Journal of Applied Pharmaceutical Science Vol. 10(05), pp 142-157, May, 2020 Available online at http://www.japsonline.com DOI: 10.7324/JAPS.2020.10519 ISSN 2231-3354



Marine sponge compounds with antiplasmodial properties: Focus on *in vitro* study against *Plasmodium falciparum*

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ARTICLE INFO

Received on: 09/10/2019 Accepted on: 18/02/2020 Available online: 06/05/2020

Key words: Bioactive compounds, drug development, antimalarial, marine natural product, *Porifera*.

ABSTRACT

Malaria continues to be a major cause of morbidity and mortality in many tropical countries. The lack of progress in drug discovery and the spread of drug resistance become the reason behind this. *Porifera* (sponges) is a potential source of novel bioactive compounds to provide future drugs against malaria. In this review, we summarized 243 isolated molecules belonging to 35 different genera that active against *Plasmodium falciparum* from published paper until March 2019. The molecules were classified into potent, good, moderate, low, and inactive based on their IC₅₀, and among observed bioactive metabolites, there were 57 marine sponge molecules reported to act as potent antiplasmodium against various strains of *P. falciparum* including drug resistance and nondrug resistance. Table 2 represents the list of isolated compounds with "potent" antimalarial activity. The class of the listed compounds includes manzamine alkaloid, guanidine alkaloids, bispyrroloiminoquinone alkaloids, pyrroloiminoquinone alkaloids, bispyrcloiminoquinone alkaloids, and sterols. With this up-to-date review, we attempt to present new perspectives for the rational discovery of novel sponge metabolites that can be used as lead compounds in antimalarial drug development.

INTRODUCTION

Malaria is the most life-threatening and infectious disease caused by *Plasmodium* parasites such as *Plasmodium falciparum*, *Plasmodium ovale*, *Plasmodium vivax*, *Plasmodium malariae*. Among those protozoans, *P. falciparum* is considered to be responsible for most severe diseases and most fatal cases. The World Health Organization (2018) stated in the year of 2017 that more than 99% of estimated malaria cases in the WHO African Region followed by the WHO regions of the Western Pacific (71.9%), the Eastern Mediterranean (69%), and Southeast Asia (62.8%) were caused by this most prevalent malaria parasite. In the same period, the WHO reported approximately 219 million cases of malaria occurred worldwide including 435,000 deaths.

Nowadays, malaria continues to be a major cause of morbidity and mortality in tropical countries. It is further aggravated by an increase in a number of multidrug-resistant strains of *Plasmodium* accompanied by a lack of progress in the development of vaccines and drug discovery. As a consequence, the search of new agent that actives against malaria becomes urgent needs (Antony and Parija 2016; Burrows *et al.*, 2011; Cui *et al.*, 2015; Dondorp *et al.*, 2000; Noedl *et al.*, 2008).

Marine ecosystems are the largest part of the biosphere. More than 70% of the Earth's surface is covered by water, and several theories believe that the life on earth originated from the ocean. In certain marine ecosystems such as coral reefs or the deep-sea floor, scientists estimate that the diversity of marine biota is even greater than the biota inhabiting tropical rainforests. Many immotile or slow-moving marine invertebrates, which usually do not have physical protection such as shells or thorns, will produce secondary metabolites as a form of defense mechanism from the environment and other creatures in the ocean (Ebada *et al.*, 2008). These compounds attract the attention of researchers from various fields such as chemistry, pharmacology, biology, and ecology. This

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statement is supported by the fact that the number of new bioactive constituents isolated from marine biota has been increasing in the past three decades (332 compounds were isolated in 1984, and 1490 new compounds were isolated in 2017) (Blunt *et al.*, 2016; Carroll *et al.*, 2019).

Exploration of secondary metabolites from marine organisms is expected to provide new active substituents against various diseases (Newman and Cragg, 2007). Several studies have managed to isolate metabolites from marine microorganisms, green, red, and brown algae, phytoplankton, *Cnidaria, Bryozoa*, molluscs, tunicates, echinoderms, mangroves, sponges, and itertidal plants which have proven to have pharmaceutical properties such as acetylcholinesterase inhibitor, radical scavenging activity, cytotoxicity, antimicrobial, anticancer, antitumor, hemolytic, anti-inflammatory, antiparasitic, antimalarial, and antifungal (Blunt *et al.*, 2016; D'Ambrosio *et al.*, 1996; Fattorusso and Taglialatela-Scafati 2009; Orhan *et al.*, 2010; Rama Rao and Faulkner 2002; Setyowati *et al.*, 2009; 2017a; 2017b).

From the perspective of drug discovery, a marine sponge is one of the invertebrate organisms which is interesting to be explored due to its potency producing new compounds (Anjum *et al.*, 2016). The lack of physical defense of sponges resulting in secondary metabolites is estimated to vary depending on their habitats. Metabolite compounds isolated from sponges are highly diverse such as alkaloids, esters, fatty acids, glycosides, ketones, lipids, macrolides, peptides, peroxides, quinones, terpenoids, and polyketides and have shown many biological activities, in which one of them is antimalaria (Blunt *et al.*, 2016; 2017; 2018; Carroll *et al.*, 2019). These kinds of compounds have been found to interfere with pathogenesis at many distinct points; therefore, this can be beneficial in developing selective antimalarial drugs (Sipkema *et al.*, 2005)

The aim of this review is to summarize compounds isolated from marine sponges which exhibit *in vitro* antiplasmodial properties, to identify the compounds with potent activity based on their IC_{50} values, and to highlight the most important functional groups of the compounds related to their potent activity against various strains of *P. falciparum*. One of the advantages of an *in vitro* study is that the study could thoroughly illustrate an effect of structural features of tested compounds to their activity with no interference from other factors such as biological system which can be found on *in vivo* study. Therefore, it can be used to generate more potent derivatives of the compounds to develop selective antimalaria drugs that work in blood-stage *P. falciparum*.

METHOD

A systematic search was accomplished to find all publications related to the theme until March 2019 in PubMed and Google Scholar. The keywords used to search the articles were "*Plasmodium falciparum*, sponge, antimalarial" or "*Plasmodium falciparum*, sponge, antiplasmodial." The data included in the review were primary articles in English about *in vitro* antimalarial study of pure compounds isolated from marine sponges against *P. falciparum* as shown in Table 1. The articles obtained were then removed if they are review articles, conference articles, and thesis, and there are no data available to be retrieved. All the synthetic compounds derived from naturally

occurring metabolites in sponge are not mentioned in this review. Variables assessed in this review include sponge species/genus, isolated compound, strain of *P. falciparum*, region/country of origin, and effect on parasite growth inhibition.

