

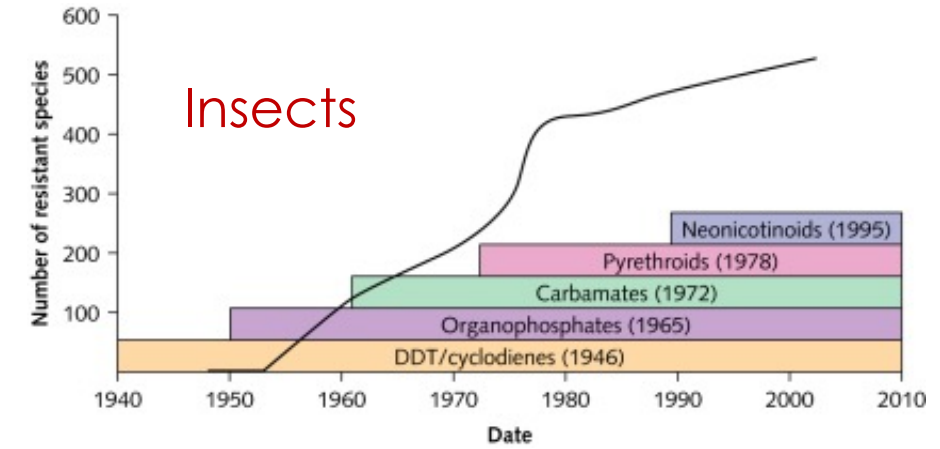
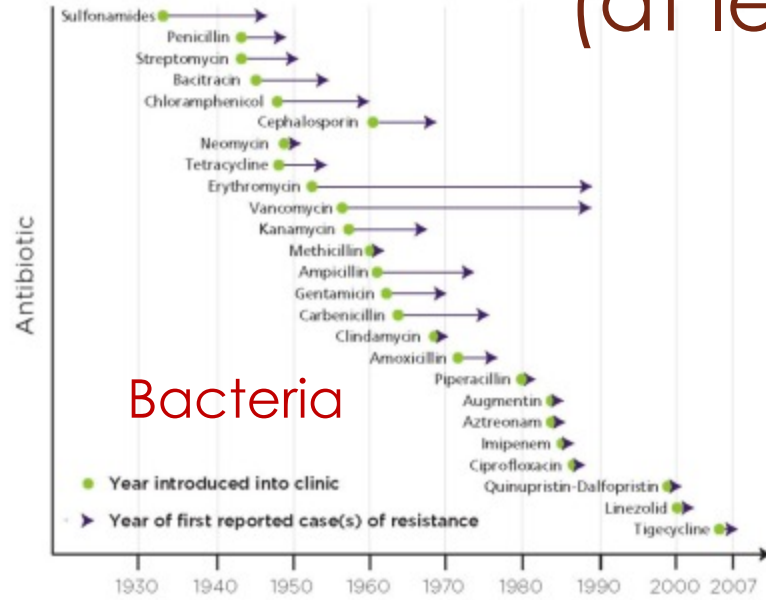
# NWO Groen III – One-Health Aspects of circularity

Monday the 27<sup>th</sup> of June 2022

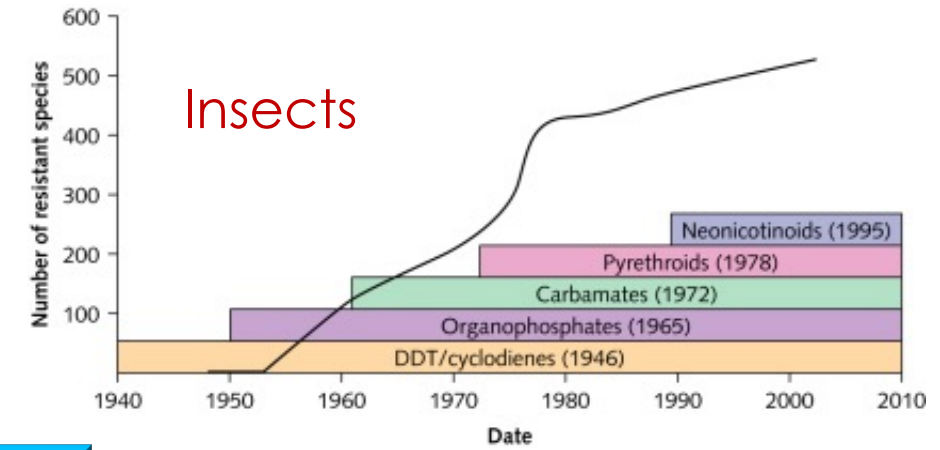
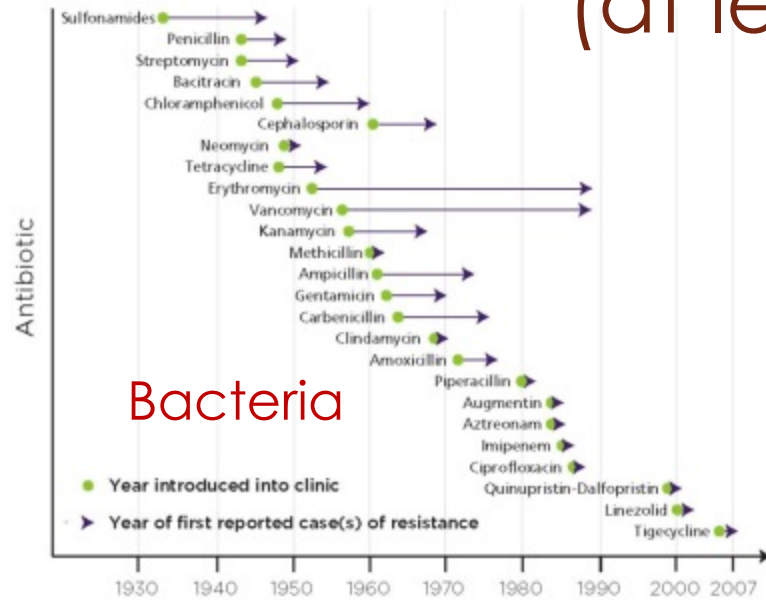
Laboratory of Genetics, Wageningen University & Research  
Radboud University Medical Center, Nijmegen  
+ PARTNERS!



# Chemical resistance selection; it will happen (at least with current practices)



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1998  
34 bp  
L98H

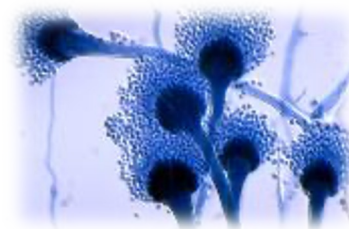
2006  
53 bp  
-

2009  
46 bp  
Y121F/T289A

2012  
92 bp  
Y121F/M172Y/T289A/G448S

**Fungi**

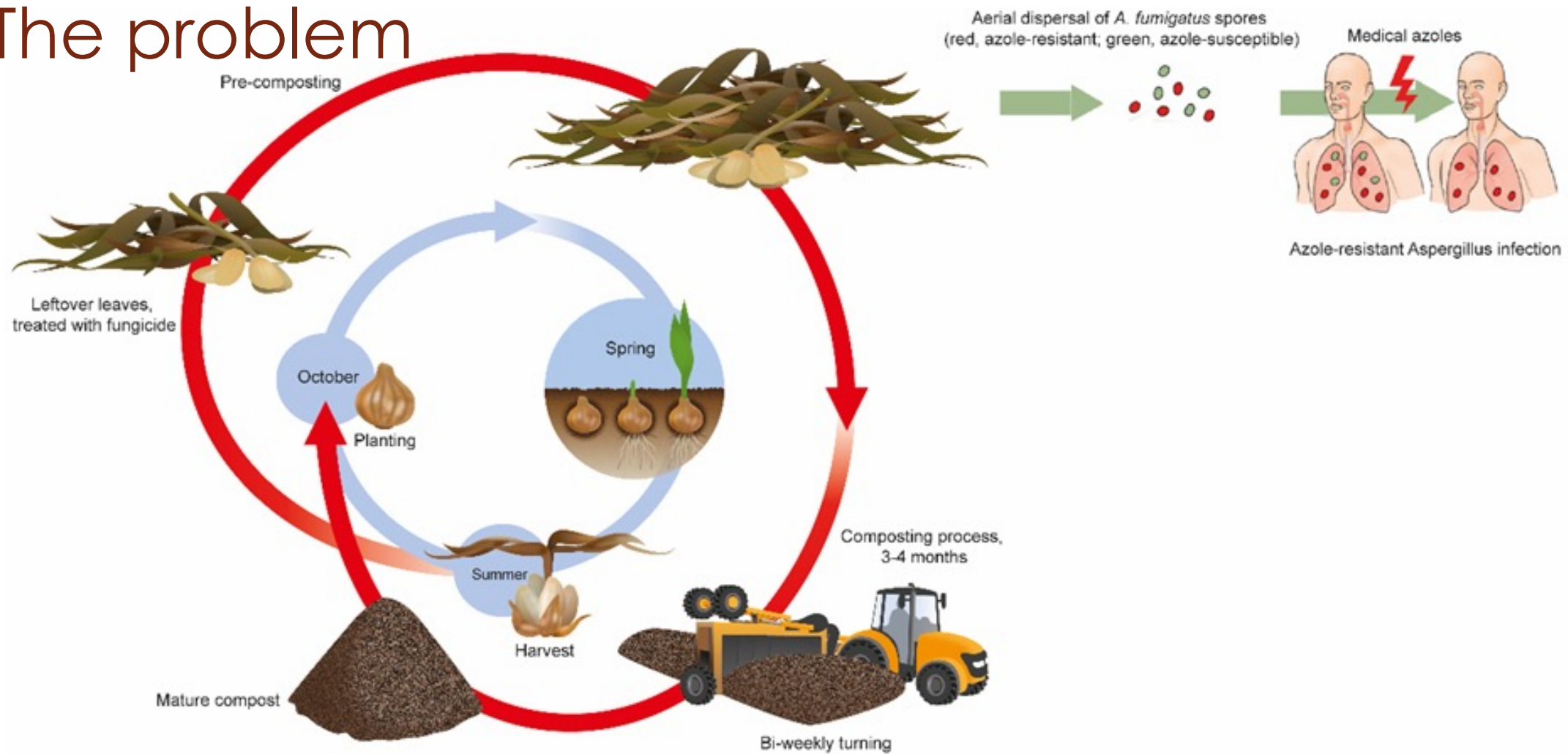
Resistance of the fungus  
*Aspergillus fumigatus*  
against fungicide of the  
azole-class



