

Sources for late blight resistance from Potato Collection in Vavilov' Institute of Plant Industry

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- Manifestation of the late blight resistance in wild tuber-bearing *Solanum* species from Collection of Vavilov' Institute of Plant Industry
- Distribution SCAR-markers for *R*-genes and their aptitude for prediction of late blight resistance in wild potato species
- *Solanum* genotypes promising for breeding program

Race non-specific late blight resistance in wild tuber-bearing *Solanum* species

SEEDLING TEST in the field

Series	Species	Number of accessions (families) tested	Distribution of families on rating scale:			
			1-2	3-4	5-6	7-9
Bulbocastana	<i>S. bulbocastanum</i>	4	1 (1) *	2 (1,6-2,8)	1 (3)	
Longipedicellata	<i>S. stoloniferum</i>	27	6 (0-1)	5 (1-2,6)	6 (0-2,7)	10 (0-2)
Demissa	<i>S. demissum</i>	24	4 (0-1)	4 (0,8-2,3)	5 (0-1)	11 (0-0,5)

* **Number of families** (standard deviation s_x min-max)

Diversity of wild *Solanum* germplasm on relation to disease.

How to find new sources of late blight resistance?

Race non-specific late blight resistance in wild tuber-bearing *Solanum* species

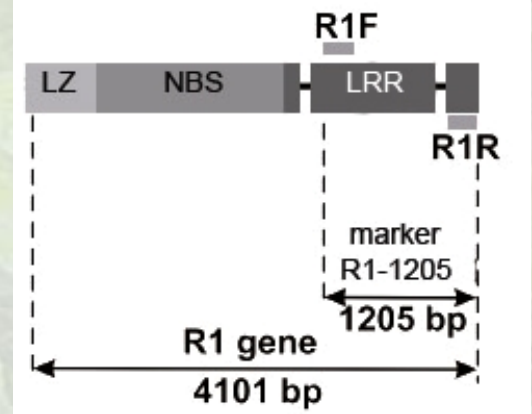
CLONAL PLANTS in the laboratory test

Series	Species	Number of genotypes (accessions) tested	Distribution of genotypes in groups with a resistance rating:		
			2-4	5-7	8-9
Bulbocastana	<i>S. bulbocastanum</i>	26 (19)	4	3	19
Pinnatisecta	<i>S. brachistotrichum</i>	2 (2)	2		
	<i>S. cardiophyllum</i>	6 (6)	2	3	1
	<i>ssp. ehrenbergii</i>	16 (13)	10	2	4
	<i>S. jamesii</i>	12 (10)	3	7	2
	<i>S. pinnatisectum</i>	10 (7)		4	6
	<i>S. stenofillidium</i>	3 (3)	3		
	<i>S. tarnii</i>	3 (3)	1	2	
Polyadenia	<i>S. polyadenium</i>	7 (7)		1	6
Tuberosa	<i>S. verrucosum</i>	12 (9)	7	4	1
Longipedicellata	<i>S. fendleri</i>	3 (3)	3		
	<i>S. hjertingii</i>	5 (4)	3	2	
	<i>S. papita</i>	2 (2)		2	
	<i>S. polytrichon</i>	9 (9)	5	3	1
	<i>S. stoloniferum</i>	33 (27)	18	5	10
Demissa	<i>S. brachycarpum</i>	4 (4)		4	
	<i>S. demissum</i>	26 (23)	3	10	13
	<i>S. hougasii</i>	4 (3)		3	1
Total	18	183 (156)			

