Research article

Media Optimization and Partial Purification of Anti-Candida Compound of Probiotics Cultures

Pashmin Anand¹, Mayur Aswani², Suyash Kathade¹ and Bipinraj Kunchiraman^{1*}

¹Department of Microbiology, Rajiv Gandhi Institute of I.T. and Biotechnology, Bharati Vidyapeeth Deemed to be University, Pune- 411046, India ²Department of Herbal Medicine, Interactive Research School for Health Affairs, Bharati Vidyapeeth Deemed to be University, Pune- 411046, India

Received: 29 March 2022, Revised: 29 May 2022, Accepted: 23 December 2022

DOI: 10.55003/cast.2022.04.23.010

Abstract

Keywords	Fungal infections in humans are mostly due to <i>Candida albicans</i> , an opportunistic commensal yeast that causes systemic and invasive
anti-candida;	candidiasis. Conventional treatment of candidiasis using antifungal compounds such as polyenes and azoles can cause various side effects.
Bacillus;	Besides, treatment becomes difficult as the organism acquires
probiotic;	resistance. Probiotics with anti-candida activity have been projected as a safe and alternative solutions to treat candidiasis. Probiotic <i>Bacillus</i>
purification	isolates previously reported showed wonderful activity against pathogenic <i>Candida</i> species. This study reports on media optimization and partial purification of anti-candida compounds from probiotic <i>Bacillus</i> isolates MW3, MW9, MW19, and MW27. Maximum growth and anti-candida activity (2.9 cm inhibition zone) by the isolate culture was observed when media had been optimized with dextrose, casein and peptone. The media composition numbers 18, 4 and 19 showed the best growth for MW3, MW9, MW19 and MW27. Fractions obtained by solvent precipitation using acetone and methanol showed activity against the pathogen, and the fractions were further purified by ultra- filtration. After ultra-filtration, it was found that the size of the active compound was between 10 to 30 kDa. The compound lost its activity after treatment with proteinase K.

1. Introduction

Candida albicans is the most prevalent cause of fungal diseases [1, 2] which range from non-life-threatening oral thrush to life-threatening invasive diseases like candidemia. Globally, 75% of women have experienced *C. albicans* infection at some point in their lifetime, and 40-50% of these

^{*}Corresponding author: Tel.: 7907610568 Fax: 020-24365713 E-mail: k.bipinraj@bharatividyapeeth.edu

have recurrent infections [3]. These infections are mostly caused by dysbiosis in the gastrointestinal (GI) tract. Apart from an increase in the pathogenic count in the GI tract, dysbiosis reduces host immunity, which favors *C. albicans* evasion and infection. Polyenes, azoles, allylamines, morpholines, antimetabolites, and echinocandins are antifungal medications that have been conventionally used to treat *C. albicans* infections [4]. However, common medications like clotrimazole, miconazole, and fluconazole interfere with normal metabolic processes, cause allergic reactions, and are toxic to the liver and kidneys [5]. It has been well established that a balanced proportion of probiotics in the intestinal tract plays a vital role in preventing pathogens as well as maintaining the proper immunity of the host.

Probiotics are live microorganisms, which can be administered in adequate amounts to confer health benefits to the host [6, 7]. Different businesses, such as dairy/functional foods/dietary supplements and pharma, have formulations with various probiotic strains that offer specific human health advantages [8, 9]. Lactobacillus and Bifidobacterium are well-known probiotic microorganisms that have been linked to a variety of health benefits. In addition to these two genera, Enterococcus, Streptococcus, and Bacillus species have been described as probiotics with potential health benefits [10, 11]. Bacillus species, in particular, have been observed to resist the intense digestive action of bile salts due to their spore-producing ability [7, 11]. Bacillus spp. are preferred on an industrial scale because they are stable during numerous industrial processes. This species has a significant commercial value due to its ability to secrete vast amounts of extracellular compounds [12], as well as its antibacterial, anti-oxidant, and immunomodulatory properties [13-15]. Due to the increase in anti-fungus resistance, it is critical for developing therapeutic or prophylactic alternatives that have a distinct mechanism of action capable of inhibiting fungal infections without harming the host organism. Hence a probiotic Bacillus with anti-fungal activity offers an alternate option for candidiasis prevention and therapy. The composition of the medium has a strong impact on the bacteria's ability to create diverse extracellular components. This is accomplished by including the appropriate carbon and nitrogen sources at manageable concentrations. Carbon is essential for growth as well as the creation of primary and secondary metabolites. The rate of carbon metabolization is directly proportional to the rate of biomass creation and metabolite production [16]. While nitrogen is responsible for protein synthesis, it also serves as an energy source in the absence of carbon and is important in a variety of metabolic processes [16, 17].

