESISTANCE -

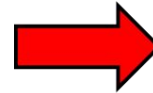


THE MUTANTS (EXCEPT $\Delta MoSLN1$) ARE **RESISTANT** TO HOG EFFECTORS (I.E. FLUDIOXONIL)

GENE INACTIVATION OF *MoHIK1* IS NOT EQUAL TO FUNGICIDE ACTION



...but...

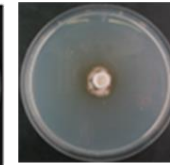


Osmosensitive

Wildtype strain



$\Delta MoHIK1$

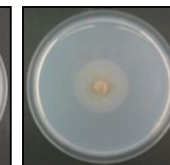


Fludioxonil resistant

Wildtype strain



$\Delta MoHIK1$

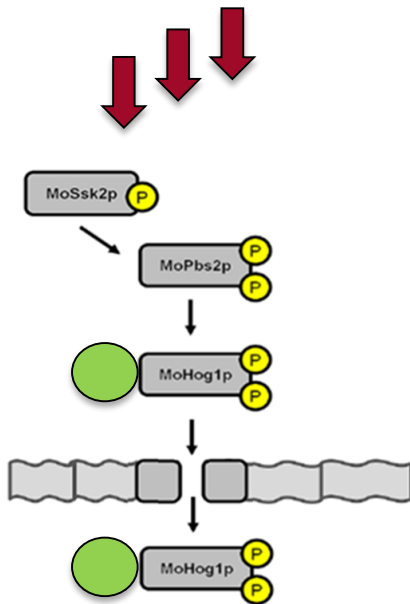


No cell death, phosphorelay is maintained through MoSln1p

MoSLN1P APPEARS TO BE ABLE TO MAINTAIN THE PHOSPHORYLATION PATTERN NEEDED FOR VITALITY!

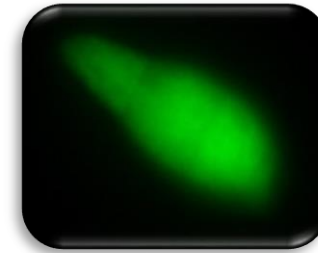
HOG1-GFP FUSION AS TOOL FOR VISUALIZING FUNGICIDE ACTIVITY

PATHWAY ACTIVATION



NUCLEAR ENVELOPE

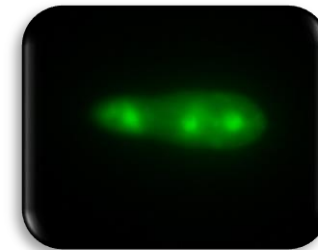
UNTREATED



- FLUDIOXONIL
- NACL
- SORBITOL

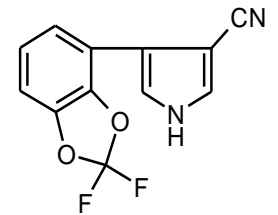
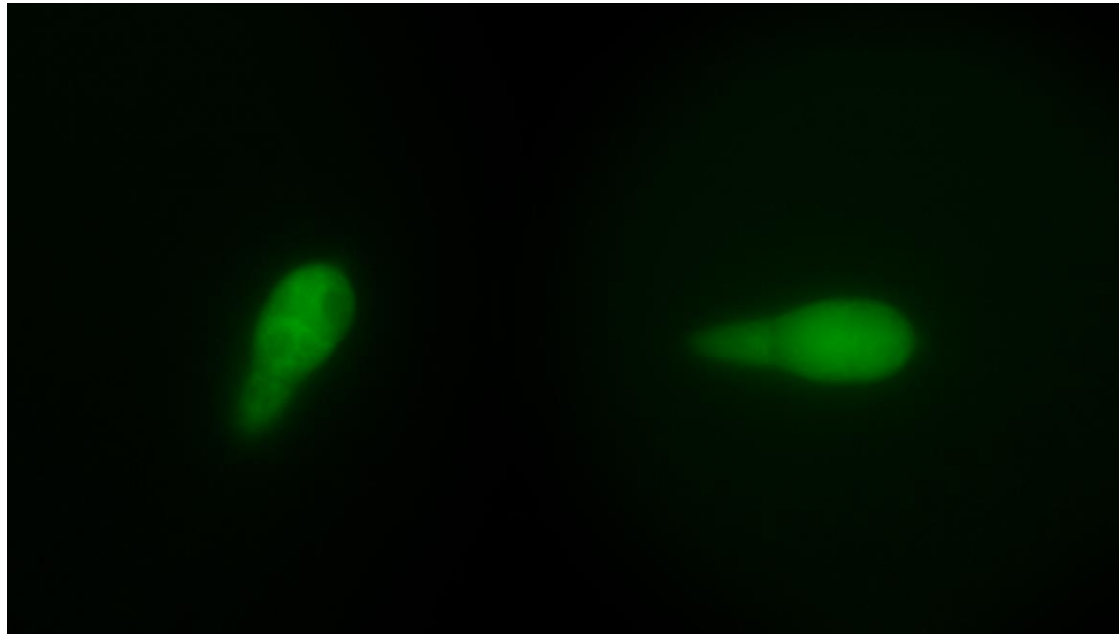


