Resistance to Stripe Rust and Eyespot Diseases of Wheat in Triticum tauschii

A. Yildirim and **S. S. Jones**, Department of Crop and Soil Sciences, Washington State University (WSU), Pullman, 99164-6420; **T. D. Murray**, Department of Plant Pathology, WSU, Pullman; **T. S. Cox**, USDA-ARS, Agronomy Department, KSU, Manhattan, KS; and **R. F. Line**, USDA-ARS, WSU, Pullman, WA

ABSTRACT

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A collection of 279 Triticum tauschii (syn. Aegilops squarrosa) accessions was evaluated for resistance to stripe rust (Puccinia striiformis) and eyespot (Pseudocercosporella herpotrichoides) diseases. Seedlings were inoculated with four different races of P. striiformis that represent all known virulences in the Pacific Northwest, and a genetically modified strain of P. herpotrichoides expressing β -glucuronidase. Seventeen percent (44) of the T. tauschii accessions were resistant to all Pacific Northwest races of stripe rust, and 45% (115) were resistant to eyespot. Thirty-nine of the 279 accessions were resistant to the stripe rust races and the eyespot pathogen. Accessions resistant to stripe rust were mainly from the Caspian Sea region of Iran and Azerbaijan, with the majority belonging to T. tauschii subsp. strangulata and T. t. subsp. meyeri. There was no clear association between resistance to eyespot and geographical origin or taxonomic subgroup.

Additional keywords: Triticum aestivum, wild wheat

Triticum tauschii (Coss.) Schmal. (syn. Aegilops squarrosa, 2n = 2x = 14, DD), a wild diploid, is the D-genome donor of hexaploid wheat (Triticum aestivum L.) (2n = 6x = 42, AABBDD) (18,26). Its close evolutionary relationship and the relative ease of hybridization with other Triticum spp. have made T. tauschii a subject of interest for the genetic improvement of hexaploid wheat (3,20). Accessions of T. tauschii are resistant to leaf rust (caused by Puccinia recondita Roberge ex Desmasz. f. sp. tritici), eyespot (caused by Pseudocercosporella herpotrichoides (Fron.) Deighton), powdery mildew (caused by Blumeria graminis (DC.) E.O. Speer f. sp. tritici Em. Marchal), tan spot (caused by Pyrenophora tritici-repentis (Died.) Drechs.), greenbug (Schizaphis graminum (Rondani)), Hessian fly (Mayetiola destructor (Say)), Karnal bunt (Tilletia indica Mitra), and wheat curl mite (Eriophyes tulipae Keifer) (3,11,17,28,-35,39), and possess genes that improve agronomic and end-use quality traits of wheat (16,32,38).

Stripe rust, caused by *Puccinia strii-formis* Westend., is an important yield-limiting foliar disease in the U.S. Pacific Northwest as well as in several other

Corresponding author: Stephen S. Jones, formerly USDA-ARS; E-mail: joness.wsu.edu

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wheat growing areas worldwide. This disease occurs on many grasses and cereal crops, including wheat, barley, Triticale, and rye (31). Cultivars with nonspecific and/or race-specific resistance are used to control stripe rust. Although race-specific resistance may become ineffective, usually within a few years of general use, resistant cultivars with this type of resistance have been effective in controlling the disease.

Eyespot, caused by Pseudocercosporella herpotrichoides (teleomorph = Tapesia yallundae Wallwork & Spooner), is another economically important disease of winter wheat in the Pacific Northwest and other temperate areas of the world (5,15,30,36). The disease has been effectively controlled with chemicals, but the discovery of isolates of the pathogen resistant to fungicides (29) has prompted plant breeders to focus on genetic control. Breeding programs have improved resistance in commercial cultivars (7,19,27), but progress is slow because only two resistance genes have been described for eyespot in wheat (25,37), and one on chromosome 4V in Dasypyrum villosum (L) (29), and these genes have not been combined in the same cultivar. Sprague (35) evaluated several wheat relatives. including T. tauschii, for eyespot resistance and suggested the use of wheat relatives in general, and Aegilops ventricosa in particular, as sources of resistance genes in breeding for resistance. Simonet (33) transferred resistance to evespot from A. ventricosa (genomes DDMVMV) into the hexaploid wheat cultivar Marne. Maia (25) then selected the line VPM-1 from this material based on its high degree of resistance to eyespot. VPM-1 has since become a widely used source of resistance in breeding programs. Identifying new sources of resistance is critical to maintaining a high level of resistance and to preventing the pathogen from circumventing resistance genes (12).

Cox et al. (3) and others (9,10,20,35) found that *T. tauschii* is a valuable source of genes for resistance to wheat pathogens. The objective of this study was to evaluate *T. tauschii* accessions for resistance to stripe rust and eyespot.

MATERIALS AND METHODS

Genetic stocks. Accessions of *T. tauschii* were obtained from the Wheat Genetic Resource Center, Kansas State University. Most of these accessions have been described previously (3,18,24). The following lines were used as controls for eyespot reaction: VPM-1, highly resistant; Cappelle-Desprez and Cerco, resistant; and Chinese Spring and Selection 101, susceptible.

