



# Genetic diversity of Belgian *Phytophthora ramorum* isolates

I. De Dobbelaere, A. Vercauteren, K. Heungens, M. Maes

*Phytophthora* & *Pythium* Workshop  
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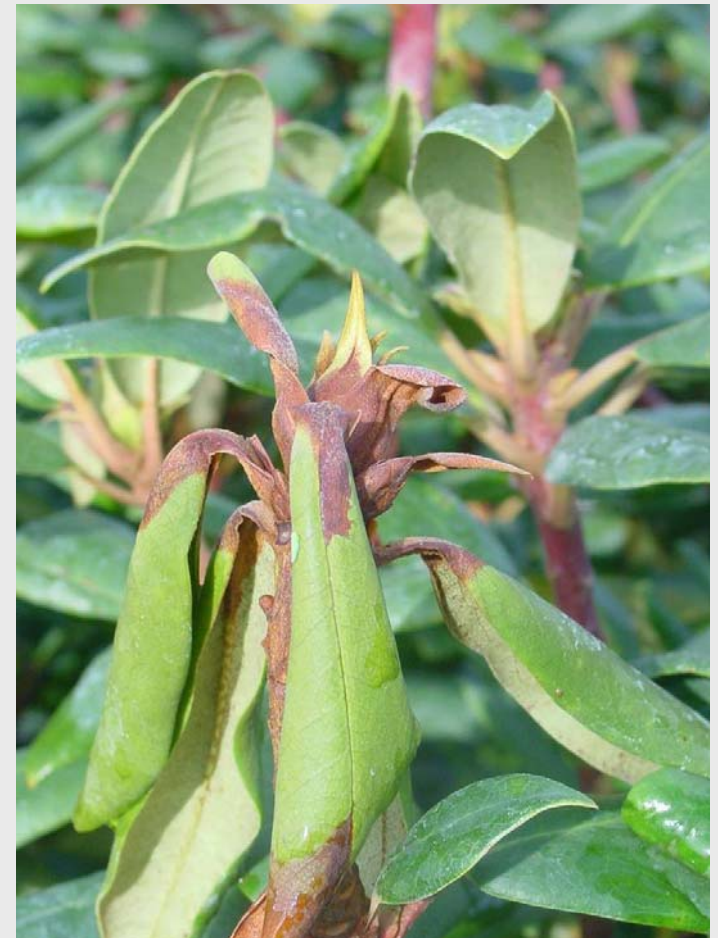
Northern part of Belgium  
(Flanders) is one of the  
largest producers of  
*Rhododendron* in Europe



# Results of the annual federal survey (FAVV)

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- ± 50 major nurseries sites inspected
- 100-500 samples annually
- 20-25% of Belgian samples positive
- Main host plants:  
*Rhododendron* (>80%) and *Viburnum*



# Global research genotyping *P. ramorum*

		<b>Number of isolates</b>	<b>Primer pairs</b>	<b>Different genotypes</b>	<b>Dominant genotype</b>
Ivors et al. 2004	AFLP	67 USA 18 EU	4	16 USA 15 EU	USA: 1
Ivors et al. 2006	SSLP	71 USA 80 EU	12	4 USA 7 EU	USA: 1 EU: 1
Prospero et al. 2007	SSLP	323 USA	10	33 USA	USA: 1 (OR)
Mascheretti et al. 2008	SSLP	292 USA	9	35 USA	USA: 3 (CA)

# Objective

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Determine the genetic diversity within the population of Belgian *P. ramorum* isolates

# Materials and methods

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## 1. Amplified Fragment Length Polymorphism (AFLP)

- 79 isolates (2002-2006)
- Protocol:
  - Adapted from Ivors et al. (2004)
  - Five pairs of selective primers
  - 2-3 replicates per *P. ramorum* isolate (reproducibility)

# Materials and methods

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## **2. SSLP (microsatellite polymorphisms)**

