Electronic Supplementary Material (ESI) for RSC Medicinal Chemistry.

Antibacterial activities of plant-derived xanthones

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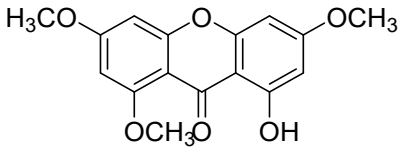
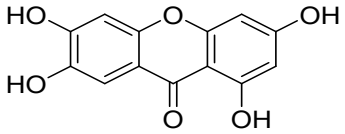
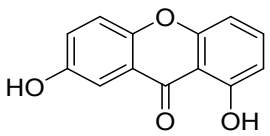
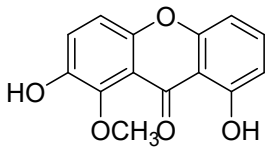
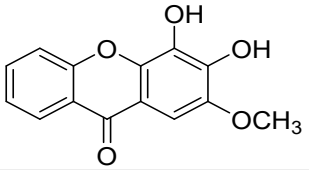
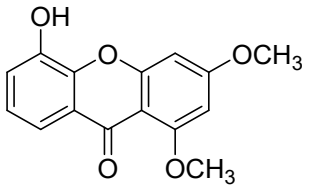
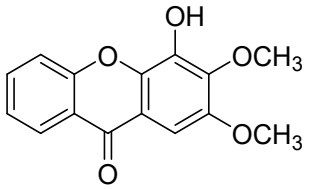
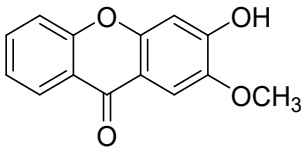
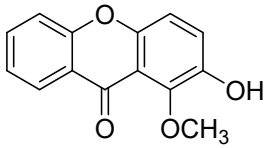
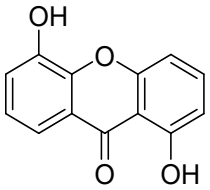
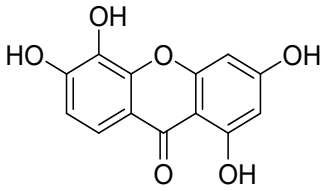
1. The detail information of all xanthones.

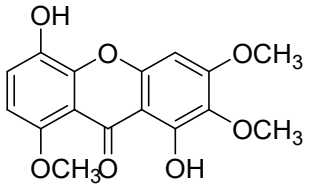
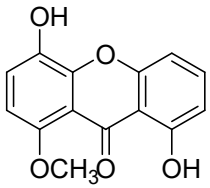
Table S1. The names, sources, structures, MIC values and bacteria species of the all xanthones.

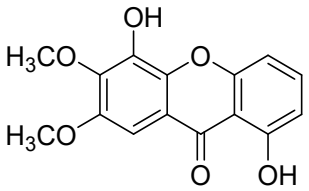
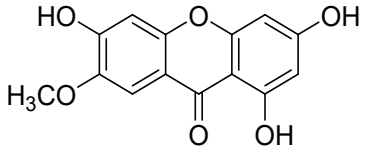
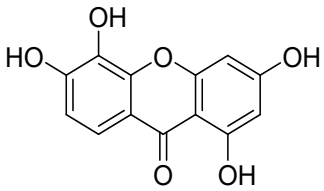
No.	Names	Sources	Structures	Strains	MICs	Ref.
Oxygen containing						
1	1-Hydroxy-3,6,8-trimethoxyxanthone	*Halenia elliptica*		*Staphylococcus aureus*	>100	1
				Bacillus subtilis	>100	
				Escherichia coli	256\>512	
				Enterobacter aerogenes	256\>512	
				Klebsiella pneumoniae	>512	
				Providencia stuartii	256\>512	
				Pseudomonas aeruginosa	>512	
2	Norathyriol	*Cratoxylum cochinchinense*, *Garcinia mangostana*		MRSA	18.75	8
				Bacillus subtilis	37.5	
				Enterococcus faecalis	300	
				VRE	>300	
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Pseudomonas aeruginosa	37.5	

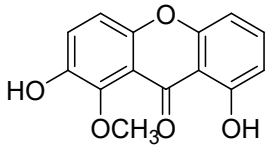
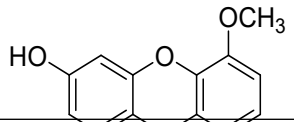
3	Euxanthone	*Garcinia dulcis*		MRSA	>128	18
				Staphylococcus ayreus	≥128	
				Escherichia coli	128	
				Staphylococcus typhimurium	\	
				Micrococcus luteus	32	
				Bacillus cereus	64	
				Bacillus subtilis	128	
				Staphylococcus aureus	>128	
				Staphylococcus epidermidis	32	
				Escherichia coli	16	
				Staphylococcus typhimurium	32	
				Pseudomonas aeruginosa	16	
4	1,7-Dihydroxy-8-methoxyxanthone	*Cratoxylum formosum*		*Bacillus subtilis*	37.5	29
				Staphylococcus. aureus	37.5	
				Enterococcus faecalis	300	
				MRSA	75	
				VRE	300	
				Salmonella typhi	>300	

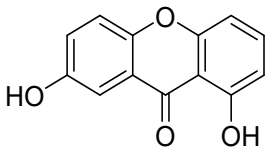
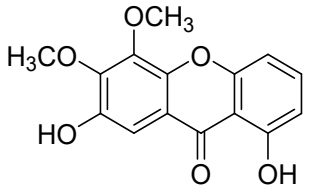
				Shigella sonei	>300	
				Pseudomonas aeruginosa	300	
5	3,4-Dihydroxy-2-methoxy-xanthone	*Kielmeyera variabilis*		MRSA	16-32\64	40
6	5-Hydroxy-1,3-dimethoxy-xanthone	*Kielmeyera variabilis*		MRSA	64-128\>512	40
7	4-hydroxy-2,3-dimethoxy-xanthone	*Kielmeyera variabilis*		MRSA	64-128\>512	40

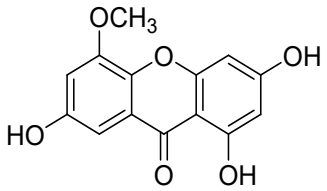
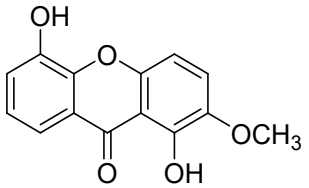
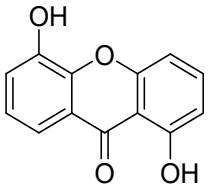
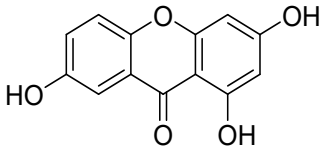
8	3-hydroxy-2-methoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\32	40
9	2-hydroxy-1-methoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\32\128	40
10	1,5-Dihydroxy-xanthone	the roots of *Garcinia propinqua*		*Bacillus cereus*	8	43
				Bacillus subtilis	32	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
11	1,3,5,6-Tetrahydroxy-xanthone	*Garcinia fusca*		*Helicobacter pylori*	2.02\16.25	46
				Micrococcus luteus	128	
				Bacillus cereus	64	
				Bacillus subtilis	64	
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	128	
				Escherichia coli	64	
				Salmonella	32	

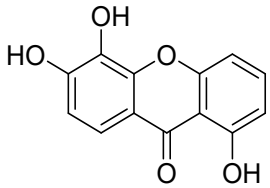
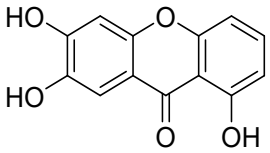
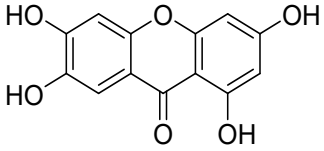
				typhimurium		
				Pseudomonas aeruginosa	128	
12	Cratosumatranone F	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	128	51
				Bacillus cereus	64	
				Bacillus subtilis	128	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	128	
				Escherichia coli	64	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	128	
13	1,5-Dihydroxy-8-methoxy-xanthone	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	32	51
				Bacillus cereus	64	
				Bacillus subtilis	128	
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	32	
				Escherichia coli	32	
				Salmonella typhimurium	32	
				Pseudomonas	16	

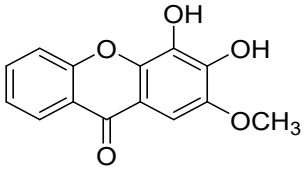
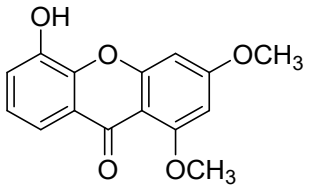
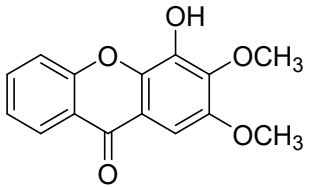
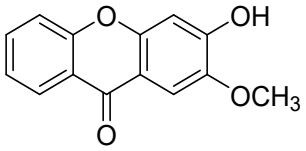
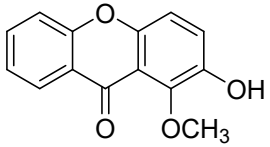
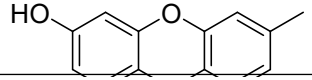
				aeruginosa		
14	1,5-Dihydroxy-6,7-dimethoxy-xanthone	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	32	51
				Bacillus cereus	16	
				Bacillus subtilis	64	
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	16	
				Escherichia coli	64	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	16	
15	1,3,6-Trihydroxy-7-methoxy-xanthone	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	128	51
				Bacillus cereus	128	
				Bacillus subtilis	128	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	128	
				Escherichia coli	64	
				Salmonella typhimurium	4	
				Pseudomonas aeruginosa	128	
16	1,3,5,6-Tetrahydroxy-	the twigs of *Cratoxylum*		*Micrococcus luteus*	128	51

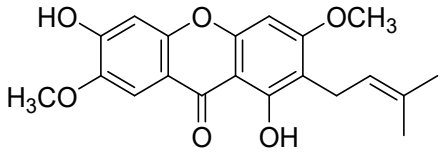
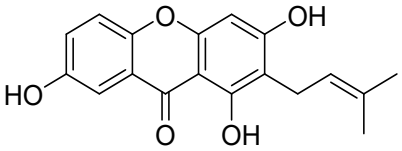
	xanthone	*sumatranum*		*Bacillus cereus*	64	
				Bacillus subtilis	64	
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	128	
				Escherichia coli	64	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	128	
17	2,8-Dihydroxy-1-methoxyxanthoe	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	64	51
				Bacillus cereus	64	
				Bacillus subtilis	128	
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	64	
				Escherichia coli	16	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	64	
18	Cratoxyarborenone F	the twigs of *Cratoxylum sumatranum*		*Micrococcus luteus*	64	51
				Bacillus cereus	64	
				Bacillus subtilis	128	

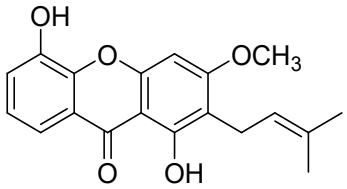
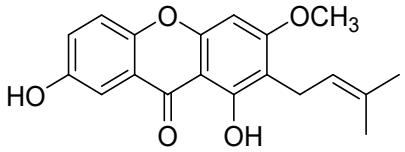
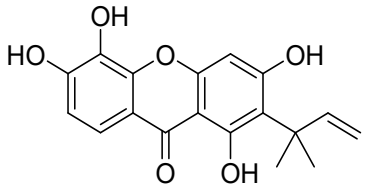
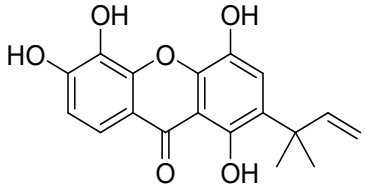
				Staphylococcus aureu	128	
				Staphylococcus epidermidis	64	
				Escherichia coli	32	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	
19	1,7-Dihydroxyxanthone	*Harungana madagascariensis*, *Garcinia succifolia*		MRSA	>25\6.25\>256	8
				Staphylococcus aureus	>25\6.25\>256	
				Bacillus subtillis	>256	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				VRE	>256	
20	1, 7-dimethoxy-5,6-trihydroxyxanthone	*Hypericum sampsonii*		MRSA	>128	63

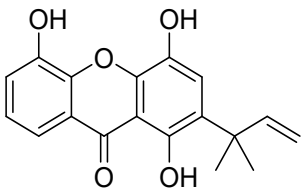
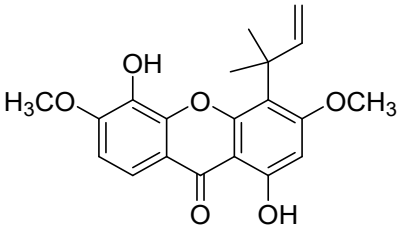
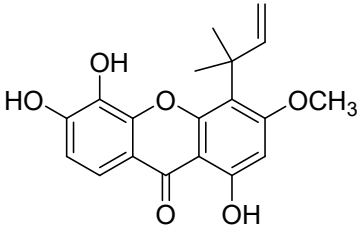
21	5-Methoxy-1,3,7-trihydroxyxanthone	*Hypericum sampsonii*		MRSA	>128	63
22	2-Methoxy-1,5-dihydroxyxanthone	*Hypericum sampsonii*		MRSA	>128	63
23	1,5-Dihydroxyxanthone	*Garcinia succifolia*		*Staphylococcus aureus*	>256	66
				Bacillus subtillis	>256	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				MRSA	>256	
				VRE	>256	
24	1,3,7-Trihydroxyxanthone	*Garcinia succifolia*		*Staphylococcus aureus*	>256	66
				Bacillus subtillis	>256	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				MRSA	>256	

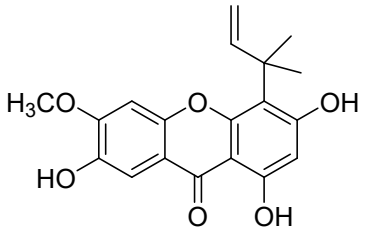
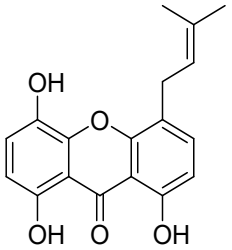
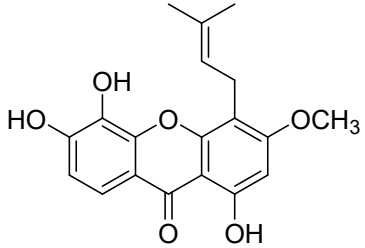
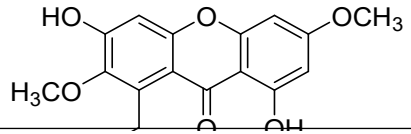
				VRE	>256	
25	1,5,6-Trihydroxyxanthone	*Garcinia succifolia*		*Staphylococcus aureus*	64	66
				Bacillus subtillis	64	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				MRSA	64	
				VRE	>256	
26	1,6,7-Trihtdroxyxanthone	*Garcinia succifolia*		*Staphylococcus aureus*	64	66
				Bacillus subtillis	128	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				MRSA	64	
				VRE	>256	
27	1,3,6,7-Tetrahydroxyxanthone	*Garcinia succifolia*		*Staphylococcus aureus*	256	66
				Bacillus subtillis	256	
				Pseudomonas aeruginosa	>256	
				Escherichia coli	>256	
				MRSA	>256	