Stripe rust reaction. Four races of P. striiformis representing the virulence of all Pacific Northwest (PNW) races (Table 1) were used for inoculation (23). Verified pure inoculum was increased on race-specific differential cultivars in isolation booths to prevent contamination. Freshly collected urediospores were used as inoculum for all tests. The seedlings were grown in a rust-free growth chamber with a diurnal temperature cycle of 10 to 25°C. Ten to 15 seedlings of each accession were uniformly inoculated with urediospores at the two-leaf stage, incubated in a dew chamber at 10°C for 18 to 24 h, and then placed in a growth chamber with a diurnal temperature cycle that gradually changed between 4 and 20°C. Day length was adjusted to 16 h with metal halide lights. Control cultivars and differential cultivars were planted in each experiment to determine viability of inoculum and purity of

Stripe rust infection type was recorded 15 to 18 days and 21 to 23 days after inoculation to reduce variability attributable to time (23). Infection types were recorded as 2, 5, and 8, representing resistant, intermediate, and susceptible, respectively, using the concept of the basic and expanded scales (23). Resistant accessions were retested to confirm their resistance.

Eyespot reaction. A genetically modified strain of *P. herpotrichoides* (P84-8) expressing β -glucuronidase (GUS) was

used to evaluate resistance (6,14). Two seeds per accession were sown in peat pots in a commercial potting mixture (55% peat:35% pumice:10% sand, wt/wt/wt) and covered with 2 cm of vermiculite. Fifty pots were then placed randomly in plastic flats (54 \times 27 \times 6 cm). Following germination, the flats were placed in a growth chamber at 12°C with 12 h light and 95 to 100% relative humidity. Two-week-old seedlings were inoculated, and disease progress was evaluated with a GUS assay 6 weeks after inoculation as described previously (6). Visual ratings of disease severity were modified from de la Peña and Murray (6) as follows: 1 = a lesion on the first leaf sheath only; 2 = a lesion on the first leaf sheath and a small lesion or speck on the second leaf sheath; 3 = alesion on the first and up to half of the second leaf sheath; and 4 = a lesion covering the entire first and second leaf sheaths. Relative resistance was determined by comparing accessions with the reactions of known control genotypes or cultivars. All resistant accessions were reevaluated in a second experiment to confirm their resistance reactions.

Experimental design and statistical analysis. Accessions were arranged in a randomized complete block design with three replicates. Analysis of variance was conducted with SAS statistical analysis software (SAS Institute, Cary, NC). GUS data were transformed by a logarithmic transformation before analysis because means and variances were not independent. Means were differentiated by Fisher's least significant difference (P = 0.01). Accessions with mean GUS values that exceeded the mean of the sum of the VPM value plus the LSD value were considered susceptible.

RESULTS

Seventeen percent of the T. tauschii accessions were resistant to all PNW races of P. striiformis, and an additional 21% exhibited resistance to specific races (Table 2). T. tauschii subsp. typica had the fewest accessions that were resistant to stripe rust (33%), whereas T. t. subsp. strangulata and T. t. subsp. meyeri had the most resistant accessions (59 and 66% resistant to all races, respectively). Eighteen percent of the accessions had an intermediate reaction to one or more CDL races. Some of the accessions had poor germination or few

seeds and therefore were not tested for all races or both diseases. Most of the stripe rust resistant accessions originated from Iran and Azerbaijan near the Caspian Sea. No resistant accessions were found among the collections from Turkey, Afghanistan, Pakistan, or China.

Forty-five percent of the accessions were resistant to eyespot. Differences among accessions were highly significant (P = 0.01). All subspecies had some resistant accessions. Two of the resistant accessions in the first experiment were determined to be susceptible in the second experiment (Table 2). The resistance of all other accessions was confirmed. T. t. subsp. strangulata and T. t. subsp. meyeri had the most resistant accessions, with 77 and 100% resistance to eyespot, respectively.

Thirty nine accessions from Turkmenistan, Iran, and Azerbaijan were resistant to both diseases, whereas most of the accessions from Afghanistan, Pakistan, and China were susceptible. Although all accessions from Turkey were highly susceptible to stripe rust, many were resistant to eyespot. Susceptibility was common in T. tauschii subsp. anathera and T. t. subsp. typica, whereas all accessions of T. t. subsp. strangulata and T. t. subsp. meyeri exhibited multiple resistance. The Caspian seacoast of Azerbaijan and Iran appeared to hold the main concentration of resistant accessions for both diseases.

DISCUSSION

Several accessions had resistance to all races of P. striiformis and to P. herpotrichoides. Resistance to both pathogens was most common in T. t. subsp. strangulata and T. t. subsp. meyeri. Cox et al. (3) found 13 of these same lines to be resistant to leaf rust, stem rust, powdery mildew, and tan spot. The occurrence of lines with multiple resistance should allow for the transfer of several beneficial genes from single crosses. Although we have not attempted to study the allelic relationships of already characterized stripe rust resistance genes in the D-genome, Yr8, Yr16, Yr18 (31,34), it is possible that some genes in T. tauschii lines will be different than genes already characterized. We also identified 32 spring-habit T. tauschii accessions among T. t. subsp. typica, mainly from Afghanistan. Generation time in the greenhouse can be greatly reduced because these lines have no vernalization require-

Another interesting result was the diversity among accessions in relation to their geographic origin. Resistance to eyespot was more dispersed geographically than that of stripe rust. T. t. subsp. strangulata from Iran had the highest frequency of resistance to stripe rust, eyespot, and other diseases studied by Cox et al. (3). T. t. subsp. strangulata is also easy to cross with wheat because it has larger heads and greater pollen shed than the other subspecies.

