

Production of extracellular Pullulanase by NP 7 bacteria using OFAT methodology

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Abstract

Pullulanase is a debranching enzyme that splits explicitly α -1,6 glycosidic bonds. It is widely used in starch saccharification, in production of glucose, fructose as well as maltotriose syrup and bioethanol production. Pullulanase is produced by submerged fermentation process that requires optimization of physical and chemical parameters to achieve higher activity and make this process cost effective. The present study aimed the optimization of physical as well as chemical parameters of production medium. The optimization includes physical parameters such as inoculum size, incubation time, temperature, pH and agitation rate while chemical parameters include carbon source, nitrogen source and different metal ions.

From the optimization study, it was found that highest pullulanase production was achieved after 48 hours of incubation at 45°C temperature, pH 7 and 120 rpm. Optimization of chemical parameters reveals that incorporation of starch (1%) and 0.2mMole/L of calcium chloride in the media gave significant increase in enzyme activity. The maximum productivity achieved after optimization was 139.33 U/ml using OFAT method which was four times higher.

Keywords: Pullulanase, saccharification, calcium chloride, starch, pH.

Introduction

Pullulanase (pullulan-6-glucanohydrolase, EC:3.2.1.41) is a debranching enzyme that specifically hydrolyses the α -1,6-glycosidic linkages in pullulan and other branched polysaccharides such as starch⁸ which can lower the dosage of glucoamylase, reduces the reaction time and efficiently improves the conversion rate^{7,23}. Pullulanase hydrolyses pullulan, an α -glucan elaborated by *Pullularia pullulans* into maltotriose units, although traces of higher polysaccharides have also been detected⁸. Bacterial Pullulanase was isolated from *Thermotoga neapolitana*, *Geobacillus thermocatenulatus*, *Bacillus acidopullulyticus*, *Thermococcus marinus* and *Klebsiella variicola*⁵ in recent years. To increase the efficiency and yield of enzymatic reactions with amylolytic enzymes like α -amylase, β -amylase or glucoamylase, addition of pullulanase is the most important in starch processing. Pullulanase enhanced the quality of sugar syrup in the starch saccharification process⁹.

In food and pharmaceutical industries, pullulanase hydrolysed the products of pullulan which are frequently used²³. Pullulanase also improves the production of cyclodextrins by pre-treating the starch prior to the cyclodextrin glucanotransferase (CGTase) reaction¹². Pullulanase also has applications in the textile, baking and detergent industries¹⁰.

Mesophilic, thermophilic and hyperthermophilic bacteria have ability to produce several pullulanase in high amounts⁹. Due to wide application of pullulanase in industry, several studies have been done to enhance enzyme production in bacterial cells by optimizing medium composition as well as physical parameters. The most common parameters are the types of carbon and nitrogen sources used in the fermentation medium. Pullulan, sucrose, maltose, glucose and starch are types of carbon sources previously used to improve pullulanase production^{13,14}.

Additionally, nitrogen sources, including yeast extract, peptone, tryptone, casein, ammonium acetate, ammonium sulphate and sodium nitrate, were found to improve the pullulanase production²⁴.

- Microorganisms require optimum production condition and nutrient to achieve maximum productivity of enzyme. Moreover, media optimization should be carried out to get specific media and condition which is cost effective as submerged fermentation is expensive.
- In the current study, media optimization was carried out using OFAT to achieve higher productivity of enzyme by pullulanase producing bacterial strain NP 7.

Material and Methods

Bacterial culture: For isolation of NP 7 bacterial strain, Soil sample (5 g) was added to 50 ml of medium containing soluble starch 10g/L, yeast extract 0.6g/L, K₂HPO₄ 5 g/L, NaCl 1 g/L, MgSO₄ 0.2 g/L, (NH₄)₂SO₄ 0.6 g/L (pH 7.0) and incubated at 37°C for 5 days. 0.2 ml of sample was spread on nutrient agar plate and incubated at 37°C for 24 to 48 hours. For screening of pullulanase producing microorganisms, a medium (pH 7.0) was required containing pullulan-5, peptone-8, NaCl-2, MgSO₄-0.1, K₂HPO₄-0.17, KH₂PO₄-0.12 (g/L). Pullulanase producers were further screened on different concentrations of pullulan using plate method⁷. The pullulanase producers were isolated at 37° C using pH 7.0. Pullulanase producing miroorganisms were preserved on 0.5% pullulan containing basal medium (Pullulan-5, Peptone-8, NaCl-2, MgSO₄-0.1, K₂HPO₄-0.17, KH₂PO₄-0.12 (g/L) pH-7.0) at 4°C.

