

# Improved Screening of the Arizona Population of *Fusarium oxysporum* f.sp. *lactucae* Final Report

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## Executive Summary

To examine genetic variation in *Fusarium oxysporum* f. sp. *lactucae* (FOL), the causal agent of Fusarium wilt of lettuce, a collection of isolates was recovered from symptomatic lettuce plants in Arizona. Additionally, isolates from California were added to the collection to determine if the Arizona isolates differed from the CA isolates. Over 150 isolates were assembled from 42 locations and production companies in AZ and CA isolates in 2021 and early 2022 (Table 1). The collection included reference isolates from California and also the original isolate of FOL first recovered from lettuce in the U.S. (KP02). Several isolates from Italy and the Netherlands, were examined including several FOL race 4 isolates. More recently, reference isolates of FOL race 2 and race 3 (designated *F. odoratissimum* and *F. cugenangense* respectively) were obtained for examination.

Isolates were tested for pathogenicity and grouped into vegetative compatibility groups (VCGs). Pathogenicity tests indicated that the isolates could be clearly identified as *Fusarium oxysporum* f.sp. *lactucae* (Table 2), and all AZ and CA isolates examined collected, including the original isolate of FOL from 2002 (KP02), belong to VCG 0300 (Table 1). Vegetative compatibility tests are a non-molecular test to assess relatedness of fungal isolates by examining the ability of two isolates to fuse. Isolates that share a VCG have a higher genetic similarity than isolates of another VCG. The four FOL races have been found to be genetically distinct, with each as a separate VCG subgroup (Fujinaga et al., 2005; Mbofung et al., 2007; Pintore et al., 2017).

A subset of isolates was examined for molecular diversity by examining sequence variation in the elongation factor target sequence (Figure 1). The majority of the isolates examined show little to no sequence variation. Interestingly, two isolates of FOL race 4 from Italy (GG41 and GG42) could be distinguished from the rest of the isolates examined based on vegetative compatibility (Table 3).

Based on the vegetative compatibility and molecular assays, the isolates representing the Arizona *Fusarium oxysporum* f.sp. *lactucae* population are highly similar. This suggests that changes in disease severity observed in fields in Arizona are due to an increase of the pathogen levels in the soil. However, changes in the native population may still occur, or material contaminated with a pathogen variant may enter the growing region.

Isolates have been collected in California that do not conform to the race 1 phenotype. Care should be taken to not introduce this pathogen variant into Arizona on plant material including seed, and in soil on equipment. Growers should continue to report unexpected disease incidence or severity, especially on cultivars with a history of tolerance or resistance to FOL. A robust pipeline for evaluating FOL populations is in development, and the pathogenicity of FOL isolates from Arizona and California continues to be examined outside of the scope of this project.

## References

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- Mbofung, G. Y., Hong, S. G., and Pryor, B. M. 2007. Phylogeny of *Fusarium oxysporum* f. sp. *lactucae* inferred from mitochondrial small subunit, elongation factor 1- $\alpha$ , and nuclear ribosomal intergenic spacer sequence data. Phytopathology 97:87-98.
- Pintore, I., Gilardi, G., Gullino, M., and Garibaldi, A. 2017. Analysis of vegetative compatibility groups of Italian and Dutch isolates of *Fusarium oxysporum* f. sp. *lactucae*. Journal of Plant Pathology (Rivista di patologia vegetale) 99:517-521.