Accessions TA 2450 and TA 2460, which are resistant to both diseases, and accessions TA 2542 and TA 2547, which are susceptible to both diseases, have produced hexaploid progenies with soft endosperm (4). The only known soft-endosperm gene in hexaploid wheat is located on the short arm of chromosome 5D (22) and probably was transferred from the original T. tauschii parents of hexaploid wheat. The existence of a soft endosperm gene in T. tauschii offers an opportunity to improve the disease resistance of the Pacific Northwest soft white wheat cultivars while simultaneously manipulating kernel softness, an important end-use quality characteristic.

When Sprague screened wild relatives of wheat in 1936 (35), he reported that Aegilops squarrosa L. var. typica was moderately resistant, but Aegilops tauschii Coss. was mostly susceptible. Although he used a different classification for his accessions, they are now considered to be different subspecies of T. tauschii. We also found that 45% of T. t. subsp. typica accessions were moderately resistant to eyespot, whereas T. t. subsp. strangulata and T. t. subsp. meyeri were highly resistant. There have been no further reports on the use of T. tauschii for eyespot resistance since the initial report of Sprague.

There was a positive correlation observed between visual scores and GUS activity (r = 0.86), although a few exceptions were observed, for example accessions TA 1618 and 1649 from Iran. In both cases, high visual disease scores were associated with low GUS activity. In this situation, the hypersensitive reaction may result in a phenotypic appearance similar to that of a susceptible plant but limited disease development due to rapid death of infected epidermal cells. Similar responses

Table 1. Virulence of Cereal Disease Laboratory (CDL) races of Puccinia striiformis on North American differential cultivars

	Differential wheat cultivara														
CDL race	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
17	V	v	v	Α	A	A	Α	Α	V	Α	V	Α	Α	Α	Α
37	v	À	v	A	A	V	Α	V	V	V	V	V	Α	Α	Α
43	V	Α	V	V	V	Α	Α	Α	Α	Α	Α	V	Α	V	Α
45	V	Α	V	Α	Α	Α	Α	Α	Α	Α	Α	V	V	Α	V

^a 1: Lemhi; 2: Chinese 166; 3: Heines VII; 4: Moro; 5: Paha; 6: Druchamp; 7: Riebesel 47-51; 8: Produra; 9: Yamhill; 10: Stephens; 11: Lee; 12: Fielder; 13: Tyee; 14: Tres; 15: Hyak. V: Virulent, A: Avirulent

Table 2. Reactions of Triticum tauschii accessions to Pseudocercosporella herpotrichoides and North American races of Puccinia striiformis

