



Biotin and Zn²⁺ Increase Xylitol Production by *Candida tropicalis*

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Abstract In this study, the medium requirements to increase the production of xylitol by using *Candida tropicalis* (CT) have been investigated. The technique of single addition or omission of medium components was applied to determine the nutritional requirements. The addition of amino acids such as Asp, Glu, Gln, Asn, Thr, and Gly had no significant effect on CT growth. However, in the absence of other metal ions, there was a higher concentration of cell growth and xylitol production when only Zn²⁺ was present in the medium. The analysis of various vitamins unveiled that biotin and thiamine were the only vitamins required for the growth of CT. Surprisingly, when only biotin was present in the medium as a vitamin, there was less growth of CT than when the medium was complete, but the amount of xylitol released was significantly higher. Overall, this study will increase the xylitol production using the single omission or addition technique.

Keywords Biotin · Thiamine · Urea · Xylitol · Zn²⁺

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Introduction

Xylitol, a five-carbon alcoholic sugar, is popular in the food industry due to its non-carcinogenicity and lack of dependence on insulin [1–4]. People with diabetes and people with low glucose-6-phosphate dehydrogenase do not require insulin and glucose-6-phosphate dehydrogenase when consuming xylitol, which makes it a suitable alternative for these people [5, 6]. Xylitol is currently manufactured from xylose, the five-carbon sugar derived from hemicellulose hydrolysates, through a chemical process using Ni/Al₂O₃ as a catalyst [7–12].

The type of media utilized during fermentation greatly influences the chemical or nutritional environment, which in turn significantly influences the productivity and the economics of a fermentation process [13–18]. The media used to promote high productivity in commercial/industrial fermentation are mainly developed from complex sources of carbon and nitrogen [19]. However, the rate of growth and the activity of metabolic processes may be affected strongly by the type and ratio of nutrients provided to the culture. Due to this inherent inconsistency of natural origin, the fermentation performance may vary from lot to lot [20].

Yeast extract (YE) is used in the fermentation of xylitol, and thus xylitol is produced using microbial fermentation using YE as one of the medium ingredients. However, the variation yields have been observed due to unknown variations of YE. On top of that, the high price of YE hinders its uses in industrial applications. Moreover, the medium cost is one of the major factors in economic xylitol production [21–26]. Therefore, replacing YE with a synthetic defined medium (SDM) is required to lower the cost of medium and performance variability while maintaining the production yield [27]. Additionally, when producing xylitol, the performance consistency of synthetic-defined

media should be comparable to that of YE. When the concentration of nitrogen is high, the recovery and purification of xylitol production become very difficult. Xylitol purification is simplified because no additional contaminants are added to the media, leading to a lower production cost [28]. Hence, it's indeed crucial for the metabolic investigation to have a precise growth medium for the microorganisms, which really supports high yield and productivity. Therefore, SDM which enables exponential growth with high xylitol production while eliminating or adding the need for a single medium component, has been developed in the current study.

Materials and Methods

Microorganism and Media

Candida tropicalis KFCC-10690 was used in this study because it is an established member of the *Candida* genus [29]. Freezing of the cell stock was done at $-70\text{ }^{\circ}\text{C}$. This medium consists of 5 g/L YE and 20 g/L glucose. For fermentation, there were 5–33% xylose, 10 g/L YE, and 0–90 g/L glucose in a complex medium that had a concentration of 5 g/L KH_2PO_4 . Finished materials were made up into separate batches of medium and dense components, which consisting of carbohydrates, basal salts, amino acids, vitamins, and metals [30–32]. The components of SDM were sterilized by a membrane filtration method [33] (Millex-GV filter; Millipore Corp., Bedford, Mass), and the working cultures of CT were propagated in SDM. Further, cultures were centrifuged and washed twice in 50 mM potassium phosphate and at the pH of 6.5 to elimination of carryover nutrients. For the inocula, 5% (vol/vol) exponentially growing cells were used.