**Table 1. Master list of isolates of *Fusarium oxysporum* f. sp. *lactucae* VCGs Updated: 06/27/2022**

Isolate <sup>a</sup>	VCG ID	Race	Collected <sup>b</sup>	Location	Company/ Contact
SS01	0300		10/26/20	AZ	1
SS05	0300		10/26/20	AZ	2
SS06			10/26/20	AZ	2
SS09	0300		10/26/20	AZ	3
SS10			10/26/20	AZ	4
SS12	0300		10/26/20	AZ	5
SS13	0300		10/26/20	AZ	5
SS14	0300		10/26/20	AZ	5
SS15	0300		10/26/20	AZ	6
SS16	0300		10/26/20	AZ	6
SS17	0300		10/26/20	AZ	6
SS18				CA	7
SS19	0300			CA	7
SS20	0300			CA	7
SS21	0300		11/04/20	AZ	8
SS22			11/04/20	AZ	8
SS23			11/04/20	AZ	8
SS24	0300		11/04/20	AZ	9
SS25	0300		11/04/20	AZ	9
SS26			11/04/20	AZ	9
SS27			11/04/20	AZ	10
SS28	0300		11/04/20	AZ	10
SS29			11/04/20	AZ	10
SS30			11/04/20	AZ	11
SS31	0300		11/04/20	AZ	11
SS32	0300		11/04/20	AZ	12
SS33	0300		11/04/20	AZ	12
SS34	0300		11/04/20	AZ	12
SS36	0300		11/04/20	AZ	13
SS37	0300		11/04/20	AZ	13
SS38			11/04/20	AZ	13
SS39	0300		11/04/20	AZ	14
SS40			11/04/20	AZ	14
SS41			11/04/20	AZ	14
SS42			11/04/20	AZ	15
SS43	0300		11/04/20	AZ	15
SS44	0300		11/04/20	AZ	15
SS45	0300		11/13/20	AZ	16
SS46	0300		11/13/20	AZ	16
SS47	0300		12/31/20	AZ	17
SS48	0300		12/31/20	AZ	17
SS49	0300		12/31/20	AZ	18
SS50	0300		12/31/20	AZ	18
SS51			12/31/20	AZ	19
SS52			12/31/20	AZ	19
SS53			12/31/20	AZ	20
SS54	0300		12/31/20	AZ	20
SS55			12/31/20	AZ	21
SS56	0300		12/31/20	AZ	21
SS57			12/31/20	AZ	22
SS58	3000		12/31/20	AZ	22
SS59			12/31/20	AZ	23
SS60			12/31/20	AZ	23
SS61	0300		03/15/21	AZ	24
SS62			03/15/21	AZ	24
SS63			03/15/21	AZ	24
SS64			03/15/21	AZ	24
SS65			03/15/21		25
SS66			03/15/21		25
SS67			03/15/21		25
SS68			03/15/21		25
SS69	0300		03/22/21		26

SS70	0300	03/22/21		26
SS71		03/22/21		26
SS72	0300	03/22/21		26
SS73		11/23/21		27
SS74		11/23/21		27
SS75	0300	11/23/21		27
SS76	0300	11/23/21		28
SS77	0300	11/23/21		28
SS78	0300	11/23/21		28
SS80				29
SS81	0300			29
SS82	0300	11/23/21		30
SS83	0300	11/23/21		30
SS84	0300	11/23/21		30
SS85	0300	11/23/21	AZ	31
SS86		11/23/21	AZ	31
SS87		11/23/21	AZ	31
JV 192	0300	01/08/20	AZ	32
JV 198	0300	01/08/20	AZ	33
JV 199	0300	01/08/20	AZ	33
JV 200	0300	01/08/20	AZ	34
JV 201	0300	01/08/20	AZ	35
JV 202	0300	01/08/20	AZ	36
Q1A1		02/26/20	AZ	36
Q1A2	0300	02/26/20	AZ	36
Q1B1	0300	02/26/20	AZ	37
Q3D1		02/26/20	AZ	37
Q3D2		02/26/20	AZ	37
Q4A1	0300	02/26/20		38
Q4D2		02/26/20		38
Q4E2		02/26/20		38
JQ11	0300	02/23/21		39
JQ12	0300	02/23/21		39
JQ21		02/23/21		39
JQ22	0300	02/23/21		40
JQ31	0300	02/23/21		40
JQ32		02/23/21		40
JQ41		02/23/21		41
JQ42	0300	02/23/21		41
JQ51	0300	02/23/21		41
JQ52		02/23/21		42
KP01		2004	Wellton, AZ	Kelly Paugh/UC Davis
KP02	0300	2002	Huron, CA	Kelly Paugh/UC Davis
KP03		2002	Yuma, AZ	Kelly Paugh/UC Davis
KP04	0300	2002	Huron, CA	Kelly Paugh/UC Davis
KP05		2002	Huron, CA	Kelly Paugh/UC Davis
KP06		2015	Davis, CA	Kelly Paugh/UC Davis
KP07	0300	2015	Watsonville, CA	Kelly Paugh/UC Davis
KP08		2016	Chualar, CA	Kelly Paugh/UC Davis
KP09		02/18/2022	Davis, CA	Kelley Paugh/UC Davis
GG11		1	Italy	Giovanna Gilardi-Agroinnova
GG12		1	France	Giovanna Gilardi-Agroinnova
GG13		1	Italy	Giovanna Gilardi-Agroinnova
GG41	0304?	4	Italy	Giovanna Gilardi-Agroinnova
GG42	0304?	4	Italy	Giovanna Gilardi-Agroinnova
GG43		4	Netherlands	Giovanna Gilardi-Agroinnova
GG44		4	Netherlands	Giovanna Gilardi-Agroinnova
OC01		11/10/2020	AZ	43
OC02	0300	12/03/2020	AZ	43
OC03	0300	12/03/2020	AZ	43

<sup>a</sup> Isolates with the nomenclature "SS" are from Dr. Stephanie Slinski, "KP" isolates are from Kelly Paugh, "GG" are isolates sent from Giovanna Gilardi-Agroinnova.

