

Supplemental figures and tables

Figure S1. Molecular models of Prothioconazole

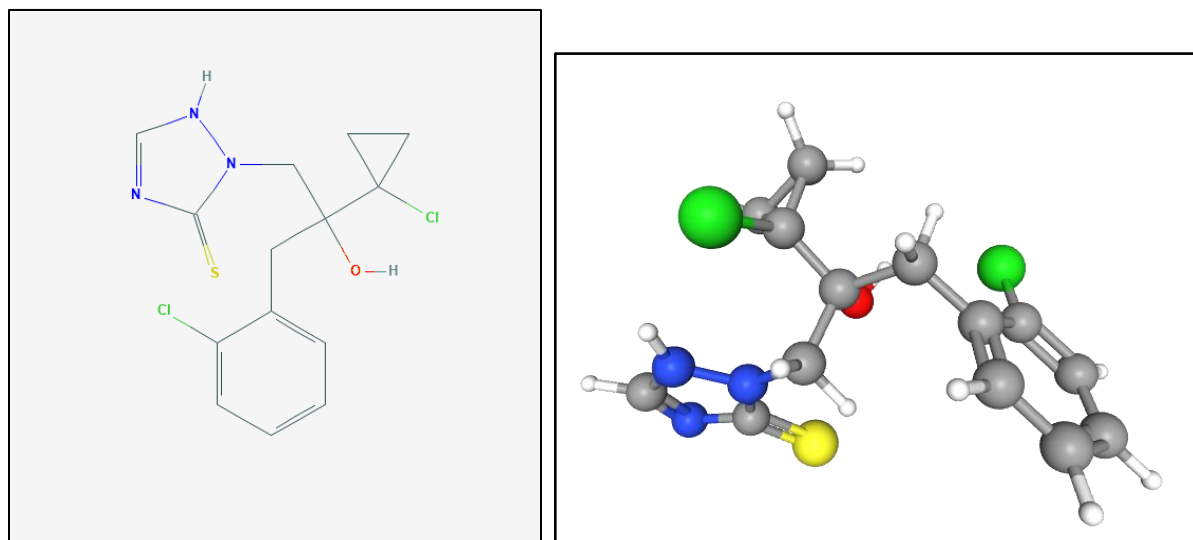


Figure S2. Nucleotide sequence of sensitive and resistant isolates CYP51A coding region

Sensitive1	ATGTTCTCACTCCTATACTACCCTCTATGGGCCTTTGCTTCCTGCCTAGT	50
Resistant1	ATGTTCTCACTCCTATACTACCCTCTATGGGCCTTTGCTTCCTGCCTAGT	50
Sensitive51	TATCATCACTCTCAACGTCTTATACCAGAAGCTCCCTCGAAATGCCAACG	100
Resistant51	TATCATCACTCTCAACGTCTTATACCAGAAGCTCCCTCGAAATGCCAACG	100
Sensitive101	AACCTCCGTTAGTGTTCCACTGGCTTCCATTCGTTGGGAATGCTGTTGCT	150
Resistant101	AACCTCCGTTAGTGTTCCACTGGCTTCCATTCGTTGGGAATGCTGTTGCT	150
Sensitive151	TATGGACTCGACCCTTATGGTTTCTTTGTGAAGTGTCGAGAAAAGCACGG	200
Resistant151	TATGGACTCGACCCTTATGGTTTCTTTGTGAAGTGTCGAGAAAAGCACGG	200
Sensitive201	CGATGTCTTCACCTTTATCCTCTTCGGTCGAAAAATCGTTGCCTGTCTTG	250
Resistant201	CGATGTCTTCACCTTTATCCTCTTCGGTCGAAAAATCGTTGCCTGTCTTG	250
Sensitive251	GTGTTGACGGCAATGACTTTGTTCTCAACAGTCGAATTCAGGACGCCAAC	300
Resistant251	GTGTTGACGGCAATGACTTTGTTCTCAACAGTCGAATTCAGGACGCCAAC	300
Sensitive301	GCCGAAGAAATCTACAGTCCATTGACAACGCCTGTCTTTGGTAGTGATGT	350
Resistant301	GCCGAAGAAATCTACAGTCCATTGACAACGCCTGTCTTTGGTAGTGATGT	350
Sensitive351	CGTATACGATTGTCCCAACTCGAAGCTCATGGAGCAAAAGAAGTTTGTCA	400