WGRC ^a				Reaction to	CDL race	b		Visual	GUS	Growth
acc. no.	Region	Subspecies	17	37	43	45	- Eyespot ^c	score	activity	habit
1604	Afghanistan	typica	ND^d	ND	ND	ND	S	4.0	3.8	Winter
1620	Afghanistan	Unknown	S	S	S	S	ND	ND	ND	Winter
1629	Afghanistan	typica	S	S	S	S	ND	ND	ND	Winter
1632	Afghanistan	typica	S	S	S	S	S	3.2	2.4	Spring
1637	Afghanistan	Unknown	S	S	S	S	S	4.0	2.6	Winter
1638	Afghanistan	typica	S	S	S	S	R	1.5	1.7	Spring
1655	Afghanistan	typica	S	S	S	S	R	1.8	1.9	Spring
1703	Afghanistan	Unknown	S	S	S	S	S	3.4	2.5	Spring
1705	Afghanistan	typica	S	S	S	S	ND	ND	ND	Winter
2387	Afghanistan	typica	S	S	S	S	S	3.6	2.3	Winter
2389	Afghanistan	typica	S	S	S	S	S	3.0	2.4	Winter
2390	Afghanistan	typica	S	S	S	S	S	3.8	2.4	Winter
2391	Afghanistan	typica	S	S	ND	I	S	4.0	2.6	Spring
394	Afghanistan	typica	S	S	S	S	S	4.0	2.6	Winter
395	Afghanistan	typica	S	S	S	I	R	2.5	1.9	Winter
2396	Afghanistan	typica	S	S	S	S	S	3.5	2.3	Winter
397	Afghanistan	typica	S	S	S	S	R	2.4	2.1	Winter
398	Afghanistan	typica	S	S	S	S	R	2.3	1.7	Winter
399	Afghanistan	typica	S	S	S	S	R	1.0	1.4	Winter
400	Afghanistan	typica	ND	ND	ND	ND	R	1.0	1.4	Winter
401	Afghanistan	typica	S	S	S	S	R	1.0	1.3	Winter
402	Afghanistan	typica	S	S	S	S	S	4.0	2.6	Winter
403	Afghanistan	typica	S	I	ND	. I	S	4.0	2.8	Spring
405	Afghanistan	anathera	S	S	S	S	S	3.8	2.6	Winter
406	Afghanistan	typica	S	S	ND	S	S	4.0	2.5	Spring
407	Afghanistan	anathera	S	S	S	S	S	3.0	2.8	Winter
408	Afghanistan	typica	S	S	S	S	ND	ND	ND	Winter
409	Afghanistan	typica	S	S	S	S	S	2.3	2.3	Spring
410	Afghanistan	typica	ND	ND	ND	ND	R	1.2	1.2	Winter
412	Afghanistan	typica	ND	ND	ND	ND	R	2.6	1.9	Winter
415	Afghanistan	typica	S	S	S	S	R	2.6	1.9	Winter
416	Afghanistan	typica	S	S	S	S	S	3.0	2.2	Winter
417	Afghanistan	anathera	S	S	S	S	S°	2.2	2.6	Winter
418	Afghanistan	typica	S	S	S	S	S	4.0	2.3	Winter
419	Afghanistan	typica	S	S	ND	S	ND	ND	ND	Spring
420	Afghanistan	typica	S	S	ND	S	S	3.0	2.3	Spring
421	Afghanistan	typica	S	S	S	S	S	4.0	2.8	Spring
422	Afghanistan	typica	S	S	S	S	S	3.1	2.5	Winter
423	Afghanistan	typica	S	S	ND	S	R	2.0	1.3	Spring
424	Afghanistan	typica	S	S	ND	S	R	1.5	1.5	Spring
426	Afghanistan	typica	S	S	ND	S	S	4.0	3.3	Spring
427	Afghanistan	typica	S	S	ND	S	S	4.0	2.6	Spring
128	Afghanistan	typica	S	S	ND	S	S	4.0	2.9	Spring
130	Afghanistan	typica	S	S	S	S	S	4.0	3.1	Winter
132	Afghanistan	typica	S	S	S	Š	Š	4.0	2.8	Winter
133	Afghanistan	typica	S	S	S	S	Š	3.6	2.8	Winter
134	Afghanistan	typica	S	S	S	S	S	3.5	2.5	Winter
135	Afghanistan	typica	S	S	I	I	Š	3.5	2.5	Winter
137	Afghanistan	typica	S	S	ND	Š	Š	4.0	2.7	Spring
38	Afghanistan	typica	S	S	ND	Š	S	4.0	2.7	Spring
39	Afghanistan	typica	S	S	S	S	Š	3.6	2.9	Winter
40	Afghanistan	typica	S	S	Š	Š	Š	4.0	3.0	Winter
42	Afghanistan	typica	S	S	ND	S	Š	4.0	2.6	Spring
43	Afghanistan	typica	S	S	ND	S	Š	3.7	2.8	Spring
44	Afghanistan	typica	S	S	S	Š	S	3.5	2.6	Winter
32	Afghanistan	typica	S	S	Š	S	S	4.0	2.7	Winter
33	Afghanistan	typica	ND	ND	ND	ND	S	3.1	2.7	Winter
35	Afghanistan	typica	ND	ND	ND	ND	S	4.0	3.0	Winter
36	Afghanistan	typica	S	S	S	S	S	4.0	3.0	
37	Afghanistan	typica	Š	Š	S	S	S	4.0		Winter
38	Afghanistan	typica	ND	ND	ND	ND	S	4.0 4.0	2.8	Winter
39	Afghanistan	typica	S	S	S	S	S S		2.9	Winter
40	Afghanistan	typica	S	S	S	S S	S S	4.0	3.0	Winter
541	Afghanistan	typica	S	S	S	S		2.4	2.4	Winter
542	Afghanistan	typica	S	S	S	S	S	4.0	2.9	Winter
143	Afghanistan	typica	S	S	S	S S	S	4.0	2.7	Winter
44	Afghanistan	typica	S	S	S	S S	S S	4.0	3.0 2.7	Winter
	J		5	J	ن			4.0	.17	Winter

^a WGRC: Wheat Genetics Resource Center, Manhattan, Kansas.

b Infection types were recorded as 2, 5, 8, representing resistant (R), intermediate (I) and susceptible (S), respectively, using basic and expanded scales.

C Disease progress was evaluated visually and with a GUS assay 6 wk after inoculation.

^d ND: Not determined.

^e Determined to be susceptible in the second experiment.