EXPLORATION OF MARINE SPONGE METABOLITES FOR ANTIPLASMODIAL ASSAY

Among marine invertebrates, a sponge is the most dominant source for discovering natural products that have been used as lead compound to develop therapeutic drugs (Perdicaris et al., 2013). However, the study done in the investigation of marine sponge metabolites for antimalarial activity is relatively low compared to those of antitumor and anticancer. From literature published until March 2019, we included 50 primary articles for the review (Table 1). We identified that 35 different genera have been studied for their antiplasmodial activities and found that the most frequently studied genera were genus Agelas, Plakortis, and Xestospongia from different locations. Although many bioactive compounds have been isolated from marine sponges (Blunt et al., 2016; 2018; Carroll et al., 2019), the evaluation of their antiplasmodial activity is still relatively low. Figure 1 shows the number of studies that have been done on the examination of *in vitro* antiplasmodium of isolated compounds from marine sponge.

Overall, the number of publications from year to year shows fluctuation pattern. The highest number of the published papers was in the year of 2010 with 10 articles, followed by six publications in 2009 and 2012. In regard to the number of publications from 2013 to March 2019, it seemed to be stuck at one to three studies each year. This indicates that exploration trend of marine sponge metabolites for antiplasmodial activity diminished from 31 published papers during the period of 1992–2010 to 21 publications during the period of 2011–March 2019. One of the reasons behind the trend is that many scientists are interested in microbiological sample investigations for marine natural product exploration including bacteria and fungi sponge associated, making the detriment of sponge-derived compounds (Carroll *et al.*, 2019; Thomas *et al.*, 2010).

Various ecological studies have shown that secondary metabolites produced by sponges often serve defensive purposes to protect them from threats such as predator attacks, microbial infections, biofouling, and overgrowth by other sessile organisms (Paul and Puglisi, 2004; Paul et al., 2006). Therefore, compounds isolated from the same sponge species are more likely to be different if their habitat is distinct due to the ecological response (Mani et al., 2012). Moreover, a review done by Qaralleh (2016) found out that among 27 species of genus Neopetrosia, there are only nine species which have been chemically studied thus far. These facts disclose significant opportunities to do the chemical constituent exploration from not only genus Neopetrosia but also the other genus. In terms of collection site of the sponges, Australia, Bahamas, Indonesia, and Thailand were the most explored site so far for the search of compounds which exhibit in vitro antiplasmodium (P. falciparum strains). Other sponges were collected from Turkey, Vanuatu, Madagascar, Caledonia, Fiji, China, Japan, Alaska, Jamaica, Solomon Island, Puerto Rico, Papua New Guinea, and others (Table 1).

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (μM)	Origin	Ref.
1	Acanthella klethra	Axisonitrile 3	D6	0.61	Pelorus Island, Queensland, Australia	(Angerhofer et al., 1992)
			W2	0.07		
		Axisothiocyanate 3	D6	46.85		
			W2	11.81		
		The eudesmane compound A ^a	D6	8.50		
			W2	2.32		
		The eudesmane compound B ^b	D6	16.17		
			W2	2.22		
		The eudesmane compound C ^c	D6	>37.96		
			W2	>37.96		
2	Acanthostrongylophora ingens	(+)-8-hydroxymanzamine A	D6	0.03	Papua New Guinea	(Samoylenko et al., 2009)
			W2	0.04		
		(+)-manzamine A	D6	0.04		
			W2	0.05		
		(+)-8-hydroxymanzamine A hydrochloride	D6	0.04		
			W2	0.06		
		(+)-manzamine A hydrochloride	D6	0.01		
			W2	0.01		
3	Acanthostrongylophora sp.	Manzamine A	D6	0.01	Knife Cape Manado, Indonesia	(Rao <i>et al.</i> , 2006)
			W2	0.01		
		(+)-8-hydroxymanzamine A	D6	0.01		
			W2	0.01		
		Manzamine Y	D6	0.74		
			W2	1.50		
		Manzamine E	D6	6.02		
			W2	8.43		
		6-hydroxymanzamine E	D6	1.36		
			W2	1.50		
		Manzamine F	D6	1.34		
			W2	2.93		
		12,34-oxamanzamine A	D6	8.97		
			W2	na		
		Ent-12,34-oxamanzamine F	D6	1.45		
			W2	1.90		
		12,28-oxamanzamine A	D6 and W2	na		
		12,28-oxa-8-hydroxy-manzamine A	D6 and W2	na		
		12,34-oxamanzamine E	D6 and W2	na		
		12,28-oxamanzamine E	D6 and W2	na		
		12,34-oxa-6-hydroxymanzamine E	D6 and W2	na		
4	Acanthostrongylophora sp.	Manzamine A N-oxide	D6	0.02	Manado, Indonesia	(Rao et al., 2004)
			W2	0.02		
		3,4-dihydromanzamine A-N-oxide	D6	2.82		
			W2	6.53		
		Manzamine J	D6	2.36		
			W2	1.36		
		6-deoxymanzamine X	D6	2.30		
		-	W2	2.48		

Table 1. Summarized data of isolated compounds which have been tested for their antiplasmodial activity.