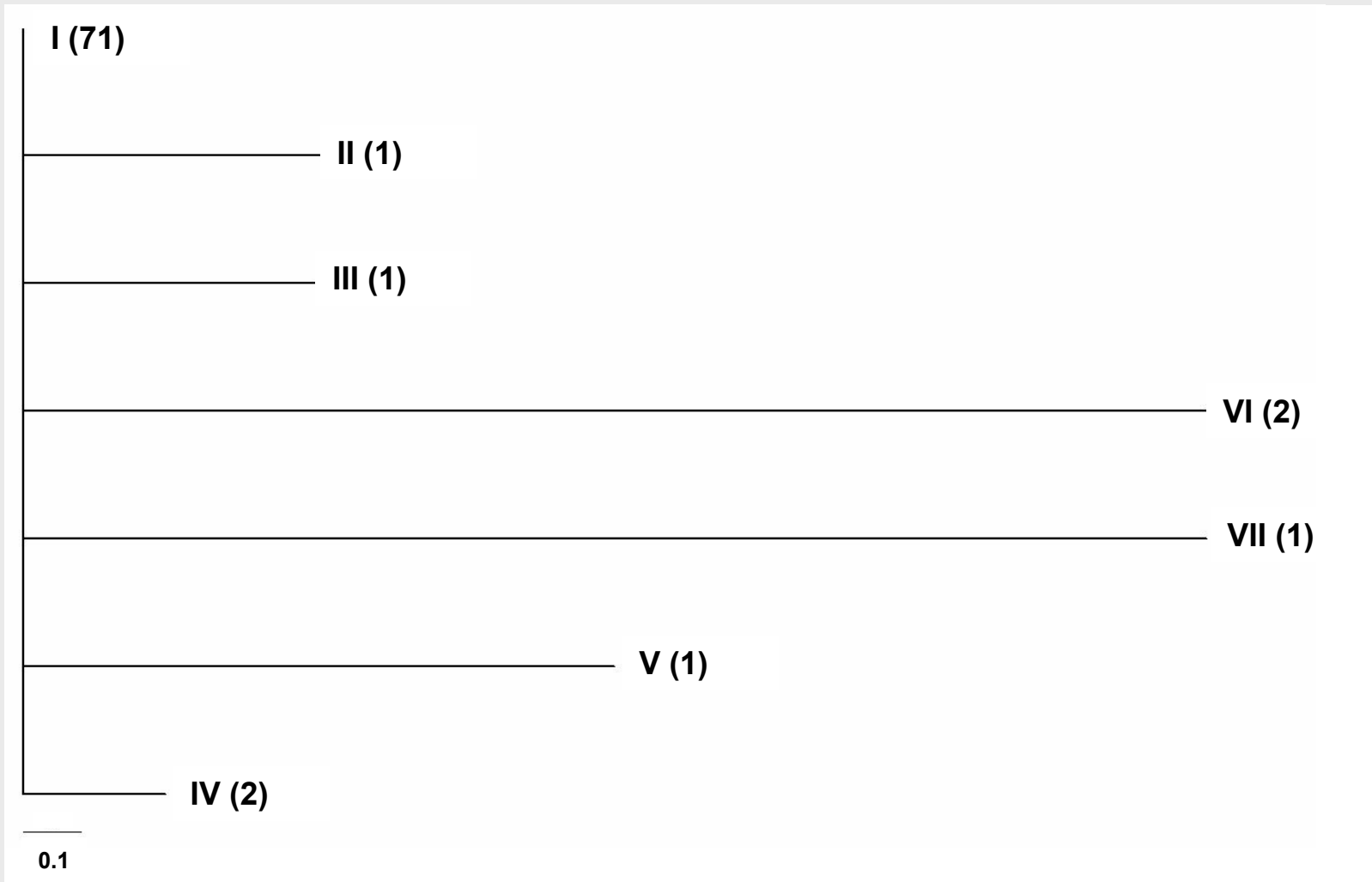
- 314 isolates (2002-2007)
- Protocol:
  - Screening of isolates with 6 selected SSR markers:
    - ✓ 3 from Ivors et al. (2006)
    - ✓ 3 from A. Vercauteren: screening of 132 candidate polymorphic SSR (screening based on polymorphisms within selected EU isolates)
  - Confirmation of polymorphic status of isolates