Pullulanase production in submerged fermentation

Seed culture: 24-hour-old actively growing bacterial culture was inoculated in 250ml conical flask containing 50ml of 0.5% pullulan containing basal broth. The inoculum medium was incubated at 37 °C in rotary shaking at 120 rpm for 24h.

Pullulanase Production: The turbidity in terms of absorbance was adjusted to ~2.0 at 660 nm and production flasks were inoculated with 1% bacterial culture. Fermentation was carried out in 250-mL flasks containing 100 mL of pullulan medium at 37°C upto 48 hrs at 120 rpm. After interval of 24 hours, samples were harvested and centrifuged at 7500 rpm for 10 min at 4°C. The supernatant was used for determination of pullulanase enzyme activity²³.

Pullulanase enzyme assay: Pullulanase activity was assayed by measuring the reducing sugar (glucose) released from pullulan. The reaction mixture (1ml) containing pullulan (0.5% w/v) in sodium phosphate buffer (0.02 M, pH 7.5) and enzyme was incubated at 40°C for 30 min. The reducing sugar was measured by the dinitrosalicylic acid method according to Miller¹⁷.

The blank contained all the assay constituents except the active enzyme which was replaced by buffer. One unit of pullulanase activity is defined as the amount of enzyme which produces 1μmol of reducing sugar with glucose as standard per min under the optimum conditions².

Characterization of crude pullulanase production: The cell free extract of the bacteria was taken as the crude pullulanase preparation. While studying each parameter, the other reaction conditions were kept constant and after every 24 hours, pullulanase activity was measured up to 72 hours.

Optimization of physical parameters

Effect of inoculum size on pullulanase production: Different concentrations of seed inoculum were inoculated in production medium to check the optimum concentration of culture to achieve higher production.

Different inoculation sizes ranging from 1%, 2 % up to 5% were used in respective production flasks and productivity was checked after every 24 hours using pullulanase assay.

Effect of incubation time on pullulanase production: It has been observed that the culture shows maximum activity at particular time of incubation. The production flask was incubated up to drop in activity and pullulanase activity was measured after every 24 hours.

Effect of Temperature on Pullulanase production: Temperature plays a critical role in production of enzyme as culture shows maximum efficacy for enzyme production at particular temperature. In the present study, production flask was incubated at various temperatures at 30°C, 35°C, 37°C, 40°C, 45°C and 50°C. Pullulanase activity was measured after every 24 hours.

Effect of pH on pullulanase production: Production medium was prepared with varying pH (4-9) in the interval of 1. The flask was incubated for optimized time and at optimized temperature. After every 24 hours, pullulanase activity was observed.

Effect of agitation rate (RPM) on pullulanase production: Agitation rate in submerged fermentation maintains the rheology of medium which is necessary to provide proper dissolved oxygen and even distribution of medium for optimum enzyme production. But high agitation rate results in damage to cell structure and yield of secondary metabolites decreases¹¹. Production flasks were incubated at different RPM ranging from 100 RPM to 180 RPM at the interval of 20 RPM and enzyme activity was measured after every 24 hours.

Optimization of chemical parameters

Effect of Crude substrate on pullulanase production: 0.5% of different crude carbon sources like corn flour, sago starch, rice bran, red gram and wheat bran were used as substrate for pullulanase production. After every 24 hours, enzyme was extracted using centrifugation and assay was carried out.

Effect of different Carbon source on pullulanase production: To check the effect of different carbon sources, 0.5% pullulan was replaced by glucose, sucrose, starch and maltose in the basal medium and enzyme activity was measured after every 24 hours.