Bacillus species have been found to produce a wide spectrum of antifungal and antibacterial proteins. *Bacillus* is non-pathogenic bacteria that can non-ribosomally generate a variety of short antibiotic peptides (<2000 kDa) with antifungal properties. *Bacillus* sp. can even generate a wide spectrum of lipopeptide antibiotics, which have been widely used in medicine and agriculture [18]. Production of antimicrobial peptides (AMPs) by *Bacillus* strains has been progressively studied in recent years, with several peptides produced by this group of bacteria discovered to be suitable for a variety of uses [19]. *Bacillus* has always had an advantage over *Lactobacillus* due to its ability to produce resistant spores, its creation of various antifungal compounds, particularly lipoproteins or bacteriocins, and the ability to manufacture vast numbers of proteins [20]. Anti-microbial peptides (AMPs) are naturally the first line of defense between the host and the environment since they have the potential to inhibit a wide range of microorganisms [21]. *Bacillus* species produce AMPs that have wide antimicrobial efficacy against harmful bacteria.

As a result, this study presents media optimization for the maximum anti-candida activity of previously identified probiotic anti-candida *Bacillus* cultures designated as MW3, MW9, MW19, and MW27 [22-24]. The partial purification and chemical characterization of an anti-candida molecule are also reported in this study.

2. Materials and Methods

2.1 Culture Collection

Bacillus isolates designated as MW3, MW9, MW19, (*Bacillus tequilensis*) and MW27 (*Bacillus subtilis*), previously isolated from Indian traditional fermented batter of Meduwada, were used for further analysis of their anti-candida components [22-24]. These cultures were grown on Sabouraud dextrose agar (SDA) slants and stored in a refrigerator for further use. *Candida albicans* (NCIM-3557) culture was obtained from NCIM and cultivated in SDA slants before being stored for future use.

2.2 Inoculum development

The inoculum was prepared by adding 24-h old cell suspension into a 100 ml conical flask containing 50 ml of sterile inoculum medium Sabouraud dextrose broth, (SDB). At 37°C, the flask was kept under shaking at 120 rpm on a rotary shaker.

2.3 Media optimization experiments

The inoculum was centrifuged after 24 h, and the cell pellet was suspended in 100 mL of different production media to get 1 OD at 600 nm. The production medium contained only dextrose as a carbon source, and the nitrogen sources were the peptic digests of animal tissue and pancreatic digest of casein [25]. To optimize biomass output, several permutations and combinations of medium components were tested. As a result, three different concentrations of media components (Table 1) were added while the other two components remained unchanged, and the pH was adjusted to 6.5. The best combination was chosen based on the maximum bacterial growth measured as optical density at 600 nm after 24 h of incubation at 37°C.

2.4 MIC

A 96-well plate was used to find out the MIC (minimum inhibitory concentration) of the supernatant after growing the culture in the optimized medium. The protein concentration of different culture supernatant was determined by the Lowry method and compared with standard BSA (Bovine Serum Albumin) [18]. The supernatant was then diluted to adjust the protein concentration from 256 μ g/ μ L to 4 μ g/ μ L by the double dilution method and was tested against *Candida albicans*. The experiment was carried out by first adding 100 μ L of *Candida* suspension (0.5 OD at 600 nm) in SDB medium, followed by 100 μ L culture supernatants produced in the optimized media. The plates were incubated at 37°C for 3 h before being measured for optical density at 600 nm with an ELISA plate reader and compared to the positive (fluconazole 1 mg/mL) and negative controls (*Candida* without any inhibitor).

