

# Use of probiotics in aquaculture of China—a review of the past decade

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## A B S T R A C T

China is the largest aquaculture producer in the world. Antibiotics were extensively used to ensure the development of the intensive aquaculture; however, the use of antibiotics causes safety- and environment-associated problems. As an alternative strategy to antibiotics, aquatic probiotics have attracted attention. The microbial organisms used as probiotics or tested as potential probiotics in Chinese aquaculture belong to various taxonomic divisions, including Actinobacteria, Bacteroidetes, Firmicutes, Proteobacteria and yeast. Moreover, the mixture of probiotic strains and synbiotics are also widely used. Studies on the mode of action of aquatic probiotics have extended our understanding of the probiotic effects, and novel mechanisms have been discovered, such as interference of quorum sensing. However, use of probiotics in Chinese aquaculture is still at an initial stage, and there are potential risks for some probiotic applications in aquaculture. Further regulation and management are required to normalize the production and usage of aquatic probiotics. In this review, we discuss species, effects, and mode of actions of probiotics in Chinese aquaculture since 2008. Challenges and future directions for research are also discussed.

## 1. Introduction

Aquaculture has a long history and started in China more than 2000 years ago [1]. It is a fast-growing and rapidly expanding industry, and it is reported that food fish provides an average of one-fifth of the total animal protein intake of the world population [2,3]. Aquaculture production in China accounts for nearly 70% of world aquaculture production, and the total production rose from 2.33 million tons in 1978 to 51.42 million tons in 2016, making China the only country in the world where aquaculture production exceeds the wild catch (FAO 2015). However, the intensive aquaculture enhanced the incidence of disease outbreaks in the farmed species, leading to massive mortalities and economic loss. For many cultured aquatic species in China, vaccines are not available. Moreover, vaccines will hardly be available in the near future due to the limitations in fish vaccine development [4]. For these species the only treatment of diseases has been through extensive use of chemical additives and veterinary medicines, especially antibiotics [5].

However, the abuse of antibiotics is linked to many secondary complications like modulation of the “normal” microbiota in the culture environment, antibiotic accumulation in edible products, and the continuously rising cost for disease treatment [6–8]. In addition, there is a general concern over the increased numbers of antibiotic resistant strains of bacteria in our environment [9]. To reduce the abuse of antibiotics in aquaculture, many alternative solutions have been suggested particularly when vaccines are not available. These include antimicrobial peptide, herb extracts, pro-, pre- and synbiotics, and among them, probiotics have received particular attention due to the high abundance, low cost and convenience in application [3].

The Food and Agricultural Organization (FAO) and World Health Organization (WHO) define probiotics as “live microorganisms which when administered in adequate amounts confer a health benefit on the host” [10]. However, the probiotics for aquatic usage are different from probiotics for terrestrial animals and human. The key factor that contributes to this difference is the intricate relationship aquatic animals

have with the ambient environment. Therefore, it is accepted that we must have a distinctive definition for aquatic probiotics as opposed to that for terrestrial animals. Many authors proposed definitions for probiotics in aquaculture [11–14]. The definitions were similar while each has its own specificity. Verschuere et al. suggested the definition of probiotics for aquatic animals as “a live microbial adjunct which has a beneficial effect on the host by modifying the host-associated or ambient microbial community, by ensuring improved use of the feed or enhancing its nutritional value, by enhancing the host response towards disease, or by improving the quality of its ambient environment” [12]. In this review, we would adhere to this definition, and will summarize live microbial adjuncts applied by dietary supplementation or as water additives in this review. Notably, some scientists define aquatic probiotics as an entire or component(s) of a microorganism [15,16]. While the functional components of probiotics are very important, it is more appropriate to define them as “probiotic effectors”, which cannot be considered as probiotics.

During the last decade, several reviews have summarized the use and beneficial effects of probiotics in aquaculture [17–20]. In particular, Qi et al. reviewed probiotic studies in Chinese aquaculture from 1988 to 2008 [18]. However, as numerous studies have been published on probiotics in Chinese aquaculture since 2008, an update is needed. Therefore, the current paper addressed the use of probiotics in Chinese aquaculture from 2008 and until today. Furthermore, issues associated with probiotics in Chinese aquaculture that merit further investigations are also discussed.

## 2. The microorganisms applied as probiotics in aquaculture of China

Probiotics used in Chinese aquaculture have a diverse origin. Most of them are either derived from the intestines of the host animals [21–24], or isolated from the environments [25–29]. When discussing probiotics, it is of interest to notice that some of the probiotic bacteria have been developed into commercial products [30–33] (Table 1). The microbial organisms used as probiotics or tested as potential probiotics in Chinese aquaculture belong to various taxonomic divisions, including Actinobacteria, Bacteroidetes, Firmicutes, Proteobacteria and yeast.

### 2.1. Actinobacteria

#### 2.1.1. *Arthrobacter*

*Arthrobacter* is a genus of Gram-negative, obligate aerobe rods or cocci, and they have been identified successfully from the gastrointestinal (GI) tract of several finfish species [34–36]. However, less information is available on probiotic *Arthrobacter*. In an early study with white shrimp (*Penaeus chinensis*) by Li et al. [37], an *Arthrobacter* XE-7 isolated from white shrimp showing antagonisms towards *Vibrio parahaemolyticus*, *Vibrio anguillarum* and *Vibrio nereis* was added to the rearing water. As this treatment revealed protection of shrimp post larvae against pathogenic vibrios, the authors suggested that *Arthrobacter* XE-7 could be probiotic in shrimp larvae culture. Furthermore, *Arthrobacter* sp. CW9 isolated from the gut of white shrimp (*Penaeus vannamei*) improved survival and growth rates, phenoloxidase activity, phagocytic activity and clearance efficiency of haemocytes of white shrimp when added to the rearing water during a breeding experiment [38].

### 2.2. Bacteroidetes

#### 2.2.1. *Flaviramulus*

*Flaviramulus ichthyenteri* Th78<sup>T</sup>, a novel species in family Flavobacteriaceae, was isolated from flounder (*Paralichthys olivaceus*) and showed strong quorum quenching (QQ) ability [39,40]. The genome of strain Th78<sup>T</sup> has been sequenced and genome information

suggests production of QQ enzyme, utilization of various nutrients available in the intestine, as well as production of digestive enzymes and vitamins, which suggests the prospect of Th78<sup>T</sup> to be used as a probiotic in aquaculture [39,40].

#### 2.2.2. *Flavobacterium*

Genus *Flavobacterium* is Gram-negative, rod-shaped, non-motile and motile bacteria, and they have been reported in the GI tract of several fish species [34,35,41–43], but several species are known to cause disease in fish [44,45]. *Flavobacterium sasangense* BA-3, an autochthonous intestinal strain isolated from common carp (*Cyprinus carpio*), has been suggested as probiotics [21]. In this study, dietary supplementation of bacterial cells or extracellular products of BA-3 positively influenced the innate immune parameters of carp (lysozyme, complement C3, total serum protein, albumin and globulin levels, respiratory burst activity, phagocytic activity by blood leucocytes and the expression of IL-1 $\beta$ , lysozyme-C, and TNF- $\alpha$  in blood). Moreover, both bacterial cells and extracellular products of *F. sasangense* BA-3 improved resistance of carp against *Aeromonas hydrophila* infection.

### 2.3. Firmicutes

#### 2.3.1. *Bacillus*

*Bacillus* species are one of the most widely used probiotics in aquaculture [15,46]. According to Gatesoupe's study, *Bacillus* strains were introduced as probiotics in aquaculture in the 90ties [46]. They can exert *in vitro* antagonistic or inhibitory activities against a variety of bacterial and fungal pathogens, provide digestive enzymes, activate growth factors, and regulate the immunity of aquatic animals [47–51].

Numerous studies have revealed that supplementation of *Bacillus* to the diet improved growth performance, digestive enzyme activities, antioxidant function, immune responses, and disease resistance of fish [28,52–58]. The *Bacillus* species used include *Bacillus subtilis* [59–63], *Bacillus cereus* [51,64,65], *Bacillus amyloliquefaciens* [56], *Bacillus baekryungensis* [66], *Bacillus licheniformis* [27,28,67], *Bacillus pumilus* [22,52,68], *Bacillus clausii* [52,69], *Bacillus coagulans* [25,26,29,58,70], and *Bacillus velezensis* [71].

*Bacillus* strains have also been used as probiotics for invertebrate aquatic species [26,60,72]. Four studies on sea cucumber (*Apostichopus japonicus*) revealed that *B. subtilis* T13, *B. baekryungensis* YD13, *B. cereus* G19, *B. cereus* BC-01, and *B. cereus* EN25 significantly enhanced growth performance, immunity and disease resistance of sea cucumber against *Vibrio splendidus* [61,64–66]. The commercial probiotic product BZT<sup>®</sup>, containing *B. subtilis* YB-1 (50%) and *B. cereus* YB-2 (50%), stimulated non-specific immune responses and enhanced growth performance of sea cucumbers, and was effective in controlling infection caused by *V. alginolyticus* [30]. Some studies using *B. subtilis* S12 and *Bacillus* PC465 revealed improved growth performance, non-specific immunity and disease-resistance of Pacific white shrimp (*Litopenaeus vannamei*) [73,74]. Two strains, *B. subtilis* DCU and *B. pumilus* BP, were reported to have potential application by controlling pathogenic vibriosis in mud crab (*Scylla paramamosain*) aquaculture [22].

Use of heat-inactivated bacilli has been investigated in two recent studies. Yan et al. demonstrated that heat-inactivated *B. pumilus* SE5 could effectively improve growth performance and remarkably up-regulated expression of TLR2 and pro-inflammatory cytokines (IL-8 and IL-1 $\beta$ ) in head kidney of grouper (*Epinephelus coioides*) [68]. More recently, Wang et al. revealed that heat-inactivated *B. clausii* DE5 improved feed utilization and the expression of TLR5, pro-inflammatory cytokines (IL-8 and IL-1) and TGF-1 in the head kidney and intestine of grouper [69].

