


THE VALIDATION AND USE OF MARKER-ASSISTED SELECTION IN NS WHEAT BREEDING PROGRAM



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From 1953 up to date **372 varieties** of small grains
have been developed at IFVC – Novi Sad:

winter wheat	230
spring wheat	31
winter barley	46
spring barley	41
winter durum	2
spring durum	2
winter oat	4
spring oat	4
winter triticale	11
winter ray	1

42 of which are registered abroad:

Hungary
Slovakia
Spain
Russia
Ukraine
Romania
Greece
Canada
Croatia
(EU) ...



Conventional breeding !

Two countries story

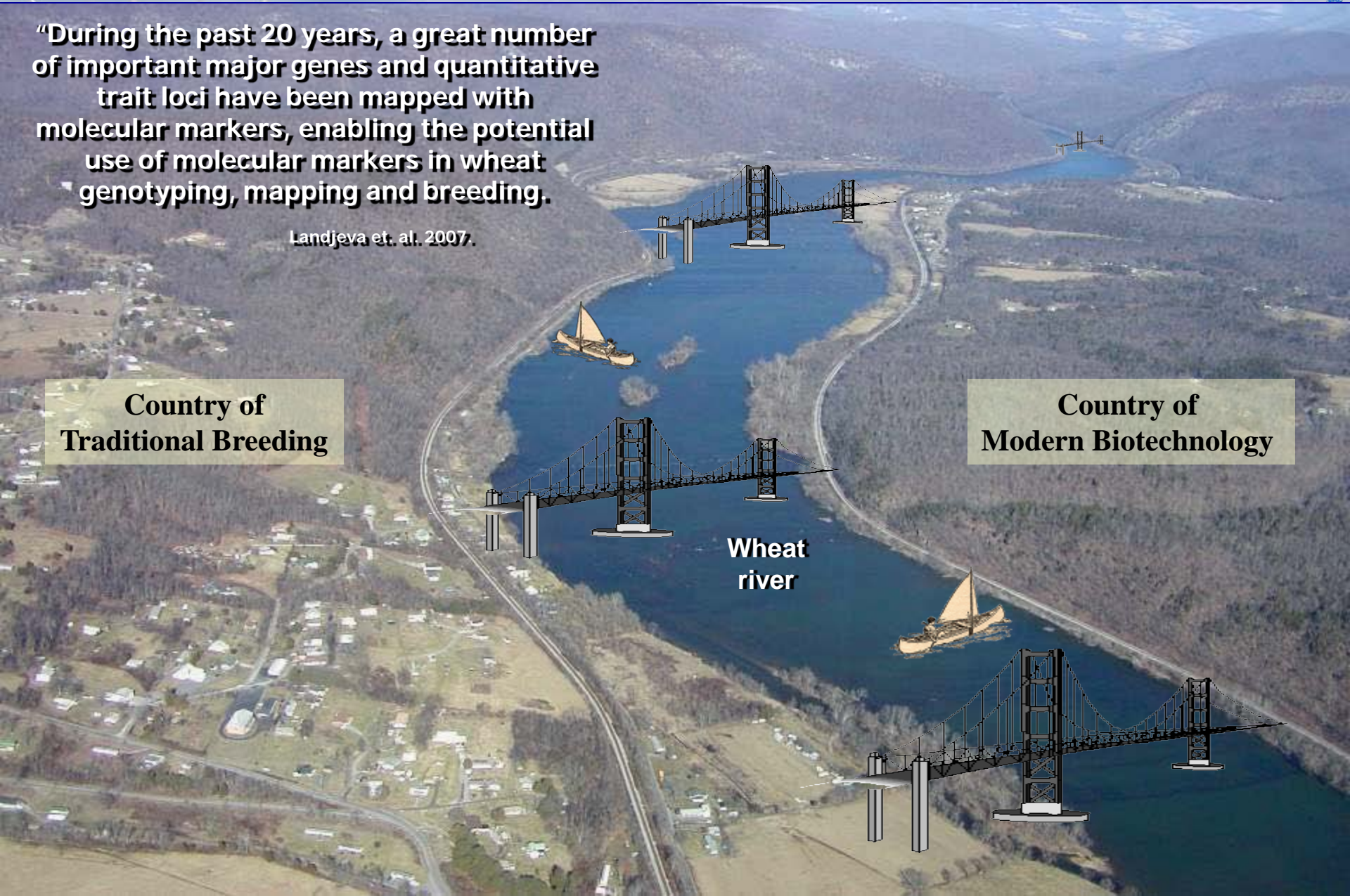
“During the past 20 years, a great number of important major genes and quantitative trait loci have been mapped with molecular markers, enabling the potential use of molecular markers in wheat genotyping, mapping and breeding.

