

Resistances to the tobacco cyst nematode complex in *Nicotiana*



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The tobacco cyst nematode complex

→ Host spectrum: Solanaceous plants

- ✓ Tomato, egg-plant, tobacco, solanum sp.

→ Affects yield & leaf quality of tobacco

→ Link with other diseases (*Fusarium* wilt)

→ Build up of populations in soils

- ✓ Short rotations of susceptible hosts

→ More difficult to detect than root-knot nematodes

- ✓ Mature cysts (females) turn brown and detach from roots
- ✓ No deformation of roots
- ✓ Likely under-estimated problem

The tobacco cyst nematode complex

→ *Globodera tabacum*

- ✓ *Globodera tabacum tabacum* GTT
 - USA: Connecticut, Massachusetts
 - Bulgaria, Italy, France...
- ✓ *Globodera tabacum solanacearum* GTS
 - USA: Virginia
 - Closely related populations in Mexico
- ✓ *Globodera tabacum virginiae* GTV
 - USA: Virginia

→ Cyst nematodes probably present in many tobacco producing countries

- Spain, Argentina, etc...

Material and methods

→ Greenhouse test (INRA Rennes)

- ✓ 14 well characterized *Globodera* populations
 - 5 plants / population in 1000 cm³ pots
- ✓ Cysts in nylon bags → 10 J2 / g of soil.
- ✓ 120 days after planting, cysts extracted (Kurt elutriator) and counted.

→ Field tests (Tabac Garonne Adour)

- ✓ 2004, 2005 & 2006: same farm, Virazeil, Lot-et-Garonne, Fr.
- ✓ Highly infested with GT but also some *Meloidogyne*
 - Likely GTT (French GT populations from adjacent localities identified as GTT, Marché 1997)
- ✓ Cyst count on roots (2') or notation
 - 6 to 15 plants / entry depending on trial.

M&M, *Globodera* populations in greenhouse test

Population name	Location / code (country)	host	Identification **
Santa Ana	Santa Ana (ME)	wild	GM
Tlaxcala	Tlaxcala (ME)	<i>Solanum</i>	
GM4 *	Huamanda (ME)		GTV
GM5 *	Huamanda (ME)		GM
GM3 *	(ME)		GTT
GV2 *	(USA)		tobacco
GV3 *	Crutchlow (USA)		GTV
GV1 *	(USA)		GTS
GS2 *	(USA)		
GS3 *	(USA)		
GS1 *	Watkins (USA)		
Connecticut *	(USA)		GTT
Aiguillon	Aiguillon (F)		
Agen	Agen (F)		

* Classified by Miller, Virginia University and modified (RAPD, AFLP) (Thiéry et al 1997, Marché et al. 2001)

* Populations from L. I. Miller collection (University of Virginia, USA).

** GM: *Globodera "mexicana"*.

Results, greenhouse test (INRA Rennes)

