2019 Flue-Cured Tobacco Production Guide Disease Resistance Tables

Table 3. Survival of selected flue-cured tobacco varieties in 2018 on-farm tests in black shank-infested fields in Virginia.

		% Healthy Plants				Black Shank
Variety	Ph_p					Yield
-	Gene	Mecklenburg	Brunswick	Mecklenburg-1	Average	Index ¹
PVH	+	87	98	100		97
1452					95	
CC 33	-	90	93	99	94	99
CC 1063	-	93	89	98	93	94
CC 143	-	95	81	97	91	97
PVH	+	77	88	98		95
1600					88	
CC 37	+	75	90	99	88	79
K 346	-	74	94	93	87	88
GL 395	-	74	87	100	87	85
NC 925	-	69	92	99	87	78
NC 196	+	77	84	96	86	83
NC 938	-	74	84	96	85	86
GF 318	+	67	90	97	85	90
PVH	+	3	55	96		48
2275					51	

¹ Yield indexes for Black Shank (race 1) = yield index without black shank (from the 2018 Virginia OVT test at the Southern Piedmont AREC, Blackstone) multiplied by the average proportional survival from the three on-farm black shank resistance tests conducted in Virginia in 2018.





Virginia Cooperative Extension programs and employment are open to all, regardless of age, color, disability, gender, gender identity, gender expression, national origin, political affiliation, race, religion, sexual orientation, genetic information, veteran status, or any other basis protected by law. An equal opportunity/affirmative action employer.

Disclaimer: Commercial products are named in this publication for informational purposes only. Virginia Cooperative Extension does not endorse these products and does not intend discrimination against other products which also may be suitable.

2019 Flue-Cured Tobacco Production Guide Disease Resistance Tables

Table 4. Flue-cured tobacco variety reactions to Black Shank.

Varieties with the Php gene¹: (Race 1)² Black Shank (Race 1) No Black (Race 1) SP 225 88 77 . NC 196 67 70 104 PVH 1452 69 68 98 CC 67 63 59 93 CC 700 58 58 100 PVH 1600⁴ 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ 89 90 102 NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 9			Relative Yield Index ³	
Varieties with the Php gene¹: (Race 1)² (Race 1) Shank SP 225 88 77 . NC 196 67 70 104 PVH 1452 69 68 98 CC 67 63 59 93 CC 700 58 58 100 PVH 1600⁴ 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ 89 90 102 NC 938 89 90 102 NC 925 83 83 99				
SP 225 88 77 NC 196 67 70 104 PVH 1452 69 68 98 CC 67 63 59 93 CC 700 58 58 58 100 PVH 16004 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2215 11 11 98 Varieties without the Php gene1 NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112		% Survival	Shank	No Black
NC 196 67 70 104 PVH 1452 69 68 98 CC 67 63 59 93 CC 700 58 58 100 PVH 1600 ⁴ 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35	Varieties with the <i>Php</i> gene ¹ :	(Race 1) ²	(Race 1)	Shank
PVH 1452 69 68 98 CC 67 63 59 93 CC 700 58 58 58 100 PVH 1600 ⁴ 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35	SP 225	88	77	
CC 67 CC 700 S8 S8 S8 S9 100 PVH 1600 ⁴ S8 S9 101 CC 37 S3 S1 96 NC 71 47 S0 106 GF 318 45 47 103 NC 299 44 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 PVH 2275 11 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 66 100 GL 395 CC 13 48 FVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35	NC 196	67	70	104
CC 700	PVH 1452	69	68	98
PVH 1600 ⁴ 58 59 101 CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 11 98 Varieties without the Php gene ¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35	CC 67	63	59	93
CC 37 53 51 96 NC 71 47 50 106 GF 318 45 47 103 NC 299 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 4	CC 700	58	58	100
NC 71	PVH 1600⁴	58	59	101
GF 318 45 47 103 NC 299 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 60 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	CC 37	53	51	96
NC 299 44 44 101 NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	NC 71	47	50	106
NC 72 38 40 101 NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	GF 318	45	47	103
NC 297 35 36 103 CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	NC 299	44	44	101
CC 27 34 36 105 PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	NC 72	38	40	101
PVH 2310 25 25 102 PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	NC 297	35	36	103
PVH 2275 11 11 98 Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	CC 27	34	36	105
Varieties without the Php gene¹ NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	PVH 2310	25	25	102
NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	PVH 2275	11	11	98
NC 938 89 90 102 NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112	Varieties without the <i>Php</i> gene	,1		
NC 925 83 83 99 CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112			90	102
CC 1063 84 83 99 CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112				
CC 143 79 82 104 K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112				
K 346 82 77 94 NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112				
NC 606 69 68 98 CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112				
CC 33 66 66 100 GL 395 64 61 95 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112			68	98
GL 395 CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35				
CC 13 48 50 104 PVH 2254 48 47 98 GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112		64	61	95
GL 26H 44 47 108 PVH 2110 43 46 108 CC 35 40 45 112		48	50	104
PVH 2110 43 46 108 CC 35 40 45 112	PVH 2254	48	47	98
CC 35 40 45 112	GL 26H	44	47	108
CC 35 40 45 112		43	46	
	CC 35	40	45	
		32	34	107

