

FRAC Classification of Fungicides

Fungal control agents by cross resistance pattern and mode of action 2021 (www.frac.info)

A: Nucleic Acids Metabolism

A1: RNA polymerase I
4: PA-fungicides (Phenylamides)

A2: adenosine-deaminase
8: hydroxy (2-amino)-pyrimidines

A3: DNA / RNA synthesis (prop.)
32: heteroaromatics

A4: DNA topoisomerase type II (gyrase)
31: carboxylic acids

B: Cytoskeleton and Motor Proteins

B1: β -tubulin assembly in mitosis
(= Methyl Benzimidazole Carbamates)
1: MBC fungicides

B2: β -tubulin assembly in mitosis*
10 N-phenyl carbamates
* negative cross-resistance to B1

B3: β -tubulin assembly in mitosis
22 benzamides and thiazole carboxamides

B4: cell division (unknown site)
20 phenylureas

B5: delocalisation of spectrin-like proteins
43 benzamides

B6: actin/myosin/fimbrin function
47 cyanoacrylates
50 aryl-phenyl-ketones

C: Respiration

C1: complex I NADH Oxido-reductase
39 pyrimidinamines, pyrazole-MET1, quinazoline

C2: complex II: succinate-dehydrogenase
7 SDHI (Succinate Dehydrogenase Inhibitors)

C3: complex III cytochrome bc1 (ubiquinol oxidase) at Qo site (cyt b gene)
11 QoI fungicides (Quinone outside Inhibitors)

C4: complex III cytochrome bc1 (ubiquinone reductase) at Qi site
21 QiI fungicides (Quinone inside Inhibitors)

C5: uncouplers of oxidative Phosphorylation
29

C6: inhibitors of oxidative phosphorylation, ATP synthase
30 organo tins

C7: ATP transport (proposed)
38 thiophene-carboxamides

C8: inhibition of complex III cytochrome bc1 (ubiquinone reductase) at Qo site stigmatellin binding sub site
45 QoSI-fungicide (stigmatellin binding)

C9: inhibition of complex III cytochrome bc1 (ubiquinone reductase) at Qo site
11 QoI fungicides (Quinone outside Inhibitors); Subgroup A

C10: inhibition of complex III cytochrome bc1 (ubiquinone reductase) at Qo site
11 QoI fungicides (Quinone outside Inhibitors); Subgroup A

D: Amino Acid and Protein Synthesis

D1: methionine biosynthesis (cgs gene) (proposed)
9 Anilino-Pyrimidines (AP fungicides)

D2: protein synthesis (ribosome, termination step)
23 enopyranuronic acid

D3: protein synthesis (ribosome, initiation step)
24 hexopyranuronic antibiotics

D4: protein synthesis (ribosome, initiation step)
25 glucopyranosyl antibiotics

D5: protein synthesis (ribosome, elongation step)
41 tetracycline antibiotics

E: Signal Transduction

E1: signal transduction (mechanism unknown)
13 azanaphthalenes

E2: osmotic signal transduction Δ MAP / histidine-kinase (os-2, HOG1)
12 phenylglyrols (PP-fungicides)

E3: osmotic signal transduction Δ MAP / histidine kinase (os-1, Daff1)
2 dicarboximides

F: Lipid Synthesis or Transport / Membrane Integrity or Function

F1: phospholipid biosynthesis Δ methyltransferase
6 phosphorothiolates & dithiolanes

F2: cell peroxidation (prop.)
14 aromatic hydrocarbons & heteroaromatics

F3: cell membrane permeability, fatty acids (prop.)
28 carbamates

F4: cell membrane permeability, fatty acids (prop.)
28 carbamates

F5: lipid homeostasis and transfer/storage
49 OSBPI Oxysterol binding protein homologue inhibition

F6: ergosterol binding
48 polylene

I: Melanin Synthesis in Cell Wall

I1: reductase in melanin biosynthesis
16.1 Melanin Biosynthesis Inhibitors: Reductase (MBI-R)

I2: dehydratase in melanin biosynthesis
16.2 Melanin Biosynthesis Inhibitors: Dehydratase (MBI-D)

I3: polyketide synthase in melanin biosynthesis
16.3 Melanin Biosynthesis Inhibitors: Polyketide synthase (MBI-P)

G: Sterol Biosynthesis in Membranes

G1: C14-demethylase in sterol biosynthesis (erg11/cyp51)
3DMI-fungicides (Demethylation_inhibitors) (SBI : Class I)

G2: Δ^14 -reductase and $\Delta^8 \rightarrow \Delta^7$ -isomerase in sterol biosynthesis (erg2, erg 24)
5 Amines ("Morpholines") (SBI : Class II)

G3: 3-keto reductase in C4-de-methylation (erg27)
17 (KRI fungicides KetoReductase Inhibitors) (SBI : Class III)

G4: squalene epoxidase in sterol biosynthesis (erg1)
18 (SBI : Class IV)

H: Cell Wall Biosynthesis

H4: chitin synthase
19 Polyoxins

H5: cellulose synthase
40 Carboxylic Acid Amides (CAA fungicides)

P: Host Plant Defence Induction

P1: salicylate related
#P01 benzothiazole BTH

P2: salicylate related
#P02 benzothiazole

P3: salicylate related
#P03 thiazole carboxamide

P4: polysaccharide elicitors
#P04 polysaccharide

P5: anthraquinone elicitors
#P05 plant extract

P6: microbial elicitors
#P06

P7: phosphonates
#P07 phosphonates

P8: salicylate related
#P08 isothiazole

Unknown Mode of Action

BM: Biologicals with Multiple Modes of Action

BM 01: plant extract

BM 02: microbial (strains of living microbes or extract, metabolites)

M: Chemicals with Multi-Site Activity

C: Respiration

C2: inhibition of complex II: Δ succinate-dehydrogenase
7 SDHI (Succinate dehydrogenase inhibitors)