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No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM)	Origin	Ref.
		Manzamine X	D6	1.64		
			W2	3.44		
		Neo-kauluamine	D6	1.46		
			W2	2.41		
		Ircinal A	D6	5.82		
			W2	7.51		
		Ircinal A	D6 and W2	na		
5	Agelas cf. mauritiana	Agelasine J	FcB1	6.60	Solomon Islands	(Appenzeller et al., 2008)
		Agelasine K	FcB1	8.30		
		Agelasine L	FcB1	18.00		
6	Agelas gracilis	Gracilioethers A	ItG	28.22	Oshima-Shinsone, Japan	(Ueoka et al., 2009)
		Gracilioethers B	ItG	1.56		
		Gracilioethers C	ItG	31.02		
7	Agelas oroides	24-ethyl-cholest-5α-7-en-3-α-ol	K1	38.82	Go"kc,eada, Turkey	(Tasdemir et al., 2007)
		4,5-dibromopyrrole-2-carboxylic acid methyl ester	K1	>176.73		
		4,5-dibromopyrrole-2-carboxylic acid (free base)	K1	>185.95		
		4,5-dibromopyrrole-2-carboxylic acid (salt)	K1	136.37		
		(E)-oroidin (free base)	K1	10.02		
		(E)-oroidin (salt)	K1	16.25		
		3-amino-1-(2-aminoimidazoyl)-prop-1-ene	K1	53.56		
		Taurine	K1	>399.52		
8	Agelas dispar	Longamide B	K1	21.19	Little San Salvador Island	(Scala et al., 2010)
9	Agelas longissima	Longamide A	K1	>64.53	Little San Salvador Island	(Scala et al., 2010)
		Agelongine	K1	32.97		
10	Genus Agelas (A. conifera, A. clathrodes, A. longissima, and A. dispar)	Sceptrin	K1	17.86	Little San Salvador Island	(Scala et al., 2010)
		Hymenidin	K1	40.43		
		Dispacamide B	K1	4.11		
		Dispacamide D	K1	>58.45		
11	Aplysinella strongylata	19-hydroxypsammaplysin E	3D7	6.40	Tulamben Bay, Bali, Indonesia	(Mudianta et al., 2012)
		Psammaplysin K	3D7	nat 10 µM		
		Psammaplysin L	3D7	nat 10 µM		
		Psammaplysin M	3D7	nat 10 µM		
		Psammaplysin N	3D7	nat 10 µM		
		19-hydroxypsammaplysin P	3D7	nat 10 µM		
		Psammaplysin T	3D7	nat 10 µM		
		Psammaplysin V	3D7	nat 10 µM		
12	Axinyssa djiferi	Axidjiferosides (mix-A, -B, -C)	FcB1	0.53	Senegalese coasts, Keur Bamboung	(Farokhi <i>et al.</i> , 2013)
13	Axinella verrucosa	Stevensine	K1	12.61	Calvi Bay, Corsica	(Scala et al., 2010)
		Spongiacidin B	K1	3,34		
		Bromoaldisin	K1	>82.08		
		Dibromopalau'amine	K1	1.48 µg/ml		
		Bromopyrrolohomoarginin	K1	>20 µg/ml		
		Manzacidin A	K1	>20 µg/ml		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (μM)	Origin	Ref.
14	Biemna laboutei	Netamine K	not available	2.40	Salary Bay, Madagascar	(Gros et al., 2014)
		Mirabilin A	not available	20.70		
15	Biemna laboutei	Netamine O	not available	16.99	Salary Bay, Madagascar	(Gros et al., 2015)
		Netamine P	not available	32.62		
		Netamine Q	not available	8.37		
		Netamine H	not available	na		
		Netamine I	not available	na		
		Netamine N	not available	na		
		Netamine C	not available	na		
		Netamine F	not available	na		
16	Callyspongia fibrosa	24 <i>S</i> -24- methyl-cholestane 3β,6β,25-triol- 25-O-acetate	3D7	54.81	The Gulf of Mannar, Western Bay of Bengal, India	(Prakasa Rao <i>et al.</i> , 2010)
			K1	54.02		
		24 <i>S</i> -24-methyl-cholestane-3β,5α,6β,25-tetraol- 25-monoacetate	3D7	30.10		
			K1	20.54		
		24S-24-methyl-cholestane-3β,6β,8β,25-tetraol- 25-O-acetate	3D7	48.46		
			K1	44,44		
		24S-24-methyl-chelestane-3β,5α,6β,12β,25- pentaol-25-O-acetate	3D7	48,48		
			K1	47,75		
17	Clathria calla	Norbatzelladine L	FcB1	0.40	Island of Martinique	(Laville et al., 2009)
		Clathriadic acid		2.30		
18	Cymbastela cantharella	Girolline	FcB1	0.21	Caledonian sponge	(Benoit-Vical et al., 2008)
			W2	0.11		
			FcM29	0.13		
			F32	0.08		
19	Cymbastela hooperi	(1S,3S,4R,7S,8S,11S,12S,13S,15R,20R)-7- Formamido-20-isocyanoisocycloamphilectane	FCR3F86	0.58	Not available	(Wright and Lang-Unnascl 2009)
			W2	1.75		
			D6	2.34		
		(1S,3S,4R,7S,8S,11S,12S,13S,15R,20R)-7,20- Diformamidoisocycloamphilectane	FCR3F86	41.05		
		(1S*,3S*,4R*,7S*, 8S*,12S*,13S*)-7- formamidocycloamphilect-11(20)-ene	FCR3F86	na		
		(1R*,3S*,4R*,7S*,8S*,12S*,13S*)-7- formamidoamphilecta-11(20),14-diene	FCR3F86	na		
		(1S*,3S*,4R*,7S*,8S*,12S*,13S*)-7- formamidoamphilecta-11(20),15-diene	FCR3F86	na		
20	Desmapsamma anchorata	sulfated polysaccharides	3D7	66.3 μg/ml	Not available	(Marques et al., 2016)
21	Diacarnus megaspinorhabdosa	Diacarnuperoxide M	W2	4.20	Xisha Islands	(Yang et al., 2010)
			D6	5.60		
		Diacarnuperoxide N	W2	3.00		
			D6	6.60		
		(+)-2, 3, 6-epihurghaperoxide	W2	1.60		
			D6	2.20		
		(+)-2,3,6-epihurghaperoxide acid	W2	4.90		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM)	Origin	Ref.
			D6	7.30		
		(-)-muqubilin A	W2	5.60		
			D6	8.60		
		Nuapapuin A	W2	5.50		
			D6	8.10		
		Diacarperoxide A	W2	1.90		
		-	D6	2.00		
22	Fascaplysinopsis reticulata	8-oxo-tryptamine	3D7	50.52	Passe Bateau,	(Campos et al., 2019)
		(E) and (Z)-6-bromo-20-demethyl-30-N- methylaplysinopsin	3D7	24.01	Mayotte	
		6,6'-bis-(debromo)-gelliusine F	3D7	na		
		6-bromo-8,1'-dihydro-isoplysin A	3D7			
				na		
		5,6-dibromo-8,1'-dihydro-isoplysin A	3D7	na		
22		tryptamine	3D7	na	нъ	QL (1. 2010)
23	Hyattella sp.	psammaplysin G	Dd2	98% iotga 40 μM	Hervey Bay, Sponge Garden, Queensland, Australia	(Yang et al., 2010)
		psammaplysin F	Dd2	1.40		
24	Hymeniacidon sp	monamphilectine A	W2	0.60	Mona Island, Puerto Rico	(Avilés and Rodríguez, 201
25	Hyrtios cf. erecta	homofascaplysin A	K1	0.04	Fiji	(Kirsch et al., 2000)
			NF54	0.07		
		fascaplysin	K1	0.16		
			NF54	0.11		
26	Hyrtios erectus	smenotronic acid	Dd2	3.51	Chuuk Island, Federated States of Micronesia	(Ju et al., 2018)
		ilimaquinone	Dd2	2.11		
		pelorol	Dd2	0.80		
27	Ircinia sp.	tryptophol	K1	31.51	Aegean Sea, Turkey	(Orhan et al., 2010)
		4-hydroxy-3-tetraprenyl-phenylacetic acid	K1	7.77		
		demethylfurospongin-4	K1	32.23		
		dorisenone D	K1	1.03		
		11β-acetoxyspongi-12-en-16-one	K1	3.02		
28	Genus Latrunculia	discorhabdins A	D6	0.05	Aleutian Islands	(Na et al., 2010)
	(later identified as Latrunculia		W2	0.05		
	(L.) hamanni sp. nov. (Kelly	discorhabdins C	D6	2.80		
	<i>et al.</i> , 2016))		W2	2		
		dihydrodiscorhabdin C	D6	0.17		
		,	W2	0.13		
29	Lendenfeldia dendyi	Four polybromidated diphenyl ethers ^d	D6	na	Papua New Guinea	(Radwan et al., 2015)
			W2	na		
30	Mycophora sp.	Crambescidin 800	FCR3	0.24	Not available	(Lazaro et al., 2006)
			3D7	0.16		
31	Monanchora arbuscula	norbatzelladine A	FcB1	0.20	island of Martinique	(Laville et al., 2009)
		dinorbatzelladine A	FcB1	0.90	-	
		dinordehydrobatzelladine B	FcB1	0.80		
		dihomodehydrobatzelladine C	FcB1	4.50		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM)	Origin	Ref.
		batzelladine A	FcB1	0.30		
		batzelladine L	FcB1	0.30		
		ptilomycalin A	FcB1	0.10		
32	Monanchora unguiculata	Unguiculin A	3D7	12.89	Mitsio Islands, Madagascar	(Campos et al., 2019)
		Ptilomycalin E	3D7	0.35		
		Ptilomycalin F	3D7	0.23		
		Ptilomycalins G + H	3D7	0.46		
		Crambescidin 800	3D7	0.52		
		Fromiamycalin	3D7	0.24		
33	New Caledonian Sponge	Alisiaquinones A	FcMC29	8.50	the Norfolk Rise (New Caledonia)	(Desoubzdanne et al., 2008
			FcB1	7.40		
			F32	9.10		
		Alisiaquinones B	FcMC29	2.60		
			FcB1	8.40		
			F32	7.10		
		Alisiaquinones C	FcMC29	0.08		
			FcB1	0.21		
			F32	0.15		
		Alisioaquinol	FcMC29	7.90		
			FcB1	6.40		
			F32	9.90		
34	Pachastrissa nux	Kabiramide J	K1	0.31	Koh-Tao, Surat- Thani Province and Chumphon Islands National Park, Chumphon Province, Thaland	(Sirirak <i>et al.</i> , 2011)
		Kabiramide K	K1	0.39		
		Kabiramide B	K1	1.67		
		Kabiramide C	K1	4.79		
		Kabiramide D	K1	1.87		
		Kabiramide G	K1	na		
35	Pachastrissa nux	Kabiramide L	K1	2.60	Chumphon Islands National Park, Thailand	(Sirirak et al., 2011)
		Kabiramide I	K1	4.50	Koh Tao, Surat Thani Province, Thailand	
36	Petrosid Ng5 Sp5	Ingamine A	D6	0.20	Not available	(Fattorusso et al., 2010)
	· ·		W2	0.16		
		22(S)-hydroxyingamine A	D6	0.47		
			W2	0.30		
		Dihydroingenamine D	D6	0.18		
			W2	0.30		
37	Plakortis cfr. simplex	Manadoperoxide A	D10	6.88	Bunaken Marine Park of Manado, Indonesia	(Fattorusso et al., 2010)
			W2	3.74		
		Manadoperoxide B	D10	6.76		
			W2	3.69		
		Manadoperoxide C	D10	4.54		
			W2	2.33		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM)	Origin	Ref.
		Manadoperoxide D	D10	10.38		
			W2	7.93		
38	Plakortis halichondrioides	Epiplakinic acid F methyl ester	W2	0.01	Mona Island, Puerto Rico	(Jiménez-Romero <i>et al.</i> , 2010)
		Epiplakinidioic acid	W2	0.95		
		Epiplakinic acid F	W2	7.93		
		Plakortolide J	W2	na		
		Plakortolide F	W2	na		
39	Plakortis lita	Thiaplakortones A	3D7	0.05	Melville Passage, Tydeman Reef, Queensland, Australia	(Davis et al., 2012)
			Dd2	0.01		
		Thiaplakortones B	3D7	0.65		
			Dd2	0.09		
		Thiaplakortones C	3D7	0.31		
			Dd2	0.17		
		Thiaplakortones D	3D7	0.28		
			Dd2	0.16		
40	Plakortis simplex	Plakortin	D10	1.26	Berry Island (Bahamas)	(Fattorusso, 2002)
			W2	0.73		
		Dihydroplakortin	D10	1.12		
			W2	0.76		
		Plakortide E	D10	na		
			W2	na		
41	Plakortis sp.	Plakortide F	D6	1.35	Discovery Bay, Jamaica	(Gochfeld and Hamann, 2001)
			W2	1.10		
		Plakortone G	D6	15.09		
			W2	17.10		
42	Genus Pseudoceratina	Psammaplysin H	3D7	0.41	Not available	(Xu et al., 2011)
	Centus I seutocentanna	Psammaplysin G	3D7	5.22		(114 07 47., 2011)
		Psammaplysin F	3D7	1.92		
13	Pseudoceratina sp.	Ceratinadin E	K1	0.90	Okinawa, Japan	(Kurimoto et al., 2018)
	i senuoceranna sp.		FCR3	0.67	Okinuwu, Jupun	(Rumitor et u., 2010)
		Ceratinadin F	K1			
		Psammaplysin F	K1 K1	>8.16 5.16		
		i sanniapiysin i'				
44	Pseudoceratina sp.	Methyl (2,4-dibromo-3,6-dihydroxyphenyl) acetate	FCR3 FcB1	3.35 12	Rowa islands, Banks Territory (Vanuatu)	(Lebouvier et al., 2009)
45	Smenospongia aurea	6'-chloroaureol	D6	9.74	Discovery Bay, Jamaica	(Hu et al., 2002)
		Isoplysin A	D6	3.54		
		6-bromo-2'-de-N-methylaplysinopsin	D6	3.45		
		6-bromoaplysinopsin	D6	1.02		
		Makaluvamine O	D6	3.52		
		Aureol	D6	na		
		Aureol acetate	D6	na		
		2'-de-N-methylaplysinopsin	D6	na		
		N-3'-methylaplysinopsin	D6 D6			
		18-5 -meuryraprysmopsin	D0	na		

