

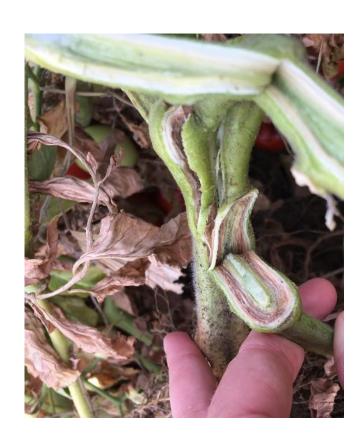
Fusarium oxysporum

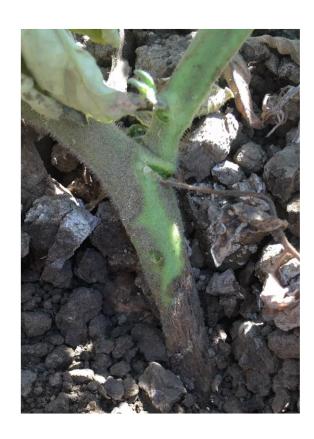
Fusarium wilt f. sp. lycopersici
Fol (race 3)

Fusarium crown and root rot f.sp. radicis-lycopersici Forl

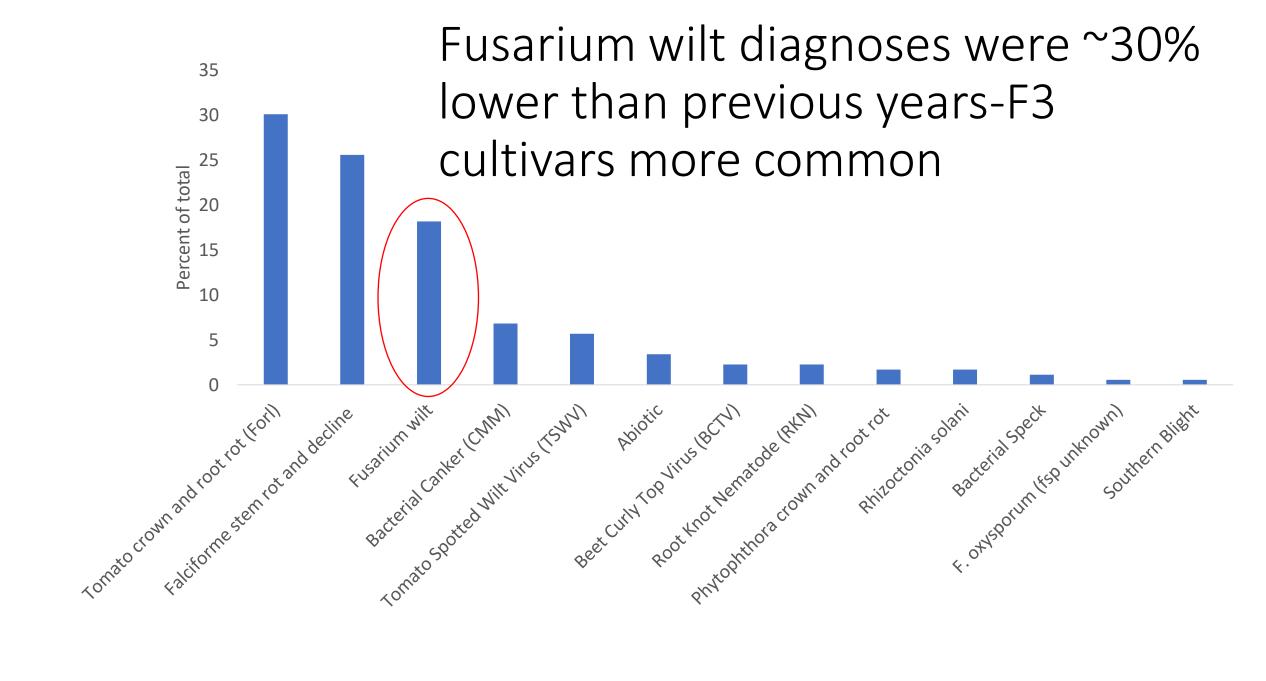


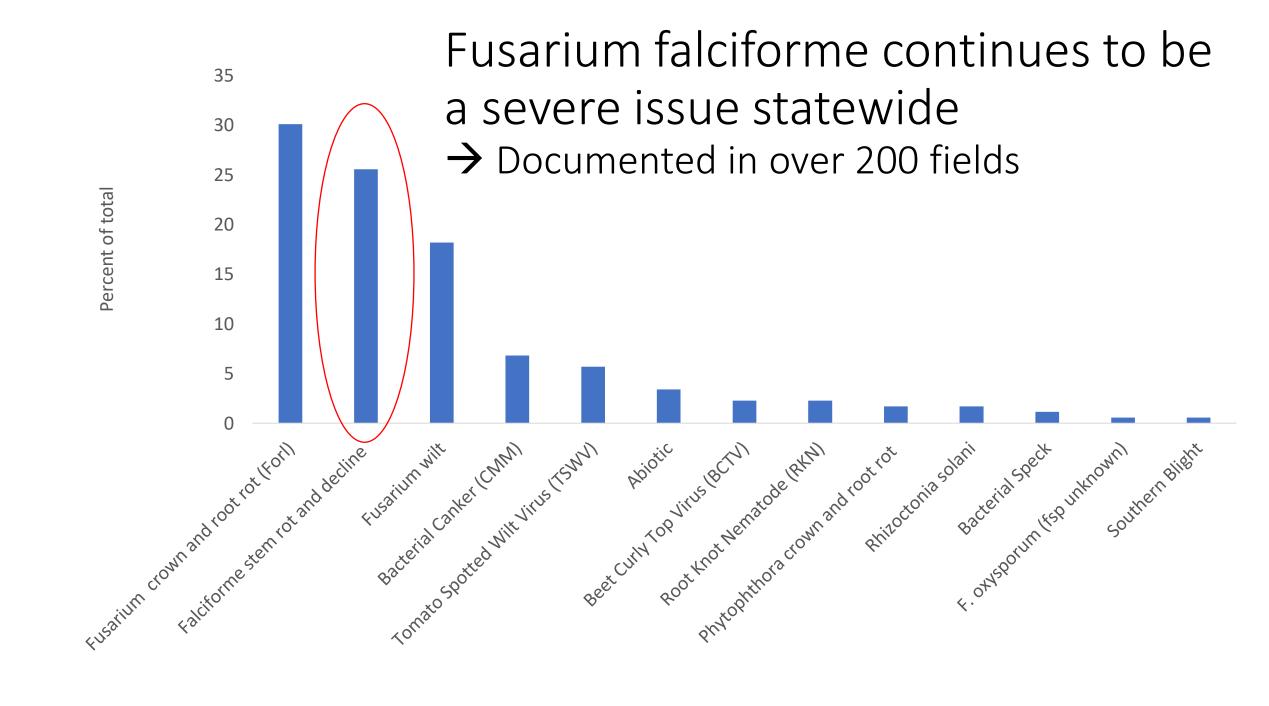
Stem rot and vine decline

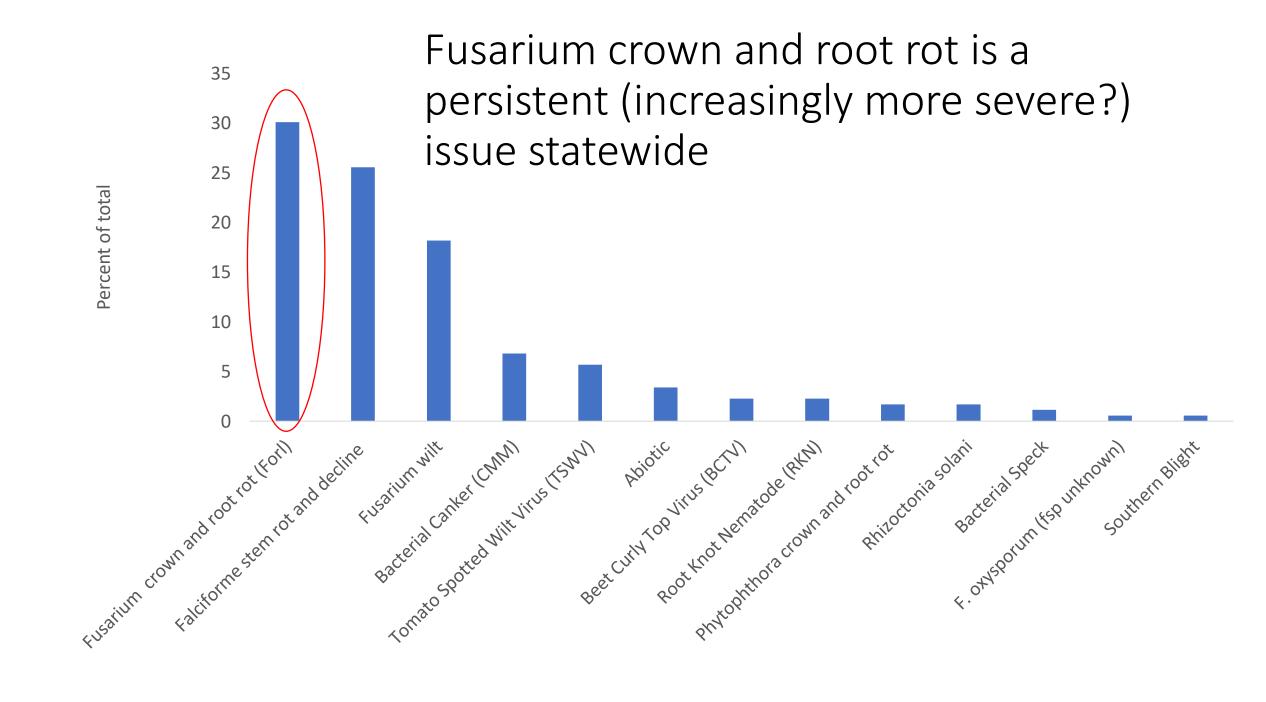




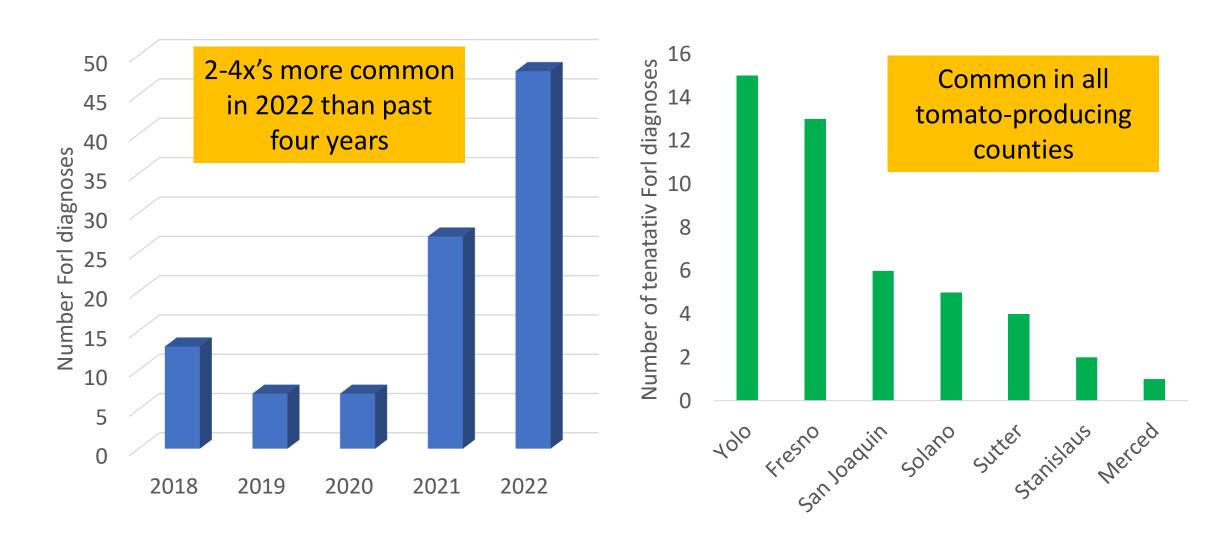








Fusarium crown and root rot: an increasing statewide issue



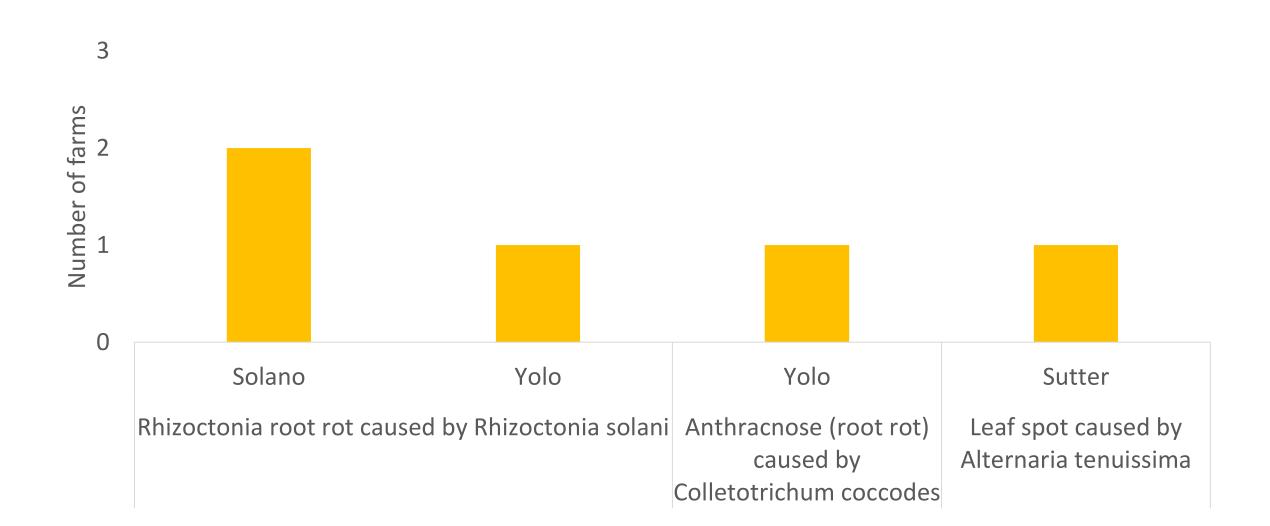
