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REVIEW

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Natural products as antivibrio agents: insight into the chemistry and biological activity

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Vibriosis causes serious problems and economic loss in aquaculture and human health. Investigating natural products as antivibrio agents has gained more attention to combat vibriosis. The present review highlights the chemical diversity of antivibrio isolated from bacteria, fungi, plants, and marine organisms. Based on the study covering the literature from 1985–2021, the chemical diversity ranges from alkaloids, terpenoids, polyketides, sterols, and peptides. The mechanisms of action are included inhibiting growth, interfering with biofilm formation, and disrupting of quorum sensing. Relevant summaries focusing on the source organisms and the associated bioactivity of different chemical classes are also provided. Further research on in vivo studies, toxicity, and clinical is required for the application in aquaculture and human health. REVIEW **Notice of the chemical products as antivibrio agents: insight int

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1. Introduction

The genus Vibrio is Gram-negative, curved-rod shape bacteria, halophilic, fermentative, motile by polar flagella, catalase, and oxidase-positive. The genus inhabits aquatic environment, freshwater, water column, sediment, and is associated with marine organisms.^{1,2} Vibrio spp. play roles as nutrient cyclers in aquatic ecosystems, take up organic material, produce polyunsaturated fatty acids to the aquatic food web, and degrade chitin.³ These groups of bacteria are responsible for several serious infections and opportunistic pathogens to aquatic animals and humans. $1,4,5$

Studies about the effects of increasing sea surface temperature on the biology and ecology of Vibrio showed that there are correlations between the escalation of the emergence of Vibrio infections and global warming. Climate change induces global warming and as a result, the rising sea surface temperature corresponds to the number and distribution of Vibrio as reported in many places worldwide. Salinity less than 25 ppt contributes to Vibrio prevalence and infection in the marine system.⁵⁻⁸

The term vibriosis is used to refer to infections by the member family of Vibrionaceae both in aquatic animals and humans.⁹ Vibriosis is one of the primary problems in aquaculture that causes severe economic losses and large-scale mortality of shrimp, fish, and shellfish.¹⁰ Comprehensive reviews are available focusing on vibriosis in fish, $11-13$ shrimps, 14 crustaceans,¹⁵⁻¹⁷ and mollusks.¹⁸

More than a hundred Vibrio species have been identified and caused infections in humans. About 14 species of Vibrio reported as causative agents of human vibriosis cause foodborne and nonfoodborne Vibrio infections such as V. cholerae, V. parahaemolyticus, V. alginolyticus, and V. vulnificus.^{9,19} Vibrio spp. infect humans worldwide and is responsible for gastroenteritis, septicemia, and invasive skin and soft tissue infection (SSTI).^{9,20} Non-foodborne Vibrio infections, caused by V. vulnificus, V. alginolyticus, and V. parahaemolyticus are fatal and often leads to the amputation or death of immunocompromised patients suffering from liver disease, alcohol abuse, or diabetes.²⁰⁻²²

A single or combinational antibiotic is the treatment for curative against vibriosis both in aquatic animals and humans.²⁰ Most *Vibrio* spp. are susceptible to most antibiotics for animals or humans. Overuse and unregulated antibiotic used in aquaculture are contributing to growing problems and concerns in antimicrobial resistance that impacts human health. Antimicrobial resistance may reduce the effectiveness of treatment options for fish and human health management.^{23,24} Multiple-antibiotic resistance of V. vulnificus and V. parahaemolyticus were reported in countries such as the United States, Italy, Brazil, Philippines, Malaysia, Indonesia, Thailand, China, India, Iran, South Africa, and Australia.²⁴–³⁰

Antibiotic resistance and the restricted choices of available antivibrio agents are the reasons for searching natural products as new antivibrio agents. The increase in the emergence of antibiotic-resistant bacterial pathogens, including Vibrio spp. is a major public health concern. Therefore, it has intensified the interest in research on the search for effective alternatives to cope with the issue of antibiotic-resistant bacteria. Attempts have been done on screening, isolation, and structure determination of antivibrio compounds from natural products. This review intends to deliver the exploration of natural products for new antivibrio compounds.

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2. Targets for antivibrio

Antibiotic resistance is becoming an important issue in the world of medicine. Newly developed antibiotics also starting to lose their effectiveness against some bacterial strains. As a result, it is critical to look for novel antimicrobial agents that are both effective against resistant microbes and long-lasting.

Quorum sensing (QS) is a small diffusible signaling molecule that trigger the expression of multiple genes that govern a wide range of activities including bioluminescence, virulence control, sporulation, host colonization, biofilm development, defense against competition, and environmental adaptability. Vibrio fischeri, V. harveyi, V. cholerae, V. anguillarum, and V. vulnificus use QS to regulate their pathogenicity. 31 Biofilm is a complex structure of microbiome having different bacterial colonies or single type of cells in a group; adhere to the surface that are embedded in a membrane structure of the extracellular polymeric substances (EPS) composed of eDNA, proteins and polysaccharides. The matrix complex are attached to the biotic or abiotic surface, showed high resistance to antibiotics.^{32,33} Biofilms formation is the key factor for accelerating *Vibrio* spp. to grow and survive by providing access to nutrients, protecting from the host immune system, defending from the predator, and antimicrobial compounds. Studies showed that biofilm is important for survival, virulence and stress resistance of Vibrio sp.^{34,35} The formation and maintenance of biofilms, as well as their resistance to antimicrobials and the host's innate immune system, are controlled by OS-regulated gene expression.³⁵⁻³⁷ **PRC Advances** Weekly Reverse Article on 2022. Transparent issues in the distance of the common and interactional constraints are allow the common and the distance article in the common and the common and the common and t

In the aquaculture system, QS regulates virulence factors and the formation of biofilm. Thus, disruption of QS is a potential strategy for preventing disease in aquaculture systems. Quorum sensing inhibitors (QSIs) or quorum quenchers inhibit both the expression of virulence-associated genes and attenuate the virulence of aquaculture pathogens.³² Quorum sensing plays a role in the formation of biofilms. Thus, fighting Vibrio spp. by interrupting quorum sensing and biofilms formation are the right strategies to combat vibriosis.38,39 Inhibiting growth, interrupting quorum sensing, and interfering biofilms formation are targets for antagonistic effects in searching for antivibrio.

3. Antivibrio from bacteria

3.1. Actinobacteria

Actinobacteria are important assets for microbial natural products with therapeutic properties for medicinal, agricultural, veterinary, and aquaculture applications including chloramphenicol, tetracycline, erythromycin, rifamycin, rapamycin, vancomycin, bleomycin, and avermectin. Actinobacteria are known to produce 70% of the antibiotics used today.^{40,41}

Screening have been carried out to obtain isolates that produce antivibrio compounds. Actinobacteria mainly Streptomyces spp. from different sources were tested for the antagonistic effect against Vibrio spp.⁴²–⁴⁷ A comprehensive review showed a list of 128 strains of Streptomyces isolated from terrestrial and marine environments that are active against Vibrio spp.⁴⁸ Most of the studies have focused on the

preliminary screening bioactivity of crude extract fermentation. To date, only a limited number of structure elucidations and identified the bioactive compounds that displayed potent antibacterial activity against Vibrio spp. Herein, we collected data on antivibrio compounds isolated from Actinobacteria presented in Fig. 1 and Table 1.

Brevibacterium casei MSI04 associated with a marine sponge Dendrilla nigra produces poly-hydroxy butyrate with the activity as antiadhesive. The inhibition activity was tested again on pathogen Vibrio spp. from shrimp aquaculture. The compound inhibited V. vulnificus and V. fischeri (96%), V. parahaemolyticus and V. alginolyticus (92%), and V. harveyi (88%).⁴⁹

Actinobacteria produce wide type of antibiotics such as nanaomycins, munumbicins and guadinomine active against Vibrio spp. Nanaomycins are quinone antibiotics produced by Streptomyces rosa var notoensis OS-3966. Nanaomycin A (1) showed bioactivity against V. parahaemolyticus K-1 and V. alginolyticus 138–2 at MIC 3.1 µg mL⁻¹ and 6.3 µg mL⁻¹, respectively Management D₁(2) has the greater activity against V₁ tively. Nanaomycin D (2) has the greater activity against V. parahaemolyticus K-1 and V. alginolyticus 138–2 at MIC less than 0.05 μ g mL⁻¹. The mechanism of action is inhibiting biosynthesis of protein DNA BNA and cellurall pentideclygan⁵⁶ thesis of protein, DNA, RNA, and cell-wall peptidoglycan.⁵⁰ Munumbicins are antibiotic peptides with broad spectrum activity against Gram-positive and negative bacteria. The peptides were isolated from endophytic Streptomyces NRRL 3052. Munumbicins A-D were tested against V. fischeri PIC 345 at a concentration of 10 µg. Munumbicin A was inactive, while munumbicins B (3), C, and D showed zone inhibition of 16, 9, and 12 cm, respectively.⁵¹ Guadinomine B (4) is an antibiotic peptide produced by Streptomyces sp. K01-0509. The compound works as an antivirulence at IC_{50} 14 nM with a novel mechanism of action as an inhibitor of the type III secretion system (TTSS) of Gram-negative bacteria including Vibrio sp.^{52,53}