Some *Bacillus* strains applied as water additives showed similar beneficial effects to the aquatic animals as those supplemented in the diet. Zhou et al. reported that probiotic, *B. coagulans* SC8168, supplemented as water additive at a certain concentration, could significantly increase survival rate and the activity of some digestive enzymes of

**Table 1**  
Bibliographic review of research published in the last 10 years (2008–2018) on the use of probiotics in aquaculture of China.

| Strains (source)  | Phylum         | Effective doses/mode of application                                      | Target host  | Duration | Probiotic effects   |  | Reference |
|---|----------------|--|--|----------|---|--|-----------|
|   |                |  |  |          | Disease resistance including immune improvements  | Nutritional modulation   |           |
|   |                |  |  |          | Productive effects and others   | Improvement of aquaculture environment   |           |
|   |                |  |  |          | Water   |  |           |
| <b>Gram-negative bacteria</b>   |                |  |  |          |   |  |           |
| <i>Aeromonas</i>  |                |  |  |          |   |  |           |
| <i>A. veronii</i> BA-1 (from common carp)                                   | Proteobacteria | Diets $10^8$ cell $g^{-1}$ (live)  | Common carp ( <i>Cyprinus carpio</i> )               | 28 days  | Serum protein, albumin, globulin ↑<br>Phagocytic capacity ↑<br>Transcriptional expression of IL-1 $\beta$ , TNF- $\alpha$ , lysozyme-C ↑<br>Mortality ↓ (80–66.66%) after <i>A. hydrophila</i> challenge ↓  |  | [21]      |
| <i>A. veronii</i> A-7 (from the intestinal tract of the healthy grass carp) | Proteobacteria | Diets $10^8$ cell $g^{-1}$ (live)  | Grass carp ( <i>Ctenopharyngodon idella</i> )        | 14 days  | Respiratory burst, phagocytic and lysozyme activities ↑<br>Serum complement C3, total serum protein, albumin, globulin ↑<br>Transcriptional expression of IL-8, IL-1, lysozyme-C, TNF- $\alpha$ ↑   | Potential pathogen bacteria ↓<br>Potential probiotics ↑<br>Cellulose-degrading intestinal bacteria ↑                                     | [23,154]  |
| <i>A. bivalvium</i> D15 (from the gut of healthy Pacific white shrimp)      | Proteobacteria | Diets $10^7$ cell $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 28 days  | Mortality ↓ (80–26.67%) after <i>A. hydrophila</i> challenge.<br>Respiratory burst and SOD activities ↑<br>Transcriptional expression of oprophenoloxidase, $\beta$ -1, 3-glucan-binding protein (LGBP) ↑<br>Mortality ↓ (43.33%–13.33%) after <i>V. harveyi</i> challenge. | Growth ↑   | [72]      |
| <i>Bdellovibrio</i>   |                |  |  |          |   |  |           |
| BA10 strain BDH12 (from sediment of Daya Bay in China)                      |                | Add to water at dose of $1 \times 10^8$ PFU $ml^{-1}$ (live)             | Abalone ( <i>Haliotis discus hannai</i> Ino)         | 90 days  |   | Growth ↑   | [119]     |
| <i>B. bacteriovorus</i> Bdm4 (from the sediment of fish ponds)              | Proteobacteria | Add to water at dose of $10^7$ PFU $ml^{-1}$ (live)                      | Grass carp ( <i>Ctenopharyngodon idellus</i> )       | 17 days  |   | Total cultivable Vibrio counts (TCVC) and bacterial counts (TCBC) ↓<br>Chemical oxygen demand (COD) ↓,<br>NH $_3$ -N ↓, sulphide ↓, DO ↑ | [120]     |
| <i>B. bacteriovorus</i> (commercial product)                                | Proteobacteria | Add to water at dose $5 \times 10^8$ PFU $m^{-3}$ (unidentified)         | Tilapia ( <i>Oreochromis nilotica</i> )              | 126 days | The death number of Tilapia caused by bacteriosis ↓   | The general bacterial population ↓<br>The number of coli group ↓   | [125]     |
| <i>B. bacteriovorus</i>   | Proteobacteria | Add to water with a final concentration of $5 \times 10^6$ PFU $ml^{-1}$ | White leg shrimp ( <i>Penaeus vannamei</i> )         | 7 days   | Control of shrimp pathogen <i>V. cholerae</i>   |  | [127]     |
| <i>Ectothiorhodospira</i>   |                |  |  |          |   |  |           |

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**Table 1 (continued)**

| Strains (source)  | Phylum         | Effective does/mode of application   | Target host   | Duration | Probiotic effects  | Disease resistance including immune improvements | Nutritional modulation | Productive effects and others   | Improvement of aquaculture environment  | Reference |
|---|----------------|--|---|----------|--|--|------------------------|---|---|-----------|
| <i>E. shaposhnikovii</i> WF (from marine shrimp pond)                           | Proteobacteria | Immersion twice at $10^9$ CFU ml <sup>-1</sup> (live)                        | Pacific white shrimp ( <i>Litopenaeus vannamei</i> )      | 10 days  |  |  |                        | Mortality ↓ (46%–74%)<br>Growth ↑   | Ammonia, nitrite, COD ↓                 | [129]     |
| <i>Flavobacterium</i>   | Proteobacteria | Diets $1 \times 10^8$ cell g <sup>-1</sup>                                   | Common carp ( <i>Cyprinus carpio</i> )                    | 28 days  | Lysozyme, complement C3, total serum protein, albumin and globulin, Respiratory burst and phagocytic activities ↑<br>Transcriptional expression of IL-1, lysozyme-C, and TNF-α ↑<br>Mortality rate ↓ (80%–66.66%) after <i>A. hydrophila</i> challenge.  |  |                        |   |   | [21]      |
| <i>Halomonas</i>  | Proteobacteria | Diets $3.68 \times 10^7$ or $7.18 \times 10^{10}$ CFU g <sup>-1</sup> (live) | Chinese whiter shrimp ( <i>Fenneropenaeus chinensis</i> ) | 6 weeks  | Hemocyte counts ↑, phenoloxidase activity ↑<br>Mortality after white spot syndrome virus challenge test ↓  |  |                        |   |   | [132]     |
| <i>Rhodopseudomonas</i>   | Proteobacteria | Immersion at $1 \times 10^7$ CFU ml <sup>-1</sup> (live)                     | Tilapia ( <i>Oreochromis niloticus</i> )                  | 40 days  | SOD and catalase (CAT) activities ↑  |  |                        | Growth ↑  |   | [25]      |
| <i>R. palustris</i> G06 (from a carp pond in Haining, China)                    | Proteobacteria | Incubation $1 \times 10^{11}$ CFU m <sup>-3</sup> (live)                     | Grass carp ( <i>Ctenopharyngodon idella</i> )             | 15 days  |  |  |                        |   | Ammonia ↓, nitrite ↓, COD ↓, nitrogen ↓ | [149]     |
| <i>Shewanella</i>   | Proteobacteria | Diets $10^9$ cell g <sup>-1</sup> (live)                                     | Abalone ( <i>Haliotis discus hannai</i> Ino),             | 4 weeks  | Haemocytes, respiratory burst activity, serum lysozyme activity, and total protein levels ↑<br>Mortality ↓ after <i>V. harveyi</i> challenge.  |  |                        |   |   | [153]     |
| <i>S. colwelliana</i> WA64<br><i>S.olleyana</i> WA65 (from GI tract of abalone) | Proteobacteria | Diets $10^9$ cell g <sup>-1</sup> (unidentified)                             | Sea cucumber ( <i>Apostichopus japonicus</i> )            | 28 days  | Total coelomocytes counts, respiratory burst activity, lysozyme, ACP, and phagocytic activities ↑<br>Respiratory burst, phagocytic and lysozyme activities ↑<br>Complement C3, total serum proteins, albumin and globulin ↑<br>Transcriptional expression of IL-8, IL-1, lysozyme-C, and TNF-α ↑<br>Mortality ↓ (80%–46.67%, 80%–33.33%) after <i>A. hydrophila</i> challenge. |  |                        | Mortality ↓ (97.6%–90.71%)  |   | [140]     |
| <i>S. japonica</i> HS7 (from the intestines of healthy sea cucumbers)           | Proteobacteria | Diets $10^8$ cells g <sup>-1</sup> of <i>S. xiamenensis</i> A-1 (live)       | Grass carp ( <i>Ctenopharyngodon idella</i> )             | 14 days  |  |  |                        | The abundance of the potential pathogenic bacteria ↓ (e.g., <i>Pseudomonas</i> and <i>Flavobacterium</i> genus), reproduction of potential probiotics ↓ (e.g., <i>Vibrio</i> , <i>Streptococcus</i> , and <i>Enterococcus</i> genus) and impact the abundance of cellulose-degrading bacteria ↑ (e.g., <i>Citrobacter</i> |   | [23]      |

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**Table 1 (continued)**

| Strains (source)  | Phylum         | Effective does/mode of application   | Target host  | Duration | Probiotic effects  | Productive effects and others                                     | Improvement of aquaculture environment | Reference |
|---|----------------|--|--|----------|--|---|--|-----------|
| <i>S. halitatis</i> D4 (from the gut of shrimp)   | Proteobacteria | Diets $10^7$ cell $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 28 days  | Respiratory burst and SOD activities $\uparrow$<br>Expression of prophenoloxidase, LPS- and LGBP protein $\uparrow$<br>Mortality $\downarrow$ (43.33%–33.33%) after <i>V. harveyi</i> challenge.   | genus) in the grass carp intestine.<br>Growth $\uparrow$          | Water                                  | [72]      |
| <i>Vibrio</i><br><i>V. tasmaniensis</i> HS10 (from the intestine of sea cucumber)                               | Proteobacteria | Diets $10^9$ cell $g^{-1}$ (unidentified)  | Sea cucumber ( <i>Apostichopus japonicus</i> )       | 28 days  | Total coelomocytes counts, respiratory burst, lysozyme, ACP, and phagocytic activities $\uparrow$ .  | Survival rate $\uparrow$ (87.60%–97.97%).                         |  | [140]     |
| <b>Gram-positive bacteria</b><br><i>Arthrobacter</i><br><i>Arthrobacter</i> sp. CW9 (from guts of white shrimp) | Actinobacteria | Added to the saline rearing water at $0, 10^5, 10^6$ and $10^7$ CFU $ml^{-1}$ every 5 days during the 24-day breeding experiment (unidentified)  | Pacific white shrimp ( <i>Penaeus vannamei</i> )     | 24 days  | Phenoloxidase, phagocytic and clearance efficiency $\uparrow$  | Survival rate $\uparrow$<br>Mean shrimp weights $\uparrow$        |  | [38]      |
| <i>Bacillus</i><br><i>Bacillus</i> PC465 (from the intestine of a healthy <i>Fenneropenaeus chinensis</i> )     | Firmicutes     | Diets $10^7$ and $10^9$ CFU $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 30 days  | Transcriptional expression of Pen-3a, peroxinectin, C-type lectin 3 ( <i>Lec-3</i> ), thioredoxin ( <i>Trx</i> ), prophenoloxidase ( <i>proPO</i> ) $\uparrow$<br>Survival rate $\uparrow$ (14.9%–45.2%, 50.7%) after white spot syndrome virus (WSSV) challenge.  | Growth $\uparrow$<br>Survival rate $\uparrow$ (35%–73.32%, 66.7%) |  | [74]      |
| <i>B. subtilis</i><br><i>B. amyloliquefaciens</i>   | Firmicutes     | Diets $10^5$ CFU $g^{-1}$ (live)   | Grass carp ( <i>Ctenopharyngodon idellus</i> )       | 52 days  | IgM, complement C3 and AKP activities $\uparrow$<br>Serum glutathione (GSH), total antioxidant capacity (T-AOC), SOD and glutathione peroxidase (GSH-Px) activities $\uparrow$<br>Malondialdehyde (MDA) and anti-superoxide anion (anti-O <sub>2</sub> <sup>-</sup> ) activity $\downarrow$<br>Globulin, IgM, myeloperoxidase and C3 activities $\uparrow$<br>T-AOC, anti-superoxide anion free radical (ASAFR) and GSH $\uparrow$ |   |  | [56]      |
| <i>B. licheniformis</i> <i>B. subtilis</i> (from the pond of grass carp)  | Firmicutes     | Group 1: added <i>Bacillus</i> preparation no. 1 with $10^6$ CFU $m^{-3}$ per 7 days in culture water (live)<br>Group 2: diet mixed with 0.5% <i>Bacillus</i> preparation no. 2, and the culture water | Grass carp ( <i>Ctenopharyngodon</i> )               | 4 weeks  |  |   |  | [28]      |

