

Genetics of Leaf Rust Resistance in the Soft Red Winter Wheat Cultivars Coker 9663 and Pioneer 26R61

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ABSTRACT

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Leaf rust, caused by the fungus *Puccinia triticina*, is an important disease of soft red winter wheat cultivars that are grown in the southern and eastern United States. The objectives of this study were to identify the leaf rust resistance genes in two soft red winter wheat cultivars, Coker 9663 and Pioneer 26R61, that have been widely grown and were initially highly resistant to leaf rust. Both cultivars were crossed with the leaf-rust-susceptible spring wheat cv. Thatcher and the F₁ plants were crossed to Thatcher to obtain backcross (BC₁) F₂ families. In seedlings, the Thatcher/Coker 9663 BC₁F₂ families segregated for three independent seedling resistance genes when tested with different leaf rust isolates. The leaf rust infection types of selected BC₁F₃ lines, when tested with different leaf rust isolates, indicated that seedling resistance genes *Lr9*, *Lr10*, and *Lr14a* were present. In field plot tests, BC₁F₄ lines that were seedling susceptible had some adult plant resistance to leaf rust. Seedlings of the Thatcher/Pioneer 26R61 BC₁F₂ families segregated for two independent resistance genes. Infection types of selected BC₁F₃ lines indicated the presence of *Lr14b* and *Lr26*. The adult plant gene *Lr13* was determined to be present in selected BC₁F₄ lines that were tested with different leaf rust isolates in greenhouse tests.

Leaf rust, caused by *Puccinia triticina* Erikss., is the most common and widespread disease of wheat in the United States and worldwide (13). Soft red winter wheat cultivars are widely grown in the southern and eastern United States and have unique grain-quality characteristics that are suited for the production of cakes, cookies, and crackers. The soft red winter wheat cultivars have traditionally relied upon leaf rust resistance genes that are effective in seedlings. The comparison of seedling leaf rust infection types (ITs) produced on soft red winter wheat cultivars with ITs produced by the same *P. triticina* races on lines of wheat that differ for single leaf rust resistance genes allows for the postulation of seedling leaf rust resistance genes in these cultivars. Seedling resistance genes *Lr1*, *Lr2a*, *Lr9*, *Lr10*, *Lr11*, *Lr14a*, *Lr18*, *Lr24*, and *Lr26* have been postulated to be present in the soft red winter wheats (3). Genes *Lr12*, *Lr13*, *Lr34*, and *Lr46* were derived from common hexaploid wheat and are most effective in the adult plant stage. Genes *Lr13* and *Lr34* are common in the hard red

spring wheats (6,10), and *Lr46* has been described in CIMMYT wheats (15). Gene *Lr12* is likely present in cv. Caldwell (1) and other soft red winter wheats. Genetic analysis to identify seedling and adult plant leaf rust resistance genes has rarely been conducted with soft red winter wheat cultivars. The expression of leaf rust resistance in adult plants in field plots is dependent upon the effectiveness of seedling and adult plant resistance genes. In most cases, gene postulation based on seedling ITs is not sufficient to determine which adult plant resistance genes may be present in a cultivar.

The objective of this study was to determine the identity of seedling and adult plant resistance genes in the soft red winter wheat cvs. Coker 9663 and Pioneer 26R61. These cultivars were grown over large areas in the southern United States and are from two important wheat breeding programs. Coker 9663 and Pioneer 26R61 were highly resistant to leaf rust in field plots in North Carolina in 2000 and were also postulated to differ for seedling leaf rust resistance genes (3). The pedigree of Coker 9663 is IN71761A431548/F1302; and the pedigree of Pioneer 26R61 is Omega 78/S76/Arthur71/3/Stadler//Redcoat/Wisconsin 1/5/Coker 747/6/Pioneer 2555sib.