Streptomyces atrovirens PK288-21 associated with seaweed Undaria pinnatifida produces two compounds 2-hydroxy-5-(3methylbut-2-enyl) benzal-dehyde (5) and 2-hepta-1,5-dienyl-3,6-dihydroxy-5-(3-methylbut-2-enyl) benzaldehyde (6) were isolated from. Compound (5) inhibited V. anguillarum and V. harveyi at MIC 65 and 20 µg mL^{-1} , respectively. While
compound (6) was active expiret *V* anguillary and *V* harvey is compound (6) was active against V. anguillarum and V. harveyi at MIC 65 and 32 μ g mL⁻¹, respectively.⁵⁴
High throughout expening of exident

High throughput screening of crude extract of marine Actinobacteria was examined targeting peptide deformylase (PDF) of V. anguillarum that catalyzes the removal of N-formyl group from N-terminal methionine following translation in prokaryotes. Extraction of fermentation broth of Streptomyces sp. NHF 165 yielded Actionin (7) that inhibited peptide deformylase (PDF) of *V. anguillarum* at IC $_{50}$ was 2.85 μ M.⁵⁵

Streptomyces leeuweenhoekii strain C34 isolated from the Chilean hyper-arid Atacama Desert soil produces a new type of antibiotic ansamycin which is active as antivibrio. Using the OSMAC approach led to isolating new 22-membered macro lactone-type polyketides called Chaxalactin A-C (8–10). Chaxalactins A (8) , B (9) , and C (10) inhibited *V. parahaemolyticus* at MIC 12.5; 20; and 12.5 μ g mL⁻¹, respectively.^{56,57} Streptomyces sp. SCSIO 01689 produces antivibrio compounds pyranosesquiterpene (11) and cyclic peptides Cyclo(D)-Pro-(D)-Ile

(12), Cyclo(D)-Pro-(D)-Leu (13), and Cyclo(D)-trans-4-OH-Pro- (D)-Phe (14). The compound 11 inhibited V. anguillarum at MIC at > 100 μ g mL⁻¹ while the cyclic peptides showed potency at concentrations of 0.05, 0.04, and 0.07 μ g mL⁻¹ for 12, 13, and 14, respectively.⁴⁸

3.2. Pseudoalteromonas

The genus Pseudoalteromonas is Gram-negative bacteria, heterotrophic, aerobic, and belongs to the γ -Proteobacteria class. This genus attracts attention due to its wide array of metabolites and ecological role in the ocean. The metabolites of Pseudoalteromonas have bioactivity including antimicrobial agents.^{58,59} Antivibrio compounds isolated from Pseudoalteromonas spp. are presented in Fig. 2 and Table 2.

Pseudoalteromonas A1-J11 from the coastal Kagoshima Bay, Japan produced bioactive quilinolinol derivatives 2-n-pentyl-4 quinolinol (15). Disk diffusion assay of the compounds was conducted against V. harveyi ATCC 14126, V. harveyi ATCC

35084, V. alginolyticus ATCC 17749, Vibrio sp. 9M-P5-1, V. fischeri VF-74, V. parahaemolyticus IFO 12711. Based on the bioassay compound 15 was active against V. harveyi ATCC 14126, V. harveyi ATCC 35084, and V. fischeri VF-74 at a concentration of $10 \mu g.^{60}$

Crude extract of Pseudomonas haloplanktis INH from scallop hatchery was tested against V. ordalii ATCC 33509, V. algiynolyticus ATCC 17749, V. anguillarum IFO 13266, and V. anguillarum (VAR). The inhibition of V. ordalii ATCC 33509 was observed at a concentration of 1 mg mL $^{-1}$. Antibacterial compounds from the ethyl acetate extract were identified as isovaleric acid (16) and 2-methyl butyric acid (17) .⁶¹

Pseudoalteromonas strain J010 associated with the surface of the crustose coralline alga Neogoniolithon fosliei, produced bioactive compounds antivibrio tetrabromopyrrole (18), 4'- $((3,4,5\text{-tribromo-1}H\text{-pyrrol-2-yl)methyl)phenol (19), and kor$ ormicins G-I (20-22) and K (23). The compounds were tested at a concentration of 200 μ g mL⁻¹ using disk diffusion assay and showed antagonistic effects to *Vibrio campbellii*, *V*. antagonistic effects to Vibrio campbellii, V.

Fig. 2 Antivibrio compounds isolated from Pseudoalteromonas spp.

coralliilyticus, V. harveyi, and V. vulnificus. The korormicins may play a role in disrupting quorum sensing.⁶²

3.3. Pseudomonas spp.

Pseudomonas aeruginosa is Gram-negative bacteria, widespread in the terrestrial and marine environment. It has been reported that Pseudomonas aeruginosa exhibited antagonistic activity to aquaculture and agriculture pathogens. Some antivibrio compounds have been identified from P. aeruginosa as seen in Fig. 3 and Table 3.

Pseudomonas MCCB 102 and 103A produces phenazine antibiotic, N-methyl-1-hydroxyphenazine (24). The compound

has bacteriostatic activity against V. harveyi at the dose of 0.5 mg L⁻¹. The toxicity in *Penaeus monodon* haemocyte at IC₅₀ was 1.4 \pm 0.31 mg L⁻¹.⁶³ Investigation of bioactivities and toxicities of ethyl acetate extract of Pseudomonas aeruginosa sp. W3 led to the isolation of 2-heptyl-4-quinolone (HHQ) (25) that was active against 18 strains of shrimp pathogenic of V. harveyi. The compound was active at MIC value 225–450 μ g mL⁻¹.⁶⁴
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Antagonistic activity of Pseudomonas aeruginosa was tested against V. parahaemolyticus, V. vulnificus, V. alginolyticus, V. fluvialis, V. mediteranii, V. nereis, and V. harveyi. Isolation of extract fermentation led to identify pyocyanin (26) as the bioactive compound responsible for the antagonistic effect at a concentration of more than 30 mg L⁻¹.⁶⁵ Pseudomonas aeruginosa PA31X produces phenazine-1-carboxylic acid (27) that is active against *V. anguillarum* C312 at 3 µg mL⁻¹.⁶⁶

3.4. Miscellaneous bacteria

A Gram-positive marine bacterium Halobacillus salinus produced two phenetylamide metabolites: N-(2'-phenylethyl)isobutyramide (28) and 3-methyl-N-(2′ -phenylethyl)-butyramid (29). The compounds are anti-quorum sensing and bioluminescence of *V. harveyi* at a concentration below 200 μ g mL⁻¹.⁶⁷
Olgie agid (20) isolated from *Vibria* and from North Chile

Oleic acid (30) isolated from Vibrio sp. from North Chile inhibited the growth of *V. parahaemolyticus*. Long-chain fatty acids such as oleic, linoleic, and linolenic have antibacterial activity through inhibition of fatty acid synthesis (Table 4).⁶⁸

	Table 2 Bioactivity of antivibrio compounds from Pseudoalteromonas								
	No. Compounds	Sources	Antivibrio activities	Mechanism of action	Ref.				
	1 $2-n$ -Pentyl-4-quinolinol (15)	Pseudoalteromonas A1-J11	<i>V. harveyi</i> ATCC 14126, <i>V. harveyi</i> Inhibition of the growth ATCC 35084, V. fischeri VF-74, V. harveyi, Dose 10 µg per disk		60				
2	• Isovaleric acid (16) , • 2-methyl butyric acid (17)	Pseudoalteromonas haloplanktis INH	V ordalii ATCC 33509, V. alginolyticus ATCC 17749, V. anguillarum IFO 13266, dose 1 mg mL^{-1}	Inhibition of the growth	61				
3	• Tetrabromopyrrole (18) , • $4'$ - $((3,4,5\text{-tribromo-1}H\text{-pyrrol-2-vl})$ methyl) phenol (19) , • korormicin G-I $(20-22)$. • korormicin K (23)	Pseudoalteromonas [010	V. campbelii, V. vulnificus, V. coralliilyticus, V. harveyi, Dose 200 disrupting of quorum sensing μ g mL ⁻¹	\bullet Inhibition of the growth, \bullet	62				

Table 2 Bioactivity of antivibrio compounds from Pseudoalteromonas

Table 3 Bioactivity of antivibrio compounds from Pseudomonas spp

Table 4 Bioactivity of antivibrio compounds from Miscellaneous bacteria

Fig. 4 Antivibrio compounds isolated from miscellaneous bacteria.