# Results AFLP

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- 5 primer pairs → 222 fragments → 13 polymorphic
- 71 isolates (90%) belong to a single genotype
- 7 genotypes identified
- no correlation between genotype and place, year





Neighbour-joining cluster analysis

# Results microsatellites

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- 5 out of 6 EU-polymorphic primer pairs show polymorphisms within BE isolates
- 233 out of 314 isolates (74%) belong to one genotype
- 26 genotypes identified

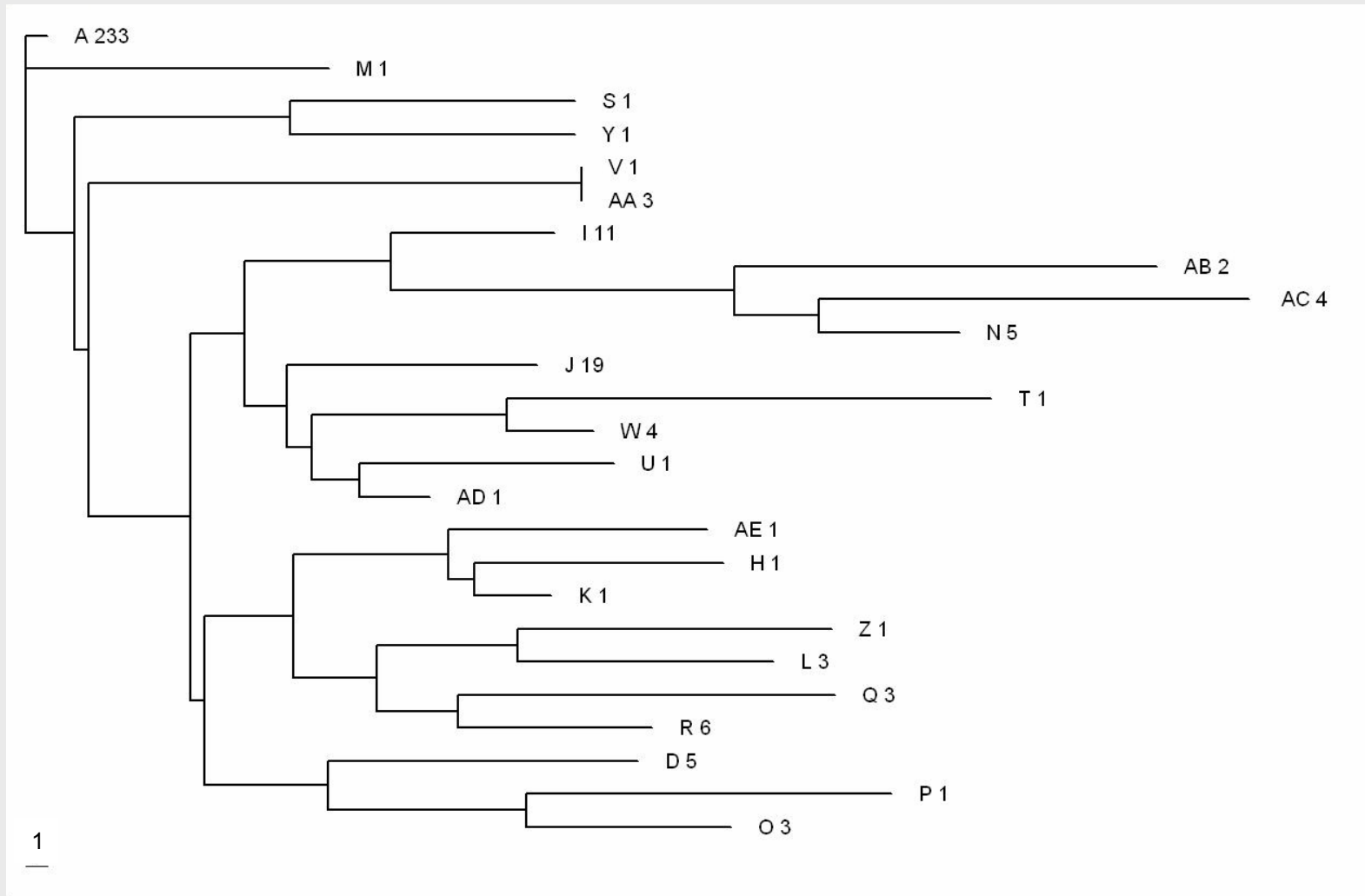
# Microsatellite primer pair code

Ivors et al.