*Aspergillus fumigatus*

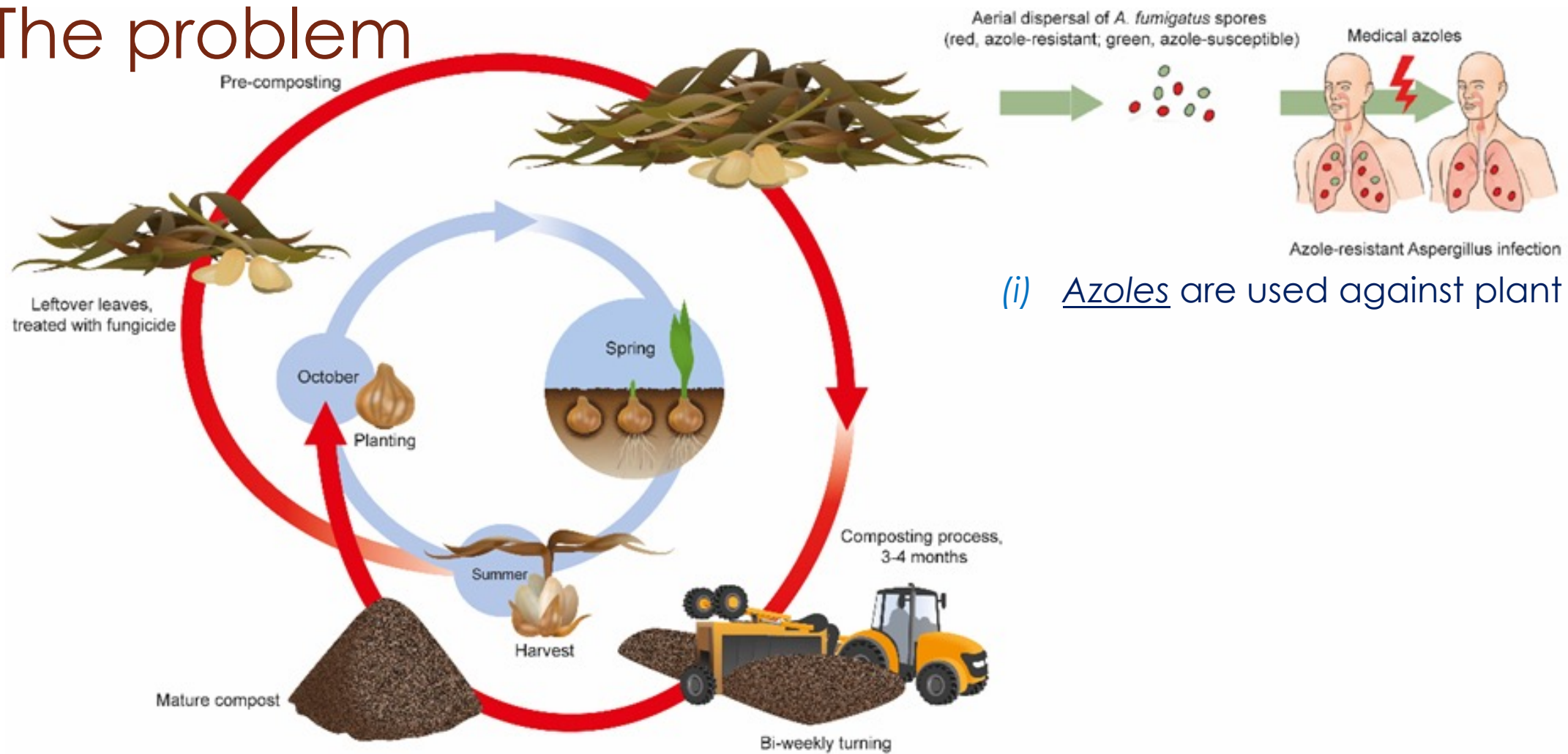
1995 2000 2005 2010

# The problem



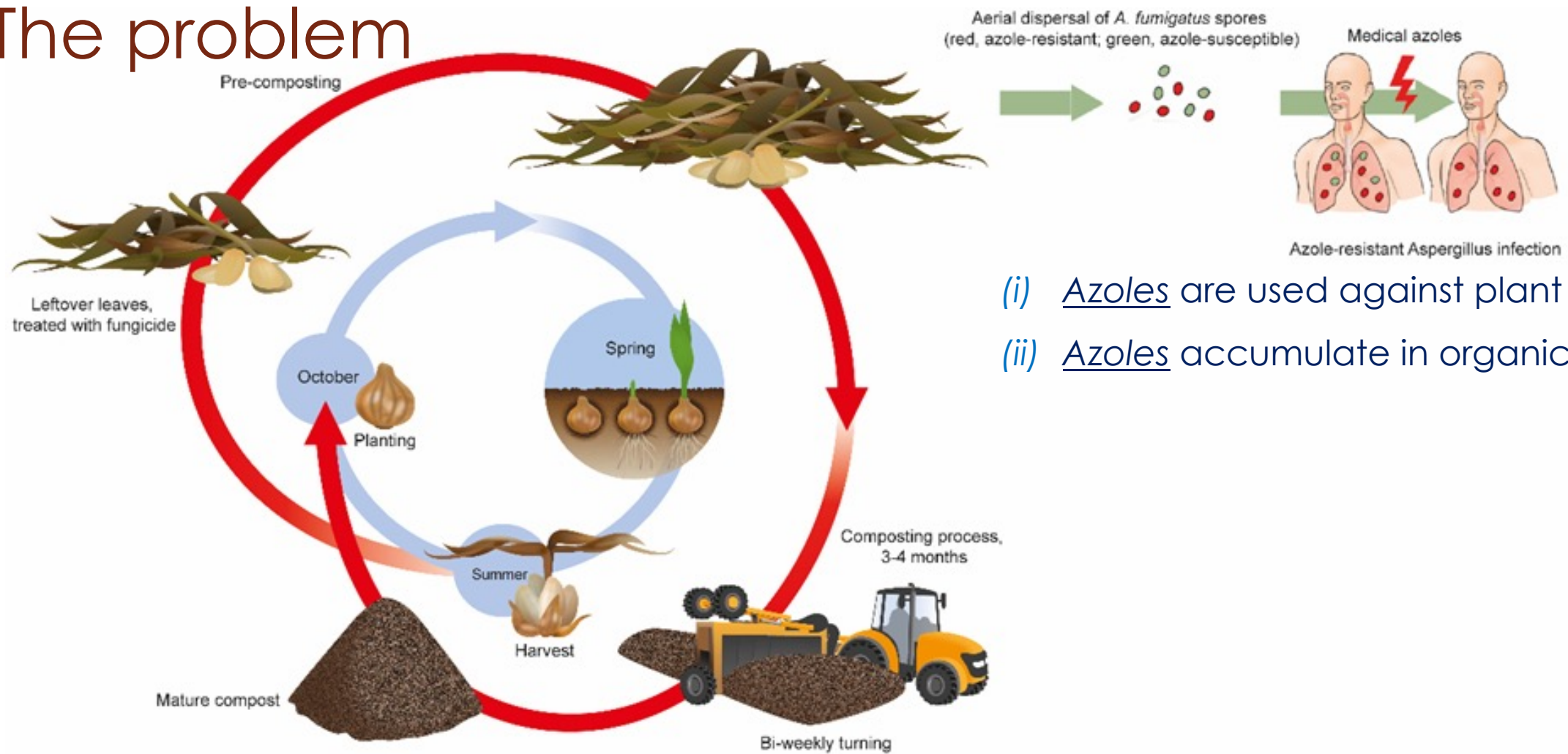


# The problem



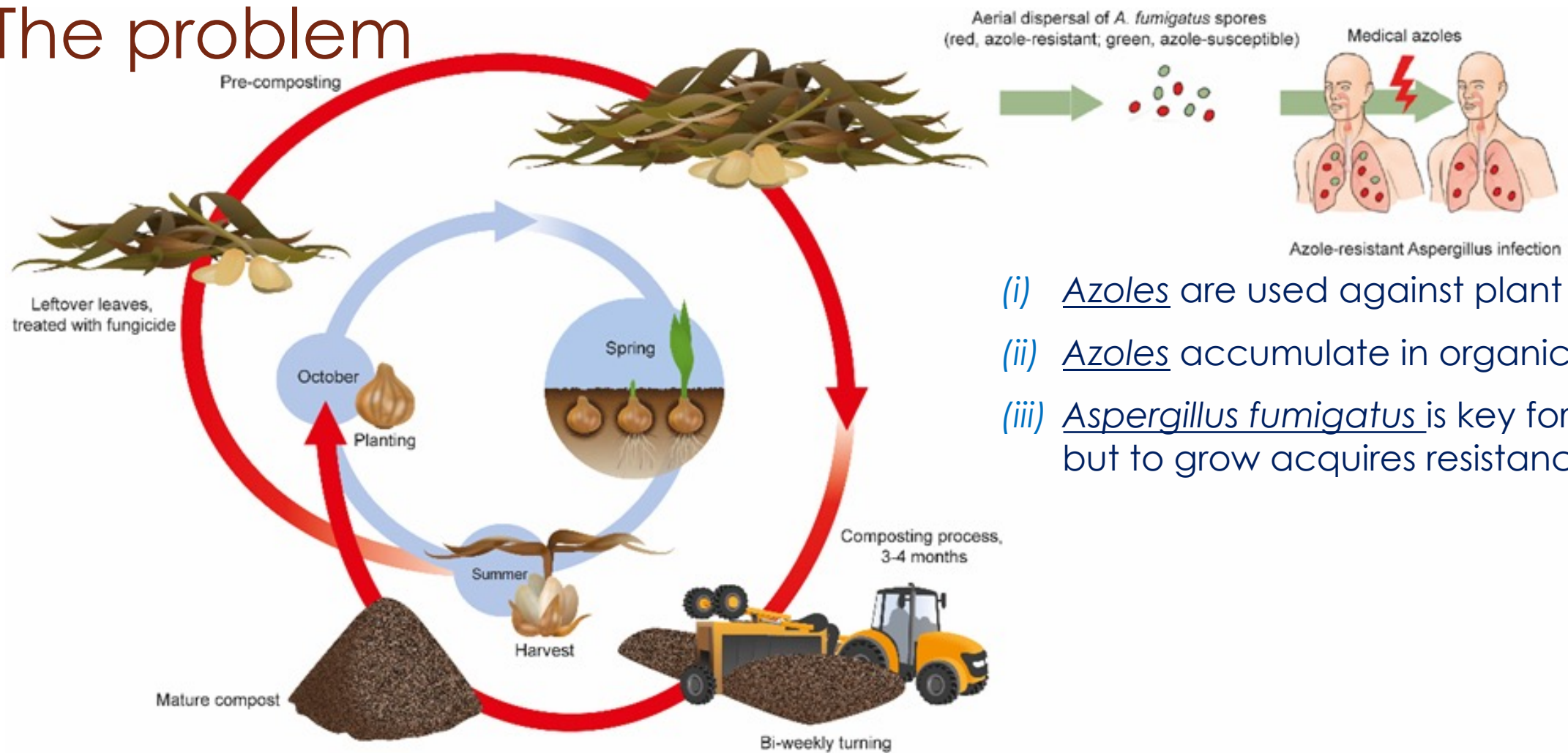
(i) Azoles are used against plant pathogens

# The problem



- (i) Azoles are used against plant pathogens
- (ii) Azoles accumulate in organic waste

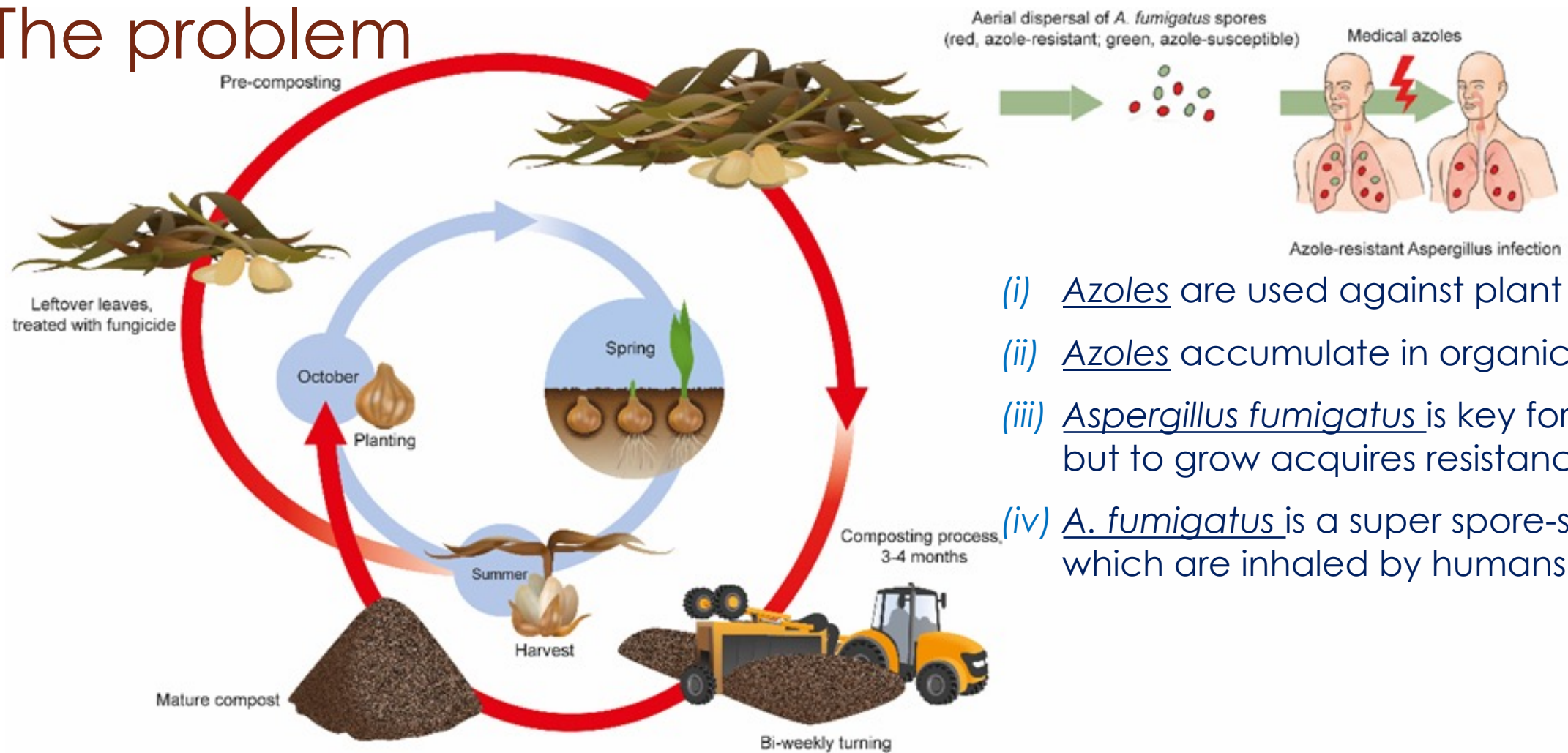
# The problem



- (i) Azoles are used against plant pathogens
- (ii) Azoles accumulate in organic waste
- (iii) Aspergillus fumigatus is key for composting but to grow acquires resistance to azoles