2.5 Anti-candida activity

The isolates cultivated in an optimized medium were tested for antimicrobial activity against *Candida albicans* NCIM 3557. The pathogen was spread on SDA plates (100 μ L, 0.5 OD at 600 nm) and incubated for 10 min at 37°C. An agar borer was used to create wells in agar and 20 μ L of 24- h old grown culture supernatant was added to the wells before incubating at 37°C for 24 h [26, 27].

Sr. No.	Peptic digestof animaltissue (g/L)	Pancreaticdigest of casein (g/L)	Dextrose(g/L)	Ор	Optical Density at 600 nm			
	(3)		-	MW3	MW9	MW19	MW27	
1	2.5	2.5	10	0.042	1.188	1.1	1.155	
2	2.5	2.5	20	0.056	0.673	0.608	0.831	
3	2.5	2.5	30	0.052	0.04	0.025	0.934	
4	2.5	5	10	0.029	1.278	1.134	0.859	
5	2.5	5	20	0.016	0.498	0.381	0.579	
6	2.5	5	30	0.027	0.082	0.066	0.147	
7	2.5	7.5	10	0.016	1.169	0.852	1.218	
8	2.5	7.5	20	0.098	0.425	0.141	0.741	
9	2.5	7.5	30	0.078	0.014	0.12	0.691	
10	5	2.5	10	0.047	1.073	1.145	1.117	
11	5	2.5	20	0.028	0.067	0.582	0.869	
12	5	2.5	30	0.048	0.019	0.034	0.416	
13	5	5	10	0.098	1.11	1.135	1.204	
14	5	5	20	0.028	0.561	0.161	0.853	
15	5	5	30	0.015	0.165	0.064	0.379	
16	5	7.5	10	0.015	1.063	1.07	1.215	
17	5	7.5	20	0.037	0.122	0.241	0.28	
18	5	7.5	30	0.451	0.119	0.144	0.576	
19	7.5	2.5	10	0.008	1.07	1.159	1.24	
20	7.5	2.5	20	0.007	0.545	0.111	0.928	
21	7.5	2.5	30	0.045	0.03	0.088	0.563	
22	7.5	5	10	0.016	0.973	0.858	1.007	
23	7.5	5	20	0.046	0.301	0.119	0.235	
24	7.5	5	30	0.324	0.059	0.248	0.216	
25	7.5	7.5	10	0.289	0.831	0.78	1.176	
26	7.5	7.5	20	0.242	0.361	0.109	0.821	
27	7.5	7.5	30	0.264	0.017	0.114	0.491	

Table 1. Composition of media optimization

2.6 Partial purification of anti-candida compound

The anti-candida compound from the supernatant was precipitated using three methods; i) equal proportion (10%) v/v of TCA and ice-cold acetone were mixed thoroughly with culture supernatant and incubated overnight at -20°C. ii) The culture supernatant was mixed thoroughly with 4 Vol of methanol followed by the addition of chloroform (1 Vol). After adding 3 Vol of distilled water, the mixture was then allowed to settle for 2-3 min before centrifuging at 6000 rpm for 2 m. The aqueous top layer was removed, and 4 Vol of methanol was added before centrifuging the mixture at 6000 rpm for 2 min. The precipitate formed were collected after removing the supernatant. iii) The culture supernatant was slowly mixed with an equal volume of acetone. The mixture was incubated at -20°C for 12 h and the pellets were collected by centrifugation at 7500 rpm at 4°C for 15 min. The pellets collected using the above methods were then air dried and washed twice with 1M Tris base buffer (pH 7). The pellets were re-suspended in a solution of 1M Tris base buffer (pH 7) and used for anti-candida activity by agar well diffusion method.

2.7 Effect of proteolytic enzyme and temperature on anti-candida activity

To check the proteinaceous nature of the anti-candida molecules, the supernatant was treated with an equal volume of proteinase K in phosphate buffer (pH 7) (1mg/mL) and incubated at 37°C, and the mixture was incubated for 2 h. Similarly, each supernatant was heated for 3 min at 80°C to check the effect of temperature on the anti-candida activity. All samples were used for checking the anti-candida activity using the agar well diffusion method [28].

