



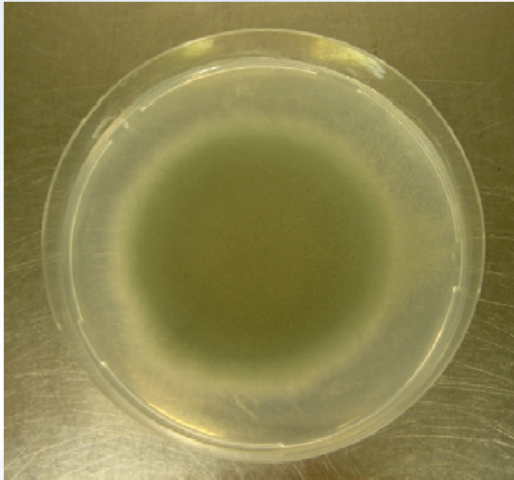
มหาวิทยาลัยมหิดล ปัญญาของแผ่นดิน  
Mahidol University Wisdom of the Land

**Phenotypic analysis of *Aspergillus fumigatus* strain lacking the NAD-dependent formate dehydrogenase encoding gene *fdh***

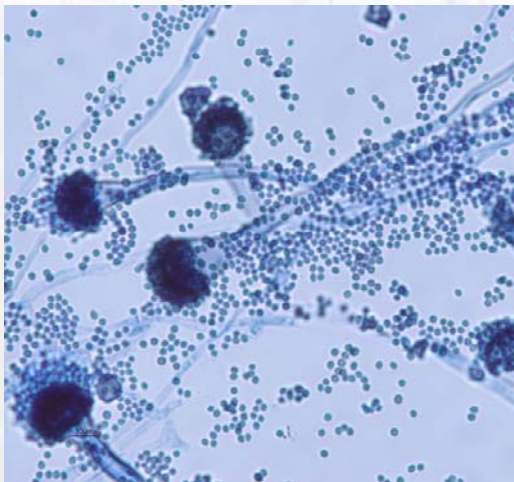
Miss Teerada Ponpinit  
Department of Microbiology  
Faculty of Medicine Siriraj Hospital  
12<sup>th</sup> December 2013



# *Aspergillus fumigatus*

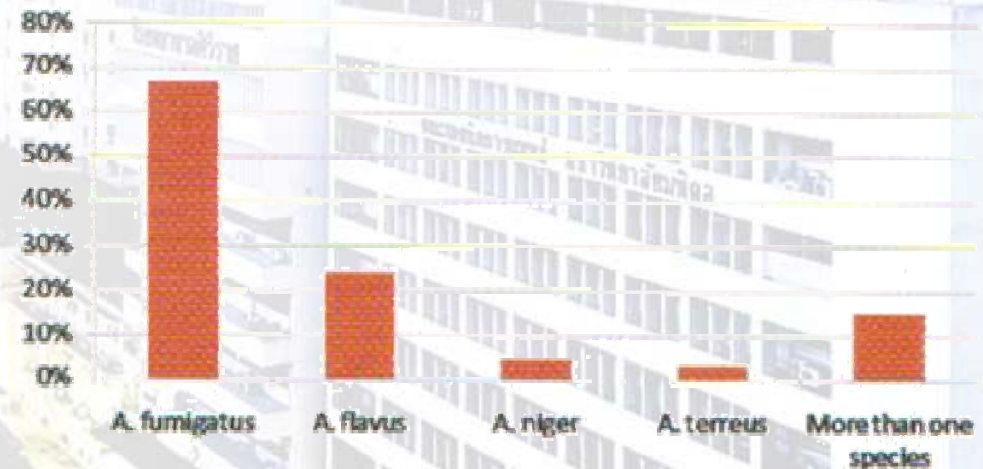


- Phylum Ascomycota
- The most common pathogenic *Aspergillus* species

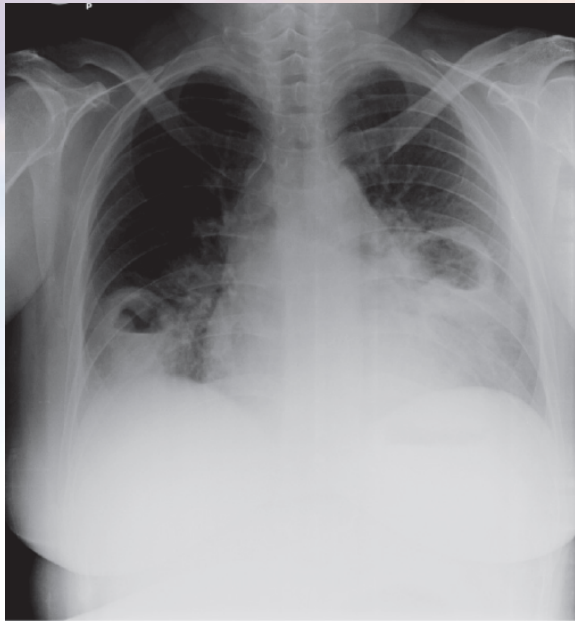


## Epidemiology in Thailand

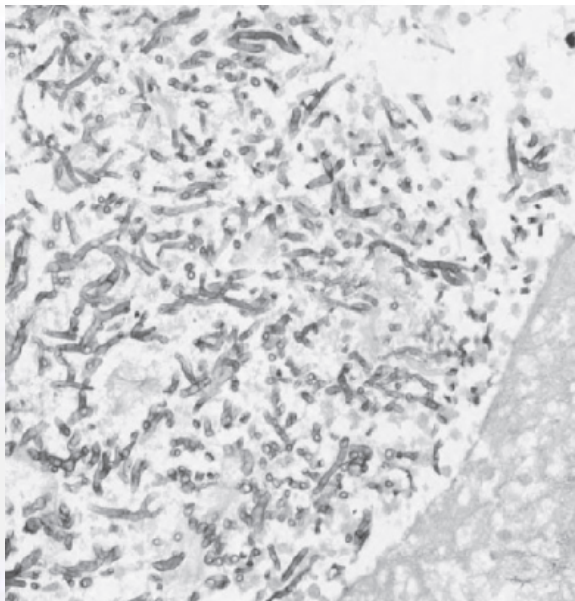
Jan 2000 - Dec 2005 at Ramathibodi Hospital



Kiertiburanakul S et al. *J Med Assoc Thai* 2007;90(5):895-902



- Common cause of invasive pulmonary aspergillosis in immunosuppressed patients
- Mortality rate of 50% to 95%



Steinbach WJ, et al. Journal of Infection. 2012;65(5):453-64.

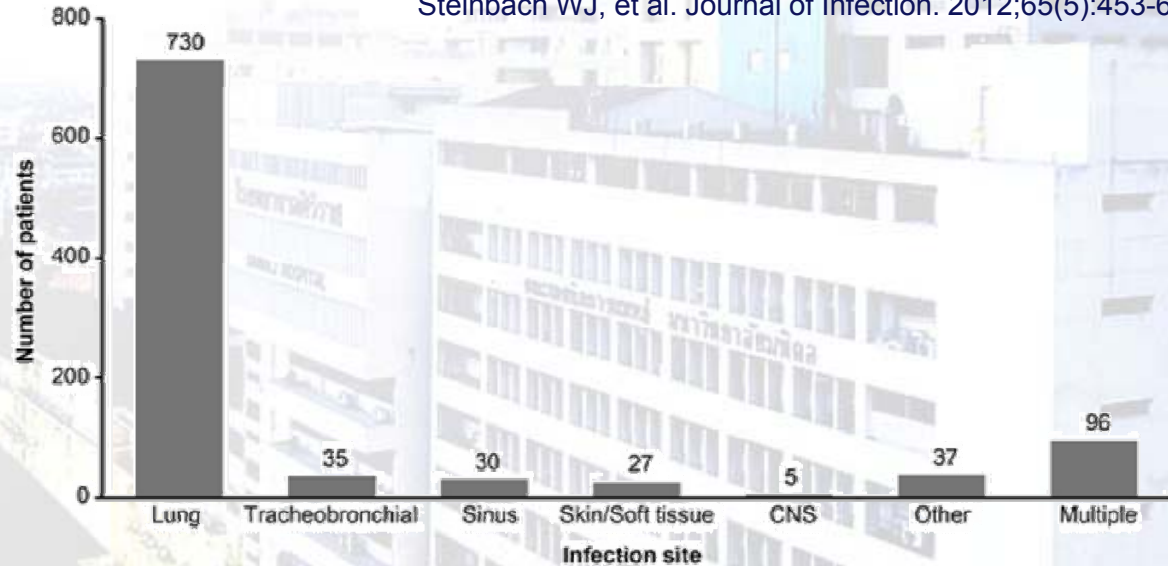
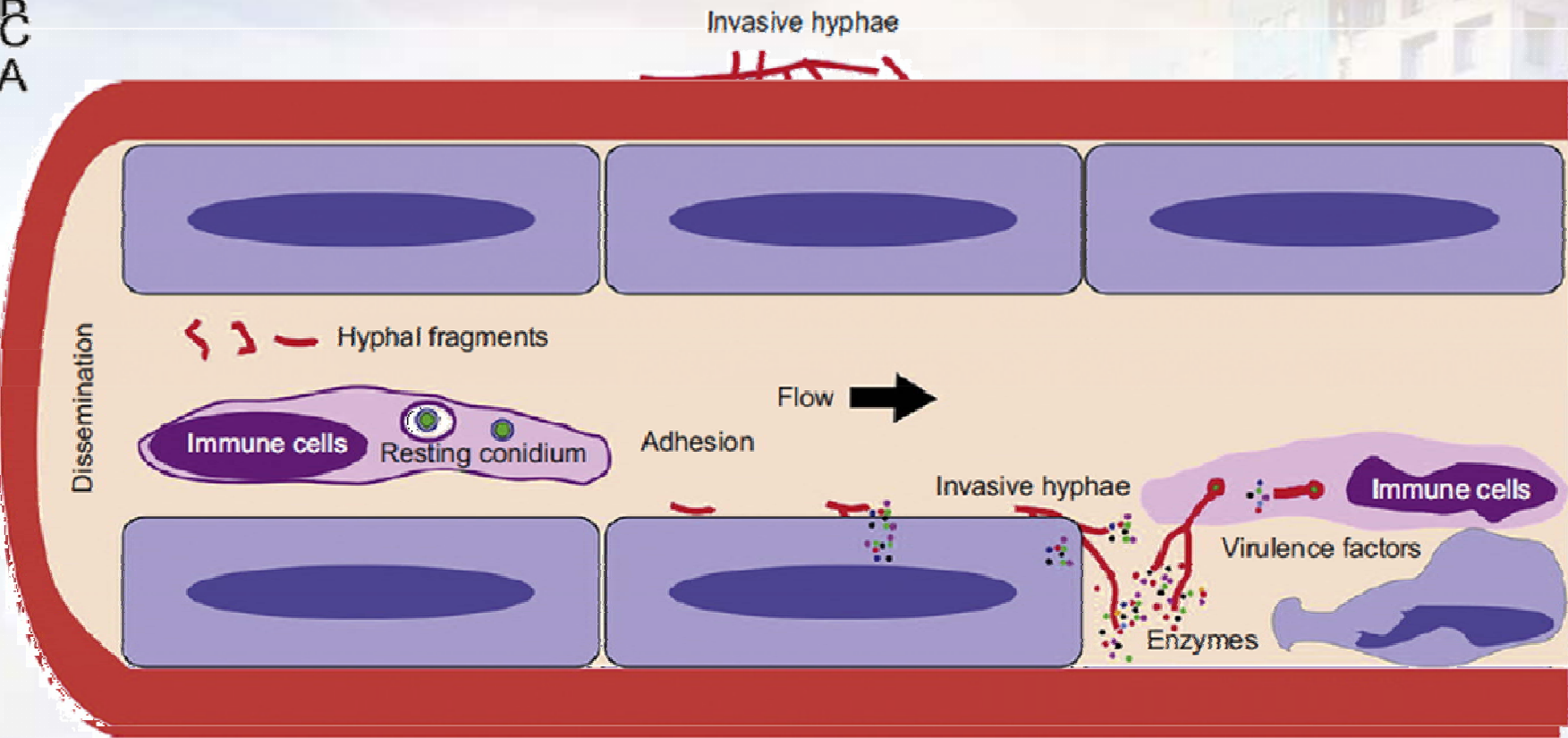