Bacillus pumilus H2 produces an antibacterial compound amicoumacin A (31) (Fig. 4) inhibited broad range species of Vibro V. natriegens, V. vulnificus, V. alginolyticus, V. harveyi, V. azareus, V. campbelli, V. fischeri.⁶⁹

4. Antivibrio from marine fungi

Since the discovery of penicillin from Penicillium chrysogenum in the twentieth century, the fungus is an important source of natural products for drug discovery. Marine fungi have been explored for their bioactive secondary metabolites and potential for anti-microbial agents.⁷⁰–⁷² To date, 38% of 22.000 bioactive microbial metabolites are from fungi.⁷³ Among those metabolites, there are only a few antivibrio agents derived from marine fungi as presented in Fig. 5 and Table 5.

The genera of Penicillium contributes diverse of antivibrio compounds. Extraction of culture Penicillium sp. AS-79

associated with sea anemone Haliplanella luciae yielded indole diterpenoids that are active against V. parahaemolyticus and V. alginolyticus. The various compounds: 6-hydroxylpaspalinine (32), paspalitrem C (33), emindole SB (34), 3-deoxo-4b-deoxypaxilline (35) , and $10,23$ -dihydro-24,25-dehydroaflavinine (36) exhibited activity against the aquatic pathogen V. parahaemolyticus. In addition, compounds 33, 34, 36 showed inhibition activity against V. alginolyticus.⁷⁴ Chemical investigation of ethyl acetate extract of culture Penicillium janthinellum yielded two indole diterpenoid penijanthine C (37) and D (38) , two new steroids penijanthoid A (39) and B (40) along with two known analogs PC-M6 and 7-hydoxy-13-dehydroxypaxilline. The compounds 37–40 were active against V. anguillarum, V. parahaemolyticus, and V. alginolyticus. Indole diterpenoid is a new class of antivibrio agents.⁷⁵

The genera of *Aspergillus* produce flourishing classes of antivibrio compounds. Deep investigation of marine-derived fungus Aspergillus sp. ZA-01 led to the isolation of new antivibrio compounds prenylxanthone derivate aspergixanthones I–K (41–43) along with known compounds (44–47). The compounds were tested against V. parahaemolyticus, V. anguillarum, and V. alginolyticus.⁷⁶ Marine fungi Aspergillus terreus EN-539 associated red algae Laurencia okamurai, produced new prenylated phenol derivative terreprenphenol A (48) along with 4-hydroxy-3-prenybenzoic acid (49), and 4-hydroxy-3-(3-methylbut-2-enyl)-benzaldehyde (50). Evaluation of antivibrio activity against V. harveyi, V. parahemolyticus, and V. vulnificus showed inhibitory activity at MIC values ranging from 4 to 64 μ g mL^{−1}.⁷⁷
The deep see sediment derived funcys depensilles versiseler SD. The deep-sea sediment-derived fungus Aspergillus versicolor SD-330 yielded one new aromatic bisabolene-type sesquiterpenoid (51) along with four known analogs, aspergoterpenin C (52), $(7S, 11S)$ -(b)-12-hydroxysydonic acid (53) , (S) -(b)-11-dehydrosydonic acid (54), and engyodontiumone I (55). All compounds exhibited inhibitory activities against V.

anguillarum, V. harveyi, and V. parahaemolyticus with the MIC values ranging from 4 to >32 μ g mL^{-1,78} Bioassay-guided
isolation has identified the bioastive compound transaiding isolation has identified the bioactive compound trypacidin (56) from a marine symbiotic fungi Aspergillus fumigatus HX-1. In vitro bacteriostatic assay confirms the MIC value at 31.25 μ g mL⁻¹.⁷⁹ The MIC of each compound is presented in Table 5.

Marine fungi associated with crab, Paraconiothyrium sp., produced a new polyketide, paraconthone A (57) together with botryosphaerone (58) and O-methylaspmenone (59). The compounds showed moderate inhibitory effects against V. anguillarum and V. parahaemolyticus at 30 μ g mL $^{-1}$.^{so}

Table 5 Bioactivity of antivibrio compounds isolated from marine fungi Table 5

A new steroid acremocholone (60) was produced by spongeassociated fungi Acremonium sp. NBUF150. Acremocholone exhibited antivibrio activity against V. scophthalmi, V. shilonii and *V. brasiliensis* at MIC of 8 µg mL⁻¹.⁸¹

5. Antivibrio from sponges

Sponges are the oldest metazoan and have been investigated extensively for bioactive metabolites. Three new alkaloids isonaamide D, di-isonaamide A, and leucettamine D, and two known compounds isonaamine A and isonaanidine from a sponge Leucetta chagosensis Dendy, 1863 from French Polynesia. The compounds were screened for quorum sensing (QS) inhibitor of V. harveyi. The result showed that Isonaamidine A (61) inhibited the QS pathway at IC₅₀ 1 µg mL⁻¹. None of the compounds of footed bostorial growth at 50 µg mL^{-1 s2} compounds affected bacterial growth at 50 μg mL^{-1 ,82}

In the searching for antimicrobial agents against V . *vulnificus* twelve pure marine compounds from a variety of sponges were screened for inhibition effect. Psammaplin A (62), a bromotyrosine derivative from the sponge Poecillastra sp., Jaspis sp., and Psammaplin aplysilla inhibited V. vulnificus in vitro and in vivo assay at $5-50 \mu$ g (Table 6).⁸³

Alkaloid aaptamin and derivates from sponge Aaptos aaptos were tested against Vibrio spp. and V. harveyi. Aaptamine (63), 9 demethylaaptamine (64), 4-N-methylaaptamine (65), 9-methoxyaaptamine (66) were active at concentration 1 mg mL^{-1} $(Fig. 6).^{84}$

6. Antivibrio from coral

Four new steroids, dendronecholones A–D (67–70), and two known analogues, $12\beta,16\beta,20$ -trihydroxycholesta-1,4-dien-3-one 16-acetate (71) and nanjiol A (72) were identified from soft coral Dendronephthya collected in waters off Zhejiang Province, China. Antivibrio assay was conducted against V. parahaemolyticus, V. scophthalmi, and V. harveyi. The MIC range from 8–>32 μ g mL⁻¹ is presented in Table 7.⁸⁵

7. Antivibrio from seaweeds

Seaweeds are well known as rich sources of primary and secondary metabolites with diverse applications for food, feed, agriculture, pharmaceutical, and cosmetics.^{86,87} Numerous substances were isolated from seaweed such as halogenated compounds,^{88,89} polyether,⁹⁰ phenolic compounds,⁹¹ and polyunsaturated fatty acid.⁹² Antimicrobial activity testing of seaweed extracts support the possibility of using seaweeds as a source of antimicrobial agents or as a health-promoting feed for aquaculture.⁹³ Bioactive compounds from seaweed can be applied in aquaculture health and disease management to control bacterial infection.⁹⁴⁻⁹⁶ Seaweeds are rich in fatty acid and the mechanism of action of fatty acid as an antibacterial agent through inhibition of the electron transport chain and normal oxidative phosphorylation in bacterial membranes.⁹⁷ Polysaccharides from seaweed have been examined for the purpose as prebiotic or immunostimulant in

Table 6 Bioactivity of antivibrio compounds isolated from sponge

Fig. 6 Antivibrio compounds isolated from sponge.

aquaculture⁹⁸ while red seaweed (Rhodophyta) are good source of antibacterial agents (Table 8).⁹⁹

Water-soluble fractions of red algae Palmaria paltata and Grateloupia turuturu were examined for the activity against V. harveyi. The NMR data suggested that the active water fraction of Palmaria paltata contains floridoside (73) (Fig. 7).¹⁶ Further structure elucidation should be done to identify principal compounds responsible for an antivibrio agent.¹⁶

Red algae Delisea pulchra produced halogenated furanones called fimbrolide (Fig. 8).¹⁰¹ Brominated furanones from marine algae inhibited biofilm formation and quorum sensing (QS) Gram-negative without affecting their growth. The structure is similar to bacterial acyl homoserine lactones (AHL).¹⁰⁰ Some marine algae produced halogenated furanones as AHL antagonists as a response to the negative impact of bacterial colonization. Fimbrolide 1 (74) and Fimbrolide 2 (75) were tested for inhibiting bioluminescence in V. harveyi and V. campbellii with the target on LuxS, PhaB, and uncharacterized IMPD protein.¹⁰¹

Extracts of Indonesian red seaweeds have been screened for bioactivity against fish pathogens including Vibrio spp. Extract of Gracilaria arcuata was active against Vibrio sp. at a concentration of 2.5 $\mu g \mu L^{-1}$. The active fraction contained hex-
adecessorie acid and starel compounds such as Expect 5 on 2.0 adecanoic acid and sterol compounds such as Ergost-5-en-3-ol