2.8 Ultra-filtration

The Sigma-Aldrich Amicon ultracentrifugal filter units, Ultra-15-MWCO-30kDa, 10kDa, and 4kDa, were used for ultrafiltration. *Bacillus* cultures were cultivated in an optimized medium for 24 h. The culture was then centrifuged twice at 10000 rpm for 10 min each time until the supernatant became cell-free. Each supernatant was then passed through 0.22 μ m filter using a syringe filter. The filtrate was collected and ultra-centrifuged at 10000 rpm for 30 min using Ultra-15-MWCO-30 kDa. The filtrate of Ultra-15-MWCO-30 kDa was filtered with an Ultra-15-MWCO-30 kDa filter, and the resulting filtrate was then filtered using an Ultra-15-MWCO-4 kDa filter. Both the filtrates and retentates were tested for anti-candida activity using the agar well diffusion method.

2.9 Germ tube inhibition assay

Germ tube inhibition was performed according to Palande *et al.* [22]. *Candida* culture (100 μ L) suspended in SDB, supplemented with 0.2% human serum was added to 96-well flat-bottom plates followed by the addition of 100 μ L of culture supernatant or partially purified compound (acetone precipitated). The plate was incubated at 37°C for 4 h. *Candida* culture with saline was used as a negative control. After the incubation period, the medium in the wells were discarded, and each well was washed with 70% ethanol followed by 25% SDS and twice with distilled water. Crystal violet (0.1%) 200 μ L was added to each well and allowed to react for 10 min. After the crystal violet treatment, the wells were washed twice with D/W (distilled water) and then with 0.25% SDS. The remaining crystal violet was eluted with 200 μ L of a mixture of isopropanol with 0.04 N HCl and 50 μ L of 0.25% SDS. Using an ELISA plate reader, the absorbance was measured at 590 nm.

2.10 SDS-PAGE

Different fractions collected in the ultra-filtration step and the cell-free supernatants were subjected to SDS- PAGE, with 12% and 5% acrylamide as the resolving gel and stacking gel, respectively. With the current flow of 120 V after the run, the gel was exposed to Coomassie staining for visualization.

3. Results and Discussion

3.1 Media optimization for anti-candida activity

The influence of nitrogen and carbon sources on the growth rate of cell production was evaluated using different production mediums [28]. The results revealed that the eupeptic digest of animal tissue, pancreatic digest of casein, and dextrose were good sources for growing *Bacillus* species. A total of 27 combinations were prepared for the experiment (as shown in Table 1) to check the growth of *Bacillus* species. Each culture produced a different pattern of growth in different media. It was

observed that except for MW3, all cultures showed the highest growth when the dextrose concentration was at a minimum (10 g/L). Hence, we can conclude that glucose concentration plays a major role in controlling the growth of these three cultures. Carbon is a major constituent in the culture media. Optimum concentration of glucose is essential for optimum growth, thus an excess carbon source can lead to a different mode of growth in the microorganisms. A similar condition was observed by Stamenković-Stojanović *et al.* [29] with *Bacillus subtilis* culture. However, with respect to the peptic digest of animal tissue (peptone) and pancreatic digest of casein, cultures showed a different pattern. For MW 3, the maximum growth was observed when the peptone and pancreatic digest of casein were highest (7.5 g/L), and for MW 9, the concentrations that gave maximum growth were 2.5 and 5 g/L. MW 19 and 27 showed maximum growth with 7.5g/L peptone and 2.5 g/L pancreatic digests of casein. Media optimization is important to check for the enhancement of the production of biomolecules responsible for inhibiting pathogenic bacteria [30]. Accordingly, the maximum growth for MW3 was observed in formulation 18, and for MW9, it was seen in formulation 4. However, MW19 and MW27 showed maximum growth in formulation 19.

3.2 MIC and anti-candida activity

The MIC values of each supernatant were measured by comparing the optical density (OD at 600 nm) of *Candida* culture exposed to different dilutions of culture supernatant after growing them in the optimized medium. The OD was measured at 0 h and 3 h of incubation at 37°C using an ELISA plate reader at 600 nm. The MIC of different cultures is shown in Table 2. The supernatant (S) and the *Bacillus* culture were tested against *Candida* cultures. The results showed that the MW3, MW9, MW19, and MW27 cultures were able to inhibit *C. albicans* NCIM 3557 with inhibition zones of 0.6 cm, 0.8 cm, 1.0 cm, and 1.5 cm, respectively (Figure 1). The lowest concentration of an antimicrobial compound that inhibits the visible growth of a microorganism is an important step to avoid the development of antibiotic resistance [31].