MATERIALS AND METHODS

Seed of Coker 9663 and Pioneer 26R61 was planted in 15-cm-diameter pots that were filled with steamed topsoil and placed on a greenhouse bench with a mix-

ture of fluorescent and incandescent lighting. After 10 days, when the plants were at the two-leaf stage, the pots were placed in a growth chamber with incandescent lighting at 10°C for 2 months of vernalization. After this period, the pots were placed in a growth cabinet with fluorescent and incandescent lighting with a 16-h day length at 18°C. Pots with seed of the leaf-rust-susceptible spring wheat cv. Thatcher (CI 1003) were placed in the growth cabinet 1 week before the Coker 9663 and Pioneer 26R61 plants were removed from the vernalization chamber and, at the same time, when the two soft red winter cultivars were placed in the growth cabinet. The Thatcher and soft red winter wheat plants were treated with Nutricot 13-13-13 NPK (Plantco Inc., Brampton, ON, Canada). At heading, the Thatcher plants were emasculated and pollen-shedding anthers from Coker 9663 and Pioneer 26R61 were used to pollinate the Thatcher female parents. The F₁ seed was harvested and then planted and vernalized for 1 month, and the F₁ plants were crossed as the male parent to Thatcher. Approximately 80 backcross (BC₁)F₁ seeds were obtained. The BC₁F₁ seed were planted in a greenhouse in 15-cm-diameter pots and selfed to obtain BC₁F₂ families.

In all, 15 to 20 seeds of each BC₁F₂ family were planted in a 3.5-cm² plastic pot and inoculated when the primary leaves were fully expanded with different leaf rust isolates. The *P. triticina* isolates were identified by a four-letter code that describes virulence to a set of 16 near-isogenic lines of Thatcher wheat that have different leaf rust resistance genes (4). ITs of the nine *P. triticina* isolates used in this study to Thatcher lines with single genes for leaf rust resistance are given in Table 1. For seedling inoculations, rust urediniospores were mixed with Soltol 170 oil (Phillips Petroleum, Borger, OK) and then spray inoculated onto plants using the equipment and methods previously described (13). After inoculation, seedling plants were allowed to dry for 1 h and then placed in a mist chamber overnight at 18°C and 100% relative humidity. The seedlings were placed on a greenhouse bench after incubation. Seedlings were fertilized with a 20-20-20 NPK solution immediately after inoculation and at 14 days after planting. The ITs on the primary leaves of individual plants were read at 10 to 12 days after inoculation. The ITs were

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classified using a 0-to-4 scale (9). ITs 0 (no visible sign of infection), ; (hypersensitive flecks), 1 (small uredinia surrounded by necrosis), and 2 (small/moderate size uredinia surrounded by chlorosis) were considered to be low (resistant) and ITs from 3 (moderate size uredinia without necrosis or chlorosis) to 4 (large uredinia) were considered to be high (susceptible). Mixtures of ITs or mesothetic responses were indicated by listing the most common IT first, followed by the less common ITs. Larger and smaller uredinia were indicated by “+” and “-”, respectively. BC₁F₂ families that had only susceptible seedlings were considered to be homozygous susceptible and families that had both resistant and susceptible plants were considered to be segregating. The ratio of segregating to homozygous susceptible families was used to estimate the number of segregating resistance genes. A χ^2 test (16) was used to determine whether the observed ratio significantly deviated from the expected ratio. The BC₁F₃ lines derived from individual BC₁F₂ plants selected for leaf rust resistance were tested for seedling resistance in the same manner as the BC₁F₂ families. An F₂ population of Thatcher/Pioneer 26R61 of approximately 120 individuals was tested for seedling resistance when the primary leaves were fully expanded.