Fermentation Conditions

Inoculation was carried out on a 500 mL flask with 100 mL of culture medium for 10 h at $30\text{ }^{\circ}\text{C}$ and 250 rpm. The resulting culture broth, diluted to a total volume of 10% (v/v), was transferred to a 500 mL flask and used to inoculate a 5-L jar fermenter, which was filled with 100 mL of production medium until it was 2.8–3.5 L of production medium (Kobiotech. Co., Republic of Korea). Complex media contains 200 g/L of xylose, 17 g/L of glucose, 1.3 g/L of KH_2PO_4 , 2.5 g/L of $(\text{NH}_4)_2\text{SO}_4$, 0.13 g/L of MgSO_4 , and 5 g/L of YE. SDM contains urea 3.1 g/L, xylose 200 g/L, glucose 17 g/L, KH_2PO_4 1.3 g/L, MgSO_4 0.13 g/L, biotin 16.5 g/L, folic acid 7.5 $\mu\text{g/L}$, thiamine 2.65 mg/L, boric acid 0.5 mg/L, copper sulfate 0.04 mg/L, potassium iodide 0.1 mg/L, ferric chloride 0.2 mg/L, manganese sulfate 0.4 mg/L, sodium molybdate

0.2 mg/L, and zinc sulfate 5.0 mg/L. Experiments in jar fermenters were conducted at $30\text{ }^{\circ}\text{C}$ in a fed-batch mode controlling the pH at 4.8. The peristaltic pump (10–50 mL/h) continuously fed the solution of xylose or the mixture of xylose and glucose, which was aerated at 0.5 vvm. The agitation was increased from 250 to 750 rpm to maintain the percentage of dissolved oxygen above 20 until the cell mass reached 14 g/L; it was then decreased to 340 rpm to limit the concentration of dissolved oxygen.

Enzyme Assay

Cultured cells were collected by centrifuging at 10,000 rpm for 15 min. Washing was carried out with 0.1 M Tris–HCl (pH 7.8), 0.5 mM EDTA, and 5 mM mercaptoethanol. Further, the cells were resuspended in a buffer [34] containing 20 mM Tris–HCl (pH 7.8), 10 mM MgCl_2 , 1 mM EDTA, 1 mM dithiothreitol, and 1 mM phenylmethylsulphonyl fluoride. Glass beads of 0.5 mm in diameter were used for the suspension. (Sigma, USA). To determine the xylose reductase (XR) activity, a decrease in the absorbance at 340 nm was measured after the addition of D-xylose, a marker for NADPH oxidation (Sigma, USA) [35].

Analytical Methods

A Bradford assay has been used to estimate protein concentration, and bovine serum albumin is being used as a standard [36]. To estimate the concentrations of xylitol, glucose, and xylose, HPLC coupled to an RI detector (Waters 410, USA) and a High-Performance Carbohydrate Column (4.6 mm \times 250 mm, Waters, USA) were used. Acetonitrile/water (85:15 v/v) was used as a mobile phase at a 1.5 mL/min flow rate.

Results and Discussion

Influence of Nitrogen Source

A defined medium with a sole carbon source and a sole nitrogen source was designed in order to investigate the effect of the substrates on xylitol production. A shake flask system was developed to explore a range of inorganic and organic nitrogen sources in culturing a defined medium containing 200 g/L xylose and 17 g/L glucose as the carbon source. Ammonia, which is an important component of nitrogen metabolism in yeast, was also tested alongside two other common nitrogen sources, urea and nitrate, and CT grew on all of the nitrogen sources, indicating that they had been consuming it. With the exception of the ammonium acetate experiment, all experiments found that the glucose

supply had been depleted after 40 h. Although inorganic nitrogen sources like ammonium tartrate, ammonium nitrate, ammonium acetate, and sodium nitrate were consumed for biomass formation, the production of xylitol was poor after 60 h of cultivation. However, urea has been found to produce xylitol similar to the level achieved by complex media by YE.

Effect of Amino Acids, Nucleic Acids, and Buffers

The cell growth and the amount of xylitol did not change regardless of whether single or multiple amino acids were added, and the same results have been observed when nucleic acids such as guanine, xanthine, adenine, and uracil were omitted from the growth medium. Moreover, no major changes were found in the growth and xylitol production of the strain when tenfold lower levels of buffers, such as phosphate, citrate, and acetate, were added. (Table 1).