Vercauteren

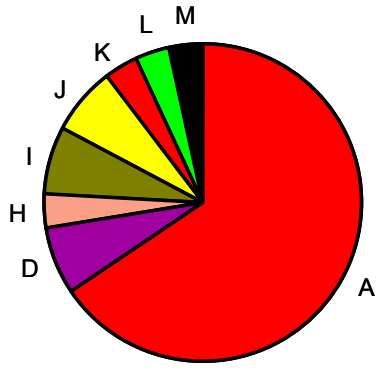
G  
e  
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	18	64	82		2199				512
A	262	398	137	242	186	202	217	265	156
D	262	400	137	242	186	202	217	265	156
H	262	398	137	248	186	202	217	265	156
I	262	398	137	242	186	202	217	249	156
J	262	398	137	242	186	202	217	260	156
K	262	398	137	238	186	202	217	265	156
L	262	398	137	244	186	202	217	260	156
M	262	398	137	242	-	-	217	265	156
N	262	398	137	242	186	202	212	249	156
O	262	400	137	242	186	202	217	260	156
P	262	400	137	244	186	202	217	260	156
Q	262	398	137	244	186	202	217	270	156
R	262	398	137	244	186	202	217	265	156
S	262	398	137	242	186	202	217	265	167
T	262	398	133	242	186	202	217	270	156
U	262	398	137	242	186	202	217	254	156
V	260	398	137	242	186	202	222	265	156
W	262	398	137	242	186	202	217	270	156
X	262	398	137	242	186	202	212	265	156
Y	262	398	137	242	186	202	217	265	171
Z	262	398	137	228	186	202	217	260	156
AA	262	398	137	242	186	202	222	265	156
AB	262	398	137	246	186	202	212	249	156
AC	262	402	137	242	186	202	212	249	156
AD	262	398	137	242	186	202	217	243	156
AE	262	398	137	246	186	202	217	265	156

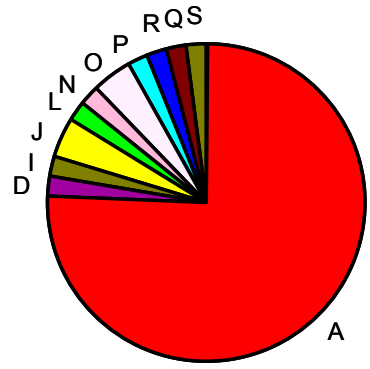


Neighbour-joining cluster analysis

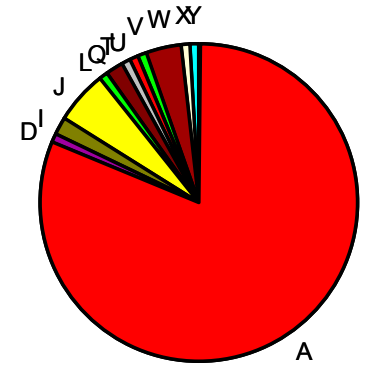
# Genotype composition of Belgian *P. ramorum* population



