

A: Nucleic Acids Metabolism

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

FRAC Classification of Fungicides

Fungal control agents by cross resistance pattern and mode of action 2019 (www.frac.info)

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

B4: cell division (unknown site)
20 phenylureas

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

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 Qo1 fungicides (Quinone outside Inhibitors)

C4: complex III cytochrome bc1 (ubiquinone reductase) at Qi site
21 Qi1 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)

D: Amino Acid and Protein Synthesis

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

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

D3: protein synthesis (ribosome, initiation step)
24 hexopyranosyl 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 \triangleright MAP / histidine-kinase (os-2, HOG1)
12 phenylpyrroles (PP-fungicides)

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

F: Lipid Synthesis or Transport / Membrane Integrity or Function

F2: phospholipid biosynthesis \triangleright methyltransferase
6 phosphorothiolates & dithiolanes

F3: cell peroxidation (prop.)
14 aromatic hydrocarbons & 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
46 plant extract

F8: ergosterol binding
48 polyene

F9: lipid homeostasis and transfer/storage
49 OSBP

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 DMI-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

NC : Not Classified

M: Chemicals with Multi-Site Activity

M01 inorganic (electrophiles)

M02 inorganic (electrophiles)

M03 dithiocarbamates & relatives (electrophiles)

M04 phthalimides (electrophiles)

M05 chloronitriles (unspecified mechanism)

M06 sulphamides (electrophiles)

M07 bi-guanidines (membrane disruptors, detergents)

M08 triazines (unspecified mechanism)

M09 anthraquinones (electrophiles)

M10 quinoxalines (electrophiles)

M11 maleimides (electrophiles)

M12 thiocarbamates (electrophiles)

Unknown Mode of Action

*Temporary status; information on mode of action and / or resistance risk is still uncertain

#27 cyanacetamide-oxime

#34 phthalamic acid

#35 benzotriazines

#36 benzene-sulfonamides

#37 pyridazines

#U06 phenyl acetamides

#U13 cyano-methylene thiazolidine

#U14 pyrimidinone-hydrazones

#U16 4-quinolyl acetate

#U17 tetrazoloxime

#U18 glucopyranosyl antibiotic

#U12 guanidines

BM : Biologicals with Multiple Modes of Action

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

BM 01: plant extract

Legend:

Mode of action group

sub-group

\triangleright target site of action (where known) or putative target site (prop.)

FRAC code no. (#) and group name

chemical (sub-) group