Landjeva et. al. 2007.

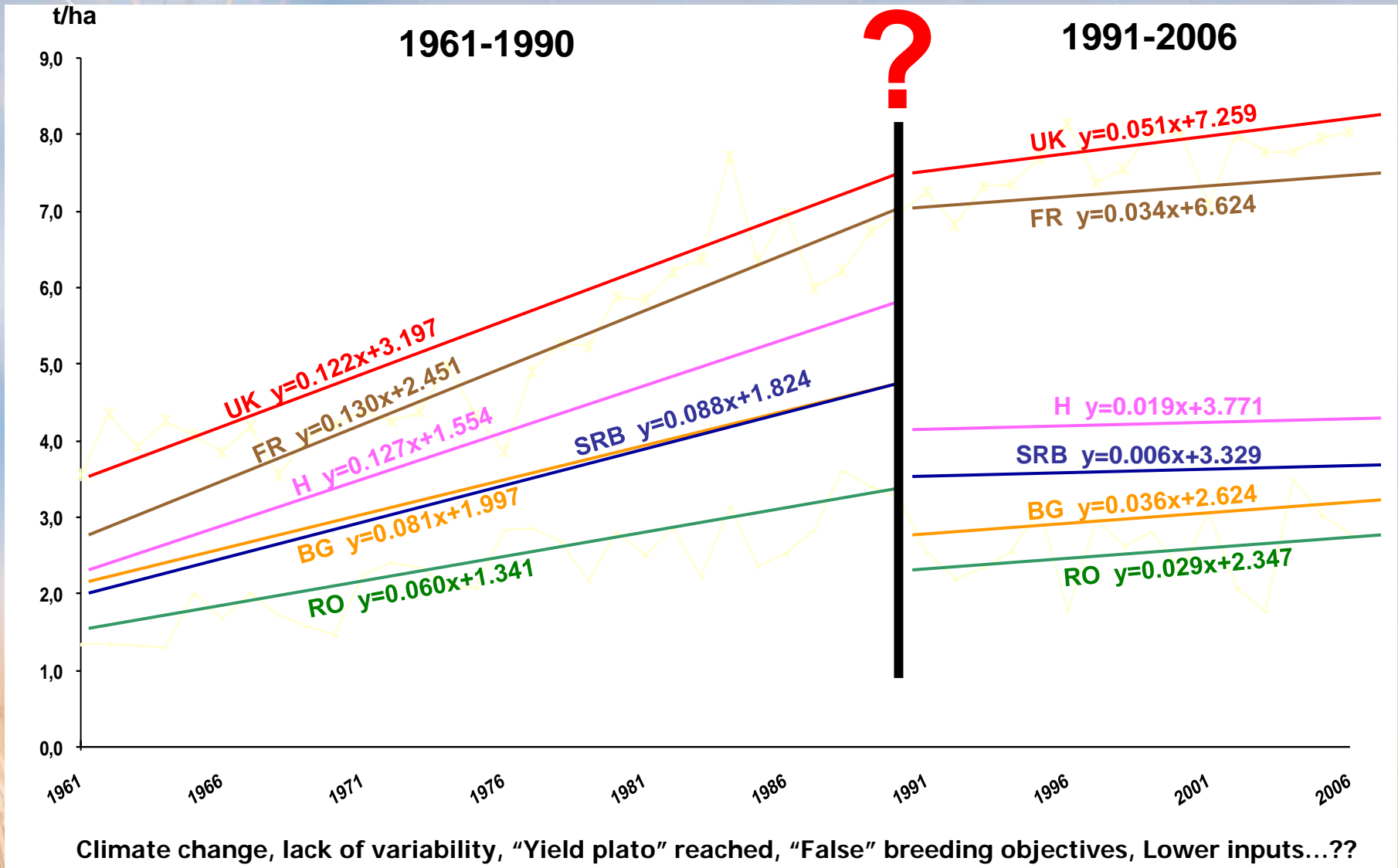
**Country of
Traditional Breeding**

**Country of
Modern Biotechnology**

**Wheat
river**

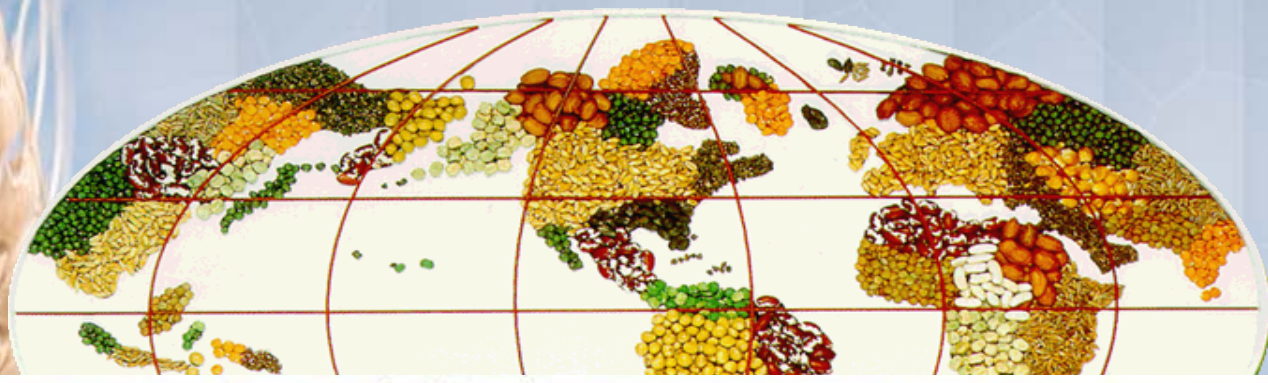


Wheat yields in UK, FR, HU, RO, BG and RS in 1961-2006 period

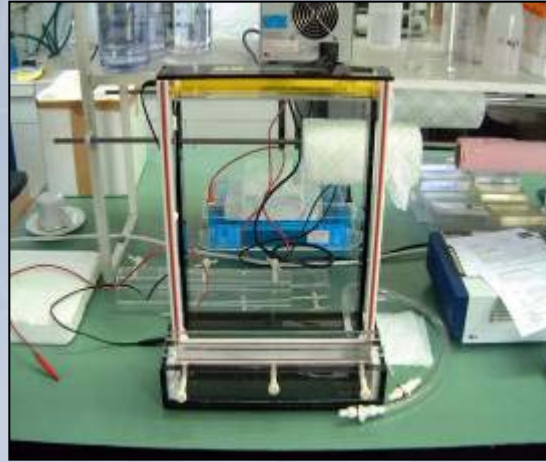


For all breeders, the most promising and challenging task is the possible use of molecular markers to enhance the selection of superior genotypes for traits that are:

1. **difficult or impossible to select phenotypically**
2. **subject to high environmental error**
3. **expensive to asses.**



The Molecular Lab was established in year 2001 at IFVC



**REDUCING COSTS AND
INCREASING BREEDING EFFICIENCY !**



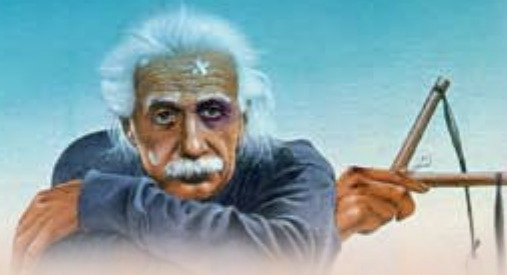
Single cross strategy (parents AxB)

Association study (set of genotypes)



Target trait (gene)

Target trait (gene)



Scientist (with sling)

Scientist (with gun)



ITMI mapping population (Opata x Synthetic)