_	Organisms	Isolated compound	Pf Strain	IC ₅₀ (μM)	Origin	Ref.
46	Spongia sp.	Squalene	K1	2.82 μM	Aegean Sea, Turkey	(Orhan et al., 2010)
		Furonospinulosin-1	K1	31.53 µM		
		Furospongine 1	K1	42.42 µM		
		2-(hexaprenylmethyl)-2-methylchromenol	K1	>34.19 µM		
		Heptaprenyl-p-quinol	K1	>33.28 µM		
		12-epi-deoxoscalarin	K1	17.37 μM		
		4-hydroxy-3-octaprenylbenzoic acid	K1	2.29 µM		
		furospinulosin-2	K1	8.30 µM		
47	Spongosorites sp.	Nortopsentin A	3D7	0.46	Lucaya, Bahamas	(Alvarado et al., 2013)
48	Stylissa caribica	Stevensin	D6	4.65	Columbus Park, Jamaica	(Mohammed et al., 2006)
		oroidin	D6	3.08		
		Stylisin 1	D6	na		
		Stylisin 2	D6	na		
		Phakellistatin 13	D6	na		
		sceptrin	D6	na		
49	Stylissa cf. massa	8-isocyanato-15-formamidoamphilect-11(20)- ene	K1	8.85	Koh-Tao, Surat- Thani Province, Thailand (10°7.569' N, 99°48.665' E)	(Chanthathamrongsiri <i>et a</i> 2012)
		8-isothiocyanato-15-formamidoamphilect- 11(20)-ene	K1	8.07		
		8-isocyano-15-formamidoamphilect-11(20)-ene	K1	0.52		
		7-formamidoamphilect-11(20),15-diene	K1	na		
50	Suberea ianthelliformis	Araplysillin I	FcB1	4.5	Anuta Paina Island (Malaita)	(Mani et al., 2012)
			3D7	4.6		
		Araplysillin II	FcB1	34.2		
		Araplysillin N20-formamide	FcB1	3.6		
			3D7	7.0		
		Araplysillin IV	FcB1	27.6		
		Araplysillin V	FcB1	50.5		
		Araplysillin VI	FcB1	37.4		
	Suberea ianthelliformis	Aerophobin I	FcB1	59.0	New Georgia Island	(Mani et al., 2012)
		Aerophobin II	FcB1	24.9		
			3D7	19.9		
		Purealidin Q	FcB1	3.6		
		Araplysillin N20-hydroxyformamide	FcB1	5.0		
			3D7	4.1		
	Suberea ianthelliformis	Aerothionin	FcB1	3.4	North West of Nggela Island	(Mani et al., 2012)
			3D7	4.2		
		Homoaerothionin	FcB1	2.8		
			3D7	4.0		
		11,19-Dideoxyfistularin 3	FcB1	2.1		
			3D7	0.9		
		11-Hydroxyfistularin 3	FcB1	2.1		
			3D7	2.6		
		Aplysinone D	FcB1	1.0		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM)	Origin	Ref.
51	Verongula rigida	Purealidin B	NF54	23.2% iotga 5 μM	Urabá Gulf, Caribbean Sea, Colombia (8°40'14"N, 77°21'28"W)	(Galeano <i>et al.</i> , 2011)
		11-hydroxyaerothionin	NF54	8.0% iotga 5 μM		
		Aeroplysinin	NF54	35.3% iotga 5 μM		
		Dihydroxyaerothionin	NF54	7.9% iotga 5 μM		
		Purealidin R	NF54	7.1% iotga 5 μM		
		3,5-dibromo-N,N,N-trimethyltyraminium	NF54	na		
		3,5-dibromo-N,N,N,O-tetramethyltyraminium	NF54	na		
		19-deoxyfistularin 3	NF54	na		
52	Xestospongia exigua	Araguspongine C	D6	1.4	Bayadha, Saudi Arabian Red Sea coast	(Orabi <i>et al.</i> , 2002)
			W2	0.58		
		(+)- Araguspongine K	D6	na		
			W2	na		
		(+)- Araguspongine L	D6	na		
			W2	na		
53	Xestospongia sp.	Kaimanol	3D7	0.36	Kaimana, West Papua, Indonesia	(Murtihapsari et al., 2019
		Saringosterol	3D7	2.50×10^{-4}		
54	Xestospongia sp.	Xestoquinone	FcB1	3	Malvoror reef, Vanuatu	(Laurent et al., 2006)
55	genus Xestospongia	Halenaquinone	FcB1	>30	South Pacific	(Longeon et al., 2010)
			3D7	>30		
		3-Ketoadociaquinone A	FcB1	1.08		
			3D7	1.67		
		3-Ketoadociaquinone B	FcB1	3.89		
			3D7	4.12		
		Tetrahydrohalenaquinone A	FcB1	>29		
			3D7	>29		
		Tetrahydrohalenaquinone B	FcB1	>29		
			3D7	>29		
		Halenaquinol sulfate	FcB1	>24		
			3D7	>24		
		Xestosaprol C methylacetal	FcB1	>21		
			3D7	>21		
		Orhalquinone	FcB1	9.22		
			3D7	10.94		
56	<i>Zyzzya</i> sp.	Tsitsikammamine C	3D7	0.01	Rodda Reef, Queensland, Australia	(Davis et al., 2012)
			Dd2	0.02		
		makaluvamines J	3D7	0.02		
			Dd2	0.02		
		makaluvamines G	3D7	0.04		
			Dd2	0.04		
		makaluvamines L	3D7	0.04		

