



**Laboratory of Plant Molecular Genetics and Cytogenetics
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Advanced panel of molecular markers identifying of stem rust resistance genes *Sr2*, *Sr15*, *Sr22*, *Sr23*, *Sr24*, *Sr25*, *Sr26*, *Sr31*, *Sr35*, *Sr36*, *Sr38*, *Sr39*, *Sr44*, *Sr45*, *Sr57*, *Lr6Ai#2* in Siberian wheat cultivars

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Introduction

- Wheat stem rust is a plant disease caused by pathogenic fungus *Puccinia graminis* f. sp. *tritici* (**Pgt**) which leads to significant damage of the crop wheat.

The aim of this study was to identify genes and gene loci for resistance to wheat stem rust in Russian wheat germplasm using advanced panel of (Sr) markers.

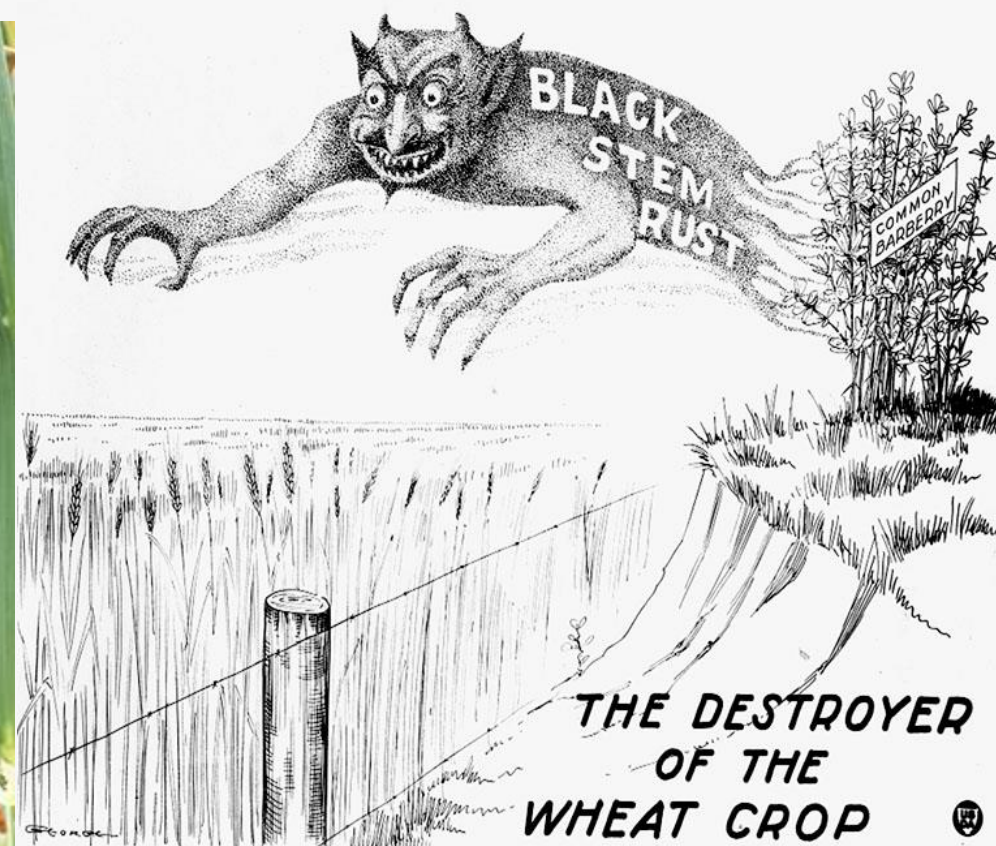


UREDINIOSPORES

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Photo by Robert Park



- Pgt** causes disruption of stem transpiration,
- reduces the photosynthetic ability of plants,
- reduces the quality of grain and its baking properties,
- reduces the strength of stems and sheaths, which leads to lodging of plants.

Materials and methods

- A total of **224** accessions of wheat germplasm was kindly received from (1) **FSBEI HE Omsk SAU** (Omsk region) - 80 varieties of spring soft wheat and 12 lines of spelt wheat; (2) **FRC IC&G SB RAS** (Novosibirsk region), consisting of 132 varieties of spring soft wheat.
- To identify resistance genes (**Sr**) in the germplasm, **18 markers** were used.