Resistant351	CGTATACGATTGTCCCAACTCGAAGCTCATGGAGCAAAGAAGTTTGTCA	400
Sensitive401	AGTTTGGCCTTACACAAAAGGCTCTCGAGTCCCATGTCCAGTTGATCGAG	450
Resistant401	AGTTTGGCCTTACACAAAAGGCTCTCGAGTCCCATGTCCAGTTGATCGAG	450
Sensitive451	CGAGAGGTTCTGGAGTACATCCAAGCTGTACCTTCATTCTCTGGAAAGTC	500
Resistant451	CGAGAGGTTCTGGAGTACATCCAAGCTGTACCTTCATTCTCTGGAAAGTC	500
Sensitive501	TGGCACAGTTGATGTATCCAAGGCAATGGCTGAGATAACCATCTTCACTG	550
Resistant501	TGGCACAGTTGATGTATCCAAGGCAATGGCTGAGATAACCATCTTCACTG	550
Sensitive551	CTGCTCGCTCTCTGCAGGGCGAAGAAGTTCGACGGAAGCTTACAGCTGAG	600
Resistant551	CTGCTCGCTCTCTGCAGGGCGAAGAAGTTCGACGGAAGCTTACAGCTGAG	600
Sensitive601	TTTGCAGCTCTGTATCATGACCTTGACCTAGGCTTCACTCCTGTAAACTT	650
Resistant601	TTTGCAGCTCTGTATCATGACCTTGACCTAGGCTTCACTCCTGTAAACTT	650
Sensitive651	CCTGTTCCCTTGGCTACCTTTGCCTCATAACCGACGTCGAGATGCTGCTC	700
Resistant651	CCTGTTCCCTTGGCTACCTTTGCCTCATAACCGACGTCGAGATGCTGCTC	700
Sensitive701	ATGCAAAGATGAGAGAGATCTACATGGACATCATTAAACGAACGAAGAAGA	750
Resistant701	ATGCAAAGATGAGAGAGATCTACATGGACATCATTAAACGAACGAAGAAGA	750
Sensitive751	GGCGTAGGGGACTTGGAGAAAGGAACTGACATGATCGCCAACCTGATGAA	800
Resistant751	GGCGTAGGGGACTTGGAGAAAGGAACTGACATGATCGCCAACCTGATGAA	800
Sensitive801	TTGCGAGTACAAAAACGGGCAGCCGATTCCGGACAAAGAGATCGCGTACA	850
Resistant801	TTGCGAGTACAAAAACGGGCAGCCGATTCCGGACAAAGAGATCGCGTACA	850
Sensitive851	TGATGATCACTCTTCTCATGGCTGGACAACACTCTTCGTCATCTGCTAGT	900
Resistant851	TGATGATCACTCTTCTCATGGCTGGACAACACTCTTCGTCATCTGCTAGT	900
Sensitive901	TCATGGATCATACTACATCTGGCTTCATCCACTGACATTGCTGAGGAACT	950
Resistant901	TCATGGATCATACTACATCTGGCTTCATCCACTGACATTGCTGAGGAACT	950
Sensitive951	CTACCAAGAGCAACTCATTAACCTTGAGTGCCGATGGTGTTCTCCCTCCCC	1000
Resistant951	CTACCAAGAGCAACTCATTAACCTTGAGTGCCGATGGTGTTCTCCCTCCCC	1000
Sensitive1001	TTCAGTACTCCGATCTCGACAAGCTTCCCCTTCTTCAGAATGTCGTCAA	1050
Resistant1001	TTCAGTACTCCGATCTCGACAAGCTTCCCCTTCTTCAGAATGTCGTCAA	1050
Sensitive1051	GAAACACTCCGTGTTTATTCTTCCATTCACTCCATTCTGCGAAAGGTTAA	1100
Resistant1051	GAAACACTCCGTGTTTATTCTTCCATTCACTCCATTCTGCGAAAGGTTAA	1100