Effect of Metal Ions on the Growth and Xylitol Production

By excluding one metal ion at a time, the metal ion requirement of CT in SDM was determined. The strain

grew well when NiCl₂ and CoCl₂ were omitted individually, and the cell growth was slightly inhibited when FeCl₂, CoCl₂, H₃BO₃, Na₂MoO₄, MnCl₂, and CuCl₂ were excluded. However, in the absence of ZnCl₂, the growth of CT was significantly decreased, and it appeared to be crucial for cell growth (Table 1). Thus, to determine the effect of Zn²⁺ on cell growth and xylitol production, a 5-L jar fermenter was used. To that end, various concentrations of ZnCl₂ were tested in the range of 0 to 10 mg/L. The fermentation conditions were provided in the “Materials and Methods” section. Xylitol gave the greatest yield and productivity at a concentration of 5 mg/L of ZnCl₂ (Table 2). Further, an SDM mixture produced by the addition of optimal concentration of ZnSO₄ (5 mg/L) was as effective as the complete metal mixture of SDM has been observed to promote cell growth and xylitol production. XR activity of supernatants obtained from cultures grown without zinc were assayed with and without ZnCl₂ added to the samples, and no significant activities were found, even when the samples were incubated with zinc for 1 h at 37 °C before the assay. This suggests that zinc does not increase the level of XR activity by an enzyme mechanism. However, it seems to play a metabolic role and is needed during growth to induce significant protease production [37].

Table 1 Nutrient requirements of *C. tropicalis* in synthetic defined medium investigated by addition or omission of a single medium component

Added medium component	OD ₆₀₀ ^a	Omitted medium component	OD ₆₀₀ ^a
None	14.4	Phosphate	11.2
L-Alanine	14.3	MgSO ₄	13.0
L-Arginine	14.7	H ₃ BO ₃	13.3
L-Asparagine	14.7	MnCl ₂	12.4
L-Leucine	14.4	ZnCl ₂	5.6
L-Glutamic acid	14.5	CuSO ₄	13.1
L-Glutamine	14.3	FeCl ₂	13.5
Glycine	14.4	NiCl ₂	14.6
L-Lysine	14.6	CoCl ₂	14.3
L-Phenylalanine	14.3	Na ₂ MoO ₄	13.5
L-Proline	14.3	Biotin	13.4
L-Serine	14.5	Inositol	15.1
L-Tryptophan	14.7	Folic acid	14.5
L-Tyrosine	14.7	ρ-Aminobenzoic acid	14.9
L-Valine	14.5	Nicotinic acid	14.4
L-Histidine	14.4	Pantothenate	14.8
L-Cysteine	14.6	Pyridoxamine	15.0
Adenine	14.8	Pyridoxine	15.2
Guanine	14.2	Riboflavin	10.5
Uracil	14.5	Thiamine	8.7
Xanthine	14.3		

Values are the means ± standard deviations of triplicate measurements

^aOD measurements were performed after 48 h of incubation

Table 2 Effect of ZnSO₄ concentration on the cell growth and xylitol production

	Conc. of ZnSO ₄ (mg/L)			
	0	1.0	5.0	10.0
Cell conc. (OD ₆₀₀)	47.9	47.7	43.4	38.5
Produced xylitol (g/L)	186	260	252	246
Yield (Volumetric, %)	54.3	77.1	77.2	74.7
Productivity (g/L h)	2.10	3.31	3.63	3.32

Effect of Vitamins on the Growth and Xylitol Production

In the absence of any individual vitamin, only riboflavin and thiamine were detected as essential nutrients for growth. In contrast, a similar OD was found when other vitamins were overlooked. Further, we found that folic acid, ρ -Aminobenzoic acid, pantothenate, inositol, niacin, and pyridoxine were unnecessary for cell growth. A single omission of the nonessential vitamins did not change the specific production of xylitol [37, 38], but when biotin was overlooked, the specific xylitol production was significantly decreased, and it emerged to be essential for xylitol production (Table 3). When CT was grown in a medium lacking vitamins except for riboflavin, biotin, and thiamine, the OD of the cultures was getting lower after 48 h. In contrast, the specific xylitol production was increased significantly (Table 3). Biotin limitation decreased the xylose consumption of CT, and the decrease became more significant as the initial concentration of biotin decreased.

Biotin acts as a prosthetic group for carboxylases, and it is unclear why its limitation results in more xylitol accumulation.