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**Table 1 (continued)**

| Strains (source)   | Phylum     | Effective does/mode of application   | Target host   | Duration           | Probiotic effects   | Disease resistance including immune improvements | Nutritional modulation | Productive effects and others   | Improvement of aquaculture environment | Reference    |
|--|------------|--|---|--------------------|---|--|------------------------|---|--|--------------|
| Probiotics product containing <i>B. subtilis</i> YB-1 (50%) and <i>B. cereus</i> YB-2 (50%)  | Firmicutes | was added $10^8$ CFU $m^{-3}$ <i>Bacillus</i> preparation no. 1 per 7 days (live)<br>Diets $10^{10}$ CFU $g^{-1}$ (live)   | Sea cucumbers ( <i>Apostichopus japonicus</i> )   | 32 days            | Phagocytic activity $\uparrow$ , superoxide anion production $\uparrow$ , lysozyme, catalase and phenoloxidase activities $\uparrow$<br>The cumulative mortality $\downarrow$ (100%–47%) after <i>V. alginolyticus</i> challenge.   |  |                        | Growth $\uparrow$   |  | [30]         |
| <i>B. pumilus</i><br><i>B. clausii</i> (from the gut of grouper <i>E. coioides</i> )   | Firmicutes | Diets $1 \times 10^8$ cell $g^{-1}$ (live)   | Grouper ( <i>Epinephelus coioides</i> )   | 60 days            | Phagocytic activity and lysozyme, complement C3, IgM and SOD activities $\uparrow$<br>Expression levels of CAT, proPO and SOD genes $\uparrow$<br>Respiratory burst activity $\uparrow$<br>Final survival rates $\uparrow$ (54.88%–76.67%, 78.33%) after <i>V. parahaemolyticus</i> challenge.                      |  |                        | Feed conversion ratio (FCR) $\uparrow$  |  | [52]         |
| <i>B. subtilis</i> DCU<br><i>B. pumilus</i> BP (from the intestine of mud crabs)   | Firmicutes | Diets $1 \times 10^5$ cell $g^{-1}$  | Juvenile mud crabs ( <i>Scylla paramamosain</i> )   | 30 days            |   |  |                        |   |  | [22]         |
| <i>B. baekryungensis</i> YD13 (from sea cucumber culturing ponds)<br><i>B. cereus</i> G19<br><i>B. cereus</i> BC-01 (from the intestine of the sea cucumber) | Firmicutes | Diets $1 \times 10^6$ CFU $g^{-1}$ (live)<br>Diets $1 \times 10^9$ CFU $kg^{-1}$ (live)  | Sea cucumber ( <i>Apostichopus japonicus</i> )<br>Sea cucumber ( <i>Apostichopus japonicus</i> Selenka) | 60 days<br>60 days | Lysozyme, ACP, AKP, SOD, CAT activities $\uparrow$  |  |                        | Growth $\uparrow$   |  | [66]<br>[64] |
| <i>B. cereus</i> EN25 (from mud of sea cucumber culturing water bodies)  | Firmicutes | Diets $1 \times 10^7$ CFU $g^{-1}$ (live)  | Juvenile sea cucumbers ( <i>Apostichopus japonicus</i> )  | 30 days            | Phagocytic, respiratory burst and AKP activities $\uparrow$<br>Transcriptional expression of Aj-p50, Aj-rel $\uparrow$<br>Phagocytosis, respiratory burst activity and total nitric oxide synthase activity $\uparrow$<br>The cumulative mortality $\downarrow$ (64.2%–33.3%) after <i>V. splendidus</i> challenge. |  |                        |   |  | [65]         |
| <i>B. cereus</i> (from the intestinal contents of tilapia)   | Firmicutes | Added to the water ( $1 \times 10^4$ CFU $ml^{-1}$ and $1 \times 10^5$ CFU $ml^{-1}$ ) or feed ( $1 \times 10^7$ cell $g^{-1}$ and $1 \times 10^8$ cell $g^{-1}$ ) | Tilapia ( <i>Oreochromis niloticus</i> )  | 42 days            | Serum lysozyme, peroxidase, AKP and total superoxide dismutase (TSOD) activities $\uparrow$   |  |                        | Affect the gut microbiota of tilapia and stimulated various potentially beneficial bacteria |  | [51]         |
| <i>B. clausii</i> DE5 (from the gut of juvenile <i>Epinephelus coioides</i> )  | Firmicutes | Diets supplemented with viable (T1) and heat-inactivated (T2) <i>B. clausii</i> DE5 at dose of $1 \times 10^6$ cell $g^{-1}$                                       | Grouper ( <i>Epinephelus coioides</i> )   | 60 days            | Serum lysozyme activity and complement C3 $\uparrow$<br>Transcriptional expression of TLR5, pro-inflammatory cytokines (IL-8, IL-1 $\beta$ ) and TGF- $\beta$ 1 $\uparrow$ in head kidney and intestine.  |  |                        | Feed intake $\downarrow$<br>FCR- $\downarrow$   |  | [69]         |
| <i>B. coagulans</i> (from the shrimp <i>Penaeus vannamei</i> ponds)  | Firmicutes | Diets viable (T-1) and dead (T-2) <i>B. coagulans</i> at dose of $1 \times 10^7$ cell $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> )  | 50 days            | Protease, amylase, and lipase $\uparrow$  |  |                        | Growth $\downarrow$<br>Survival rate $\downarrow$   |  | [26]         |

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**Table 1 (continued)**

| Strains (source)  | Phylum     | Effective does/mode of application   | Target host  | Duration           | Probiotic effects  | Productive effects and others  | Reference |
|---|------------|--|--|--------------------|--|--|-----------|
| <i>B. coagulans</i> (from common carp aquaculture ponds)  | Firmicutes | Diets $1 \times 10^7$ cell $g^{-1}$ , $2 \times 10^7$ cell $g^{-1}$ , $4 \times 10^7$ cell $g^{-1}$ (live) | Common carp ( <i>Cyprinus carpio</i> )                   | 45 days            | Lysozyme, myeloperoxidase, and respiratory burst activities ↑  | Growth ↑   | [29]      |
| <i>B. coagulans</i> , (from a commercial product provided by Suwei Microbial Research Co., Ltd (Jiangsu, China) that contained <i>B. coagulans</i> $2 \times 10^{10}$ CFU $g^{-1}$ ). | Firmicutes | Diets 125, 250, 500, and 1000 mg $kg^{-1}$ (live)  | Juvenile gibel carp ( <i>Carassius auratus gibelio</i> ) | 8 weeks            | Plasma lysozyme activity, superoxide dismutase, and superoxide anion radical scavenging activity ↑<br>Transcriptional expression of heat shock protein 70 (HSP70) ↑<br>The cumulative mortality ↓ after <i>A. hydrophila</i> challenge.  | Growth ↑<br>FCR ↓  | [58]      |
| <i>B. coagulans</i> SC8168 (from the pond sediment of shrimp)   | Firmicutes | Diets supplemented $1 \times 10^6$ CFU $ml^{-1}$ (live)  | Larvae shrimp ( <i>Penaeus vannamei</i> )                | ontogenetic stages |  | Survival rate ↑ (generally by 7.00–13.10%) over the controls.<br>Growth ↑  | [70]      |
| <i>B. coagulans</i> B16 (from a carp)   | Firmicutes | Added to the water of tanks at final concentration of $1 \times 10^7$ CFU $ml^{-1}$ every 2 days           | Tilapia ( <i>Oreochromis niloticus</i> )                 | 40 days            | SOD, CAT and respiratory burst activities ↑<br>Concentrations of serum protein and globulin ↑  |  | [25]      |
| <i>B. licheniformis</i> BSK-4 (from a grass carp culture pond)  | Firmicutes | Diets $3 \times 10^8$ CFU $m^{-3}$ (live)  | Grass carp ( <i>Ctenopharyngodon idellus</i> )           | 18 days            |  | Nitrite, nitrate and total nitrogen levels ↓, ammonia level ↑<br>Alterations of the microbial composition in grass carp water. | [27]      |
| <i>B. licheniformis</i> (commercial product, AlCare*, Zoetis, Shanghai, China)  | Firmicutes | Diets $4.4 \times 10^6$ CFU $g^{-1}$ (live)  | Juvenile Tilapia ( <i>Oreochromis niloticus</i> )        | 10 weeks           | Complement C3 ↑<br>Survival rate ↑   | Growth ↑   | [67]      |
| <i>B. pumilus</i> SE5 (from the gut of juvenile grouper <i>Epinephelus coioides</i> )   | Firmicutes | Diets live and heat-inactivated <i>B. pumilus</i> SE5 at dose of $1 \times 10^8$ CFU $g^{-1}$              | Grouper ( <i>Epinephelus coioides</i> )                  | 60 days            | Phagocytic activity, serum complement C3 and IgM levels, SOD activity ↑<br>Transcriptional expression of TLR2 and pro-inflammatory cytokines (IL-8 and IL-1β) ↑<br>Phenoloxidase and lysozyme activities ↑<br>Total antioxidant capacity, SOD, GSH-Px activity and superoxide anion activity ↑<br>Phagocytosis, respiratory burst activity and total nitric oxide synthase (i-NOS) activities ↑<br>The cumulative mortality ↓ (56.2%–20.0%) after <i>V. splendens</i> challenge. | Growth ↑   | [68]      |
| <i>B. subtilis</i> (from the gut contents of farm-reared shrimp)  | Firmicutes | Diets <i>B. subtilis</i> at dose of $5 \times 10^4$ CFU $g^{-1}$ (live)                                    | Pacific white shrimp ( <i>Litopenaeus vannamei</i> )     | 40 days            |  | Growth ↑   | [60]      |
| <i>B. subtilis</i> T13 (from intestine of healthy sea cucumbers)  | Firmicutes | Diets $1 \times 10^9$ CFU $g^{-1}$ (live)  | Juvenile sea cucumbers ( <i>Apostichopus japonicus</i> ) | 30 days            |  | Growth ↑   | [61]      |

(continued on next page)