Sensitive1101	AAGACCTATGCAAGCAACTGGATCACCTTACACCATCACCACAGACAAGG	1150
Resistant1101	GAGACCTATGCAAGCAACTGGATCACCTTACACCATCACCACAGACAAGG	1150
Sensitive1151	TTCTCCTCGCTTCACCAACTGTTACAGCGTTGAGTGAAGAACTTCACA	1200
Resistant1151	TTCTCCTCGCTTCACCAACTGTTACAGCGTTGAGTGAAGAACTTCACA	1200
Sensitive1201	GACGCCCAAAGATGGAATCCTCATCGGTGGGATAACAAACCCAGGAGGA	1250
Resistant1201	GACGCCCAAAGATGGAATCCTCATCGGTGGGATAACAAACCCAGGAGGA	1250
Sensitive1251	GGCCGTGACGGACGATGTCATTGACTACGGCTACGGCGCTGTTTCTAAAG	1300
Resistant1251	GGCCGTGACGGACGATGTCATTGACTACGGCTACGGCGCTGTTTCTAAAG	1300
Sensitive1301	GAACGAAGAGCCCATACTTACCCTTTGGCGCTGGTCGGCATCGCTGCATC	1350
Resistant1301	GAACGAAGAGCCCATACTTACCCTTTGGCGCTGGTCGGCATCGCTGCATC	1350
Sensitive1351	GGGGAGAAGTTTGCTTATGTCAACTTGGGCGTTATCGTCGCGACTTTGGT	1400
Resistant1351	GGGGAGAAGTTTGCTTATGTCAACTTGGGCGTTATCGTCGCGACTTTGGT	1400
Sensitive1401	GCGCAACTTCAGACTGTGCGACTCTTGATGGCAAGCCTGGTGTTCCGGCAA	1450
Resistant1401	GCGCAACTTCAGACTGTGCGACTCTTGATGGCAAGCCTGGTGTTCCGGCAA	1450
Sensitive1451	CTGACTACACTTCTCTCTTCTCAAGGCCAGCCCAACCTGCATACATAAAC	1500
Resistant1451	CTGACTACACTTCTCTCTTCTCAAGGCCAGCCCAACCTGCATACATAAAC	1500
Sensitive1501	TGGGAGCGCAGGAGGGCTTAA	1521
Resistant1501	TGGGAGCGCAGGAGGGCTTAA	1521

Figure S3. Amino Acid sequence of sensitive and resistant isolates

Sensitive1	MFSLLYYPLWAFASCLVIIITLNVLYQKLPRNANEPLVFWHLFPVGNVA	50
Resistant1	MFSLLYYPLWAFASCLVIIITLNVLYQKLPRNANEPLVFWHLFPVGNVA	50
Sensitive51	YGLDPYGFFVKCREKHGDVFTFILFGRKIVACLVDGNDFVLNSRIQDAN	100
Resistant51	YGLDPYGFFVKCREKHGDVFTFILFGRKIVACLVDGNDFVLNSRIQDAN	100
Sensitive101	AEEIYSPLTTPVFGSDVVYDCPNSKLMEQKKFVKFGLTQKALESHVQLIE	150
Resistant101	AEEIYSPLTTPVFGSDVVYDCPNSKLMEQKKFVKFGLTQKALESHVQLIE	150
Sensitive151	REVLEYIQAVPSFSGKSGTVDVSKAMAEITIFTAARSLQGEEVRRKLTAE	200
Resistant151	REVLEYIQAVPSFSGKSGTVDVSKAMAEITIFTAARSLQGEEVRRKLTAE	200