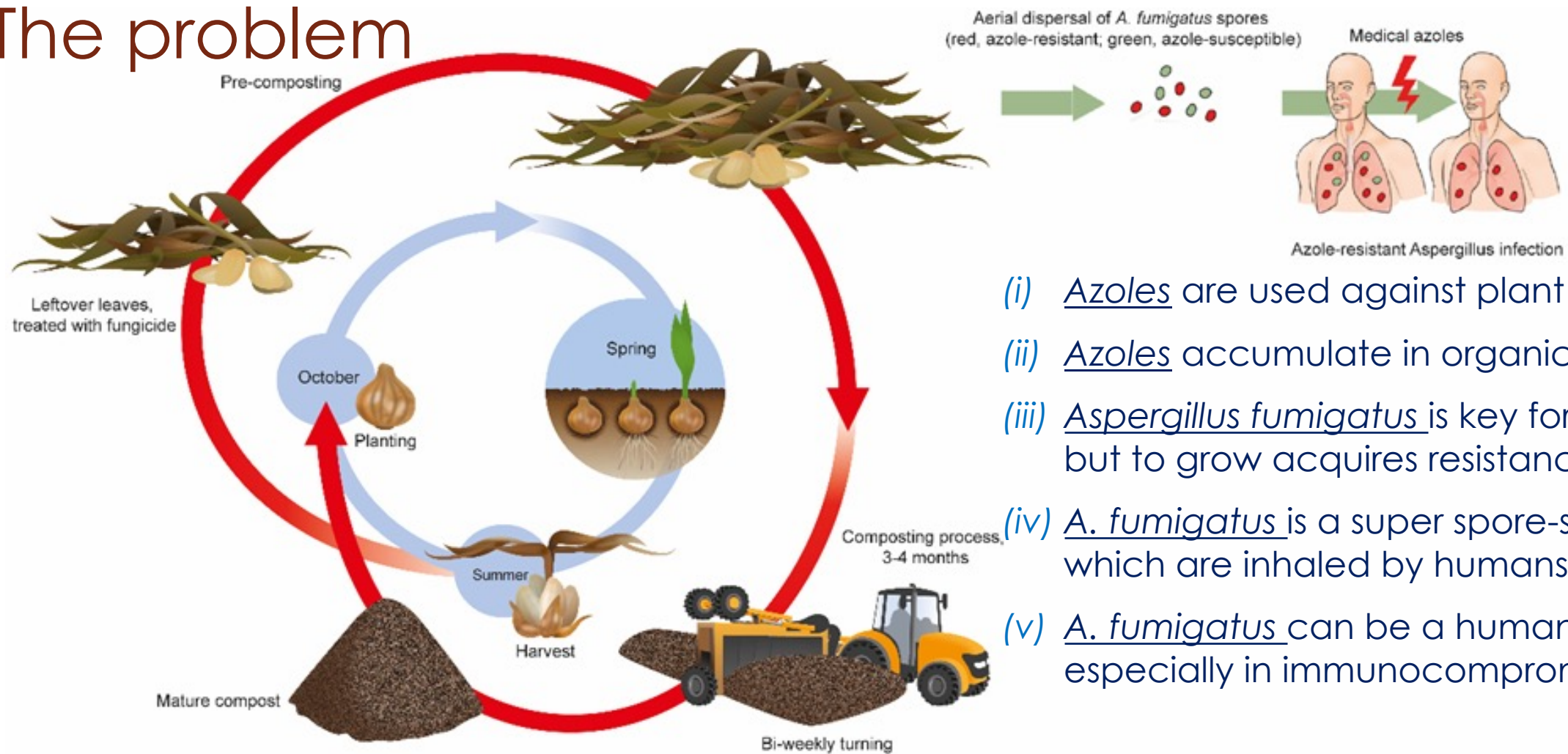


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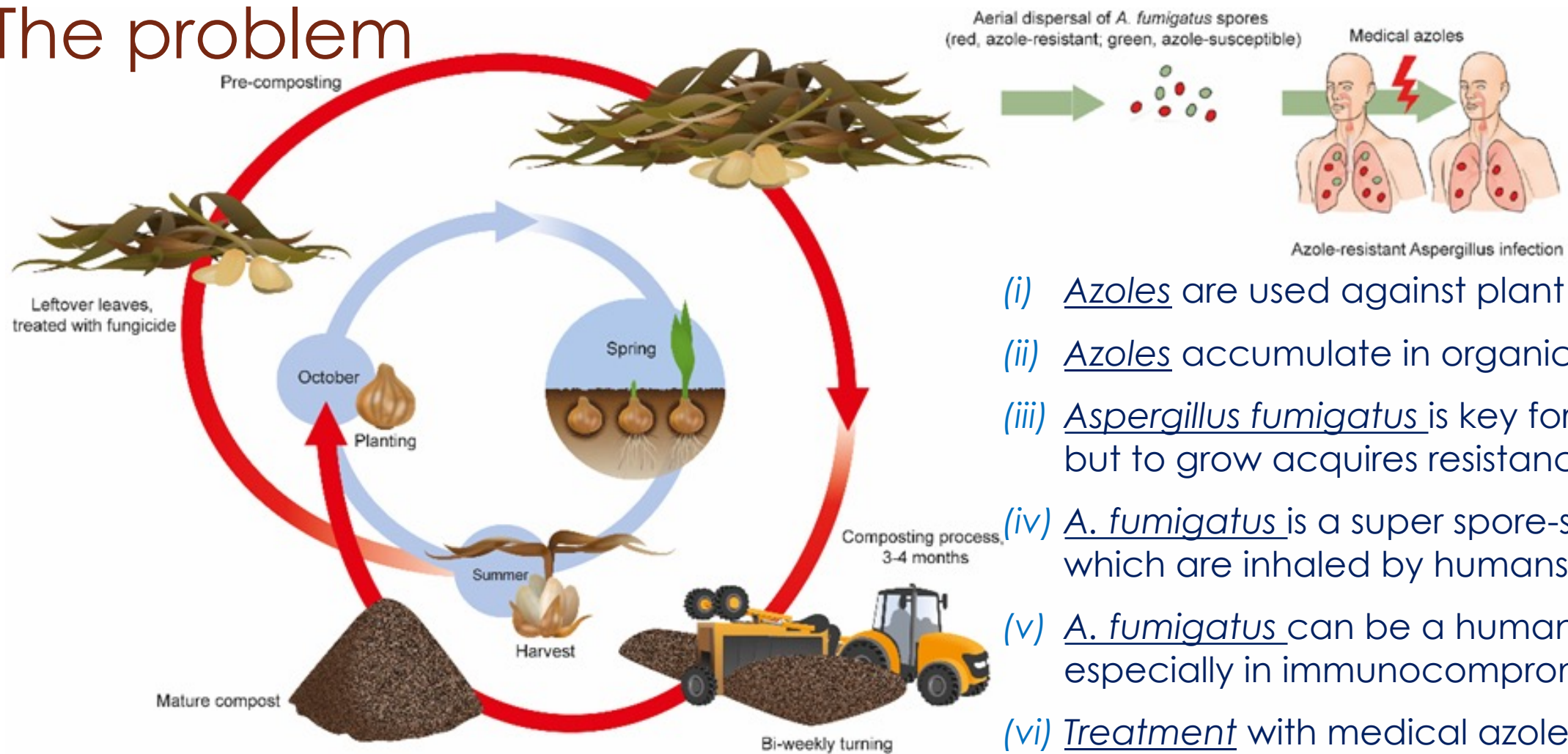
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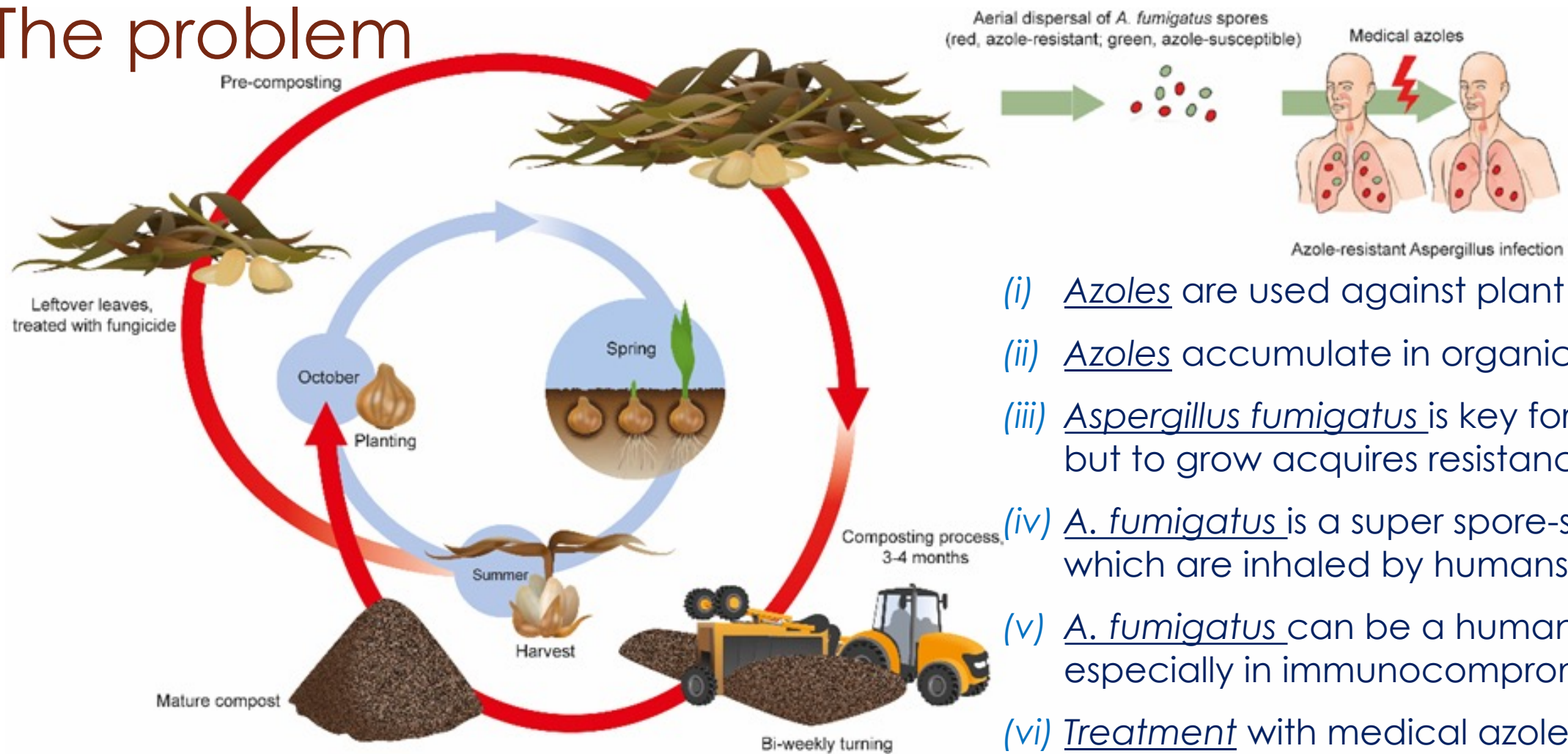