Effect of different concentration of pullulan and starch on pullulanase production: The optimum concentration of carbon source is necessary in production medium to achieve optimum growth and productivity. In the present study, varying concentrations of pullulan and starch ranging from 0.1% to 1% with interval of 0.2% were studied and enzyme activity was measured up to 72 hours.

Effect of Nitrogen source on pullulanase production: In the basal medium, peptone was used as nitrogen source which was replaced by yeast extract, tryptone, casein, ammonium acetate, ammonium sulphate and sodium nitrate affecting the growth of bacteria by its crucial need to full the metabolic pathways¹⁸. To check effect of nitrogen source on pullulanase production, enzyme was extracted after every 24 hours and activity was measured.

Effect of different concentration of Peptone on pullulanase production: The optimum concentration of peptone is necessary in production medium that improves and boosts growth of bacteria by increasing the production of essential enzymes¹⁸. The effect of different peptone concentration was tested by adding 0.4 to 2 % peptone with interval of 0.4% in production medium. In the present study, varying concentrations of peptone from 0.4 to 2 % with interval of 0.4% were checked with the combination of pullulan and starch.

Effect of different metal ions on pullulanase production:

The effect of different ions (Ca^{+2} , Cu^{+2} , Co^{+2} , Mn^{+2} , Mg^{+2} , Zn^{+2} , Ni^{+2} , Fe^{+2}) on the pullulanase production was studied at 0.2 mM concentration. These ions were included in the production medium as the CaCl_2 , CuCl_2 , CoCl_2 , MnCl_2 , MgCl_2 , ZnSO_4 , NiSO_4 and FeSO_4 salts and after every 24 hours, pullulanase activity was measured.

Results and Discussion

Bacterial culture: Total 57 bacterial strains were isolated from enriched media and screened for the pullulanase producing capabilities at 37° C and pH 7.5 on plates containing pullulan. Pullulan-degrading microorganisms were identified by the presence of clear zones (haloes) around colonies, after immersing the culture in ethanol (99 % (v/v)) for 3 h^{2,26}. After primary screening, 8 bacterial strains were screened which were selected for secondary screening. In secondary screening, NP 7 bacterial strain gave 33.33 U/ml maximum pullulanase activity among other cultures based on which NP 7 bacterial strain was selected for future work and microscopic observation of culture is shown in fig. 1.

Optimization of physical parameters

Effect of inoculum size: In the present study, production medium inoculated with 3 % inoculum shows maximum productivity at 24 hours which was similar to *Bacillus cereus* FDTA 13¹⁹. Above 3% inoculum activity starts decreasing as rheology of media is not maintained due to overgrowth of culture (Graph 1). An inoculum size of less than 3% was found to be insufficient for growth and enzyme production.

Effect of incubation time: Study of incubation time is necessary for any experiment as it gives an idea of length of entire fermentation process, moreover, it gave an idea of actual time when culture is producing maximum enzyme. In the submerged fermentation, NP 7 shows 41.34 U/ml pullulanase activities at 48 hours which was maximum (Graph 2). *Bacillus cereus* also shows maximum extracellular pullulanase production at 48 h with 0.55U/ml activity²⁵.

Effect of incubation Temperature: In the present experiment, productivity of pullulanase was checked in range of temperature between 30°C to 50°C. The maximum yield of pullulanase was observed at 45°C (35.50 U/ml) (Graph 3). In contrast, *Bacillus* sp. and *Klebsiella pneumonia* were reported to exhibit optimum pullulanase production at 50°C temperature by Waleed et al²⁵ and Abdul-Hadi et al¹. Further increase in temperature suppressed the growth of bacteria and consequently enzyme production also gets inhibited.

Effect of pH: NP 7 bacterial culture shows maximum pullulanase enzyme production at 7 pH (Graph 4). Different organisms have different pH optima and decrease and increase in pH result in poor microbial growth. Enzyme production stimulates at neutral pH and also increases the cell growth. Neutral pH was also found to be optimal for pullulanase production in *Bacillus safensis*²¹.