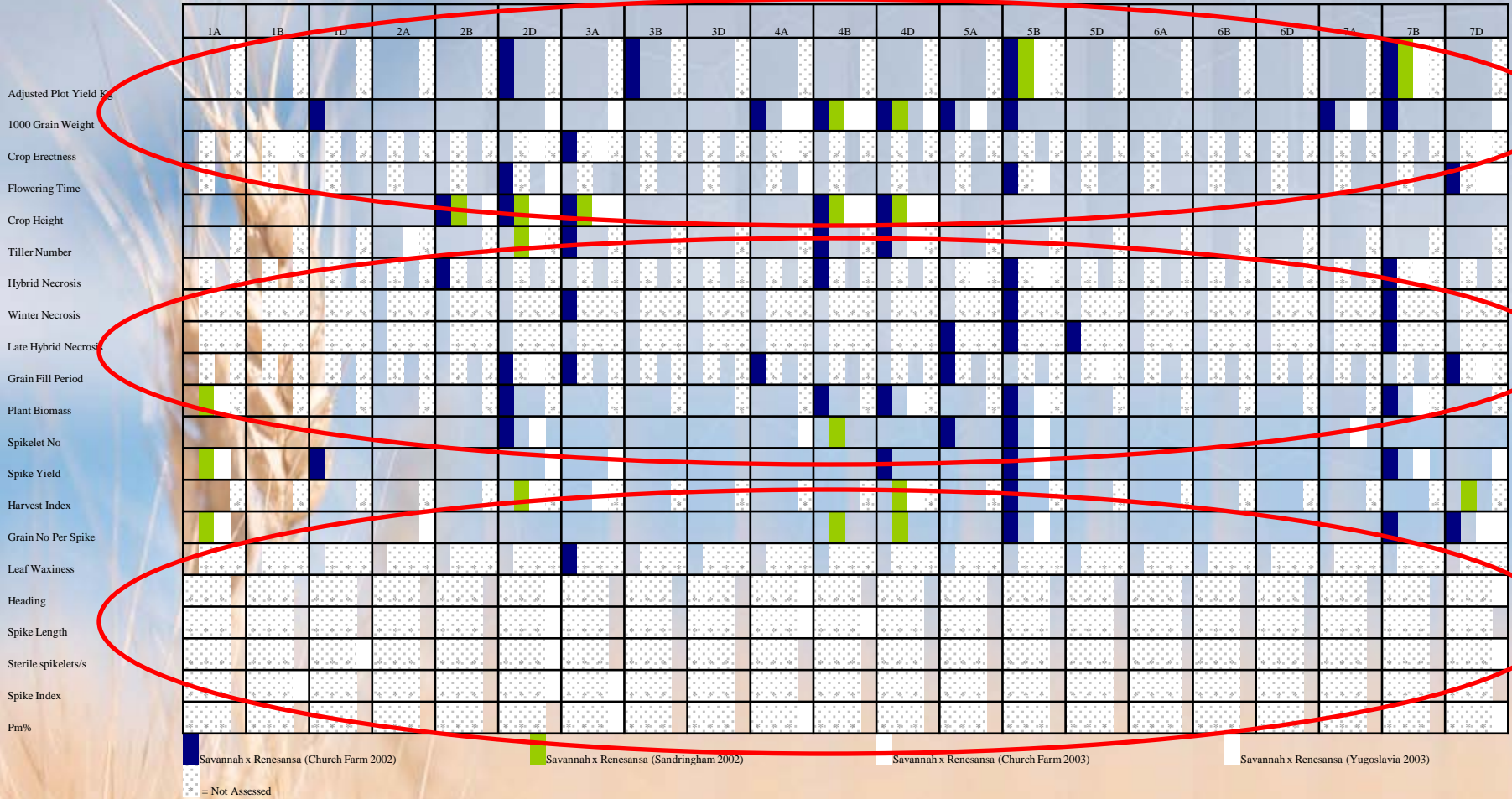
Savanahh / Renesansa DH population (single cross)



Detected significant QTL's in Savannah x Renesansa mapping population



QTL found using marker regression (>0.05) for Savannah x Renesansa comparing sites and years



Where there are only two markers in a linkage group marker regression can not be carried out, in these cases

significant (& interesting) QTL found using single marker analysis have been included.

* - Denotes where no assessment was made



POTENTIAL USES OF MICROSATELLITES IN MARKER – ASSISTED SELECTION (MAS) FOR IMPROVED GRAIN YIELD IN WHEAT



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Genetic (Core) Collection - 603 genotypes – 38 countries