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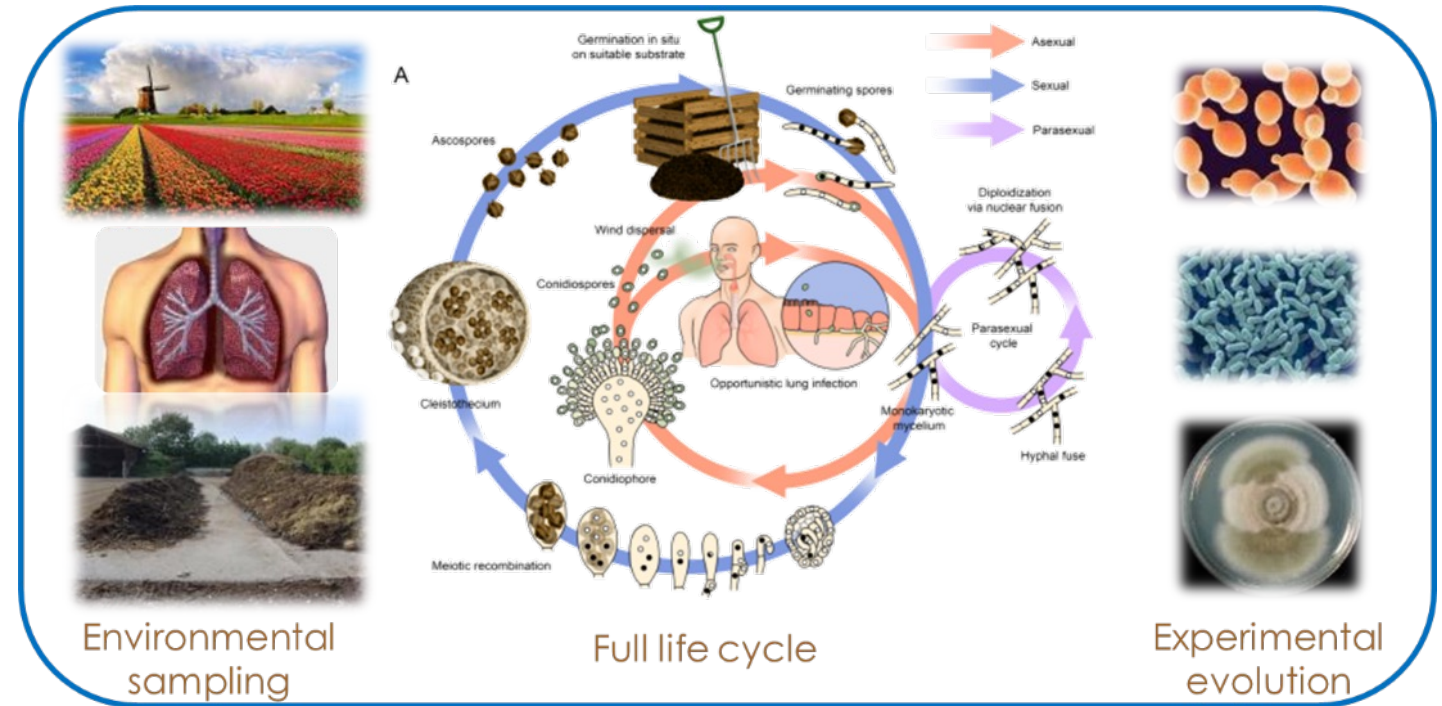
**When rest-streams, microbes, and chemicals meet  
a potential (One-)health problem emerges**

# Disentangling AMR evolution

Separating the communalities from the specifics

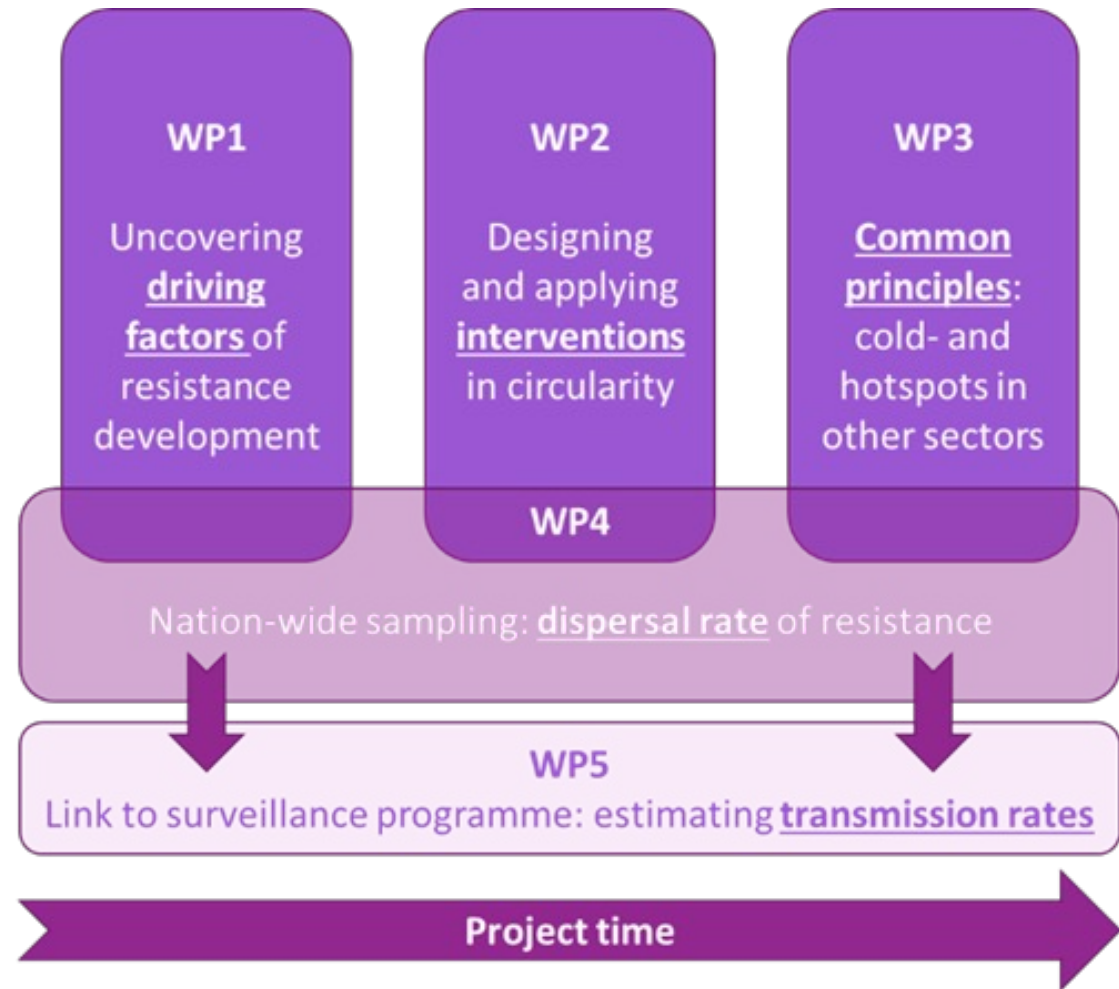
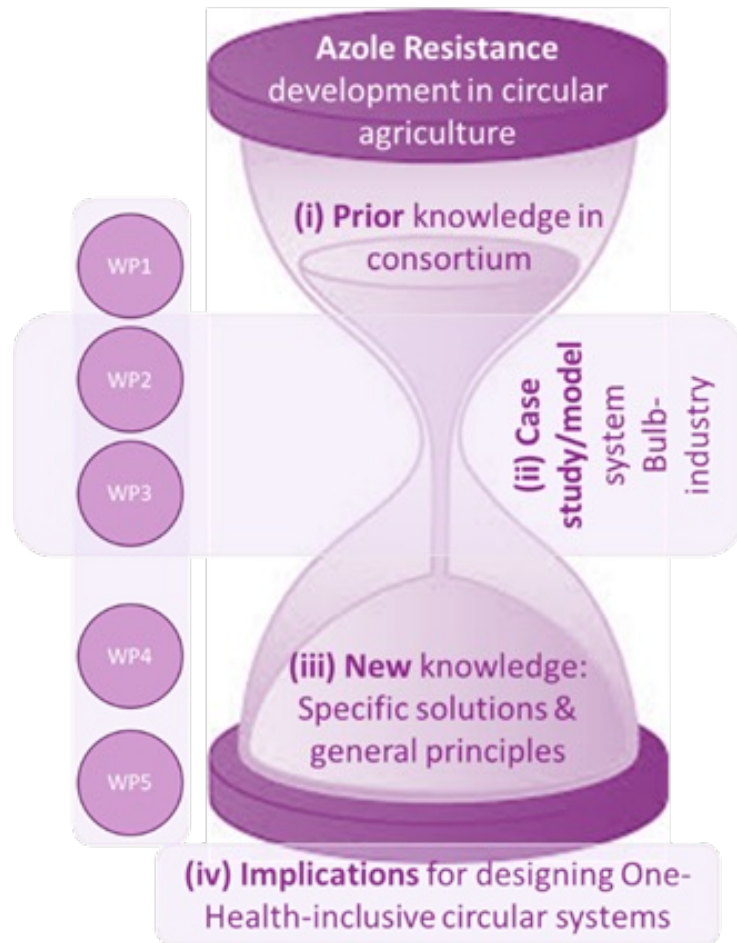
## Relevant factors

- Resistance mechanisms
- Ecological settings
- Biology of the species
- Nature of the selection pressure





# Scientific approach



# It is NOT just the Dutch bulb sector

Genetic analysis of the cyp51A promotor region and structural gene of 41 representative azole-resistant A. fumigatus isolates.								
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B-wood	NG_HAB_BW_200817_2	552000	0.03	0.03	Colony 1	ITR	34	L98H*
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Strawberry	DB_AB3_20201228_1B	43200000	0.13	0.13	Colony 1	ITR	34	L98H*
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Strawberry	DB_AB2_20201228_1A	54000000	0.03	0.01	Colony 1	ITR	0	None
					Colony 2	ITR	46	Y112F/T289A/S363P/I364V/G448S*
Onion	DA2_UI_20201210_1A	13800000	0.78	0.45	Colony 1	ITR	46	Y112F/T289A/S363P/I364V/G448S*
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Onion	EM_UI_20210106_1A	10800	0.19	0.19	Colony 1	ITR	46	Y112F/T289A/S363P/I364V/G448S*
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Potato	OB_AA1_20210108_1B	7560000	0.21	0.25	Colony 1	ITR	34	L98H*
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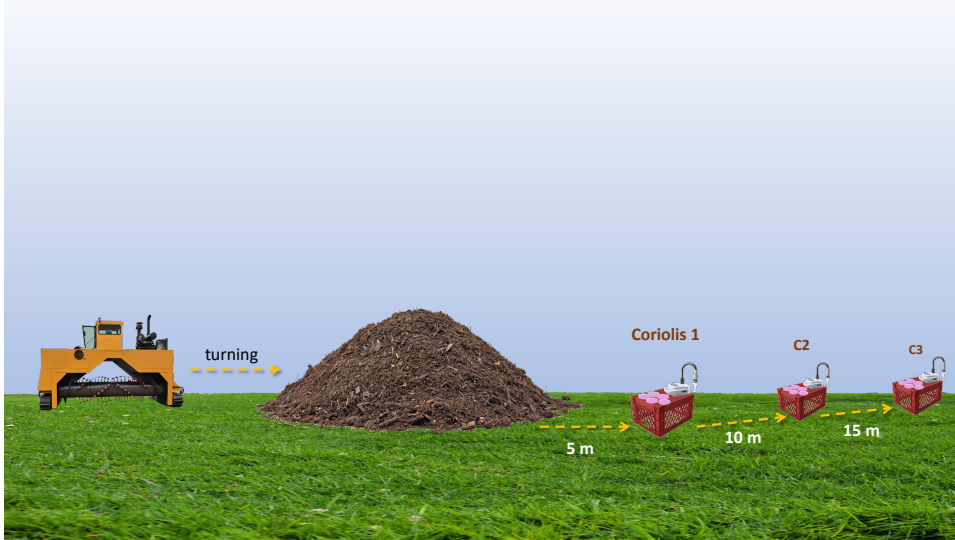
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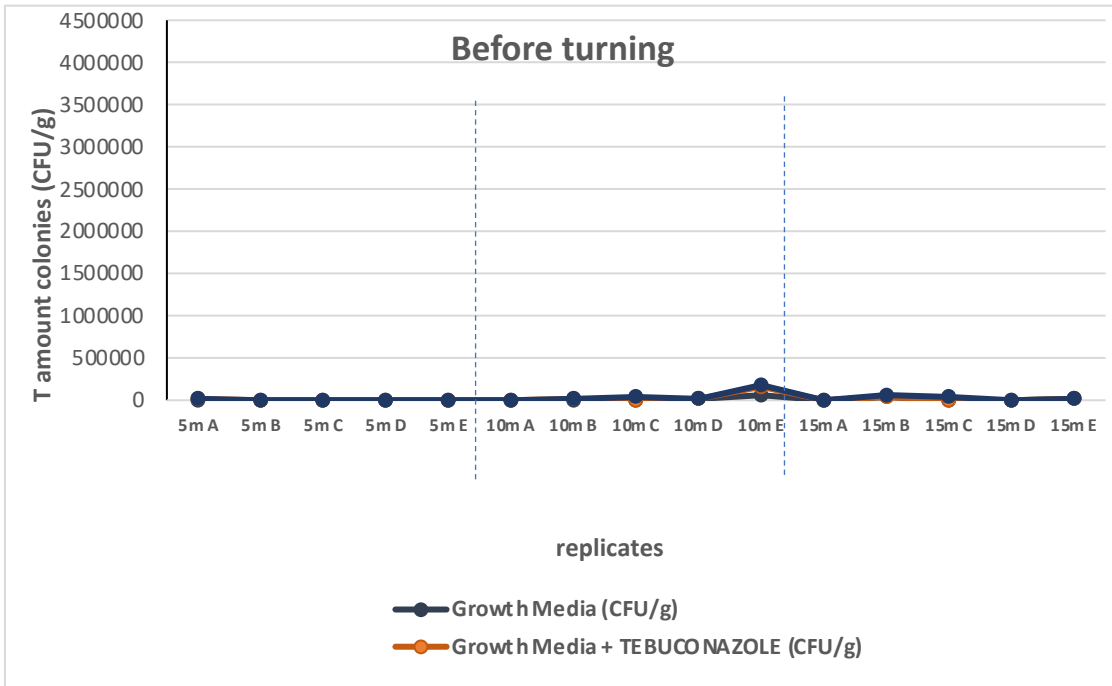
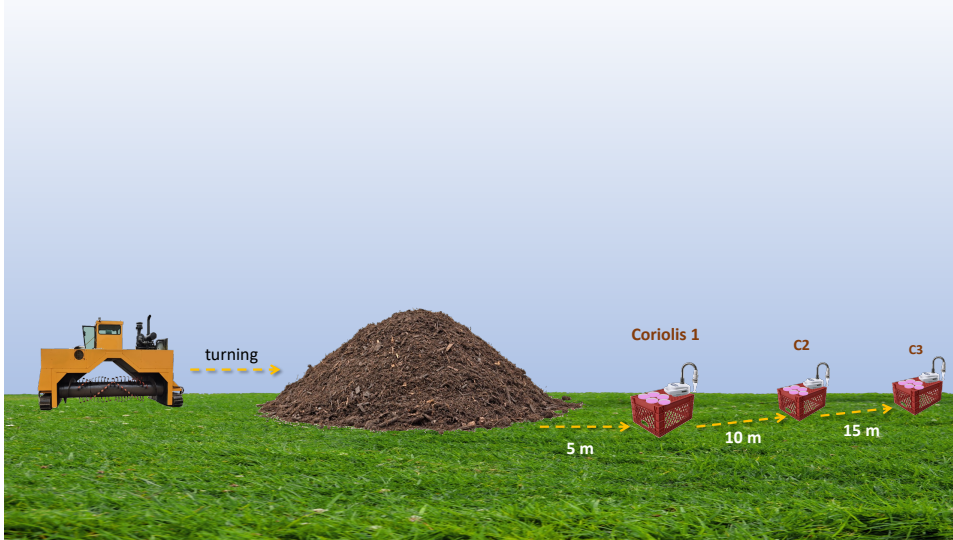
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- Similar tandem repeat-mediated resistance mechanisms were found as reported in clinical azole-resistant isolates

