

Mode of Action of Fungicides

FRAC classification on mode of action 2017 (www.frac.info)

A: Nucleic Acid Synthesis

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

A2: adenosin-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
1: MBC fungicides (= Methyl Benzimidazole Carbamates)

B2: β -tubulin assembly in mitosis*
10 N-phenyl carbamates

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

B4: cell division (prop.)
20 phenylureas

B5: delocalisation of spectrin-like proteins
43 benzamides

B6: actin/myosin/fimbrin function e.g. in vesicle trafficking
47 cyanocrylates

C: Respiration

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

C: Respiration

C1: inhibition of complex I NADH Oxido-reductase
39 pyrimidinamines pyrazole-MET1

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

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

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

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

C7: ATP production (prop.)
38 thiophene-carboxamides

C5: uncouplers of oxidative phosphorylation
29

D: Amino Acid and Protein Synthesis

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

D2: protein synthesis
24 enopyranonic acid

D3: protein synthesis
24 hexopyranosyl antibiotics

D4: protein synthesis
25 glucopyranosyl antibiotics

D5: protein synthesis
41 tetracycline antibiotics

E: Signal Transduction

E1: signal transduction (mechanism unknown)
13 azanaphthalenes

E3: osmotic signal transduction β MAP / histidine kinase (os-1, Daf1)
2 dicarboximides

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

F: Lipid Synthesis or Transport / Membrane Integrity or Function

F2: phospholipid biosynthesis β methyltransferase
6 phosphorothiolates and dithiolanes

F3: lipid peroxidation (prop.)
14 aromatic hydrocarbons and heteroaromatics

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

F6: microbial disrupters of pathogen cell membranes
44 Microbial (Bacillus sp.)

F7: cell membrane disruption (prop.)
46 plant extract

F8: ergosterol binding
48 polyene

F9: lipid homeostasis and transfer/storage
Oxysterol binding protein homologue inhibition

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)
3 DMF-fungicides (Demethylation Inhibitors) (SBI: Class I)

G2: Δ^{14} -reductase and $\Delta^8 \rightarrow \Delta^7$ -isomerase in sterol biosynthesis (erg2, erg24)
5 Amines ("Morpholines") (SBI: Class II)

G3: 3-keto reductase in C4-de-methylation (erg27)
17 (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)

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

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

P: Host Plant Defence Induction

P01: salicylic pathway
benzothiazole BTH

P02: benzothiazole
probenazole

P03: thiazole carboxamide
isotianil

P04: polysaccharide
laminarin

P05: plant extract
Reynoutria sachalinensis (Giant Knotweed Extract)

P06: Bacillus cereus group
Bacillus mycodens (Isolate J)

M: Chemicals with Multi-Site Activity

M01/M2 inorganics
Sulphur, Copper preparations

M03 dithiocarbamates & relatives
tetram, mancozeb, metiram, propineb, thiram, ziram, zineb

M04 phthalimides
anilazine, captan, captafol, folpet, guazatine

M05 chloronitriles
chlorothalonil

M06 sulphamides
tolpocarb

M07 guanidines
fenoxanil

M08 triazines
anilazine

M09 anthraquinones
dithianon

M10 quinoxalines
tricyclazole

M11 maleimides
fenoxanil

BM: Biologicals with Multiple Modes of Action

BM01: polypeptide lectin
Extract from the cotyledons of lupine plantlets (BLAD)

BM02: Trichoderma spp. metabolites
Trichoderma atroviride (strain SC1)

Unknown Mode of Action

27 cyanoamide-oxime
cyanoamide-oxime

33 ethyl-phosphonates
ethyl-phosphonates

33 phosphorous acid
phosphorous acid

34 tediolthalamid
tediolthalamid

36 benzene-sulfonamides
benzene-sulfonamides

37 diclozoxazine
diclozoxazine

42 methasulfocarb
methasulfocarb

44 cyflufenamid
cyflufenamid

46 cyflufenamid
cyflufenamid

47 cyflufenamid
cyflufenamid

48 metarfenone
metarfenone

49 dafconil
dafconil

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NC: Not Classified

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Mineral oils, organic oils, potassium bicarbonate, material of biological origin

Legend:

- mode of action group
- sub-group
- target site of action (where known) or putative target site (-prop.)
- FRAC code no. (#) and group name
- chemical (sub-) group

C: Respiration

C2: inhibition of complex II: succinate-dehydrogenase inhibitors

thiazole carboxamides