**Pheno data base
(54 traits, 3-8 years)**

26 TRAITS (SRB+UK)

603 GENOTYPES

The “worst” 48

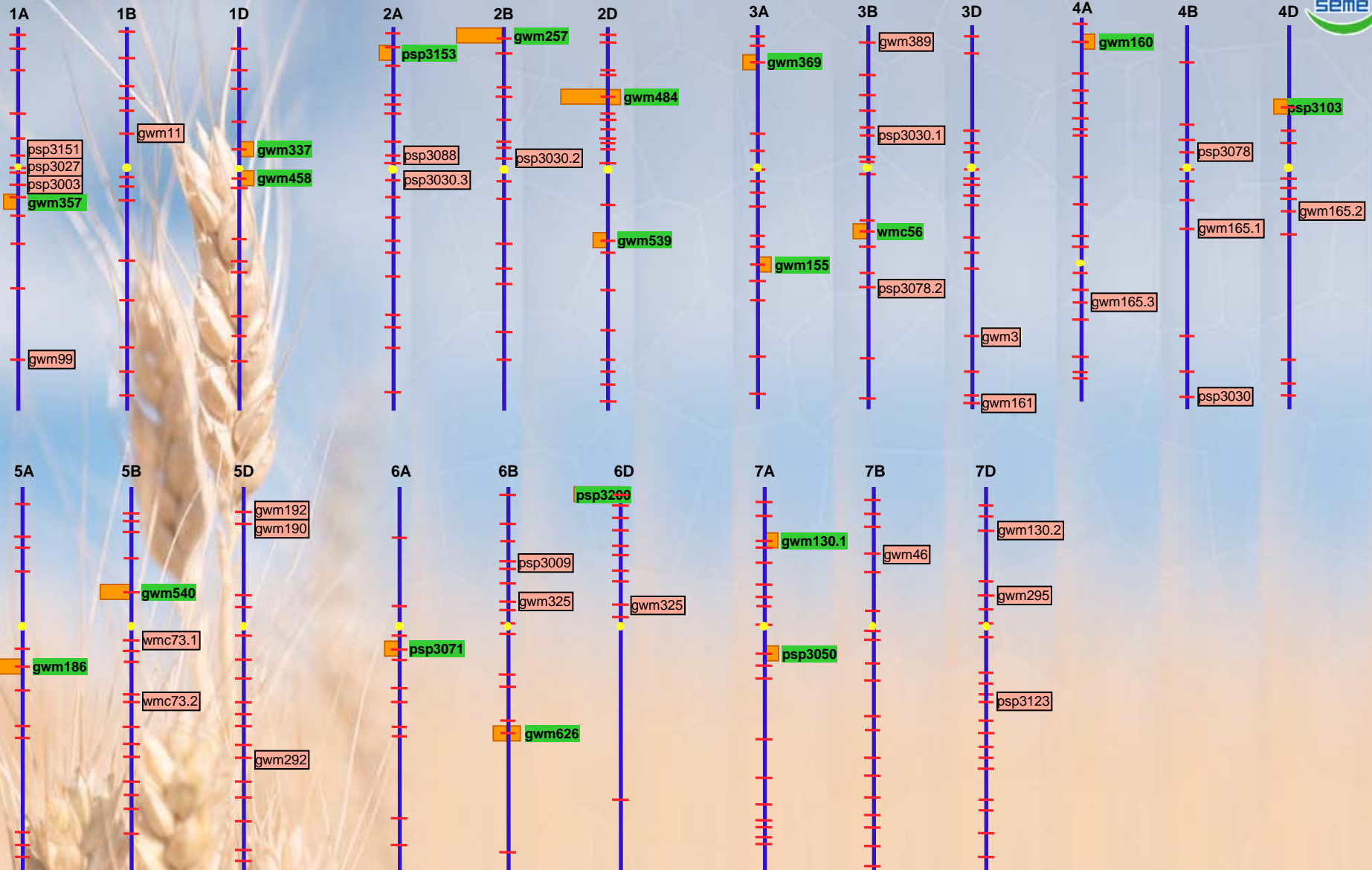
The “best” 48

**96 genotypes representing
overall phenotypic variation**

96 wheat “delegates” for molecular screening

Acciaio	L - 1	Purdue 5392
Ai-bian	Lambriego Inia	Red Coat
Al KanTzao	Lr 10	Renesansa
Ana	Lr 12	Rusalka
Avalon	Magnif 41	Siete Cerros
Bankuty 1205	Mex.17 bb	Saitama 27
BCD 1302/83	Mex.3	Sava
Benni multifloret	Mexico120	Semillia Eligulata
Bezostaya 1	Minister Dwarf	Slavija
Brigand	Mina	Sofija
Cajeme 71	Mironovska 808	Sonalika
Capelle Desprez	Nizija	Suwwon 92
Centurk	Norin 10/Brev.14	Szegedi 768
Ching-Chang 6	Norin 10	Tibet Dwarf
Cook	Nov. Crvena	Timson
Don.polupatuljasta	Nova Banatka	TJB 990-15
Durin	NS 22/92	Tom Thumb
F 4 4687	NS 33/90	Tr. compactum
Florida	NS 46/90	Tr sphaerococcum
Gala	NS 55-25	Triple Dirk B
HAYS 2	NS 559	Triple DirkB (cont.)
Helios	NS 602	Triple Dirk S
Highbury	NS 63-24	UC 65680
Hira	NS 66/92	UPI 301
Holly E	NS 74/95	Vel
Hope	NS 79/90	Vireo “S”
Inia 66	Peking 11	WWMCB 2
INTRO 615	Phoenix	ZG 1011
Ivanka	PKB Krupna	ZG 987/3
Kite	Pobeda	ZG K 238/82
L 1/91	Purd./Loras	ZG K 3/82
L 1A/91	Purdue 39120	ZG K T 159/82

SSRs used for evaluation of the 96 “delegates”