No.	Compounds	Sources	Antivibrio Activities	Mechanism of action	Ref.
$\mathbf{1}$	Dendronecholone A (67)	Dendronephthya	V. scophthalmi MIC 32 μ g mL ⁻¹ , V. parahemolyticus MIC >32 µg mL ⁻¹ , V. harveyi MIC 32 μ g mL ⁻¹	Inhibition of the growth	85
2	Dendronecholone B (68)	Dendronephthya	V. scophthalmi MIC 8 μ g mL ⁻¹ , V. parahemolyticus MIC >32 µg mL ⁻¹ , V. harveyi MIC 8 µg mL ⁻¹	Inhibition of the growth	85
3	Dendronecholone C (69)	Dendronephthya	V. scophthalmi MIC 32 μ g mL ⁻¹ , V. parahemolyticus MIC 8 µg mL ⁻¹ , V. harveyi MIC >32 µg mL $^{-1}$	Inhibition of the growth	85
$\overline{4}$	Dendronecholone D (70)	Dendronephthya	V. scophthalmi MIC 16 μ g mL ⁻¹ , V. parahemolyticus MIC >32 µg mL ⁻¹ , V. harveyi MIC >32 µg mL ⁻¹	Inhibition of the growth	85
5	12β,16β,20-Trihydroxycholesta-1,4-dien- 3-one 16-acetate (71)	Dendronephthya	<i>V. scophthalmi</i> MIC 8 μ g mL ⁻¹ , <i>V.</i> parahemolyticus MIC > 32 µg mL ⁻¹ , V. harveyi MIC >32 µg mL ⁻¹	Inhibition of the growth	85
6	Nanjiol A (72)	Dendronephthya	V. scophthalmi MIC 8 μ g mL ⁻¹ , V. parahemolyticus MIC 8 µg mL ⁻¹ , V. harveyi MIC 8 μ g mL ⁻¹	Inhibition of the growth	85

Table 7 Bioactivity of antivibrio compounds isolated from coral

Table 8 Bioactivity of antivibrio compounds isolated from seaweed

(76), Stigmast-5-en-3 β -ol (77). The MIC of the active fraction was 1.25 µg µL^{−1}.¹⁰² Extract of Indonesian seaweed *Gracilaria edulis*
showed inhibition against *V* fluvialis and *V* campbelii. Eurther showed inhibition against V. fluvialis and V. compbelii. Further analysis showed that the active fraction contained sterol

cholest-8-en-3-ol (78) and long-chain fatty acids such as pentadecanoic acid (79), hexadecanoic acid (80), 13-octadecenoic acid (81), 10-octadecenoic acid (82), eicosanoic acid (83). The active

Fig. 7 Antivibrio compounds isolated from coral.

Fig. 8 Antivibrio compounds isolated from seaweed.

Table 9 Bioactivity of antivibrio from plants

No.	Compounds	Sources	Antivibrio Activities	Mechanism of action	Ref.
	Capsaicin (86)	Capsicum annum	V. chloreae	Inhibition of toxin	112
\mathcal{D}_{α}	Curcumin (87)	Curcuma longa	V. harveyi reduce bioluminescence 88%	Interfere the production of QS- dependent virulence factors in Vibrio spp., inhibition of bacterial adhesion and RTX toxin binding	113
3	Piperidine (88)	Piper bettle	<i>Vibrio</i> spp., MIC ₉₀ 2–6 mg mL ⁻¹	Inhibition of the growth	114
$\overline{4}$	Chlogenic acid (89)	Piper bettle	Vibrio spp. MIC ₉₀ 5-16 mg mL ⁻¹	Inhibition of the growth	114
5	Eugenyl acetate (90)	Piper bettle	Vibrio spp. MIC ₉₀ 5-20 mg mL ⁻¹	Inhibition of the growth	114
6	Punicalagin (91)	Punica granatum Linn	V. anguillarum MIC 25 mg mL ⁻¹	Inhibition of the growth	115

fraction showed inhibition against V . fluvialis at MIC 2.5 $μg mL^{-1,103}$

Ethanolic extract of Gracilaria fischeri exhibited anti-quorum sensing activity in V. harveyi and V. parahaemolyticus at concentrations of 5, 10, and 100 μ g mL⁻¹. The extract also
reduced the luminoscenes of *V* harvay ¹⁰⁴ Eurther investigation reduced the luminescence of *V. harveyi*.¹⁰⁴ Further investigation showed G. fisheri contains N-benzyl cinnamamide (84) and α resorcylic acid (85) and which are responsible for antivibrio activity.¹⁰⁵

8. Antivibrio from plants

Plants are well known as a source of bioactive compounds and are used in traditional medicine. Various plant extracts containing phenolic, alkaloid, flavonoid, and polysaccharide have been tested and used in aquaculture as an immunostimulant, antioxidant, prebiotic, antibacterial, and antifungal.106,107 Plant extracts have been screened as sources for antivibrio agents.108,109 Phytochemicals can be used to interfere with bacterial quorum sensing to counteract the biofilm resistance. Medicinal plants are rich resources for screening bioactive QS.¹¹⁰ Antivibrio compounds identified from plants are shown in Table 9.

The essential oil from aromatic plants Mentha longifolia, M. pulegium, Eugenia caryophyllata, Thymus vulgaris, and Rosmarinus officinalis were tested against V. alginolyticus, V. parahaemolyticus, V. vulnificus, and V. fluvialis strains. Results showed variable activity and essential oils of T. vulgaris yielded the highest zone of growth inhibition against V. parahaemolyticus. 111

One of the approaches in the screening of natural products as antivibrio is targeting the production of virulence factors such as capsaicin and curcumin. Extract methanol of Capsicum annum containing capsaicin was reported to inhibit CT (cholera toxin) production in *V. cholerae*. The transcriptions of *ctxA*, *tcpA*, and toxT genes were repressed by capsaicin (86). On the contrary, capsaicin significantly enhanced the transcription of the hns gene, the product of which is known to regulate negatively the transcription of *ctxAB*, *tcpA*, and *toxT* genes. These results suggest that capsaicin might act as a potent repressor for CT production possibly by enhancing the transcription of hns.¹¹² Curcumin (87) from Curcuma longa reduced 88% of bioluminescence of *V. harveyi* and inhibited components of biofilms

Fig. 9 Antivibrio compounds isolated from plants.

and virulence factor in V. parahaemolyticus, V. vulnificus, V. harveyi. 113

Three compounds piperidine (88), chlorogenic acid (89), and eugenyl acetate (90) isolated from Piper bettle were reported as bactericidal against several pathogenic Vibrio spp. The MIC range 0.6 to 16 mg mL^{-1} . Piperidine has the strongest inhibition effect on Vibrio spp. compare to chlorogenic acid and eugenyl acetate (Fig. 9).¹¹⁴

Punicalagin (91) from pomegranate (Punica granatum Linn.) was reported against *V. anguillarum* at MIC 25 µg mL⁻¹.¹¹⁵

9. Conclusions and perspective

Climate change and global warming will impact increasing cases of vibriosis in the future. *Vibrio* spp. cause serious problems in aquaculture with consequent huge economic losses. Moreover, vibriosis threatens human health through seafood contamination and contact with seawater during wound events. To date, an effective vaccine to prevent vibriosis has not been available yet. Efforts have been done to prevent vibriosis in aquaculture with probiotics, prebiotics, and immunostimulants. The rising incidence of Vibrio resistance to antimicrobial agents and the limited option of antibiotics have driven the search for new antivibrio agents.

Different stages of work have been performed ranging from the preliminary screening to an in-depth characterization of antivibrio compounds. This review provides proof that natural products are promising as a source of antivibrio agents. Screening of natural products from different sources has been carried out to discover antivibrio agents. Fig. 10 summarizes the exploration of natural resources to discover antivibrio agents. Natural product compounds exhibit bioactivity against Vibrio spp. through mechanism of action inhibiting the growth, disrupting quorum sensing, and interfering with biofilm formation.

This review shows that natural products as antivibrio are produced by prokaryotes and eukaryotes living in terrestrial and marine environments (Fig. 11). Based on data on this review, marine fungi demonstrated prolific sources of antivibrio and contribute 36% of bioactive antivibrio. Actinobacteria and sponges are well-known as sources of bioactive compounds for decades, but their compounds account for only 16% and 7%, respectively for antivibrio. The type classes of natural antivibrio derived from natural product compounds are alkaloid, polyketide, peptide, sterol, terpene, organic acid, and fatty acid.

Fig. 11 The structure type of antivibrio compounds derived from natural resources.

Fig. 12 The biological sources of natural products with antivibrio activity.

Fig. 10 Summary of the chemistry of natural products as antivibrio and their mechanism of actions.

Polyketide and alkaloid are the major class of antivibrio compounds and count about 28% and 23%, respectively presented in this review (Fig. 12). The alkaloids are produced by fungi and sponges, while polyketides were produced by mostly all organisms except coral. Antivibrio from coral and seaweed are mostly sterol.