Irrigation shifts may in part underly increased occurrence



■ Grower ■ DI 100% 90% 80% 70% 60% 50% 40% 30% 20% Any Stem Rot stem rot pootrot pootrot

watered (at least 100% ET)

Putative new diseases detected in 2022



Upcoming: new UC IPM tomato disease diagnosis field guide

Diagnosing vine decline
and rot diseases of
tomatoes in the field

UC Davis 2022 Vegetable Disease Field Day Cassandra Swett, Bob Gilbertson

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Diagnóstico de decaimiento foliar y enfermedades de pudrición de tomates en el campo

UC Davis 2022 Vegetable Disease Field Day Cassandra Swett, Bob Gilbertson Traducido por: Johanna Del Castillo <u>Múnera</u> Department of Plant Pathology UC Davis



urly top disease (CTD)-beet urly top virus (BCTV)

ector: beet leafhopper

ymptoms

Observed early in the season often in fields near foothills Plants are stunted and dull-green Leaves: dull-green to yellow, crumple, curl upward or even roll, and swollen purple veins

Fruits: small and ripen prematurely

TD is sporadic but can cause economic oss in bad years; no resistant varieties ut known risk factors

Tomato necrotic spot diseasetomato necrotic spot (ToNSV)

A windborne pollen-transmitted virus introduced to tomato via thrips feeding Symptoms

- Generally seen early in the season
- Leaves are distorted and show brown necrotic spots and stems are necrotic
- Not economically important: tomato plants recover (defense response) and there is little within field spread
- More common in Northern Counties in 2022
- Detected with RT-PCR test



- TSWV (Sw5 gene)
 - Known to have resistance breaking strains (Gilbertson, Turini)