# Some highlights: air-sampling during turning



- Air-sampling around organic waste-heaps

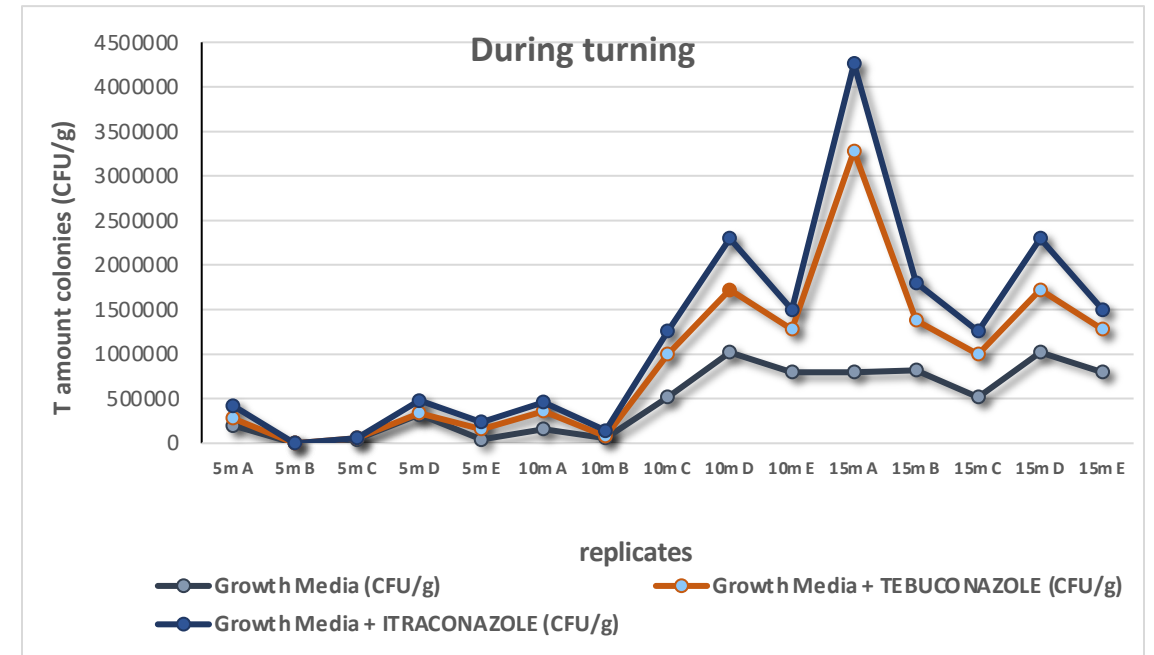
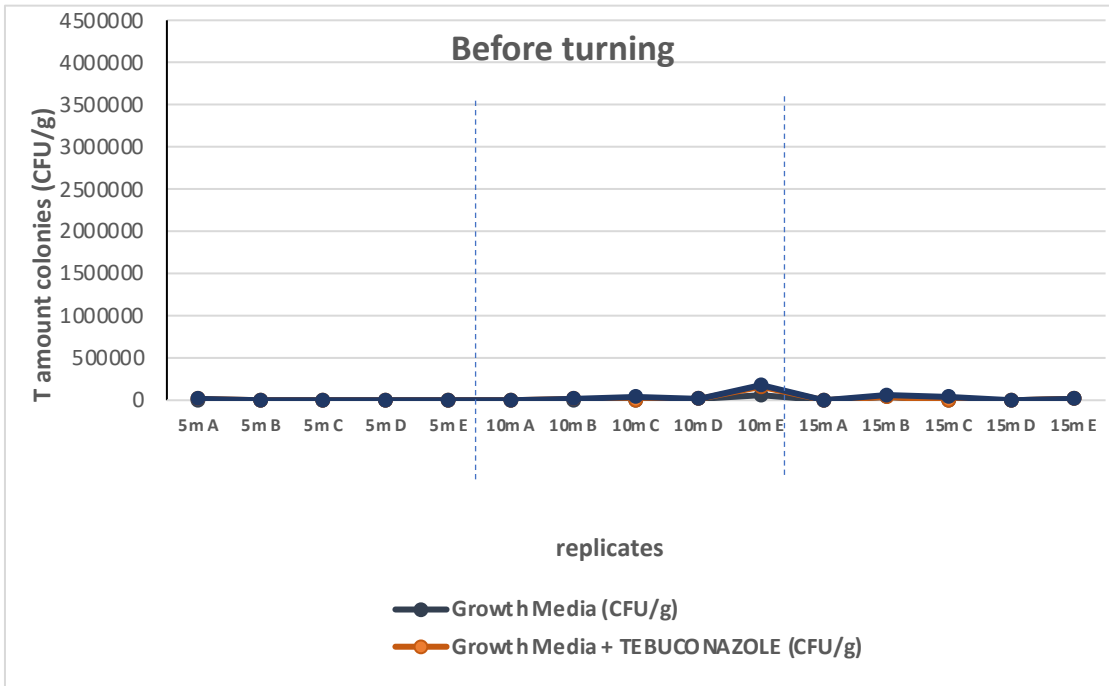
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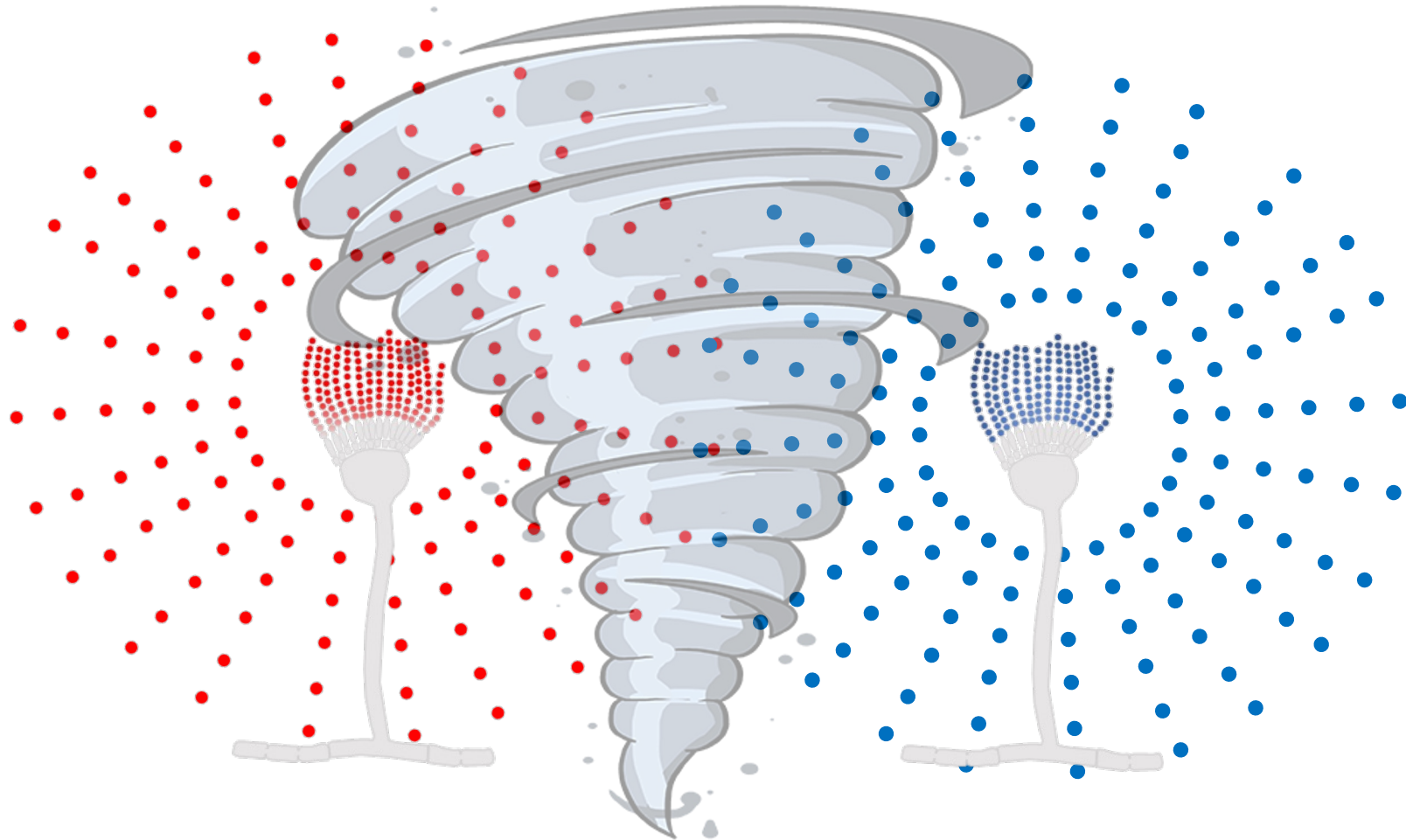


# Some highlights: air-sampling during turning



- Air-sampling around organic waste-heaps
- No spores trapped when undisturbed
- Profusive spore release from heaps when turned

# How to air sample?



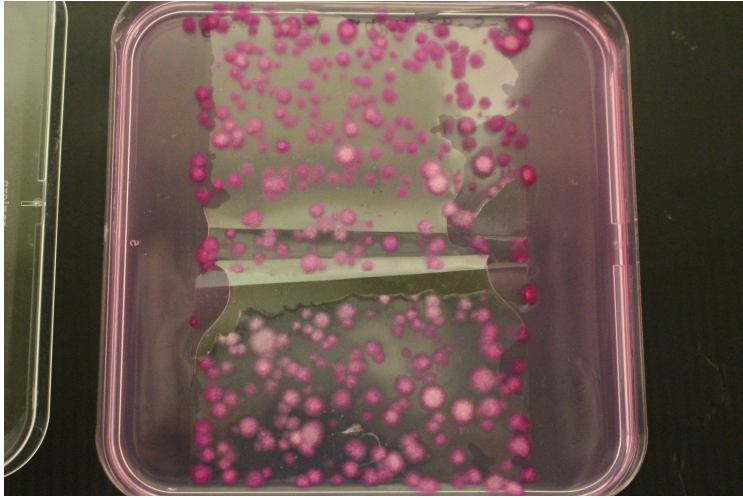
# How to air sample?