This review summarizes that nature has provided a plethora of natural products with extraordinary chemistry and bioactivity against Vibrio spp. Further research and development of promising compounds are necessary for application in aquaculture and human health. Future efforts are necessary to evaluate the biological activities in vivo, toxicity, and mechanisms of action. Biofilms is the leading cause of multidrug resistance among microorganisms including Vibrio spp. Thus, study and examination of antivibrio compounds as inhibitor of biofilm formation is needed. The clinical study of antivibrio compounds has not been reported yet. Post compounds are the busine of a method on the common Access Article is the common access Article is the busine of the common Access Article is likely expected. And the expected on the same of the common accept and the c

Author contribution

NK and MU wrote the manuscript, DCR helped supervise the project and reviewed the manuscript, MU contributed to collect data and references. NK received the funding and DCR provided lab access to do research on antivibrio.

Conflicts of interest

The authors report there are no competing interests to declare.

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References

- 1 B. Austin, Vet. Microbiol., 2010, 140(3), 310–317, DOI: [10.1016/j.vetmic.2009.03.015](https://doi.org/10.1016/j.vetmic.2009.03.015).
- 2 C. N. Johnson, Microb. Ecol., 2013, 65(4), 826–851, DOI: [10.1007/s00248-012-0168-x](https://doi.org/10.1007/s00248-012-0168-x).
- 3 F. L. Thompson, T. Iida and J. Swings, Microbiol. Mol. Biol. Rev., 2004, 68(3), 403–431, DOI: [10.1128/MMBR.68.3.403-](https://doi.org/10.1128/MMBR.68.3.403-431.2004) [431.2004](https://doi.org/10.1128/MMBR.68.3.403-431.2004).
- 4 A. Newton, M. Kendall, D. J. Vugia, O. L. Henao and B. E. Mahon, Clin. Infect. Dis., 2012, 54, 391–395, DOI: [10.1093/cid/cis243](https://doi.org/10.1093/cid/cis243).
- 5 L. Vezzulli, R. R. Colwell and C. Pruzzo, Microb. Ecol., 2013, 65(4), 817–825, DOI: [10.1007/s00248-012-0163-2](https://doi.org/10.1007/s00248-012-0163-2).
- 6 L. Vezzulli, I. Brettar, E. Pezzati, P. C. Reid, R. R. Colwell, M. G. Hofle and C. Pruzzo, ISME J., 2012, 6, 21-30, DOI: [10.1038/ismej.2011.89](https://doi.org/10.1038/ismej.2011.89).
- 7 C. Baker-Austin, J. A. Trinanes, N. G. H. Taylor, R. Hartnell, A. Siitonen and J. Martinez-Urtaza, Nat. Clim. Change, 2013, 3, 73–77, DOI: [10.1038/nclimate1628](https://doi.org/10.1038/nclimate1628).
- 8 L. Vezzulli, E. Pezzati, I. Brettar, M. Hofle and C. Pruzzo, Microbiol. Spectrum, 2014, 3, 1, DOI: [10.1128/](https://doi.org/10.1128/microbiolspec.VE-0004-2014) [microbiolspec.VE-0004-2014](https://doi.org/10.1128/microbiolspec.VE-0004-2014).
- 9 J. M. Janda, A. E. Newton and C. A. Bopp, Clin. Lab. Med., 2015, 35(2), 273–288, DOI: [10.1016/j.cll.2015.02.007](https://doi.org/10.1016/j.cll.2015.02.007).
- 10 M. Y. Ina-Salwany, N. Al-Saari, A. Mohamad, F.-A. Mursidi, A. Mohd-Aris, M. N. A. Amal, H. Kasai and M. Zamri-Saad, J. Aquat. Anim. Health, 2019, 31, 3–22, DOI: [10.1002/](https://doi.org/10.1002/aah.10045) [aah.10045](https://doi.org/10.1002/aah.10045).
- 11 A. E. Toranzo, B. Magarinos and J. L. Romalde, Aquaculture, 2005, 246, 37–61, DOI: [10.1016/j.aquaculture.2005.01.002](https://doi.org/10.1016/j.aquaculture.2005.01.002).
- 12 I. Frans, C. W. Michiels, P. Bossier, K. A. Willems, B. Lievens and H. Rediers, J. Fish Dis., 2011, 34, 643–661, DOI: [10.1111/](https://doi.org/10.1111/j.1365-2761.2011.01279.x) [j.1365-2761.2011.01279.x](https://doi.org/10.1111/j.1365-2761.2011.01279.x).
- 13 N. Mohamad, M. N. A. Amala, I. S. Yasin, M. Z. Saad, N. S. Nasruddin, N. Al-Saarif, S. Minog and T. Sawab, Aquaculture, 2019, 512, 734289, DOI: [10.1016/](https://doi.org/10.1016/j.aquaculture.2019.734289) [j.aquaculture.2019.734289](https://doi.org/10.1016/j.aquaculture.2019.734289).
- 14 U. Khimmakthong and P. Sukkarun, Microb. Pathog., 2017, 113, 107–112, DOI: [10.1016/j.micpath.2017.10.028](https://doi.org/10.1016/j.micpath.2017.10.028).
- 15 B. Austin and X. H. Zhang, Lett. Appl. Microbiol., 2006, 43(2), 119–124, DOI: [10.1111/j.1472-765X.2006.01989.x](https://doi.org/10.1111/j.1472-765X.2006.01989.x).
- 16 N. Garcia-Bueno, P. Decottignies, V. Turpin, J. Dumay, C. Paillard, V. Stiger-Pouvreau, N. Kervarec, Y.-F. Pouchus, A. A. Marrin-Atucha and J. Fleurence, Aquat. Living Resour., 2014, 27, 83–89, DOI: [10.1051/alr/2014009](https://doi.org/10.1051/alr/2014009).
- 17 J. Dubert, J. L. Barja and J. L. Romalde, Front. Microbiol., 2017, 8, 762, DOI: [10.3389/fmicb.2017.00762](https://doi.org/10.3389/fmicb.2017.00762).
- 18 R. Beaz-Hidalgo, S. Balboa, J. L. Romalde and M. J. Figueras, Environ. Microbiol. Rep., 2010, 2(1), 34–43, DOI: [10.1111/j.1758-2229.2010.00135](https://doi.org/10.1111/j.1758-2229.2010.00135).
- 19 C. Baker-Austin, J. D. Oliver, M. Alam, A. Ali, M. K. Waldor, F. Qadri and J. Martinez-Urtaza, Nat. Rev. Dis. Prim., 2018, 4, 8, DOI: [10.1038/s41572-018-0005-8](https://doi.org/10.1038/s41572-018-0005-8).
- 20 R. Finkelstein and I. Oren, Curr. Infect. Dis. Rep., 2011, 13, 470–477, DOI: [10.1007/s11908-011-0199-3](https://doi.org/10.1007/s11908-011-0199-3).
- 21 A. M. Dechet, A. Y. Patricia, N. Koram and J. Painter, Clin. Infect. Dis., 2008, 46, 970–976, DOI: [10.1086/529148](https://doi.org/10.1086/529148).
- 22 S.-P. Heng, V. Letchumanan, C.-Y. Deng, N.-S. S. Mutalib, T. M. Khan, L.-H. Chuah, K.-G. Chan, B.-H. Goh, P. Pusparajah and L.-H. Lee, Front. Microbiol., 2017, 8, 997, DOI: [10.3389/fmicb.2017.00997](https://doi.org/10.3389/fmicb.2017.00997).
- 23 S. M. Aly and A. Albutti, J. Aquacult. Res. Dev., 2014, 5, 4, DOI: [10.4172/2155-9546.1000247](https://doi.org/10.4172/2155-9546.1000247).
- 24 J. E. M. Watts, H. J. Schreier, L. Lanska and M. S. Hale, Mar. Drugs, 2017, 5, 158, DOI: [10.3390/md15060158](https://doi.org/10.3390/md15060158).
- 25 R. H. Rebouças, O. V. Sousa, A. S. Lima, F. R. Vasconcelos, P. B. Carvalho and R. H. S. F. Viera, Environ. Res., 2011, 11, 21–24, DOI: [10.1016/j.envres.2010.09.012](https://doi.org/10.1016/j.envres.2010.09.012).
- 26 C. Scarano, C. Spanu, G. Ziino, F. Pedonese, A. Dalmasso, V. Spanu, S. Virdis and E. P. De Santis, New Microbiol., 2014, 37(3), 329–337.
- 27 R. X. Wang, J. Y. Wang, Y. C. Sun, B. L. Yang and A. L. Wang, Mar. Pollut. Bull., 2015, 101(2), 701–706, DOI: [10.1016/](https://doi.org/10.1016/j.marpolbul.2015.10.027) [j.marpolbul.2015.10.027](https://doi.org/10.1016/j.marpolbul.2015.10.027).
- 28 Q. Yu, M. Niu, M. Yu, Y. Liu, D. Wang and X. Shi, Food Control, 2016, 60, 263–268, DOI: [10.1016/](https://doi.org/10.1016/j.foodcont.2015.08.005) [j.foodcont.2015.08.005](https://doi.org/10.1016/j.foodcont.2015.08.005).
- 29 S. Elmahdi, L. V. DaSilva and S. Parveen, Food Microbiol., 2016, 57, 128–134, DOI: [10.1016/j.fm.2016.02.008](https://doi.org/10.1016/j.fm.2016.02.008).
- 30 C. H. Kang, Y. Shin, S. Jang, Y. Jung and J. S. So, Environ. Sci. Pollut. Res., 2016, 23(20), 21106–21112, DOI: [10.1007/](https://doi.org/10.1007/s11356-016-7426-2) [s11356-016-7426-2](https://doi.org/10.1007/s11356-016-7426-2).
- 31 D. L. Milton, Int. J. Med. Microbiol., 2006, 296(2), 61–71, DOI: [10.1016/j.ijmm.2006.01.044](https://doi.org/10.1016/j.ijmm.2006.01.044).
- 32 Y. Deng, Y. Liu, J. Li, X. Wang, S. He, X. Yan, Y. Shi, W. Zhang and L. Ding, Eur. J. Med. Chem., 2022, 239, 114513, DOI: [10.1016/j.ejmech.2022.114513](https://doi.org/10.1016/j.ejmech.2022.114513).
- 33 J. Zhao, M. Chen, C. S. Quan and S. D. Fan, J. Fish Dis., 2015, 38(9), 771–786, DOI: [10.1111/jfd.12299](https://doi.org/10.1111/jfd.12299).
- 34 F. H. Yildiz and K. L. Visick, Trends Microbiol., 2009, 17(3), 109–118, DOI: [10.1016/j.tim.2008.12.004](https://doi.org/10.1016/j.tim.2008.12.004).
- 35 L. A. Hawver, S. A. Jung and W. L. Ng, FEMS Microbiol. Rev., 2016, 40, 738–752, DOI: [10.1093/femsre/fuw014](https://doi.org/10.1093/femsre/fuw014).