Figure 1 Invasive aspergillosis patients: breakdown by infection site.



# Invasive aspergillosis development

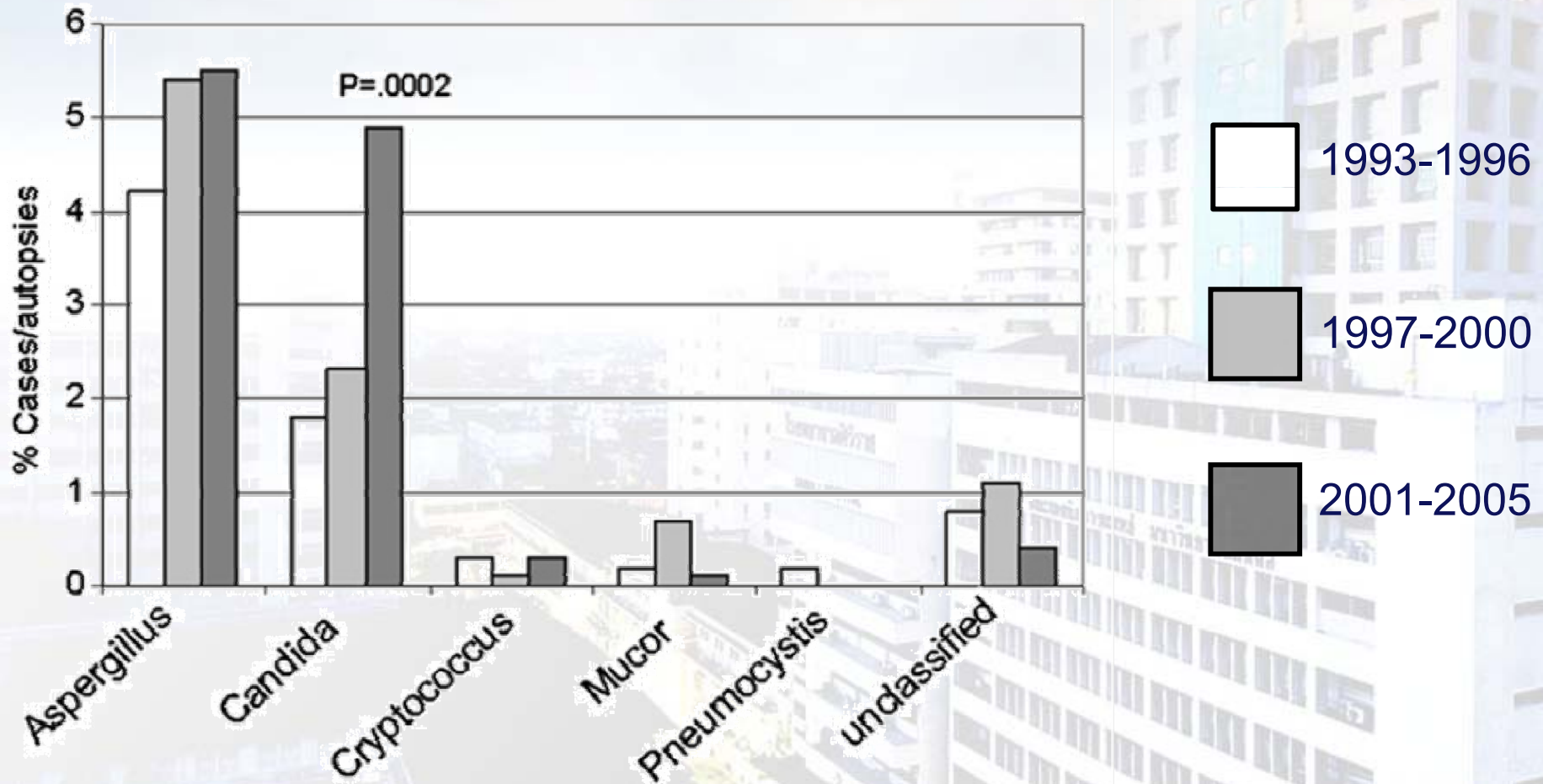
B  
C  
A



Invasion of disseminated hyphal fragments through epithelial cells  
dissemination of hyphal fragments