## A promising new method

- Incubate at 48C for three days

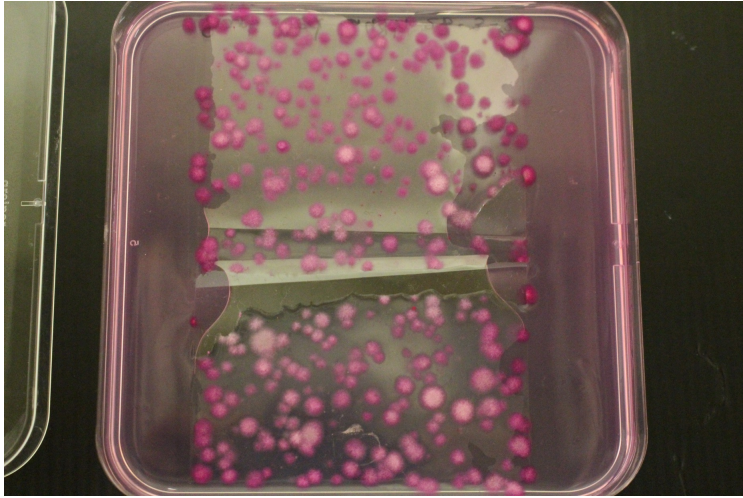


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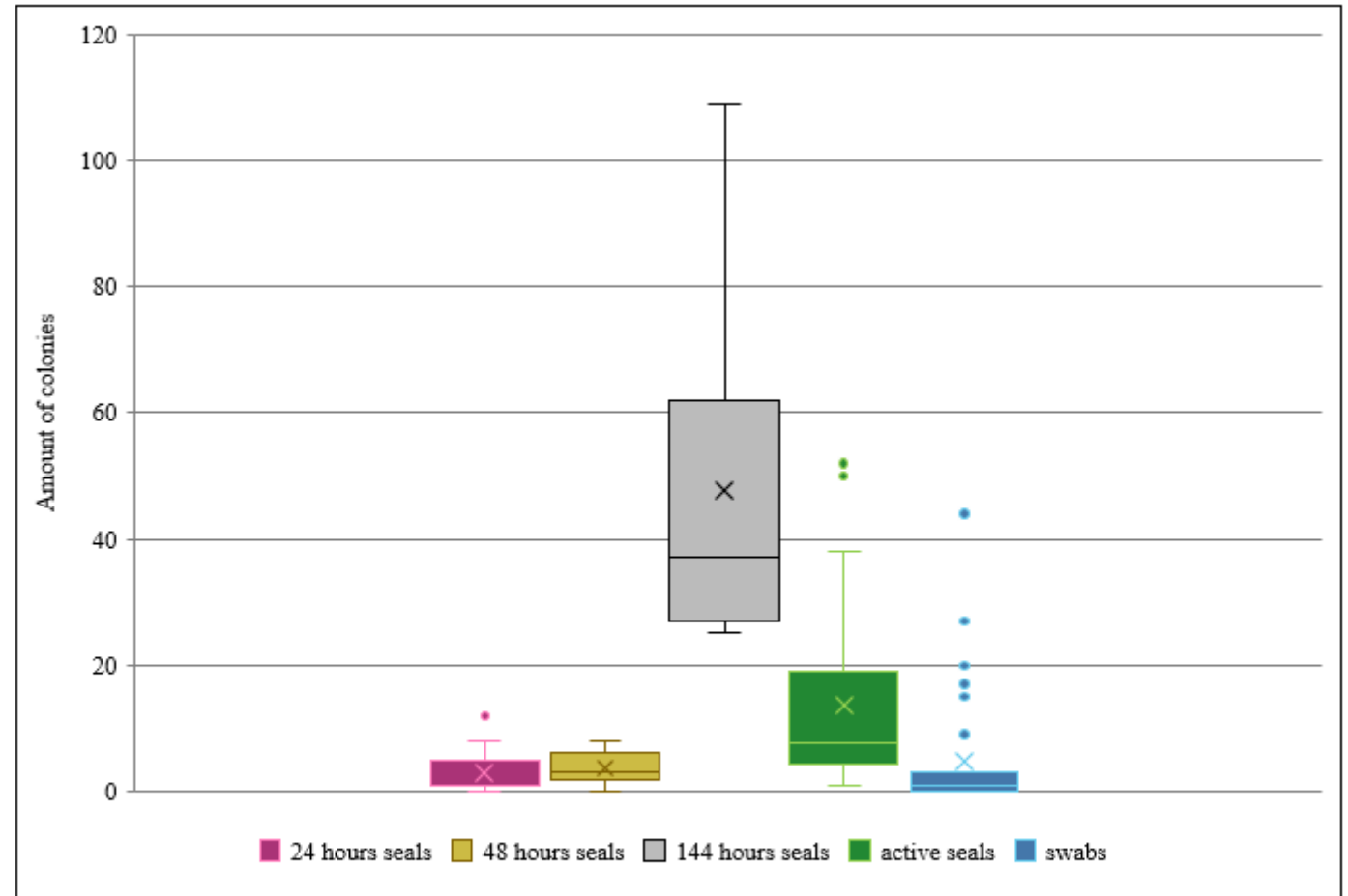
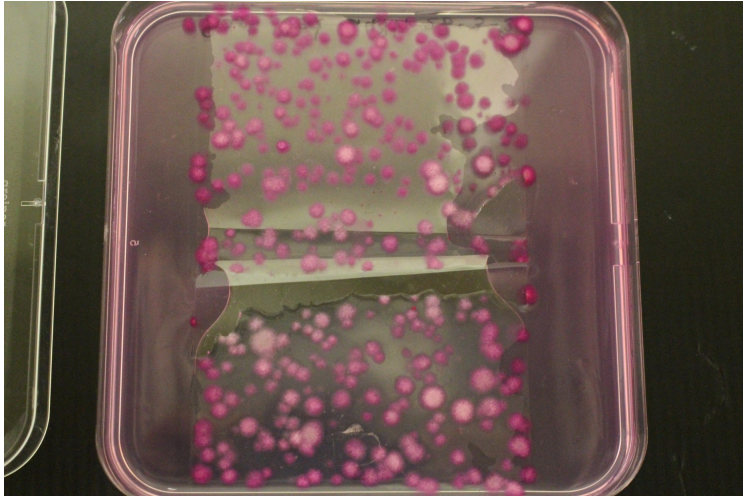


Figure: Colony counts of different sampling methods. The amount of colonies for respectively 24 hours, 48 hours, 144 hours seals, active seals and swabs.



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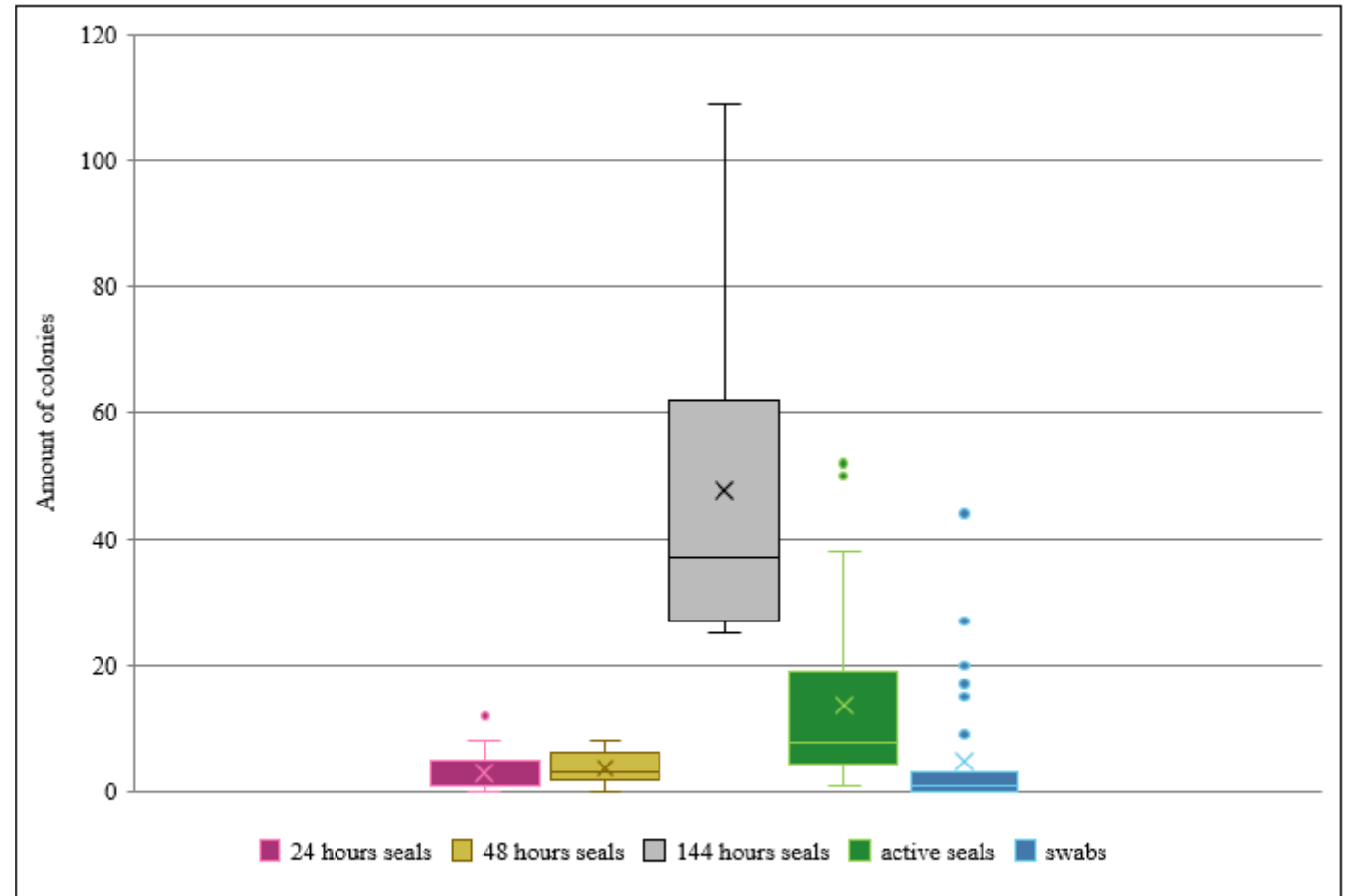


Figure: Colony counts of different sampling methods. The amount of colonies for respectively 24 hours, 48 hours, 144 hours seals, active seals and swabs.