- 36 K. Papenfort and B. L. Bassler, Nat. Rev. Microbiol., 2016, 14(9), 576–588, DOI: [10.1038/nrmicro.2016.89](https://doi.org/10.1038/nrmicro.2016.89).
- 37 B. Vu, M. Chen, R. J. Crawford and E. P. Ivanova, Molecules, 2009, 14(7), 2535–2554, DOI: [10.3390/molecules14072535](https://doi.org/10.3390/molecules14072535).
- 38 S. E. Rossiter, M. H. Fletcher and W. M. Wuest, Chem. Rev., 2017, 117, 12415–12474, DOI: [10.1021/](https://doi.org/10.1021/acs.chemrev.7b00283) [acs.chemrev.7b00283](https://doi.org/10.1021/acs.chemrev.7b00283).
- 39 J. F. M. Ong, H. C. Goh, S. C. Lim, L. M. Pang, J. S. F. Chin, K. S. Tan, Z.-X. Liang, L. Yang, E. Glukhov, W. H. Gerwick and L. T. Tan, Mar. Drugs, 2019, 17, 72, DOI: [10.3390/](https://doi.org/10.3390/md17010072) [md17010072](https://doi.org/10.3390/md17010072).
- 40 O. Geniloud, Nat. Prod. Rep., 2017, 34, 1203–1232, DOI: [10.1039/c7np00026j](https://doi.org/10.1039/c7np00026j).
- 41 S. W. Behie, B. Bonet, V. M. Zacharia, D. J. McClung and M. F. Traxler, Front. Microbiol., 2017, 7, 2149, DOI: [10.3389/fmicb.2016.02149](https://doi.org/10.3389/fmicb.2016.02149).
- 42 J. L. You, L. X. Cao, G. F. Liu, S. N. Zhou, H. M. Tan and Y. C. Lin, World J. Microbiol. Biotechnol., 2005, 21(5), 679– 682, DOI: [10.1007/s11274-004-3851-3](https://doi.org/10.1007/s11274-004-3851-3).
- 43 J. L. You, X. L. Xue, L. X. Cao, X. Lu, J. Wang, L. Zhang and S. Zhou, Appl. Microbiol. Biotechnol., 2007, 76(5), 1137–1144, DOI: [10.1007/s00253-007-1074-x](https://doi.org/10.1007/s00253-007-1074-x).
- 44 N. Augustine, S. Kerkar and S. Thomas, Curr. Microbiol., 2012, 64(4), 338–342, DOI: [10.1007/s00284-011-0073-4](https://doi.org/10.1007/s00284-011-0073-4).
- 45 L. T.-H. Tan, K.-G. Chan, L.-H. Lee and B.-H. Goh, Front. Microbiol., 2016, 7, 79, DOI: [10.3389/fmicb.2016.00079](https://doi.org/10.3389/fmicb.2016.00079).
- 46 M. Bodhaguru, S. Prakash, R. Ramasubburayan, N. K. Ahila, L. Mariselvam, G. Immanuel, A. Palavesa and E. Kannapiran, Microb. Pathog., 2019, 134, 103597, DOI: [10.1016/j.micpath.2019.103597](https://doi.org/10.1016/j.micpath.2019.103597).
- 47 G. Raissa, D. E. Waturangi and D. Wahjuningrum, BMC Microbiol., 2020, 20, 343, DOI: [10.1186/s12866-020-02022-z](https://doi.org/10.1186/s12866-020-02022-z).
- 48 L. T.-H. Tan, L.-H. Lee and B.-H. Goh, Prog. Microbes. Mol. Biol., 2019, 2, 1.
- 49 G. S. Kiran, A. N. Lipton, S. Priyadharshini, K. Anitha, L. E. Suárez, M. V. Arasu, K. C. Choi, J. Selvin and N. A. Al-Dhabi, Microb. Cell Fact., 2014, 13, 114, DOI: [10.1186/s12934-014-0114-3](https://doi.org/10.1186/s12934-014-0114-3).
- 50 M. Hayashi, T. Unemoto, S. Minami-Kakinuma, H. Tanaka and S. Omura, J. Antibiot., 1982, 35, 1078–1085, DOI: [10.7164/antibiotics.35.1078](https://doi.org/10.7164/antibiotics.35.1078).
- 51 U. F. Castillo, G. A. Strobel, E. J. Ford, W. M. Hess, H. Porter, J. B. Jensen, H. Albert, R. Robinson, M. A. M. Condron, D. B. Teplow, D. Stevens and D. Yaver, Microbiology, 2002, 148(9), 2675–2685, DOI: [10.1099/00221287-148-9-2675](https://doi.org/10.1099/00221287-148-9-2675).
- 52 M. Iwatsuki, R. Uchida, H. Yoshijima, H. Ui, K. Shiomi, A. Matsumoto, Y. Takahashi, A. Abe, H. Tomoda and S. Omura, J. Antibiot., 2008, 61, 222–229, DOI: [10.1038/](https://doi.org/10.1038/ja.2008.32) [ja.2008.32](https://doi.org/10.1038/ja.2008.32). Review Water Articles. Articles. Published on December 2022. Downloaded on 2022. Downloaded on 2022. Downloaded on 5/29/2024 12:43:56 PM. This article is licensed under a creative Commons Attribution-Non-Commons Attributi
	- 53 T. C. Holmes, A. E. May, K. Zaleta-Rivera, J. G. Ruby, P. Skewes-Cox, M. A. Fischbach, J. L. DeRisi, M. Iwatsuki, S. Omura and C. Khosla, J. Am. Chem. Soc., 2012, 134(42), 17797–17806, DOI: [10.1021/ja308622d](https://doi.org/10.1021/ja308622d).
	- 54 J. Y. Cho and M. S. Kim, Fish. Sci., 2012, 78(5), 1065–1073, DOI: [10.1007/s12562-012-0531-3](https://doi.org/10.1007/s12562-012-0531-3).
	- 55 N. Yang and C. Sun, Front. Microbiol., 2016, 7, 1467, DOI: [10.3389/fmicb.2016.01467](https://doi.org/10.3389/fmicb.2016.01467).
	- 56 M. E. Rateb, W. E. Houssen, W. T. A. Harrison, H. Deng, C. K. Okoro, J. A. Asenjo, B. A. Andrews, A. T. Bull, M. Goodfellow, R. Ebel and M. Jaspars, J. Nat. Prod., 2011, 74(9), 1965–1971, DOI: [10.1021/np200470u](https://doi.org/10.1021/np200470u).
	- 57 J. F. Castro, V. Razmilic, J. P. Gomez-Escribano, B. Andrews, J. Asenjo and M. Bibb, Antonie van Leeuwenhoek, 2018, 111(8), 1433–1448, DOI: [10.1007/s10482-018-1034-8](https://doi.org/10.1007/s10482-018-1034-8).
	- 58 J. P. Bowman, Mar. Drugs, 2007, 5, 220–241, DOI: [10.3390/](https://doi.org/10.3390/md504220) [md504220](https://doi.org/10.3390/md504220).
	- 59 C. Offret, F. Desriac, P. L. Chevalier, J. Mounier, C. Jegou and Y. Fleury, Mar. Drugs, 2016, 14, 129, DOI: [10.3390/](https://doi.org/10.3390/md14070129) [md14070129](https://doi.org/10.3390/md14070129).
	- 60 C. S. Castillo, M. I. Wahid, T. Yoshikawa and T. Sakata, Fish. Sci., 2008, 74, 174–179, DOI: [10.1111/j.1444-](https://doi.org/10.1111/j.1444-2906.2007.01507.x) [2906.2007.01507.x](https://doi.org/10.1111/j.1444-2906.2007.01507.x).
	- 61 G. Hayashida-Soiza, A. Uchida, N. Mori, Y. Kuwahara and Y. Ishida, J. Appl. Microbiol., 2008, 105(5), 1672–1677, DOI: [10.1111/j.1365-2672.2008.03878.x](https://doi.org/10.1111/j.1365-2672.2008.03878.x).
	- 62 J. Tebben, C. Motti, D. Tapiolas, P. Thomas-Hall and T. Harder, Mar. Drugs, 2014, 12, 2802–2815, DOI: [10.3390/](https://doi.org/10.3390/md12052802) [md12052802](https://doi.org/10.3390/md12052802).
	- 63 R. Preeta, S. Jose, S. Prathapan and K. K. Vijayan, Aquacult. Res., 2010, 41(10), 1452–1461, DOI: [10.1111/j.1365-](https://doi.org/10.1111/j.1365-2109.2009.02436.x) [2109.2009.02436.x](https://doi.org/10.1111/j.1365-2109.2009.02436.x).
	- 64 P. Rattanachuay, D. Kantachote, M. Tantirungkij, T. Nitoda and H. Kanzak, World J. Microbiol. Biotechnol., 2011, 27, 869–880, DOI: [10.1007/s11274-010-0529-x](https://doi.org/10.1007/s11274-010-0529-x).
	- 65 P. Priyaja, P. Jayesh, N. S. Correya, B. Sreelakshmi, N. S. Sudheer, R. Philip and I. S. B. Singh, J. Coastal Life Med., 2014, 2(1), 76–84, DOI: [10.12980/JCLM.2.2014J30](https://doi.org/10.12980/JCLM.2.2014J30).
	- 66 L. Zhang, X. Tian, S. Kuang, G. Liu, C. Zhang and C. Sun, Front. Microbiol., 2017, 8, 289, DOI: [10.3389/](https://doi.org/10.3389/fmicb.2017.00289) [fmicb.2017.00289](https://doi.org/10.3389/fmicb.2017.00289).
- 67 M. E. Teasdale, J. Liu, J. Wallace, F. Akhlaghi and D. C. Rowley, Appl. Environ. Microbiol., 2009, 75(3), 567– 572, DOI: [10.1128/AEM.00632-08](https://doi.org/10.1128/AEM.00632-08).
- 68 Y. Leyton, J. Borquez, J. Darias, M. Cueto, A. R. Díaz-Marrer and C. Riquelme, Mar. Drugs, 2011, 9(10), 2155–2163, DOI: [10.3390/md9102155](https://doi.org/10.3390/md9102155).
- 69 X. Y. Gao, Y. Liu, L. L. Miao, E. W. Li, T. T. Hou and Z. P. Liu, AMB Express, 2017, 7, 23, DOI: [10.1186/s13568-017-0323-3](https://doi.org/10.1186/s13568-017-0323-3).
- 70 P. Bhadury, B. Mohammad and P. C. Wright, J. Ind. Microbiol. Biotechnol., 2006, 33(5), 325–337, DOI: [10.1007/](https://doi.org/10.1007/s10295-005-0070-3) [s10295-005-0070-3](https://doi.org/10.1007/s10295-005-0070-3).
- 71 J. F. Imhof, Mar. Drugs, 2016, 14, 19, DOI: [10.3390/](https://doi.org/10.3390/md14010019) [md14010019](https://doi.org/10.3390/md14010019).
- 72 D. Tasdemir, Fungal Biol. Biotechnol., 2017, 4, 5, DOI: [10.1186/s40694-017-0034-1](https://doi.org/10.1186/s40694-017-0034-1).
- 73 H. J. Shin, Mar. Drugs, 2020, 18(5), 230, DOI: [10.3390/](https://doi.org/10.3390/md18050230) [md18050230](https://doi.org/10.3390/md18050230).
- 74 X.-Y. Hu, L.-M. Meng, X. Li, S.-Q. Yang, X.-M. Li and B.-G. Wang, Mar. Drugs, 2017, 15, 137, DOI: [10.3390/](https://doi.org/10.3390/md15050137) [md15050137](https://doi.org/10.3390/md15050137).