Distribution SCAR-markers for *R*-genes across wild potato species

SCAR marker R1-1205 for *R1* gene (*S. demissum*),

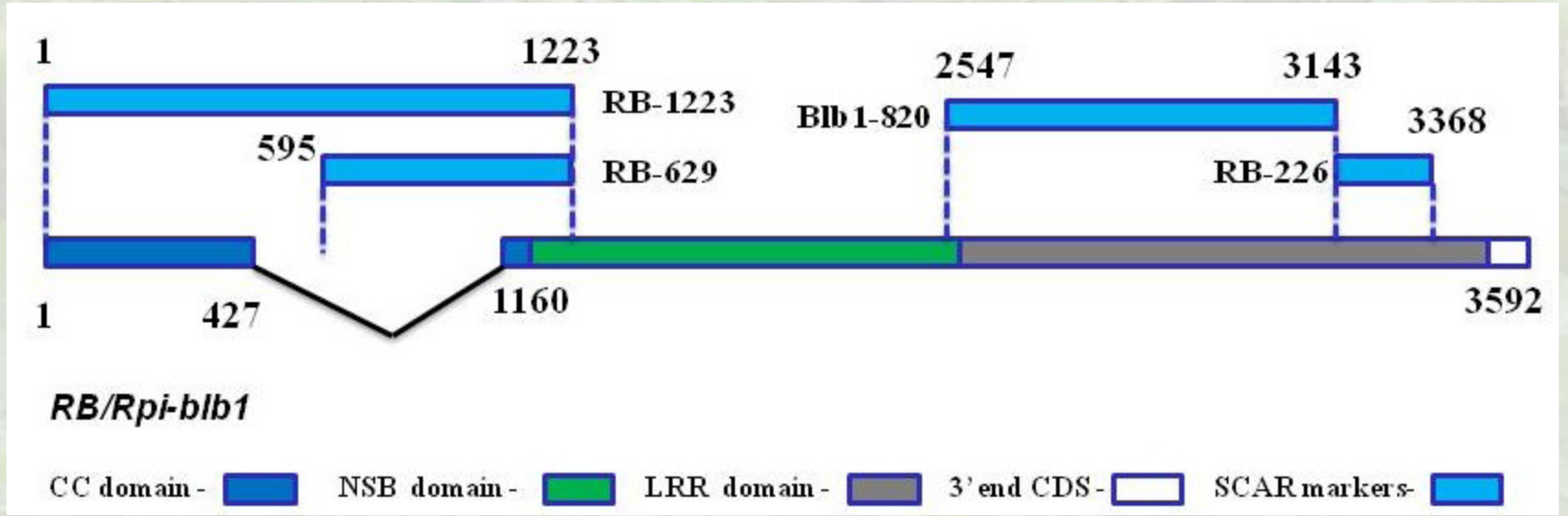
for more details of markers for *R1*- gene see the poster by E. Sokolova et al. at this meeting



Frequency of the marker R1-1205 in 18 wild potato species

Series	Species	Number of genotypes tested	F
Demissa	<i>S. brachycarpum</i>	4	0,75
	<i>S. demissum</i>	36	0,42
	<i>S. hougasii</i>	3	0,66
Longipedicellata	<i>S. polytrichon</i>	12	0,08
	<i>S. stoloniferum</i>	33	0,15
Polyadenia	<i>fen, hjt, pta</i>	10	0
	<i>pld</i>	8	0
Pinnatisecta	<i>bst, jam, pnt, sph, trn, cph, ehr</i>	51	0
Bulbocastana	<i>blb</i>	27	0
Verrucosa	<i>ver</i>	10	0

SCAR markers for *RB/Rpi-blb1* gene (*S. tuberosum*)



for more details of markers for *RB/Rpi-blb1* gene see the poster by O. Fadina et al. at this meeting

Distribution of markers RB-629 and RB-226 (*S. bulbocastanum*) across wild potato species