No	Organisms	Isolated compound	Pf Strain	IC ₅₀ (µM) Origin	Ref.
			Dd2	0.02	
		makaluvamines K	3D7	0.40	
			Dd2	0.30	
		Damirone A	3D7	1.88	
			Dd2	0.36	
		Damirone B	3D7	12.25	
			Dd2	3.80	





Figure 1. Distribution of conducted studies about marine sponge metabolite exploration for *in vitro* antiplasmodium.

CLASSIFICATION OF ANTIPLASMODIAL ACTIVITY OF ISOLATED COMPOUND FROM SPONGES

In this review, we give an overview of the bioactive metabolites recently isolated from marine sponges that have shown activity in in vitro study against P. falciparum. To compare the IC₅₀ values, the units in μ g/ml and nM were converted to μ M. All the isolated compounds were then classified based on their IC₅₀ values by following the definition of Batista et al. (2009), who grouped compounds into potent activity: $IC_{50} < 1 \mu M$, good activity: IC₅₀ of 1–20 µM, moderate activity: IC₅₀ of 20–100 µM, low activity: IC₅₀ of 100–200 μ M, and inactive: IC₅₀ >200 μ M (Batista et al., 2009). To be noted, the mechanism of the in vitro continuous cultures of *P. falciparum* approach is only related to the inhibition of growth in erythrocytic stages of the parasite (Chin et al., 1979). Consequently, this IC₅₀-based classification would exclude compounds that may have other specific mechanism of action. It would be wise to re-evaluate "not active compounds" with other assay or holistic approach such as the reverse pharmacology technique (Simoes-Pires et al., 2014).