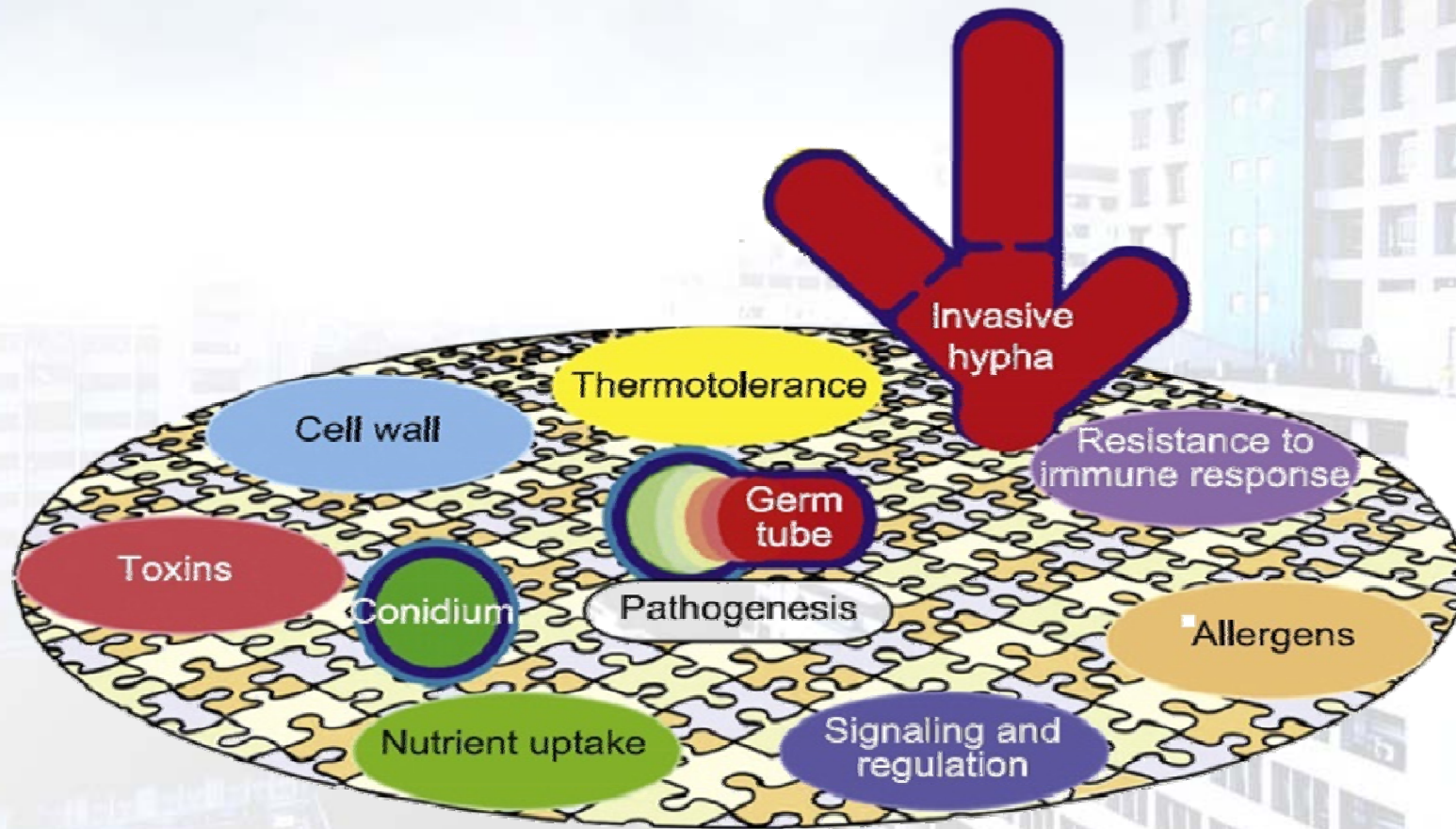


# Prevalence of fungal pathogens



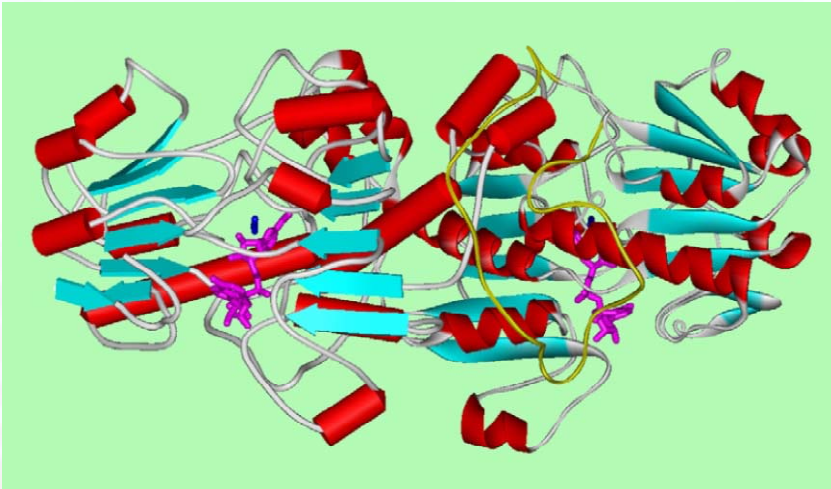


# Factors that cause *A. fumigatus* to be a successful pathogen





# NAD<sup>+</sup>-dependent formate dehydrogenase (NAD<sup>+</sup>-dependent FDH)



- Widespread enzyme in mammals, plants, bacteria, yeasts, and fungi
- Plays a crucial role in respiratory chain and oxidative phosphorylation





```

Q07103      Q07103      -MVKVLIA      6
Q4wDJ0      Q4wDJ0      MVLIRSLSRHLRPPATSFSLTKGTLSPTSSSPFRAASLGGSSISGARTLTASANLQGKVLIM 60
BAA36181    BAA36181    MAAMWRRAAARQLVDR      AVGSRAAHTSAG-SKKIVG      33
P33160      P33160      MAKVLCVLYDDPVDG      YPKTYARDDDLKLDHYGG      34
  
```

```

      : . . * ::* . . **:*::*::** :.* ***:* :* ::::: * :..
Q07103      Q07103      VLYDGGKHGEEVP-ELLGTIQNEIGLRKWLIEDQHTLVTTCKDKGENSTFDKLELDAETI      65
Q4wDJ0      Q4wDJ0      VLYDGGGEHAKQQP-GLLGTTENELGLRWIEEOGHTLVTTSDKDKGENSTFDKLEIVDAEVI 119
BAA36181    BAA36181    VFYQAGEYADKNP-NFVGCVEGALGIRDWLESKGGHHYIVTDDKREGFNSELEKHIEDMHVL 92
P33160      P33160      QTLPTPKAIDFTPGQLLGSVSGELGLRKYLESNGHTLVVTSDKDKGPDVFEREIVDADV      94
  
```

```

      *: ** *.*:*:*: :** . * :***:***:* * :* . :*** ** * :***
Q07103      Q07103      ITTPFHYPGYLTAERLARAKKLELAVTAGIGSDHVDLNAANKTNGGITVAEVTGNSVVSVA 125
Q4wDJ0      Q4wDJ0      ITTPFHYPGYLTAERLAKAKKLELAVTAGVSDHVDLNAANKTNGGITVAEVTGCNVVSVA 179
BAA36181    BAA36181    ITTPFHPAYVTAEKIKKAKTPELILLTAGIGSDHIDLPA--AAAGLTVARVTGSNTVSVA 150
P33160      P33160      ISQPFWPAYLTPERIAKRNKLELAVTAGIGSDHVDLQSE--IDRNVTVAEVTYCNLSIVA 152
  
```

```

      * . :* ** *:*:*:*: :* *:* * . . :*** ** * . :***
Q07103      Q07103      EHVLMTILVLVRFVPAHEQIQEGRDVAAEAANKNEFDLEGKVVGTGVRIGERVLRLR      184
Q4wDJ0      Q4wDJ0      EHVVMTILALVRFVPAHEQIRNGEWDVAAVAKNEFDLENKVVGTVAVGRIGERVLRLR      238
BAA36181    BAA36181    EDELMRILILLRNFLPGYQQVVKGGEWVAGIAHRAVDLEGKTVGTGAGRYGRLLLR      209
P33160      P33160      EHVMMILSLVRNYLPSHEWARKGGWNIADCVSHAYDLEAMHVTVAAGRIGLAVLR      211
  
```

```

      **: : * * * : . * * . : * . *****:* ** * . :*:*
Q07103      Q07103      KPFDCKELLYDYQPLSAEKAEIGORRVADLEEMLAQCDDVVTINCPLEHKTQGLFKNKEL 244
Q4wDJ0      Q4wDJ0      KPFDCKELLYDYQPLRPEVEKEIGCRVENLEEMLAQCDDVVTINCPLEHSTRGLFKNKEL 298
BAA36181    BAA36181    KPFNCN-LLYHDLRQINPELEKEIGAFEEFDLAMLPKCDVVVINTPLTEKTRGMFNKEK 268
P33160      P33160      APFDVH-LHYTDRHRLPESVEKELNLTWHTATREDMYPVCDVVTLNCPLEHETEHEMINDET 270
  
```

```

      : :*:* :***:*** : : * * .***: * .*****:*** *
Q07103      Q07103      ISKMKKGSWLVNTARGAIVVKEDVAEALKSGHLRGYGGDVWFPPQAPQDHPHRYAKNPFG 304
Q4wDJ0      Q4wDJ0      ISKMKKGSWLVNTARGAIVVKEDVAEAVKSGHLRGYGGDVWFPPQAPKDHPLRYVQGPWG 358
BAA36181    BAA36181    IAKMKKGVIIVNNARGAIMDTQAVADACSSGHIAGYGGDVWFPPQAPKDHPRWYMPN--- 324
P33160      P33160      LKLFKRGAYIVNTARGKLCRDAVARALESGRLAGYAGDVWFPPQAPKDHPRWTMPY--- 326
  
```

```

      :.*:***:***: * * ** * * . : : : . * . * * *
Q07103      Q07103      GGNAMVPHMSGTSLDAQRYAAGTRATIESYLSGKHDIYRPEDLIVYGGDYATKSYGERER 364
Q4wDJ0      Q4wDJ0      GGNAMVPHMSGTSLDAQRYAAGTRATILESYSYFSGRHDIYRPEDLIVKDG DYVTKAYGQRQK 417
BAA36181    BAA36181    --HMTPHISGTTIDAQLRYAAGVKDMLDRYFKG-EEFPVENIYVKEGELASQYK----- 376
P33160      P33160    --NGMTPHISGTTILTAQRYAAGTREILECFEFG-RPIRDEYLVIVQGGALAGTG AHSYSK 384
  
```

```

Q07103      Q07103      AKRAAAAAKSA----- 375
Q4wDJ0      Q4wDJ0      -----
BAA36181    BAA36181    -----
P33160      P33160      GNATGGSEEAARFKKAV 401
  
```

- Q07103 = *Neurospora crassa*
- Q4wDJ0 = *Aspergillus fumigatus*
- BAA36181 = *Hordeum vulgare* (Barley)
- P33160 = *Pseudomonas* sp. 101





## Previous studies of NAD<sup>+</sup>-dependent FDH

- In barley, *fdh* was expressed under anaerobic conditions and iron deficiency. (Suzuki K, et al. Plant Physiol. 1998;116:725-32.)
- In potato, FDH transcripts is strongly increased under various stresses including hypoxia, chilling, drought, dark and wounding. (Hourton- Cabassa C, et al. Plant Physiol. 1998;116:627-35.)
- In *Neurospora crassa*, *fdh* mRNA is found only during conidiation and early germination and is not detectable during mycelial growth. (Chow CM, et al. J Bacteriol. 1993;175(12):3703-9.)
- In *A. fumigatus*, the expression of *fdh* also increased significantly 48h after biofilm production. (Bruns S. et al. Proteomics. 2010;10(17):3097-107.)

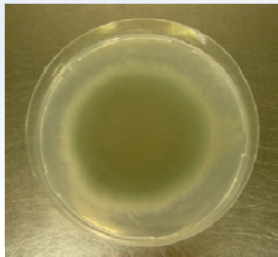


# Objective

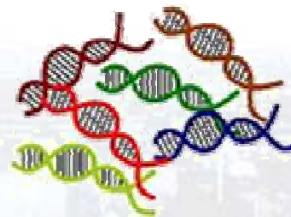
To clarify the physiological role of *fdh* in *Aspergillus fumigatus*



# Generation of $\Delta fdh$ strain



*A. fumigatus* WT strain AF293

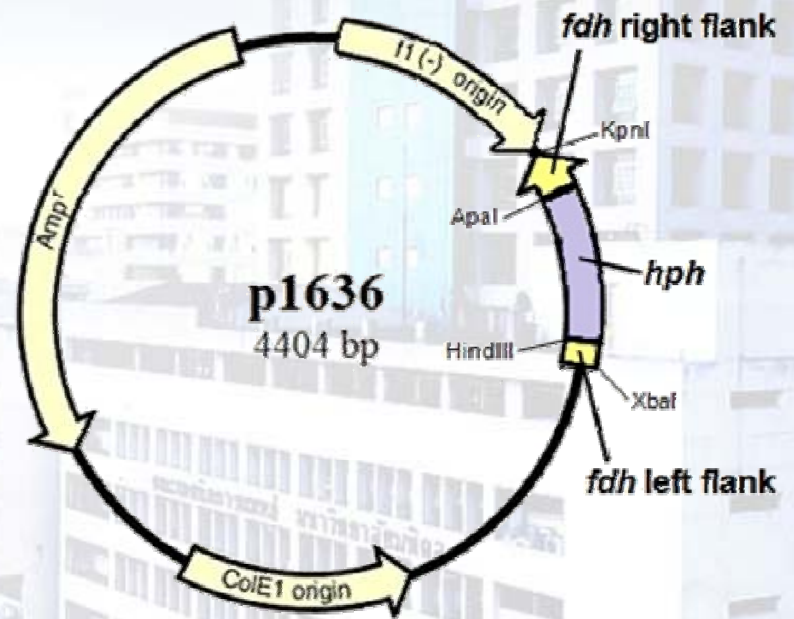


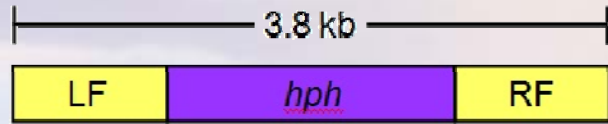
Extraction



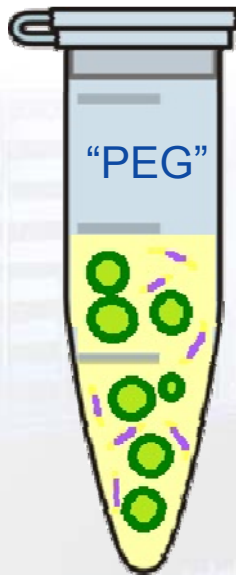
PCR

*fdh* gene (AFUA\_6G04920)