- TSWV (SW5 gene)
- Root knot nematode (Mi gene)

Root knot nematode resistance breaking common statewide-2021 survey (Hodson, Swett)

Statewide:

- RKN recovered from 27 fields planted to resistant cultivars
- 100% of tested isolates (18) were resistance-breaking (controlled temp)

%	Root	gal	ling
/0	NOOL	Sui	שו ווו

	County	Isolate	Celebrity' (<i>Mi</i> +)	Rutgers' (Mi-)
M. incognita	Yolo	139	33	31
	Yolo	140	44	32
	Yolo	143	23	27
	Yolo	144	35	33
	Yolo	145	19	28
	Yolo	146	30	37
	Yolo	213	6	7
	Yolo	R-R	43	44
	Solano	212	4.6	12
	Fresno	157	30	32
	Fresno	158	26	28
	Fresno	208	24	20
	Fresno	207	28	44
	Fresno	151	34	44
	Merced	183	28	38
	Sutter	A-S	46	37
M. javanica	Yolo	184	19	29
	Stanislaus	C-L	0	1

Controls

M. incognita M. incognita M. javanica M. javanica

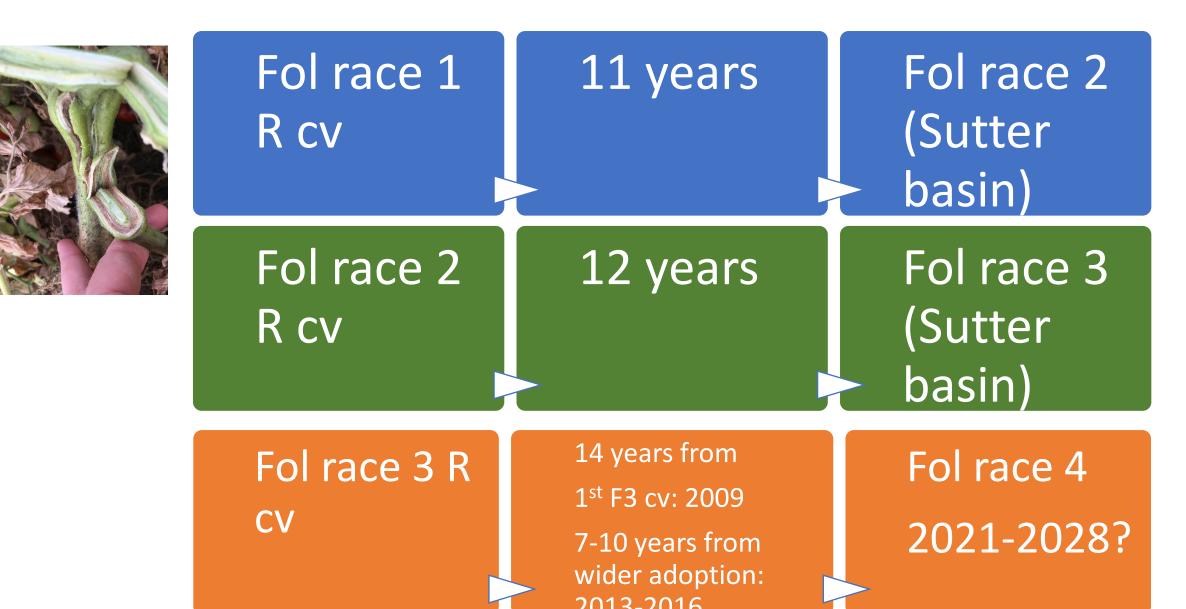
Hr3	36	29
13	0	47
VW5	33	25
VW4	5	42



- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
 - Resistance breaking is widespread
 - This is not a temperature-related issue-GH studies controlled for temperature
 - 60% of RB-RKN-infested fields also had one or more Fusarium disease
 - There are likely RKN interactions with Fusarium diseases

- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
- Fusarium wilt-race 3 (I3 gene, F3 cultivars)
 - No resistance-breaking race 4 known to be present in the state
 - Monitoring efforts aim to detect early emergence of race 4