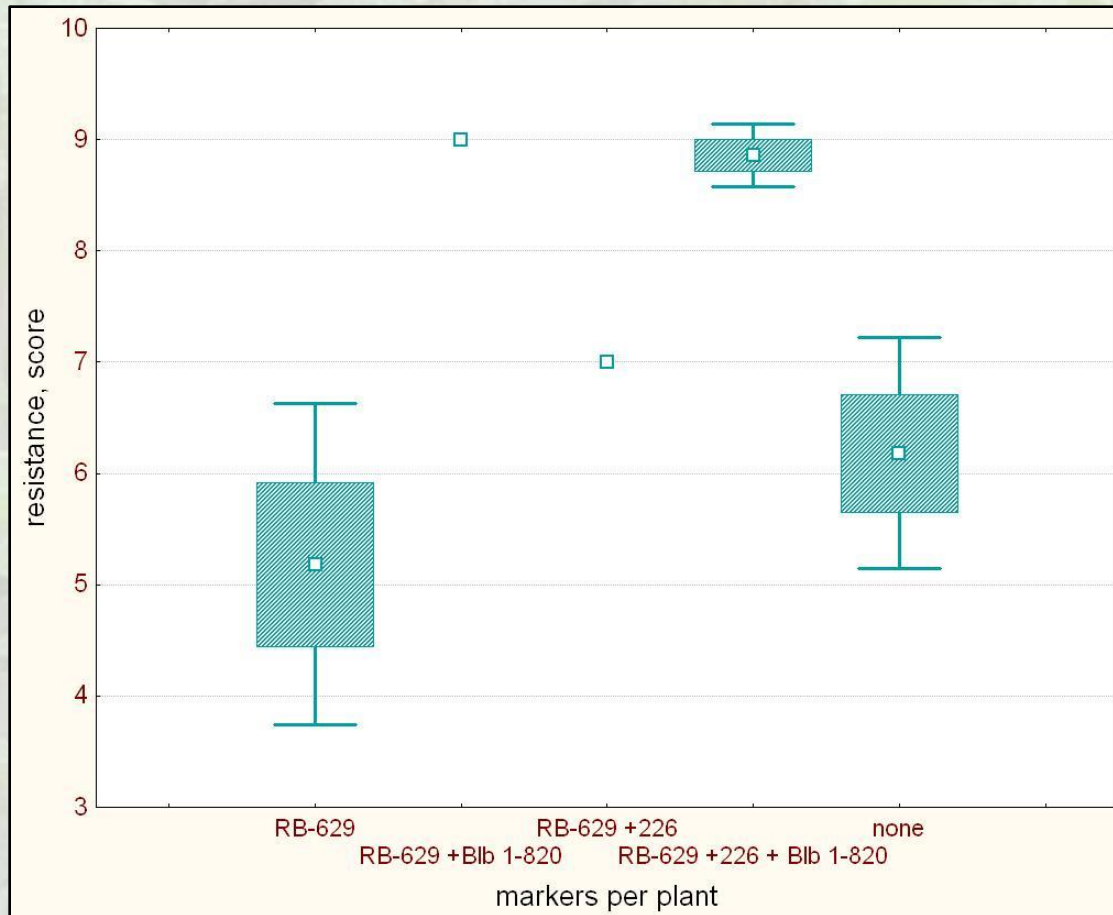
Series	Species	Number of genotypes tested \ frequencies of SCAR-markers:	
		RB-629	RB-226
Bulbocastana	<i>S. bulbocastanum</i>	22 \ 0,68	11 \ 0,45
Pinnatisecta	<i>S. brachistotrichum</i>	2 \ 1,0	n.d.
	<i>S. cardiophyllum</i>	4 \ 0,75	5 \ 0,20
	<i>ssp. ehrenbergii</i>	13 \ 0,31	16 \ 0,06
	<i>S. jamesii</i>	8 \ 0,62	3 \ 0
	<i>S. pinnatisectum</i>	9 \ 0,78	3 \ 0
	<i>S. stenofillidium</i>	3 \ 0,67	2 \ 0
Polyadenia	<i>S. polyadenium</i>	5 \ 0,40	1 \ 0
Tuberosa	<i>S. verrucosum</i>	6 \ 0,17	4 \ 0,25
Longipedicellata	<i>S. fendleri</i>	3 \ 1,0	n.d.
	<i>S. hjertingii</i>	6 \ 0,67	1 \ 0
	<i>S. papita</i>	2 \ 0,50	n.d.
	<i>S. polytrichon</i>	10 \ 0,90	n.d.
	<i>S. stoloniferum</i>	24 \ 0,50	9 \ 0,33
Demissa	<i>S. brachycarpum</i>	4 \ 0	n.d.
	<i>S. demissum</i>	20 \ 0,35	3 \ 0,33
	<i>S. hougasii</i>	5 \ 0,40	2 \ 0,50

Association of potato late blight resistance with the presence of markers of the *RB/Rpi-blb1* gene

	RB-629	RB-226	Blb 1 - 820	RB-226+ Blb 1-820
resistance, score	0,13	0,43	0,51	0,49
markers per plant	0,95	0,74	0,74	0,78
RB-629		0,51	0,51	0,54
RB-226			0,85	

Spearman correlation coefficient is calculated for $n=41$, $p < 0,05$

Segregation of *Solanum* genotypes according to results of phytopathological and marker analysis for ***RB/Rpi-blb1*** gene



ANOVA (n=41): each of three groups of genotypes of wild potato species are different in resistance to late blight significantly: **F=3,24, p=0,002.**