Effect of agitation rate: Present study shows that 120 RPM was favorable for maximum yield of pullulanase. On further increase of RPM, production of pullulanase was decreased (Graph 5). For proper balance of mixing of nutrients and their physical availability to the organism, agitation is necessary. The uptake of nutrients from the outer environment may be hindered, as too little time is available for contact with substrates at higher agitation rate. This may lead to reduced uptake of substrates and other nutrients resulting in poor growth and production of enzymes. For the production of amylopullulanase from *Geobacillus thermoleovorans*, agitation rates in the range of 100 to 250 rpm were checked for maximum production^{3,20}.

Optimization of chemical parameters

Effect of crude carbon substrate: In the present study, different crude substrate such as corn flour, sago starch, rice bran, red gram and wheat bran were studied with 0.5% liquid production medium. The result reveals that the maximum productivity (123.00 U/ml) was obtained in control production (Graph 6) after 48 hours.

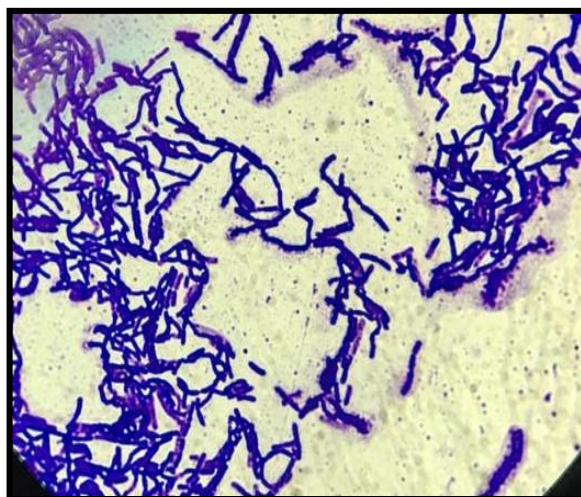
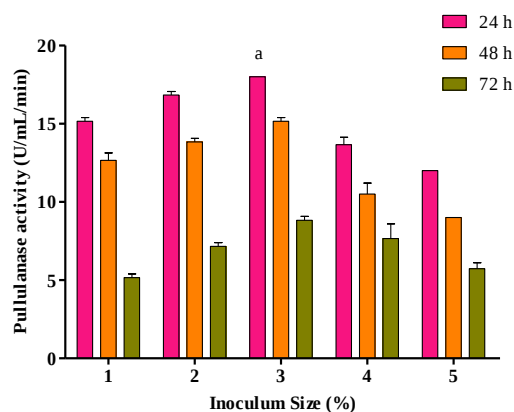
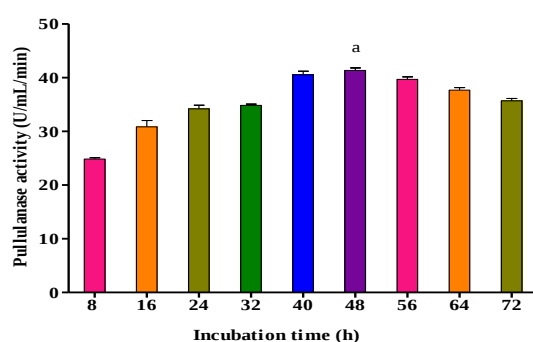


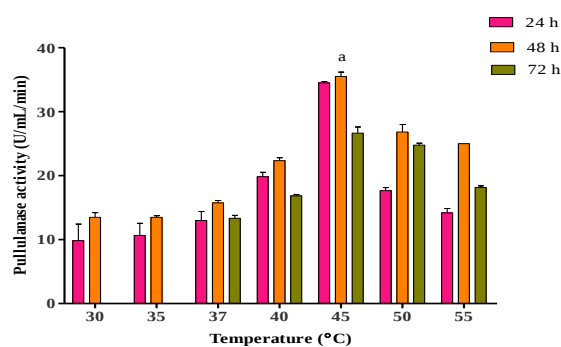
Figure 1: Microscopic observation of NP 7 strain



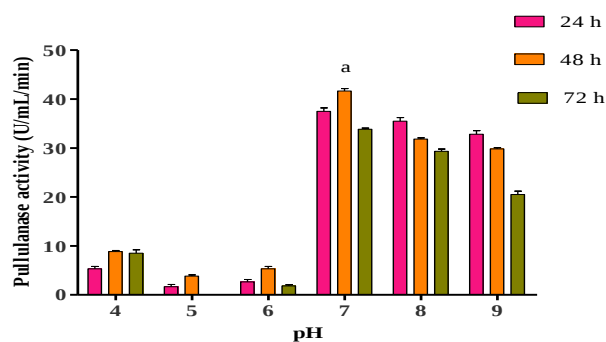
**Graph 1: Effect of Inoculum Size: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs**