3.3 Effect of proteolytic enzyme

The supernatants of the MW27, MW3, and MW9 cultures completely lost their anti-candida activity when they were treated with proteolytic enzyme, while the supernatant of the MW19 culture showed a reduction in anti-candida activity from a 1.5 cm zone of inhibition to 0.6 cm. This indicates that the activity of MW27, MW3, and MW9 may be due to protein-based inhibitors and MW19 may include both protein and other types of molecules (Figure 2). Moreover, supernatants MW19 and MW9 did not lose activity even after treatment with high temperature, indicating the heat stability of the anti-candida molecule. Antimicrobial proteins produced by the cultures are excellent candidates to improve the safety of probiotics. Some bacteriocin-like compounds are sensitive to proteolytic enzymes but resistant to heat at 80°C [32, 33] (Figure 2).

Culture	MIC (Concentration of proteins, μg/μL)		
MW3	64		
MW9	32		
MW19	64		
MW27	128		

Table 2. MIC of <i>Bacillus</i> cultures on selected media f	ormu	lations
--	------	---------



Figure 1. Inhibition of *C. albicans* NCIM 3557 by *Bacillus* cultures
(A) MW3, MW9, MW19, and MW27 showed zones of inhibition of 0.5 cm, 0.6 cm, 0.7 cm, and 1.2 cm against *C. albicans* NCIM 3557. (A and B) Negative control supernatant, (S): MW(S)9, MW(S)19, and MW(S)27 showing zones of inhibition of 0.8 cm, 0.2 cm, and 0.9 cm against *C. albicans* NCIM 3557



Figure 2. Inhibition of *C. albicans* NCIM 3557 by supernatant of MW19 and MW9 before and after treatment, (A) *Bacillus* MW19 and (B) MW9 with the effect proteinase K and temperature against *C. albicans* NCIM 3557

(S: Supernatant, PK: proteinase K, S+P: Supernatant with proteinase K)

3.4 Solvent precipitation method

MW19 supernatant precipitated with different solvents, acetone, methanol, and a mixture of methanol with chloroform precipitates showed activity against *Candida* species. Methanol precipitate showed 0.6 cm inhibition zone, the mixture of methanol with chloroform precipitate showed 0.3 cm and acetone precipitate showed 0.5 cm zone of inhibition (Figure 3).

3.5 Partial purification of protein by ultrafiltration method

The ultra-filtration results showed that the filtrates of 30 kDa and 10 kDa retentate has anti-candida activity. The zone of inhibition was 0.7 cm for the 30 kDa retentate and 0.5 cm for the 10 kDa retentate. There was no activity found in the 10 kDa filtrate and 4 kDa filtrate, as well as in the retentates. This indicates that active molecule was retained between the 30 to the 10 kDa fractions (Figure 4).



Figure 3. Zone of inhibition of *C. albicans* NCIM 3557 by various fractions of solvent precipitation

(A) MW19S: Supernatant, M: Methanol, MW19 M+P: Methanol precipitated MW19M+C+P: Methanol chloroform precipitate, MW19M+C: Methanol and TCA

(B) AC19 20: Acetone (20%) with precipitated of 19, AC19 30: Acetone (30%) with precipitated of 19, AC19 60: Acetone (60%) with precipitated of 19, AC19 100: Acetone (100%) with precipitated of 19, AC100: Acetone (100%)



Figure 4. Zone of inhibition of *C. albicans* NCIM 3557 by different ultrafiltration fraction S: supernatant, 1: 30 kDa retentate, 2: 10 kDa retentate, 3: 10 kDa filtrate, 4: 4 kDa retentate, and 5: 4 kDa filtrate

3.6 Germ tube inhibition

Germ tube formation is a very important stage in which the *Candida* culture is in the virulence stage, and so it is important to inhibit the formation of germ tubes in *C. albicans*. Germ tube inhibition assays were carried out with bacterial suspension precipitates and extracted compounds. The bacterial suspensions of MW9, MW19, MW27, MW3, and the acetone precipitates, methanol precipitates, extracted compounds, and supernatants were capable of maximum inhibition of germ tube formation (Table 3).