Sensitive201 FAALYHDLDLGFTPVNFLFPWLPLPHNRRRDAAHAKMREIYMDIINERRR 250
 |||||
 Resistant201 FAALYHDLDLGFTPVNFLFPWLPLPHNRRRDAAHAKMREIYMDIINERRR 250

Sensitive251 GVGDLKGTDMIANLMNCEYKNGQPIPDKEIAHMMITLLMAGQHSSSSAS 300
 |||||
 Resistant251 GVGDLKGTDMIANLMNCEYKNGQPIPDKEIAHMMITLLMAGQHSSSSAS 300

Sensitive301 SWIILHLASSTDIAEELYQEQLINLSADGVLPPPLQYSDLDKPLQLNVVK 350
 |||||
 Resistant301 SWIILHLASSTDIAEELYQEQLINLSADGVLPPPLQYSDLDKPLQLNVVK 350

Sensitive351 ETLRVHSSIHSILRKVKRPMQATGSPYTITTDKVLLASPTVTALSEEHFT 400
 |||||
 Resistant351 ETLRVHSSIHSILRKVKRPMQATGSPYTITTDKVLLASPTVTALSEEHFT 400

Sensitive401 DAQRWNPHRWDNKPQEEAVTDDVIDYGYGAVSKGTKSPYLPFGAGRHRCI 450
 |||||
 Resistant401 DAQRWNPHRWDNKPQEEAVTDDVIDYGYGAVSKGTKSPYLPFGAGRHRCI 450

Sensitive451 GEKFAYVNLGVIVATLVRNFRLLSTLDGKPGVPATDYTSLFSRPAQPAYIN 500
 |||||
 Resistant451 GEKFAYVNLGVIVATLVRNFRLLSTLDGKPGVPATDYTSLFSRPAQPAYIN 500

Sensitive501 WERRRA 506
 |||||
 Resistant501 WERRRA 506

Table S1. List of primers used in this study.

Primer name	Sequence	Assay	Conditions	expected size (bp)	Source
FCYPA1upF	GCTTACGATCGGAGAAGAACA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	336	This study
FCYPA1upR	AGGCCCATAGAGGGTAGTATAG				
FCYPA2upF	CCTTTGCTCCTGCCTAGTT	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	347	This study
FCYPA2upR	GAGCTTCGAGTTGGGACAAT				
FCYPB1upF	GTGTTTGACCGTTGTGTTGAG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 57 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	364	This study
FCYPB1upR	CATGGACGGTTCCTGGAAATA				
FCYPB2upF	TATTTCCAGGAACCGTCCATG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 60 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	393	This study
FCYPB2upR	CGTTCCGTTTCGAAGGATGA				
FCYPC1upF	CCAGTACATAATAGCAGGAGTG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	349	This study
FCYPC1upR	AAATTGGTCGCTCTGACTCAC				
FCYPC2upF	GTGAGTCAGAGCGACCAATTT	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 59 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	408	This study
FCYPC2upR	GGTGTCTGGGATGAGGATTTG				
FCypA1F	TGAGGACGCGAATCCTTCTG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 59 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	692	This study
FCypA1R	TGCTCCATGAGCTTCGAGTT				