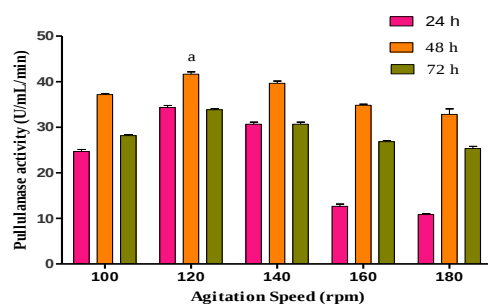
**Graph 2: Effect of Incubation time: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs.**



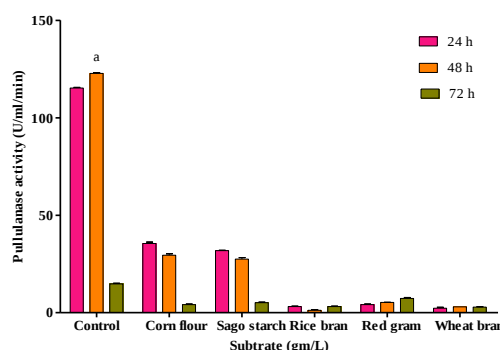
**Graph 3: Effect of Temperature: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs**



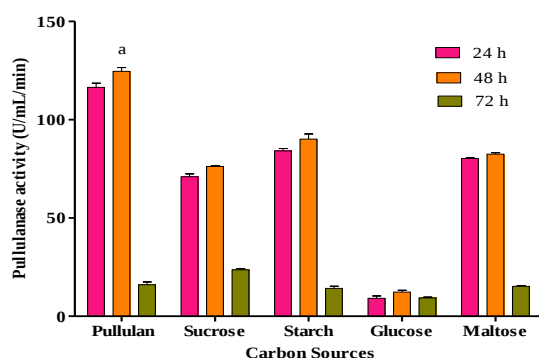
**Graph 4: Effect of pH: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs**



Graph 5: Effect of agitation rate: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs.



Graph 6: Effect of crude substrate: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs



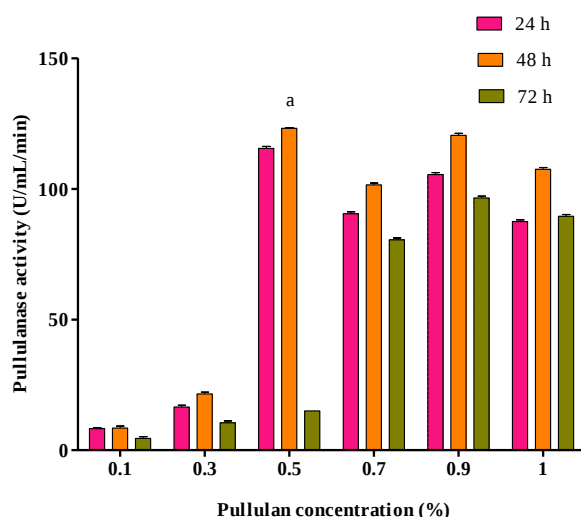
Graph 7: Effect of different carbon sources: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests.
^a highly significant at a time interval of 24 hrs

Effect of different Carbon source: In present study with NP 7, among different carbon sources, 0.5% starch gave 127.50 U/ml enzyme activities and pullulan gave 125.00 U/ml activities. Other carbon sources do not show any significant increase in enzyme activity (Graph 7). Maximum production of pullulanase by *Bacillus cereus* FDTA 13 was observed with soluble starch which increased the yield 1.8-fold as compared to control. For production of pullulanase, soluble starch is a good source of carbon because it can induce its release in case of cell associated pullulanase¹⁹.