MW3	MW9	MW19	MW27	Extracted compound		Acetone	Methanol	Clotrimazole	Ethanol
					<30kDa				
				<10kDa					
97%	99%	99%	99%	89%	80%	77%	77%	30%	0%

-	<u> </u>			
Table 3	(form to	1ha 1n	hihition	norcontogo
I ADDE J		псш		DELCEIIIA9E
				percence

3.7 SDS PAGE

By visual inspection of the patterns of proteins, all the examined partially filtered compound bands were observed. Although the different fractions have shown anti-candida activity, the SDS PAGE (Figure 5) showed that the anti-candida molecule may be between 30 to 10 kDa size. Further studies are required to understand the actual molecular weight of the anti-candida molecule.



Figure 5. SDS PAGE of cell free supernatant of *Bacillus* culture after partial purification Well (1) ladder: contains the perfect protein marker from 10 kDa -250 kDa, Well (2) 30 kDa retentate, Well (3) 30 kDa filtrate, Well (4) 10 kDa retentate, and Well (5) un-inoculated media

4. Conclusions

The main contribution of this study was the design of cost-effective production media that could be explored for use by industries for the production of probiotic *Bacillus* culture with anti-candida activity. The growth media containing lower carbon content showed high viable counts and were suitable for culturing *Bacillus* isolates. The media composition number 18 showed the best growth for MW3, while composition number 4 showed the best growth for MW 9. Composition number 19 showed the best growth for MW19 and MW27. A lower sugar concentration in the media was similar to or better than in MRS and SDB. Moreover, the active ingredient was partially purified using solvent precipitation and ultra-filtration methods after growing the culture in the optimized medium. The activity of the compound produced by MW19 was stable even after proteinase K and high-

temperature treatment. Even the partially purified compound showed anti-candida activity and could inhibit the formation of germ tubes in *Candida albicans*. Ultra-filtration results indicated that the active compound was between 30 to 10 kDa in size. However, further studies are required to more precisely calculate the molecular weight of the active molecule.

5. Acknowledgements

The authors are indebted to BV- Rajiv Gandhi Institute of IT and Biotechnology, Bharati Vidyapeeth Deemed University (BVDU), Pune, for the permission to undertake this work.

References

- [1] Calderone, R.A., Fonzi, W.A. and Fonzi, W.A., 2001. Virulence factors of *Candida albicans*. *Trends in Microbiology*, 9(7), 327-335.
- [2] Falagas, M.E., Apostolou, K.E. and Pappas, V.D., 2006. Attributable mortality of candidemia: a systematic review of matched cohort and case-control studies. *European Journal of Clinical Microbiology and Infectious Diseases*, 25(7), 419-425.
- [3] Douglas, L.J., 2003. *Candida* biofilms and their role in infection. *Trends in Microbiology*, 11(1), 30-36, DOI: 10.1016/S0966-842X (02)0002-1.
- [4] Gibson, G. and Wang, X., 1994. Regulatory effects of bifidobacteria on the growth of other colonic bacteria. *Journal of Applied Bacteriology*, 77(4), 412-420.
- [5] Sobel, J., 2013. Factors involved in patient choice of oral or vaginal treatment for vulvovaginal candidiasis. *Patient Preference and Adherence*, 31, 31-34.
- [6] FAO/WHO, 2002. *Guidelines for the Evaluation of Probiotics in Food*. London, Ontario: Food and Agriculture Organization of the United Nations and World Health Organization.
- [7] Kathade, S.A., Aswani, M.A., Anand, P.K., Jagtap, S. and Bipinraj, N.K., 2020. Isolationof *Lactobacillus* from donkey dung and its probiotic characterization. *Korean Journal of Microbiology*, 56(2), 160-169.
- [8] Terpou, A., Papadaki, A., Lappa, I.K., Kachrimanidou, V., Bosnea, L.A. and Kopsahelis, N., 2019. Probiotics in food systems: Significance and emerging strategies towards improved viability and delivery of enhanced beneficial value. *Nutrients*, 11(7), DOI: 10.3390/nu11071591.
- [9] Kathade, S.A., Aswani, M.A., Anand, P.K. and Kunchiraman, B.N., 2020. Probiotic characterization and cholesterol assimilation ability of *Pichia kudriavzevii* isolated from the gut of the edible freshwater snail '*Pila globosa*'. *Egyptian Journal of Aquatic Biology and Fisheries*, 24(7), 23-39.
- [10] Majeed, M., Majeed, S., Nagabhushanam, K., Natarajan, S. and Ali, F., 2016. Evaluation of the stability of *Bacillus coagulans* MTCC 5856 during processing and storage of functional foods. *International Journal of Food Science and Technology*, 51(4), 894-901.
- [11] Kathade, S., Aswani, M., Anand, P.K. and Nirichan, B., 2020. Probiotic characterization and cholesterol assimilation ability of *Pichia kudriavzevii* isolated from the gut of the edible freshwater snail '*Pila globosa*,'. *Egyptian Journal of Aquatic Biology and Fisheries*, 24(7), 23-39.
- [12] Varela, H., Ferrari, M.D., Belobrajdic, L., Weyrauch, R. and Loperena, L., 1996. Effect of medium composition on the production by a new *Bacillus subtilis* isolate of protease with promising unhairing activity. *World Journal of Microbiology and Biotechnology*, 12(6), 643-645.