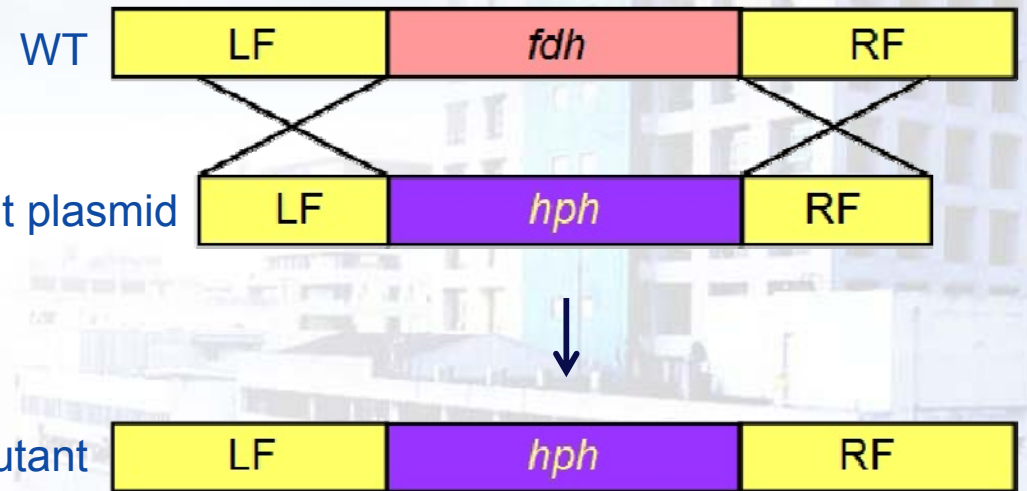




Final product



Protoplast transformation



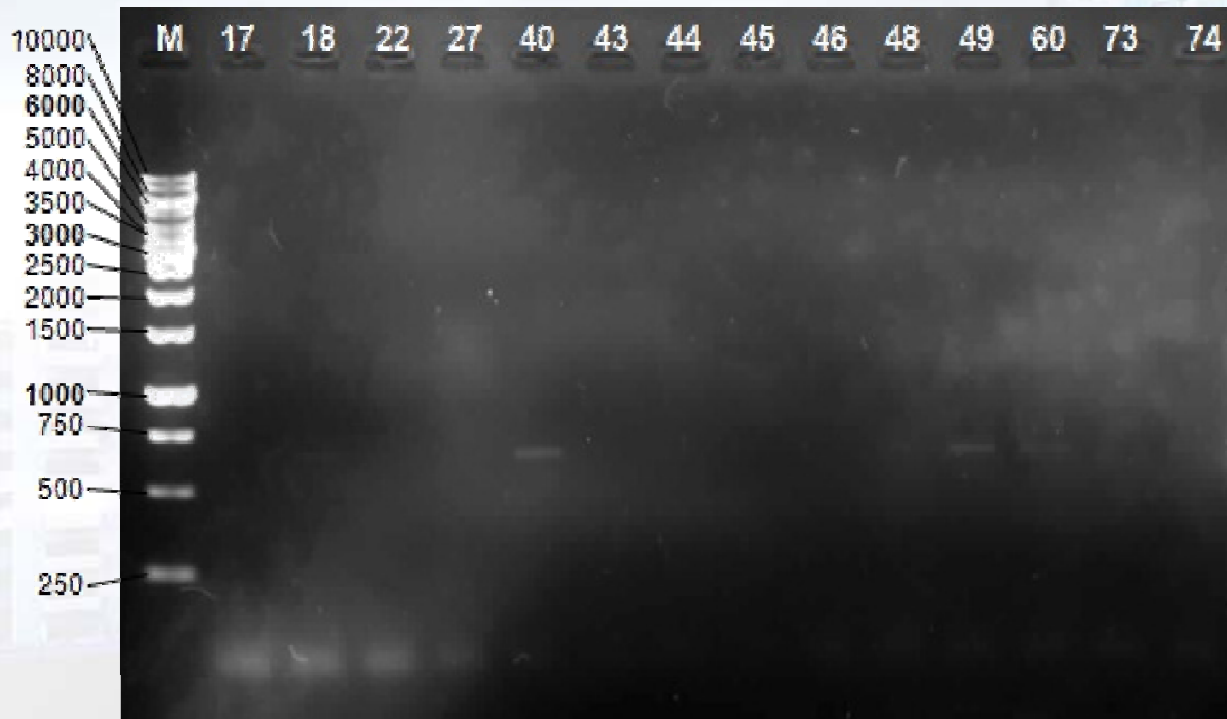
Knock-out plasmid

Knock-out mutant

Homologous recombination



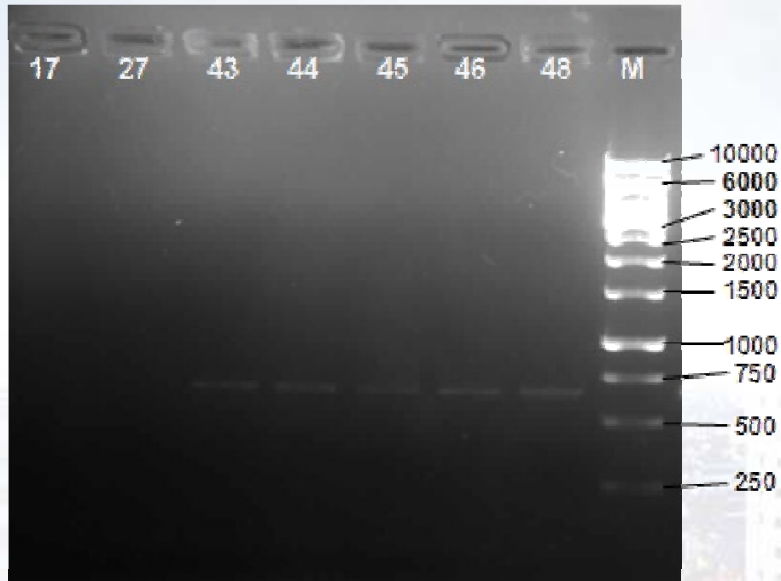
# Screening and confirmation of $\Delta fdh$ strain