Good correspondence host spectrum / classification from molecular data

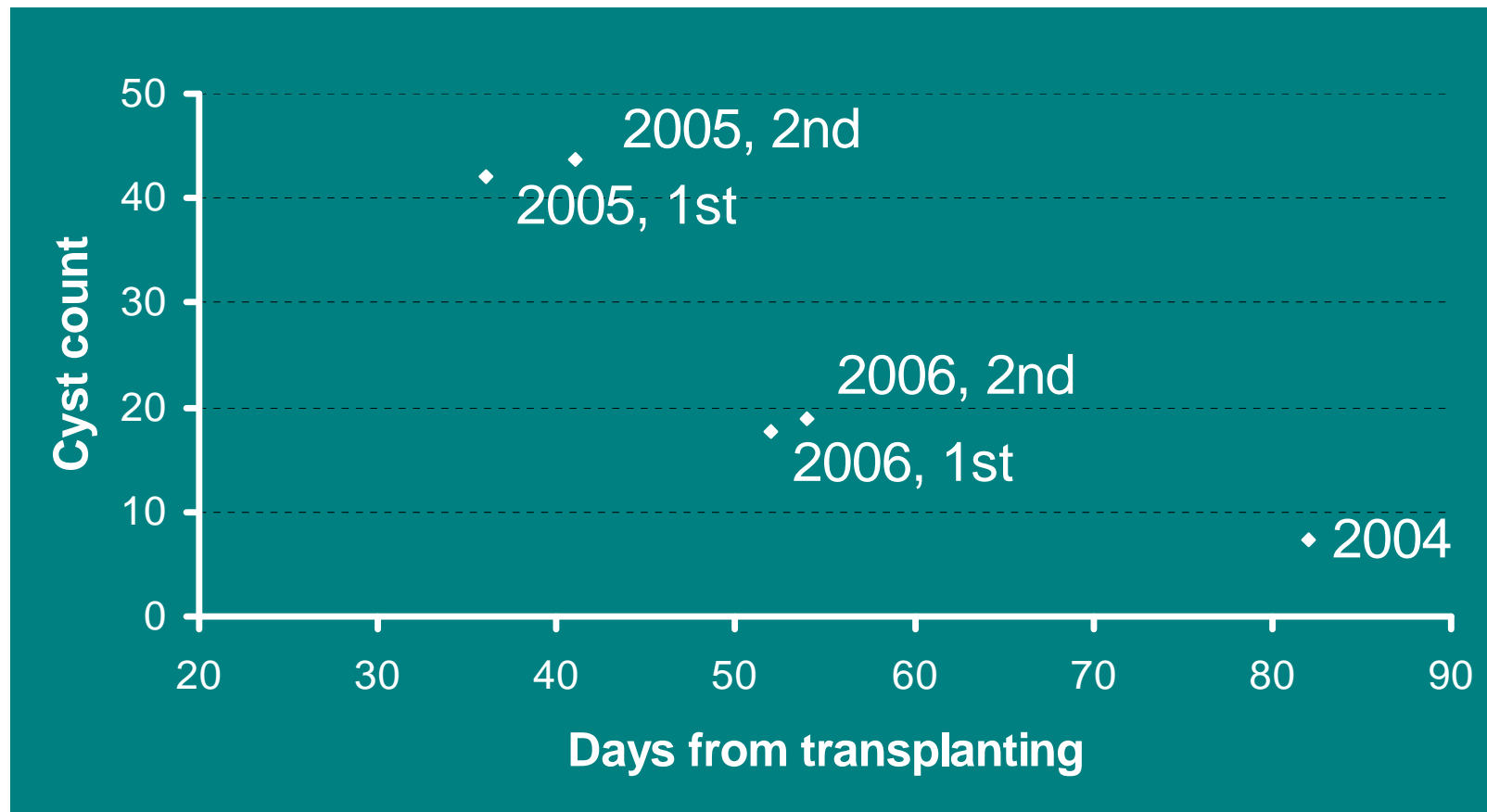
Mean cyst number		Host						Non host				
		<i>bigelovi</i>	<i>sylvestris</i>	<i>debneyi</i>	Samsun	Cocker 254	PBD6	<i>benthamiana</i>	<i>longiflora</i>	<i>repanda</i> x	<i>sylvestris</i>	<i>repanda</i>
Population	Group											
Santa Ana, Tlaxcala, GM5	GM	48	1	1	0	2	0	0	0	0	0	0
GM3, Connecticut	GTT (USA)	44	140	40	85	12	0	2	0	0	0	0
Aiguillon, Agen	GTT (Fr)	38	21	25	43	2	22	1	0	0	0	0
GM4, GV2, GV3	GTV	40	74	16	35	17	8	0	7	0	0	0
GS1, GS2, GS3, GV1	GTS	61	145	52	172	142	6	12	0	0	0	0

Resistance of *N. repanda* and *N. longiflora* effective against GTT and GTS from different origins,

GTT Fr: specific ability to multiply on PBD6 → ancient presence of GTT ?

Results, field tests

→ Highest cyst counts 40 days after transplanting



Results, field tests, cultivar screening

2004

Cultivar		Mean count (9 plants)	NK group **
Coker 371	Ph	1,3	a
Burley 64	WF res	3,9	b
Burley 21	WF res	4,6	b
K 326		6,9	bc
Kutsaga 110	WF res	7,7	bc
TN 90	WF res	7,9	bc
NC 567		13,3	bc
K 394		12,7	c

** based on log. transformed data

Ph: from *N. plumbaginifolia* origin, res. to PPN race 0.
WF res: from *N. longiflora* origin, Wildfire res.