# Population genetics of resistance mutations

Sex and recombination rate affect the dynamics of genetic variation

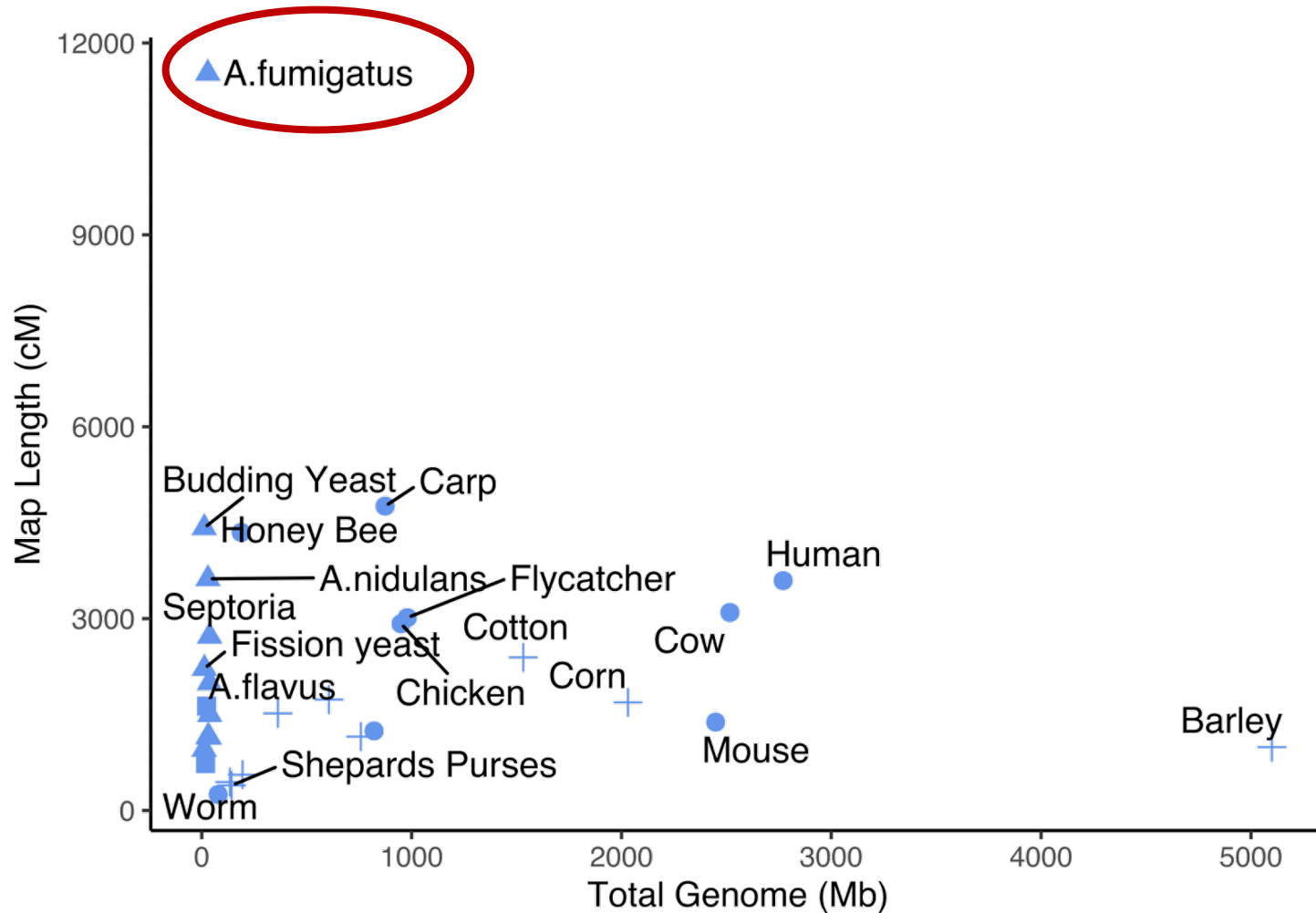
# Population genetics of resistance mutations

Sex and recombination rate affect the dynamics of genetic variation

- Cross between fertile strains AfIR964 and AfIR974
- 14,113 segregating markers
- 195 offspring from several cleistothecia; 90X short read sequencing

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Sex and recombination rate affect the dynamics of genetic variation

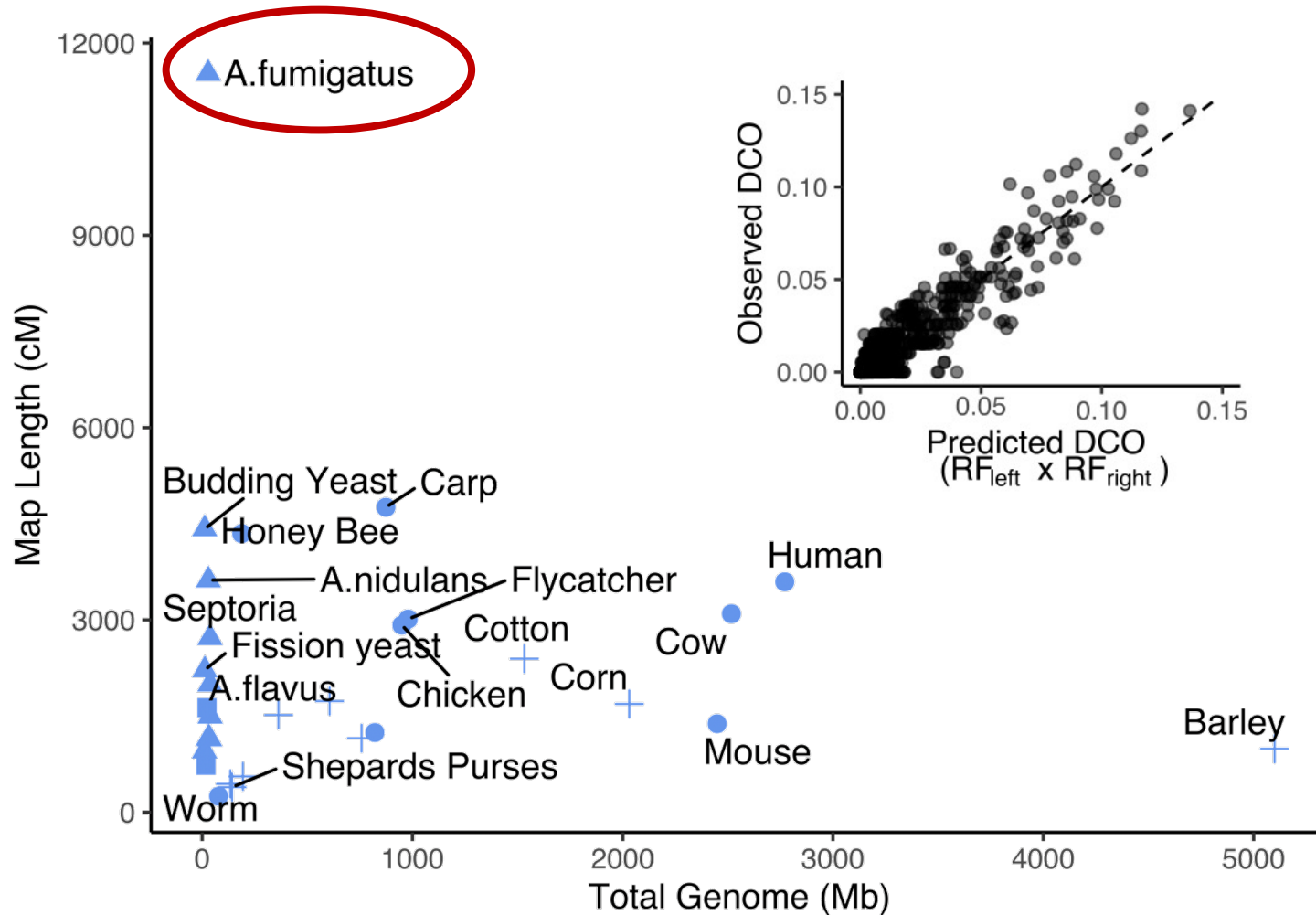


- Cross between fertile strains AfIR964 and AfIR974
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- Map length 11,966 cM
- 0.422 cM/kb
- ~ 29 cross-overs per chromosome



# Population genetics of resistance mutations

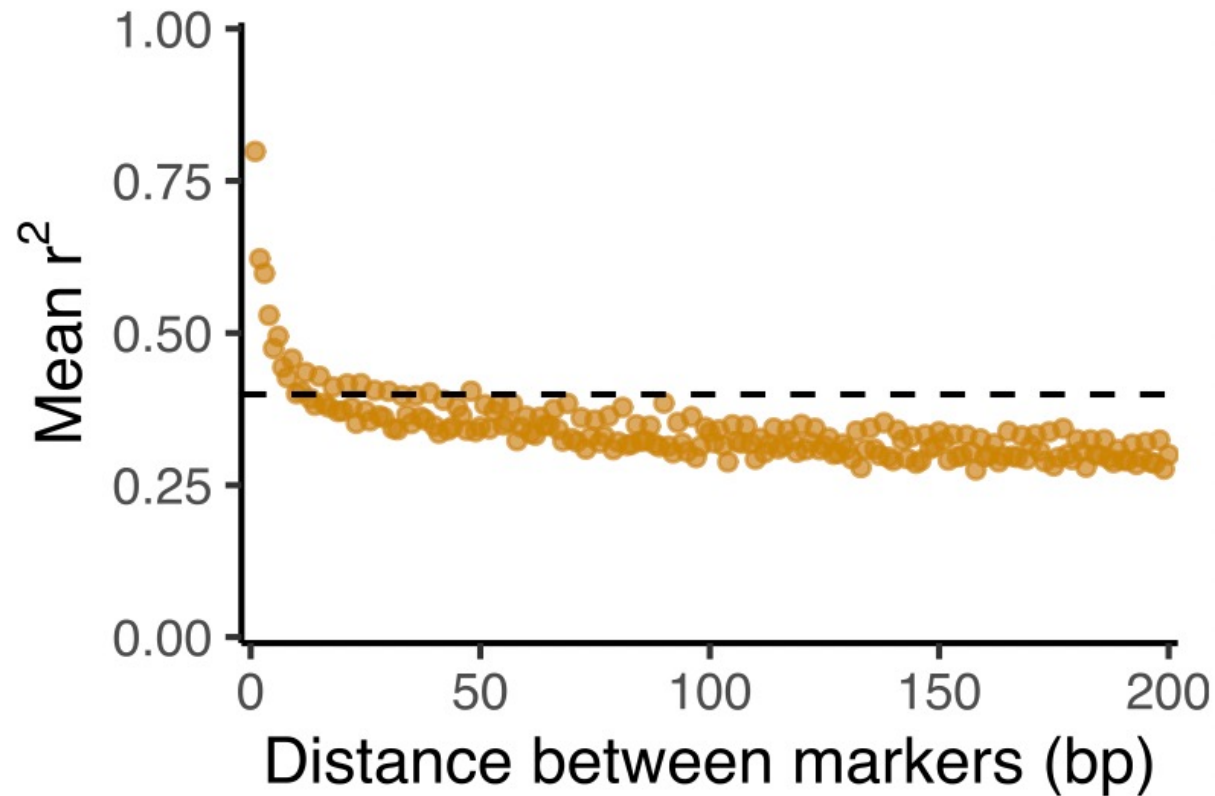
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- ~ 29 cross-overs per chromosome
- As for *A. nidulans*, no evidence for cross-over interference

# Population genetics of resistance mutations

High recombination rate, rapid decay of linkage disequilibrium (LD)



- Using published\* data from 175 sequenced individuals
- Very rapid decay of LD consistent with high recombination rates in *A. fumigatus*

\*A. E. Barber *et al.*, Nat Microbiol. 6, 1526–1536 (2021)

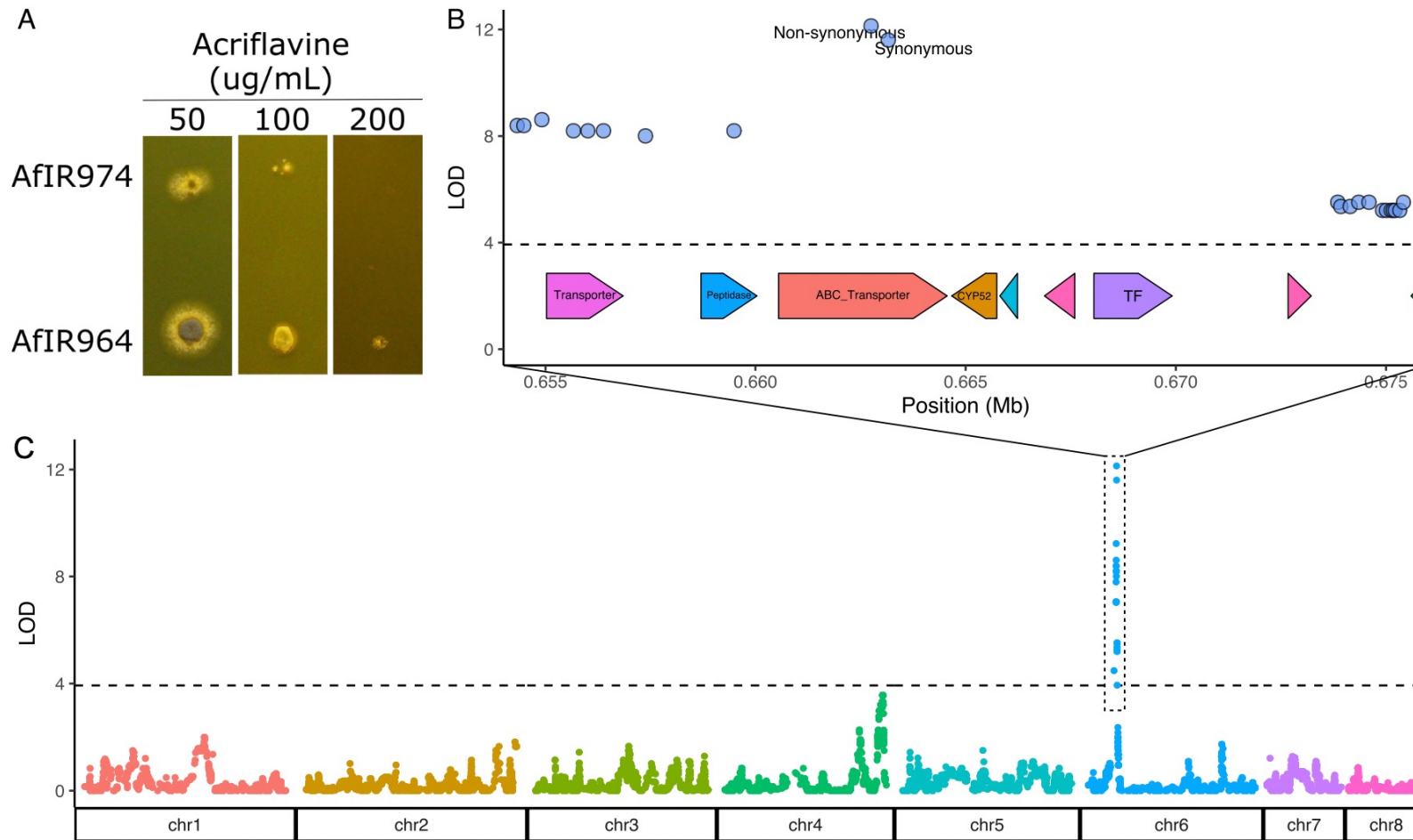
# Population genetics of resistance mutations