< 1 MINUTE



➔ MOHOG1-GFP FUSION PROTEIN CAN BE USED AS A SUITABLE TOOL FOR „**REAL-TIME**“ VISUALIZATION OF FUNGICIDE ACTION!

HOG1-GFP FUSION AS TOOL FOR VISUALIZING FUNGICIDE ACTIVITY



FLUDIOXONIL

PART II:

„TAKE HOME“-MESSAGE

MUTANT STRAINS WITH INACTIVATED HOG SIGNALING ARE:

- **SENSITIVE** TO ENVIRONMENTAL STRESSES
- **RESISTANT** TO HOG-EFFECTORS (I.E. FLUDIOXONIL)

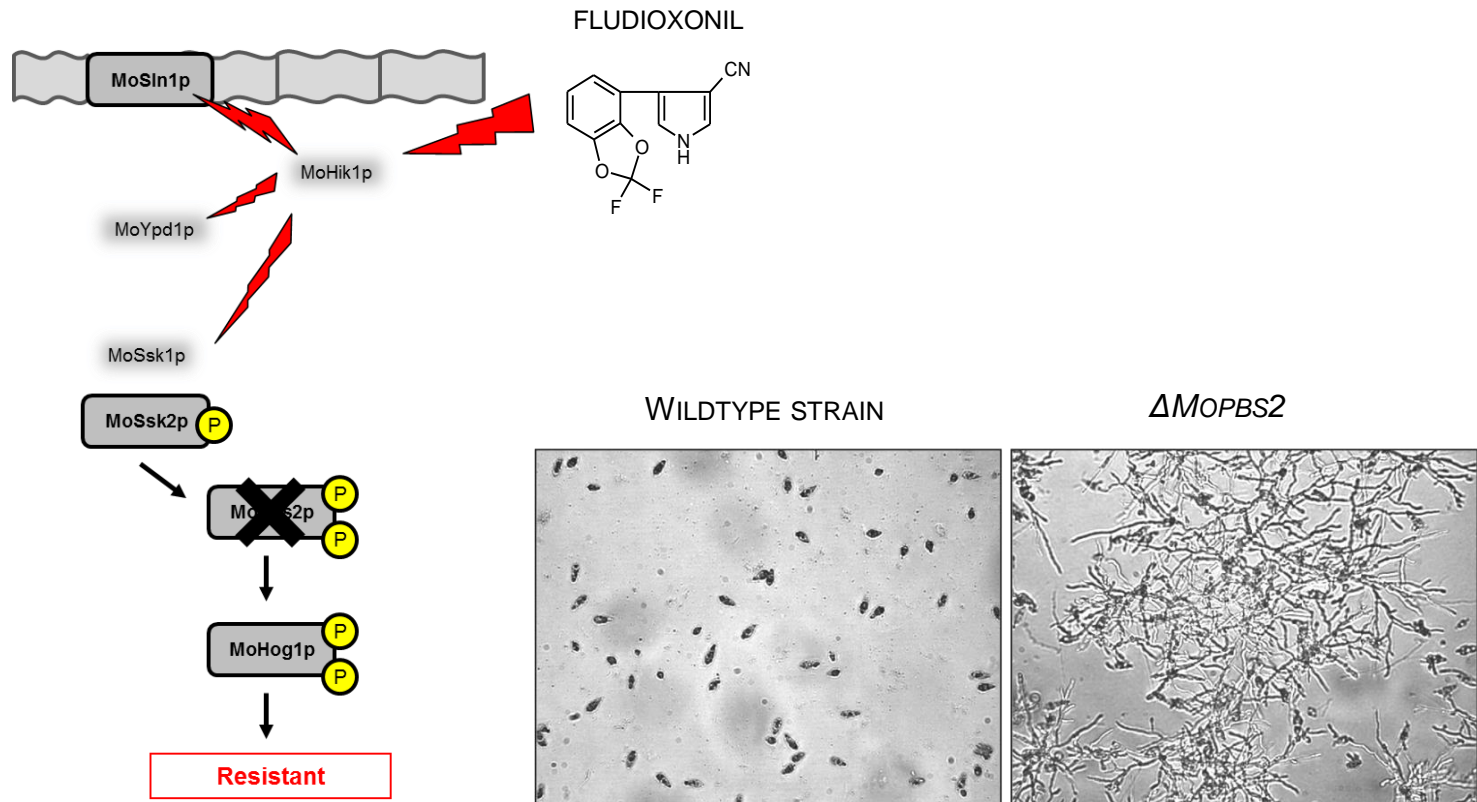
PATHWAY ACTIVITY CAN BE VISUALIZED IN „REAL-TIME“

PART III:

A TARGET BASED *IN VIVO* TEST SYSTEM

A TARGET BASED *IN VIVO* TEST SYSTEM

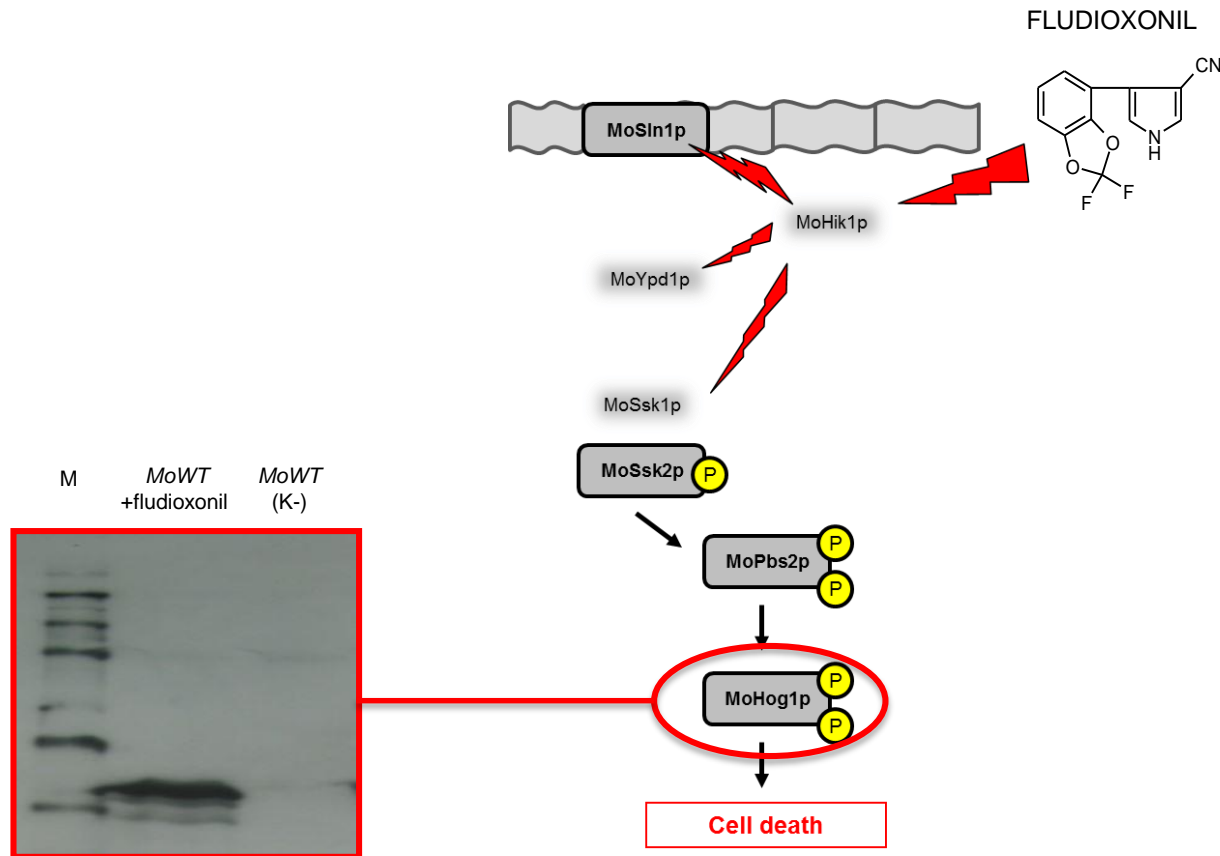
(1) USE OF “LOSS OF FUNCTION” MUTANTS



THE MUTANT STRAINS ARE RESISTANT! AS A CONSEQUENCE WE CAN ASSUME THE TARGET LOCATION IS LOCALIZED WITHIN THE HOG PATHWAY!

A TARGET BASED *IN VIVO* TEST SYSTEM

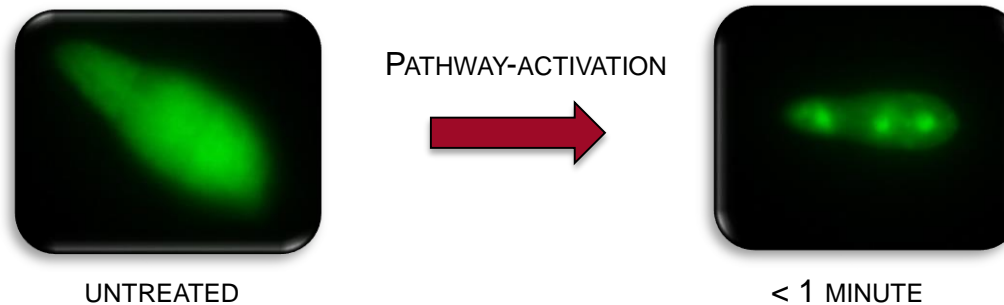
(2) WESTERN BLOTS TO SHOW PATHWAY ACTIVATION



WESTERN BLOTS CAN PROVIDE FURTHER EVIDENCE OF „HOG-ACTIVITY“!

A TARGET BASED *IN VIVO* TEST SYSTEM

(3) GFP-TOOL FOR „REAL-TIME“ VISUALIZATION



**Pest Management
Science**



Research Article | [Full Access](#)

Visualizing fungicide action: an *in vivo* tool for rapid validation of fungicides with target location HOG pathway

Stefan Bohnert , Hendrik Neumann, Eckhard Thines, Stefan Jacob 

THE *MoHOG::GFP*-MUTANT AS EXCELLENT TOOL FOR VISUALIZATION OF „HOG-ACTIVITY“!

SUMMARY

- **PART II: FUNGAL GENETICS**

- “LOSS OF FUNCTION”-MUTANTS
- BIOLOGICAL CHARACTERIZATION OF MUTANT STRAINS
- GFP-TAGGING FOR PATHWAY VISUALIZATION

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ACKNOWLEDGEMENT

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M.SC. STEFAN BOHNERT

imP

Institute of Molecular
Physiology



Mikrobiologie
und Weinforschung


INSTITUT FÜR BIOTECHNOLOGIE
UND WIRKSTOFF-FORSCHUNG



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Carl Zeiss Stiftung



Deutsche
Forschungsgemeinschaft