

Host Resistance:

Two major types:

'Specific recognition'

race specific
qualitative
major
monogenic
differential
'R-gene'
vertical

'Concert' of defense responses

non-race specific
quantitative
minor
polygenic
general
field resistance
horizontal

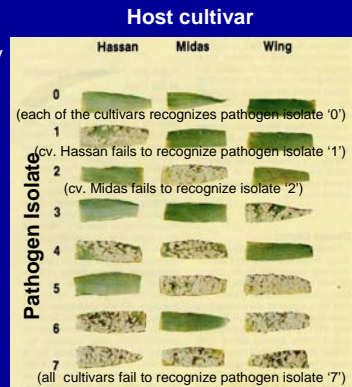
Genetics of resistance continued:

Race-specific resistance:

Resistance characterized by **specific recognition** (genetic interactions) between host genotypes and pathogen genotypes.

The ranking of cultivars from least to most resistant depends on the pathogen genotype (race) used.

This type of resistance is commonly 'all or nothing' (qualitative)



Genetics of resistance continued:

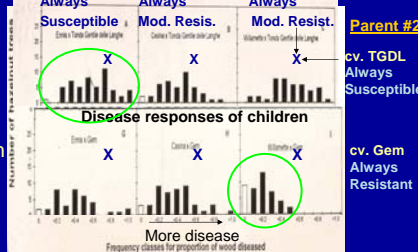
Non race-specific ('horizontal') resistance:

Host cultivars show the same relative level of resistance to all pathogen isolates

When effective, this type of resistance commonly involves multiple genes, each with a small effect (quantitative)

Example: Parent #1: cv. Ennis, cv. Casina, cv. Willamette

Disease responses in 2-yr-old seedlings of European hazelnut where the parent cultivars (crossed with each other) varied in level of non-race specific resistance



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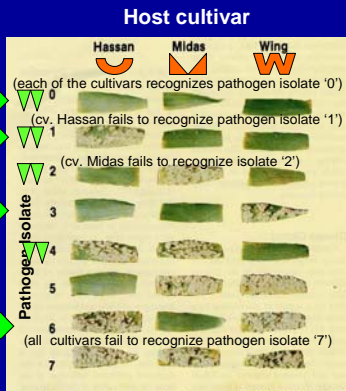
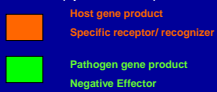


Fig. 1. The seven possible race combinations corresponding to the three resistance phenotypes—Hassan (R₁), Midas (R₂), and Wing (R₃)—include three simple races, three intermediate races, and one complex race.

Race-specific resistance is based on the

Gene-for-Gene Hypothesis :

The product of a single gene in the host **specifically recognizes** the product of a gene in the pathogen (i.e., an 'effector'). During an infection event, an interaction between these products results in incompatibility (i.e., a hypersensitive response and programmed cell death).

Gene-for-Gene Hypothesis

Host: 'R'-gene = recognizer
this gene produces a product that 'recognizes'

usually an R-gene is a dominant gene

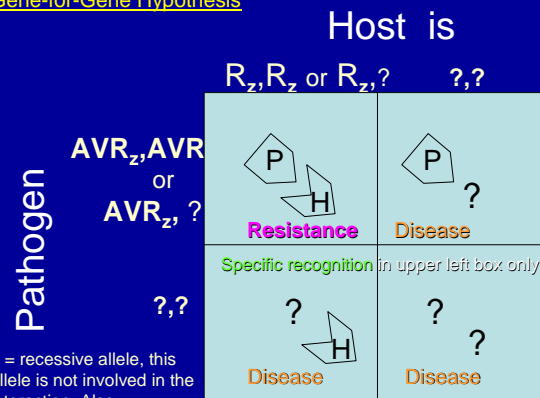
'dominant' means that receiving a copy of the gene from only one parent is sufficient to trigger the recognition response in the plant

Pathogen: 'AVR' gene = avirulence gene

this gene produces an elicitor (effector) – it 'elicits the HR'
an AVR-gene is dominant

dominant means that receiving a copy of the gene from only one parent is sufficient to elicit a recognition response in the host