- 75 X.-C. Guo, L.-L. Xu, R.-Y. Yang, M.-Y. Yang, L.-D. Hu, H.-J. Zhu and F. Cao, Front. Chem., 2019, 7, 80, DOI: [10.3389/fchem.2019.00080](https://doi.org/10.3389/fchem.2019.00080).
- 76 A. Zhu, X.-W. Zhang, M. Zhang, W. Li, Z.-Y. Ma and H.-J. Zhu, Mar. Drugs, 2018, 16(9), 312, DOI: [10.3390/](https://doi.org/10.3390/md16090312) [md16090312](https://doi.org/10.3390/md16090312).
- 77 H. L. Li, X. M. Li, S. Q. Yang, L. H. Meng, X. Li and B. G. Wang, Mar. Drugs, 2019, 17(11), 605, DOI: [10.3390/](https://doi.org/10.3390/md17110605) [md17110605](https://doi.org/10.3390/md17110605).
- 78 X. D. Li, X. Li, X. M. Li, X. L. Yin and B. G. Wang, Nat. Prod. Res., 2021, 35(22), 4265–4271, DOI: [10.1080/](https://doi.org/10.1080/14786419.2019.1696792) [14786419.2019.1696792](https://doi.org/10.1080/14786419.2019.1696792).
- 79 X. Xu, S. Guo, H. Chen, Z. Zhang, X. Li, W. Wang and L. Guo, 3 Biotech, 2021, 11(4), 1–7, DOI: [10.1007/s13205-](https://doi.org/10.1007/s13205-021-02754-3) [021-02754-3](https://doi.org/10.1007/s13205-021-02754-3).
- 80 X. Wei, Y. Ding and F. An, Nat. Prod. Commun., 2021, 17(3), 1–5, DOI: [10.1177/1934578X221075986](https://doi.org/10.1177/1934578X221075986).
- 81 Y.-P. Feng, H.-K. Wang, J.-L. Wu, P. Shao, W.-L. Zhou, Q.-L. Lai, H.-W. Lin, B. Naman, T.-T. Wang and S. He, Chem. Biodiversity, 2022, 19(4), e202200028, DOI: [10.1002/](https://doi.org/10.1002/cbdv.202200028) [cbdv.202200028](https://doi.org/10.1002/cbdv.202200028).
- 82 T. Mai, F. Tintillier, A. Lucasson, C. Moriou, E. Bonno, S. Petek, K. Magre, A. A. Mourabit, D. Saulnier and C. Debitus, Lett. Appl. Microbiol., 2015, 61, 31–317, DOI: [10.1111/lam.12461](https://doi.org/10.1111/lam.12461).
- 83 B. C. Lee, A. Lee, J. H. Jung, S. H. Choi and T. S. Kim, Mol. Med. Rep., 2016, 14, 2691–2696, DOI: [10.3892/](https://doi.org/10.3892/mmr.2016.5522) [mmr.2016.5522](https://doi.org/10.3892/mmr.2016.5522).
- 84 H. Mohamad, R. Rosmiati, T. S. T. Muhammad, Y. Andriani, K. Bakar, N. Ismail, J. Saidin, J. Latip, N. Musa and A. Parenrengi, Nat. Prod. Commun., 2017, 12(8), 1227–1230, DOI: [10.1177/1934578X1701200819](https://doi.org/10.1177/1934578X1701200819).
- 85 T. Wang, J. Zou, T. Li, P. Shao, W. Zhou, Q. Lai, Y. Feng, C. B. Naman, X. Yan and S. He, Aquaculture, 2022, 549, 737727, DOI: [10.1016/j.aquaculture.2021.737727](https://doi.org/10.1016/j.aquaculture.2021.737727).
- 86 S. L. Holdt and S. Kraan, J. Phycol., 2011, 23, 543–597, DOI: [10.1007/s10811-010-9632-5](https://doi.org/10.1007/s10811-010-9632-5).
- 87 A. Leandro, L. Pereira and A. M. M. Gonçalves, Mar. Drugs, 2019, 18(1), 17, DOI: [10.3390/md18010017](https://doi.org/10.3390/md18010017).
- 88 M. Kladi, C. Vagias and V. Roussis, Phytochem. Rev., 2004, 3, 337–366, DOI: [10.1007/s11101-004-4155-9](https://doi.org/10.1007/s11101-004-4155-9).
- 89 M. T. Cabrita, C. Vale and A. P. Rauter, Mar. Drugs, 2010, 8, 2301–2317, DOI: [10.3390/md8082301](https://doi.org/10.3390/md8082301).
- 90 F. C. Pacheco, Mar. Drugs, 2010, 8, 1178–1188, DOI: [10.3390/md8041178](https://doi.org/10.3390/md8041178).
- 91 J. Cotas, A. Leandro, P. Monteiro, D. Pacheco, A. Figueirinha, A. M. M. Gonçalves, G. J. da Silva and J. Pereira, Mar. Drugs, 2020, 18, 384, DOI: [10.3390/](https://doi.org/10.3390/md18080384) [md18080384](https://doi.org/10.3390/md18080384).
- 92 H. Pereira, L. Barreira, F. Figueiredo, L. Custódio, C. V. Duarte, C. Polo, E. Rešek, A. Engelen and J. Varela, Mar. Drugs, 2012, 10, 1920–1935, DOI: [10.3390/](https://doi.org/10.3390/md10091920) [md10091920](https://doi.org/10.3390/md10091920).
- 93 S. Thanigaivel, N. Chadrasekaran, A. Mukherjee and J. Thomas, Aquaculture, 2015, 448, 82–86, DOI: [10.1016/](https://doi.org/10.1016/j.aquaculture.2015.05.039) [j.aquaculture.2015.05.039](https://doi.org/10.1016/j.aquaculture.2015.05.039).
- 94 I. N. Vatsos and C. Rebours, J. Appl. Phycol., 2015, 27, 2017– 2035, DOI: [10.1007/s10811-014-0506-0](https://doi.org/10.1007/s10811-014-0506-0).
- 95 S. Thanigaivel, N. Chadrasekaran, A. Mukherjee and J. Thomas, Aquaculture, 2016, 464, 529–536, DOI: [10.1016/](https://doi.org/10.1016/j.aquaculture.2016.08.001) [j.aquaculture.2016.08.001](https://doi.org/10.1016/j.aquaculture.2016.08.001).
- 96 V. Zammuto, M. G. Rizzo, A. Spanò, D. Spagnuolo, A. Di Martino, M. Morabito, A. Manghisi, G. Genovese, S. Guglielmino, G. Calabrese, F. Capparucci, C. Gervasi, M. S. Nicolo and C. Gugliandolo, Algal Res., 2022, 63, 102646, DOI: [10.1016/j.algal.2022.102646](https://doi.org/10.1016/j.algal.2022.102646). RSC Advances Article. Published on 5/2022. Downloaded on 5/2022. Downloaded on 5/2022. Downloaded on 5/29/2024 12:23:56 PM. Common Maccess Articles. 2008. The published is like the same of the same of the same of the same
	- 97 E. Shannon and N. Abu-Ghannam, Mar. Drugs, 2016, 14, 81, DOI: [10.3390/md14040081](https://doi.org/10.3390/md14040081).
	- 98 K. Wongprasert, T. Rudtanatip and J. Praiboon, Fish Shellfish Immunol., 2014, 36, 52-60, DOI: [10.1016/](https://doi.org/10.1016/j.fsi.2013.10.010) [j.fsi.2013.10.010](https://doi.org/10.1016/j.fsi.2013.10.010).
	- 99 N. Kasanah, T. Triyanto, D. S. Seto, W. Amelia and A. Isnansetyo, Indones. J. Chem., 2015, 15(2), 201–209, DOI: [10.22146/ijc.21215](https://doi.org/10.22146/ijc.21215).
	- 100 G. Brackman and T. Coenye, Curr. Pharm. Des., 2015, 21(1), 5–11, DOI: [10.2174/1381612820666140905114627](https://doi.org/10.2174/1381612820666140905114627).
	- 101 W. Zhao, N. Lorenz, K. Jung and S. A. Sieber, Angew. Chem., Int. Ed., 2016, 55(3), 1187-1191, DOI: [10.1002/](https://doi.org/10.1002/anie.201508052) [anie.201508052](https://doi.org/10.1002/anie.201508052).
	- 102 A. P. A. Wijnana, N. Kasanah and T. Triyanto, Nat. Prod. J., 2018, 8, 1–6, DOI: [10.2174/1573401313666170925161408](https://doi.org/10.2174/1573401313666170925161408).
	- 103 N. Kasanah, W. Amelia, A. Mukminin, T. Triyanto and A. Isnansetyo, Nat. Prod. Res., 2019, 33(22), 3303–3307, DOI: [10.1080/14786419.2018.1471079](https://doi.org/10.1080/14786419.2018.1471079).
	- 104 K. Karnjana, C. Soowannayan and K. Wongprasert, Fish Shellfish Immunol., 2019, 88, 91-101, DOI: [10.1016/](https://doi.org/10.1016/j.fsi.2019.01.058) [j.fsi.2019.01.058](https://doi.org/10.1016/j.fsi.2019.01.058).
	- 105 K. Karnjana, S. Nobsathian, C. Soowannayan, W. Zhao, Y. J. Tang and K. Wongprasert, Mar. Drugs, 2020, 18(2), 80, DOI: [10.3390/md18020080](https://doi.org/10.3390/md18020080).
	- 106 M. Reverter, N. Bontemps, D. Lecchini, B. Banaigs and P. Sasal, Aquaculture, 2014, 433, 50–61, DOI: [10.1016/](https://doi.org/10.1016/j.aquaculture.2014.05.048) [j.aquaculture.2014.05.048](https://doi.org/10.1016/j.aquaculture.2014.05.048).
- 107 P. Elumalai, A. Kurian, S. Lakshmi, C. Faggio, M. A. Esteban and E. Ringø, Rev. Fish. Sci., 2021, 29(1), 33–35, DOI: [10.1080/23308249.2020.1779651](https://doi.org/10.1080/23308249.2020.1779651).
- 108 E. Sanchez, S. Garcia and N. Heredia, Appl. Environ. Microbiol., 2010, 76, 6888–6894, DOI: [10.1128/AEM.03052-](https://doi.org/10.1128/AEM.03052-09) [09](https://doi.org/10.1128/AEM.03052-09).
- 109 M. Aminzare, M. Hashemi, Z. Abbasi, Z. M. Mohseni and E. Amiri, J. Appl. Pharm. Sci., 2018, 8, 170–177, DOI: [10.7324/JAPS.2018.8126](https://doi.org/10.7324/JAPS.2018.8126).
- 110 A. Borges, A. C. Abreu, C. Dias, M. J. Saavedra, F. Borges and M. Simões, Molecules, 2016, 21(7), 877, DOI: [10.3390/](https://doi.org/10.3390/molecules21070877) [molecules21070877](https://doi.org/10.3390/molecules21070877).
- 111 M. Snoussi, H. Hajlaoui, E. Noumi, D. Usai, L. A. Sechi, S. Zanetti and A. Bakhrouf, World J. Microbiol. Biotechnol., 2008, 24(12), 3071–3076, DOI: [10.1007/s11274-008-9848-6](https://doi.org/10.1007/s11274-008-9848-6).
- 112 S. Chatterjee, M. Asakura, N. Chowdury, S. B. Neogi, N. Sugimoto, S. Haldar, S. P. Awasthi, A. Hinenoya, S. Aoki and S. Yamasaki, FEMS Microbiol. Lett., 2010, 306(1), 54–60, DOI: [10.1111/j.1574-6968.2010.01931.x](https://doi.org/10.1111/j.1574-6968.2010.01931.x). Review Watch