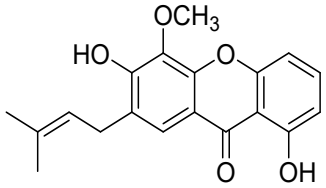
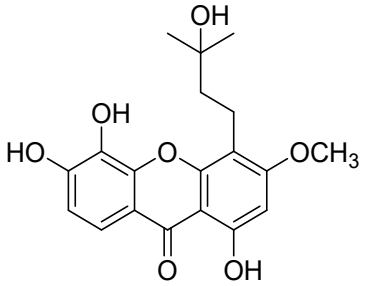
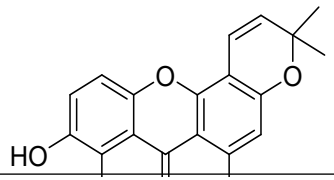
				VRE	>256	
28	3,4-dihydroxy-2-2methoxy-xanthone	*Kielmeyera variabilis*		MRSA	16\32\64	40
29	5-hydroxy-1,3-dimethoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\128\>512	40
30	4-hydroxy-2,3-dimethoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\128\>512	40
31	3-hydroxy-2-methoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\32	40
32	2-hydroxy-1-methoxy-xanthone	*Kielmeyera variabilis*		MRSA	64\128\32	40
33	Questin	*Cassia obtusefolia*		*Micrococcus luteus*	25	18
				Staphylococcus	25	

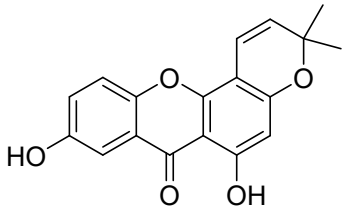
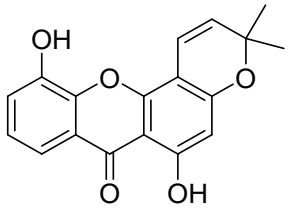
				aureus		
				Bacillus megaterium	12.5	
				Bacillus anthracis	12.5	
				Bacillus paratyphosum	25	
				MRSA	>100	
				Pseudomonas aeruginosa	50	
				Escherichia coli	50	
				Enterobacter aerogenes	50	
Monoprenyl group						
34	Cowagarcinone B	*Garcinia cowa*		MRSA	>128	1
				Staphylococcus aureus	>128	
				Escherichia coli	128	
				Salmonella typhimurium	128	
35	1,3,7-trihydroxy-2-prenylxanthone	*Cudrania cochinchinensis*, the leaves of *Garcinia goudotiana*		*Bacillus subtilis*	6.25	15,44, 60
				Staphylococcus aureus	6.25	
				Enterococci faecalis	6.25\39	
				Mycobacterium smegmatis	39	

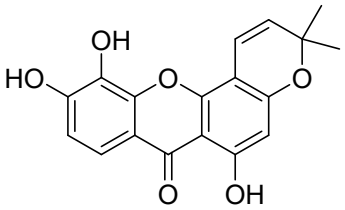
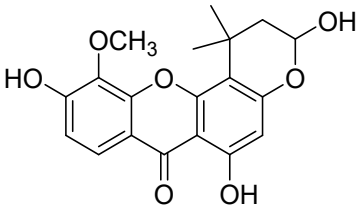
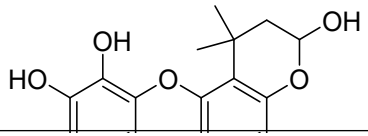
				Staphylococcus lugdunensis	78	
				MRSA	6.25	
				Micrococcus lutes	6.25	
36	1,5-dihydroxy-2-isoprenyl-3-methoxyxanthone	*Garcinia mangostana*		MRSA	25	54
				Staphylococcus aureus	12.5	
				Escherichia coli	25	
37	1,7-dihydroxy-2-isoprenyl-3-methoxyxanthone	*Garcinia mangostana*		MRSA	25	54
				Staphylococcus aureus	12.5	
				Escherichia coli	25	
38	Cudraxanthone S	*Cudrania cochinchinensis*		*Bacillus subtilis*	3.13	44,60
				Staphylococcus aureus	3.13	
				Micrococcus lutes	12.5	
				Enterococci faecalis	12.5\6.25\25	
				MRSA	3.13-6.25	
39	Subelliptenone F	*Garcinia subelliptica*		MRSA	25	54
				Staphylococcus aureus	25\12.5	
				Escherichia coli	25	

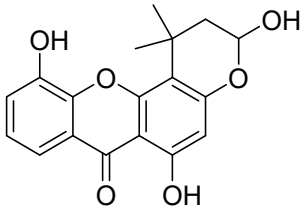
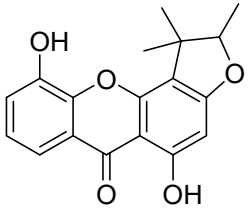
40	12-b-dihydroxy-des-D-garcigerin	*Garcinia subelliptica*		MRSA	12.5\3.13\6.25	54
				Staphylococcus aureus	25\6.25	
41	Vieillardiixanthone B	the root of *Cratoxylum formosum*		*Bacillus subtilis*	300	29
				Staphylococcus. aureus	300	
				Enterococcus faecalis	300	
				MRSA	300	
				VRE	300	
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Pseudomonas aeruginosa	>300	
42	Isocudraniaxanthone B	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	

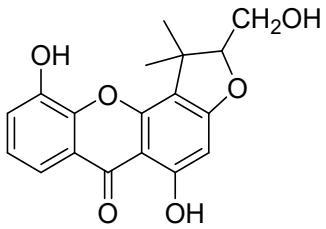
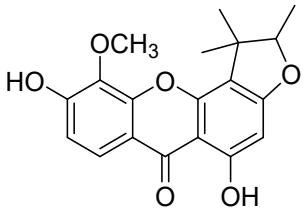
				Pseudomonas aeruginosa	>128	
43	Hypericumxanthone A	*Hypericum sampsonii*		MRSA	32	63
44	Caroxanthone	*Garcinia nobilis*		*Escherichia coli*	256\128\>512	21
				Enterobacter aerogenes	>512	
				Klebsiella pneumoniae	>512	
				Providencia stuartii	>512	
				Pseudomonas aeruginosa	>512	
45	Dulxanthone A	*Garcinia mangostana*		MRSA	3.9	38
				Vibrio vulnificus	15.6	
				Vibrio rotiferianus	15.6	
				Vibrio campbellii	31.2	
46	Garcicowanone A	*Garcinia cowa*		*Bacillus cereus*	0.25	2
				Bacillus subtilis	2	
				Micrococcus luteus	4	

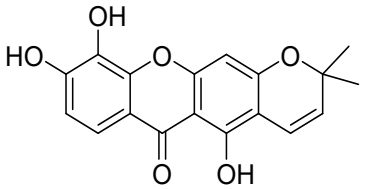
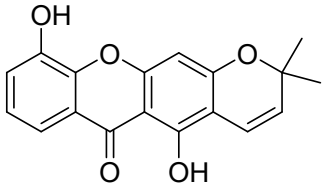
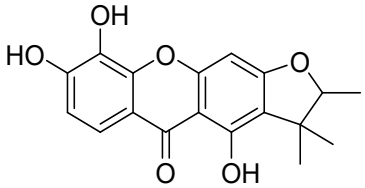
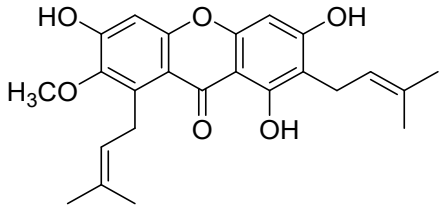
				Staphylococcus aureus	64	
				Escherichia coli	64	
				Pseudomonas aeruginosa	128	
				Salmonella typhimurium	64	
				Staphylococcus epidermidis	4	
47	Globulixanthone D	the root bark of *Symphonia globulifera*		*Staphylococcus aureus*	0.80	11
				Bacillus subtilis	12.5	
				Vibrio anguillarium	>100	
				Escherichia coli	>100	
48	Nigrolineaxanthone T	the bark of *Garcinia mangostana*		MRSA	62.4	38
				Vibrio vulnificus	>125	
				Vibrio rotiferianus	>125	
				Vibrio campbellii	>125	
49	Globulixanthone C	the root bark of *Symphonia globulifera*		*Staphylococcus aureus*	14.05	11
				Bacillus subtilis	8.24	
				Vibrio anguillarium	\	

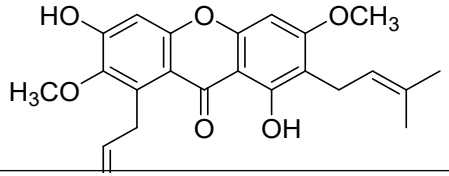
				Escherichia coli	>100	
50	Nigrolineaxanthone F	the stem bark *Garcinia nigrolinaeta*		MRSA	2	20
51	6-deoxyiso-jacareubin	*Vismia laurentii*		*Citrobacter freundii*	19.53	32
				Enterobacter aerogens	\	
				Enterobacter cloacae	\	
				Escherichia coli	\	
				Klebsiella pneumoniae	39.06	
				Morganella morganii	78.12	
				Proteus mirabilis	78.12	
				proteus vulgaris	39.06	
				Pseudomonas aeruginosa	\	
				Shigella dysenteriae	\	
				Shigella flexneri	\	
				Salmonella typhi	\	
				Streptococcus faecalis	\	
				Staphylococcus aureus	39.06	

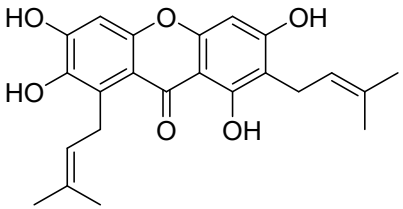
				Bacillus cereus	78.12	
				Bacillus megaterium	\	
				Bacillus stearothermophilus	1.22	
				Bacillus subtilis	4.88	
52	Isojacareubin	*Garcinia fusca*		*Helicobacter pylori*	2.54	46
53	Cratosumatranone D	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	8	51
				Bacillus cereus	8	
				Bacillus subtilis	16	
				Staphylococcus aureu	32	
				Staphylococcus epidermidis	8	
				Escherichia coli	64	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	16	
54	Cratosumatranone E	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	

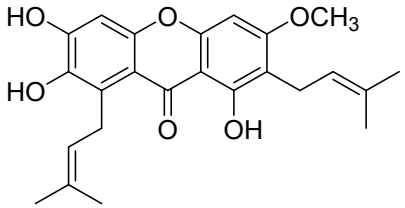
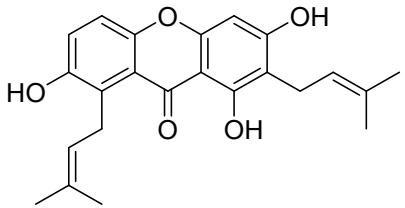
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	
55	Pruniflorone N	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	16	51
				Bacillus cereus	32	
				Bacillus subtilis	32	
				Staphylococcus aureu	8	
				Staphylococcus epidermidis	16	
				Escherichia coli	32	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	16	
56	Pancixanthone B	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	32	51
				Bacillus cereus	16	
				Bacillus subtilis	128	
				Staphylococcus aureu	>128	
				Staphylococcus	32	

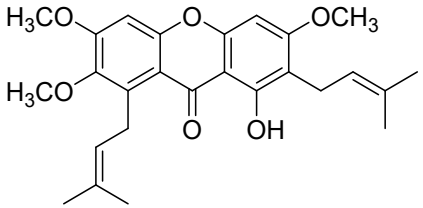
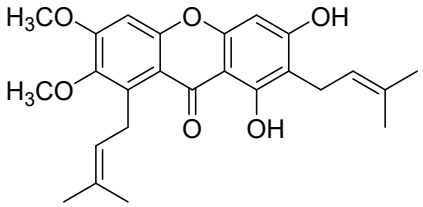
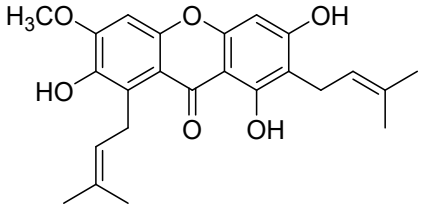
				epidermidis		
				Escherichia coli	32	
				Salmonella typhimurium	32	
				Pseudomonas aeruginosa	16	
57	Pruniflorone M	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	
58	5-*O*-methyl-2-deprenylrheedixanthone B	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	

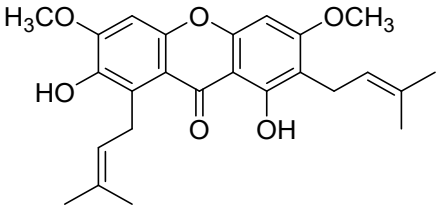
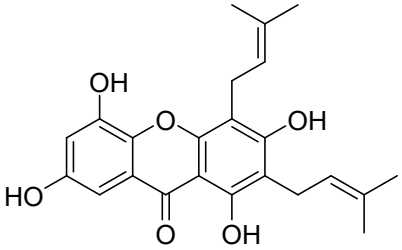
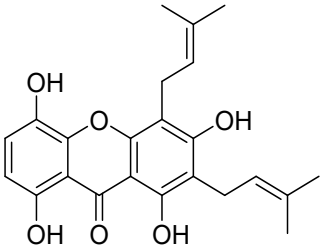
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	
59	Jacareubin	the twigs of *Garcinia staudtii*		MRSA	\	19
60	6-deoxyjacareubin	the stem bark *Garcinia nigrolinaeta*		MRSA	4	20
61	Toxyloxanthone C	*Cudrania cochinchinensis*		*Bacillus subtilis*	3.13	44,60
				MRSA	3.13-6.25	
				Enterococci faecalis	12.5-25	
				MRSA	3.13-6.25	
				Micrococcus luteus	12.5	
Biprenyl group						
62	α-mangostin	the fruits, inflorescences, flowers of *Garcinia cowa*; the twigs of *Garcinia staudtii*; *Tetragonula laeviceps* propolis; *Garcinia fusca*; *Garcinia mangostana*		*Micrococcus luteus*	1\6.25	1-3, 8,13, 19,27,37, 46-47, 54-
				Staphylococcus aureus	64\4\3.13\25\12.5\8\794\64\128	
				Staphylococcus epidermidis	0.5\1.56	