As shown in Figure 2, among observed bioactive metabolites, there were 57 different compounds that have potent activity, 101 with good activity, and 26 compounds with moderate activity against various strains of *P. falciparum*. Some of the compounds could not be classified because, in the highest tested concentration, their activity was low or inactive and some reports use inhibition concentration instead of IC₅₀, making it incomparable. In regard to the dependency of IC₅₀ to plasmodium



Figure 2. Classification of the isolated compounds activity according to their IC_{50} values.

strains, it seems that antiplasmodial activity of some isolated compounds did not depend on chloroquine/drug sensitivity of the strain (Fattorusso *et al.*, 2010; Longeon *et al.*, 2010; Mani *et al.*, 2012).

The class of compounds which exhibit potent antiplasmodial activity includes manzamine alkaloid (Rao *et al.*, 2004; 2006; Samoylenko *et al.*, 2009), guanidine alkaloids (Campos *et al.*, 2017; Laville *et al.*,2009), bispyrroloiminoquinone alkaloid (Davis *et al.*, 2012), pyrroloiminoquinone alkaloids (Na *et al.*, 2010), ingamine alkaloids (Ilias *et al.*, 2012), sesquiterpenoids (Angerhofer *et al.*, 1992), diterpene formamides (Wright and Lang-Unnasch, 2009), aminoimidazole (Benoit-Vical *et al.*, 2008), β -galactosyl ceramides (Farokhi *et al.*, 2013), β -lactam (Avilés and Rodríguez, 2010), meroterpene (Desoubzdanne *et al.*, 2008), trisoxazole macrolides (Sirirak *et al.*, 2011), peroxides, thiazine alkaloids (Davis *et al.*, 2012), bromotyrosine alkaloids (Kurimoto *et al.*, 2018; Xu *et al.*, 2011), and sterols (Murtihapsari *et al.*, 2019).

FUNCTIONAL GROUP IN POTENT ANTIPLASMODIAL ACTIVITY

Some marine isonitriles show various biological activities such as antimalarial, antitubercular, antifouling, and antiplasmodial effect. Marine isonitriles differ from terrestrial isonitriles in terms of their biosynthetic pathways. Most of the marine compounds containing isonitrile were derived from terpenoid, whereas terrestrial isonitriles originate from α -amino acids (Emsermann

Table 2. List of isolated compounds with potent antiplasmodial activity basedon IC_{so} measurement.