FCypA2F	TCCCAACTCGAAGCTCATGG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	759	This study
FCypA2R	CCAGGTGCTTGCATAGGTCT				
FCypA3F	CCTATGCAAGCACCTGGATCA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	678	This study
FCypA3R	GTGGAATTGTGCAAATAGGGCA				
FCypA4F	TCCACCTCTACTGTTGCGAA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	680	This study
FCypA4R	CGACCCCGCTTATACCAAGG				
FCypA5F	CCTTGGTATAAGCGGGGTCTG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	712	This study
FCypA5R	ACCTACCCGCTTCTTGTTT				
FOCYP51Bpyes2-F	GGGTCTCCTCCAAGAACTT	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	692	Zheng et al. (2018) & This study
FCypB1R	CGAACCTCGGCGGAGATAAT				
FCypB2F	CGTGCCTATTATCTCCGCCG	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	672	This study
FCypB2R	ACAGGCATGGGAGACTTGAC				
FCypB3F	GTCAAGTCTCCCATGCCTGT	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	444	This study & Zheng et al. (2018)
FoCYP51Bpyes2-R	CTACTGCTGGCGTCTCTC				
FCypC1F	AGCACATTCGCAACCCTGTA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	675	This study
FCypC1R	TTCATCACGCCGAAGCCATA				
FCypC2F	TTATGGCTTCGGCGTGATGA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	691	This study
FCypC2R	CCGGCAGTCCAGGTATCTTTT				
FCypC3F	AAGATACCTGGACTGCCGGA	PCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	505	This study
FCypC3R	CACTGTTGGGACGCATCTA				
FCAq1F	TGGCTACCTTTGCCTCATAAC	qPCR	95 °C for 3 mins, 35 cycles of 95 °C for 30s, 58 °C for 30s, and 72 °C for 1 min, then 72 °C for 6 mins	120	This study
FCAq1R	GTCAGTTCTTTCTCCAAGTCC				
Fn-1	TACCACTTGTTGCCTCGGC	qPCR	95°C for 2 mins, 40 cycles of 95°C for 10s and 60°C for 40s and a temperature ramp of 0.2°C/s	327	Zhang et al. (2006)
Fn-2	TTGAGGAACGCGAATTAAC				

Table S2. Resistant phenotype and the relative expression of the FON mutants

Resistant Group	EC ₅₀	Resistance Factor	Relative Expression
Highly Resistant 1	47.679	9.562767	23.10286712836
Highly Resistant 2	172.228	34.5430113	38.0546276800871
Highly Resistant 3	184.78	37.0605106	46.6886510156855
HR mean	108.33	21.7272709	35.94871527
Intermediately Resistant 1	26.365	5.28791191	12.5099142900575
Intermediately Resistant 2	36.45	7.31061594	27.8576180254759

Intermediately Resistant 3	33.321	6.68304619	14.1232479406504
IR mean	31.77	6.37196895	18.16359342
Sensitive 1	4.217	0.84578511	8.75434961008591
Sensitive 2	6.969	1.39774163	11.79415374
Sensitive 3	4.079	0.81810706	4.642816
S mean	4.9859	1	8.39710634

Supplementary Table S3. Raw results from the greenhouse bioassay

Treatment	Plant 1	Plant 2	Plant 3	Plant 4	Plant 5	Plant 6	Plant 7	Plant 8	Plant 9	Plant 10	Plant 11	Plant 12	Plant 13
0 (NC)	0	0	3	3	0	3	0	0	3	3	3	7	9
1 (S)	9	5	5	3	3	3	3	0	5	3	7	5	3
2 (HR)	5	3	5	5	9	0	3	5	9	3	9	5	5
3 (IR)	3	3	5	3	3	5	0	3	3	5	9	9	7
4 (S-Pro)	7	0	0	3	0	3	0	3	3	0	3	9	5
5 (HR-Pro)	7	3	3	3	3	3	0	0	5	3	3	3	9
6 (IR-Pro)	9	3	3	3	0	0	0	3	0	0	7	7	7
0 (NC)	0	0	3	3	0	3	0	0	3	3	3	7	9
Treatment	Plant 14	Plant 15	Plant 16	Plant 17	Plant 18								
0 (NC)	9	0	0	0	3								
1 (S)	3	0	5	0	3								
2 (HR)	7	3	3	3	5								
3 (IR)	9	0	0	5	5								
4 (S-Pro)	3	3	0	3	3								
5 (HR-Pro)	3	5	3	3	5								
6 (IR-Pro)	3	3	3	0	3								
0 (NC)	9	0	0	0	3								