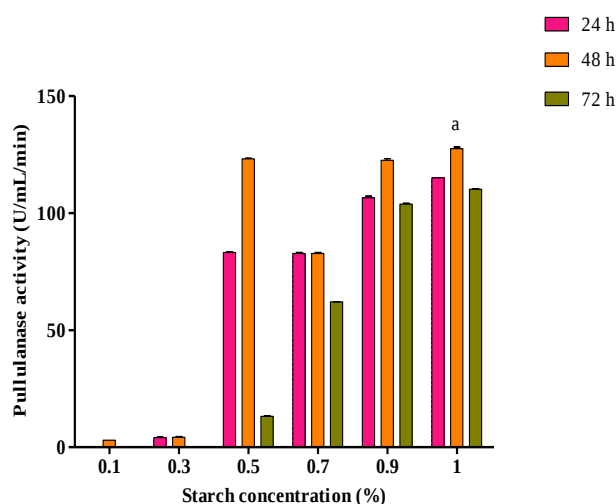
Effect of different concentrations of pullulan and starch in production medium: Different concentrations (0.1 to 1.0%) of pullulan and starch were studied and it was found that 0.5% pullulan gave maximum productivity of 123.00 U/ml (Graph 8) and 1% starch gave maximum productivity of 127.50 U/ml (Graph 9) which is similar. So, 1% starch

can be used further for pullulanase production. Among the carbon sources, pullulan and starch were found to support pullulanase production. *B. cereus* showed high enzyme yield (1.16 U/ml) when pullulan was used as carbon source. Similar observation was noticed for an enhancement of pullulanase synthesis by pullulan in *B. cereus* FDTA-13¹⁹ and in *B. halodurans*^{4,25}.

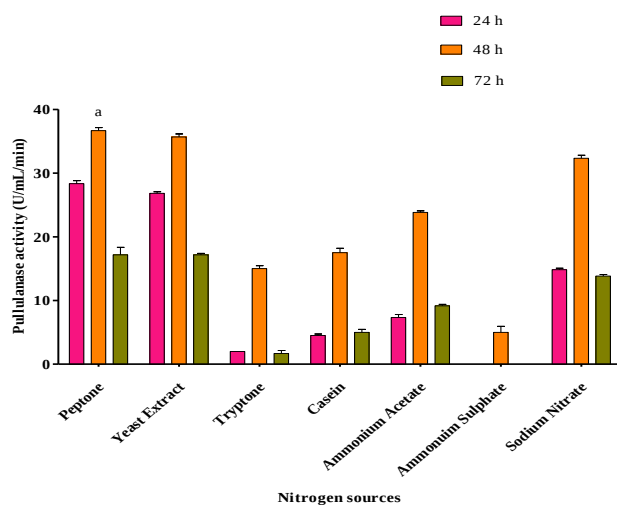
Effect of different nitrogen source: In present study with NP 7, it was observed that peptone showed highest productivity of pullulanase enzyme compared to other nitrogen sources (graph 10). Peptone can be easily degraded and absorbed because of low molecular weight¹⁵. Among the different nitrogen sources tested, peptone was more preferred for extracellular pullulanase production. Similar observation was noticed for *Clostridium thermosulfurogenes* SV9⁴.



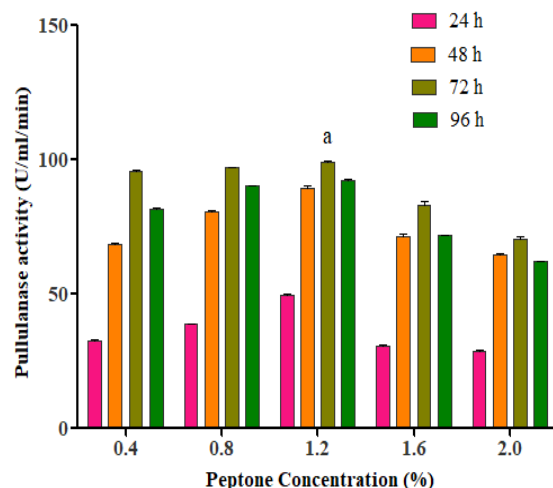
Graph 8: Effect of different Pullulan concentration: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests. ^a highly significant at a time interval of 24 hrs