1Axisonitrile 3D6 and W22 $(+)$ -8-hydroxymanzamine AD6 and W23 $(+)$ -manzamine AD6 and W24 $(+)$ -8-hydroxymanzamine A hydrochlorideD6 and W25 $(+)$ -manzamine A hydrochlorideD6 and W26Manzamine AD6 and W27Manzamine AD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(IS,3S,4R,7S,8S,1IS,12S,13S,1SR,20R)-7. Formanido-20-isseyanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415Fascaplysin AK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin F3D728Ptilomycalin F3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide KK132Iabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W2 <th>No.</th> <th>Isolated Compound</th> <th>P. falciparum strain</th>	No.	Isolated Compound	P. falciparum strain
C)manzamine AD6 and W24(+)-8-hydroxymanzamine A hydrochlorideD6 and W25(+)-manzamine A hydrochlorideD6 and W26Manzamine AD6 and W27Manzamine A N-oxideD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(15,35,47,58,51,15,125,135,158,20R)-7- Formanido-20-isocyanoisocycloamphilectameFCB313Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdin S AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinordatzelladine AFcB122Dinordatzelladine AFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin G + H3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromanine AD6 and W236Epiplakinicici acidW237Epiplakinici acid F methyl esterW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone C3D7 and Dd230Dihydroplakotrine B3D7 and Dd	1	Axisonitrile 3	D6 and W2
11114(+)-manzamine A hydrochlorideD6 and W26Manzamine AD6 and W27Manzamine AD68Manzamine A N-oxideD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB112(IS,3S,4R,7S,8S,1IS,12S,15S,20R)-7- Formanido-20-isocyanoisocycloamphilectaneFCB113Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolD217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinordehydrobatzelladine BFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D729Fromiamycalin G + H3D730Alisiaquinone CFcM22; FcB1; and F3231Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroegotamine DD6 and W236Epiplakiniciacid F methyl esterW237Epiplakortine A3D7 and Dd238Thiaplakortone D <td>2</td> <td>(+)-8-hydroxymanzamine A</td> <td>D6 and W2</td>	2	(+)-8-hydroxymanzamine A	D6 and W2
S(+)-manzamine A hydrochlorideD6 and W26Manzamine A M-oxideD6 and W27Manzamine A N-oxideD6 and W29AxidjiferosidesFeB110Norbatzelladine LFeB111GirollineFeB112(IS,3S,4R,7S,8S,1IS,12S,13S,1SR,20R)-7- Formamido-20-isoeyanoisoeycloamphilectameFCB3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolD6 and W217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFeB121Dinordebtzelladine AFeB122Dinordebtydrobatzelladine BFeB123Batzelladine AFeB124Batzelladine IFeB125Ptilomycalin F3D728Ptilomycalin F3D729Fromiamycalin3D730Alisiaquinone CFeMC29; FeB1; and F3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinicia cidW237Epiplakinicia cidW238Thiaplakortone A3D7 and Dd239Thiaplakortone D3D7 and Dd239Thiaplakortone D3D7 and Dd230	3	(+)-manzamine A	D6 and W2
6Manzamine AD6 and W27Manzamine A N-oxideD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(IS,3S,4R,7S,8S,1IS,12S,13S,15R,20R)-7- Formamido-20-isocyanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415Fascaplysin AK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 80FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine FSD725Ptilomycalin F3D726Ptilomycalin G + H3D727Ptilomycalin G + H3D728Ptilomycalin G + HD6 and W229FromiamycalinM230Alisiaquinone CFc3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroegotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakindioic acidW238Thiaplakortone C3D7 and Dd239Thiaplakortone C3D7 and Dd230Th	4	(+)-8-hydroxymanzamine A hydrochloride	D6 and W2
7Manzamine YDo8Manzamine A N-oxideD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(15,35,4R,75,38,115,125,135,15R,20R)-7 Formamido-20-isocyanoisocycloamphilectameW213Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine IFcB124Batzelladine IFcB125Pitlomycalin F3D726Pitlomycalin G + H3D727Pitlomycalin G + H3D728Pitlomycalin G + H3D730Alisiaquinone CFcB1; and F3231Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroegotamine DD6 and W236Epiplakinidioic acidW237Epiplakindioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone D3D7 and Dd230Thiapla	5	(+)-manzamine A hydrochloride	D6 and W2
8Manzamine A N-oxideD6 and W29AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(IS,3S,4R,7S,8S,1IS,12S,1SR,20R)-7. Formamido-20-isocycloamphilectaneFcB1; W2; FcM29; and F3213Monamphilectine AW214Homofascaplysin AK1 and NF5415Fascaplysin AK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbetzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KM23422(S)-hydroxyingamine AD6 and W235Dihydroegotamine DD6 and W236Epiplakinici acid F methyl esterW237Epiplakindioic acid3D7 and Dd238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone C3D7 and Dd230Thiaplakortone A	6	Manzamine A	D6 and W2
9AxidjiferosidesFcB110Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(1S,3S,4R,7S,8S,11S,12S,13S,15R,20R)-7. Fornamido-20-isocycanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415Fascaplysin AK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine A3D6 and W235Dihydrocegotamine DD6 and W236Epiplakinici acid T methyl esterW237Epiplakinicia caid T methyl esterW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone C3D7 and Dd234Thiaplakortone C<	7	Manzamine Y	D6
Norbatzelladine LFcB111GirollineFcB1; W2; FcM29; and F3212(IS,3S,4R,7S,8S,1IS,12S,1SR,20R)-7. Fornamido-20-isocyanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D729FromiamycalinJD730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroegtoamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakindioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone A3D7 and Dd230Thiaplakortone B3D7 and Dd23422(S)-hydroxyingamine A3D7 and Dd2	8	Manzamine A N-oxide	D6 and W2
11GirollineFcB1; W2; FcM29; and F3212(15,35,48,75,85,115,125,135,15R,20R)-7- Formamido-20-isocyanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin G + H3D727Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinici acid F methyl esterW238Thiaplakortone A3D7 and Dd239Thiaplakortone D3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone C3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D7 <td>9</td> <td>Axidjiferosides</td> <td>FcB1</td>	9	Axidjiferosides	FcB1
and F3212(1S,3S,4R,7S,8S,11S,12S,15R,20R)-7. Formamide-20-isocyanoisocycloamphilectaneFCR3F8613Monamphilectine AW214Homofascaplysin AK1 and NF5415Fascaplysin AK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinordehydrobatzelladine BFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFceNC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinicio acid3D7 and Dd238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd234DihydroplakortinW235DihydroplakortinW236Epiplakinidoic acid3D7 and Dd239Thiaplakortone D3D7 and Dd240Thiaplakortone D <td>10</td> <td>Norbatzelladine L</td> <td>FcB1</td>	10	Norbatzelladine L	FcB1
Formamido-20-isocyanoisocycloamphilectane13Monamphilectine AW214Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinic acid F methyl esterW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone C3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	11	Girolline	
14Homofascaplysin AK1 and NF5415FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine AFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd239Thiaplakortone D3D7 and Dd230Thiaplakortone C3D7 and Dd231Kabtrini di F methyl esterW234Piplakinidioic acidW235DihydroplakortinW236Fiplakinidioic acid3D7 and Dd237Epiplakinidioic acid3D7 and Dd238Thiaplakortone D3D7 and Dd	12		FCR3F86
15FascaplysinK1 and NF5416PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin F3D726Ptilomycalin F3D727Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinidioic acidW237Epiplakinidioic acid3D7 and Dd239Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd239Thiaplakortone M3D7 and Dd241Thiaplakortone M3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	13	Monamphilectine A	W2
16PelorolDd217Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinidioic acidW237Epiplakinidioic acid3D7 and Dd239Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Paammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	14	Homofascaplysin A	K1 and NF54
17Discorhabdins AD6 and W218Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakindioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone C3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	15	Fascaplysin	K1 and NF54
18Dihydrodiscorhabdin CD6 and W219Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroogtamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinicio acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	16	Pelorol	Dd2
19Crambescidin 800FCR3 and 3D720Norbatzelladine AFcB121Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F2231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinicio acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	17	Discorhabdins A	D6 and W2
10Norbatzelladine AFcB120Norbatzelladine AFcB121Dinordehydrobatzelladine BFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakindioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	18	Dihydrodiscorhabdin C	D6 and W2
21Dinorbatzelladine AFcB122Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinici acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	19	Crambescidin 800	FCR3 and 3D7
22Dinordehydrobatzelladine BFcB123Batzelladine AFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin E3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	20	Norbatzelladine A	FcB1
12Entroting Definition in23Batzelladine IFcB124Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin F3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinici acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	21	Dinorbatzelladine A	FcB1
24Batzelladine LFcB125Ptilomycalin AFcB126Ptilomycalin E3D727Ptilomycalin G + H3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinicia caidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	22	Dinordehydrobatzelladine B	FcB1
25Ptilomycalin AFcB126Ptilomycalin E3D727Ptilomycalin F3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidoic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	23	Batzelladine A	FcB1
26Ptilomycalin E3D727Ptilomycalin F3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone M3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	24	Batzelladine L	FcB1
27Ptilomycalin F3D728Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	25	Ptilomycalin A	FcB1
28Ptilomycalin G + H3D729Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone C3D7 and Dd240Thiaplakortone D3D7 and Dd241Thiaplakortone M3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	26	Ptilomycalin E	3D7
29Fromiamycalin3D730Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	27	Ptilomycalin F	3D7
30Alisiaquinone CFcMC29; FcB1; and F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	28	Ptilomycalin G + H	3D7
F3231Kabiramide JK132Kabiramide KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	29	Fromiamycalin	3D7
32Kabinande FKI32Kabinande KK133Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	30	Alisiaquinone C	
33Ingamine AD6 and W23422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	31	Kabiramide J	K1
3422(S)-hydroxyingamine AD6 and W235Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	32	Kabiramide K	K1
35Dihydroergotamine DD6 and W236Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	33	Ingamine A	D6 and W2
36Epiplakinic acid F methyl esterW237Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	34	22(S)-hydroxyingamine A	D6 and W2
37Epiplakinidioic acidW238Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	35	Dihydroergotamine D	D6 and W2
38Thiaplakortone A3D7 and Dd239Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	36	Epiplakinic acid F methyl ester	W2
39Thiaplakortone B3D7 and Dd240Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	37	Epiplakinidioic acid	W2
40Thiaplakortone C3D7 and Dd241Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	38	Thiaplakortone A	3D7 and Dd2
41Thiaplakortone D3D7 and Dd242PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	39	Thiaplakortone B	3D7 and Dd2
42PlakortinW243DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	40	Thiaplakortone C	3D7 and Dd2
43DihydroplakortinW244Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	41	Thiaplakortone D	3D7 and Dd2
44Psammaplysin H3D745Ceratinadin EFCR346Nortopsentin A3D7	42	Plakortin	W2
45Ceratinadin EFCR346Nortopsentin A3D7	43	Dihydroplakortin	W2
46 Nortopsentin A 3D7	44	Psammaplysin H	3D7
	45	Ceratinadin E	FCR3
47 8-isocyano-15-formamidoamphilect-11(20)-ene K1	46	Nortopsentin A	3D7
	47	8-isocyano-15-formamidoamphilect-11(20)-ene	K1

No.	Isolated Compound	P. falciparum strain
48	11,19-Dideoxyfistularin 3	3D7
49	Araguspongine C	W2
50	Kaimanol	3D7
51	Saringosterol	3D7
52	Tsitsikammamine C	3D7 and Dd2
53	Makaluvamine J	3D7 and Dd2
54	Makaluvamine G	3D7 and Dd2
55	Makaluvamine L	3D7 and Dd2
56	Makaluvamine K	3D7 and Dd2
57	Damirone A	Dd2

et al., 2016). Axisonitrile-3 (1) is a sesquiterpene derived from chloroform fraction of sponge *Acanthella klethra* containing isonitrile group which appears to be crucial for activity since the corresponding isothiocyanate derivative compound 2 (moderate activity) is less active than 1 (potent activity) (Angerhofer *et al.*, 1992). The eudesmane compounds 3 and 4 which contain isothiocyanate still showed good antiplasmodial activity, whereas the reversal of the stereochemical configuration between 4 and 5 exhibits a significant change on their antiplasmodial effect (see Figure 3).