Xylitol Production Using a 5-L Jar Fermenter

Finally, a pH-controlled fed-batch culture experiment was carried out to compare the cell growth and xylitol production by CT in the complex medium and the SDM. The composition of the SDM met the nutritional requirements of the strains and took advantage of the beneficial effects in the downstream process. Fermentation of CT in complex medium containing is represented in Fig. 1. During the glucose consumption, pH decreased from 6.4 to 4.8; thereafter, pH increased to 6.8 until the end of the fermentation process. In the SDM, CT grew up with a minimal growth rate of 0.18 h⁻¹ and a final OD of 47.7 and produced 260 g of xylitol per liter with a conversion yield of 81.5% when grown in a pH-controlled fed-batch culture. On the other hand, in the complex medium, the growth rate was 0.23 h⁻¹, and the final OD was 47.1, while the xylitol production was 251 g L⁻¹ with the conversion yield of 78.1% (Table 4). Further, It has been observed that in both the complex medium and the SDM, xylitol production continued after growth had come to an end. Still, beyond the stationary growth phase, more xylitol production was observed in the SDM than in the complex medium. Moreover, the addition to SDM of the ten amino acids (Gln, Leu, Ile, Val, Met, His, Arg, Trp, Pro, and Phe) did not increase the xylitol production of CT.

Table 3 Effect of individual and multiple omission of essential vitamins on the xylitol production by *C. tropicalis* in defined medium

Omission	Xylitol (g/L)	Specific xylitol production (g/g of dry cell weight) ^a
None	114	10.6
ρ -Aminobenzoic acid	117	10.8
Biotin	2.8	0.27
Calcium pantothenate	118	11.0
Folic acid	114	10.5
Inositol	112	10.2
Niacin	118	10.8
Pyridoxine	113	10.2
Riboflavin	108	10.1
Thiamine	103	9.90
All essential vitamins ^b except riboflavin, biotin, and thiamine	119	11.2

^aThe specific xylitol production was calculated from a standard curve of OD₆₀₀ against cell dry weight

^bAll essential vitamins: ρ -Aminobenzoic acid, biotin, calcium pantothenate, folic acid, inositol, niacin, pyridoxine hydrochloride, riboflavin, thiamine

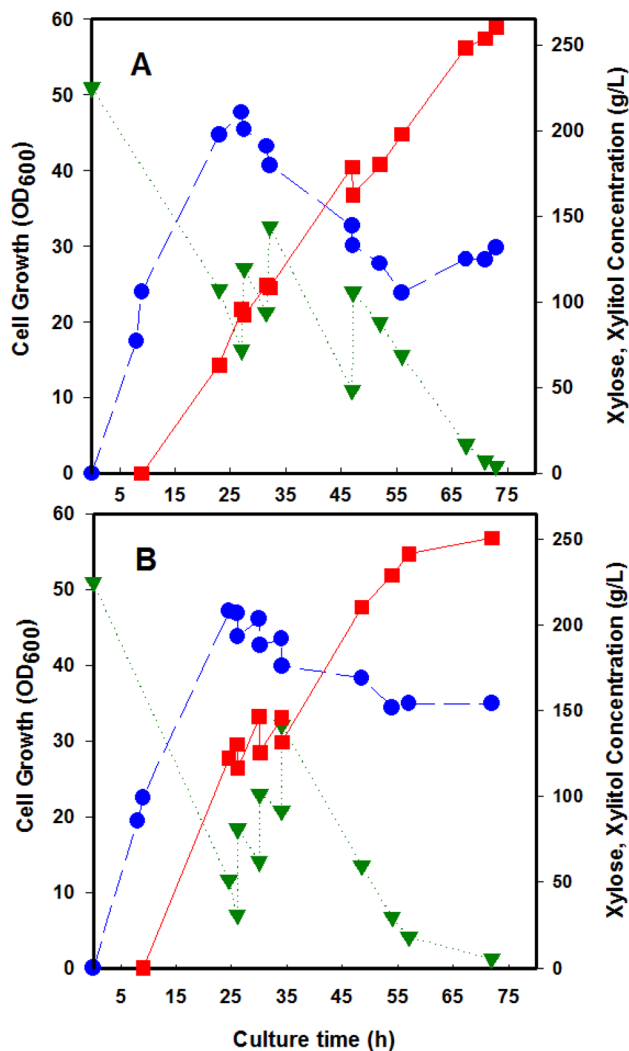


Fig. 1 Xylitol production using SDM (A) and complex medium (B) containing urea and yeast extract as a nitrogen source, respectively. Blue circle: cell growth, green inverted triangle: xylose, red square: xylitol