High recombination rates and low LD facilitates genetic mapping

- Parental strains differ in Acriflavine resistance
- Offspring phenotyped on 50µg/mL acriflavine

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High recombination rates and low LD facilitates genetic mapping



- Parental strains differ in Acriflavine resistance
- Offspring phenotyped on 50µg/mL acriflavine
- Single locus on chromosome 6 between positions 657kb and 675kb
- Presumed causal variant (Phe to Cys non-synonymous substitution)



# Population genetics of resistance mutations

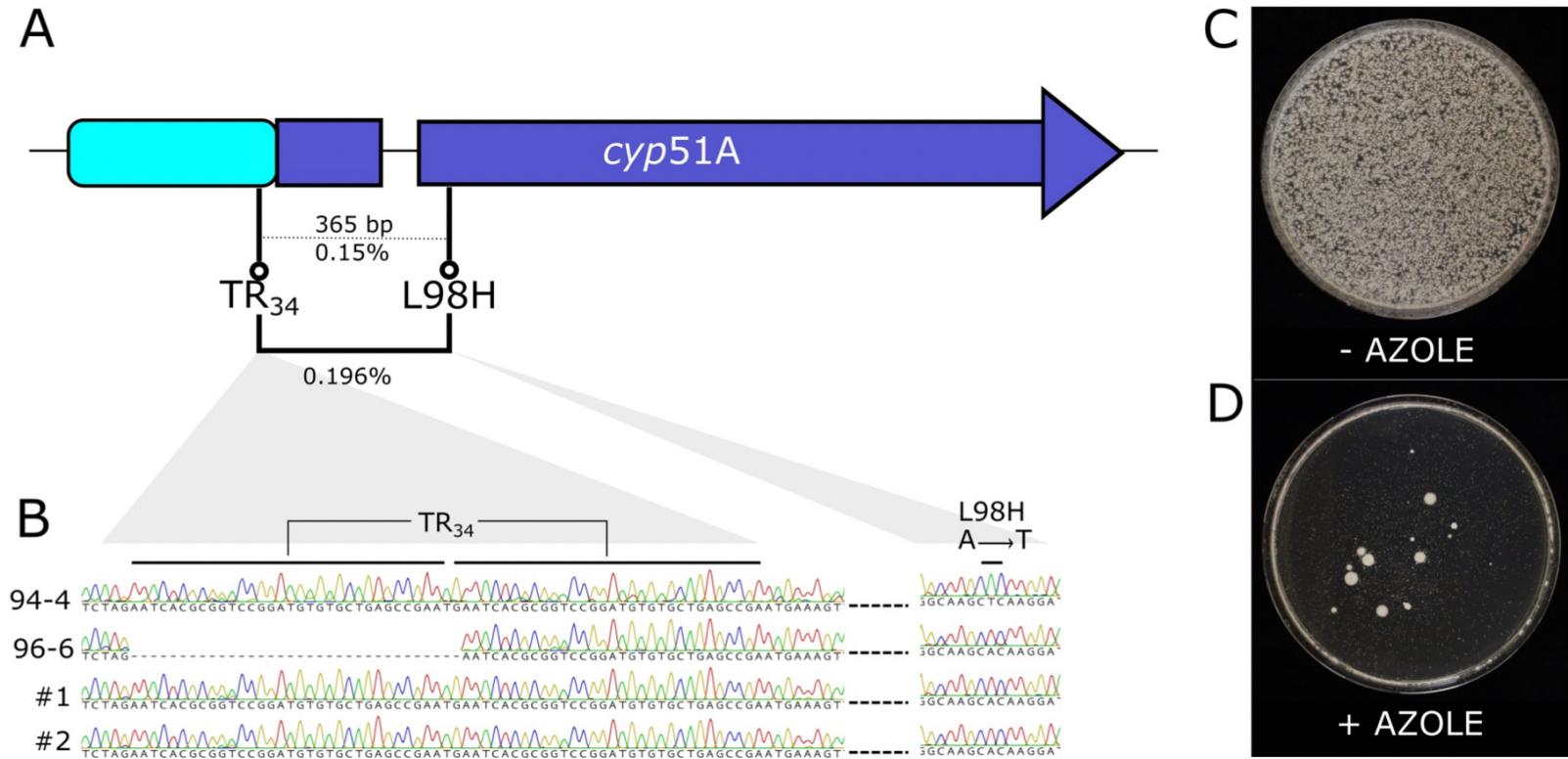
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# Linking the environment to the clinic: the smoking gun

Whole genome sequencing of a unbiased balanced set of samples

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Whole genome sequencing of a unbiased balanced set of samples

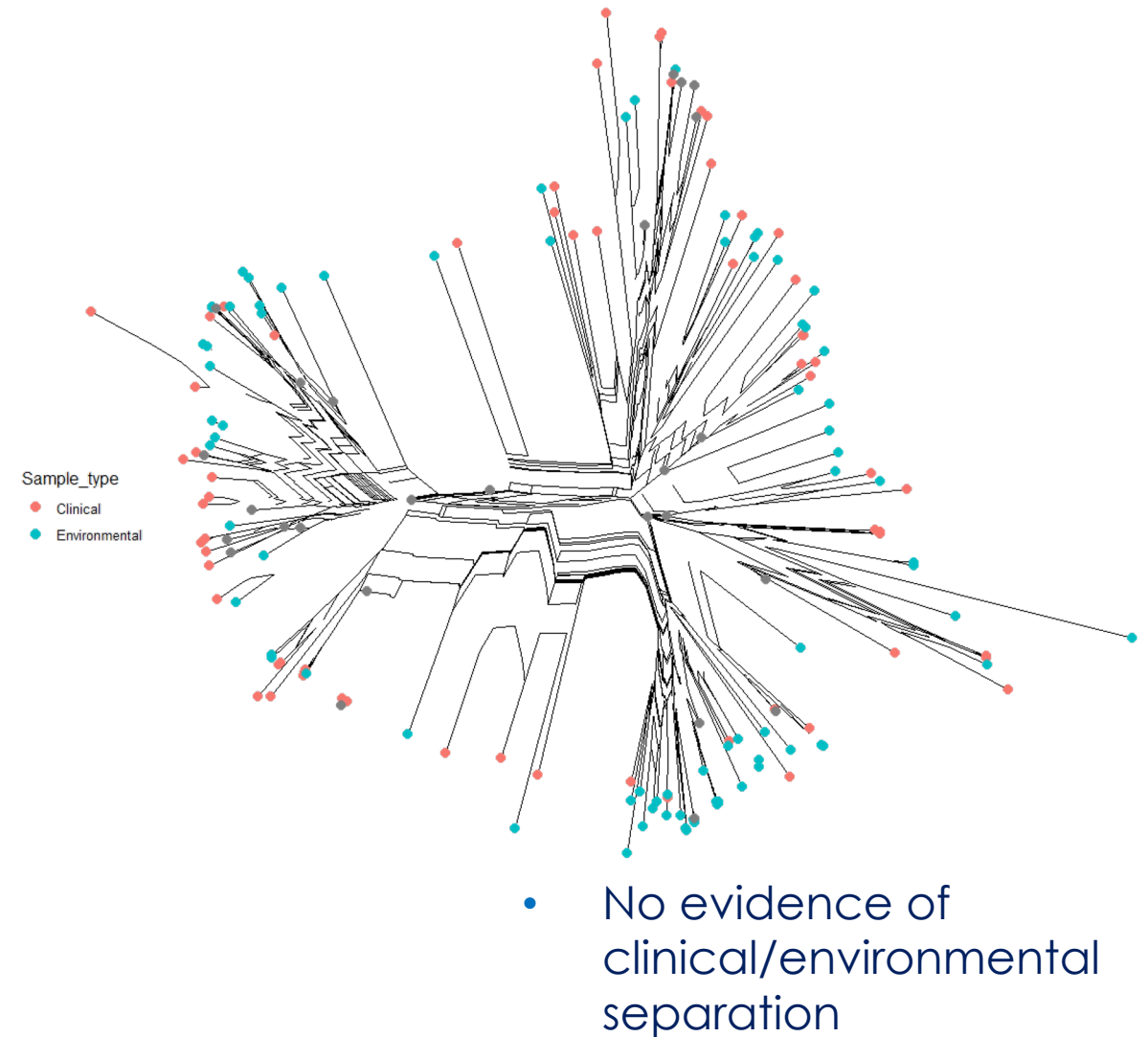
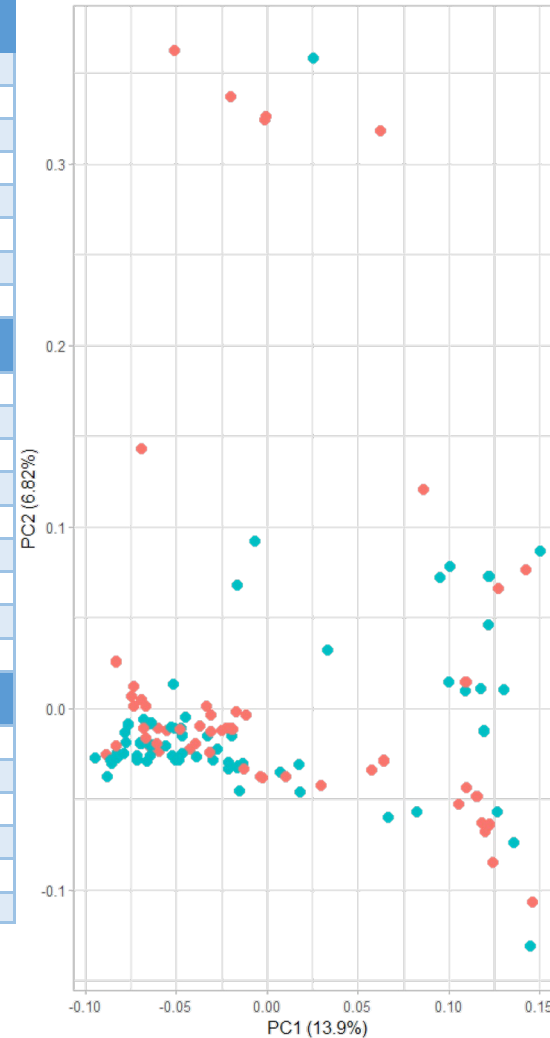
Haplotype		
	Environmental	Clinical
WT	26	26
WT R	4	6
TR34	24	33
TR46	31	15
TR92	0	2
unclear		3
Total	85	83
Location		
	Environmental	Clinical
Location A	31	
Location B	11	
Location C	40	
Other farm	3	
AmsterdamUMC		38
RadboudUMC		34
Other hospital		11
Total	85	83
Year		
	Environmental	Clinical
2016	38	18
2017	47	24
2018		31
2019		10
Total	85	83



# Linking the environment to the clinic: the smoking gun

Whole genome sequencing of a unbiased balanced set of samples

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  - Compensatory mutations and competition with wildtypes

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- Whole genome sequencing analysis to confirm causal link between resistance in environment and in the clinic
- Resistant *A. fumigatus* spores are ubiquitously present
  - Prevention of resistance development seems a lost cause
  - Prevention of transmission should be the new target



# Acknowledgements

## People



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Ben Auxier



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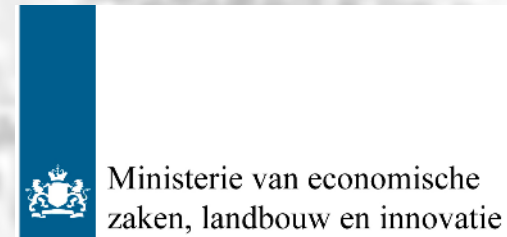


Jianhua Zhang



Bas Zwaan

## Funding





Thank you

I am happy to take  
questions and  
comments