¹ Varieties with the *Php* gene possess very high resistance to race 0 of the black shank pathogen. Resistance to race 0 in varieties without the *Php* gene is similar to or higher than that to race 1. ² Average % Survival near 2nd harvest without a soil fungicide. Results are averages from 10 field

experiments conducted in 2010-2015 and 2017 by Clemson and by North Carolina State Universities as part of the Regional Flue-Cured Tobacco Variety Evaluation Program.

³ Relative Yield Index = yield of each cultivar relative to the yield of all other cultivars in the experiment(s). Yield indexes for "No Black Shank" = average relative yield from the 2010-2018 Virginia OVT tests conducted at the Southern Piedmont AREC, Blackstone. Yield indexes for "Black Shank (race 1)" = yield index without black shank multiplied by the average proportional survival near 2nd harvest.

2019 Flue-Cured Tobacco Production Guide Disease Resistance Tables

Table 5. Performance of selected flue-cured tobacco varieties in 2018 Virginia Tech on-farm tests for resistance to Granville Wilt.

	% Healthy Plants					Granville	
					4-test	Wilt Yield	
Variety	Baskerville-1	Alberta	Dolphin	Baskerville-2	Average	Index 1	
PVH 1600	89	94	93	81	89	96	
PVH 1452	84	93	94	84	89	90	
GL 939	78	94	94		89		
PVH 1063	84	93	89	86	88	89	
CC 27	82	79	92	91	86	95	
NC 606	75	89	90	84	85	82	
GL 395	77	89	85	81	83	81	
CC 37	78	91	89	74	83	75	
NC 938	73	76	91	93	83	84	
CC 143	63	84	87	89	81	86	
NC 299	60	88	86	84	80	70	
NC 196	61	81	88	87	79	77	
CC 33	55	82	90	83	78	81	
NC 925	42	91	86	77	74	66	
CC 35	3	16	55	71	36	44	

¹ Yield Index with Granville Wilt = proportion of plants surviving for each variety multiplied by the relative yield for that cultivar in the 2018 Virginia OVT test at the Southern Piedmont AREC, Blackstone.

2019 Flue-Cured Tobacco Production Guide Disease Resistance Tables

Table 6. Reactions of flue-cured tobacco varieties to Granville Wilt.