**Table 1** (continued)

| Strains (source)  | Phylum     | Effective does/mode of application   | Target host  | Duration | Probiotic effects | Disease resistance including immune improvements  | Nutritional modulation  | Productive effects and others             | Improvement of aquaculture environment                  | Reference |
|---|------------|--|--|----------|-------------------|---|---|---|---|-----------|
| <i>B. subtilis</i> SCO2 (from a pond containing grass carp)   | Firmicutes | Diets $1 \times 10^9$ CFU $m^{-3}$ per 7 days in culture water (live)                    | Grass carp ( <i>Ctenopharyngodon idellus</i> )       | 15 days  |                   |   |   |   | Ammonia, nitrite and total nitrogen ↓                   | [62]      |
| <i>B. subtilis</i> FY99-01 (commercial probiotic)   | Firmicutes | Add to water with $5 \times 10^4$ CFU $mL^{-1}$ (live)                                   | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 56 days  |                   |   |   |   | Levels of pH, nitrite and soluble reactive phosphorus ↓ | [32]      |
| <i>B. subtilis</i> B2 (from sediment in healthy sea cucumber <i>A. postichopus japonicus</i> )              | Firmicutes | Diets supplemented according to the weight of sea cucumber by 0.1%, 0.2% and 0.3% (live) | Sea cucumbers ( <i>Apostichopus japonicus</i> )      | 7 days   |                   | ● AKP, TSOD and PO activities ↑   | Activity of amylase and protease ↑  |   |   | [59]      |
| <i>B. subtilis</i> HAINUP40 (from natural pond water in Hainan University)                                  | Firmicutes | Diets $10^8$ CFU $g^{-1}$ (live)   | Tilapia ( <i>Oreochromis nilotica</i> )              | 4 weeks  |                   | Respiratory bursts, serum lysozyme, T-AOC and SOD activities ↑  | protease and amylase activities ↑<br>The relative percent survival (RPS%) ↓   | Growth ↑                                  |   | [63]      |
| <i>B. subtilis</i> FY99-01 (from the gastrointestinal tract (GIT) of healthy grass carp)                    | Firmicutes | Diets $10^9$ CFU $g^{-1}$ (live)   | Crucian ( <i>Carassius auratus</i> )                 | 4 weeks  |                   | Activities of ACP, AKP, and GSH-PX ↑<br>Transcriptional expression of IFN- $\gamma$ , TNF- $\alpha$ , IL-1, IL-4, IL-10 ↓ and IL-12↓<br>Survival rate ↑ (30%–80%) after <i>A. hydrophila</i> challenge.   |   |   |   | [71]      |
| <i>Clostridium</i> Miyarisan  | Firmicutes | Diets $1.0 \times 10^8$ CFU $g^{-1}$ (live)  | Silver pomfret ( <i>Pampus argenteus</i> )           | 60 days  |                   |   | Lipase, protease and Amylase activities ↑   | Growth ↑                                  |   | [31]      |
| <i>C. butyricum</i> (from Pharmaceutical Co. Ltd, Tokyo, Japan)   | Firmicutes | Diets $1.0 \times 10^9$ CFU $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 56 days  |                   | T-AOC, lysozyme activity, inducible nitric oxide synthase (iNOS) activity ↑<br>Transcriptional expression of heat shock protein 70 (HSP70), Toll and immune deficiency (Imd) ↓<br>Serum phenoloxidase and acid phosphatases activities ↑, immunoglobulin M ↑<br>Transcriptional expression of prophenoloxidase, lipopolysaccharide and $\beta$ -1,3-glucan binding protein, lysozyme, crustin, and superoxide dismutase ↑ | Intestine SCFA content ↑ and body crude protein content ↓, modulated intestine digestive capacity ↑                     | Growth ↑<br>Intestine epithelium height ↑ |   | [33]      |
| <i>C. butyricum</i> (from healthy chicken intestine)  | Firmicutes | Diets $1.0 \times 10^3$ , $1.0 \times 10^5$ CFU $g^{-1}$ (live)                          | <i>Mitichthys mityi</i>                              | 8 weeks  |                   |   |   | Growth ↑                                  |   | [85]      |
| <i>C. butyricum</i> (from Zhongke Biotic Co., Ltd, China)   | Firmicutes | Diets $5.0 \times 10^5$ CFU $kg^{-1}$ (live)   | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 56 days  |                   |   | Transcriptional expression of $\alpha$ -amylase, lipase, trypsin, fatty acid-binding protein, and fatty acid synthase ↑ |   |   | [87]      |
| <i>C. butyricum</i> CBG01 (provided by the Microbial Culture Collection Center, Lab of Aquaculture Ecology, | Firmicutes | Diets $10^{11}$ and $10^{12}$ CFU $g^{-1}$ (live)  | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 42 days  |                   |   | AKP, ACP, lysozyme and total nitric oxide synthase (TNOS) activities ↑<br>The respiratory burst activity of hemolymph ↑ |   |   | [88]      |

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**Table 1 (continued)**

| Strains (source)  | Phylum     | Effective does/mode of application                              | Target host                                     | Duration | Probiotic effects  | Nutritional modulation   | Productive effects and others | Improvement of aquaculture environment | Reference |
|---|------------|---|---|----------|--|--|-------------------------------|--|-----------|
| Ocean University of China   |            |   |   |          |  |  |                               |  |           |
| <i>C. butyricum</i> (from Zhongke Biotic Co., Ltd, China)                                       | Firmicutes | Diets $1.0 \times 10^9$ CFU g <sup>-1</sup> (live)              | Kuruma shrimp ( <i>Marsupenaeus japonicus</i> ) | 56 days  | Transcriptional expression of Toll, Imd and Relish ↑<br>The cumulative mortality ↓ after <i>V. parahaemolyticus</i> challenge.<br>T-AOC, catalase, peroxidase activities ↑, MDA content ↑<br>Transcriptional expression of HSP70, ferritin and metallothionein ↑ | Amylase, lipase, alanine aminotransferase and aspartate aminotransferase activities ↑<br>Intestine SCFA and body crude protein content ↑ |                               |  | [89]      |
| <i>C. butyricum</i> (from Zhongke Biotic Co., Ltd, China)                                       | Firmicutes | Diets $10^8$ CFU g <sup>-1</sup> (live)                         | Kuruma shrimp ( <i>Marsupenaeus japonicus</i> ) | 56 days  |  |  |                               |  | [90]      |
| <i>Enterococcus faecium</i> MM4 (from the gut of healthy juvenile <i>Epinephelus coioides</i> ) | Firmicutes | Diets $1.0 \times 10^8$ CFU g <sup>-1</sup> (live)              | Grouper ( <i>Epinephelus coioides</i> )         | 60 days  | Complement C3 levels ↑   | Activity of hepatopancreatic protease, intestinal lipase ↓ and amylase ↑   |                               |  | [94]      |
| <i>E. faecalis</i> LC3 (from intestine of marine animals)                                       | Firmicutes | Diets $1.0 \times 10^9$ CFU g <sup>-1</sup> (live)              | Sea cucumber ( <i>Apostichopus japonicus</i> )  | 30 days  | AKP activity ↑<br>Transcriptional expression of NF-κappa-B transcription factor p65 (Rel), CASP2, HSP 90, HSP70 ↓ at 15 d and ↑ at 30 d.<br>The survival rate ↑ (48.1%–61.1%) after <i>V. splendidus</i> challenge.  |  | Growth ↑                      |  | [98]      |
| <i>E. faecium</i> NRW-2 (from the intestine of health <i>Acanthogobius hasta</i> )              | Firmicutes | Diets $1.0 \times 10^7$ CFU g <sup>-1</sup> (live)              | White shrimp ( <i>Penaeus vannamei</i> )        | 4 weeks  |  | Activities of hepatopancreatic, intestinal lipase and amylase ↑  |                               |  | [96]      |
| <i>Lactobacillus pentosus</i> HC-2 (from the intestine of health <i>Acanthogobius hasta</i> )   | Firmicutes | Diets $1.0 \times 10^7$ CFU g <sup>-1</sup> (live)              | White shrimp ( <i>Penaeus vannamei</i> )        | 4 weeks  |  | Hepatopancreatic, intestinal lipase and amylase activities ↑   |                               |  | [96]      |
| <i>L. plantarum</i> L-137 (HK L-137) (from the Zhuanghe Jintuo Aquaculture Farming)             | Firmicutes | Diets 0.05 g and 0.25 g HK L-137 kg <sup>-1</sup> (heat-killed) | Sea cucumber ( <i>Apostichopus japonicus</i> )  | 60 days  | Lysozyme, phagocytic, SOD and AKP activities ↑   |  | Growth ↑                      |  | [109]     |
| <i>L. plantarum</i> CCFM639 (from the Culture Collections)                                      | Firmicutes | Diets $10^8$ CFU g <sup>-1</sup> (live)                         | Tilapia ( <i>Oreochromis nilotica</i> )         | 4 weeks  |  |  | Growth ↑                      |  | [110]     |

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**Table 1** (continued)

| Strains (source)   | Phylum     | Effective doses/mode of application   | Target host  | Duration | Probiotic effects   | Disease resistance including immune improvements | Nutritional modulation | Productive effects and others         | Improvement of aquaculture environment | Reference |
|--|------------|---|--|----------|---|--|------------------------|---------------------------------------|--|-----------|
| of Food Microbiology, Jiangnan University)   |            |   |  |          |   |  |                        |                                       | Water                                  |           |
| <i>L. plantarum</i> CCFM8661 (from the Culture Collections of Food Microbiology, Jiangnan University)  | Firmicutes | Probiotic was administered at $10^8$ CFU $g^{-1}$ in fish diet twice daily (live) | Tilapia ( <i>Oreochromis nilotica</i> )              | 4 weeks  | Al-induced oxidative stress, recovered the activities of TAOC, SOD, CAT and GPx. Pb-induced oxidative stress, blood daminelevulinic acid dehydratase activity $\uparrow$ , reversed alterations in innate immune status $\uparrow$ . The frequencies of the nuclear abnormalities in peripheral blood erythrocytes. |  |                        | Pb accumulation $\downarrow$          |  | [111]     |
| <i>L. delbrueckii</i> (from Angel Company, Wuhan, China)   | Firmicutes | Diets $1.0 \times 10^6$ and $1.0 \times 10^7$ CFU $g^{-1}$ (live)                 | Huanghe carp ( <i>Cyprinus carpio</i> Huanghe var)   | 8 weeks  | Transcriptional expression of TNF- $\alpha$ , IL-8, IL-1, NF- $\kappa$ B p65, IL-10 and TGF- $\beta$ $\uparrow$   |  |                        |                                       |  |           |
| <i>L. plantarum</i> (commercial probiotics)  | Firmicutes | Cell-free extract of <i>L. plantarum</i> (dead)                                   | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 45 days  | The survival rate $\uparrow$ after <i>A. hydrophila</i> challenge. Significantly improved the resistance of <i>L. vannamei</i> against the stress of acute low salinity, as indicated by higher survival rate as well as higher transcript levels of ProPo, SOD and Lys gene.                                       |  |                        |                                       |  | [106]     |
| <i>L. plantarum</i> (provide by Xinhailisheng Biological Technology Co., Ltd., South China Sea Fisheries Research Institute, Chinese Academy of Fishery Sciences, China) | Firmicutes | Cell-free extract of <i>L. plantarum</i> (dead)                                   | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 15 days  |   |  |                        |                                       |  | [107]     |
| <i>L. plantarum</i> LL11 (from marine fish)  | Firmicutes | Diets $1.0 \times 10^9$ CFU $g^{-1}$ (live)                                       | Sea cucumber ( <i>Apostichopus japonicus</i> )       | 30 days  | AKP, ACP and SOD activities $\uparrow$  |  |                        | Growth $\uparrow$<br>FCR $\downarrow$ |  | [98]      |
| <i>L. penosus</i> (from shrimp farming pond in Zhangpu, Fujian province, China)  | Firmicutes | Diets $1.0 \times 10^7$ CFU $g^{-1}$ (live)                                       | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 28 days  | Heat shock proteins (HSP60, HSP70 and HSP90) $\uparrow$<br>The survival rate $\uparrow$ (48.1%–66.7%) after <i>V. splendidus</i> challenge.<br>The cumulative mortality $\downarrow$ after <i>Vibrio</i> strains challenge.   |  |                        | Growth $\uparrow$                     |  | [108]     |
|  | Firmicutes | Diets $10^5$ CFU $g^{-1}$ and $10^5$ CFU $g^{-1}$ (live)                          | Abalone ( <i>Haliotis discus hannai</i> Ino),        | 8 weeks  | The total number of blood lymphocytes, lysozyme   |  |                        |                                       |  | [112]     |

(continued on next page)