107 P. Elimands, A. Kurian, S. Adelains, C. Pagelo, M. A. Elimands, S. Commons, S. Tankins, A. Chorontown, S. B. Regist

101 P. E. Suredies, A. G. Commons A. This article. Published on 5. This article is lic
	- 113 I. A. S. V. Packiavathy, P. Sasikumar, S. K. Pandian and A. V. Ravi, Appl. Microbiol. Biotechnol., 2013, 97, 10177– 10187, DOI: [10.1007/s00253-013-4704-5](https://doi.org/10.1007/s00253-013-4704-5).
	- 114 E. Acosta-Smith, N. Leon-Sicairos, S. Tiwari, H. Flores-Villasensor, A. Canizalez-Roman, R. Kumavath, P. Ghosh, V. Azevedo and D. Barh, Pathogen, 2019, 8, 64, DOI: [10.3390/pathogens8020064](https://doi.org/10.3390/pathogens8020064).
	- 115 Z. Shan, N. Guan, Y. Yang, T. Jin, X. Xia and W. Liu, Aquaculture, 2021, 533, 736109, DOI: [10.1016/](https://doi.org/10.1016/j.aquaculture.2020.736109) [j.aquaculture.2020.736109](https://doi.org/10.1016/j.aquaculture.2020.736109).