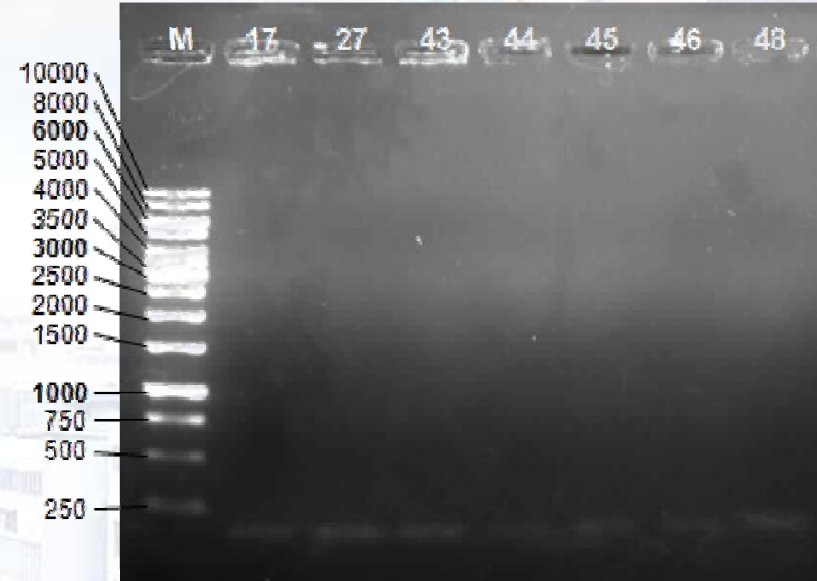
**Colony PCR (*A. fumigatus*)**



# PCR



*fdh* ORF

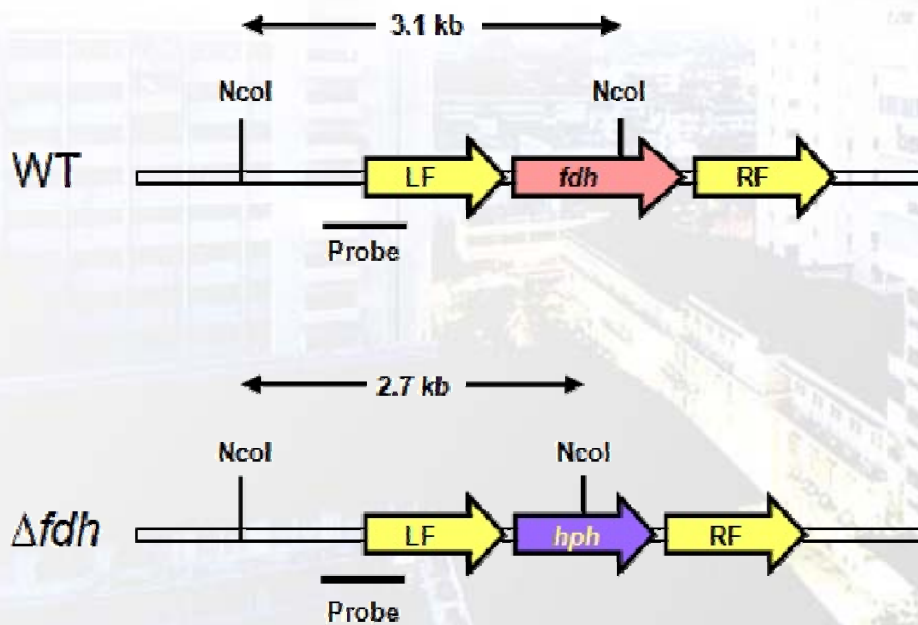
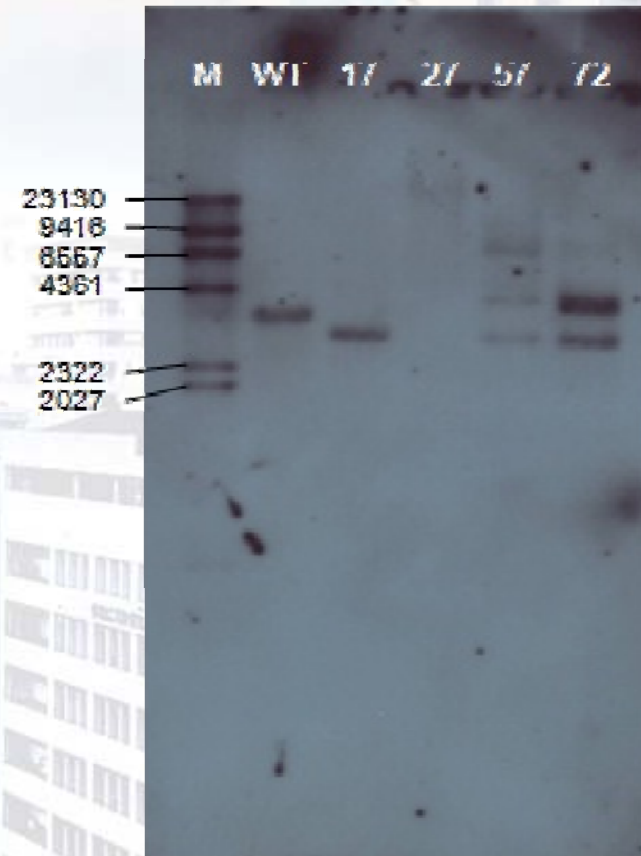
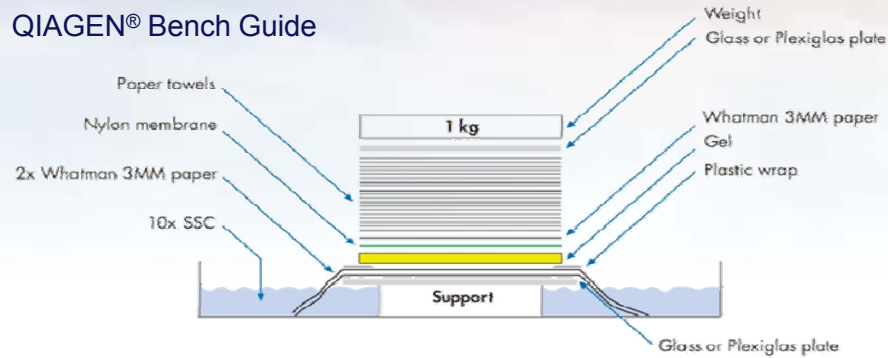


Tubulin gene



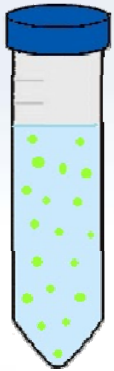
# Southern blot analysis

QIAGEN® Bench Guide





# Radial hyphal growth analysis



$10^6$  conidial/ml of  
WT or  
 $\Delta fdh$  strains



200 mM  
 $\text{CaCl}_2$

0.8 M  
NaCl

1 M  
Sorbital

200 mM  
 $\text{MgCl}_2$

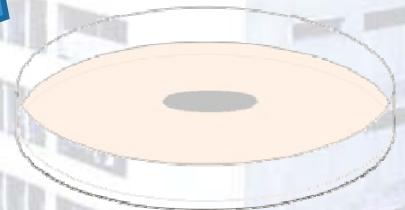
0.6 M  
KCl

10 mM  
 $\text{MnCl}_2$

GMM  
pH4

GMM

GMM  
pH9

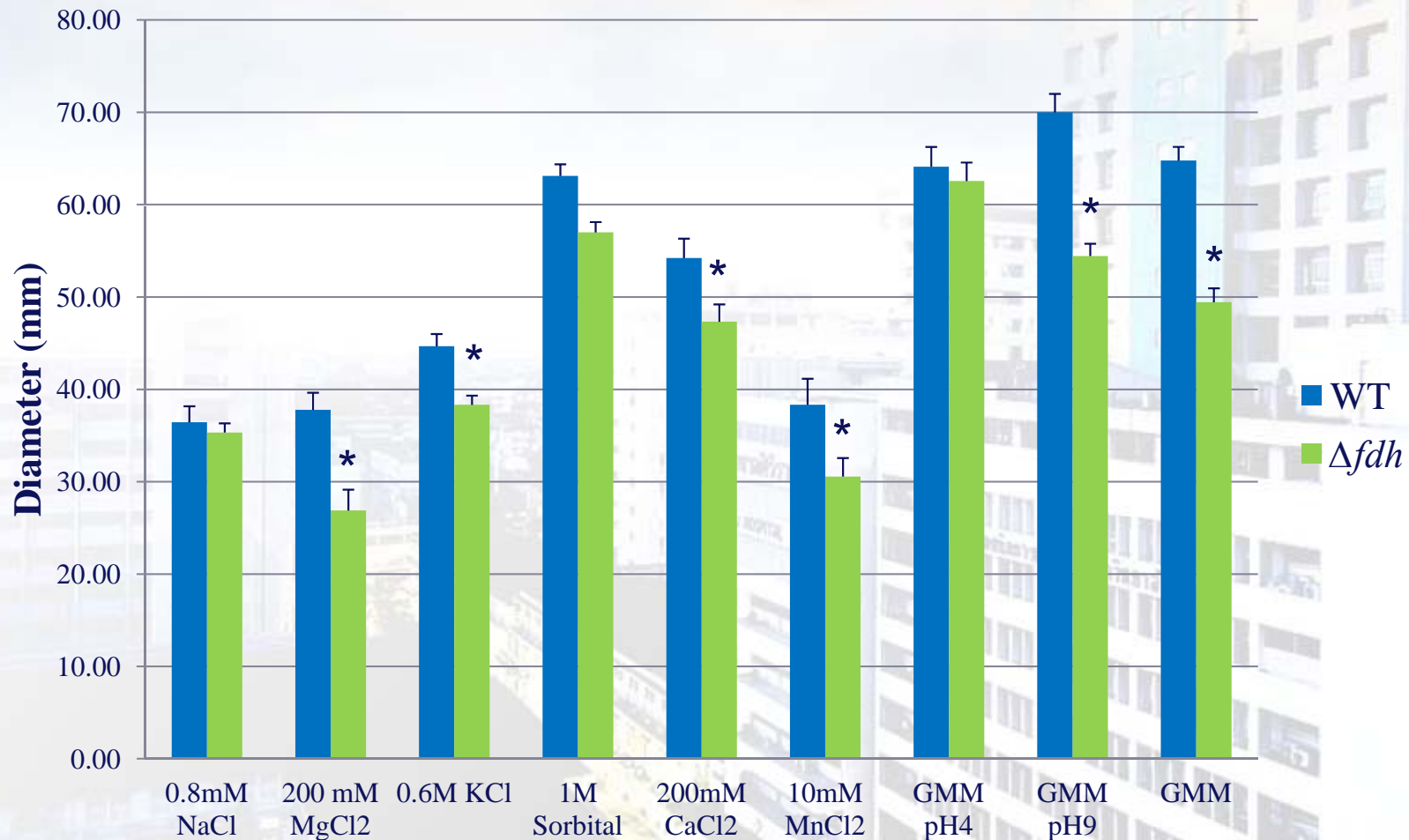


Every 24 h for  
5 days





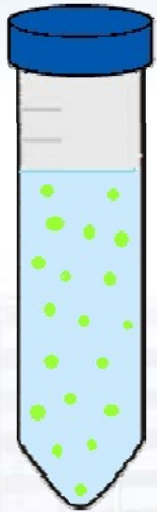
## Radial hyphal growth of wild-type and $\Delta fdh$ under various unfavorable conditions



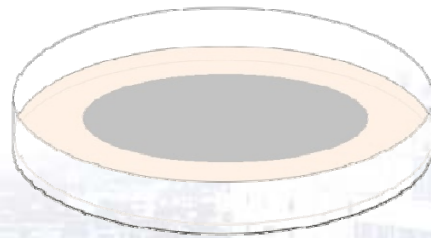
A p-value  $\leq 0.05$  was considered statistically significant.