Table 1 (continued)

| Strains (source)  | Phylum         | Effective does/mode of application  | Target host   | Duration | Probiotic effects  | Nutritional modulation                 | Productive effects and others                               | Improvement of aquaculture environment  | Reference |
|---|----------------|---|---|----------|--|--|---|---|-----------|
| <i>L. pentosus</i> (from the intestinal tracts of healthy abalone)  |                |   |   |          | Disease resistance including immune improvements   |  |   | Water   |           |
| <i>Lactococcus</i><br><i>L. lactis</i> MM1 (from the gut of juvenile grouper<br><i>Epinephelus coioides</i> ) | Firmicutes     | Diets $1.0 \times 10^8$ CFU g <sup>-1</sup> (live)  | Juvenile grouper ( <i>Epinephelus coioides</i> )        | 60 days  | activity, acid phosphatase, and SOD ↓. Expression levels of peroxidase (TPx) ↑, MDA content ↓<br>The cumulative mortality rate ↓   | Hepatopancreatic protease activities ↑ | Survival rate, Shell length-specific growth rate, and FCR ↑ |   | [95]      |
| <i>L. lactis</i> RQ516 (from fresh milk)  | Firmicutes     | Add to water at final concentration of $1 \times 10^7$ CFU ml <sup>-1</sup> every 2 days (live) | Tilapia ( <i>Oreochromis niloticus</i> )                | 40 days  | Concentrations of serum protein and globulin ↑<br>Respiratory burst activity ↑<br>lysozyme content (LC) ↑,<br>myeloperoxidase (MPO) ↑ and<br>SOD activities ↑  |  | Growth ↑  |   | [114]     |
| <i>L. lactis</i> LH8 (from marine fish)   | Firmicutes     | Diets $1.0 \times 10^9$ CFU g <sup>-1</sup> (live)  | Sea cucumber ( <i>Apostichopus japonicus</i> )          | 30 days  | The Rel, HSP90 and HSP60 were down-regulated firstly at 15 d and up-regulated at 30 d, and only CASP2 showed up-regulation at both timepoints with 5.86 fold at 15 d and 10.29 fold at 30 d.<br>The survival rate (48.1%–64.8%) ↑ after of <i>V. splendidus</i> challenge. |  | Growth ↑  |   | [98]      |
| <i>Paracoccus</i><br><i>P. marcusii</i> DB11  | Proteobacteria | Diets $1.0 \times 10^8$ CFU g <sup>-1</sup> (live)  | Juvenile sea cucumber ( <i>Apostichopus japonicus</i> ) | 60 days  | SOD, CAT, lysozyme, ACP, and alkaline phosphatase (ALP) activities ↑<br>The cumulative mortality ↓ (40%–0%) after <i>V. splendidus</i> challenge.  |  |   |   | [64,134]  |
| <i>Pseudomonas</i><br><i>P. stutzeri</i> SC221-M (from a grass carp pond)                                     | Proteobacteria | Diets $3.0 \times 10^9$ CFU g <sup>-1</sup> (live)  | Grass carp ( <i>Ctenopharyngodon idellus</i> )          | 6 days   |  |  |   | Improve both the water quality and microbial community structure in experimental aquaculture system.    | [136]     |
| <i>P. stutzeri</i> F11 (from a grass carp pond)   | Proteobacteria | Add to water at dose of $1 \times 10^5$ CFU ml <sup>-1</sup> (dead)                             | Grass carp ( <i>Ctenopharyngodon idellus</i> )          |          |  |  |   | Ammonia-N <sub>i</sub> , nitrite-N <sub>i</sub> , and total N levels, and alter the microbial community | [137]     |

(continued on next page)

**Table 1 (continued)**

| Strains (source)  | Phylum         | Effective does/mode of application                                | Target host  | Duration | Probiotic effects  | Productive effects and others  | Improvement of aquaculture environment | Reference |
|---|----------------|---|--|----------|--|--|--|-----------|
| <i>Pseudoalteromonas</i><br><i>P. monas</i> sp. BC228 (from healthy <i>A. japonicus</i> )   | Proteobacteria | Diets $10^5$ , $10^7$ and $10^9$ CFU $g^{-1}$ (live)              | Juvenile sea cucumber ( <i>Apostichopus japonicus</i> )  | 45 days  | Lyozyme and phenoloxidase activities ↑<br>Mortality ↓<br>The mortality ↓ (100%–11%) after <i>V. alginolyticus</i> challenge.   | Intestinal trypsin and lipase activities ↑   | Water<br>structure of the water.       | [141]     |
| <i>P. piscicida</i> SW-1 (from seawater from a clam farm)   | Proteobacteria | Add to water at $10^7$ , $10^8$ , or $10^9$ CFU $ml^{-1}$ (live)  | Clams  | 15 days  |  |  |  | [142]     |
| <i>Psychrobacter</i><br><i>Psychrobacter</i> sp. (from the gut of <i>Epinephelus coioides</i> )                                   | Proteobacteria | Diets $1.0 \times 10^8$ CFU $g^{-1}$ (live)                       | Grouper ( <i>Epinephelus coioides</i> )                  | 60 days  | Phagocytic activity and phagocytic index ↑   | Hepatopancreatic protease and lipase activities ↑<br>Intestinal amylase activity ↑ |  | [144]     |
| <i>Weissella</i><br><i>Weissella confusa</i> LS13   | Firmicutes     | Diets $1.0 \times 10^9$ CFU $g^{-1}$ (live)                       | Sea cucumber ( <i>Apostichopus japonicus</i> )           | 30 days  | ACP activity, HSP60 and Rel ↑<br>Survival rate ↑<br>(48.1%–63.0%) after <i>V. splendidus</i> challenge.  | Growth ↑   |  | [98]      |
| <b>Yeast</b><br><i>Hanseniaspora opuntiae</i> C21 (from the intestine of healthy sea cucumber)                                    | Ascomycota     | Diets $1.0 \times 10^4$ CFU $g^{-1}$ (live)                       | Juvenile sea cucumbers ( <i>Apostichopus japonicus</i> ) | 50 days  | Phagocytic, coelomocytes, lysozyme, phenoloxidase, T-NOS, SOD, AKP and ACP activities ↑<br>The cumulative mortality ↓ (50%–0%) after <i>V. splendidus</i> challenge.   | Trypsin and lipase activities ↑  |  | [218]     |
| <i>Meischnikowia</i> sp. C14 (from the intestine of sea cucumber <i>Apostichopus japonicus</i> )                                  | Ascomycota     | Diets $10^4$ , $10^5$ , $10^6$ CFU $g^{-1}$ (live)                | Sea cucumber ( <i>Apostichopus japonicus</i> )           | 45 days  | Phagocytic, lysozyme, phenoloxidase, T-NOS, SOD and AKP activities ↑<br>The cumulative mortality ↓ (30%–0%) after <i>V. splendidus</i> challenge.  | Growth ↑   |  | [164]     |
| <i>Rhodotorula benthamica</i> D30 (from local sea mud)  | Basidiomycota  | Diets $1.0 \times 10^6$ and $1.0 \times 10^7$ CFU $g^{-1}$ (live) | Juvenile sea cucumber ( <i>Apostichopus japonicus</i> )  |          | Lyozyme, phagocytic and total nitric oxide synthase activities ↑   | Amylase, cellulase activity and alginase activities ↑                              |  | [167]     |
| <i>Rhodotorula</i> sp. C11 (from the intestine of healthy sea cucumbers collected from the waters around Bailanzi, Dalian, China) | Basidiomycota  | Diets $1.0 \times 10^4$ CFU $g^{-1}$ (live)                       | Sea cucumber ( <i>Apostichopus japonicus</i> )           |          | The cumulative prevalence and mortality of sea cucumbers fed diets supplemented with <i>Rhodotorula</i> sp. C11 were zero, which were lower than those of animals fed the control diet. In control animals, the prevalence of disease was 70% and the resulting mortality was 30%. | Growth ↑   |  | [166]     |
|   | Basidiomycota  |   |  | 2 weeks  |  | Growth ↑   |  | [169]     |

(continued on next page)

Table 1 (continued)

| Strains (source)  | Phylum        | Effective does/mode of application   | Target host  | Duration | Probiotic effects   | Productive effects and others | Improvement of aquaculture environment | Reference |
|---|---------------|--|--|----------|---|-------------------------------|--|-----------|
| <i>Rhodospiridium pallidigenum</i> (from coastal water at Zhanjiang, China)                   |               | Dry yeast at 1 g/100 g diet and (b) live yeast at $10^8$ yeast cells (hemocytometer counts)/1 g diet                                 | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) |          | Transcriptional expression of manganese superoxide dismutase (SODMn) and CAT          |                               | Water                                  |           |
| <i>Rhodospiridium pallidigenum</i> (from coastal water at Zhanjiang, China r)                 | Basidiomycota | Diets dry yeast at 1 g/100 g diet and live yeast at $10^8$ yeast cells (hemocytometer counts)/1 g diet.                              | Pacific white shrimp ( <i>Litopenaeus vannamei</i> ) | 6 weeks  | actin ↑<br>T-AOC, CAT, SOD and GPX activities ↑                                       | Growth ↑                      |  | [168]     |
| <i>Saccharomyces cerevisiae</i> (obtain as commercial preparation, Actisaf (Lesaffre, France) | Ascomycota    | Diet was incorporated with 1 g kg <sup>-1</sup> of the yeast for a final concentration of $10^7$ CFU g <sup>-1</sup> of feeds (live) | Tilapia ( <i>Oreochromis niloticus</i> )             | 8 weeks  | Transcriptional expression of HSP70, TNF $\alpha$ and TNF $\beta$ ↓<br>AKP activity ↓ | Growth ↑                      |  | [159]     |

shrimp larvae [70]. Zhou et al. demonstrated that the superoxide dismutase (SOD) activity of tilapia (*Oreochromis mossambicus*) was increased significantly after treatment with *B. subtilis* B10 and *B. coagulans* B16 as water additives [25]. Furthermore, *Bacillus* has also been frequently reported to improve the water quality when applied as water additives. The *Bacillus* genus has nitrification and de-nitrification functions [75–77]. Liang et al. reported that supplementation of *B. licheniformis* BSK-4 removed nitrogen and modulated the microbial community in grass carp (*Ctenopharyngodon idellus*) pond water [27]. Wu et al. showed that *B. subtilis* FY99-01 improved the water quality by reducing the levels of pH, nitrite and soluble reactive phosphorus when added to the culture water of Pacific white shrimp. In addition, weekly addition of FY99-01 showed influence on the bacterial community of culture water. It increased the abundance of beneficial microalgae (Bacillariophyta and Chlorophyta), and decreased the abundance of Vibrionaceae on day 84 post initial addition [32].

### 2.3.2. *Clostridium butyricum*

*Clostridium butyricum* is Gram-positive, obligate anaerobic and endospore-forming probiotic, which can provide short-chain fatty acids (SCFAs) especially butyric acid for the regeneration and repair of the intestinal epithelium, and regulation of intestine micro-ecological environment [78,79]. Besides, *C. butyricum* can also adapt to lower pH and higher temperatures and is resistant to many antibiotics [80,81]. *C. butyricum* has been used as probiotic for the prevention of human and veterinary intestinal diseases [82], as well as in some aquaculture studies [33,83–85]. *C. butyricum* and its culture supernatants have preventive and therapeutic effect on *Salmonella enteritidis* and *V. parahaemolyticus* infection in fish intestinal epithelial cells (FIECs) [83], and heat-killed *C. butyricum* retain interesting immunomodulating properties in *Miichthys miiuy* [86]. Furthermore, diets supplemented with *C. butyricum* improved growth performance, immune response, and digestive enzyme activity in silver pomfret (*Pampus argenteus*) [84]. Later, Duan et al. reported that dietary *C. butyricum* had beneficial effect on the intestine health of *L. vannamei*, and promoted growth, increased epithelial cells height and SCFA content, enhanced intestine immune function against ammonia stress in *L. vannamei* [33,87,88]. Furthermore, they also revealed that *C. butyricum* could improve the growth performance, increase intestine antioxidant capacity of *M. japonicus* against high temperature stress [89], and modulate intestine digestive and metabolic capacities, improve intestine SCFA content and body crude protein content in *M. japonicus* [90].