In the field plot tests, 50 seeds of each genotype were planted in 2-m rows spaced 30 cm apart, perpendicular to rows of a mixture of wheat cvs. Thatcher, Morocco (PI 278386), Max (CI 15093), and Little Club (CI 4066) that are susceptible to leaf rust. The spreader rows and BC₁F₂ families of Thatcher/Coker 9663 and Thatcher/Pioneer 26R61 were inoculated in St. Paul, MN with a mixture of isolates MBBJ, THBJ, MCRK, TNRJ, and TDBG in 2005. In 2008, the spreader rows and selected Thatcher/Coker 9663 BC₁F₂ lines in St. Paul and Crookston, MN were inoculated with races MCDS, TDBG, MFPS, THBJ,

and TNRJ. The adult plants were rated for leaf rust severity using the modified Cobb scale (11). Leaf rust response in the adult plants was rated as previously described (13). The field plots were rated for leaf rust when the susceptible cv. Thatcher had a leaf rust severity of 70 to 80% with a susceptible response.

For the greenhouse evaluation of adult plants, 4 seeds of each genotype were planted in a 15-cm pot and grown in a greenhouse at 18 to 25°C with a 16-h light period. Flag leaves of adult plants were inoculated in the same manner as in the seedling tests, with a mixture of urediniospores and oil. ITs were read 14 days after inoculation using the same IT scale as for the seedling tests. BC₁F₂ plants from both crosses were evaluated for IT as adult plants, and seed from plants with low IT were selected and advanced by single-seed descent to BC₁F₄ lines. BC₁F₄ lines were tested for IT with different leaf rust isolates to postulate the identity of adult plant leaf rust resistance genes.

RESULTS

Seedlings of Thatcher/Coker 9663 BC₁F₂ families segregated in a two-gene ratio when tested with isolate BBBB and

for a single-gene ratio when tested with isolates THBJ, PNMR, and TLGF (Table 2). Segregation of BC₁F₂ families to isolates THBJ, PNMR, and TLGF combined fit a 7:1 ratio that indicated that three different genes in Coker 9663 conditioned resistance to these isolates. Seedlings of Thatcher/Pioneer 26R61 BC₁F₂ families segregated for a single gene to isolates BBBB and SBDG and all were homozygous susceptible to isolate THBJ. All BC₁F₂ families that were homozygous susceptible to isolate SBDG were also homozygous susceptible to isolate BBBB, which indicated that the same gene in Pioneer 26R61 conditioned resistance to these isolates. Eighteen selected Thatcher/Pioneer 26R61 BC₁F₂ families that segregated or were homozygous susceptible to isolate BBBB segregated independently ($P > 0.05$) for a different single gene when tested with isolate MCDS.

Seedling BC₁F₂ plants from each cross were selected on the basis of low IT to each tested isolate and grown to maturity to obtain BC₁F₃ lines for further seedling tests. Approximately 10 BC₁F₂ plants were selected from different families for each isolate. In the Thatcher/Coker 9663 BC₁F₂ lines (Table 3), lines 6 and 9 had low IT to

Table 2. Segregation of Thatcher*2/Coker 9663 and Thatcher*2/Pioneer 26R61 BC₁F₂ families for leaf rust resistance in seedlings

Cross, leaf rust isolates	BC ₁ F ₂ families		Expected ratio	χ^2	P
	Segregating	Susceptible			
Thatcher*2/Coker 9663					
BBBB	61	23	3:1	0.25	0.61
THBJ	46	41	1:1	0.28	0.59
PNMR	38	50	1:1	1.64	0.20
TLGF	43	43	1:1	0.00	1.00
THBJ+PNMR+TLGF	73	11	7:1	0.03	0.87
Thatcher*2/Pioneer 26R61					
BBBB	44	40	1:1	0.19	0.66
THBJ	0	83	0:1
SBDG	37	39	1:1	0.05	0.82
MCDS	11	7	1:1	0.89	0.35

Table 1. Seedling infection types (ITs) of 17 Thatcher (Tc) near-isogenic lines of wheat with single genes for leaf rust resistance to nine isolates of *Puccinia triticina*