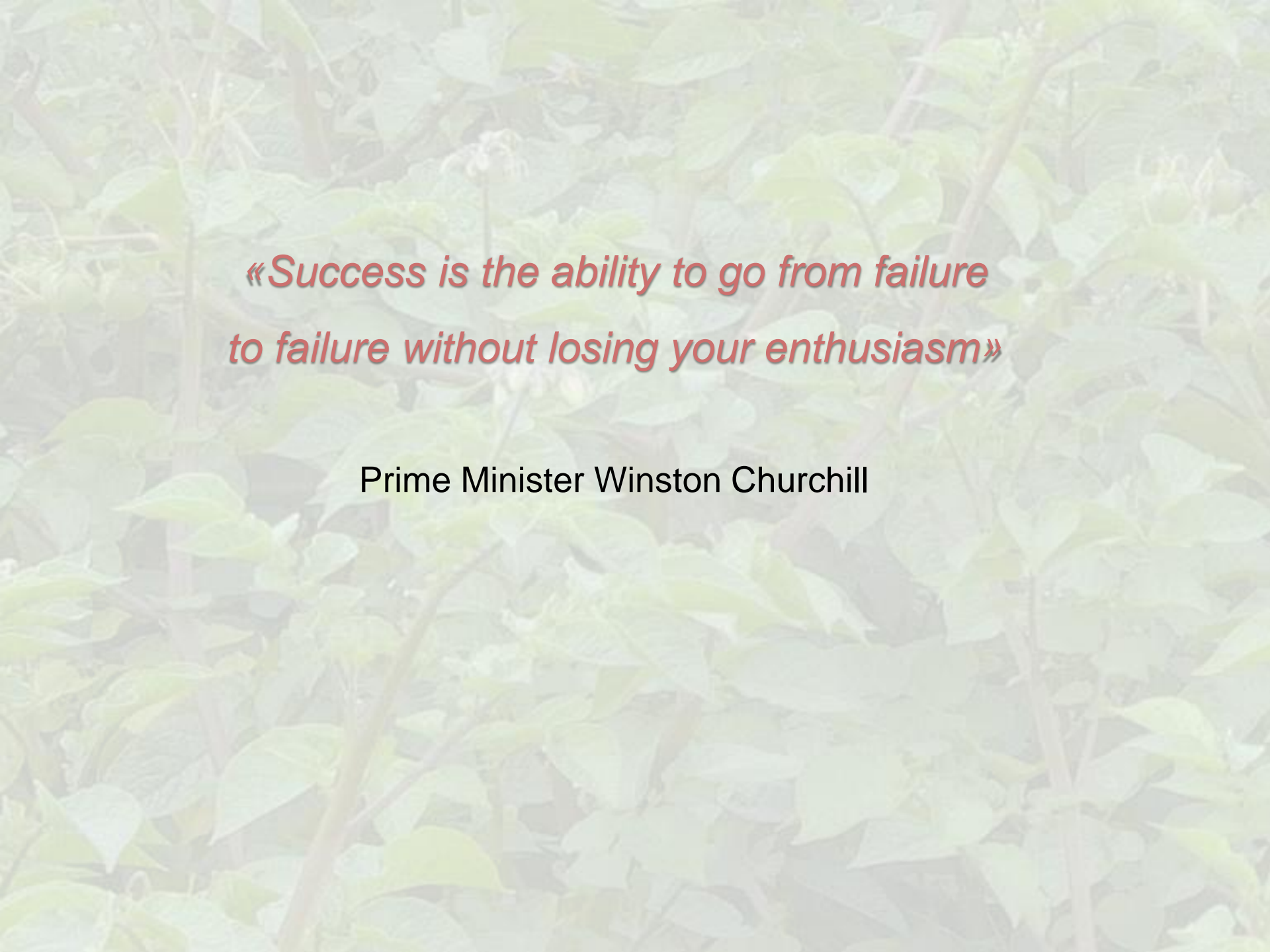
SCAR- markers for the *R* genes as predictors of late blight resistance in wild potato species

the germplasm of *Solanum* resistant to LB and carrying markers for *R1* or(and) *RB/Rpi-blb 1* gene:

<i>S. bulbocastanum</i>	<i>S. stoloniferum</i>	<i>S. demissum</i>	<i>S. hougasii</i>
<ul style="list-style-type: none">• K-21266• K-24862• K-24866 (2)	<ul style="list-style-type: none">• K-23652• K-24420• K-24976• K-24981	<ul style="list-style-type: none">• K-18487	<ul style="list-style-type: none">• K-24389• i-599618

the germplasm of *Solanum* resistant to LB and non-carrying markers for *R1* and *RB/Rpi-blb 1* gene:

<i>S. bulbocastanum</i>	<i>S. pinnatisectum</i>	<i>S. verrucosum</i>	<i>S. polyadenium</i>
<ul style="list-style-type: none">• K-21274• K-24866	<ul style="list-style-type: none">• K-24239	<ul style="list-style-type: none">• K-24990• K-24995	<ul style="list-style-type: none">• K-24408



*«Success is the ability to go from failure
to failure without losing your enthusiasm»*

Prime Minister Winston Churchill