### 2.3.3. *Enterococcus*

*Enterococcus faecium* strain has been proposed as probiotics for human use due to the antagonistic effect on enteroaggregative *Escherichia coli* [91]. Sun et al. isolated an *E. faecium* MM4 from the whole intestinal tract of healthy juvenile grouper [92]. This strain exhibited *in vitro* antagonism against *Vibrio metschnikovi*, *Vibrio harveyi* and *Staphylococcus aureus*, and improved the feed efficiency and immune response of groupers when added in feed [93–95]. Sha et al. revealed that the amylase and lipase activities were significantly increased in the intestines and hepatopancreas of *L. vannamei* fed with *E. faecium* NRW-2 [96]. Cui et al. identified two strains, *E. faecalis* LS1-2 and *E. faecium* Z1-2, which showed significant antimicrobial activities against shrimp pathogens and were suggested to be good candidates of shrimp probiotics [97]. Recently, the probiotic strain *E. faecalis* LC3 isolated from marine fish significantly improved the alkaline phosphatase (AKP) activity and up-regulated transcriptional expression of Rel, CASP2, HSP 90, HSP70 of sea cucumber [98].

### 2.3.4. *Lactic acid bacteria (LAB)*

Bacteria belonging to LAB are classified in phylum Firmicutes, class Bacilli and order Latobacillales. They can be isolated from intestine of numerous fish species, and their potential as probiotics was recently discussed in the comprehensive review of Ringø et al. [99].

2.3.4.1. *Lactobacillus lactis* *Lactobacillus* supplementation was shown to improve growth and disease resistance of tilapia, and was recommended for commercial aquaculture production systems [100]. Wang et al. indicated that adding *Lactobacillus* into basal diet promoted growth performance, increased digestive enzyme activities, and enhanced non-specific immunity in shrimp [101]. Huang et al. reported that dietary supplementation of *Lactobacillus* improved antioxidant activity, immune capacity (serum acid phosphatase and lysozyme activity) and resistance of tilapia against *A. hydrophila* in a challenge trial [102]. Dietary optimal levels of *Lactobacillus delbrueckii* at  $10^6$ – $10^7$  CFU g<sup>-1</sup> could effectively enhance immunity, disease resistance against *A. hydrophila* antioxidant capability and growth performance in *Cyprinus carpio* Huanghe var [103]. Qin et al. demonstrated that *Lactobacillus casei* BL23 significantly increased fecundity in zebrafish (*Danio rerio*) [104]. The administration of *Lactobacillus plantarum* could effectively improve the growth performance and anti-stress capacity of Pacific white shrimp [105–107]. Zheng et al. also revealed that dietary addition of *Lactobacillus pentosus* AS13 effectively enhance the growth performance, feed utilization, digestive enzymes and disease resistance of Pacific white shrimp [108]. Dietary supplementation of heat-killed *L. plantarum* L-137 (HK L-137) and *L. plantarum* LL11 improved growth, digestive enzyme activities and non-specific immune parameters of sea cucumber [98,109]. Yu et al. reported that *L. plantarum* CCFM639 significantly enhanced feed utilization and growth performance, while alleviated aluminum toxicity in tilapia [110]. Moreover, Zhai et al. reported that dietary *L. plantarum* CCFM8661 alleviated Pb-induced toxicity in Nile tilapia (*Oreochromis niloticus*) [111]. Gao et al. showed that *L. pentosus* supplemented to diets enhanced food intake and growth of abalones (*Haliotis discus hannai*), and improved the non-specific immunity (total number of blood lymphocytes, lysozyme activity, acid phosphatase (ACP), superoxide dismutase and transcriptional expression of Mn-superoxide dismutase (Mn-SOD) and thioredoxin peroxidase (TPx) and resistance of abalones against *V. parahaemolyticus* infection [112]. Xie et al. reported that a *L. plantarum* screened from the aquaculture environment has high nitrite removal ability, and suggested that *L. plantarum* could be promising microorganism for water purification in aquaculture [113].

2.3.4.2. *Lactococcus lactis* In a study with tilapia, Zhou et al. revealed that *Lactococcus lactis* RQ516 enhanced growth and protected the fish against *A. hydrophila* infection [114]. *L. lactis* MM1 exhibited *in vitro* antagonistic activity against fish pathogens, such as *S. aureus*, *V. parahaemolyticus*, *H. rveyi*, and *V. metschnikovi* [93,115], and improved the feed utilization and immune responses (serum complement component 3 (C3) level and serum lysozyme activity) of juvenile grouper [95]. *L. lactis* LH8, isolated from marine fish, showed positive effects on immune response in sea cucumber [98].

2.3.4.3. *Leuconostoc lactis* *Leuconostoc lactis* isolated from the intestinal tract of black porgy (*Sparus microcephalus*), was evaluated as a potential probiotic strain for aquaculture [116]. The results showed that *Leu. lactis* had high tolerance to bile, low pH, trypsinase and pepsin conditions. Moreover, *in vitro* studies revealed that it co-aggregates with some pathogens, such as *V. parahaemolyticus*, *Listeria monocytogenes*, *E. coli* O157, *Salmonella typhimurium*, *Shigella*, *S. aureus* and *Proteus bacillus vulgaris*. Although the strain was not tested *in vivo*, the author speculated that *Leu. lactis* might be a putative probiotic strain in marine aquaculture.

#### 2.3.5. *Weissella confusa*

*Weissella confusa* strain LS13 isolated from marine animals (no specification given) by Li et al. showed inhibitory effects against some fish pathogens (*Vibrio splendidus*, *V. anguillarum*, *V. parahaemolyticus* and *S. aureus*), but dietary *W. confusa* (WC group) showed no effect on growth performance, AKP, lysozyme and SOD activities in sea

cucumber [98]. However, ACP activity was significantly higher in the WC group vs. control. In WC group, heat shock protein 60 (HSP60) was the most up-regulated gene at day 30, but first it was down regulated at day 15. Similar expression pattern was noticed for NF-kappa-B transcription factor p65 (Rel), in contrast to heat shock protein 90 (HSP90) and nitric oxide synthase (NOS), which were down regulated at all-time points.

### 2.4. Proteobacteria

#### 2.4.1. *Aeromonas*

Although *Aeromonas* species have typically been considered as opportunistic pathogens in the aquatic environment, several *Aeromonas* species are also commensal members in fish intestinal microbiota [117], and some of *Aeromonas* strains have been tested as potential probiotics in aquaculture. Dietary supplemented *Aeromonas veronii* BA-1 enhanced cellular (respiratory burst activity, phagocytic capacity) and humoral immune responses (serum protein, albumin, globulin values and lysozyme activity) and disease resistance of common carp against *A. hydrophila* infection [21]. Part of the immunostimulatory activity of *Aeromonas* spp. seems to reside in their extracellular products [23,72].

#### 2.4.2. *Bdellovibrio*

*Bdellovibrio* and like organisms (BALOs) are a group of motile and parasitic proteobacteria that prey on other Gram-negative bacteria for growth, reproduction, and survival [118]. *Bdellovibrio* was approved for animal use by the Chinese Ministry of Agriculture in 1994, and BALOs have been tested as probiotic/biocontrol agents in aquaculture for many years [119]. *Bdellovibrio* and BALOs are known to (i) reduce the total bacterial number and/or *Vibrio* population in the rearing water [119–125], (ii) improve water quality [120], (iii) reduce bacterial infection in fish [125–127], and (iv) promote survival and growth [119,123–125].

#### 2.4.3. *Ectothiorhodospira*

*Ectothiorhodospira* is a genus of photosynthetic purple sulfur bacteria, spiral cells of red color, depositing sulfur globules outside the cells [128]. *Ectothiorhodospira shaposhnikovii* WF was isolated from a marine shrimp pond and act both as bioremediation agent and nutrient source and can benefit white shrimp larvae at an appropriate dose [129].

#### 2.4.4. *Halomonas*

*Halomonas* species demonstrated to play an important ecological role because of their abundance in hypersaline environments [130], and they are also very interesting for biotechnological purposes for producing exoenzymes and other substances of commercial interest [130,131]. Zhang et al. isolated a *Halomonas* sp. B12 from the intestine of shrimp (*Fenneropenaeus chinensis*). Dietary inclusion of the strain improved the resistance of shrimp to white spot syndrome virus (WSSV) in a challenge trial [132].

#### 2.4.5. *Paracoccus*

*Paracoccus marcusii* DB11 was isolated from healthy sea cucumber intestines and culture ponds [133]. It has been shown to effectively reduce the concentrations of COD, ammonia, and nitrite in sea cucumber feed leachate [133] and inhibit the growth of *V. splendidus*, which causes skin ulceration syndrome in sea cucumbers. *P. marcusii* DB11 stimulated the immune system of juvenile sea cucumber and enhanced their resistance against *V. splendidus* [134]. Moreover, *P. marcusii* DB11 supplementation showed a positive effect on the growth performance and immune response in coelomocytes and the intestine of sea cucumbers [64].

#### 2.4.6. *Pseudomonas*

*Pseudomonas stutzeri* is distributed widely in the environment,

occupying diverse ecological niches, and has been proposed as a model organism for denitrification studies [135]. *P. stutzeri* strain SC221-M improved both water quality and microbial community structure in farmed carp system [136]. Fu et al. showed that the addition of *P. stutzeri* F11 to an experimental grass carp aquaculture system decreased nitrogen levels and modulated water microbial community [137]. Moreover, the natural compounds produced by *Pseudomonas aeruginosa* also showed antagonistic activities against *Vibrio* pathogens, suggesting the potential of *P. aeruginosa* strains as probiotics to treat vibriosis [138,139].

#### 2.4.7. *Pseudoalteromonas*

*Pseudoalteromonas* is a genus of marine bacteria. In a study with sea cucumber, 224 bacterial strains were isolated from the intestine [140]. One of the strains, *Pseudoalteromonas selya kovii* HS1, improved total coelomocytes counts, respiratory burst activity, lysozyme activity, ACP activity, and phagocytic activity of sea cucumber when supplemented to the basal diet. Ma et al. revealed that marine *Pseudoalteromonas* sp. BC228 improved the digestive enzyme activities (trypsin and lipase activities), immune response (phagocytic and lysozyme activities) and disease resistance of juvenile sea cucumber against *V. splendidus* infection in a challenge trial [141]. Furthermore, *P. piscicida* SW-1, isolated from seawater in a clam farm, was proved to protect clams from the infection of *V. alginolyticus* (V.-MP-1), indicating that SW-1 could be used as a probiotic to protect farmed clams [142].

#### 2.4.8. *Psychrobacter*

Genus *Psychrobacter* belongs to the family Moraxellaceae and has been reported as commensal members in the GI tract of several fish species [36,92,143], and can inhibit *in vitro* growth of *Vibrio anguillarum* and *Aeromonas salmonicida* [143]. In an *in vitro* study, *Psychrobacter* sp. SE6, isolated from the gut of fast-growing grouper showed antagonistic activity against *V. harveyi*, *V. metschnikovi* and *Vibrio alginolyticus* [92]. In a subsequent 60-day feeding trial, dietary supplementation of *Psychrobacter* sp. SE6 significantly improved feed conversion ratio and complement 4 (C4) of juvenile grouper [144]. Further, Yang et al. evaluated the effect of dietary *Psychrobacter* sp. SE6 on the autochthonous (adherent) microbial diversity in fore-, mid- and hindgut of grouper, and revealed that samples from the probiotic group displayed different DGGE patterns compared with control group. Total number of bands and Shannon index of the fore-, mid- and hindgut samples in the probiotic group were significantly higher than those in the control group. Some potentially beneficial bacteria were stimulated, while some potentially harmful species, such as *Staphylococcus saprophyticus*, were suppressed [145]. Furthermore, Xia et al. showed that both the viable and heat-inactivated *Psychrobacter* sp. SE6 revealed positive effects on feed utilization and immunity in grouper [146].