Line	IT produced by <i>P. triticina</i> isolates ^a								
	BBBB	THBJ	PNMR	TLGF	SBDG	MCDS	MCRK	TCTD	MBRJ
TcLr1	;	3+	3+	3+	3+	3+	3+	3+	3+
TcLr2a	0;	3+	;1+	3+	3+	0;	0;	3+	0;
TcLr2c	0;	3+	3+	3+	3+	0;	0;	3+	0;
TcLr3a	;	3+	3+	3+	;	3+	3+	3+	3+
TcLr9	0;	;1=	3+	3+	;	;	0;	0	0;
TcLr16	;1	2+3	1+	;1-	;1-	2	1+	2	1+
TcLr24	;	;	3+	;	;	0;	0;	;	;
TcLr26	;	3+	;	;	;1=	3+	3	3+	;
TcLr3ka	;2-	;2	3+	;2-	;	2	3+	3	3+
TcLr11	;2-	;2	;2-	3+	;2-	2-	3+	3	3+
TcLr17	;2-	2	;	2+	3+	3	2=	3	2
TcLr30	;2-	;2=	3	;2	;	;2-	3+	3	3+
TcLrB	2	2	3	2-	;1-	3+	2	2	2
TcLr10	;	3+	3+	;	;	3	3+	;	3+
TcLr14a	3+	3+	;12+	3+	;12+	3+	3+	3	3+
TcLr14b	3+	3+	3	3+	3+	;	3+	;	3+
TcLr18	;2-	;1=	3	3+	;1-	;	3+	;2-	;2-

^a Infection types as described by Long and Kolmer (9).

isolates TLGF and BBBB, which indicated that these lines likely had gene *Lr10*; lines 15 and 16 had high IT to isolates TLGF and PNMR, which indicated that these lines likely had *Lr9*; and lines 38 and 43 had low IT to PNMR, which indicated that these lines likely had *Lr14a*. In the Thatcher/Pioneer 26R61 BC₁F₃ lines, lines 1 and 18 had low IT to isolate MCDS, which indicated that these lines likely have *Lr14b*, and lines 3 and 13 had high IT to isolates THBJ and MCDS, which indicated that these lines likely have *Lr26*. Seedlings of Thatcher/Pioneer 26R61 F₂ segregated 82 with IT ; (fleck) and 32 with IT 3⁺ which fit a single-gene 3:1 ratio ($\chi^2 = 0.57$; $P = 0.50$) when tested with isolate MCDS that has high IT to *Lr26*. This single-gene segregation was likely due to *Lr14b*.

The Thatcher/Coker 9663 BC₁F₂ families and the Thatcher/Pioneer 26R61 BC₁F₂ families that were homozygous susceptible to isolate BBBB were tested for segregation of leaf rust resistance as adult plants in inoculated field plot tests. The BC₁F₂ families from both crosses varied for leaf rust resistance, with rust severity varying from 10 to 80% in individual plants within families. However the

BC₁F₂ families also varied considerably for maturity. Families with later maturity had lower leaf rust severity compared with families with earlier maturity. Because segregation for maturity confounded the expression of adult plant resistance in the field plot tests, all subsequent adult plant tests were conducted in greenhouse experiments with specific isolates of leaf rust. Coker 9663 and Pioneer 26R61 were negative when tested with the PCR based markers for *Lr34* (8) and *Lr46* (E. Lagudah, CSIRO Canberra, *personal communication*), which indicated that both cultivars lack these two adult plant leaf rust resistance genes.