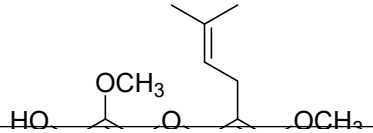
				Bacillus cereus	250\8\3.13\0.5	55
				MRSA	32\3.13\8\2\25\12.5\6.25\16\2.34	
				Listeria monocytogenes	0.78	
				Streptococcus pyogenes	0.78	
				Escherichia coli	12.5\64\250\25\128	
				Serratia marcescens	12.5	
				Helicobacter pylori	19\76.1	
				Bacillus subtilis	8\2.34\0.25	
				Salmonella typhimurium	64\12.5\128	
				Pseudomonas aeruginosa	2.34\128\12.5	
				VRE	3.13\150	
				Enterococcus faecalis	150	
				Salmonella typhi	18.75	
				Shigella sonei	150	
63	β-mangostine	the fruits, inflorescences and flowers of *Garcinia cowa*; *Garcinia mangostana*; *Garcinia fusca*		MRSA	8\6.25\4.68\>200\64	1-2, 8, 13, 27, 46-47, 54-55
				Staphylococcus aureus	12.5\6.25\25\64\>128\128	

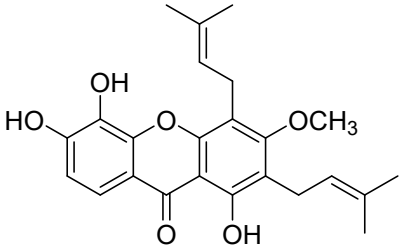
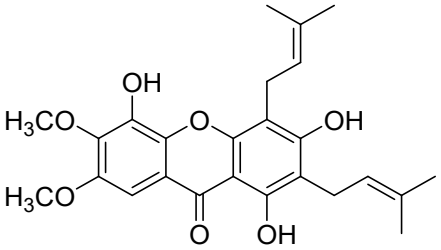
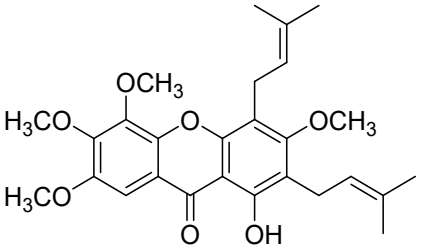
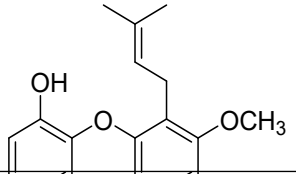
				Escherichia coli	>128\25	
				Bacillus cereus	0.25\16	
				Bacillus subtilis	4\2..34	
				Micrococcus luteus	16	
				Pseudomonas aeruginosa	128\9.37	
				Salmonella typhimurium	64	
				Staphylococcus epidermidis	2	
				Enterococcus faecalis	300	
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Helicobacter pylori	18.3\183\147.3	
				VRE	>200	
64	γ-mangostin	*Tetragonula laeviceps* propolis; *Garcinia mangostana*		MRSA	4.68\3.13\6.25\12.5	8,37, 54-55
				Bacillus subtilis	18.75	
				Enterococcus faecalis	9.37	
				Salmonella typhi	18.75	
				Shigella sonei	18.75	
				Pseudomonas aeruginosa	18.75\25	
				Bacillus cereus	25	

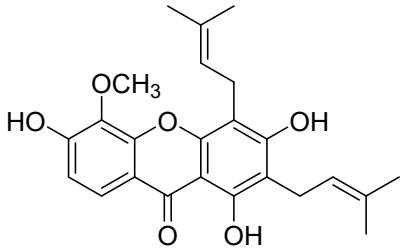
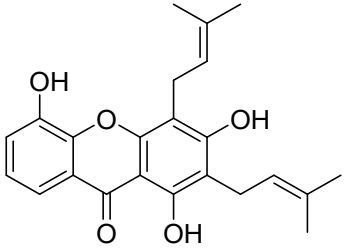
				Listeria monocytogenes	12.5	
				Micrococcus luteus	25	
				Staphylococcus epidermidis	25	
				Streptococcus pyogenes	6.25	
				Escherichia coli	25	
				Salmonella typhimurium	25	
				Serratia marcescens	25	
				Staphylococcus aureus	6.25\25	
				VRE	6.25	
65	Dulxisxanthone			MRSA	9.37	8
				Bacillus subtilis	18.75	
				Enterococcus faecalis	300	
				VRE	300	
				Salmonella typhi	150	
				Shigella sonei	150	
				Pseudomonas aeruginosa	18.75	
66	6-deoxy-γ-mangostin	Calophyllum; *Garcinia mangostana*		*Staphylococcus aureus*	250	16,55

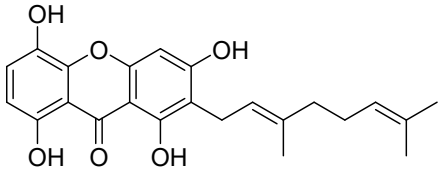
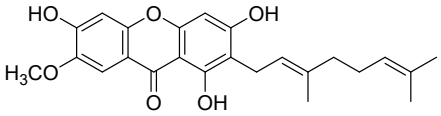
				MRSA	>250	
				VRE	>250	
67	Fuscaxanthone C	the fruits and flowers of *Garcinia cowa*; *Garcinia mangostana*		*Bacillus subtilis*	128	1,47, 55
				Pseudomonas aeruginosa	128	
				Bacillus cereus	>128	
				Staphylococcus aureus	128	
				Escherichia coli	64	
				Salmonella typhimurium	128	
				MRSA	>200	
				VRE	>200	
68	Cowaxanthone B	the fruits of *Garcinia cowa*		MRSA	18.75\128	13
				Bacillus subtilis	75	
				Enterococcus faecalis	75	
				VRE		
				Salmonella typhi	150	
				Shigella sonei	150	
				Pseudomonas aeruginosa	18.75	
				Staphylococcus aureus	128	
69	6-methoxy-γ-mangostin			MRSA	>200	55

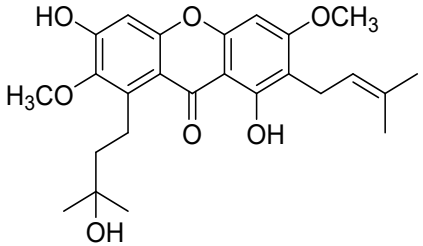
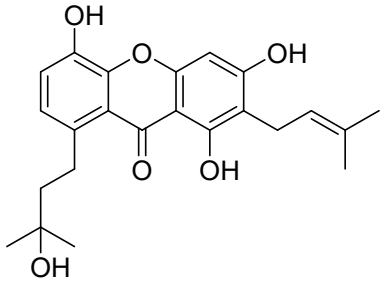
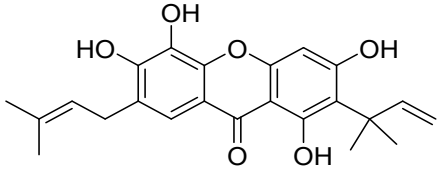
				Bacillus subtilis	75	
				Enterococcus faecalis	75	
				VRE	>200	
				Salmonella typhi	37.5	
				Shigella sonei	75	
				Pseudomonas aeruginosa	4.67	
70	3,6-di-*O*-methyl-γ- Mangostin	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	128	47
				Bacillus subtilis	128	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	128	
71	Staudtiixanthone A	the twigs of *Garcinia staudtii*		MRSA	16	19
72	Gartanin	the twigs of *Garcinia staudtii*; *Tetragonula laeviceps* propolis; the bark of *Garcinia mangostana*; the		MRSA	32\12.5\>125\>128\25	19, 37-38, 42, 54

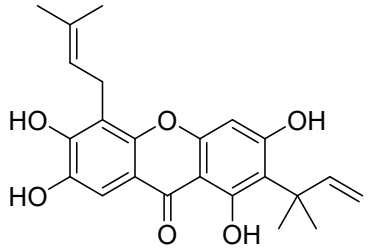
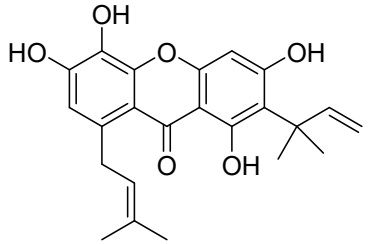
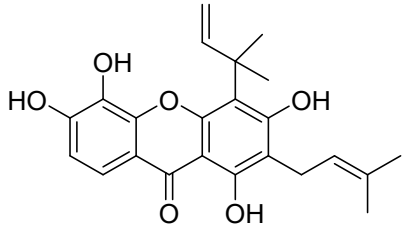
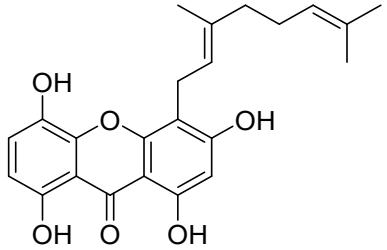
		twigs of *Garcinia propinqua*; *Garcinia mangostana*		*Bacillus cereus*	25	
				Listeria monocytogenes	12.5	
				Micrococcus luteus	25	
				Staphylococcus aureus	12.5\128\>125	
				Staphylococcus epidermidis	25	
				Streptococcus pyogenes	6.25	
				Escherichia coli	25\128	
				Pseudomonas aeruginosa	25	
				Salmonella typhimurium	25	
				Serratia marcescens	25	
				Vibrio vulnificus	62.4	
				Vibrio rotiferianus	125	
				Vibrio campbellii	125	
				Staphylococcus typhimurium	128	
73	Doitunggarcinone C	the twigs of *Garcinia propinqua*		MRSA	128	42
				Staphylococcus aureus	128	

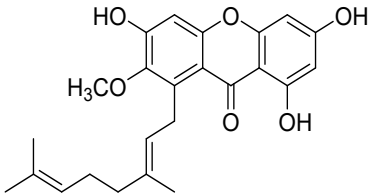
				Escherichia coli	128	
				Staphylococcus typhimurium	\	
74	Dulxanthone B	the twigs of *Garcinia propinqua*		MRSA	16	42
				Staphylococcus aureus	32	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
75	Globuliferin	the seeds of *Symphonia globulifera*		*Staphylococcus aureus*	0.58	25
				Streptococcus feacalis	8.25	
				Klebsiella pneumonia	15.24	
				Escherichia coli	21.2	
76	3,5-Di-*O*-methylglobuliferin	the seeds of *Symphonia globulifera*		*Staphylococcus aureus*	\	25
				Streptococcus feacalis	\	
				Klebsiella pneumonia	\	
				Escherichia coli	\	
77	8-hydroxy-cudraxanthone G			*Escherichia coli*	>512\64\128	21
				Enterobacter aerogenes	>512\64\128	
				Klebsiella	>512	

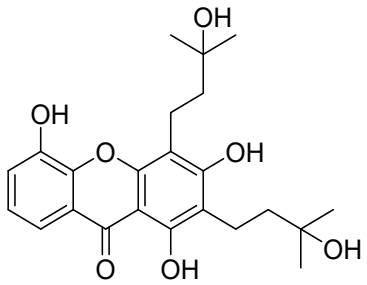
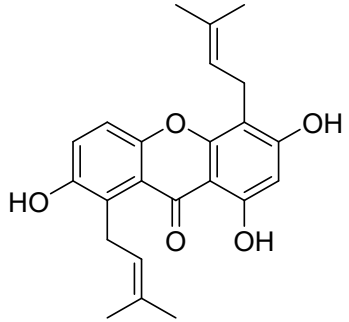
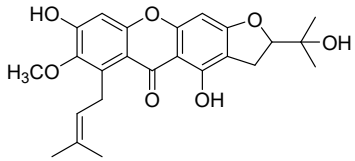
				pneumoniae		
				Providencia stuartii	>512	
				Pseudomonas aeruginosa	>512\256	
78	Doitunggarcinone D	the roots of *Garcinia propinqua*		*Bacillus cereus*	64	43
				Bacillus subtilis	16	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
79	8-Deoxygartanin	the leaves of *Garcinia nigrolineata*		*Staphylococcus aureus*	16\128	48
				Micrococcus luteus	>128	
				Streptococcus mutans	128	
				Staphylococcus epidermidis	128	
				Bacillus cereus	128\3.13	
				MRSA	125\16	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	>128	
				Escherichia coli	>128	
				Shigella flexneri	128	

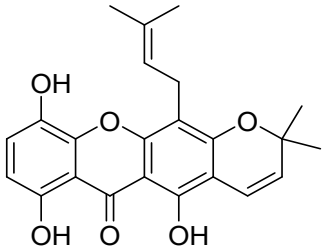
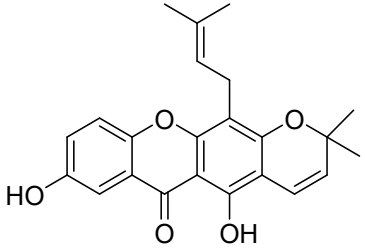
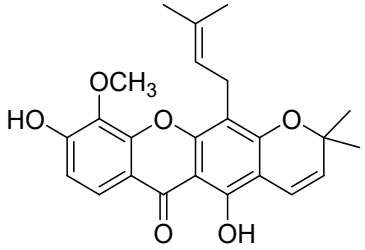
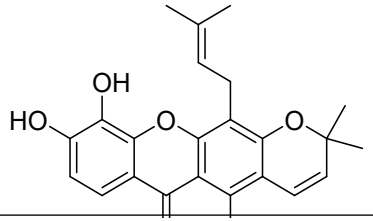
				Vibrio vulnificus	>125	
				Vibrio rotiferianus	>125	
				Vibrio campbellii	125	
80	Smeathxanthone A	*Garcinia smeathmannii*		*Escherichia coli*	128\64\32	21,53
				Enterobacter aerogenes	>512\256	
				Klebsiella pneumoniae	>512	
				Providencia stuartii	>512	
				Pseudomonas aeruginosa	>512\256	
				Staphylococcus aureus	32	
				Enterococcus faecalis	16	
				Bacillus cereus	64	
				Salmonella typhimurium	256	
81	Cowaxanthone	the fruits and flowers of *Garcinia cowa*; *Garcinia fusca*;		MRSA	16	1, 46-47
				Staphylococcus aureus	16\>128	
				Escherichia coli	128\>128	
				Salmonella typhimurium	128	
				Helicobacter pylori	0.77\3.2\25.64	

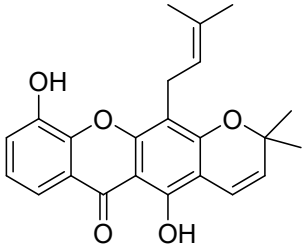
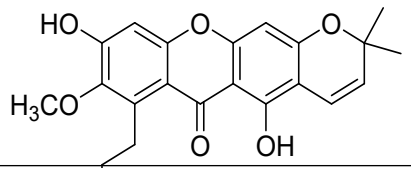
				Bacillus cereus	64	
				Bacillus subtilis	64	
				Pseudomonas aeruginosa	128	
82	Pruniflorone C	*Cratoxylum formosum*		MRSA	18.75	8
				Bacillus subtilis	9.37	
				Enterococcus faecalis	150	
				VRE	>300	
				Salmonella typhi	300	
				Shigella sonei	300	
				Pseudomonas aeruginosa	18.75	
83	Nigrolineaxanthone N	*Garcinia nigrolineata*		MRSA	128	20
84	Isoalvaxanthone	*Cudrania cochinchinensis*		*Enterococci faecalis*	6.25\12.5	44,60
				Bacillus subtilis	12.5	
				Staphylococcus aureus	6.25	
				MRSA	3.13-6.25	