#### 2.4.9. *Rhodospseudomonas*

A photosynthetic purple non-sulfur bacterial strain, *Rhodospseudomonas palustris* G06, improved the growth performance of juvenile white shrimp when administered as a water additive [147]. *R. palustris* G06 furthermore increased growth and immune function of tilapia when added to the water [25]. Wang also reported that dietary supplemented *R. palustris* improved growth performance of grass carp fingerlings [148]. Moreover, Zhang et al. reported that adding *R. palustris* to grass carp water significantly reduced the nitrogen levels as it modulates the microbial community [149].

#### 2.4.10. *Shewanella*

*Shewanella* species are facultatively anaerobic Gram-negative rods, most of which are found in extreme aquatic habitats [150]. *Shewanella* bacteria are a normal component of the surface and intestinal microbiota of fish, and are implicated in fish spoilage [151]. The use of *Shewanella* as probiotics has been reported in various aquatic species. Jiang et al. isolated *Shewanella colwelliana* WA64 and *Shewanella*

*olleyana* WA65 from the gut of abalone [152]. Dietary supplementation of WA64 and WA65 promoted growth, enhanced cellular and humoral immune responses of abalone, and decreased mortality of juvenile abalone after challenge with *V. harveyi* [153]. Similarly, *S. japonica* HS7 isolated from the intestine of healthy sea cucumber enhanced growth, survival, and cellular and humoral immune responses of sea cucumber when added in diet at  $10^9$  cell  $g^{-1}$  [140]. Hao et al. showed that *S. haliotis* D4 improved growth, innate immunity (respiratory burst, SOD and ACP activities; expression of prophenoloxidase and beta-1, 3-glucan-binding protein) and disease resistance of Pacific white shrimp [72]. The same authors further showed that dietary administration of *S. xiamenensis* A-1 and *S. xiamenensis* A-2 improved the composition of intestinal microbial community and enhanced the immunity in grass carp [23,154].

#### 2.4.11. *Vibrio*

Although isolates of *Vibrio* spp. are commonly believed to be pathogenic for aquatic animals [155], some avirulent strains have been suggested to have probiotic properties [156,157]. It has been reported that *Vibrio* sp. V33 isolated from healthy sepia has strong antagonistic activity against pathogenic *V. splendidus* Vs [158]. Chi et al. isolated a *V. tasmaniensis* strain HS10 from the intestine of sea cucumber, and dietary supplementation of the strain improved the humoral and cellular immune response of sea cucumber [140].

### 2.5. Non-bacterial candidates

#### 2.5.1. Yeast

2.5.1.1. *Saccharomyces*. *Saccharomyces cerevisiae* is the most commonly used yeast probiotic in aquaculture. Ran et al. proved that live yeast, *Saccharomyces cerevisiae*, significantly increased gut microvilli length and trypsin activity, decreased intestinal *hsp70* expression, and enhanced resistance of tilapia (*Oreochromis mossambicus*) against *A. hydrophila* infection in a challenge trial. Apart from live yeast, the fermentation product of *S. cerevisiae* has also been widely used, and some have been developed as commercial products [159]. He et al. showed that supplementation of the *S. cerevisiae* fermentation product DVAQUA<sup>®</sup> improved the immunity of hybrid tilapia (*Oreochromis mossambicus*) [160]. In a study with common carp, Huang et al. reported that dietary supplementation of  $1\text{ g kg}^{-1}$  *Saccharoculture* (a Korean-made *S. cerevisiae* culture product containing  $10^7$  CFU  $g^{-1}$  *B. amyloliquefaciens* spores) improved the posterior intestinal microvillus length and general welfare of the fish [161].

2.5.1.2. *Marine yeasts*. A series of studies have used marine yeasts as probiotics in sea cucumber. Li et al. isolated *Hanseniaspora opuntiae* C21 from the intestine of healthy sea cucumber. The strain showed *in vitro* antagonism against the growth of the pathogen *Shewanella marisflavi* AP629 [162]. Diet supplementation with *H. opuntiae* C21 at  $10^4$  and  $10^5$  CFU  $g^{-1}$  improved growth, immunity, and resistance against *V. splendidus* in juvenile sea cucumber [163]. The marine yeast *Metschnikowia* sp. C14, isolated from the intestine of sea cucumber, stimulated the immune system of juvenile sea cucumber and enhanced resistance against *V. splendidus* infection [164]. Furthermore, *Metschnikowia* sp. C14 was also reported to improve the growth and activity of intestinal digestive enzymes (trypsin and lipase) of juvenile sea cucumber [165]. The marine yeast *Rhodotorula* sp. C11, isolated from the intestine of the sea cucumber, inhibited growth of the pathogen *V. splendidus* NB13 *in vitro* [166], and further use revealed that *Rhodotorula* sp. C11 effectively colonized the intestine of Japanese spiky sea cucumbers and improved growth and disease resistance against *V. splendidus* in a challenge trial. Similarly, Wang et al. used a *Rhodotorula benthica* D30 isolated from sea mud, and supplemented the strain in feed at  $10^6$  CFU  $g^{-1}$  and  $10^7$  CFU  $g^{-1}$ . Results showed that *R. benthica* D30 can increase growth, digestive enzymes activity, immunity, and disease resistance of sea cucumber against *V.*

*splendidus* infection [167].

Yang et al. (2010) isolated the marine red yeast *Rhodospiridium paludigenum* from coastal water in Zhejiang, China. Dietary inclusion of *R. paludigenum* at  $10^8$  CFU  $g^{-1}$  enhanced growth and antioxidant capacity (SOD, catalase, glutathione peroxidase, total antioxidant capacity) of Pacific white shrimp [168]. Later, the same authors studied the expression of genes related with antioxidant function in Pacific white shrimp fed dry or live *R. paludigenum*, and the results showed a significant up-regulation of genes coding for manganese superoxide dismutase (SODMn), catalase, glutathione peroxidase, and ferritin, which are all important elements of the antioxidant defense system [169].

## 2.6. Mixture of probiotic strains

In his review “Probiotics in man and animals”, Fuller stated; “Probiotic preparations may consist of single strains or may contain any number up to eight strains” [170]. Most probiotic studies have used single administration in Chinese aquaculture, but inclusion of multiple probiotics have gained interest during the last decade [30,72,136,154,171–173]. The advantage by using multiple-strain preparations is that they may further improve the overall beneficial effect of the probiotic formulation and are active against wider range of conditions and species.

A dietary commercial probiotic product (Qingdao Master Biotechnology Co. Ltd., Qingdao, China) containing *B. subtilis*  $7.0 \times 10^9$  CFU  $g^{-1}$ , *B. licheniformis*  $3.0 \times 10^9$  CFU  $g^{-1}$ , *Lactobacillus* spp.  $5.0 \times 10^8$  CFU  $g^{-1}$  and *Arthrobacter* spp.  $1.0 \times 10^8$  CFU  $g^{-1}$  significantly increased the specific growth rate (SGR), innate immunity, and resistance against *V. harveyi* in cobia (*Rachycentron canadum*) [171]. A mixed preparation of *B. cereus* BSC24 and *P. stutzeri* SC221-M was more efficient in removing nitrogen from grass carp culture water than supplementation of SC221-M or BSC24 alone, i.e., 53.9% for the mixture versus 24.5% and 26.6% for SC221-M and BSC24, respectively [136]. Hao et al. revealed that shrimps fed *S. haliotis* D4, *B. cereus* D7 and *Aeromonas bivalvium* D15 at a ratio of 2:1:1, dosed at  $10^7$  cell  $g^{-1}$ , showed better growth performance and disease resistance compared with those fed single probiotics [72]. Wu et al. demonstrated that dietary supplementation of a mixture of *S. xiamenensis* A-1, *S. xiamenensis* A-2, and *A. veronii* A-7 at  $10^8$  cell  $g^{-1}$ , at a ratio of 4:2:1, resulted in more improved innate immunity and disease resistance compared to single fed probiotics [154]. Li et al. reported that a dietary probiotics mixture of *B. subtilis* YB-1 and *B. cereus* YB-2 stimulated non-specific immune responses and enhanced the growth performance and resistance against *V. alginolyticus* infection vs. the control group [30]. Dietary combination of *B. cereus*, *L. acidophilus* and *C. butyricum* at  $3.0 \times 10^9$  CFU  $kg^{-1}$  significantly improved final weight, SGR, food consumption, food conversion efficiency and apparent digestibility coefficient, and enhanced the activity of digestive enzymes (pepsin, trypsin, amylase and lipase), anti-oxidant enzymes (superoxide dismutase, catalase and glutathione peroxidase), and lysozyme of hybrid grouper (*Epinephelus lanceolatus* ♂ × *fuscoguttatus* ♀) compared to control [173]. Liu et al. proved that dietary supplementation with 3% complex-probiotics-preparation (BSCP, including *B. amyloliquefaciens* V4CGMCC 10149 and *Rhodotorula mucilaginosa* CGMCC 1013) improved the growth performance, enhanced the activities of some digestive enzymes, and promoted the nonspecific immunity of Atlantic salmon (*Salmo salar* L.) [174].

## 2.7. Synbiotics

Synbiotics refers to nutritional supplements combining a mixture of probiotics and prebiotics in a form of synergism. Gibson and Roberfroid defined synbiotics to ‘characterize some colonic foods with interesting nutritional properties that make these compounds candidates for classification as health-enhancing functional ingredients’ [175]. In aquaculture the first finfish study on synbiotics was published by Rodriguez-Estrada et al. [176], and since then several studies have emerged [177–181].

In Chinese aquaculture several studies have used synbiotics. Zhang et al. revealed that dietary supplementation of *B. subtilis* ( $10^7$  CFU  $g^{-1}$ ) and 0.25% fructooligosaccharide (FOS) significantly increased SGR, total coelomocytes counts (TCC), phagocytosis, and disease resistance of sea cucumbers against *V. splendidus* infection [182]. Moreover, the counts of total viable bacteria were increased by supplementation of *B. subtilis* ( $10^7$  CFU  $g^{-1}$ ) and 0.25% FOS, whereas *Vibrio* counts were decreased. Dietary *B. licheniformis* ( $10^7$  CFU  $g^{-1}$ ) and 0.3% FOS significantly enhanced the innate immunity and antioxidant capability of triangular bream (*Megalobrama terminalis*), and improved its resistance against *A. hydrophila* in a challenge trial [183]. In a study with juvenile ovate pompano, Zhang et al. showed that dietary administration of *B. subtilis* ( $10^7$  CFU  $g^{-1}$ ) and FOS (0.2% and 0.4%) had significant interaction on enhancing SGR, respiratory burst activity, lysozyme activity and disease resistance against *V. vulnificus* compared to control. Supplementation with *B. licheniformis* and xylo-oligosaccharide at 0.1% of diet increased weight gain by 20.9% with a 8.9% lower food conversion ratio in grass carp juveniles compared to controls [184]. The synbiotics fed fish also revealed higher intestinal activity of trypsin (18.8%), amylase (36.8%) and lipase (19.8%) activities. Zhang et al. showed significant improvement of final weight, WG, SGR, survival rate as well as protease and  $Na^+$ ,  $K^+$ -ATPase activities of triangular bream (*Megalobrama terminalis*) fed *B. licheniformis* and FOS vs. control fed fish [185].