In all, 19 Thatcher/Coker 9663 BC₁F₂ families and 22 Thatcher/Pioneer 26R61 BC₁F₂ families that were susceptible to isolate BBBB in the seedling tests were tested as adult plants with isolate BBBB in greenhouse tests. In the Thatcher/Coker 9663 BC₁F₂ families, the IT ranged from 2⁺ to 3⁺, with no hypersensitive low ITs present. Some of the individual BC₁F₂ plants had relatively few uredinia compared with adult plants of Thatcher. In the Thatcher/Pioneer 26R61 BC₁F₂ families, the IT ranged from ;12 (hypersensitive

flecks with small uredinia surrounded by chlorosis and necrosis) to 3⁺. In total, 11 Thatcher/Coker 9663 BC₁F₂ plants that had fewer uredinia compared with Thatcher were selected and advanced by single-seed descent to BC₁F₄ lines, as were the 11 Thatcher/Pioneer 26R61 BC₁F₂ plants that had low IT of ;1–2. The 11 BC₁F₄ plants that were selected for adult plant resistance from both crosses were tested as adult plants in greenhouse tests with isolates BBBB, TCTD, MCRK, and THBJ that differ for IT to the Thatcher lines with genes *Lr12* and *Lr13*, and results for two lines from each cross are shown in Table 4. The Thatcher/Coker 9663 BC₁F₄ lines 36-2 and 38 had IT of 3 to 3⁺ to all four isolates, which indicated the lack of any hypersensitive adult plant resistance. Seed from lines 36-2 and 38 were advanced to BC₁F₅ and evaluated for adult plant resistance in inoculated field plots in St. Paul and Crookston, MN. Lines 36-2 and 38 both had lower severity of uredinia surround by chlorosis compared with rows of the susceptible control Thatcher at both locations. The low IT of Thatcher/Pioneer 26R61 BC₁F₄ lines 76-1 and 74 to isolates BBBB and TCTD indicated that these

Table 3. Seedling leaf rust infection types (ITs) of Thatcher*2/Coker 9663 BC₁F₃ and Thatcher*2/Pioneer 26R61 BC₁F₃ lines to six isolates of *Puccinia triticina*

Line	IT produced by <i>P. triticina</i> isolate ^a						Gene
	TLGF	PNMR	BBBB	MBRJ	THBJ	MCDS	
Thatcher*2/Coker 9663 F ₃							
6	;	3 ⁺	;	3 ⁺	-	-	<i>Lr10</i>
9	;	3 ⁺	;	3 ⁺	-	-	<i>Lr10</i>
15	3 ⁺	3 ⁺	0	0	-	-	<i>Lr9</i>
16	3 ⁺	3 ⁺	0	0	-	-	<i>Lr9</i>
38	3 ⁺	;1	3 ⁺	3 ⁺	-	-	<i>Lr14a</i>
43	3 ⁺	;12	3 ⁺	3 ⁺	-	-	<i>Lr14a</i>
Thatcher*2/Pioneer 26R61 F ₃							
1	3	3 ⁺	2*3	3 ⁺	3 ⁺	;	<i>Lr14b</i>
18	3 ⁺	3 ⁺	2*3	3 ⁺	3 ⁺	;	<i>Lr14b</i>
3	;	;1 ⁻	;	;1 ⁻	3 ⁺	3 ⁺	<i>Lr26</i>
13	;	;	;	;1 ⁻	3 ⁺	3 ⁺	<i>Lr26</i>
Thatcher <i>Lr9</i> RL 6010	3 ⁺	3 ⁺	;	0;	0;	0	...
Thatcher <i>Lr10</i> RL 6004	;	3 ⁺	;	3 ⁺	3	3	...
Thatcher <i>Lr14a</i> RL6013	3 ⁺	;1 ⁻	3 ⁺	3 ⁺	3	3	...
Thatcher <i>Lr14b</i> RL6056	3 ⁺	3 ⁺	3 ⁺	3 ⁺	3 ⁺	;	...
Thatcher <i>Lr26</i> RL6078	;	;	;	;	2*3	3	...

^a ITs as described by Long and Kolmer (9).