Table 2. (continued from preceding page)

WGRC ^a					CDL raceb		_	Visual	GUS activity	Growth habit
icc. no.	Region	Subspecies	17	37	43	45	- Eyespot ^c	score		
545	Afghanistan	typica	I	I	I	I	S	4.0	3.0	Winter
546	Afghanistan	typica	S	S	S	S	S	4.0	2.7	Winter
547	Afghanistan	typica	S	S	S	S	ND	ND	ND	Winter
548 540	Afghanistan	typica	S	S	S	S	S	4.0	2.5	Winter
549 550	Afghanistan	typica	S	S	S	S	S	3.2	2.7	Winter
550	Afghanistan	typica	S	S	S	S	S	4.0	2.8	Winter
553	Afghanistan	typica	S	S	S	S	S	4.0	3.1	Winter
555	Afghanistan	typica	S	S	S	S	S	4.0	2.8	Winter
558	Afghanistan	typica	S	S	S	S	S	4.0	2.6	Winter
559	Afghanistan	typica	S	S	S	S	R	1.0	1.1	Spring
587	Afghanistan	typica	S	S	S	S	S	4.0	2.4	Winter
640 700	Armenia	Unknown	S	S	S	S	ND	ND	ND	Winter
700 566	Armenia	Unknown	S	S	S	S	S	2.8	4.5	Winter
567	Armenia	anathera	S S	S	S	I	S	3.0	2.4	Winter
568	Armenia Armenia	typica	S R	S R	S	S	R	2.7	2.0	Winter
569	Armenia	typica typica	S	S	R S	R	ND	ND	ND	Winter
570	Armenia		S	S		S	S	4.0	2.9	Winter
570 571	Armenia	typica typica	S I	S	S	S	ND	ND	ND	Winter
574	Armenia	typica typica	S	S	S S	S S	R	2.5 4.0	1.9	Winter
575	Armenia	**	S	S			S		2.6	Winter
522	Armema Azerbaijan	<i>typica</i> Unknown	S S	S S	S S	S S	ND R	ND	ND	Winter
524	Azerbaijan Azerbaijan	Unknown	S R	S R	S R	S R	R R	1.8	0.7	Winter
556	Azerbaijan	typica	K I	R R	R R	R R		1.6 2.0	0.7	Winter
557	Azerbaijan Azerbaijan	typica typica	I	R R	R R	R R	R R		1.8	Winter
558	Azerbaijan	typica typica	ND	ND	R ND	ND	S	1.5 4.0	1.6	Winter Winter
559	Azerbaijan	strangulata	R	R	R	R R	S R	4.0 1.5	2.6	Winter
660	Azerbaijan	Unknown	I	S	R S	S	S	3.6	1.7 2.6	Winter
662	Azerbaijan	typica	I	R	R	R	S	3.0	2.6	Winter
664	Azerbaijan	Unknown	S	I	I	I	R R	1.3	1.6	Winter
665	Azerbaijan	Unknown	R	R	R	R	R R	1.0	1.0	Winter
566	Azerbaijan	Unknown	ND	ND	ND	ND	R R	1.3	1.5	Winter
667	Azerbaijan	typica	ND	ND	ND	ND	R R	1.5	1.3	Winter
668	Azerbaijan	typica	R	R	R	R	R	1.0	1.3	Winter
669	Azerbaijan	Unknown	S	S	S	S	R	2.0	1.3	Winter
570	Azerbaijan	typica	I	R	R	R	R	2.9	2.0	Winter
571	Azerbaijan	typica	R	R	R	R	S	2.5	2.0	Winter
572	Azerbaijan	Unknown	I	S	I	S	R	1.5	1.3	Winter
574	Azerbaijan	typica	S	I	Ī	S	S	3.0	2.5	Winter
575	Azerbaijan	typica	Ĭ	Ř	R	R	R	1.0	1.2	Winter
576	Azerbaijan	Unknown	Ś	Š	Š	S	S	3.1	2.4	Winter
577	Azerbaijan	Unknown	R	R	R	R	R	3.0	2.7	Winter
578	Azerbaijan	typica	S	R	R	R	R	1.0	1.1	Winter
580	Azerbaijan	typica	S	Š	Ĭ	R	R	1.5	1.8	Winter
581	Azerbaijan	Unknown	S	R	Î	I	R	1.0	1.1	Winter
582	Azerbaijan	typica	R	R	R	Ř	R	1.5	1.6	Winter
584	Azerbaijan	Unknown	I	R	R	R	R	2.0	1.6	Winter
85	Azerbaijan	Unknown	ND	ND	ND	ND	R	1.5	1.4	Winter
586	Azerbaijan	typica	R	R	R	R	R	1.5	1.4	Winter
587	Azerbaijan	typica	ND	ND	ND	ND	R	1.8	1.7	Winter
560	Azerbaijan	typica	R	R	R	R	ND	ND	ND	Winter
62	Azerbaijan	typica	R	R	R	R	ND	ND	ND	Winter
501	China	typica	S	S	S	S	S	3.3	2.8	Winter
02	China	Unknown	S	S	S	S	S	2.9	2.7	Winter
603	China	Unknown	S	S	S	S	S	3.5	3.0	Winter
598	Dagestan	Unknown	Š	S	Ī	S	R	3.0	2.1	Winter
599	Dagestan	typica	S	S	S	S	S	4.0	3.0	Winter
369	Dagestan	typica	S	S	S	I	ND	ND	ND	Winter
577	Georgia	typica	Ī	S	S	Š	ND	ND	ND	Winter
578	Georgia	typica	Š	S	S	S	ND	ND	ND	Winter
579	Georgia	typica	R	R	I	R	S	4.0	2.7	Winter
80	Georgia	typica	ND	ND	Š	S	Š	4.0	3.0	Winter
81	Georgia	typica	S	S	S	S	S	3.8	2.5	Winter
85	Georgia	typica	Ĭ	S	S	ND	ND	ND	ND	Winter
586	Georgia	typica	Š	S	S	S	S	4.0	2.8	Winter
599	Iran	meyeri	S	S	S	S	R	1.3	1.4	Winter
500	Iran	Unknown	S	I	I	I	R	1.3	1.5	Winter
518	Iran	Unknown	S	I	Ĭ	Ī	R	3.3	1.7	Winter
519	Iran	meyeri	S	I	R	R	ND	S.S ND	ND	Winter
541	Iran	meyeri strangulata	I	R	R	I	R	1.5	1.2	Winter
542	Iran	strangulata strangulata	R	R	R	R	R R	1.5		
542 543	Iran	typica	R R	I I	R	R R	R R	1.5	1.6	Winter Winter
544	Iran	typica typica	R	R	R	R R	R R	2.0	1.3 1.7	
	11411	iypica	1	1/	1	IV.	I.	∠.∪	1./	Winter