Zhang et al. revealed that a mixture of  $10^8$  CFU  $g^{-1}$  *B. licheniformis*, *B. subtilis* and 0.2% isomaltooligosaccharide (IMO) improved immune activities (phenoloxidase, lysozyme, NOS activity, superoxide dismutase activities) and disease resistance against *V. alginolyticus* infection of shrimp (*Penaeus japonicus*) compared to the control group [186]. Hu et al. showed that combined use of *Bacillus* and molasses increased the diversity of the microbial community, and promoted the formation and development of a beneficial microbial community structure and inhibited pathogens in Pacific white shrimp [187]. Dietary *B. coagulans* ( $10^9$  CFU  $g^{-1}$ ) and 0.2% chitosan oligosaccharides (COS) had a synergistic effect in enhancing immunity (total leukocyte count, respiratory burst activity, phagocytic activity, lysozyme, SOD) and disease resistance against *A. veronii* infection of common carp (*Cyprinus carpio* koi) [188]. Feeding sea cucumber *B. licheniformis* WS-2 ( $10^9$  CFU  $g^{-1}$ ) and alginate oligosaccharides (AOS) improved growth, activity of digestive enzymes (amylase, protease and alginate lyase), non-specific immune response, intestinal *Bacillus* and *Lactococcus* levels, and resistance against *V. splendidus* infection [189]. Ye et al. revealed that diets supplemented with FOS, Maltoligosaccharide (MOS) and *B. clausii* ( $10^7$  cell  $g^{-1}$ ) improved growth performance and health benefits of the Japanese flounder (*Paralichthys olivaceus*) compared to the control [190]. Geng et al. demonstrated that combination of 1.0 g *B. subtilis*  $kg^{-1}$  and 6.0 g chitosan  $kg^{-1}$  is the optimal inclusion level of cobia (*Rachycentron canadum*) for growth, innate immunity (lysozyme, ACP, phagocytosis and respiratory burst), and resistance against *V. harveyi* infection [191].

## 3. Mode of action

### 3.1. Production of digestive enzymes

Higher enzyme activities in the digestive tract enhance digestive capabilities and growth performance of the host. It is widely accepted that the activity of digestive enzymes is a useful comparative indicator for food utilization, digestive capacity, and growth performance of the host [177]. Gao et al. demonstrated a likely correlation between higher growth rate and higher intestinal digestive enzyme activity in probiotic fed silver pomfret (*Pampus argenteus*). They suggested the main cause was better utilization and digestion of the diets [31]. Previous studies have shown that many *Bacillus* spp. with exo-enzyme activities can significantly improve the host's growth performance [63,67,192,193], which may be attributed to the production of digestive enzymes.

### 3.2. Production of antibacterial substances

*In vitro* antagonism against aquatic pathogens has been an important selection criterion for candidate probiotics for aquaculture use. The nature of antimicrobial compounds that are responsible for the antagonistic activity of probiotics have been investigated in several studies in the past decade. Yang et al. reported that two strains of LAB (*Lactococcus lactis* MM1 and MM4) can secrete hydrogen peroxide and bacteriocin-like substances, which have strongly inhibitory activities against pathogens, i.e., *V. metschnikovi*, *V. harveyi* and *S. aureus* that infects orange-spotted grouper [115]. Xu et al. purified lipopeptides N3 produced by *B. amyloliquefaciens* M1, which displayed strong anti-*Vibrio* action on the whole cells and cell membrane. Further, this antibacterial actions of lipopeptide N3 could be a consequence of its ability to form ion-conducting channels in bacterial cell membranes by exploiting its detergent-like action on cell membranes, also called membrane active properties [194]. In recent study, Gao et al. revealed that *B. pumilus* H2 had considerable anti-*Vibrio* activity and the major mechanism appeared to involve disruption of cell membranes and consequent cell lysis relating to an anti-*Vibrio* substance named amicoumacin A [195]. Gao et al. also identified several anti-*A. salmonicida* compounds belonging to the iturin, macrolactin, and diffidin groups from cell-free supernatant of *B. velezensis* V4. These compounds contributed collaboratively to *in vitro* growth inhibition of *A. salmonicida*. The diversity of the compounds was related to the versatility of their mode of action, especially the membrane disruptive effect of iturin group members [196]. *C. butyricum* and its spent culture supernatants inhibited growth and adherence to fish intestine epithelial cells (FIECs) of *Salmonella enteritidis* and *V. parahaemolyticus*. It was hypothesized that these antibacterial functions were related to the organic acids produced by *C. butyricum* which lowered the pH in the intestine [83].

### 3.3. Immune stimulation

Numerous publications have shown that probiotics stimulate the immune system [99,197]. However, the mechanisms underlying the immune stimulation effect of probiotics have been less investigated. Sun et al. showed that feeding viable and heat-inactivated *Psychrobacter* sp. SE6 to grouper had interesting effects on the intestinal immune responses. The expression of TLR2 and TLR5, adaptor MyD88 and cytokines (IL-1 $\beta$ , IL-8 and TGF- $\beta$ 1) was upregulated when fed the viable SE6. In fish fed heat-inactivated SE6, only TLR2 was upregulated but not MyD88 and cytokines. This suggested that a MyD88-independent TLR2 signaling pathway may be involved in the probiotic recognition in grouper [198]. In a recent study, Qin et al. revealed that *L. casei* BL23 protected zebrafish larvae from *A. veronii* infection. Further data indicated that *L. casei* BL23 enhanced host immune responses, which may be attributable to exopolysaccharide-protein complex (EPSP) from BL23 via TLR1/TLR2 pathways [104].

Moreover, SCFAs and their salts showed immune stimulatory effect when used as feed additives in fish [199,200]. Therefore, the involvement of SCFAs in the immunostimulatory effect of aquatic probiotics deserves more investigation. Moreover, isolation and characterization of SCFAs-producing commensal microbes in the intestine of fish may bring about novel aquatic probiotics.

### 3.4. Interference of quorum sensing

Quorum sensing (QS) is a cell density dependent process that enables bacteria to communicate with each other based on the production, secretion and sensing of the auto-inducer molecules and then subsequently regulate virulence associated gene expression [201]. Interrupting quorum sensing may represent a novel alternative approach to combat bacterial pathogens. Some bacteria can produce quorum quenching (QQ) enzymes. The approach has been recommended as a promising non-antibiotic strategy for bacterial disease therapy

[201–203]. *N*-Acyl-homoserine lactones (AHLs) are the main signal molecules produced by many Gram-negative bacteria, such as the aquatic pathogens *Vibrio* spp. and *Aeromonas* spp. [204]. The probiotic strain *Bacillus* sp. QSI-1, isolated from the gut of a healthy crucian (*Carassius auratus gibelio*), has been reported to protect fish from *A. hydrophila* infection [205]. Zhou et al. demonstrated that quorum quenching probiotics *Bacillus* sp. QSI-1 can modulate the gut microbiota community by degrading AHLs [50]. The authors also revealed a reduction in the abundance of the fish pathogen *A. hydrophila* in the GI tract. These results indicated that QSI-1 protect fish from *A. hydrophila* infection by quorum quenching. These results provide new insight into the mechanisms of probiotics, and suggest that quorum quenching probiotics can be used as an alternative strategy to combat bacterial infection in aquaculture instead of antibiotics.

### 3.5. Competition for adhesion sites

Competition of adhesion sites has been reported in several studies. Zhou et al. evaluated the GI tract adhesion property of ten *Lactobacillus* strains [206]. They observed that while having similar antibacterial activity *in vitro*, the highly-adhesive strain *L. plantarum* JCM 1149<sup>T</sup> conferred stronger resistance against *A. hydrophila* infection in zebrafish compared with the less-adhesive strain *L. acidophilus* JCM 1132<sup>T</sup>. Further, an *ex vivo* intestinal sac experiment showed that *L. plantarum* JCM 1149<sup>T</sup> can compete for intestinal adhesion sites with *A. hydrophila* and alleviated gut mucosa damage caused by *A. hydrophila* [207]. These results highlight GI tract adhesion (adhesion to mucosa) as the favorable criterion in the selection of dietary probiotics in aquaculture, which may confer the probiotics disease resistance property through competition for adhesion sites with the pathogens.

## 4. Regulation and safety issues

According to the catalogue of feed additives authorized by Chinese Ministry of Agriculture (2013), probiotics permitted to be supplemented in animal feed include bacteria, and fungi (for example yeast). The permitted bacterial genera include *Lactobacillus*, *Bacillus*, *Enterococcus*, *Pediococcus*, etc., while yeast permitted for use as feed additive mainly includes *Saccharomyces cerevisiae*. For further regulation, an industry standard is needed to normalize the production and usage of related products. Notably, there is no specific regulation in China for probiotics used to improve pond water and sediments, and the commercial probiotic products for this purpose are not well managed in the current stage, which deserves attention.

Although probiotics are used as alternatives of antibiotics, much attention has been directed to the production of inhibitory substances in the selection and study of aquaculture probiotics. It has been indicated that the antagonistic effects of *Bacillus* strains against aquatic pathogens were attributed to the production of antibiotics diffidin and surfactin [208], implying that the use of some *Bacillus* strains as probiotics in aquaculture might induce the same problems as antibiotics usage. As proposed by Gatesoupe, the risk to select probiotic-resistant pathogens should not be underestimated [46]. Moreover, the risk of transferring antibiotic resistance from probiotics to pathogenic bacteria should not be neglected, as some probiotic *Lactobacillus* strains have been reported to transfer antibiotic resistance genes *in vitro* and in rodent models [209,210]. In addition, previous studies have revealed that administration suspension (including cessation) of *Lactobacillus* strains may lead to gut dysbiosis and increase pathogen susceptibility of hybrid tilapia [211]. Furthermore, He et al. showed that *L. rhamnosus* GG, a well-known probiotic for humans, induced injury to the mucosa of zebrafish when applied as a probiotic strain for disease protection [212]. Similarly, a *Lactobacillus plantarum* strain originally isolated from Sabalan cheese induced damaged epithelial cells and disorganized microvilli in the intestine of beluga (*Huso huso*) [213]. Both of the studies indicated that a probiotic that is safe for human and terrestrial

animals may not be safe for aquatic animals. Consequently, the safety of potential aquaculture probiotics should be carefully evaluated in aquatic animal models.

## 5. Selection criteria and perspectives

The probiotics for aquatic usage -are different from probiotics for terrestrial animals, as certain influencing factors are fundamentally different from terrestrial probiotics. Therefore, a selection criteria list specific for aquatic probiotics is highly important. Previous reviews have proposed selection criteria for aquatic probiotics [13,14,214–216]. Following on from these papers we propose the basic criteria for aquatic probiotics: 1) Safety. The strain must be safe for both the aquatic host and human, with no plasmids encoded antibiotic resistance genes. 2) Adaptability. The strain should survive the intestinal tract of aquatic host. Colonization and persistence in the intestine are preferred quality but are not required. For water additive, the strain should survive in the aquatic environment. 3) Function. The strain should benefit the aquatic host. The beneficial effect can be on growth, digestion, immunity, disease resistance, or general welfare. For water additive, the strain should be able to improve water quality or sediments. 4) Convenience. The strain should be convenient for storage and administration. Based on these criteria, an authorized catalogue for probiotics permitted for aquatic animal use will promote the regulation of this field.

Although many studies about probiotics in aquaculture have emerged in China, the approach has been generally empirical. Moreover, the mechanisms underlying the beneficial effects of probiotics in aquatic systems are rarely studied, and the potential harmful effects of some probiotics have been generally neglected. Further studies are warranted to elucidate the mechanism of the beneficial effects and potential harmful effects of the probiotics, which may guide the selection and function-oriented manipulation/design of more efficacious and safe aquaculture probiotics. In addition, given the potential drawbacks of probiotics, administration of the probiotic effector ingredients might be a better way to obtain the health benefits associated with probiotic consumption [217]. The probiotic effectors include the structural components of the probiotic cells and secretory components such as metabolites, enzymes and functional proteins. In-depth study of well-accepted probiotic strains will promote the finding of probiotic effectors, and further topics involving the probiotic effectors may include structure-activity relationship (SAR) as well as synergy among different effectors.

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