- [13] Vinderola, G., Perdigón, G., Duarte, J., Farnworth, E. and Matar, C., 2006. Effects of the oral administration of the exopolysaccharide produced by *Lactobacillus kefiranofaciens* on the gut mucosal immunity. *Cytokine*, 36(5-6), 254-260.
- [14] Lefevre, M., Racedo, S.M., Ripert, G., Housez, B., Cazaubiel, M., Maudet, C. and Urdaci, M.C., 2015. Probiotic strain *Bacillus subtilis* CU1 stimulates immune system of elderly during common infectious disease period: a randomized, double-blind placebo-controlled study. *Immunity and Ageing*, 12(1), DOI: 10.1186/s12979-015-0051-y.
- [15] Shobharani, P., Padmaja, R.J. and Halami, P.M., 2015. Diversity in the antibacterial potential of probiotic cultures *Bacillus licheniformis* MCC2514 and *Bacillus licheniformis* MCC2512. *Research in Microbiology*, 166(6), 546-554.
- [16] Marwick, J.D., Wright, P.C. and Burgess, J.G., 1999. Bioprocess intensification for production of novel marine bacterial antibiotics through bioreactor operation and design. *Marine Biotechnology*, 1(6), 495-507.
- [17] Singh, V., Khan, M. and Khan, S., 2009. Optimization of actinomycin V production by *Streptomyces triostinicus* using artificial neural network and genetic algorithm. *Applied Microbiology and Biotechnology*, 82(2), 379-385.
- [18] Sharma, D., Shekhar, S.K., Alok, K. and Godheja, J., 2020. Isolation, characterization, production and purification of fibrinolytic enzyme natto kinase from *Bacillus subtilis*. *International Journal of Pharmaceutical Sciences and Research* 11(4),1768-1776, DOI: 10.13040/IJPSR.0975-8232.
- [19] Lei, J., Sun, L., Huang, S., Zhu, C., Li, P., He, J., Mackey, V., Coy, D.H. and He, Q., 2019. The antimicrobial peptides and their potential clinical applications. *American Journal of Translational Research*, 11(7), 3919-3931.
- [20] Silva, D.R., Sardi, J.C.O., Pitangui, N.S., Roque, S.M., Silva, A.C.B. and Rosalen, P.L., 2020. Probiotics as an alternative antimicrobial therapy: Current reality and future directions. *Journal of Functional Foods*, 73, DOI: 10.1016/j.jff.2020.104080.
- [21] Sarkar, T., Chetia, M. and Chatterjee, S., 2021. Antimicrobial peptides and proteins: From nature's reservoir to the laboratory and beyond. *Frontiers in Chemistry*, 9, DOI: 10.3389/fchem.2021.691532.
- [22] Palande, V., Meora, R., Sonavale, R., Makashir, M., Modak, M., Kapse, N., Dhakephalkar, P. and Ranjekar, P., 2015. Inhibition of pathogenic strains of *Candida albicans* and non-albicans by *Bacillus* species isolated from traditional Indian fermented food preparations. *International Journal of Current Microbiology and Applied Sciences*, 4(3), 691-699.
- [23] Anand, P., Aswani, M., Kathade, S., Kale, A.B. and Kunchiraman, B.N., 2020. Probiotic characterization of anti-candida *Bacillus*. *Research Journal of Biotechnology*, 15(11), 22-29.
- [24] Li, Y., Wang, Y., Liu, Y., Li, X., Feng, L. and Li, K., 2022. Optimization of an economical medium composition for the coculture of *Clostridium butyricum* and *Bacillus coagulans*. *AMB Express*, 12(1), DOI: 10.1186/s13568-022-01354-5.
- [25] Chelliah, R., Ramakrishnan, S.R., Prabhu, P.R. and Antony, U., 2016. Evaluation of antimicrobial activity and probiotic properties of wild-strain *Pichia kudriavzevii* isolated from frozen idli batter. *Yeast*, 33(8), 385-401.
- [26] Aswani, M.A., Kathade, S.A., Anand, P.K., Kunchiraman, B.N., Dhumma, P.R. and Jagtap, S.D., 2021. Probiotic characterization of cholesterol-lowering *Saccharomyces cerevisiae* isolated from frass of *Pyrrharctia isabella* caterpillars. *Applied Food Biotechnology*, 8(3), 189-198.
- [27] Baindara, P., Mandal, S.M., Chawla, N., Singh, P.K. and Pinnaka, A.K., 2013. Characterization of two antimicrobial peptides produced by a halotolerant *Bacillus* subtilis strain SK. DU. 4 isolated from a rhizosphere soil sample. *AMB Express*, 3(1), DOI: 10.1186/2191-0855-3-2.
- [28] Jawan, R., Abbasiliasi, S., Tan, J.S., Mustafa, S., Halim, M. and Ariff, A.B., 2020. Influenceof