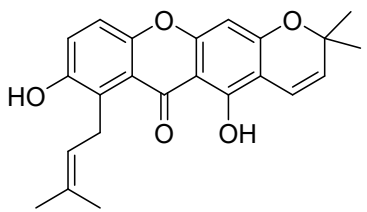
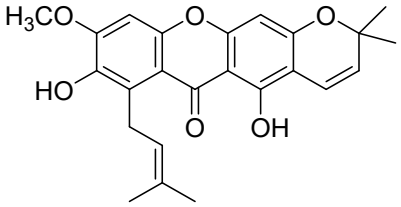
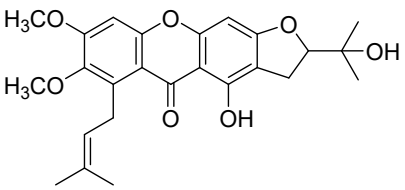
				Micrococcus lutes	6.25	
85	Cudraxanthone L	the roots of *Garcinia propinqua*		*Bacillus cereus*	8	43
				Bacillus subtilis	1	
				Escherichia coli	128	
				Staphylococcus typhimurium	64	
86	Alvaxanthone	*Cudrania cochinchinensis*		*Enterococci faecalis*	12.5\6.25	44,60
				Bacillus subtilis	12.5	
				Staphylococcus aureus	6.25	
				MRSA	6.25\12.5	
				Micrococcus lutes	12.5	
87	Gerontoxanthone Ⅰ	*Cratoxylum* species		*Enterococci faecalis*	6.25\3.13	32
				Bacillus subtilis	6.25\2.3	
				Staphylococcus aureus	3.13\1.1	
				MRSA	3.13\6.25	
				Micrococcus lutes	6.25	
				Pseudomonas aeruginosa	>50	
				Salmonella faecalis	4.6	
				Salmonella typhi	1.1	
88	Cheffouxanthone	*Garcinia smeathmannii;* the twigs of		MRSA	64	14,53

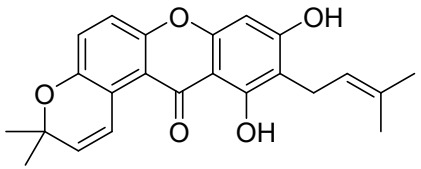
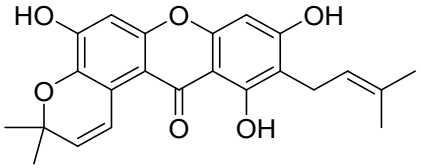
		Garcinia hombroniana		*Staphylococcus aureus*	16\64	
				Enterococcus faecalis	8	
				Bacillus cereus	32	
				Pseudomonas aeruginosa	128	
				Escherichia coli	64	
				Salmonella typhimurium	\	
89	Rubraxanthone	the fruits and flowers of *Garcinia cowa*; the stem bark of Allanblackia monticola; *Garcinia dioica*		*Bacillus cereus*	2\1000\0.313-1.25	2-3, 12,47,54
				Bacillus subtilis	1	
				Micrococcus luteus	2	
				Staphylococcus aureus	>128\1000	
				Escherichia coli	64\794	
				Salmonella typhimurium	64	
				Staphylococcus epidermidis	4	
				Staphylococcus ayreus	12	
				Vibrio anguillarium	\	
				Candida tropicalis	\	
				Pseudomonas aeruginosa	64	

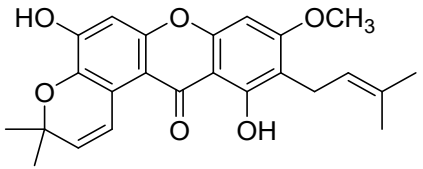
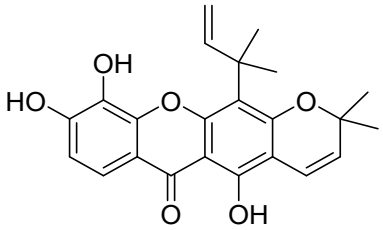
				MRSA	0.313\1.25	
90	Nigrolineaxanthone Z	*Garcinia nigrolinaeta*		*Micrococcus luteus*	>128	4
				Streptococcus mutans	128	
				Staphylococcus epidermidis	128	
				Bacillus cereus	128	
				Staphylococcus aureus	128	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	>128	
				Escherichia coli	128	
				Shigella flexneri	128	
91	Gerontoxanthone H (Cudraxanthone H)	*Cudrania cachinchinensis*		*Enterococci faecalis*	1.56	44,60
				Bacillus subtilis	1.56	
				Staphylococcus aureus	1.56	
				MRSA	1.56	
				Micrococcus lutes	1.56	
92	Dulcisxanthone G	the seeds of *Garcinia dulcis*		MRSA	>128	18
				Staphylococcus ayreus	>128	

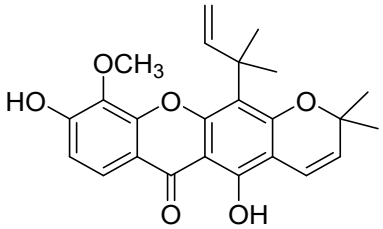
93	Morusignin			*Escherichia coli*	64\128	21
				Enterobacter aerogenes	>512	
				Klebsiella pneumoniae	>512	
				Providencia stuartii	>512	
				Pseudomonas aeruginosa	128\64	
94	Cudraxanthone Q	the roots of *Garcinia propinqua*		*Bacillus cereus*	128	43
				Bacillus subtilis	128	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
95	5-*O*-methylxanthone V_1	the twigs of *Garcinia propinqua*		MRSA	>128	42
				Staphylococcus aureus	>128	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
96	Xanthone V_1	the roots of *Cratoxylum formosum*		*Pseudomonas aeruginosa*	9.3	32
				Bacillus subtilis	1.1	
				Staphylococcus aureus	1.1	

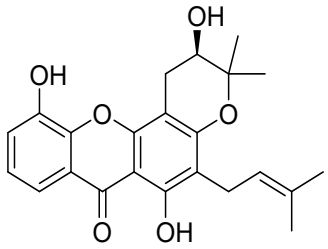
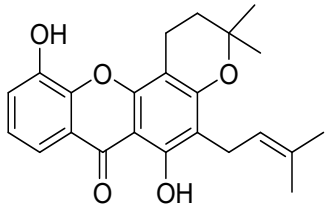
				Streptococcus faecalis	1.1	
				Salmonella typhi	1.1	
97	Toxyloxanthone A (Trapezifolixanthone)	*Garcinia tetralata*		*Micrococcus luteus*	128	4
				Streptococcus mutans	128	
				Staphylococcus epidermidis	128	
				Bacillus cereus	128	
				Staphylococcus aureus	128	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	128	
				Escherichia coli	>128	
				Shigella flexneri	128	
				Bacillus cereus	16	
				Bacillus subtilis	4	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
98	9-hydroxy-calabaxanthone	the bark of *Garcinia mangostana*; the inflorescences of *Garcinia cowa*		MRSA	>125\4	27, 38
				Vibrio vulnificus	>125	
				Vibrio rotiferianus	125	
				Vibrio campbellii	125	

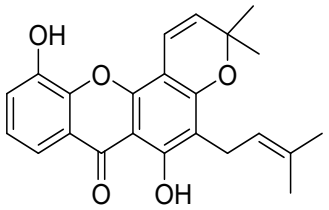
				Staphylococcus aureus	4	
99	Demethylcalabaxanthone	the twigs of *Garcinia staudtii*		MRSA	128	19
100	Garciniacowone E	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	128	47
				Bacillus subtilis	128	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	128	
101	6-*O*-methylmangostanin	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	128	47
				Bacillus subtilis	128	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	

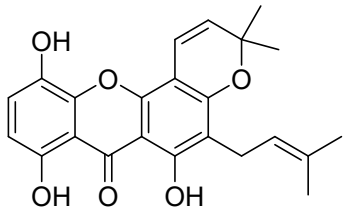
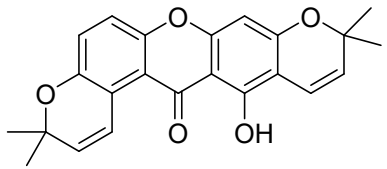
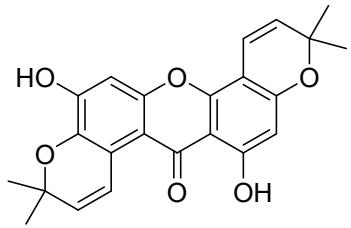
102	Calothwaitesixanthone	Calophyllum		*Staphylococcus aureus*	\	16
103	Garcinone B	the twigs of *Garcinia staudtii*; the root of *Cratoxylum formosum*; *Tetragonula laeviceps* propolis		MRSA	128\150\6.25	19, 29, 37
				Bacillus subtilis	75	
				Staphylococcus aureus	75\6.25	
				Enterococcus faecalis	75	
				VRE	75	
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Pseudomonas aeuruginosa	>300\25	
				Bacillus cereus	3.13	
				Listeria monocytogenes	6.25	
				Micrococcus luteus	3.13	
				Staphylococcus epidermidis	3.13	
				Streptococcus pyogenes	1.56	
				Escherichia coli	25	
				Salmonella	25	

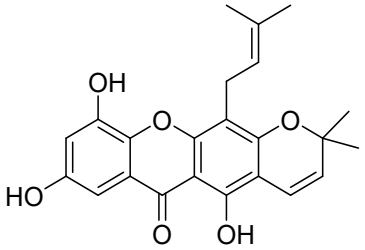
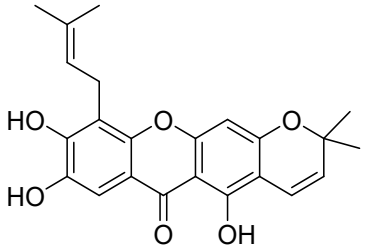
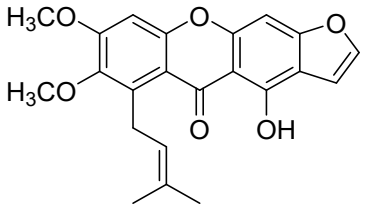
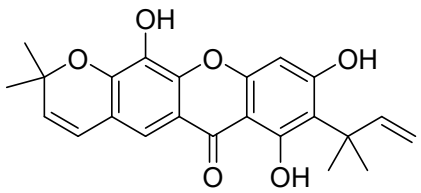
				typhimurium		
				Serratia marcescens	25	
104	Cowaxanthone D	the fruits and flowers of *Garcinia cowa*		MRSA	128	1-2, 47
				Staphylococcus aureus	>128	
				Bacillus cereus	64	
				Bacillus subtilis	128	
				Micrococcus luteus	64	
				Escherichia coli	>128	
				Pseudomonas aeruginosa	128	
				Salmonella typhimurium	64	
				Staphylococcus epidermidis	32	
105	Macluraxanthone	*Garcinia propinqua*		MRSA	4	29,32, 42, 43
				Staphylococcus aureus	4	
				Escherichia coli	128	
				Staphylococcus typhimurium	\	
				Bacillus cereus	4	
				Bacillus subtilis	4	
				Escherichia coli	128	

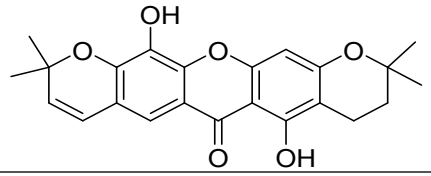
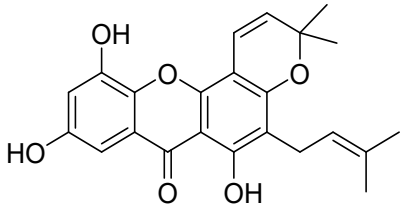
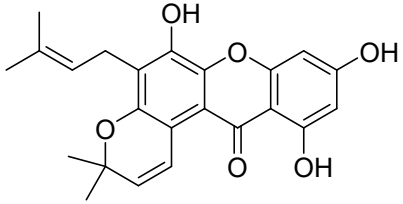
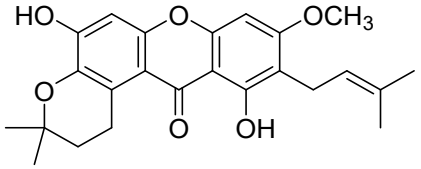
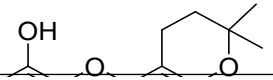
				Micrococcus luteus	>128	
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureus	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Staphylococcus typhimurium	>128	
				Pseudomonas aeruginosa	>128	
106	10-*O*-methylmaclura-xanthone	*Garcinia bracteata*		MRSA	4	42
				Staphylococcus aureus	2	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
				Micrococcus luteus	>128	
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureus	>128	
				Staphylococcus	>128	

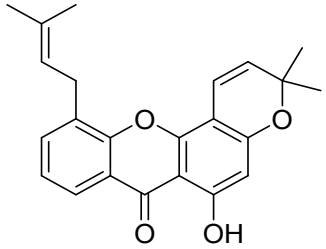
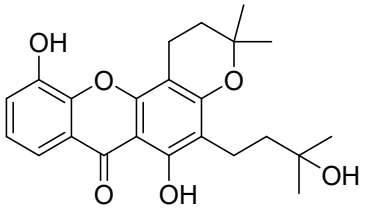
				epidermidis		
				Escherichia coli	>128	
				Staphylococcus typhimurium	>128	
				Pseudomonas aeruginosa	>128	
107	Nigrolineaxanthone X	*Garcinia nigrolineata*		*Micrococcus luteus*	>128	4
				Streptococcus mutans	>128	
				Staphylococcus epidermidis	128	
				Bacillus cereus	128	
				Staphylococcus aureus	128	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	>128	
				Escherichia coli	>128	
				Shigella flexneri	>128	
108	Nigrolineaxanthone Q	*Garcinia nigrolineata*		*Micrococcus luteus*	8	4
				Streptococcus mutans	32	
				Staphylococcus epidermidis	128	
				Bacillus cereus	32	

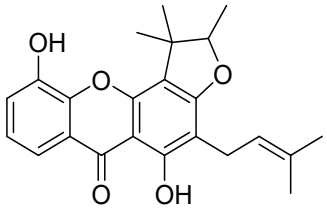
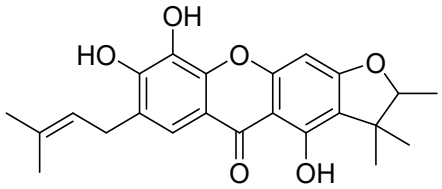
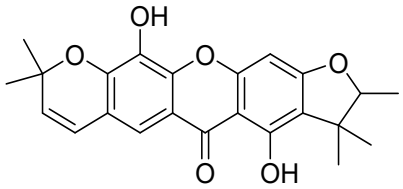
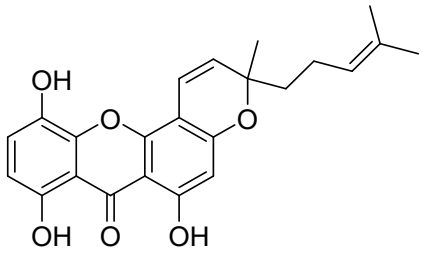
				Staphylococcus aureus	128	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	128	
				Escherichia coli	128	
				Shigella flexneri	128	
109	Ananixanthone	the leaves of *Garcinia nigrolineata*; *Garcinia smeathmannii*		*Micrococcus luteus*	32	4,48, 53
				Streptococcus mutans	32	
				Staphylococcus epidermidis	128	
				Bacillus cereus	32	
				Staphylococcus aureus	128\32	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	128	
				Escherichia coli	>128	
				Shigella flexneri	>128	
				MRSA	32	
				Enterococcus faecalis	2	
				Pseudomonas	64	