Table 4. Adult plant infection types (ITs), field rust severity, and response of Thatcher*2/Coker 9663 BC₁F₄ and Thatcher*2/Pioneer 26R61 BC₁F₄ lines and isogenic lines of Thatcher wheat with leaf rust resistance genes to four isolates of *Puccinia triticina*

Line	IT produced by <i>P. triticina</i> isolate ^a				Field rust severity and response ^b		Gene ^c
	BBBB	TCTD	MCRK	THBJ	St. Paul	Crookston	
Thatcher*2/Coker 9663 BC ₁ F ₄							
36-2	3 ⁺	33 ⁺	33 ⁺	3 ⁺	30 MS	50 MRMS	APR
38	3 ⁺	33 ⁺	3 ⁺	3 ⁺	50 MS	50 MS	APR
Thatcher*2/Pioneer 26R61 BC ₁ F ₄							
76-1	;2	0;	3	3 ⁺	<i>Lr13</i>
74	0;	0;	3 ⁺	3 ⁺	<i>Lr13</i>
Thatcher	3 ⁺	3 ⁺	3 ⁺	3 ⁺	80S	70S	...
Thatcher <i>Lr13</i> RL4031	;1 ⁻	;2 ⁻	3 ⁺	3 ⁺	70MRS	50S	...

^a ITs as described by Long and Kolmer (9).

^b Rust severity measured on a modified Cobb scale (11). MR = moderately resistant, medium to large uredinia surrounded by necrosis; MS = moderately susceptible, medium to large uredinia surrounded by chlorosis; and S = susceptible, large uredinia without necrosis or chlorosis.

^c APR = adult plant resistance.

lines likely have *Lr13*. Isolate TCTD also produces a low IT of ;1 to *Lr14b*. The low IT of 0; of the BC₁F₄ lines 76-1 and 74 to TCTD is likely due to the presence of both *Lr13* and *Lr14b*. The Thatcher line with *Lr13* conditioned a small degree of resistance relative to Thatcher in field plots at St. Paul and had slightly lower severity compared with Thatcher in plots at Crookston. Lines 76-1 and 74 both had high IT to isolate MCRK, which has a low IT to the adult plant gene *Lr12*, which indicated that neither of these lines have *Lr12*.

DISCUSSION

Coker 9663 was determined to likely have *Lr9*, *Lr10*, *Lr14a*, and some undefined adult plant resistance. The field leaf rust resistance of Coker 9663 was directly related to the effectiveness of *Lr9* and *Lr10*. In the 1990s, many soft red winter wheat cultivars were postulated to have *Lr9* (3); however, relatively few current soft red winter wheat cultivars have this gene (J. A. Kolmer, unpublished data). Virulence to *Lr9* increased steadily in the *P. triticina* population in the southeastern states starting in 1998, reaching nearly 50% of isolates in 2001 (4). However, in the late 1990s, many of the isolates with virulence to *Lr9* were race TLGF (2), which is avirulent to *Lr10*. During this time, Coker 9663 would have been resistant to TLGF in part due to the effectiveness of *Lr10*. Gene *Lr14a* has also been postulated to be in the soft red winter wheat cultivars (1) (J. A. Kolmer, unpublished data); however, virulence to this gene in the *P. triticina* population in the southeast has been very common and *Lr14a* would not be expected to condition any effective resistance. Coker 9663 was widely grown in the late 1990s throughout the mid-south and southeastern states. In 2000, Coker 9663 was highly resistant to leaf rust in field plots in North Carolina (3). However, by 2003, as races with virulence to *Lr9* and *Lr10* increased, Coker 9663 was considered to be susceptible to leaf rust and was no longer widely grown. Although Coker 9663 has some adult plant nonhypersensitive resistance, by itself it was not sufficient to condition a useful level of resistance for a released cultivar.