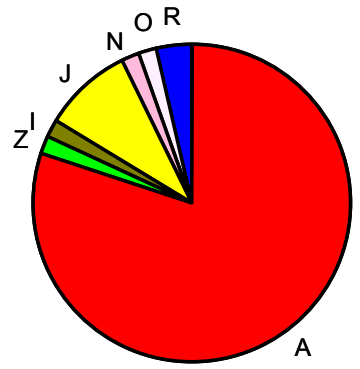
**2002 (29)**



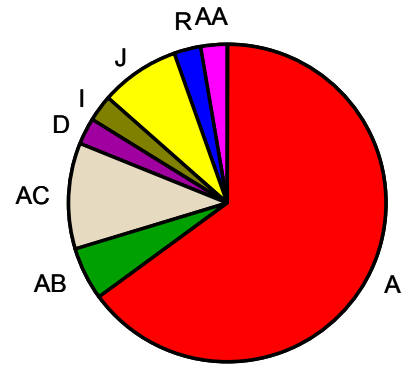
**2003 (49)**



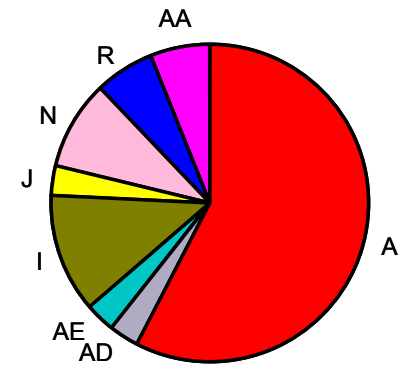
**2004 (111)**



**2005 (55)**



**2006 (37)**



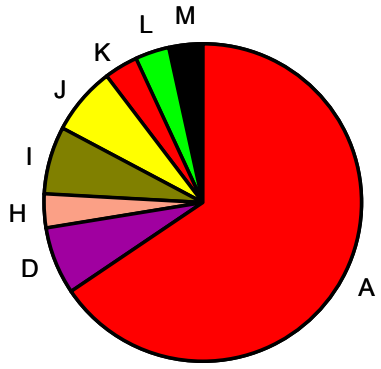
**2007 (33)**

# Resistance to fungicide metalaxyl-M

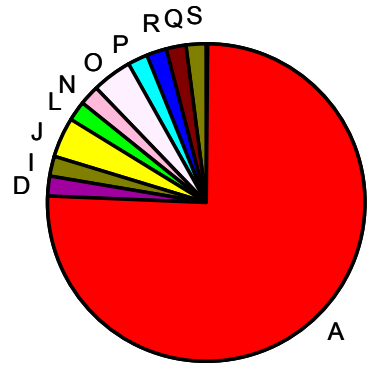
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Year	Percentage resistant isolates
2002	29
2003	17
2004	56
2005	95
2006	84
2007	59

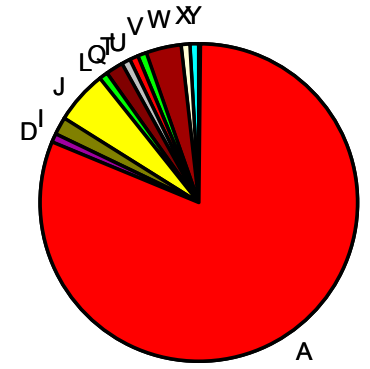
# Genotype composition of Belgian *P. ramorum* population