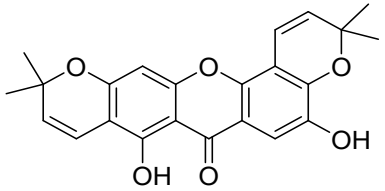
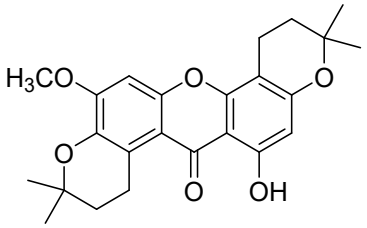
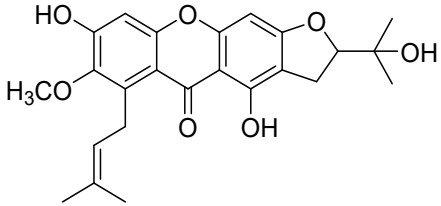
				aeruginosa		
110	Morusignin J	the seeds of *Garcinia dulcis*		*Staphylococcus aureus*	>128	18
				MRSA	>128	
				Escherichia coli	128	
				Staphylococcus typhimurium	\	
111	Thwaitesixanthone	*Calophyllum thwaitesii*		*Staphylococcus aureus*	\	16
112	Brasilixanthone	the stem bark of *Garcinia nigrolineata*; the root of *Cratoxylum formosum*		*Bacillus subtilis*	300	29
				Staphylococcus. aureus	300	
				Enterococcus faecalis	>300	
				MRSA	300	
				VRE	2	
				Salmonella typhi	300	
				Shigella sonei	>300	
				Pseudomonas aeruginosa	>300	
				Bacillus subtilis	>300	

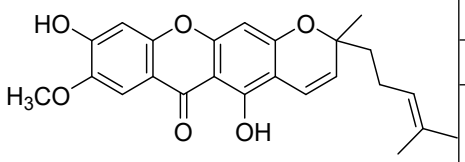
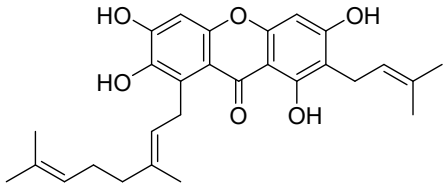
113	Staudtiixanthone B	the twigs of *Garcinia staudtii*		MRSA	32	19
114	Caloxanthone A	*Calophyllum inophyllum*		MRSA	>25	54
				Staphylococcus aureus	6.25\25	
115	Garciniafuran	the stem bark of *Allanblackia monticola*		*Staphylococcus aureus*	\	54
				Vibrio anguillarium	\	
				Candida tropicalis	\	
116	Gerontoxanthone B	*Cudrania cochinchinensis*		*Bacillus subtilis*	>25	60
				Staphylococcus aureus	>25	
				Staphylococcus aureus	12.5->25	
				Micrococcus lutes	>25	

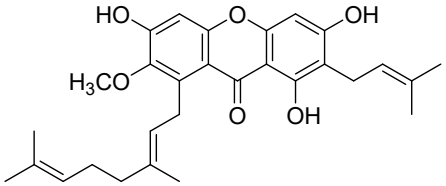
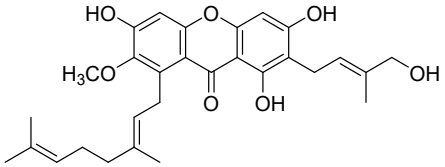
117	Nigrolineaxanthone G	the stem bark *Garcinia nigrolinaeta*		MRSA	4	20
118	Staudtiixanthone C	*Garcinia staudtii*		MRSA	128	19
119	Hypericumxanthone B	*Hypericum sampsonii*		MRSA	32	63
120	3-*O*-methylmangostenone	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	128	47
				Bacillus subtilis	\	
				Staphylococcus aureus	128	
				Escherichia coli	>128	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	>128	
121	Nigrolineaxanthone Q	*Garcinia nigrolineata*		*Micrococcus luteus*	8	4
				Streptococcus mutans	32	

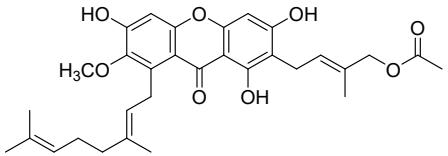
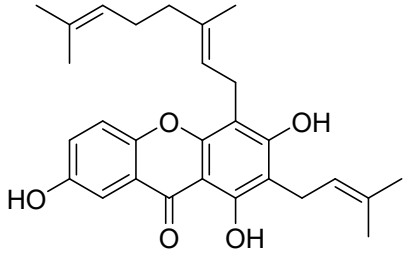
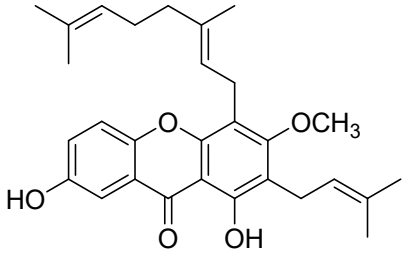
				Staphylococcus epidermidis	128	
				Bacillus cereus	32	
				Staphylococcus aureus	128	
				Salmonella typhimurium	128	
				Pseudomonas aeuruginosa	128	
				Escherichia coli	128	
				Shigella flexneri	128	
122	Laurentixanthone A	the roots of Vismia laurentii		*Candida glabrata*	1	17
123	Nigrolineaxanthone Y	*Garcinia nigrolinaeta*		*Micrococcus luteus*	32	4
				Streptococcus mutans	128	
				Staphylococcus epidermidis	128	
				Bacillus cereus	64	
				Staphylococcus aureus	128	
				Salmonella	>128	

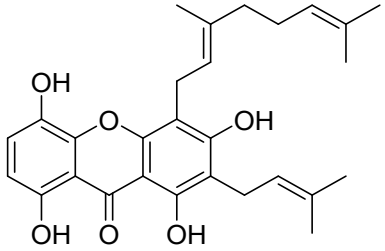
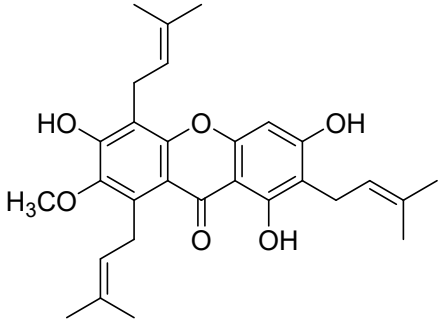
				typhimurium		
				Pseudomonas aeuruginosa	128	
				Escherichia coli	128	
				Shigella flexneri	128	
124	Caloxanthone L	the roots of *Garcinia propinqua*		*Bacillus cereus*	64	43
				Bacillus subtilis	128	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
125	Gerontoxanthone G	*Cudrania cochinchinensis*		*Bacillus subtilis*	12.5	60
				Staphylococcus aureus	12.5	
				Staphylococcus aureus	6.25-12.5	
				Micrococcus lutes	12.5	
126	Gerontoxanthone A	*Cudrania cochinchinensis*		*Bacillus subtilis*	>25	60
				Staphylococcus aureus	>25	
				Staphylococcus aureus	>25	
				Micrococcus lutes	>25	
127	Bangangxanthone A	the twigs of *Garcinia hombroniana*		MRSA	32	14
				Staphylococcus	200	

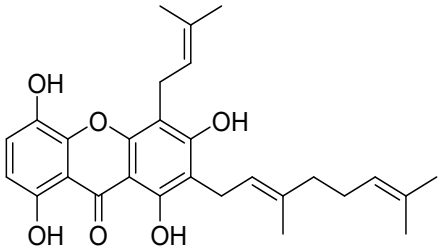
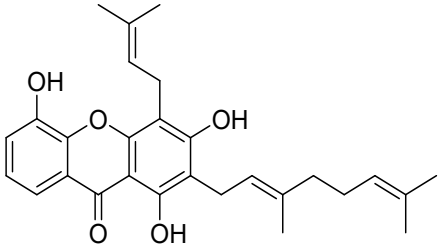
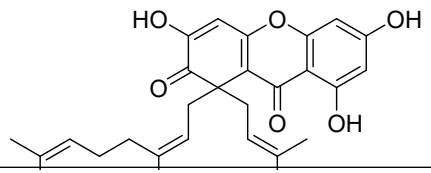
				aureus		
128	Nigrolineaxanthone I	the stem bark *Garcinia nigrolinaeta*		MRSA	4	20
129	Staudtiixanthone D	the twigs of *Garcinia staudtii*		MRSA	64	19
130	Mangostanin	*Tetragonula laeviceps* propolis; the fruits of Garcinia cowa		*Bacillus cereus*	12.5\4\8	13,37,47
				Listeria monocytogenes	0.78	
				Bacillus subtilis	2	
				Micrococcus luteus	25\4	
				Staphylococcus aureus	12.5\64\4	
				Staphylococcus epidermidis	25	
				Streptococcus pyogenes	3.13	
				MRSA	12.5\4	
				Escherichia coli	25	

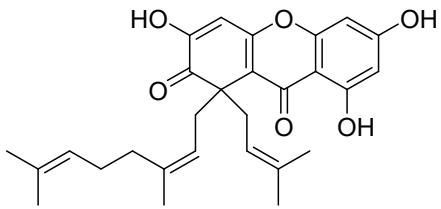
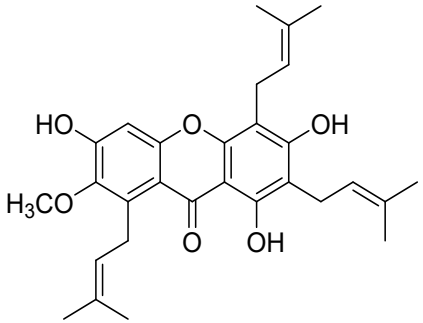
				Bacillus subtilis	8	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	128	
131	Garciniacowone C	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	32	47
				Bacillus subtilis	64	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	128	
Triprenyl group						
132	Norcowanin	*Garcinia cowa*; the root of *Cratoxylum formosum*		*Escherichia coli*	64	1,29
				Salmonella typhimurium	128	
				Bacillus subtilis	4.7	
				Staphylococcus aureus	9.37\8	
				Enterococcus faecalis	9.37	
				MRSA	18.75\16	
				VRE	75	

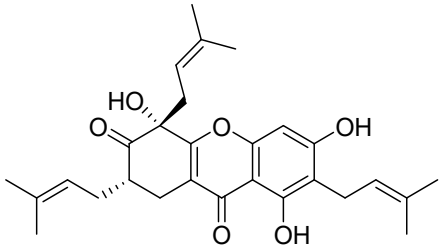
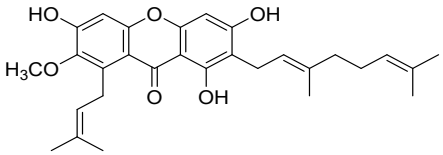
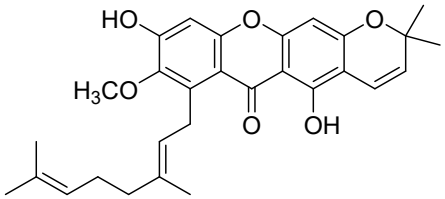
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Pseudomonas aeruginosa	300	
133	Cowanin	the fruits, inflorescences of *Garcinia cowa*; *Garcinia fusca*		*Staphylococcus aureus*	32\128\>128	1-2, 13,27,46
				MRSA	2\>128	
				Escherichia coli	128\>128	
				Bacillus cereus	32	
				Bacillus subtilis	4	
				Micrococcus luteus	4	
				Pseudomonas aeruginosa	128	
				Salmonella typhimurium	64	
				Staphylococcus epidermidis	2	
				Helicobacter pylori	29.89\3.73	
134	Cowanol	the fruits, inflorescences of *Garcinia cowa*, *Garcinia fusca*		MRSA	2\>128	1,13,27,46
				Staphylococcus aureus	8\>128	
				Escherichia coli	128	
				Salmonella typhimurium	128	

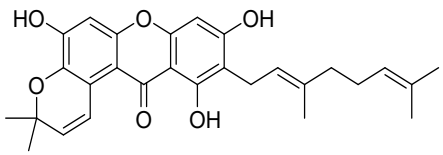
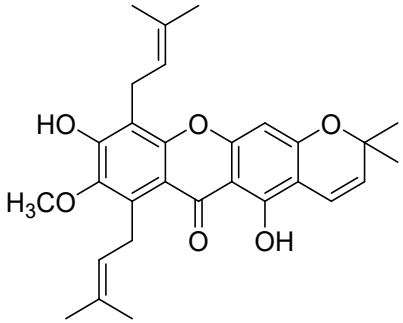
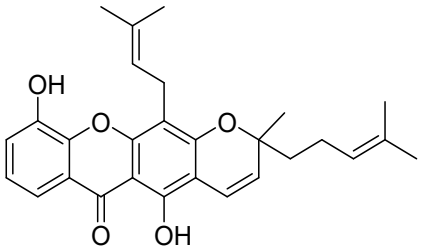
				Helicobacter pylori	3.84\30.87	
135	Cowagarcinone E	*Garcinia cowa*		MRSA	8	1-2
				Staphylococcus aureus	>128	
				Escherichia coli	128	
				Salmonella typhimurium	128	
136	Cochinchinone A	the root of *Cratoxylum formosum*		*Bacillus subtilis*	150	29
				Staphylococcus. aureus	150	
				Enterococcus faecalis	150	
				MRSA	9.37	
				VRE	150	
				Salmonella typhi	>150	
				Shigella sonei	>150	
				Pseudomonas aeruginosa	4.7	
137	Pruniflorone L	the root of *Cratoxylum formosum*		*Bacillus subtilis*	>300	29
				Staphylococcus. aureus	>300	
				Enterococcus faecalis	>300	
				MRSA	>300	
				VRE	>300	
				Salmonella typhi	>300	