<sup>b</sup> Date collected correlates with the day the lettuce sample was taken from the field.

**Table 2. Mean disease severity in greenhouse inoculations tests with *Fusarium oxysporum* f. sp. *lactucae* on the cultivar Grizzly.**

Isolate	Mean Disease Severity (Scale – 0.0 - 4.0) <sup>1</sup>					
	Test 1 06/09/20	Test 2 06/19/20	Test 3 09/01/20	Test 4 09/25/20	Test 5 11/16/20	Test 6 12/20/20
H <sub>2</sub> O*	1.6	1.3	0.0	1.0	0.0	0.0
BR35B**	0.7	1.0	0.1	0.6	0.1	0.7
JV 202	3.6	2.7	0.9	2.1	2.8	3.1
Q3D1	2.0	1.9	1.7	2.9	2.2	3.1
Q3D2	3.3	1.9	1.6	1.6	2.1	3.1
Q4A1	3.3	3.1	2.3	2.4	2.8	3.1
JV 192	3.5	2.8	1.8	1.9	3.8	3.2
JV 199	2.5	2.5	0.2	2.6	3.4	3.4
Q4D2	0.8	2.1	1.8	2.4	1.2	3.4
JV 198	3.3	1.9	1.7	2.8	3.3	3.7
Q1A2	3.2	1.5	1.7	2.5	3.3	3.7
Q1B1	3.8	2.9	1.1	1.2	2.1	3.7
Q4E2	3.4	2.7	1.0	3.0	1.5	3.8
JV 200	2.8	2.6	0.6	1.2	3.4	4.0
JV 201	3.3	2.1	3.0	2.6	2.8	4.0
Q1A1	3.0	3.1	2.1	2.2	2.7	4.0

<sup>1</sup>The mean disease severity was 4-5 weeks post inoculation. Disease was rated on a scale of

0 – 4 with zero indicating no symptoms; 4 = Severe Fusarium wilt symptoms.

\* H<sub>2</sub>O water negative control.

\*\* BR35B is an isolate of *F. oxysporum* f. sp. *coriandrii* used as a negative control.

**Table 3. Reference isolates and passport information for isolates of FOL collected and characterized by the Correll Lab.**

Isolate (Correll Lab Code)	Received As Code	Collected	Location	VCG	Pathogenicity
JV198		01/08/20	AZ	0300	+
JV199		01/08/20	AZ	0300	+
JV201		01/08/20	AZ	0300	+
JV202		01/08/20	AZ	0300	+
244141*	F9501		Japan	Race 2	
744085*	N3		Japan	Race 3	
744086*	M7		Japan	Race 3	
Q1A2		02/26/20	AZ	0300	+
GG41	FL3/19		Italy	0304?	
GG42	FL6/19		Italy	0304?	
GG43	PD015/0470896		Netherlands	0304?	
GG44	PD015/04750888		Netherlands	0304?	
HS					
HS					
HS					

\* Isolate designation is the MAFF number listed by Naro Genebank and can be found at

[https://www.gene.affrc.go.jp/databases-micro\\_search\\_en.php](https://www.gene.affrc.go.jp/databases-micro_search_en.php)

References for Japanese isolates:

Race 2: Mbofung et al, 2007

Race 3: Fujinaga et al, 2001



Fig. 1. Phylogenetic tree based on Neighbor Joining (NJ) analysis of the elongation factor sequences with the isolates of *Fusarium oxysporum* f. sp. *lactucae* (FOL); the out-group BR35B is *F. oxysporum* f. sp. *coriandrii* (FOC) from cilantro. The isolate of 244121 is race 2, and the isolates of 744085 and 744086 are race 3 from Japan. Bootstrap values were labeled on the branch of the tree. Scale bars represents the average number of substitutions per site over time, and the numbers in bold on the figure are the disease severity ratings from pathogenicity tests on lettuce on a scale of 0.0 (no disease) to 4.0 (severe disease).