2006

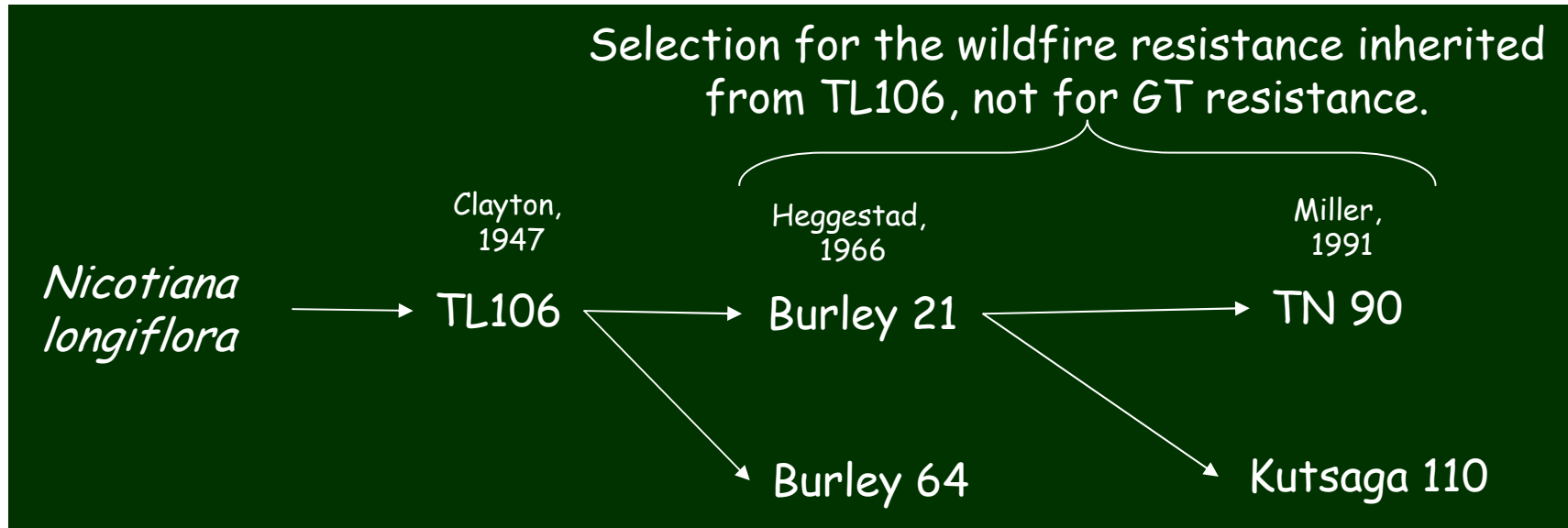
Cultivar		average class *	NK groups
TL 106	WF res	0	a
RGH 51	Ph	0,2	ab
NC 71	Ph	0,3	ab
KYR **		0,8	abc
Coker 254		1	bc
K394		1,3	c
PBD6		1,7	c
Hicks		2,5	d
TN 90	WF res	2,5	d
Bright Cospaia		2,7	d

* 0: none, 1: 1-3, 2: 3-10, 3: 11-30, 4: 31-100, 5: > 100 cysts. ** Breeding line.

→ GTT resistance associated with the Ph gene ?

- ✓ Also observed in Virginia (USA) with GTS (Johnson C.S. 2001)