Genes	Chr.	Origin	Marker name	Source
<i>Sr2</i>	3BS	<i>Triticum turgidum</i>	Xgwm533 Sr2_ger9_3p	Mago et al., 2011
<i>Sr15/Lr20/Pm1</i>	7A	<i>T. aestivum</i>	Xsts638	Hu et al., 1997
<i>Sr22</i>	7A	<i>T. monococcum</i>	CFA2019	MAS*
<i>Sr23/Lr16</i>			XTaLr16_RGA266585	Harrison et al., 2016
<i>Sr24/Lr24</i>	3DL	<i>Agropyron elongatum</i>	Sr24#12	MAS
<i>Sr25/Lr19</i>	7D	<i>Thinopyrum ponticum</i>	Xwmc221	Gupta et al., 2006
<i>Sr26</i>	6AL	<i>Ag. elongatum</i>	Sr26#43 BE518379	MAS
<i>Sr31/Lr26/Yr9</i>	1AS, 1BS	<i>Secale cereale</i>	PrCEN-2	Li et al., 2016
<i>Sr35</i>	3AL	<i>T. monococcum</i>	NL9	
<i>Sr36</i>	2BL	<i>T. timophevii</i>	STM773-2	
<i>Sr38/Lr37/Yr17</i>	2AS	<i>T. ventricosum</i>	Ventriup+Ln2	
<i>Sr39/Lr35</i>	2B	<i>Aegilops speltoides</i>	BE500705	MAS
<i>Sr44</i>	7DS	<i>T. intermedium</i>	Xbe404728	
<i>Sr45</i>	1DS	<i>Ae. tauschii</i>	cssu45	
<i>Sr57/Lr34/Yr18/Pm38</i>	7D	hexaploid wheat	csLV34	
<i>Sr6Ai/Sr6Ai#2</i>	6D	<i>T. intermedium</i>	TNAC1752_TaqI	Salina et al., 2015

*<http://maswheat.ucdavis.edu/protocols/StemRust/index.htm>

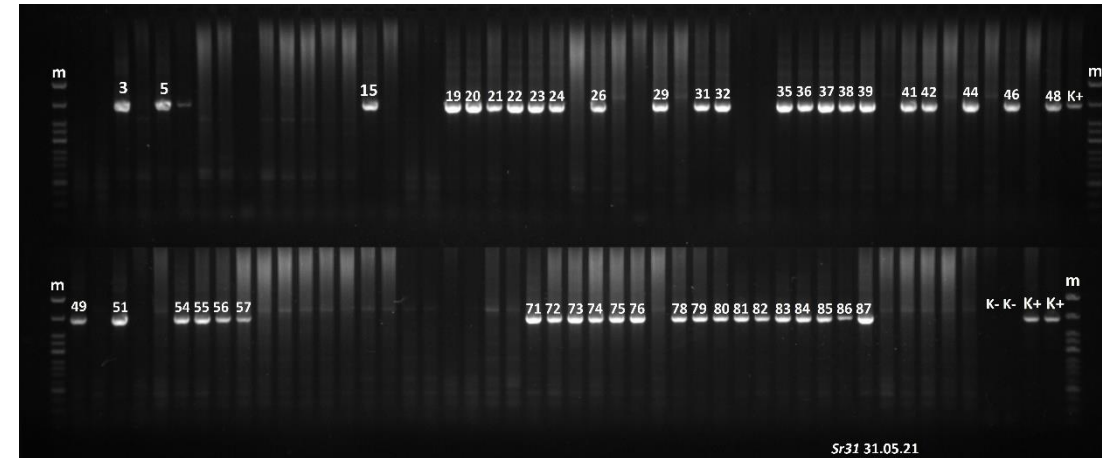
Results and Discussion

Working panel of DNA markers of stem rust resistance genes tested on wheat germ plasma.

for example: *Sr31/Lr26/Yr9* - **PrCEN-2** (5'-aatgatcttccacgacgacg-3', 5'-cctcgttgggaaatggtgca-3'), was designed according to nucleotides 1140–2090 of the pAWRC.1 sequence (GenBank accession No. AF245032).