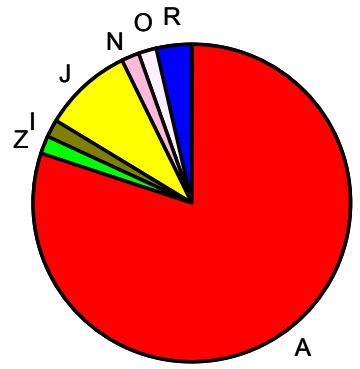
**2002 (29)**



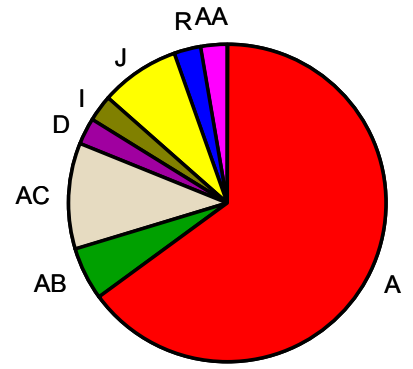
**2003 (49)**



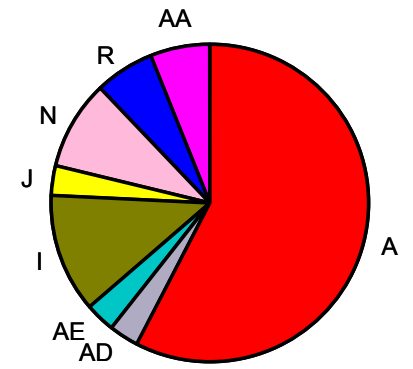
**2004 (111)**



**2005 (55)**

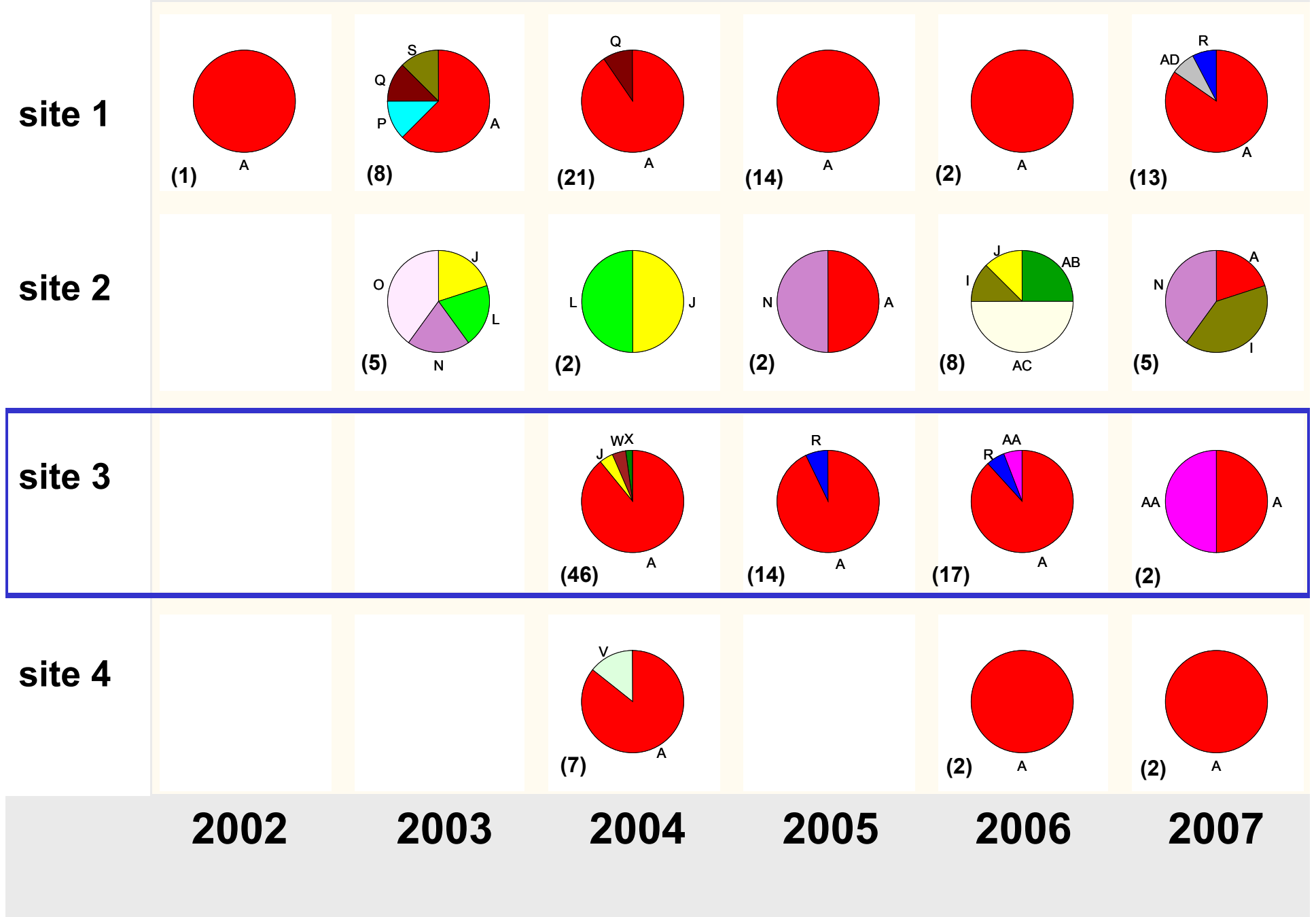


**2006 (37)**



**2007 (33)**

# Local occurrence of