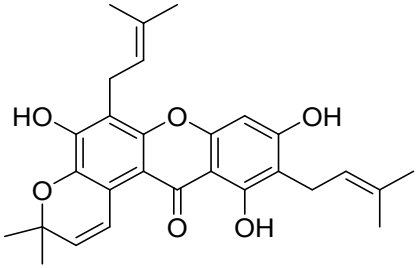
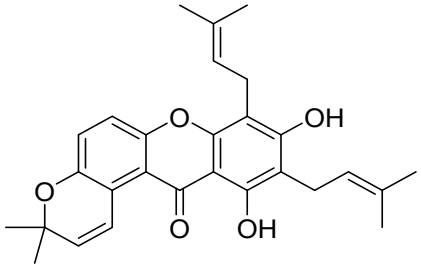
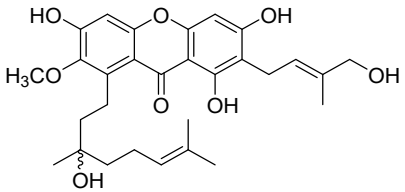
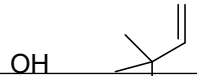
				Shigella sonei	>300	
				Pseudomonas aeruginosa	>300	
138	1,3,5,8-tetrahydroxy-2-(3-methybut-2-enyl)-4-(3,7-dimethyl-octa-2,6-dienyl)-xanthone	*Garcinia smeathmannii*		*Staphylococcus aureus*	64	53
				Enterococcus faecalis	8	
				Bacillus cereus	256	
				Pseudomonas aeruginosa	16	
				Escherichia coli	32	
				Salmonella typhimurium	64	
139	7-*O*-methyl-garcinone E	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	32	13,47
				Bacillus subtilis	32	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	128	
				Pseudomonas aeruginosa	128	
				MRSA	64	
140	Garcinone E	the bark of *Garcinia mangostana*; *Garcinia mangostana*		MRSA	3.9\25\62.4	38,54
				Staphylococcus	25\12.5	

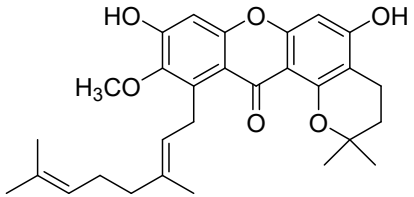
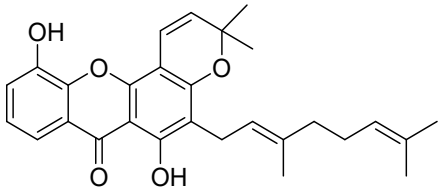
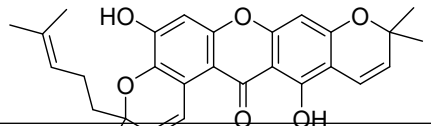
				aureus		
				Escherichia coli	25	
				Vibrio vulnificus	15.6	
				Vibrio rotiferianus	15.6	
				Vibrio campbellii	31.2	
141	4-prenyl-2-(3,7-dimethyl-2.6-octadienyl)-1,3,5,8-tetrahydroxy-xanthone			*Escherichia coli*	64\128	21
				Enterobacter aerogenes	128\256\64	
				Klebsiella pneumoniae	512	
				Providencia stuartii	64\256	
				Pseudomonas aeruginosa	64\128	
142	Formoxanthone A	the roots of *Cratoxylum formosum*		*Pseudomonas aeruginosa*	>50	32
				Bacillus subtilis	18.7	
				Staphylococcus aureus	37.5	
				Streptococcus faecalis	>50	
				Salmonella typhi	>50	
143	Garcinianone B	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	2	47
				Bacillus subtilis	2	
				Staphylococcus aureus	64	

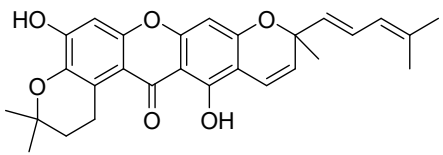
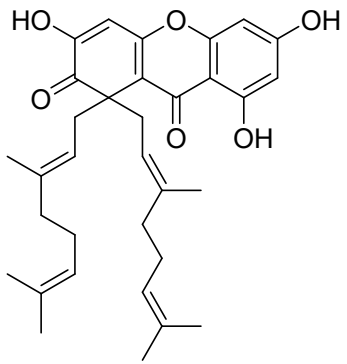
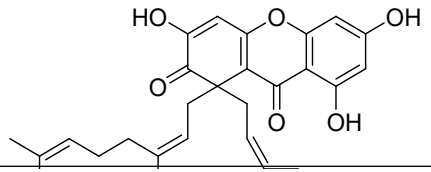
				Escherichia coli	64	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	
144	Garcinianone A	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	4	47
				Bacillus subtilis	2	
				Staphylococcus aureus	64	
				Escherichia coli	64	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	
				Staphylococcus aureus	16	
				MRSA	26	
145	mangostanaxanthone Ⅱ	*Garcinia mangostana*		*Escherichia coli*	794	3
				Bacillus cereus	1000	
				Staphylococcus aureus	1000	

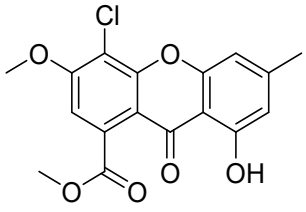
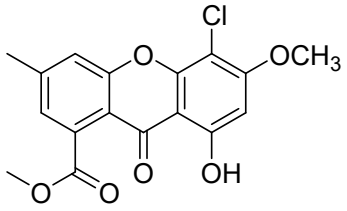
146	Garciniacowone	*Garcinia cowa*		MRSA	2	1
				Staphylococcus. aureus	2	
				Escherichia coli	128	
				Salmonella typhimurium	128	
147	2-(3,7-Dimethyl-2,6-octadien-1-yl)-1,3,6-trihydroxy-7-methoxy-8-(3-methyl-2-buten-1-yl)-9*H*-xanthen-9-one	*Cratoxylum cochinchinense*		MRSA	>128	2
				Staphylococcus aureus	>128	
				Escherichia coli	>128\64	
				Salmonella typhimurium	128	
				Bacillus cereus	>128	
				Bacillus subtilis	128	
				Pseudomonas aeruginosa	128	
148	Fuscaxanthone A	the fruits and flowers *Garcinia cowa*; *Garcinia fusca*		MRSA	>128\8	1-2, 46-47
				Staphylococcus aureus	>128	
				Micrococcus luteus	64	
				Escherichia coli	>128\64	
				Salmonella typhimurium	64	
				Staphylococcus	64	

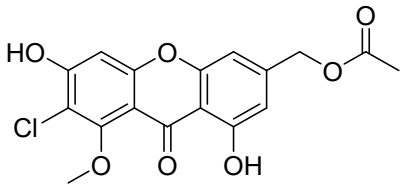
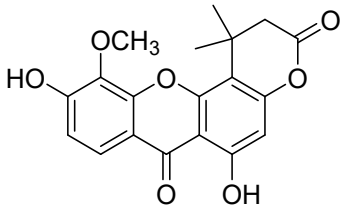
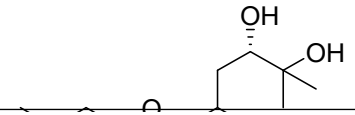
				epidermidis		
				Helicobacter pylori	3.70\7.76\29.76	
				Bacillus cereus	64	
				Bacillus subtilis	64	
				Pseudomonas aeruginosa	128	
149	Allanxanthone B	the stem bark of *Allanblackia monticola*		*Staphylococcus aureus*	\	12
				Vibrio anguillarium	\	
				Candida tropicalis	\	
150	Cowaxanthone C	the fruits of *Garcinia cowa*		*Staphylococcus aureus*	128	13
				MRSA	128	
151	Pruniflorone K	the root of *Cratoxylum formosum*		*Bacillus subtilis*	300	29
				Staphylococcus. aureus	>300	
				Enterococcus faecalis	>300	
				MRSA	>300	
				VRE	>300	

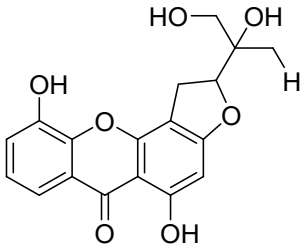
				Salmonella typhi	>300	
				Shigella sonei	>300	
				Pseudomonas aeruginosa	>300	
152	Tovophyllin A	the stem bark of Allanblackia monticola		*Staphylococcus aureus*	\	12
				Vibrio anguillarium	\	
				Candida tropicalis	\	
153	Batukinaxanthone	Calophyllum		*Staphylococcus aureus*	\	16
154	Fuscaxanthone I	*Garcinia fusca*		*Helicobacter pylori*	7.99\3.99\320	46
155	Cratosumatranone C	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	

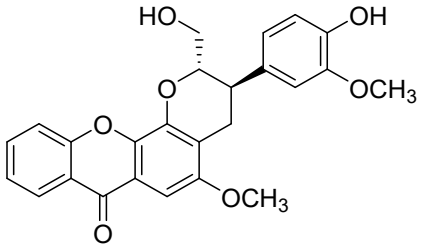
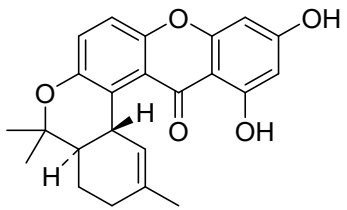
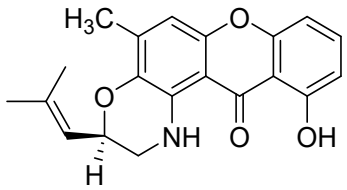
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	
156	Fuscaxanthone G	*Garcinia fusca*		*Helicobacter pylori*	7.46\3.72\29.87	46
157	Formoxanthone B	the roots of *Cratoxylum formosum*		*Pseudomonas aeruginosa*	>50	32
				Bacillus subtilis	4.6	
				Staphylococcus aureus	2.3	
				Streptococcus faecalis	18.7	
				Salmonella typhi	4.6	
158	Garciniacowone D	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	64	47
				Bacillus subtilis	128	
				Staphylococcus	>128	

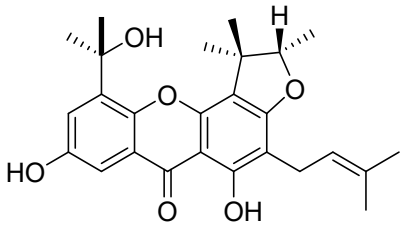
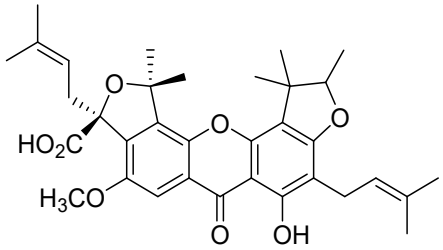
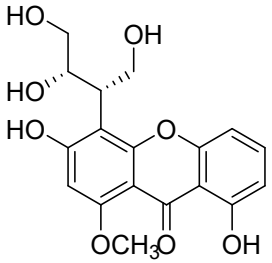
				aureus		
				Escherichia coli	>128	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	
159	Garcicowanone B	*Garcinia cowa*		*Bacillus cereus*	>128	2
Tetraprenyl group						
160	Garciniacowone A	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	8	47
				Bacillus subtilis	8	
				Staphylococcus aureus	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	128	
161	Garciniacowone B	the fruits and flowers of *Garcinia cowa*		*Bacillus cereus*	8	47
				Bacillus subtilis	8	
				Staphylococcus aureus	64	

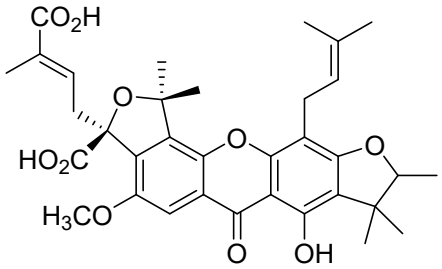
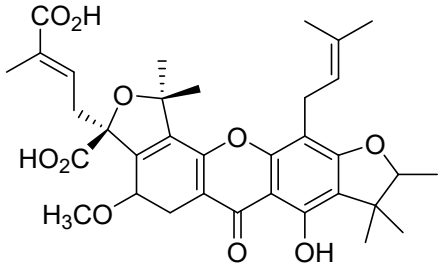
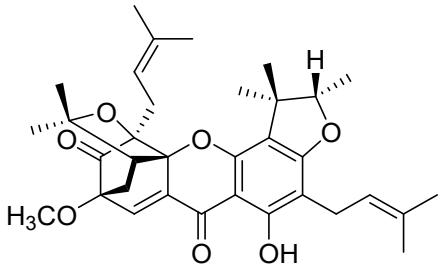
				Escherichia coli	64	
				Salmonella typhimurium	64	
				Pseudomonas aeruginosa	129	
Others						
162	Chloroisosulochrin dehydrate			MRSA	50	41
				Staphylococcus aureus ATCC 25923	50	
				Escherichia coli ATCC 25922	50	
				Baciluus cereus ATCC 11778	50	
				Vibrio alginolyticus ATCC 17749	50	
				Vibrio parahaemolyticus ATCC 17802	50	
163	4-chloro-1-hydroxy-3-methoxy-6-methyl-8-methoxycarbonyl-xanthen-9-one	fungus *Penicillium citrinum* HL-5126		MRSA	>17.4	41
				Staphylococcus aureus	>17.4	
				Escherichia coli	>17.4	
				Baciluus cereus	>17.4	
				Vibrio alginolyticus	>17.4	

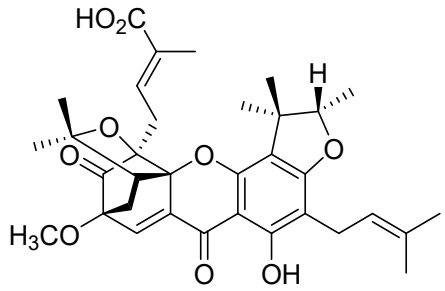
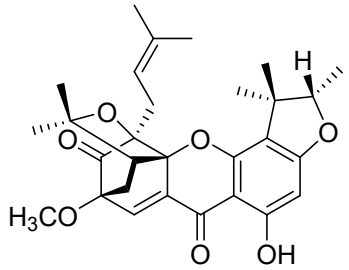
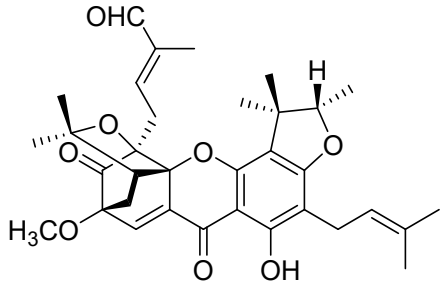
				Vibrio parahaemolyticus	>17.4	
164	2′-acetoxy-7-chloro-citreorosein	fungus *Penicillium citrinum* HL-5126		MRSA	>18.2	41
				Staphylococcus aureus	8.30	
				Escherichia coli	>18.2	
				Baciluus cereus	>18.2	
				Vibrio alginolyticus	>18.2	
				Vibrio parahaemolyticus	3.64	
165	Neriifolone B	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	
166	14-Hydroxyltajixanthone hydrate	the fungus *Emericella* sp. XL029		*Micrococcus luteus*	25	62
				Staphylococcus. aureus	25	