Conclusions

In this work, we developed a synthetic and cheap medium to allow reproducible xylitol production without variation in yields and productivity. The technique of single addition or omission of medium components revealed that the amino acids such as Asp, Glu, Gln, Asn, Thr, and Gly were slightly affecting the growth of CT. Further, we observed that the amount of cell growth and xylitol production was more significant when Zn^{2+} ion was present in the medium and other metal ions were not. In addition, it has been observed that CT required only biotin and thiamine as individual vitamins. Surprisingly, when only biotin was present in the medium as a vitamin, the amount of xylitol production was significantly greater than in the complete medium.

Table 4 Comparison of xylitol production between the SDM and complex medium containing urea and yeast extract as a nitrogen source, respectively

	Medium	
	SDM	Complex
Initial pH of medium	5.6	5.0
Culture time (h)	73	72
Maximum cell conc. (OD_{600})	47.7	47.1
Added conc. of xylose (g/L)	300	300
Final conc. of		
Xylitol (g/L)	260	251
Xylose (g/L)	4.3	5.2
Glycerol (g/L)	9.5	11.0
Yield (Volumetric, %)	81.5	78.1
Productivity (Volumetric, g/L h)	3.35	3.26

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References

- Makinen KK (1979) Xylitol and oral health. *Adv Food Res* 25:137–157. [https://doi.org/10.1016/S0065-2628\(08\)60236-0](https://doi.org/10.1016/S0065-2628(08)60236-0)
- Ylikahri R (1979) Metabolic and nutritional aspects of xylitol. *Adv Food Res* 25:159–180. [https://doi.org/10.1016/S0065-2628\(08\)60237-2](https://doi.org/10.1016/S0065-2628(08)60237-2)
- Radhakrishnan R, Lee I-J (2017) Foliar treatment of *Bacillus methylotrophicus* KE2 reprograms endogenous functional chemicals in sesame to improve plant health. *Indian J Microbiol* 57:409–415. <https://doi.org/10.1007/s12088-017-0666-0>
- Soares LCS, Chandel AK et al (2016) Screening of yeasts for selection of potential strains and their utilization for in situ microbial detoxification (ISMD) of sugarcane bagasse hemicellulosic hydrolysate. *Indian J Microbiol* 56:172–181. <https://doi.org/10.1007/s12088-016-0573-9>
- Pepper T, Olinger PM (1988) Xylitol in sugar-free confections. *Food Technol* 42:98–106
- Gao H, Kim TS, Mardina P, Zhou P, Wen F, Lee J-K (2016) Rare sugar production by coupling of NADH oxidase and L-arabinitol dehydrogenase. *RSC Adv* 6:66609–66616. <https://doi.org/10.1039/C6RA11614K>
- Dhiman SS, Haw JR, Kalyani D, Kalia VC, Kang YC, Lee J-K (2015) Simultaneous pretreatment and saccharification: green technology for enhanced sugar yields from biomass using a fungal consortium. *Bioresour Technol* 179:50–57. <https://doi.org/10.1016/j.biortech.2014.11.059>
- Prabhu P, Doan TN, Tiwari M, Singh R, Kim SC, Hong MK, Kang YC, Kang LW, Lee J-K (2014) Structure-based studies on the metal binding of two-metal-dependent sugar isomerases. *FEBS J* 15:3446–3459. <https://doi.org/10.1111/febs.12872>