genotypes





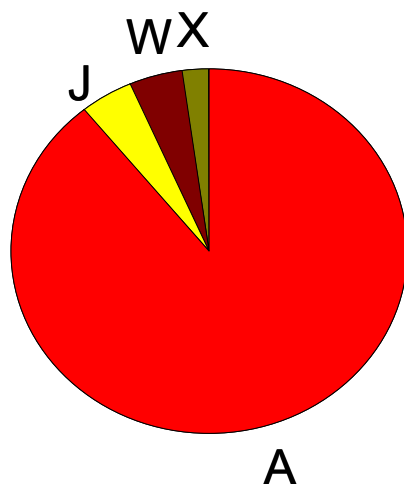


2004-2005: 1399 samples were collected → 331 *P. ramorum* isolates found

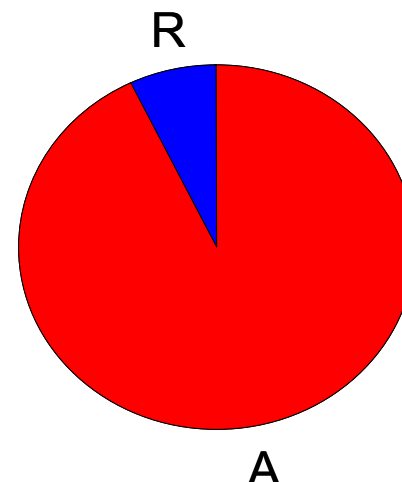
2004

2005

Federal survey

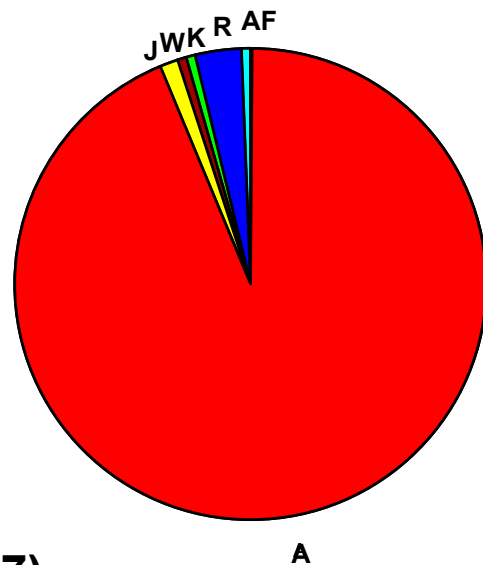


(46)

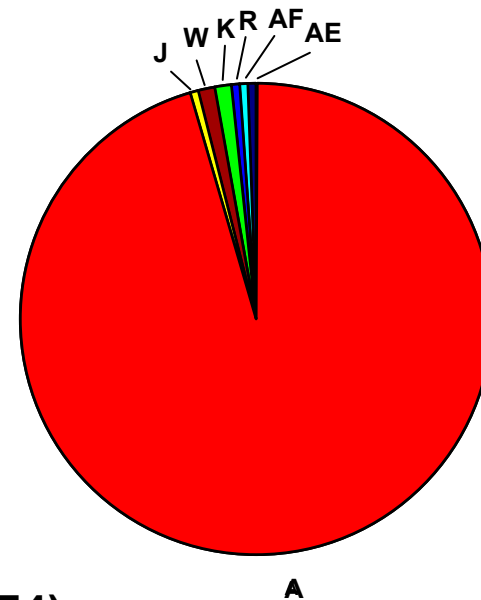


(14)