Genes	Marker name	Marker type	Annealing temperature (°C)
<i>Sr2</i>	Xgwm533*	SSR	62
	Sr2_ger9_3p**	KASP	X
<i>Sr15/Lr20/Pm1</i>	Xsts638	STS	61
<i>Sr22</i>	CFA2019	SSR	60
<i>Sr23/Lr16</i>	XTaLr16_RGA266585	RGA	52
<i>Sr24/Lr24</i>	Sr24#12	AFLPs	q.v. MAS***
<i>Sr25/Lr19</i>	Xwmc221	SSR	60
<i>Sr26</i>	Sr26#43	STS	61
	BE518379	STS	61
<i>Sr31/Lr26/Yr9</i>	PrCEN-2		60
<i>Sr35</i>	NL9		61
<i>Sr36</i>	STM773-2	STS	q.v. MAS***
<i>Sr38/Lr37/Yr17</i>	Ventriup+Ln2	STS	65
<i>Sr39/Lr35</i>	BE500705	EST	61
<i>Sr44</i>	Xbe404728	CAPS	q.v. MAS***
<i>Sr45</i>	cssu45	SSR	60
<i>Sr57/Lr34/Yr18/Pm38</i>	csLV34	STS	54
<i>Sr6Ai/Sr6Ai#2</i>	TNAC1752_TaqI	CAPS	56

***<http://maswheat.ucdavis.edu/protocols/StemRust/index.htm>



Note:

*The KASP marker **Sr2_ger9_3p** was developed for **non-*Sr2*** wheat samples showing either the **null allele** or a **G allele** and an **A allele** associated with the presence of the *Sr2* gene.

** The microsatellite locus **Xgwm533** is tightly linked to the ***Sr2*** gene. However, there are two different Xgwm533 loci on chromosome 3BS, one of which does not carry the ***Sr2*** gene but gives an amplification.

Among the Omsk germplasm we observed the varieties carrying the following resistance genes: ***Sr2*, *Sr15*, *Sr23*, *Sr24*, *Sr25*, *Sr31*, *Sr38*, *Sr57***. The genes ***Sr2*, *Sr15*, *Sr22*, *Sr23*, *Sr25*, *Sr31*, *Sr44*, *Sr57*** and ***Lr6Ai#2*** in various combinations were identified in the varieties of the Institute of Cytology and Genetics.

Genes	FSBEI HE Omsk SAU	FRC IC&G SB RAS
<i>Sr2</i> (SSR-marker)	29	63
<i>Sr2</i> (KASP-marker)	absent	absent
<i>Sr15/Lr20/Pm1</i>	1	3
<i>Sr22</i>	23	59
<i>Sr23/Lr16</i>	4	3
<i>Sr24/Lr24</i>	4	absent
<i>Sr25/Lr19</i>	7	4
<i>Sr26</i>	absent	absent
<i>Sr31/Lr26/Yr9</i>	45	10
<i>Sr35</i>	absent	absent
<i>Sr36</i>	absent	absent
<i>Sr38/Lr37/Yr17</i>	7	absent
<i>Sr39/Lr35</i>	absent	absent
<i>Sr44</i>	50	2
<i>Sr45</i>	absent	absent
<i>Sr57/Lr34/Yr18/Pm38</i>	20	11
<i>Sr6Ai/Sr6Ai#2</i>	absent	6

The genes singly or in combination (FSBEI HE Omsk SAU)						
Genes	<u>0</u>	<u>1</u>	<u>2</u>	<u>3</u>	<u>4</u>	<u>5</u>
	5	27	32	15	11	2
%	5.4	29,4	34.8	16.3	11.9	2.2

The genes singly or in combination (FRC IC&G SB RAS)						
Genes	<u>0</u>	<u>1</u>	<u>2</u>	<u>3</u>	<u>4</u>	<u>5</u>
	31	56	33	9	3	0
%	23.5	42.4	25	6.8	2.7	