N. longiflora resistance factors in TL 106



GTS Virginia	R		R	(Hayes et al. 1997)	R
GT Virazeil	R	R	Some Res.		S

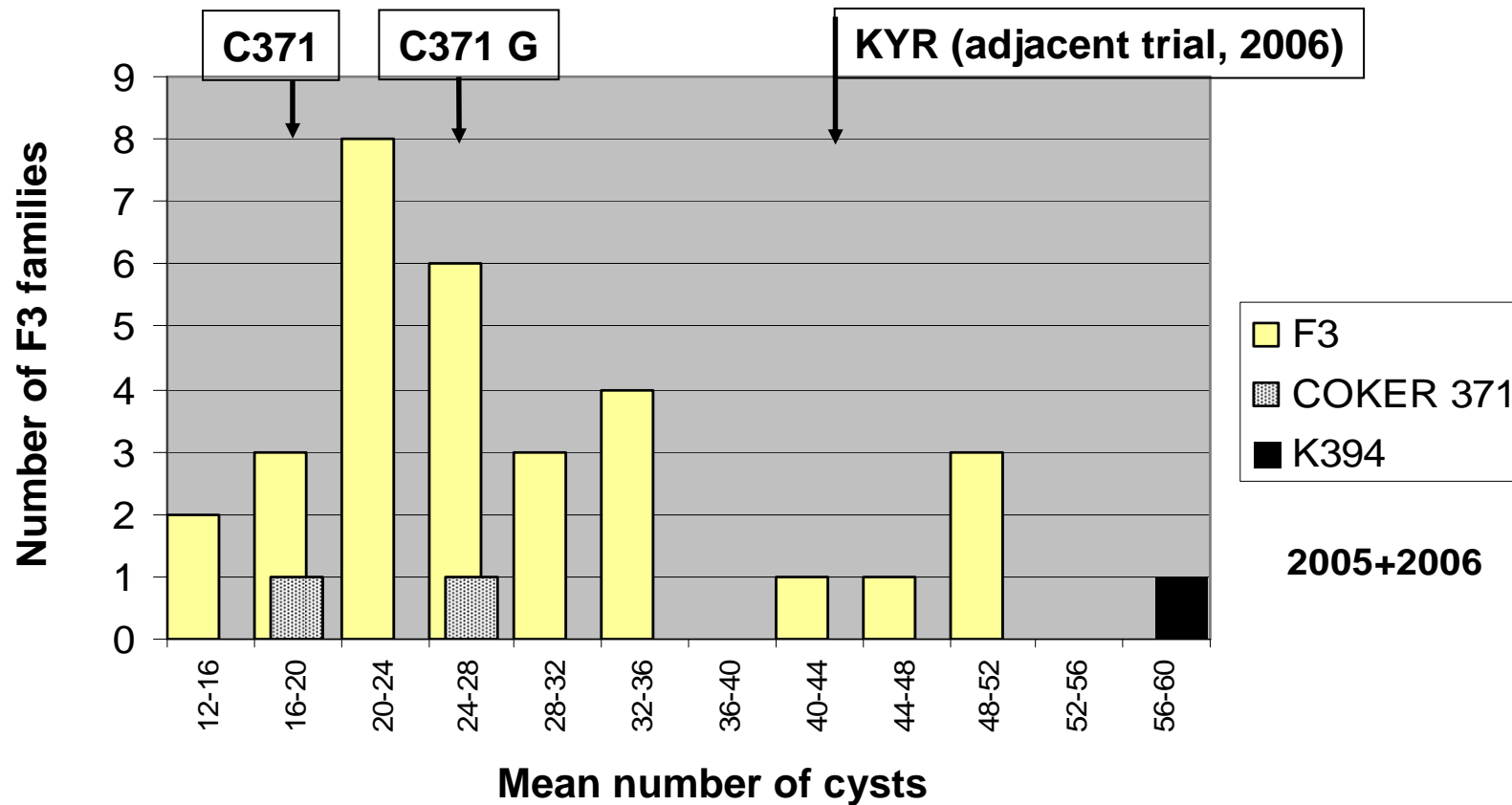
→ **GT Virazeil: loss of *N. longiflora* resistance factors through breeding cycles from TL 106 to TN 90 and Kutsaga 110 ?**

✓ Factors not required for resistance to GTS, Virginia ?

Inheritance of the resistance from Coker 371 Gold

→ F3 families from Coker 371 Gold x KYR

✓ KYR: breeding line with K326 parentage.



Inheritance of the resistance from Coker 371 Gold

→ F3 families from Coker 371 Gold x KYR

- ✓ The bimodal distribution fits with the hypothesis of one dominant gene from Coker 371 Gold conferring partial resistance to GT Virazeil.



Conclusions

→ *Nicotianae*

- ✓ Widest resistance spectrum: *N. repanda*
 - Dominant factors (*N. repanda* x *sylvestris*)
- ✓ *N. longiflora* res. to every GTS and GTT populations.
- ✓ *N. bigelovi* is a host for all GT and GM populations.

→ *N. tabacum*.

- ✓ Introgressed resistance factors efficient against GT Virazeil
 - from *N. plumbaginifolia* (in Coker 371 Gold).
 - Likely one dominant gene.
 - Associated with Ph.
 - from *N. longiflora* (in TL 106).
 - Loss of some of these factors through breeding cycles when selection for wildfire resistance ?