Advanced lines – yield trials



8 t/ha

5 t/ha

9 t/ha

7 t/ha

10 t/ha

8 t/ha

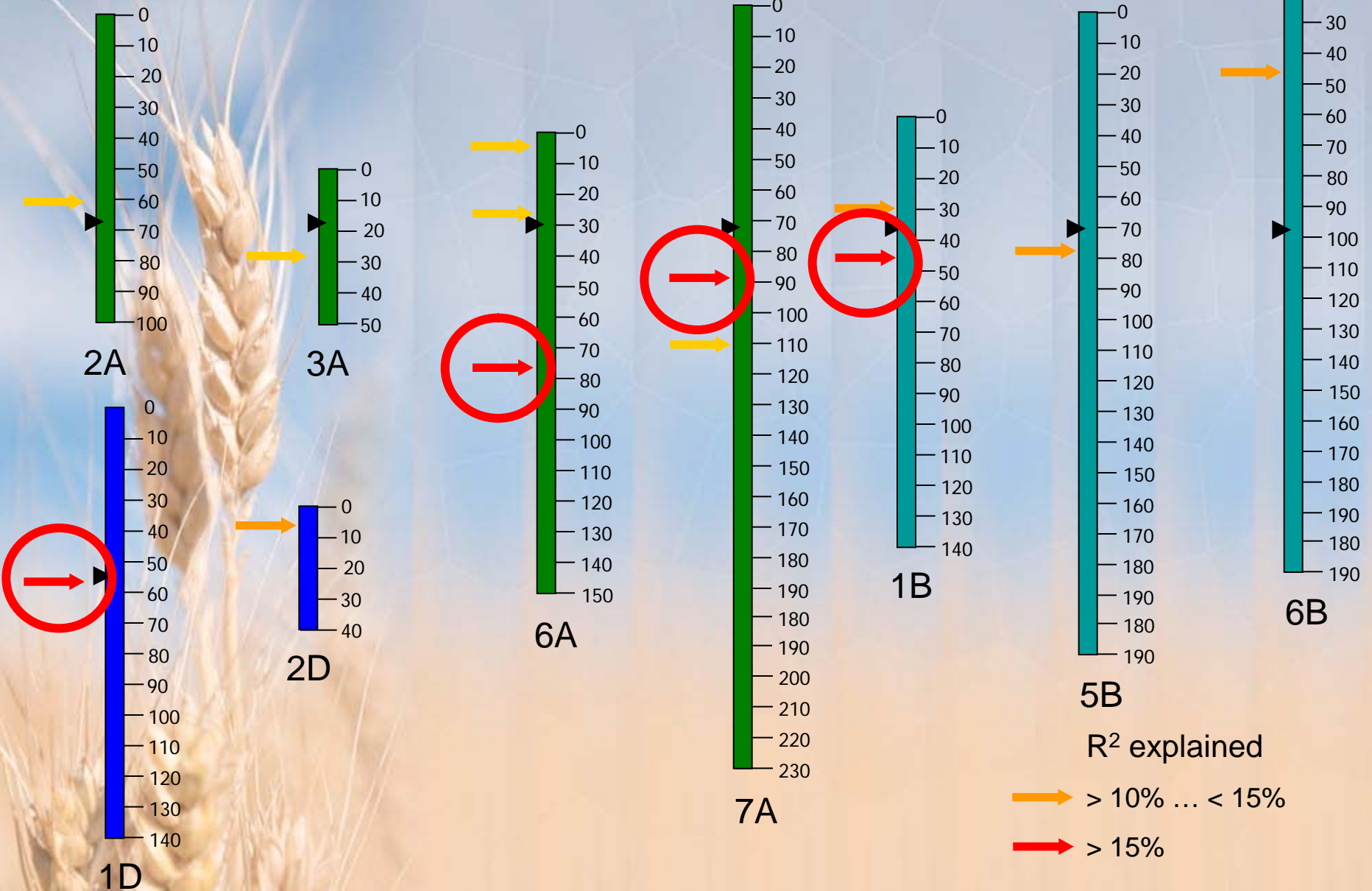
7 t/ha

Distribution of presumed “high-yielding” alleles in the 20 LYG and 20 HYG



Primer															Yield (t/ha) Mean = 7.84	Alleles No. Mean = 8.7	Rht8 locus
Genotype	GWM 46.2	GWM 155	GWM 190	GWM 192.2	GWM 295	GWM 337	GWM 484	GWM 539	GWM 540	PSP 3088	PSP 3103.1	PSP 3103.2	PSP 3153.2	PSP 3200			
NS 66/92															8.91	10	175
NS 79/90															8.88	10	192
Mina															8.41	6	192
NS 559															8.06	8	192
Centurk															8.05	6	200
Pobeda															8.03	14	192
BCD 1302/83															8.01	7	192
NS 33/90															7.88	9	192
Ana															7.83	7	165
NS 55-25															7.77	10	192
NS 46/90															7.70	7	192
Sofija															7.66	9	175
UC 65680															7.66	8	192
Slavija															7.56	9	192
Sava															7.52	7	192
Nizija															7.44	9	192
Renesansa															7.38	10	192
Triple dirk B															7.38	9	null
Nova Banatka															7.32	9	ND
Rusalka															7.26	9	192
<i>Alleles No.</i>	12	7	17	20	14	13	7	16	11	16	8	8	8	17			

Going from DARTs to SSRs !



R^2 explained

→ > 10% ... < 15%

→ > 15%



**We are going further – chasing important QTLs
for the rest of the genotypes from the Core Collection (app.500)
for 20 traits (data - 8 years, 3 field plot reps., on 1,2 m², ..)**