Table 2. (continued from preceding page)

WGRC ^a				Reaction to	CDL raceb			Visual	GUS	Growth
icc. no.	Region	Subspecies	17	37	43	45	Eyespot ^c	score	activity	habit
645	Iran	Unknown	I	I	R	R	S	2.8	2.3	Winter
549	Iran	meyeri	R	R	R	R	R	3.2	2.0	Winter
706	Iran	strangulata	R	R	R	R	R	2.4	2.0	Winter
715	Iran	typica	R	R	R	R	S	2.2	2.2	Winter
717	Iran	Unknown	S	S	S	R	S	4.0	2.8	Winter
718	Iran	Unknown	R	R	R	R	S	3.5	2.6	Winter
377	Iran	strangulata	R	R	R	R	R	1.6	1.7	Winter
377.1	Iran	strangulata	R	R	R	R	R	1.0	1.5	Winter
445	Iran	typica	R	I	R	R	ND	ND	ND	Winter
446	Iran	typica	R	I	S	R	R	1.0	1.4	Winter
447	Iran	typica	I	I	I	R	S	2.7	2.2	Winter
448	Iran	typica	S	I	ND	R	S	4.0	2.6	Winter
449	Iran	strangulata	R	R	R	R	R	1.0	1.4	Winter
450	Iran	strangulata	R	R	R	R	R	1.0	1.5	Winter
452	Iran	strangulata	R	R	R	R	S	3.7	2.4	Winter
452.1	Iran	strangulata	R	I	R	I	R	1.7	1.9	Winter
453	Iran	strangulata	R	R	R	R	R	2.5	2.0	Winter
454	Iran	strangulata	R	I	R	I	S	3.9	2.2	Winter
455	Iran	strangulata	R	S	I	S	S	3.6	2.4	Winter
457	Iran	typica	R	R	R	R	R	1.1	1.2	Winter
458	Iran	typica	S	S	I	S	ND	ND	ND	Winter
159	Iran	typica	R	R	R	I	R	1.0	1.3	Winter
460	Iran	typica	R	R	ND	R	R	1.1	1.6	Winter
463	Iran	strangulata	S	S	S	I	R	1.5	1.5	Winter
464	Iran	strangulata	S	S	I	S	S	2.8	2.3	Winter
465	Iran	strangulata	I	S	I	I	R	2.0	1.8	Winter
466	Iran	strangulata	I	I	I	I	ND	ND	ND	Winter
467	Iran	strangulata	R	R	R	R	ND	ND	ND	Winter
468	Iran	strangulata	R	I	R	I	R	2.6	1.9	Winter
469	Iran	strangulata	R	R	R	R	S	4.0	2.4	Winter
470	Iran	strangulata	R	R	ND	R	R	1.7	1.9	Winter
471	Iran	typica	R	I	I	R	R	2.5	2.0	Winter
472	Iran	typica	R	R	R	R	R	2.1	1.9	Winter
473	Iran	meyeri	R	R	ND	R	R	2.1	1.3	Winter
474	Iran	typica	R	R	I	R	S	4.0	3.0	Winter
475	Iran	typica	R	R	I	R	S	3.5	2.3	Winter
476	Iran	typica	R	R	I	R	R	3.0	2.0	Winter
477	Iran	typica	I	I	I	I	R	2.5	2.0	Winter
478	Iran	typica	I	R	I	I	R	2.0	1.6	Winter
479	Iran	typica	S	S	S	S	S	3.5	2.3	Winter
480	Iran	typica	R	R	R	R	R	1.0	1.0	Winter
481	Iran	meyeri	R	R	R	R	R	3.5	2.1	Winter
483	Iran	typica	R	R	R	R	R	1.5	1.5	Winter
485	Iran	typica	R	R	R	R	S	4.0	2.6	Winter
486	Iran	typica	S	S	S	S	S	3.2	2.7	Winter
487	Iran	typica	S	S	S	S	S	3.6	2.6	Winter
488	Iran	typica	R	R	I	R	S	4.0	3.0	Winter
489	Iran	typica	S	S	S	S	S	4.0	3.0	Winter
490	Iran	typica	R	R	R	R	R	1.0	1.2	Winter
492	Iran	typica	S	S	S	S	R	1.1	1.5	Winter
493	Iran	typica	S	S	S	S	R	1.6	1.5	Winter
494	Iran	typica	S	ND	ND	S	R	1.0	1.3	Winter
195	Iran	typica	R	R	R	R	R	1.5	1.4	Winter
196	Iran	typica	ND	ND	ND	ND	R	1.0	1.2	Winter
497	Iran	typica	ND	ND	ND	ND	R	1.6	1.3	Winter
498	Iran	typica	ND	ND	ND	ND	R	1.7	1.4	Winter
499	Iran	typica	ND	ND	ND	ND	R	1.2	1.3	Winter
511	Iran	typica	R	S	R	S	R	1.0	1.2	Winter
512	Iran	typica	S	S	R	S	R	1.0	1.3	Winter
513	Iran	typica	S	S	S	S	S	3.2	2.4	Winter
515	Iran	typica	S	S	S	S	S	2.5	2.4	Winter
517	Iran	typica	S	S	S	S	ND	ND	ND	Winter
519	Iran	typica	S	S	S	S	S	4.0	2.8	Winter
520	Iran	typica	S	S	S	S	S	3.7	2.4	Winter
521	Iran	typica	ND	ND	ND	ND	S	3.9	2.4	Winter
523	Iran	typica	ND	ND	ND	ND	S	3.6	2.9	Winter
524	Iran	typica	R	R	R	R	R	1.0	1.2	Winter
525	Iran	typica	R	R	R	R	R	1.0	1.4	Winter
527	Iran	meyeri	S	S	S	I	R	1.1	1.3	Winter
528	Iran	typica	R	R	R	R	R	1.0	1.3	Winter
529	Iran	meyeri	R	R	R	R	R	1.7	1.3	Winter
530	Iran	meyeri	R	R	R	R	R	1.9	1.5	Winter
374	Pakistan	anathera	S	S	S	I	S	4.0	2.5	Spring
	*	**				-	_			ed on next p