Pioneer 26R61 was determined to likely have *Lr13*, *Lr14b*, and *Lr26*. In the 1990s (3) and currently (J. A. Kolmer, unpublished data), *Lr26* was postulated to be present in a small number of soft red winter wheat cultivars. In the southeastern states, virulence to *Lr26* has varied between less than 5% in 2001 to 70% in 2007 (5). *Lr13* is an adult plant gene that conditions race-specific resistance. In the southeastern United States, most of the current races of *P. triticina* have virulence to *Lr13* (2), which has limited its effectiveness. *Lr13* may also be present in other soft red winter wheats. Roelfs (12) indi-

cated that the Brazilian cvs. Frondosa and Supreza were sources of *Lr13* for the soft red winter wheats and that Redcoat and Atlas 66 may have *Lr13*. *Lr14b* is not widely found in any wheat class in North America and, because nearly all isolates of *P. triticina* are virulent to this gene, it is unlikely that *Lr14b* contributes to the effective resistance in Pioneer 26R61. Although virulence to *Lr13* and *Lr26* is common in the *P. triticina* population in the United States, it would appear that the field resistance of Pioneer 26R61 is due to these two resistance genes, because no other effective seedling or adult plant resistance was detected in this cultivar. Pioneer 26R61 was widely grown in the southern states in the late 1990s and continues to be grown in Georgia. The leaf rust resistance in Pioneer 26R61 has remained relatively stable, because it has been rated as resistant to moderately resistant in recent field plot tests at various locations (2007–2008 Uniform Southern Winter Wheat Nurseries: <http://www.ars.usda.gov/main/docs.htm?docid=2925>), although, in some locations, it was rated as moderately susceptible. Isolates of *P. triticina* with virulence to *Lr13* and *Lr26* may not be evenly distributed across the southeastern region, which may explain why Pioneer 26R61 has been considered resistant and susceptible at different locations in the same year.

In a previous study (3), Coker 9663 was postulated to have *Lr2a*, *Lr9*, and *Lr10* on the basis of ITs with different *P. triticina* isolates. In that study, an isolate that was virulent to *Lr9* was also low to *Lr2a* and *Lr14a*. Similarly, in the same study, Pioneer 26R61 was postulated to have *Lr11* and *Lr26*. In the previous study, the isolates that were virulent to *Lr26* were also virulent to *Lr11*. Direct postulation of combinations of *Lr* genes in wheat cultivars by IT of different *P. triticina* isolates is limited by the virulence phenotypes of the available isolates. In this study, BC₁F₂ families and BC₁F₃ derived lines that segregated for single genes were used to identify the leaf rust resistance genes, rather than by direct postulation of the wheat cultivars.

Lr9 has been an important component of leaf rust resistance in soft red winter wheats. This gene was derived from *Triticum umbellulata* L. and was initially used in cvs. Riley 67 and Arthur 71, released in 1967 and 1971, respectively (14). Although derived from a wild relative of wheat, *Lr9* has not provided long-lasting resistance because races of *P. triticina* with virulence to this gene were found soon after the release of these cultivars. However, virulence to *Lr9* often declines rapidly after the replacement of wheat cultivars with this gene, suggesting some fitness cost associated with virulence to *Lr9*. Similarly, *Lr26* located on the 1B-1R wheat-rye translocation has not provided long-lasting resistance because leaf rust

races with virulence to this gene have increased in the United States and other countries after cultivars with the wheat-rye translocation were released. However, similar to *Lr9*, races with virulence to *Lr26* often gradually decline after the removal of cultivars with this gene. Thus, the effectiveness of *Lr9* and *Lr26* may be somewhat cyclical, depending on whether races with virulence to these genes are increasing or decreasing.

Development of soft red winter wheat cultivars that have long-lasting resistance will require combinations of leaf rust resistance genes that have conditioned effective resistance over a number of years. In the hard red spring wheats, combinations of *Lr16*, *Lr23*, and *Lr34* have provided effective, durable resistance despite the relatively poor resistance conditioned by the genes individually (6,10). Effective adult plant leaf rust resistance has been characterized in the winter wheats Caldwell (1) and CI13227 (17). These adult plant resistances could be combined with sources of *Lr34* such as Sturdy, Bezostaya1, and Sumai 3 (7) to develop cultivars with highly effective adult plant resistance that may prove to be durable over time.

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