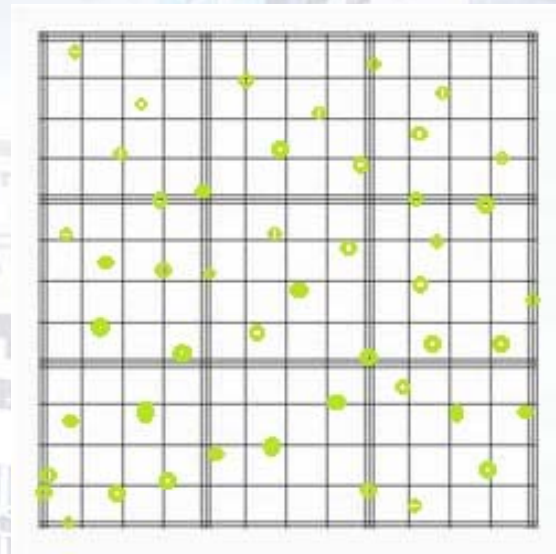
# Conidiation analysis



$10^6$  conidial/ml of  
WT or  
 $\Delta fdh$  strains

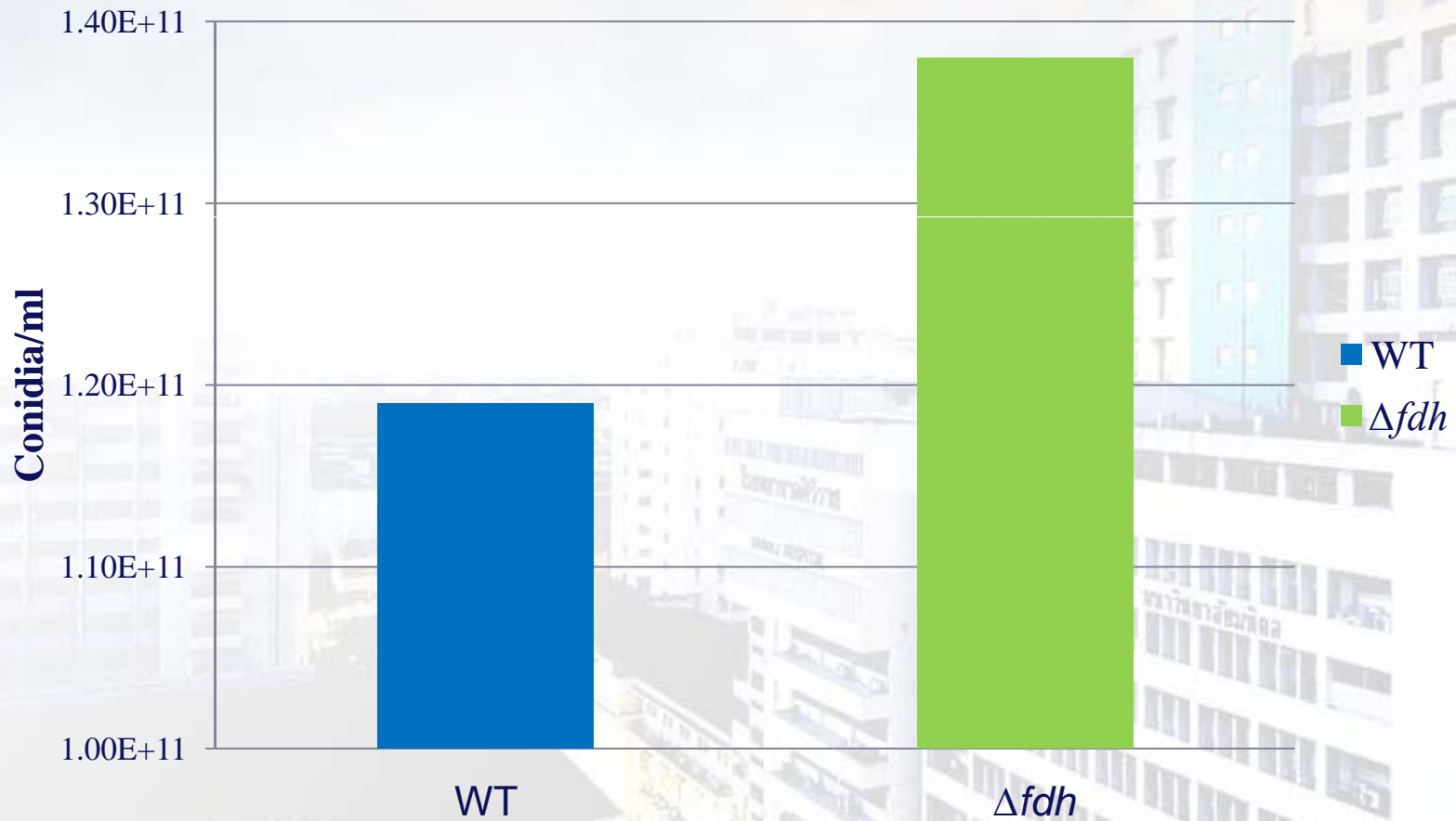


GMM agar, 37°C  
for 72 h



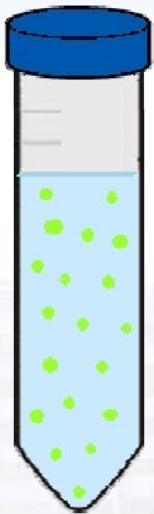


## Conidiation of wild-type and $\Delta fdh$





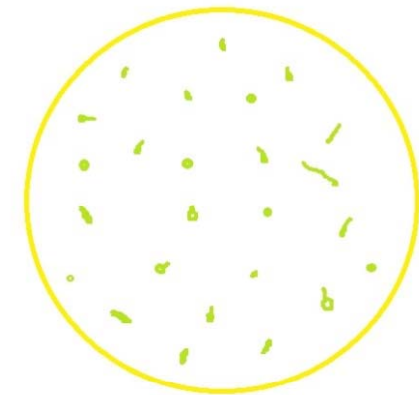
# Germination analysis



$10^7$  conidia/ml of  
WT or  
 $\Delta fdh$  strains



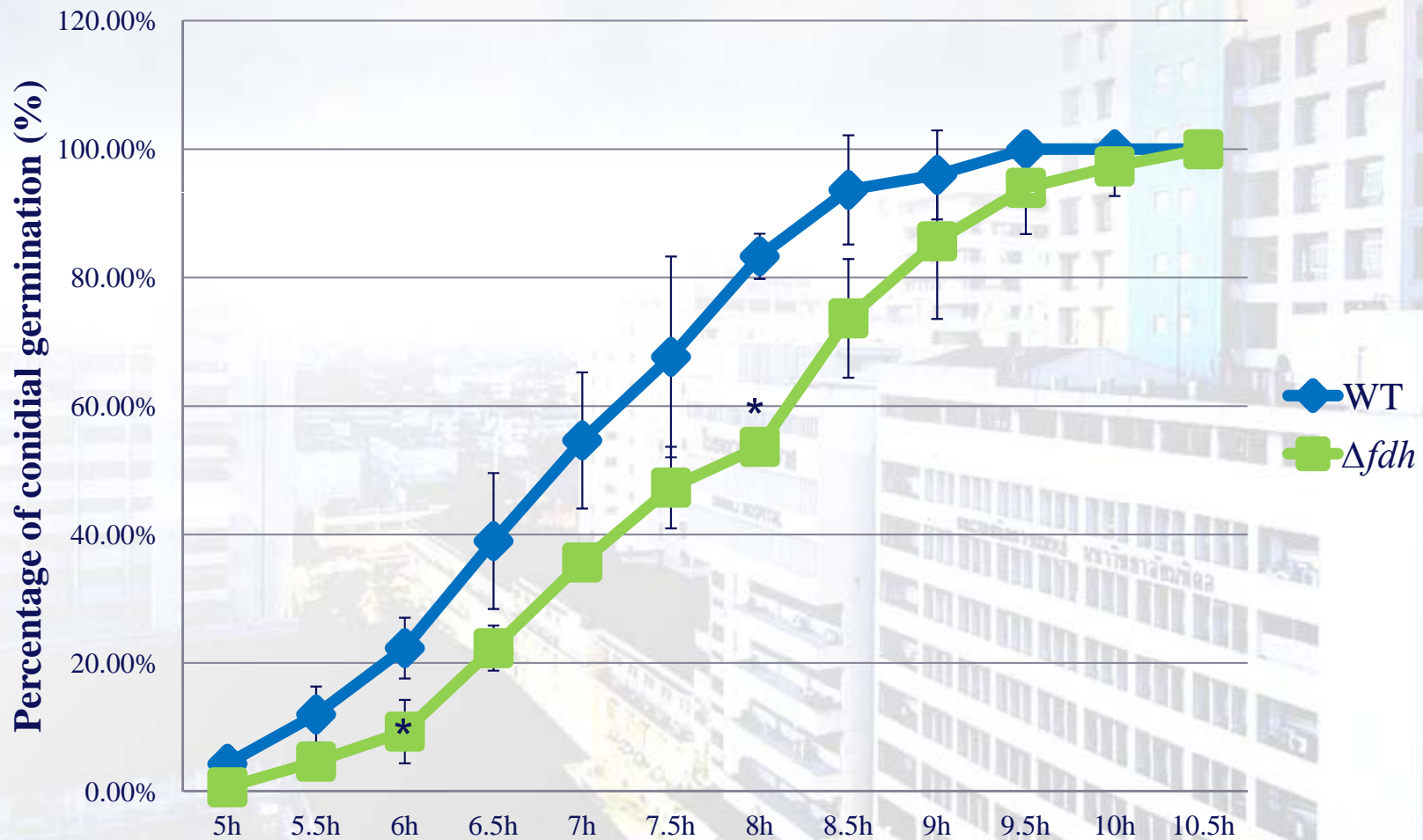
37°C, 300 rpm  
for 5 h



Determined every 30 min  
until 100% germination



## Percentage of germination of wild-type and $\Delta fdh$ over the experimental time period



A p-value  $\leq 0.05$  was considered statistically significant.



# References

1. Kiertiburanakul S, Thibbadee C, Santanirand P. Invasive Aspergillosis in a Tertiary-Care Hospital in Thailand. *J Med Assoc Thai.* 2007;90(5):895-902.
2. Rittayamai N, Tscheikuna J. Invasive Pulmonary Aspergillosis with Cerebral Dissemination in Chronic Refractory Idiopathic Thrombocytopenic Purpura. *Thai journal of tuberculosis, chest disease and critical care.* [Case report]. 2009;30(3):166-71.
3. Steinbach WJ, Marr KA, Anaissie EJ, Azie N, Quan S-P, Meier-Kriesche H-U, et al. Clinical epidemiology of 960 patients with invasive aspergillosis from the PATH Alliance registry. *Journal of Infection.* 2012;65(5):453-64.
4. Abad A, Victoria Fernández-Molina J, Bikandi J, Ramírez A, Margareto J, Sendino J, et al. What makes *Aspergillus fumigatus* a successful pathogen? Genes and molecules involved in invasive aspergillosis. *Revista Iberoamericana de Micología.* 2010;27(4):155-82
5. Lehrnbecher T, Frank C, Engels K, Kriener S, Groll AH, Schwabe D. Trends in the postmortem epidemiology of invasive fungal infections at a university hospital. *Journal of Infection.* 2010;61:259-65.



## References

6. Suzuki K, Itai R, Nakanishi K, Nishizawa NK, Yoshimura E, Mori S. Formate Dehydrogenase, an Enzyme of Anaerobic Metabolism, Is Induced by Iron Deficiency in Barley Roots. *Plant Physiol.* 1998;116:725-32.
7. Hourton- Cabassa C, Ambard-Bretteville F, Morean F, Davy de Virville J, Rémy R, Colas des Francs-Small C. Stress Induction of Mitochondrial Formate Dehydrogenase in Potato Leaves. *Plant Physiol.* 1998;116:627-35.
8. Chow CM, RajBhandary UL. Developmental regulation of the gene for formate dehydrogenase in *Neurospora crassa*. *J Bacteriol.* 1993;175(12):3703-9.
9. Schmit JC, Brody S. Biochemical Genetics of *Neurospora crassa* Conidial Germination. *Bacteriol Rev.* 1976;40(1):1-41.
10. Bruns S, Seidler M, Albrecht D, Salvenmoser S, Remme N, Hertweck C, et al. Functional genomic profiling of *Aspergillus fumigatus* biofilm reveals enhanced production of the mycotoxin gliotoxin. *Proteomics.* 2010;10(17):3097-107.