Validation of 20 candidate markers which we believe could be of value for the NS wheat breeding program

(yield and yield components, pathogen resistance, drought and heat stress tolerance, etc.).



1	WMC89	4A
2	WMC31	7D
3	WMC48	4A
4	WMC420	4A
5	gwm295	7D
6	gwm337	1D
7	gwm539	2D
8	gwm428	7D
9	gwm174	5D
10	gwm11	1B



11	gwm194	4D
12	gwm601	4A
13	psp 3071	6A
14	psp3094	7A
15	psp3200	6D
16	psp3103	4D
17	gwm18	1B
18	gwm46	7B
19	gwm95	2A
20	gwm261	2D



- 20 highest yielders from Core Collection 96 set
- 20 lowest yielders from Core Collection 96 set
- 20 lowest yielding advanced lines
- 20 highest yielding advanced lines
- 16 registered and most popular Serbian varieties



PLAN OF ACTIONS - STARTING DATE 15. OCTOBER 2009.

- 1. Routine screening for major genes (Lr, Pm, FHB, Rht, Ppd..)**
- 2. Validation and use of mm for desirable QTLs from biparental populations**
- 3. Validation and use of mm for desirable QTLs from association study (CC)**
- 4. Additional screening for desirable QTLs in 500 genotypes from CC (SSRs)**
- 5. Along with these, new markers of potential value for the NS wheat breeding program will be continually included in the validation process in order to create the best and most up-to-date collection of markers which can be used for MAS.**

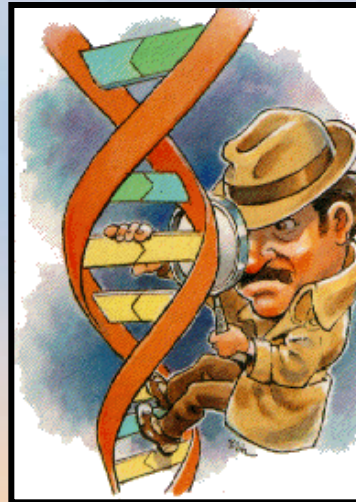


MOLECULAR

Be
Rational
Effective
Enthusiastic
Decisive
Inventive
Natural
Genuine

2009

????



THANKS