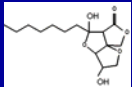
Gene-for-Gene Hypothesis



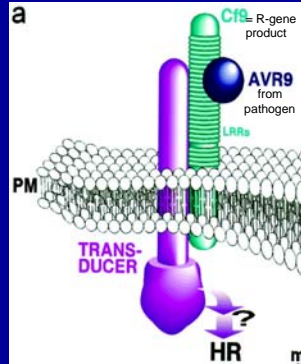
? = recessive allele, this allele is not involved in the interaction. Also sometimes written as 'r' in host, and 'avr' in pathogen

Hypothesized physical model for how R-gene and Avr-gene products interaction on plasma membrane of host cells

Pathogen protein (AVR) interacts with host R-gene protein (Cf9), which triggers the cell death response (hypersensitivity)



Example of AVR gene product from bacterial pathogen



From: MHAJ Joosten and PJGM de Wit Annual Review of Phytopathology Vol. 37:335

What if the pathogen is a necrotrophic fungus that produces a host selective toxin?

		Host	
		S,S or S,?	?, ?
Pathogen	Tox, Tox OR Tox, ?	Disease!! Disease!!	Resistant
	?, ?	Resistant	Resistant

Gene-for-gene II
Tox = gene for toxin
S = gene for receptor
Specific recognition in upper left box only

How many races?

Three cultivars, each with a different R gene yields 8 races (2^3)

Ten cultivars, each with a different R gene could discern a possible 1024 races (2^{10}) !!

The set of cultivars used to characterize races is called a 'differential set'

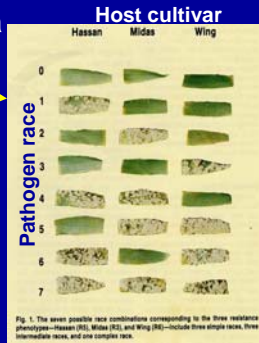


Fig. 1. The seven possible race combinations corresponding to the three resistance phenotypes—Hassan (H), Midex (M), and Wing (W)—include three simple races, three intermediate races, and one complete race.

A common test question: 😊

For each of the boxes, indicate whether or not the races of a rust pathogen will succeed in infecting the host cultivars

Avirulence (A) or lack-of-avirulence (? or 'a') genes possessed

		Race: A ₁ A ₂ A ₃ A ₄ A ₁ a ₂ A ₃ A ₄ a ₁ A ₂ A ₃ A ₄ a ₁ a ₂ A ₃ A ₄ a ₁ a ₂ a ₃ a ₄				
Resistance (R) or lack-of-resistance (? or 'r') genes in each cultivar	r ₁ r ₂ r ₃ r ₄					
	R ₁ r ₂ r ₃ r ₄					
	r ₁ R ₂ r ₃ r ₄					
	R ₁ R ₂ r ₃ r ₄					
	R ₁ R ₂ R ₃ R ₄					

'R', 'A' = active product present '?', 'a', 'r' = active product absent

A common test question: 😊

For each of the boxes, indicate whether or not the races of a rust pathogen **will succeed** in infecting the host cultivars

Avirulence (A) or lack-of-avirulence (? or 'a') genes possessed

Resistance (R) or lack-of-resistance (? or 'r') genes in each cultivar	Race:					
	A ₁ A ₂ A ₃ A ₄	A ₁ a ₂ A ₃ A ₄	a ₁ A ₂ A ₃ A ₄	a ₁ a ₂ A ₃ A ₄	a ₁ a ₂ a ₃ a ₄	
	r ₁ r ₂ r ₃ r ₄	yes	yes	yes	yes	yes
	R ₁ r ₂ r ₃ r ₄	no	no	yes	yes	yes
	r ₁ R ₂ r ₃ r ₄	no	yes	no	yes	yes
	R ₁ R ₂ r ₃ r ₄	no	no	no	yes	yes
R ₁ R ₂ r ₃ R ₄	no	no	no	no	yes	

'R', 'A' = active product present 'r', 'a', '?' = active product absent