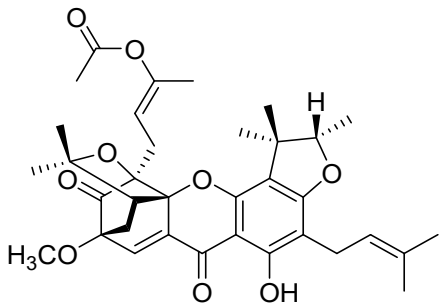
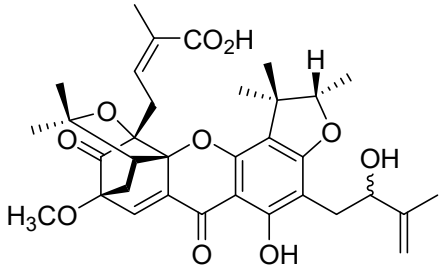
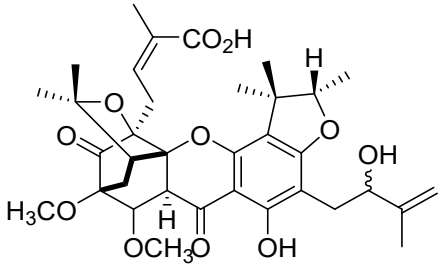
				Bacillus megaterium	25	
				Bacillus anthracis	25	
				Bacillus paratyphosum	25	
				MRSA	>100	
				Pseudomonas aeruginosa	50	
				Bacillus cereus	50	
				Enterobacter aerogenes	50	
167	*O*-demethyl-3′,4′-deoxypsorospermin-3′,4′-diol	*Vismia laurentii*		*Citrobacter freundii*	\	32
				Enterobacter aerogens	\	
				Enterobacter cloacae	19.53	
				Escherichia coli	\	
				Klebsiella pneumoniae	78.12	
				Morganella morganii	39.06	
				Proteus mirabilis	19.53	
				proteus vulgaris	\	
				Pseudomonas aeruginosa	\	
				Shigella dysenteriae	\	
				Shigella flexneri	39.06	

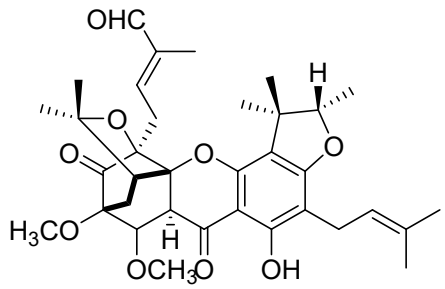
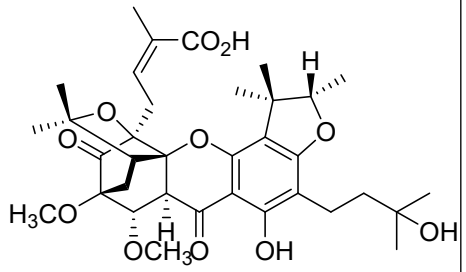
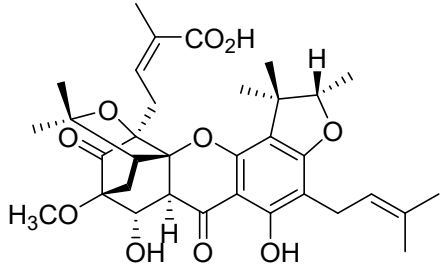
				Salmonella typhi	\	
				Streptococcus faecalis	78.12	
				Staphylococcus aureus	19.53	
				Bacillus cereus	\	
				Bacillus megaterium	\	
				Bacillus stearothermophilus	\	
				Bacillus subtilis	1.22	
168	Kielcorin	*Kielmeyera variabilis*		MRSA	>512	40
169	Calozeyloxanthone	Calophyllum		*Staphylococcus aureus*	4.1-8.3	16
				VRE	6.25	
				VSE	12.5	
170	Fusarioxazin	*Fusarium oxysporum*		*Staphylococcus aureus*	5.3	22
				Bacillus cereus	3.7	

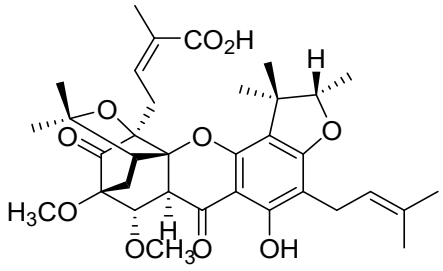
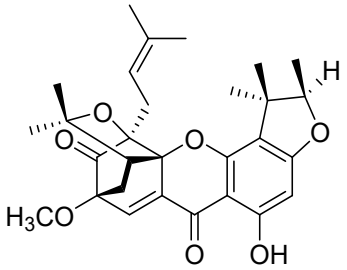
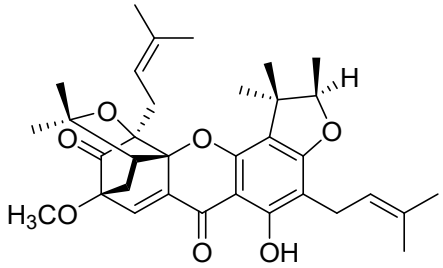
171	Scortechinone U	the fruits of *Garcinia scortechinii*		MRSA	>127.95	34-35
172	Scortechinone V	the fruits of *Garcinia scortechinii*		MRSA	35.20	34-35
173	Isosecosterigmatocystin	fungus *Aspergillus nidulans*		*Escherichia coli*	\	36
				Micrococcus luteus	\	
				Vibrio vulnificus	\	
				Vibrio anguillarum	\	
				Vibrio alginolyticus	\	
				Ed ictalurid	16	
				Vibrio parahaemolyticus	\	

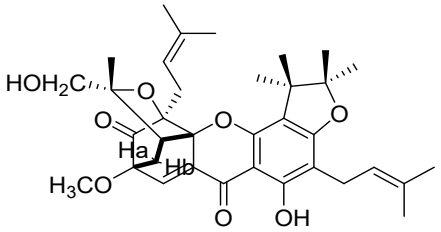
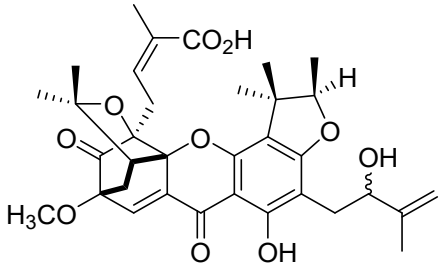
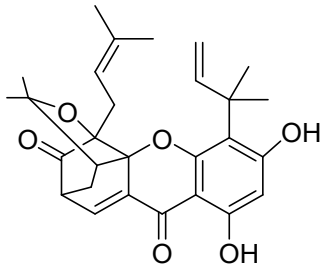
174	Scortechinone W	*Garcinia scortechinii*		MRSA	34.86	34
175	Scortechinone X	*Garcinia scortechinii*		MRSA	>127.73	34
Caged xanthones						
176	Scortechinone L	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>64	49
				Staphylococcus aureus	>64	

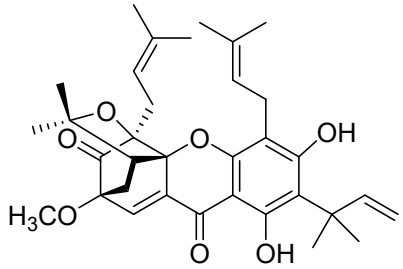
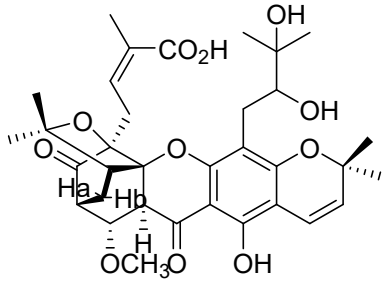
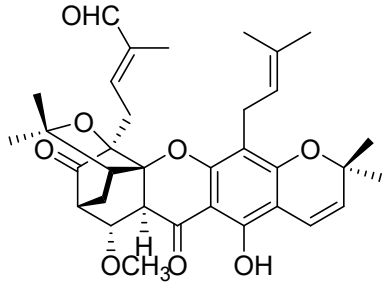
177	Scortechinone F	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	16\4	49
178	Scortechinone E	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>256	49
179	Scortechinone H	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>64\4	49

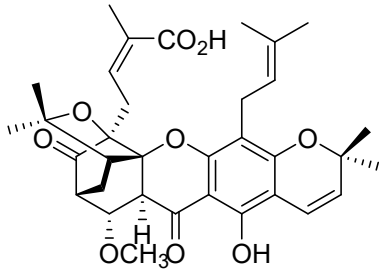
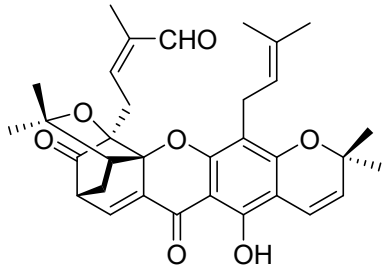
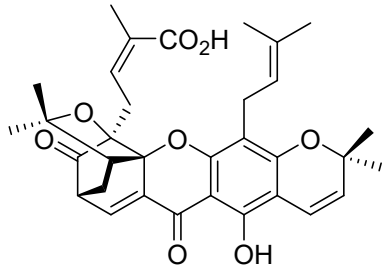
180	Scortechinone G	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>64	49
181	Scortechinone C	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	32	49
182	Scortechinone S	the fruits of *Garcinia scortechinii*		MRSA	64	34-35

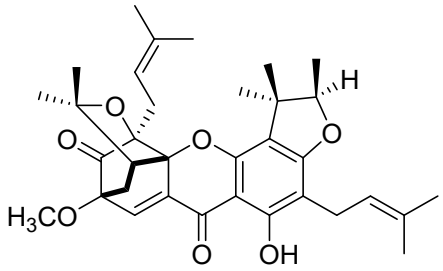
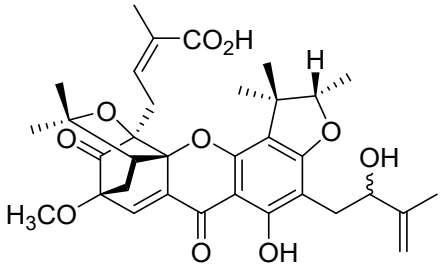
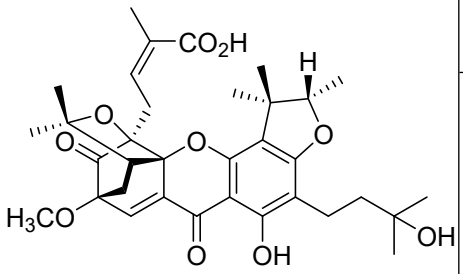
183	Scortechinone T	the fruits of *Garcinia scortechinii*		MRSA	>63.87	34-35
184	Scortechinone O	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>128	49
185	Scortechinone P	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	32\16	49

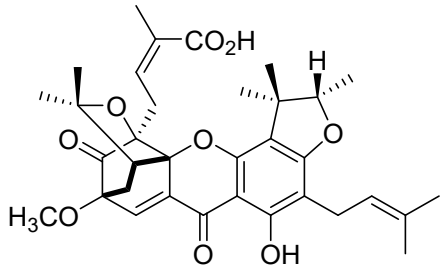
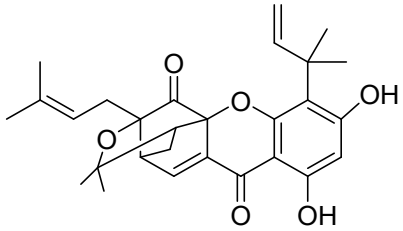
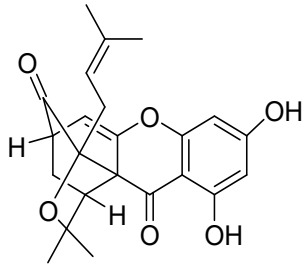
186	Scortechinone I	the stem bark and fruits of *Garcinia scortechinii*		*MRSA*	7.99	34-35, 49
187	Scortechinone D	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	>256	49
188	Scortechinone A	the stem bark of *Garcinia scortechinii*		*MRSA*	128.2	49

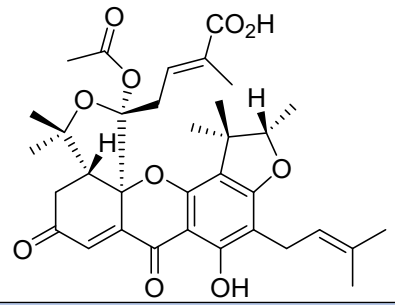
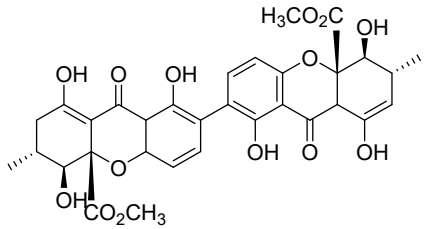
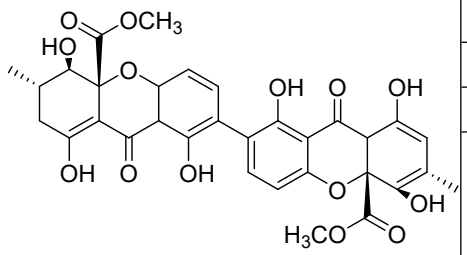
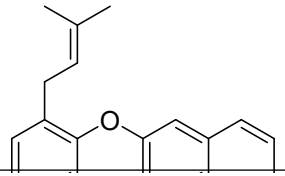
189	Scortechinone Q	the fruits of *Garcinia scortechinii*		MRSA	>130.45	34-35
190	Scortechinone R	the fruits of *Garcinia scortechinii*		MRSA	>127.7	34-35
191	Bractatin	the roots of *Garcinia propinqua*		*Bacillus cereus*	8	43
				Bacillus subtilis	1	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	

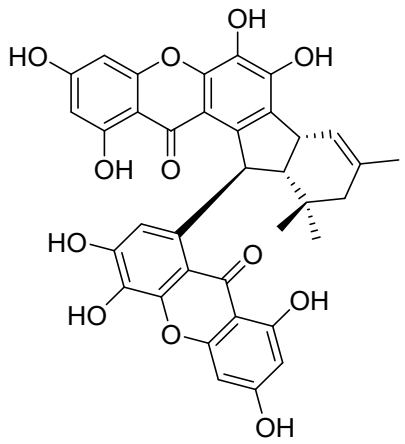
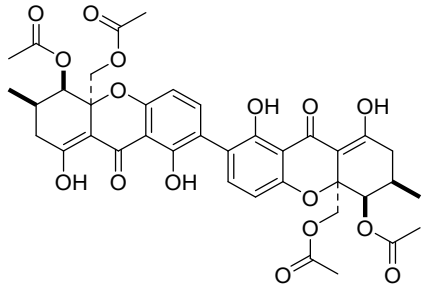
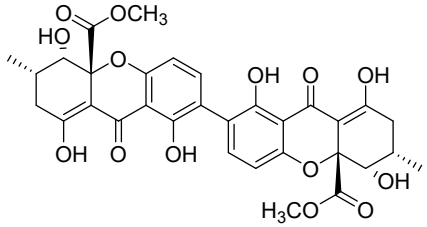
192	Scortechinone J	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	32\8	49
193	Hanburinone	the fruits of *Garcinia hanburyi*		MRSA	124.85	52
194	Isomoreollin B	the fruits of *Garcinia hanburyi*		MRSA	115.25	52