Fusarium wilt resistance-breaking race timeline



Fusarium wilt in resistant cultivars

17 F3 fields in 5 years had Fol All were Fol race 3

Efforts are opportunistic: lack rigorous surveys

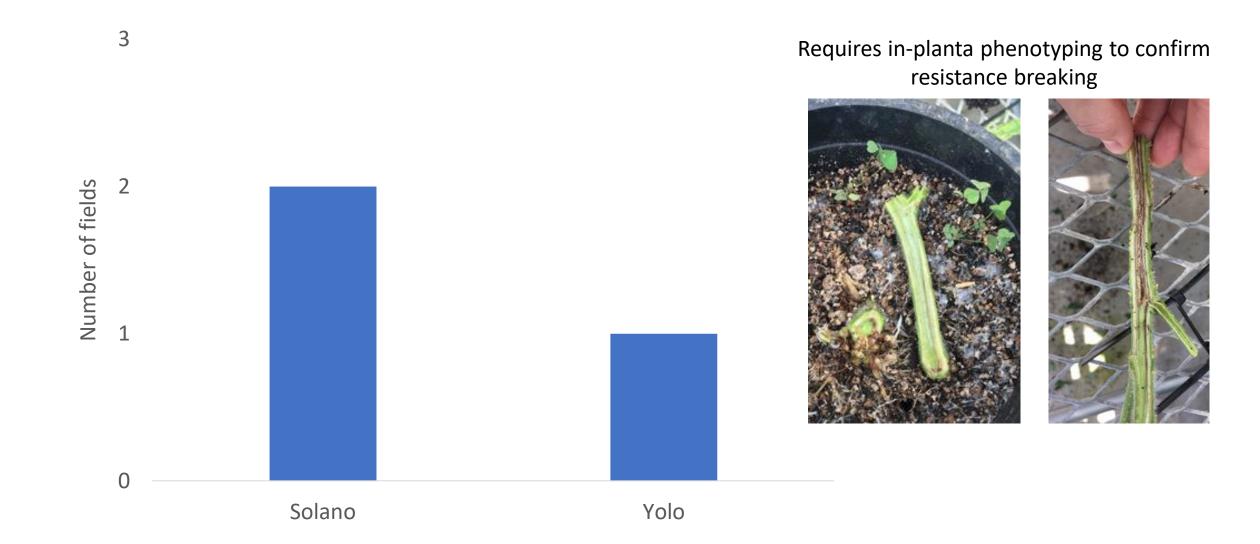
		No fields (percent)					
		Fol					
Year	Total	R1	R2	R3	R4	Forl	Non-Path
2017	2	0	0	2 (100%)	0	0	0
2018	11	0	0	11 (100%)	0	0	0
2019	0	0	0	0	0	0	0
2020	2	0	0	2 (100%)	0	0	0
2021	2	0	0	2 (100%)	0	0	0
Total	17	0	0	17 (71%)	0	0	0

- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
- Fusarium wilt race 3 (I3 gene, F3 cultivars)
 - No resistance breaking race 4 detected; have maybe five years.
 - Early detection critical to mitigate spread, preserving efficacy of F3 cultivars and providing time for development of new resistant cultivars

- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
- Fusarium wilt race 3 (I3 gene, F3 cultivars)
- Fusarium crown and root rot (Frl gene, Fr cultivars)

Potential Forl resistance-breaking detected for the first time

3 fields, H5522: the primary Fr CV in 2022



- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
- Fusarium wilt race 3 (I3 gene F3 cultivars)
- Fusarium crown and root rot (Frl gene, Fr cultivars)
 - Might have possible Forl race 2 (Frl resistance breaking race)
 - Testing is needed-may be non-pathogenic F. oxysporum

Diseases managed by single gene resistance

- TSWV (SW5 gene)
- Root knot nematode (Mi gene)
- Fusarium wilt race 3 (I3 gene F3 cultivars)
- Fusarium crown and root rot (Frl gene, Fr cultivars)

Early identification of resistance breaking is critical
Use of sanitation and other management methods will be important
to slow spread statewide

Enabling the breeding industry to develop new resistant materials