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 $\textbf{Table 2.} \ (continued \ from \ preceding \ page)$

WGRC ^a			Reaction to CDL raceb				_	Visual	GUS	Growth	
icc. no.	Region	Subspecies	17	37	43	45	Eyespot ^c	score	activity	habit	
379	Pakistan	typica	S	S	S	S	S°	2.7	2.2	Winter	
380	Pakistan	typica	S	S	S	S	S	2.7	2.3	Winter	
381	Pakistan	anathera	S	S	S	S	S	4.0	2.9	Winter	
382	Pakistan	anathera	S	ND	ND	ND	S	3.5	2.6	Spring	
383	Pakistan	typica	S	S	S	S	R	1.7	1.2	Spring	
384	Pakistan	Unknown	S	S	ND	S	S	2.8	2.5	Spring	
385	Pakistan	typica	S	S	ND	S	ND	ND	ND	Spring	
386	Pakistan	anathera	S	S	ND	I	ND	ND	ND	Spring	
652	Tadshikistan	Unknown	I	S	S	S	ND	ND	ND	Winter	
653	Tadshikistan	typica	S	S	S	S	S	3.8	2.5	Winter	
704	Tadshikistan	Unknown	S	S	S	S	R	2.7	1.9	Winter	
582	Turkey	Unknown	S	S	S	S	S	2.5	2.7	Winter	
584	Turkey	typica	S	S	S	S	S	3.6	2.9	Winter	
586	Turkey	Unknown	S	S	S	S	S	1.6	2.2	Winter	
587	Turkey	Unknown	S	S	S	S	S	4.0	3.2	Winter	
588	Turkey	Unknown	S	S	S	S	S	2.2	2.2	Winter	
589	Turkey	Unknown	S	S	S	S	R	1.0	1.8	Winter	
590	Turkey	Unknown	S	S	S	S	S	2.8	2.3	Winter	
591	Turkey	Unknown	S	S	S	S	R	2.0	2.0	Winter	
593	Turkey	Unknown	S	S	S	S	S	2.2	2.9	Winter	
594	Turkey	Unknown	S	S	S	S	R	1.0	1.5	Winter	
595	Turkey	Unknown	S	S	S	S	S	3.9	3.6	Winter	
596	Turkey	Unknown	S	S	S	S	S	3.2	2.4	Winter	
634	Turkey	Unknown	S	S	S	S	S	4.0	2.9	Winter	
500	Turkey	typica	ND	ND	ND	ND	S	2.4	2.9	Winter	
503	Turkey	typica	ND	ND	ND	ND	R	1.7	2.0	Winter	
504	Turkey	typica	ND	ND	ND	ND	R	1.0	1.8	Winter	
505	Turkey	typica	ND	ND	ND	ND	R	1.0	1.6	Winter	
507	Turkey	typica	S	S	S	S	S	3.7	2.3	Winter	
508	Turkey	typica	S	S	S	S	R	1.0	1.3	Winter	
509	Turkey	typica	S	S	S	S	S	3.0	2.5	Winter	
510	Turkey	typica	S	S	Š	Š	Ř	1.0	1.5	Winter	
614	Turkmenistan	typica	Ř	R	Ř	Ř	R	1.7	1.8	Winter	
615	Turkmenistan	typica	R	R	R	R	R	1.2	1.8	Winter	
616	Turkmenistan	typica	S	S	S	S	R	1.5	1.6	Winter	
617	Turkmenistan	Unknown	S	Š	Š	Š	R	2.0	1.5	Winter	
623	Turkmenistan	Unknown	Š	I	Ĭ	Ī	R	1.7	1.3	Winter	
626	Turkmenistan	Unknown	R	Ś	Ī	Ř	R	1.8	1.3	Winter	
692	Turkmenistan	Unknown	S	I	Î	I	Š	3.7	2.8	Winter	
693	Turkmenistan	Unknown	S	Ř	Ř	Ř	Ř	1.0	1.6	Winter	
598	USSR	Unknown	S	S	S	S	R	2.1	1.7	Winter	
625	USSR	Unknown	S	S	S	S	S	3.6	2.6	Winter	
577	Unknown	Unknown	Š	S	S	S	S	3.8	3.4	Winter	
578	Unknown	Unknown	S	S	S	S	S	3.0	3.4	Winter	
579	Unknown	Unknown	S	S	S	S	S	3.6	3.6	Winter	
580	Unknown	Unknown	S	S	S	S	S	3.7	3.7	Winter	
581	Unknown	Unknown	S	S	S	S	S	4.0	3.7	Spring	
597	Unknown	Unknown	S	S	S	S	S	3.8	3.1	Winter	
605	Unknown	strangulata	I	R	R R	R	R R	3.8 1.5			
621	Unknown	Unknown	S						2.0	Winter	
635	Unknown	onknown strangulata	S I	S S	S I	S I	S R	3.5 2.0	2.4	Winter	
651	Unknown	Unknown	R	S R	R	R	R R	2.0	1.7	Winter	