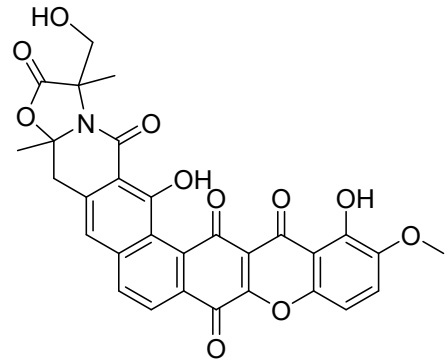
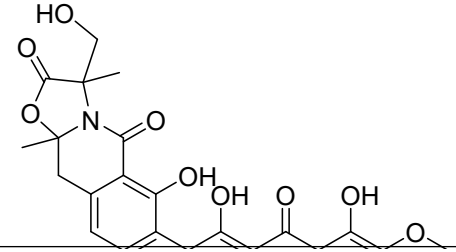
195	Moreollin acid	the fruits of *Garcinia hanburyi*		MRSA	14.80	52
196	Morellin	the fruits of *Garcinia hanburyi*		MRSA	108.85	52
197	Morellic acid	the fruits of *Garcinia hanburyi*		MRSA	14	52

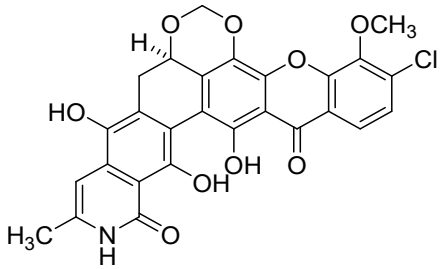
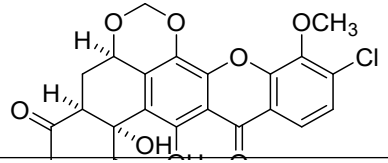
198	Scortechterpene A	the fruits of *Garcinia scortechinii*		MRSA	128	34,35
199	Scortechinone M	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	32	49
				Staphylococcus aureus	32	
200	Scortechinone N	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	32	49
				Staphylococcus aureus	32	

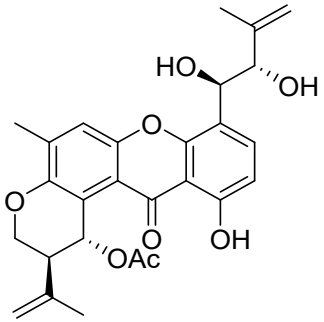
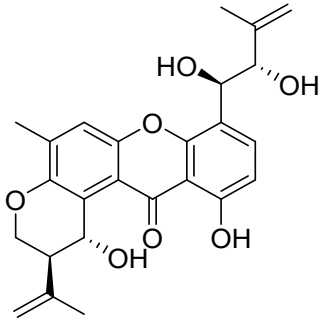
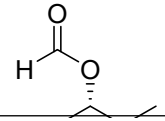
201	Scortechinone B	the fruits of *Garcinia scortechinii*		MRSA	2	34,35
202	Neobractatin	the roots of *Garcinia propinqua*		*Bacillus cereus*	32	43
				Bacillus subtilis	1	
				Escherichia coli	128	
				Staphylococcus typhimurium	128	
203	Cochinchinoxanthone	the roots of *Cratoxylum sumatranum*		*Micrococcus luteus*	>128	51
				Bacillus cereus	>128	
				Bacillus subtilis	\	
				Staphylococcus aureu	>128	
				Staphylococcus epidermidis	>128	
				Escherichia coli	>128	
				Salmonella typhimurium	>128	
				Pseudomonas aeruginosa	>128	

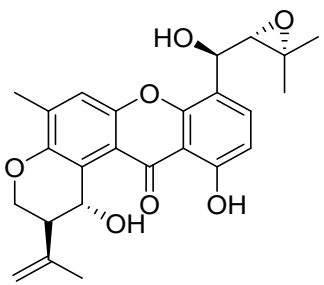
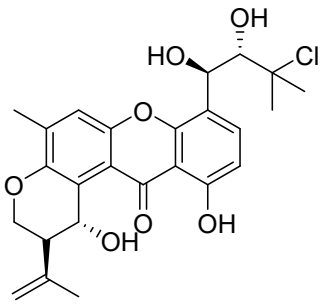
204	Scortechinone K	the stem bark of *Garcinia scortechinii*		*Staphylococcus aureus*	128	49
Bi-xanthones						
205	Secalonic acid A			*Staphylococcus aureus*	12.5	70
				Escherichia coli	25	
				Sarcina ventriculi	12.5	
				Pseudomonas aeruginosa	12.5	
206	Secalonic acid D			*Bacillus subtillis*	24.4	69
				Escherichia coli	24.4	
				Micrococcus luteus	24.4	
				Pseudoalteromonas nigrifaciens	97.5	
207	Globulixanthone E	the root bark of *Symphonia globulifera*		*Staphylococcus aureus*	4.51	16
				Bacillus subtilis	3.12	
				Vibrio anguillarium	5.56	

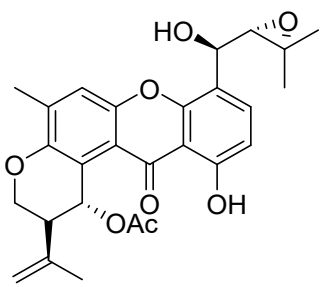
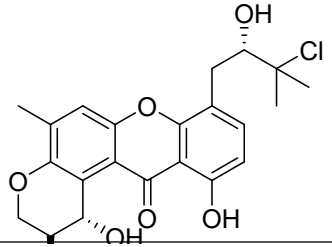
				Escherichia coli	>100	
208	Garmoxanthone	the bark of *Garcinia mangostana*		MRSA	3.9	38
				Vibrio vulnificus	15.6	
				Vibrio rotiferianus	15.6	
				Vibrio campbellii	31.2	
209	Dicerandrol C			*Staphylococcus aureus*	1/1.33	71
				Stapholococcus saprophyticus	2/2.66	
210	Secalonic acid B			*Bacillus subtillis*	24.4	72
				Escherichia coli	24.4	
				Micrococcus luteus	24.4	
				Pseudoalteromonas nigrifaciens	97.5	

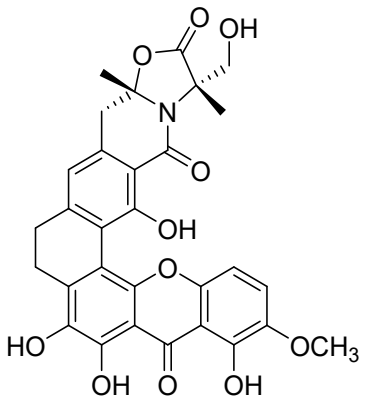
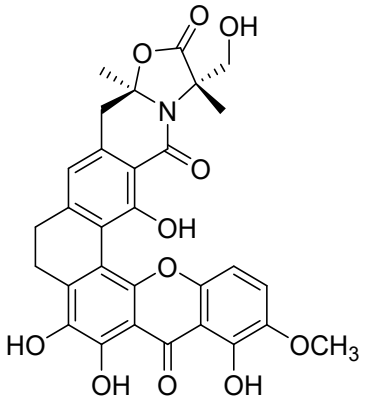
Other xanthone derivatives						
211	Citreamicin δ	*Streptomyces vinaceus*		*Staphylococcus aureus*	≤0.5\1\2	24
				VRE	≤0.25	
				Streptococcus pneumoniae	≤0.06	
				Haemophilus influenzae	1	
				Hemophilius parainfluenzae	1	
				Escherichia coli	>64	
				Klebsiella pneumoniae	>64	
				Moraxella catarrhalis	4	
				Serratia marcescens	>64	
				Pseudomonas aeruginosa	>64	
212	Citreamicin ε	*Streptomyces vinaceus*		*Staphylococcus aureus*	0.13\0.12\0.25	24
				VRE	0.5	
				Streptococcus pneumoniae	0.25\0.12	
				Haemophilus	>32	

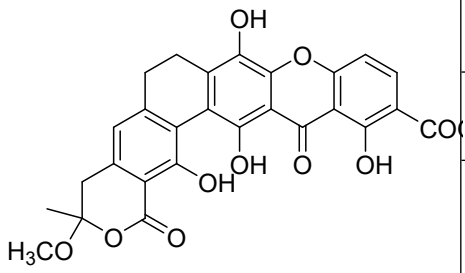
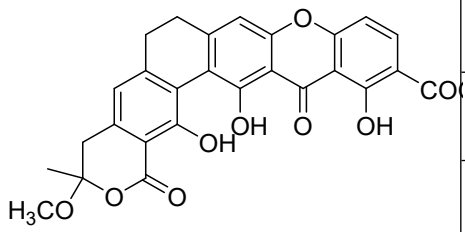
				influenzae		
				Hemophilius parainfluenzae	>32	
				Escherichia coli	>32	
				Klebsiella pneumoniae	>32	
				Moraxella catarrhalis	0.5	
				Serratia marcescens	>32	
				Pseudomonas aeruginosa	>32	
213	Xantholipin B	*Streptomyces flocculus* CGMCC 4.1223		*Bacillus mycoides*	0.08	26
				Mycobacterium smegmatis	5.0	
				Staphylococcus aureus	0.025	
				Pseudomonas aeruginosa	>30	
				Klebsiella pneumoniae	>30	
				Escherichia coli	>30	
214	Xantholipin	*Streptomyces flocculus* CGMCC 4.1223		*Bacillus mycoides*	0.08	26
				Mycobacterium smegmatis	5.0	
				Staphylococcus	0.025	

				aureus		
				Pseudomonas aeruginosa	>30	
				Klebsiella pneumoniae	>30	
				Escherichia coli	>30	
215	Aspergixanthone G	fungus *Aspergillus* sp.		*Micrococcus lysodeikticus*	0.78	39
				Bacillus anthraci	12.5	
				Salmonella typhi	6.13	
				Enterobacter aerogenes	6.13	
216	Aspergixanthone H	fungus *Aspergillus* sp.		*Micrococcus lysodeikticus*	6.13	39
				Bacillus anthraci	12.5	
				Salmonella typhi	6.13	
				Enterobacter aerogenes	6.13	
217	Varixanthone	the fungus of *Emericella variecolor*		*Escherichia coli*	12.5	62
				Proteus sp.	12.5	
				Bacillus subtilis	12.5	

				Staphylococcus. aureus	12.5	
218	14-Hydroxyltajixanthone	the fungus *Emericella* sp. XL029		*Micrococcus luteus*	25	62
				Staphylococcus. aureus	25	
				Bacillus megaterium	12.5	
				Bacillus anthracis	12.5	
				Bacillus paratyphosum	12.5	
				MRSA	50	
				Pseudomonas aeruginosa	50	
				Bacillus cereus	50	
				Enterobacter aerogenes	25	
220	14-Hydroxyl-15-chlorotajixanthone hydrate	the fungus *Emericella* sp. XL029		*Micrococcus luteus*	25	62
				Staphylococcus. aureus	25	
				Bacillus megaterium	25	
				Bacillus anthracis	25	
				Bacillus paratyphosum	25	
				MRSA	>100	

				Pseudomonas aeruginosa	50	
				Escherichia coli	50	
				Enterobacter aerogenes	50	
221	14-Methoxyltajixanthone-25-acetate	the fungus *Emericella* sp. XL029		*Micrococcus luteus*	25	62
				Staphylococcus. aureus	25	
				Bacillus megaterium	12.5	
				Bacillus anthracis	25	
				Bacillus paratyphosum	25	
				MRSA	>100	
				Pseudomonas aeruginosa	50	
				Escherichia coli	50	
				Enterobacter aerogenes	50	
222	15-Chlorotajixanthone hydrate	the fungus *Emericella* sp. XL029		*Micrococcus luteus*	50	62
				Staphylococcus. aureus	25	
				Bacillus megaterium	12.5	
				Bacillus anthracis	50	
				Bacillus	25	

				paratyphosum		
				MRSA	>100	
				Pseudomonas aeruginosa	50	
				Escherichia coli	50	
				Enterobacter aerogenes	50	
223	Citreamicin θ A	*Streptomyces caelestis*		*Staphylococcus haemolyticus*	0.5	64
				Staphylococcus aureus	1\0.25	
				Bacillus subtillis	0.25	
224	Citreamicin θ B	*Streptomyces caelestis*		*Staphylococcus haemolyticus*	0.5	64
				Staphylococcus aureus	1\0.25	
				Bacillus subtillis	0.25	

225	Citreaglycon A	*Streptomyces caelestis*		*Staphylococcus haemolyticus*	8.0	64
				Staphylococcus aureus	16.0\8	
				Bacillus subtillis	8.0	
226	Dehydrocitreaglycon A	*Streptomyces caelestis*		*Staphylococcus haemolyticus*	8.0	64
				Staphylococcus aureus	16.0	
				Bacillus subtillis	8.0	

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2. The correlation between log P and MIC values of simple xanthones.

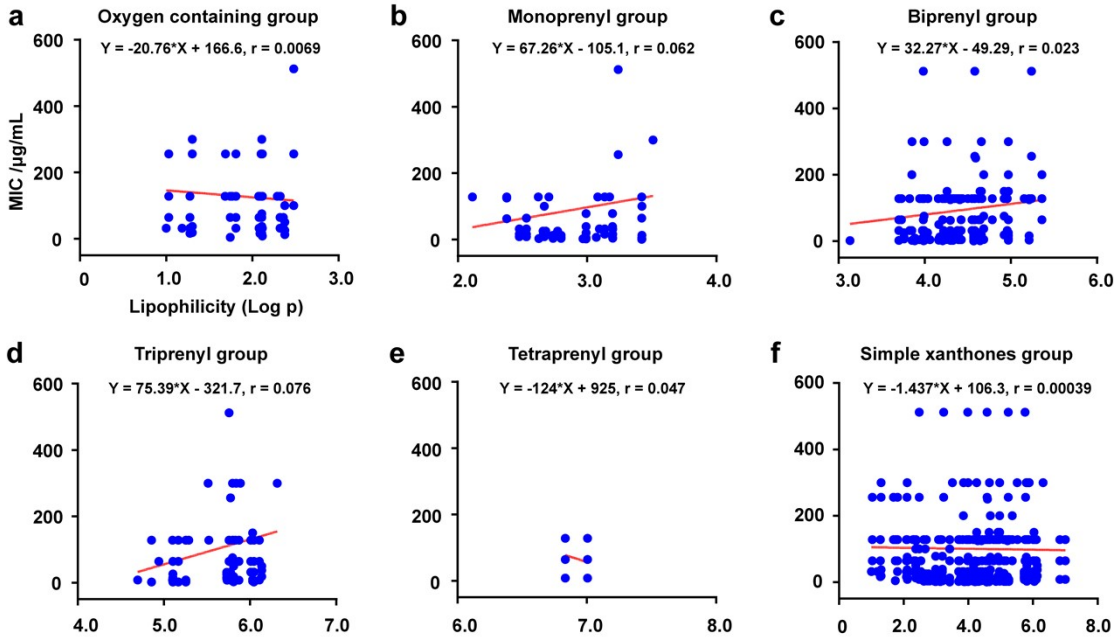

Fig. S1 The correlation between log P and MIC values of simple xanthones.