# Acknowledgement

Dr.Nadthanan Pinchai



Siriraj Graduate Thesis Scholarship

Department of Microbiology, Faculty of Medicine  
Siriraj Hospital, Mahidol University







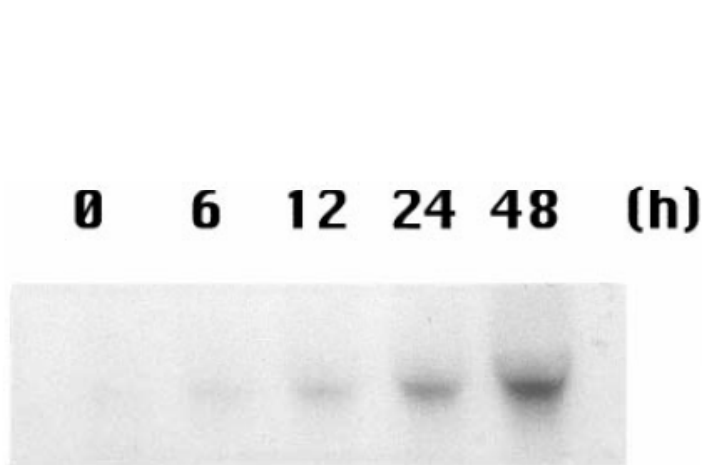
**Thank you for your attention**



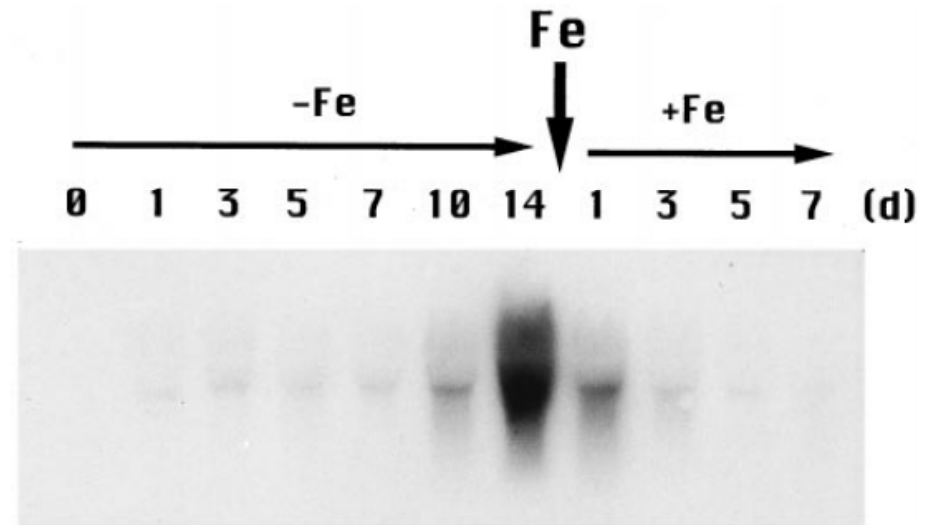
Plant Physiol. (1998) 116: 725-732

## Formate Dehydrogenase, an Enzyme of Anaerobic Metabolism, Is Induced by Iron Deficiency in Barley Roots<sup>1</sup>

Kazuya Suzuki, Reiko Itai, Koichiro Suzuki, Hiromi Nakanishi, Naoko-Kishi Nishizawa, Etsuro Yoshimura, and Satoshi Mori\*



**Figure 8.** Expression of *Fdh* under anaerobic conditions. Each lane was loaded with 10  $\mu$ g of RNA. Total RNA was isolated after 0, 6, 12, 24, and 48 h of anaerobic treatment.



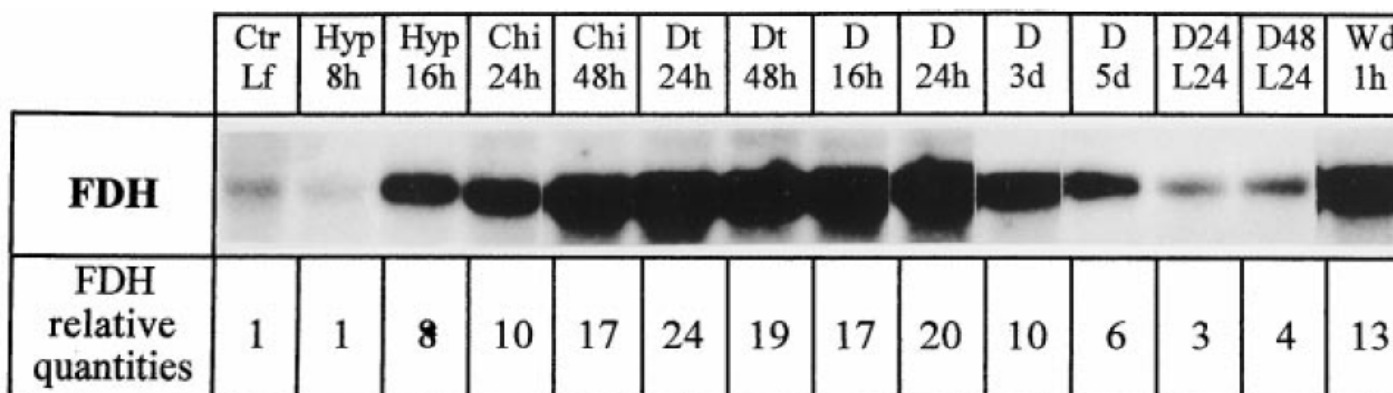
**Figure 7.** Expression of *Fdh* during Fe deficiency in barley roots. Each lane was loaded with 10  $\mu$ g of RNA. Total RNA was isolated after 0, 1, 3, 5, 7, 10, and 14 d of Fe deficiency and 1, 3, 5, and 7 d after Fe resupply.



Plant Physiol. (1998) 116: 627–635

## Stress Induction of Mitochondrial Formate Dehydrogenase in Potato Leaves<sup>1</sup>

Cécile Hourton-Cabassa, Françoise Ambard-Bretteville, François Moreau, Jacques Davy de Virville, René Rémy, and Catherine Colas des Francs-Small\*



**Figure 4.** RNA expression of FDH and SHMT in potato leaves under various stress treatments. Each lane was loaded with 10  $\mu\text{g}$  of RNA. Ctr Lf, Control leaves; Hyp 8h and Hyp 16h, 8 and 16 h of hypoxia, respectively; Chi 24h and Chi 48h, 24 and 48 h of chilling, respectively; Dt 24h and Dt 48h, 24 and 48 h of drought treatment, respectively; D 16h, D 24h, D 3d, and D 5d, 16 h, 24 h, 3 d, and 5 d of dark treatment, respectively; D24 and L24, 24 h of dark treatment followed by 24 h under a 16-h photoperiod; D48 and L24, 48 h of dark treatment followed by 24 h under a 16-h photoperiod; Wd 1h, 1 h after wounding. The autoradiographs were scanned using the control leaf sample as the reference for both probes, and the figures were corrected according to the RNA loadings.



JOURNAL OF BACTERIOLOGY, June 1993, p. 3703-3709  
0021-9193/93/123703-07\$02.00/0  
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Vol. 175, No. 12

## Developmental Regulation of the Gene for Formate Dehydrogenase in *Neurospora crassa*

C. MING CHOW AND UTTAM L. RAJBHANDARY\*

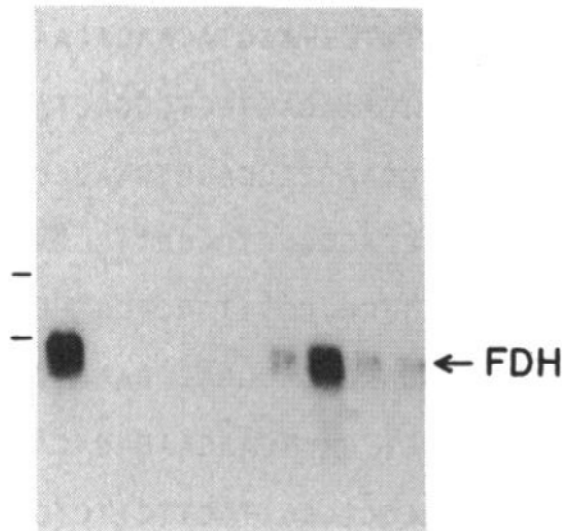


FIG. 2. Northern blot analysis of *N. crassa* total RNA. Membranes were probed for mRNA for FDH



RESEARCH ARTICLE

# Functional genomic profiling of *Aspergillus fumigatus* biofilm reveals enhanced production of the mycotoxin gliotoxin

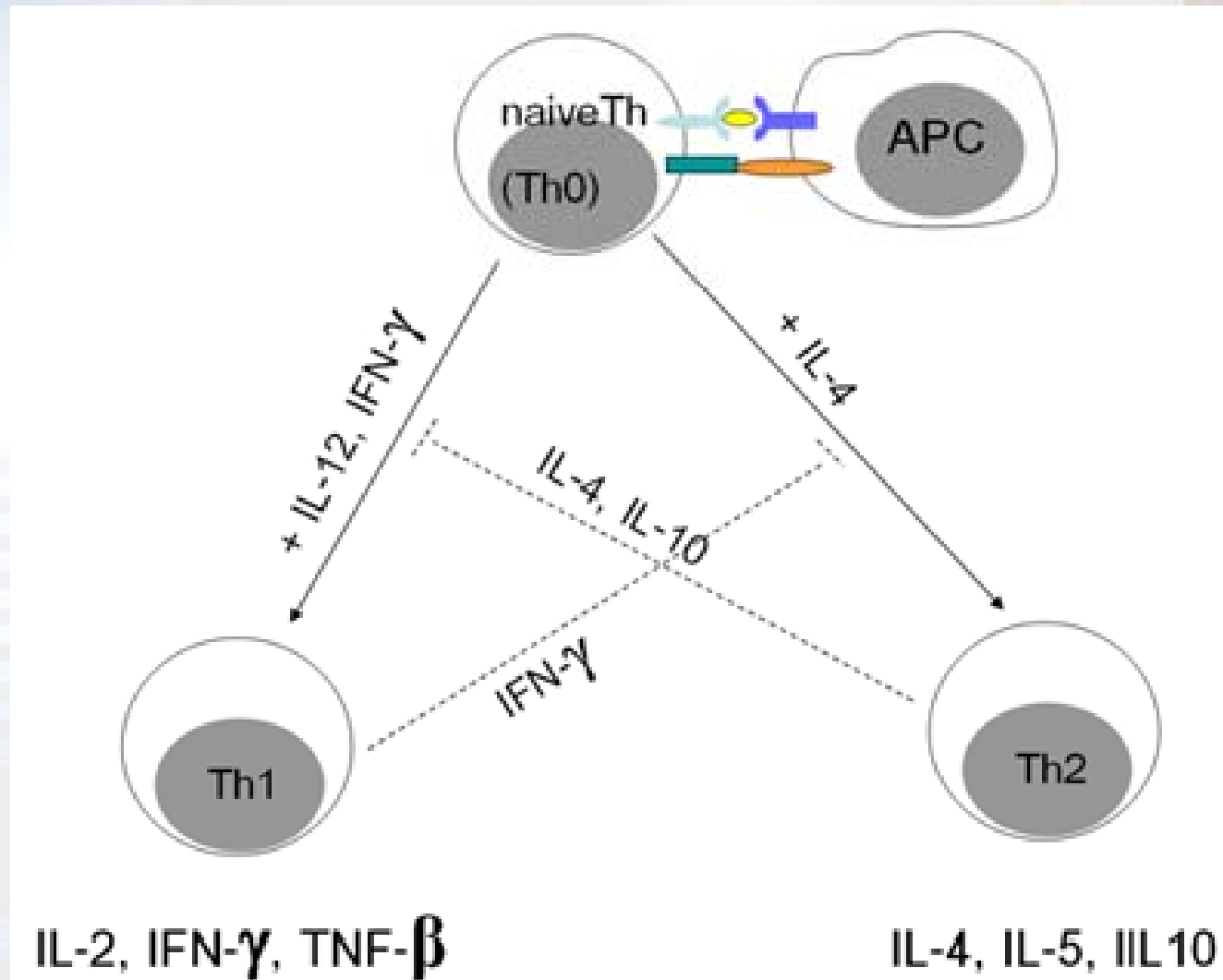
Sandra Bruns<sup>1,2\*</sup>, Marc Seidler<sup>3\*</sup>, Daniela Albrecht<sup>4</sup>, Stefanie Salvenmoser<sup>3</sup>, Nicole Remme<sup>5</sup>, Christian Hertweck<sup>5</sup>, Axel A. Brakhage<sup>1,2</sup>, Olaf Kniemeyer<sup>1,2\*\*</sup> and Frank-Michael C. Müller<sup>3</sup>