Project

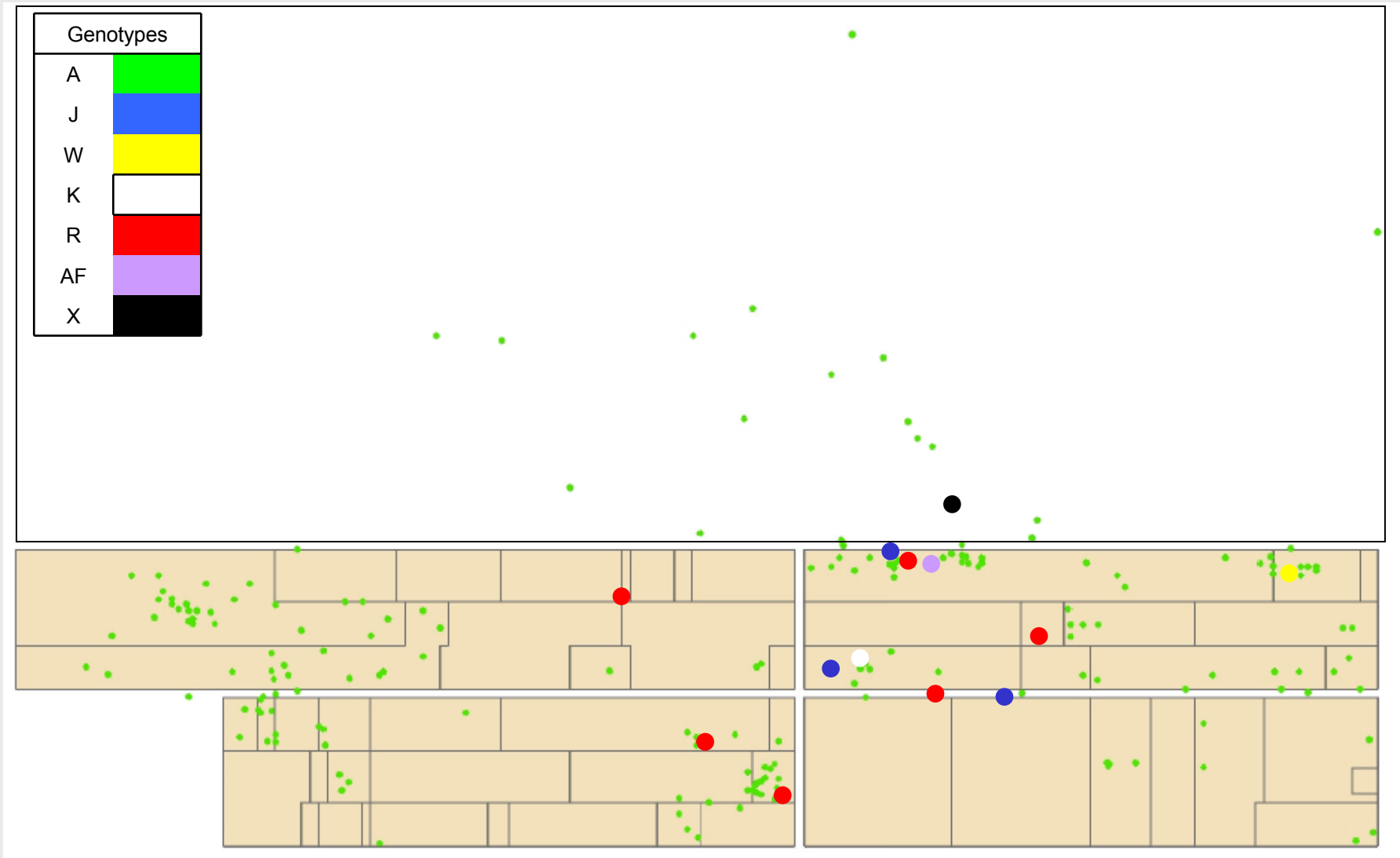


(157)



(174)

# Field surveyed in 2004



# Conclusions

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- Multiple *P. ramorum* genotypes in Belgium (based on AFLP and SSLP)
- One genotype is dominant in Belgium (AFLP: 90% – SSLP: 74%)
  - founder genotype
- Other genotypes (6/79 isolates with AFLP – 25/314 isolates with SSLP)
  - mostly single isolates
- Evolution: some genotypes disappear, new genotypes arise
- Bottleneck in 2004-2005 : metalaxyl-M resistance
- Local specialization: 15/26 genotypes found at single nurseries
- Application to study of epidemiology → dispersal of genotypes in a field

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