9. Jagtap SS, Dhiman SS, Kim TS, Li J, Lee J-K (2013) Enzymatic hydrolysis of aspen biomass into fermentable sugars by using lignocellulases from *Armillaria gemina*. *Bioresour Technol* 133:307–314. <https://doi.org/10.1016/j.biortech.2013.01.118>
10. Jagtap SS, Dhiman SS, Jeya M, Kim I-W, Lee J-K (2013) Characterization of a β -1,4-glucosidase from a newly isolated strain of *Pholiota adiposa* and its application to the hydrolysis of biomass. *Biomass Bioenergy* 54:181–190. <https://doi.org/10.1016/j.biombioe.2013.03.032>
11. Hui G, Tiwari M, Jeya M, Lee J-K (2012) Characterization of H₂O-forming NADH oxidase from *Streptococcus pyogenes* and its application in L-rare sugar production. *Bioorg Med Chem Lett* 22:1931–1935. <https://doi.org/10.1016/j.bmcl.2012.01.049>
12. Jeya M, Nguyen N-P-T, Moon H-J, Kim S-H, Lee J-K (2010) Conversion of woody biomass into fermentable sugars by cellulase from *Agaricus arvensis*. *Bioresour Technol* 101:8742–8749. <https://doi.org/10.1016/j.biortech.2010.06.055>
13. Kim JH, Lim B-C, Yeom S-J, Kim Y-S, Kim HJ, Lee J-K, Lee SH, Kim SW, Oh DK (2008) Differential selectivity of the *Escherichia coli* cell membrane shifts the equilibrium for the enzyme-catalyzed isomerization of galactose to tagatose. *Appl Environ Microbiol* 74:2307–2313. <https://doi.org/10.1128/AEM.02691-07>
14. Pagolu R, Singh R, Shanmugam R, Kondaveeti S, Patel SKS, Kalia VC, Lee J-K (2021) Site-directed lysine modification of xylanase for oriented immobilization onto silicon dioxide nanoparticles. *Bioresour Technol* 331:125063. <https://doi.org/10.1016/j.biortech.2021.125063>
15. Fernandes AMO, Garcia NFL, Fonseca GG, Leite RSR, Da Paz MF (2020) Evaluation of the fermentative capacity of *Saccharomyces cerevisiae* CAT-1 and BB9 strains and *Pichia kudriavzevii* BB2 at simulated industrial conditions. *Indian J Microbiol* 60:494–504. <https://doi.org/10.1007/s12088-020-00891-6>
16. Veerasamy M, Venkataraman K et al (2018) Point of care tuberculosis sero-diagnosis kit for wild animals: combination of proteins for improving the diagnostic sensitivity and specificity. *Indian J Microbiol* 58:81–92. <https://doi.org/10.1007/s12088-017-0688-7>
17. Xue D, Yao D, You X, Gong C (2020) Green synthesis of the flavor esters with a marine *Candida parapsilosis* esterase expressed in *Saccharomyces cerevisiae*. *Indian J Microbiol* 60:175–181. <https://doi.org/10.1007/s12088-020-00856-9>
18. Kalia VC, Patel SKS, Cho B-K, Wood TK, Lee J-K (2021) Emerging applications of bacteria as anti-tumor agents. *Sem Cancer Biol*. <https://doi.org/10.1016/j.semcancer.2021.05.012>
19. Miller TL, Churchill BW (1986) Substrates for large-scale fermentations. In: Demain AL, Solomon NA (eds) *Manual for industrial microbiology and biotechnology*. American Society for Microbiology, Washington, DC, pp 122–136
20. Kumar V, Patel SKS, Gupta RK, Otari SV, Hui G, Lee J-K, Zhang L (2019) Enhanced saccharification and fermentation of agricultural waste using an immobilized enzyme cocktail. *Biotechnol J* 14:1800468. <https://doi.org/10.1002/biot.201800468>
21. Tiwari M, Moon H-J, Jeya M, Lee J-K (2010) Cloning and characterization of a thermostable xylitol dehydrogenase from *Rhizobium etli* CFN42. *Appl Microbiol Biotechnol* 87:571–581. <https://doi.org/10.1007/s00253-010-2478-6>
22. Zhang Y-W, Tiwari M, Jeya M, Lee J-K (2011) Covalent immobilization of recombinant *Rhizobium etli* CFN42 xylitol dehydrogenase onto modified silica nanoparticles. *Appl Microbiol Biotechnol* 90:499–507. <https://doi.org/10.1007/s00253-011-3094-9>
23. Kalia VC, Gong C, Patel SKS, Lee J-K (2021) Regulation of plant mineral nutrition by signal molecules. *Microorganisms* 9:774. <https://doi.org/10.3390/microorganisms9040774>