688	Unknown Unknown		K S	R S					1.1 ND	Winter	
689	Unknown Unknown	typica typica	S S	S S	S	S	ND	ND	ND	Spring	
689 691		typica			S	S	S	3.5	2.8	Spring	
695	Unknown	meyeri strangulata	R	R	R	R	R	1.8	1.8	Winter	
695 696	Unknown	strangulata	R	R	R	R	R	1.8	1.8	Winter	
696 697	Unknown	meyeri	R	R	R	R	R	3.3	2.0	Winter	
	Unknown	anathera	S	S	S	S	S	4.0	2.9	Winter	
707	Unknown	Unknown	R	R	R	R	R	2.5	2.2	Winter	
708	Unknown	Unknown	S	S	I	S	S	4.0	2.7	Winter	
709	Unknown	Unknown	S	S	I	S	S	3.8	2.5	Winter	
712	Unknown	Unknown	S	S	S	S	S	3.4	2.5	Winter	
118	Unknown	anathera	S	S	S	S	S	4.0	2.6	Spring	
370	Unknown	typica	S	S	I	I	S	4.0	2.9	Winter	
370.1	Unknown	typica	S	S	R	S	S	4.0	3.0	Winter	
373	Unknown	typica	S	S	S	I	R	2.1	1.8	Spring	
	Percent resistan	ce:	23	24	24	26	45				
/PM-1							R	1.0	1.5	Winter	
Cappelle-D	esprez						R	2.0	1.7	Winter	
Cerco							R	2.3	1.7	Winter	
Selection 1							S	3.7	2.4	Winter	
Chinese Sp							S	4.0	2.7	Spring	
-	LSD $(P = 0.01)$							0.7	0.7		

were identified by Strausbaugh and Murray (36).

The French cultivar Cappelle-Desprez was the first wheat cultivar reported to be resistant to eyespot (37). Although the gene conferring the resistance is located on chromosome 7A (21), the source of this gene is unknown. After VPM-1 was selected (25), it was used extensively to transfer resistance into wheat cultivars because its resistance is more effective than the resistance of Cappelle-Desprez. Genetic analysis of VPM-1 revealed that the resistance gene was located on chromosome 7D (8,13). Allan et al. (1,2) used VPM-1 to produce Madsen and Hyak, which were the first eyespot-resistant cultivars in the United States. The only cultivar thought to contain both the 7A and 7D resistance genes is Rendezvous (12); however, this genotype still sustains yield loss when disease is severe (12).

It is not clear at this point whether the new eyespot resistance genes in T. tauschii are allelic to, or separate genes from, those in commercial use. However, based on the GUS assay, accessions TA 1622 and TA 1624 from Azerbaijan were significantly (P = 0.01) more resistant to eyespot than VPM-1. These two accessions warrant further study.

Wheat breeding programs utilize both race-specific and nonspecific resistances to stripe rust; therefore it is important to continue the characterization of new genes even if they may be circumvented. There is concern that the eyespot pathogen may circumvent existing resistance genes and render them ineffective in the future. It is clear from the variation in these accessions that T. tauschii offers a diverse source of resistance for stripe rust and eyespot.

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