The manzamines are a group of marine alkaloids characterized by a fused and bridged tetra- or pentacyclic ring system attached to a β -carboline moiety. Since manzamine was isolated in different genus of sponges, it is thought that manzamine is actually produced by associated microorganism. An interesting review had been done by Fattorusso and Taglialatela-Scafati (2009) who described the key role of the eight member rings as well as other functional groups that affect the antiplasmodial activity of manzamines; therefore, we will not discuss it in this review.

A mixture of new glycosphingolipids named axidjiferoside A (6), axidjiferoside B (7), and axidjiferoside C (8) shows a potent antiplasmodial activity (Figure 3). Compounds 6, 7, and 8 were isolated from Senegal marine sponge *Axinyssa djiferi* (Farokhi *et al.*, 2013). These compounds contain sphingolipid structure which are found in ceramide analogs, PPMP (d,1-threo-1-phenyl-2-palmitoylamino-3-morpholino-1-propanol), and PDMP (1-phenyl-2-decanoylamino-3-morpholino-1-propanol). These analogs are known to inhibit the parasite sphingomyelin synthase activity and block parasite development by preventing the formation of the tubovesicular network that extends from the parasitophorous vacuole to the red cell membrane and delivers essential extracellular nutrients to the parasite (Labaied *et al.*, 2004; Zhang *et al.*, 2010).

Bioactive guanidine alkaloids including norbatzelladine A (9), dinorbatzelladine A (10), batzelladine A (11), dinordehydrobatzelladine B (12), norbatzelladine L (13), and batzelladine L (14) are potent against the growth of *P. falciparum*. The aromatization in the tricyclic core of 11 (compared to 9 and 8) did not change the antimalarial activity. Batzelladine A, with one bicyclic and one tricyclic guanidine core, has similar properties with 9, 13, and 14 in terms of the activity against *P. falciparum* strain FcB1, where 13 and 14 have two tricyclic guanidine cores. The reduction of bicyclic core in dihomodehydrobatzelladine C seems to affect its activity to be less active than 9–12 (Figure 3).



Figure 3. Structure of antimalarial compounds (Angerhofer *et al.*, 1992; Benoit-Vical *et al.*, 2008; Farokhi *et al.*, 2013; Mudianta *et al.*, 2012; Wright and Lang-Unnasch 2009; Xu *et al.*, 2011).

Girolline (15), 2-aminoimidazole derivative, isolated from *Cymbastela cantharella* showed a potent activity against *P. falciparum* strains, whereas its analogs 5-deazathiogirollines (16 and 17) were considered to be inactive (Benoit-Vical *et al.*, 2008). This indicates that imidazole ring in 15 plays an important role in the antiplasmodial activity.

Sponge Cymbastela hooperi sp. nov. described by Soest et al. (1996) produces a plethora of chemical compounds structurally related to diterpene isonitrile derivatives which exhibit significant in vitro antimalarial activity. (1S, 3S,4R,7S,8S,11S,12S,13S,15R,20R)-7-Formamido-20isocyanoisocycloamphilectane (18), (1S,3S,4R,7S,8S,11S,12S,1 3S,15R,20R)-7,20-Diformamidoisocycloamphilectane (19), and (1S*,3S*,4R*,7S*,8S*,12S*,13S*)-7-Formamidocycloamphilect-11(20)-ene (20) were new diterpene formamides which were isolated from C. hooperi (Figure 3). Compound 18 is a unique molecule since it contains both formamide and isonitrile functionalities where such a feature is rarely found in natural product. Based on its IC50 against P. falciparum FCR3F86, this substituent is classified into potent (Wright and Lang-Unnasch, 2009). The lack of isonitrile in the structure of 19 decreases the activity to be moderate. This finding is supported by the activity of compound 1 that possesses isonitrile too (Angerhofer et al., 1992).

Psammaplysin H (21) derived from sponge genus *Pseudoceratina* is also included in the potent activity group against *P. falciparum* 3D7 with IC₅₀ 0.41 μ M. This activity is more likely caused by the presence of quaternary amine in the R group at C-20 (see Figure 3). However, the secondary amine at the same position in psammaplysin F (22) reduced antimalarial activity 4-fold lower than compound 21. In addition, when the alkyl amine is substituted with a urea at C-20 in Psammaplysin G (23), the activity decreased to have IC₅₀ 5.99 μ M (Xu *et al.*, 2011). Consistently, the loss of amine substituent in psammaplysin K (24) dispelled the antiplasmodial activity (Mudianta *et al.*, 2012).

CONCLUSION

Data presented in the review indicate that marine sponges could be used as sources for lead compounds in drug discovery program including the development of non-resistance antimalarial drugs in this case. The summarized "potent" isolated compounds highlight the most promising candidates which include manzamine alkaloids, guanidine alkaloids, bispyrroloiminoquinone alkaloid, pyrroloiminoquinone alkaloids, ingamine alkaloids, sesquiterpenoids, diterpene formamides, aminoimidazole, β -galactosyl ceramides, β -lactam, meroterpene, trisoxazole macrolides. peroxides, thiazine alkaloids. bromotyrosine alkaloids, and sterols. A holistic approach for their pharmacological evaluation is still needed since in vitro P. falciparum assay could only evaluate a specific mechanism of action for antiplasmodium. To reproduce the compounds for their further evaluation, the possibility of bioengineering or/and bacterial fermentation could be worth.

ACKNOWLEDGMENT

The author would like to acknowledge the funding support from UGM No: 3040/UN1/DITLIT/DIT-LIT/LT/2019.

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How to cite this article:

Hikmawan BD, Wahyuono S, Setyowati EP. Marine sponge compounds with antiplasmodial properties: Focus on *in vitro* study against *Plasmodium falciparum*. J Appl Pharm Sci, 2020; 10(05):142–157.