24. Patel SKS, Gupta RK, Kalia VC, Lee J-K (2021) Integrating anaerobic digestion of potato peels to methanol production by methanotrophs immobilized on banana leaves. *Bioresour Technol* 323:124550. <https://doi.org/10.1016/j.biortech.2020.124550>
25. Hong JH, Kim JH, Park GD, Lee JY, Lee J-K, Kang YC (2021) A strategy for fabricating three-dimensional porous architecture comprising metal oxides/CNT as highly active and durable bifunctional oxygen electrocatalysts and their application. *Chem Eng J* 414:128815. <https://doi.org/10.1016/j.cej.2021.128815>
26. Kalia VC, Patel SKS, Shanmugam R, Lee J-K (2021) Polyhydroxy alkanooates: trends and advances towards biotechnological applications. *Bioresour Technol* 326:124737. <https://doi.org/10.1016/j.biortech.2021.124737>
27. Kalyani D, Tiwari M, Li J, Kim SC, Kalia VC, Kang YC, Lee J-K (2015) A highly efficient recombinant laccase from the yeast *Yarrowia lipolytica* and its application in the hydrolysis of biomass. *PLoS ONE* 10:e0120156. <https://doi.org/10.1371/journal.pone.0120156>
28. Kumar P, Ray S, Patel SKS, Lee J-K, Kalia VC (2015) Bio-conversion of crude glycerol to polyhydroxyalkanoate by *Bacillus thuringiensis* EGU45 under high nitrogen concentration. *Int J Biol Macromol* 78:9–16. <https://doi.org/10.1016/j.ijbiomac.2015.03.046>
29. Demirbas A (2021) Comparison study of synthesized red (or blood) orange peels and juice extract-nanoflowers and their antimicrobial properties on fish pathogen (*Yersinia ruckeri*). *Indian J Microbiol*. <https://doi.org/10.1007/s12088-021-00945-3>
30. Matthews CB, Kuo A, Love KR, Love JC (2018) Development of a general defined medium for *Pichia pastoris*. *Biotechnol Bioeng* 115:103–113. <https://doi.org/10.1002/bit.26440>
31. Patel SKS, Ray S, Prakash J, Wee JH, Kim S-Y, Lee J-K, Kalia VC (2019) Co-digestion of biowastes to enhance biological hydrogen process by defined mixed bacterial cultures. *Indian J Microbiol*. 59:154–160. <https://doi.org/10.1007/s12088-018-00777-8>
32. Patel SKS, Kim JH, Kalia VC, Lee J-K (2019) Antimicrobial activity of amino-derivatized cationic polysaccharides. *Indian J Microbiol* 59:96–99. <https://doi.org/10.1007/s12088-018-0764-7>
33. Ausubel F, Brent R, Kingston RE, Moore DD, Seidman JG, Smith JA, Struhl K (1986) *Short protocols in molecular biology*, 3rd edn. Wiley, New York, p 836
34. Chiang C, Knight SG (1966) D-Xylose reductase and xylitol dehydrogenase from *Penicillium chrysogenum*. *Methods Enzymol* 9:188–193. [https://doi.org/10.1016/0076-6879\(66\)09044-X](https://doi.org/10.1016/0076-6879(66)09044-X)
35. Singh RK, Singh R, Sivakumar D, Kondaveeti S, Kim T, Li J, Sung BH, Cho B-K, Kim DR, Kim SC, Kalia VC, Zhang Y-HPJ, Zhao H, Kang YC, Lee J-K (2018) Insights into cell-free conversion of CO₂ to chemicals by a multienzyme cascade reaction. *ACS Catal* 8:11085–11093. <https://doi.org/10.1021/acscatal.8b02646>
36. Zhu Y-H, Liu C-Y, Cai S, Guo L-B, Kim I-W, Kalia VC, Lee J-K, Zhang Y-W (2019) Cloning, expression, and characterization of a highly active alcohol dehydrogenase for production of ethyl (S)-4-chloro-3-hydroxybutyrate. *Indian J Microbiol* 59:225–233. <https://doi.org/10.1007/s12088-019-00795-0>
37. Kondaveeti S, Patel SKS, Woo J, Wee JH, Kim S-Y, Al-Raoush RI, Kim I-W, Kalia VC, Lee J-K (2020) Characterization of cellobiohydrolases from *Schizophyllum commune* KMJ820. *Ind J Microbiol* 60:160–166. <https://doi.org/10.1007/s12088-019-00843-9>
38. Lee J-K, Patel SKS, Sung BH, Kalia VC (2020) Biomolecules from municipal and food industry wastes: an overview. *Bioresour Technol* 298:122346. <https://doi.org/10.1016/j.biortech.2019.122346>

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