		Relative Yield Index ³	
		With	
	% Survival ²	Granville	No Granville
Varieties with the <i>Php</i> gene ¹ :	2010,2012-2017	Wilt	Wilt
CC 37	80	77	96
CC 27	74	77	105
CC 67	78	72	93
PVH 1452	68	67	98
PVH 2275 ⁴	57	56	98
NC 297	55	56	103
NC 196	54	56	104
GF 318	53	54	103
NC 72	52	53	101
NC 299	50	51	101
CC 700	48	48	100
PVH 1600	47	47	101
PVH 1118	45	45	100
NC 71	36	38	106
PVH 2310	36	37	102
Varieties without the Php gene1			
NC 606	74	73	98
CC 1063	70	70	99
K 346	70	65	94
GL 939	69	65	95
CC 143	61	63	104
CC 33	62	62	100
NC 938	56	57	102
GL 395	58	56	95
CC 13	53	56	104
PVH 2110	52	56	108
GL 26H⁴	52	55	108
PVH 2254 ⁴	54	53	98
NC 925⁴	47	47	99
K 326	37	39	107
CC 35	19	22	112

¹ Varieties with the *Php* gene possess very high resistance to race 0 of the black shank pathogen. Resistance to race 0 in varieties without the *Php* gene is similar to or higher than that to race 1. ² Average % Survival near 2nd harvest without soil fumigation. Results are averages from 5 field experiments conducted in 2010 and 2012-2017 by Clemson University as part of the Regional Flue-Cured Tobacco Variety Evaluation Program.

³ Relative Yield Index = yield of each cultivar relative to the yield of all other cultivars in the experiment(s). Yield indexes for "No Granville Wilt" = average relative yield from the 2010-2018 Virginia OVT tests at the Southern Piedmont AREC, Blackstone. Yield indexes for "with Granville Wilt" = yield index without Granville wilt multiplied by average % Survival.

⁴ Ratings based on limited data available.

2019 Flue-Cured Tobacco Production Guide Disease Resistance Tables

	Resistance Rating ¹				Nematodes		
	Black Shank			Root-	t-Knot		Tobacco
	Ph gene		Granville	М.	Other	Tobacco	Mosaic
Variety	(race 0 only)2	Race 1	Wilt	incognita	species ³	Cyst	Virus
CC 13	-	48	53	+	+	-	-
CC 27	+	34	74	+	-	+	+
CC 33	-	66	62	+	+	-	-
CC 35	-	40	19	+	+	-	-
CC 37	+	53	80	+	+	+	+
CC 67	+	63	78	+	-	+	+
CC 143	-	79	61	+	-	-	-
CC 700	+	58	48	+	-	+	-
CC 1063	-	84	70	+	-	-	-
GF 318	+	45	53	+	-	+	-
GL 26H	-	44	52	+	-	-	+
GL 395	-	64	58	+	-	-	-
K 326	-	32	37	+	-	-	-
K 346	-	82	70	+	-	-	-
NC 71	+	47	36				
NC 72	+	38	52	+	-	+	-
NC 196	+	67	54	+	-	+	-
NC 297	+	35	55	+	-	+	+
NC 299	+	44	50	+	-	+	-
NC 606	-	69	74	+	-	-	-
NC 925	-	83	47	+	-	-	-
NC 938	-	89	56	+	-	-	-
PVH 1118	+	61	45	+	-	+	-
PVH 1452	+	69	68	+	-	+	-
PVH 1600	+	58	47	+	-	+	-
PVH 2110	-	43	52	+	-	-	-
PVH 2254	-	48	54	+	-	-	+
PVH 2275	+	11	57	+	+	+	+
PVH 2310	+	25	36	+	-	-	+

¹Resistance rating = average % plants still alive near 2nd harvest, without a soil fungicide or fumigant. See Tables 3 and 4 for more detailed information.

² Varieties with the *Php* gene are almost immune to race 0 of the black shank pathogen; resistance to race 0 without the *Php* gene is at least as high as resistance to race 1.

³"Other species" of root-knot nematode include *Meloidogyne arenaria* (peanut root-knot nematode) or *M. javanica* (Javanese root-knot nematode. These other species are now common in Virginia.

⁴ Ratings based on limited data available.