culture conditions and medium compositions on the production of bacteriocin-like inhibitory substances by *Lactococcus lactis* Gh1. *Microorganisms*, 8(10), DOI: 10.3390/microorganisms8101454.

- [29] Stamenković-Stojanović, S., Karabegović, I., Beškoski, V., Nikolić, N. and Lazić, M., 2020. *Bacillus* subtilis NCIM2063 batch cultivation: The influence of the substrate concentration and oxygen transfer rate on the biomass yield. *Advanced Technologies*, 9(1), 44-49, DOI: 10.5937/savteh2001044S.
- [30] Palla, M.S., Guntuku, G.S., Sahu, P.K., Kota, P. and Panda, J., 2020. Statistical optimization of anticandida metabolite production process using *Streptomyces hydrogenans* strain from mangrove soils. *SN Applied Sciences*, 2(11), DOI: 10.1007/s42452-020-03734-7.
- [31] Andrews, J.M., 2001. Determination of minimum inhibitory concentrations. *Journal of Antimicrobial Chemotherapy*, 48 (Suppl. 1), 5-16, DOI: 10.1093/jac/48.suppl_1.5.
- [32] Ullivarri, M.F., Arbulu, S., Garcia-Gutierrez, E. and Cotter, P.D., 2020. Antifungal peptides as therapeutic agents. *Frontiers in Cellular and Infection Microbiology*, 10, DOI: 10.3389/fcimb.2020.00105.
- [33] Kathade, S.A., Aswani, M.A., and Anand, P.K., Kale, A., Shrivastava, P., Sharma, S., Badat, U., Mohite, J., Debbarma, J., Sangma, A., Wajravad, B., Jagtap, S. and Niricharan, K.B., 2022. Isolation, characterization, and diversity of probiotic microorganisms from different postpartum milk of various animals. *International Journal of Health Sciences and Research*, 12(3), 223-234.