**Table 1.** Biofilm- (biofilm *versus* planktonic grown) and time-dependent (48 *versus* 24 h), differentially synthesized proteins of *A. fumigatus*

Putative function	Fold change <sup>a)</sup>			
	Biofilm-dependent changes		Time-dependent changes in biofilm	
	24+/24–	48+/48–	48+/24+	48–/24–
<b>Formate catabolic process</b>				
803 NAD-dependent formate DH (AFUA_6G04920)	n.s.	n.s.	13.0	2.7
807 “–”	n.s.	n.s.	11.4	3.1
814 “–”	n.s.	n.s.	34.0	2.9
815 “–”	n.s.	n.s.	20.7	2.5
817 “–”	n.s.	n.s.	3.2	0.9

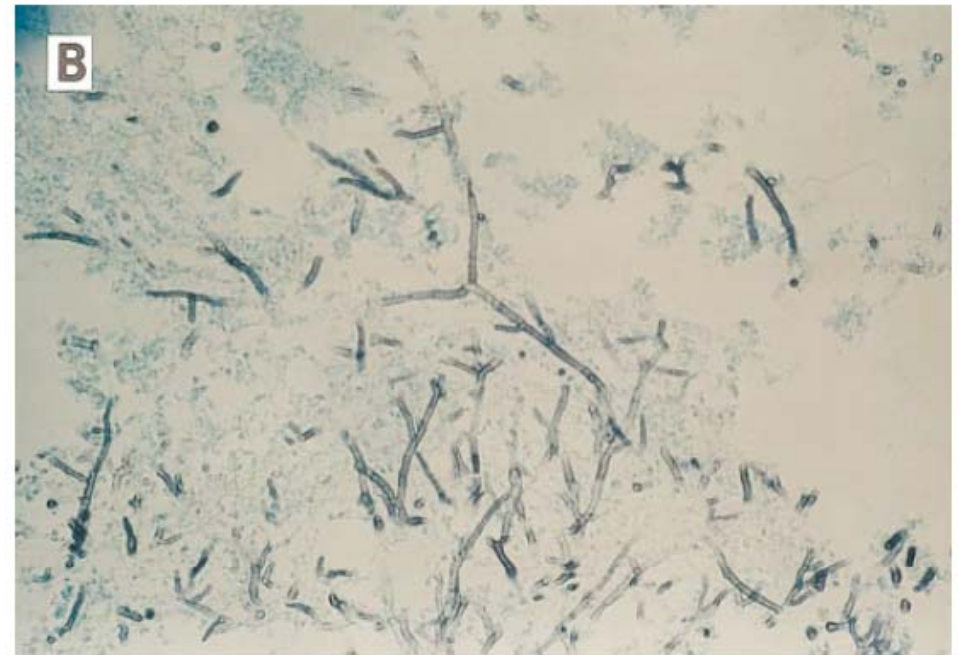
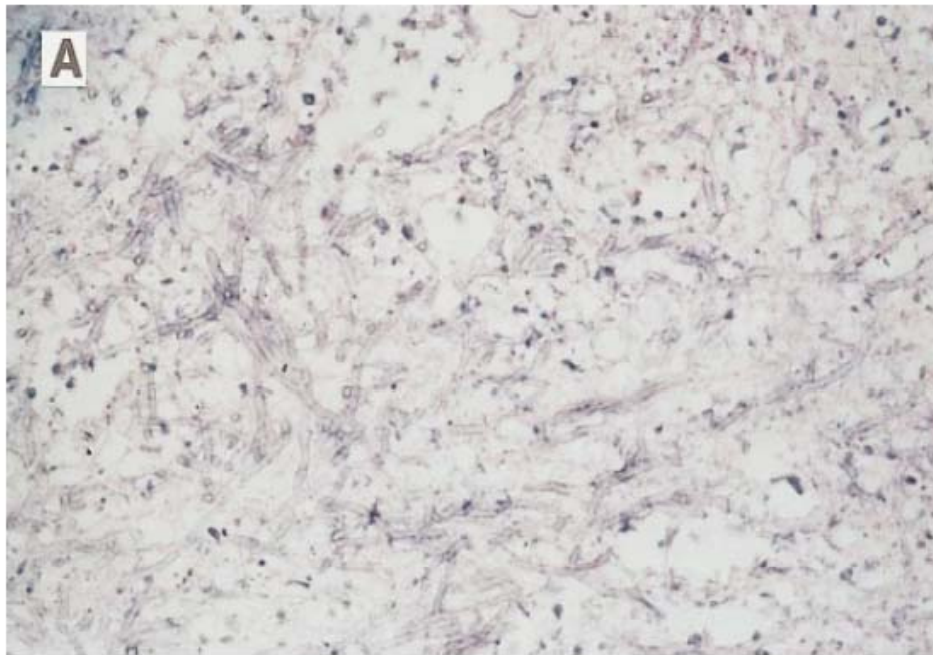


# Th1/Th2 cytokines





## Characteristic pattern of *A. fumigatus*



**Figure 2.** *A*, histological appearance under high power of an aspergillus brain abscess showing an inflammatory cell infiltrate surrounding *Aspergillus* hyphae. (Stain, hematoxylin-eosin; original magnification,  $\times 400$ .) *B*, Section of the same abscess showing the characteristic radiating pattern of *Aspergillus* hyphae with septae and branching at  $45^\circ$  angles. (Stain, methenamine silver; original magnification,  $\times 400$ .)



# Chest radiograph of AIDS patient with invasive pulmonary aspergillosis

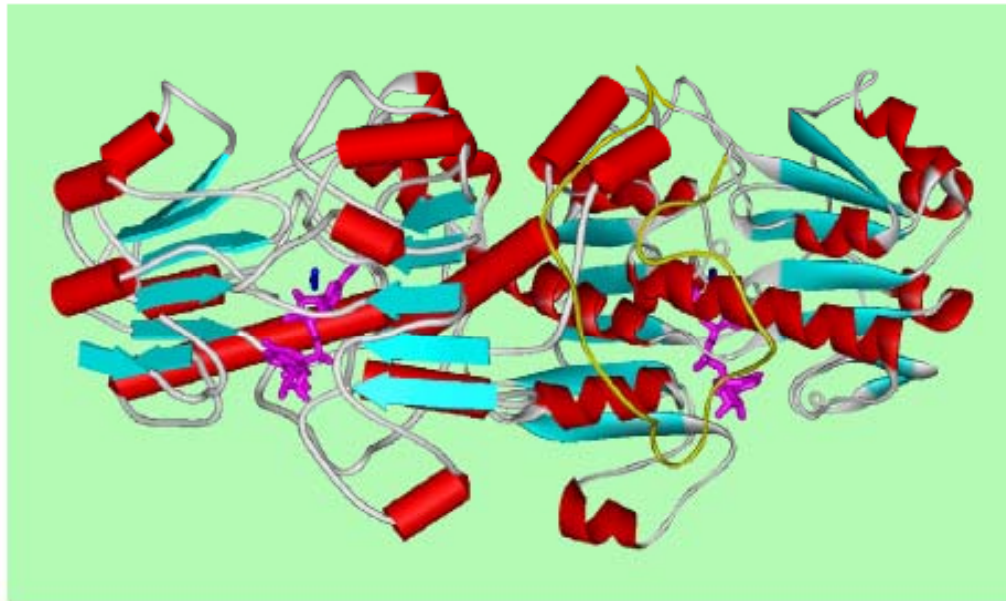


**Figure 3.** Chest radiograph of a patient with late-stage AIDS and probable invasive pulmonary aspergillosis, illustrating diffuse bilateral disease. The patient had cytomegalovirus retinitis and was receiving chemotherapy for Kaposi's sarcoma; he was symptomatic with a high fever and cough, and cultures of both sputum and bronchoalveolar lavage fluid yielded *Aspergillus fumigatus*.





# Structure of NAD<sup>+</sup>-dependent FDH



A



B

**Figure 2.** Structure of PseFDH. A) FDH ternary complex with NAD<sup>+</sup> (magenta) and formate (blue) occupying azide binding site.  $\alpha$ -helices are depicted as red cylinders (left subunit) or helices (right subunit) while  $\beta$ -strands as cyan arrows (left) or strips (right). A long loop comprising a/a residues 12-47 present in bacterial FDHs but absent in the enzymes from other species is shown in yellow. B) Representation of the structure of the FDH subunit. Numbering of structural elements from [17].