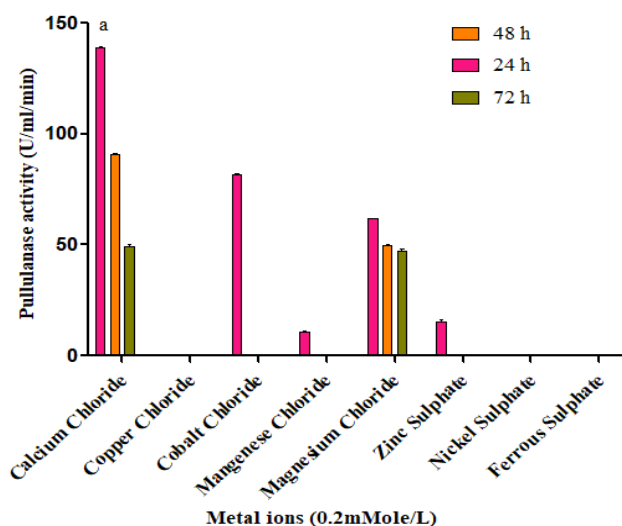
Graph 9: Effect of different Starch concentration: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests. ^a highly significant at a time interval of 24 hrs



Graph 10: Effect of different nitrogen sources: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests. ^a highly significant at a time interval of 24 hrs



Graph 11: Effect of different Peptone Concentration: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests. ^a highly significant at a time interval of 24 hrs.



Graph 12: Effect of different mineral salts: $p < 0.0001$, p value summary based on Tukey's multiple comparison tests. ^a highly significant at a time interval of 24 hrs

Effect of different concentrations of nitrogen source in production medium: Different concentrations of peptone ranging from 0.4 to 2.0% (interval of 0.4%) were studied and it was found that 1.2% peptone gave maximum pullulanase activity (Graph 11) which is same as original production medium which was similar to *Bacillus halodurans* and *Bacillus licheniformis* for production of extracellular pullulanase¹⁸.

Effect of different metal ions: Pullulanase is known to be metalloenzyme. Supplementation of salts of certain metal ions provided good growth of bacteria and thereby better enzyme production. Present medium contains metal ions. Mineral salts played an indirect role in production of enzyme. From the result, it was found that 0.2 mM/L concentration of CaCl_2 showed maximum activity of enzyme than the control which contains MgSO_4 (Graph 12) which was similar to thermophilic *Bacillus strain* 3183. For

stabilization and maintenance of pullulanase activity, Ca^{+2} might be required²⁰.

Therefore, CaCl_2 acts as activator for enzyme production and NiSO_4 and FeSO_4 act as inhibitors. FeSO_4 and BaCl_2 have the highest inhibition effect on pullulanase production by *B. cereus* KKSJ 1981¹⁵. Nair et al¹⁹ observed similar results for pullulanase production from *Bacillus* sp. and *B. cereus* respectively.

Conclusion

NP 7 has shown 4 times increase in productivity of pullulanase after optimizing physical and chemical parameters. After optimization of physical parameters, NP 7 gave maximum pullulanase production using 3% inoculum at 45°C temperature and at 7 pH after 48 hours of incubation in submerged condition with 120 RPM. Starch, peptone and CaCl_2 maximize production of enzyme at the concentrations of 1%, 1.2 % and 0.2 mM/L respectively. Optimized

parameter can be used to set up production of pullulanase at pilot or large scale production.

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References

1. Abdul-Hadi S. and Al-Bayyar A., Extraction, Purification and Characterization of Extracellular Pullulanase By *Klebsiella Pneumoniae* Isolated From Soil, *Pakistan Journal of Biotechnology*, **16(1)**, 2312–7791 (2019)
2. Ara K., Saeki K., Igarashi K., Takaiwa M., Uemura T., Hagihara H., Kawai S. and Ito S., Purification and Characterization of an Alkaline Amylopullulanase with Both α -1,4 and α -1,6 Hydrolytic Activity from Alkalophilic *Bacillus* Sp. KSM-1378, *Biochimica et Biophysica Acta (BBA) -General Subjects*, **1243(3)**, 315–24 (1995)
3. Arabacı N. and Arıkan B., An amylopullulanase (ApuNP1) from *Geobacillus thermoleovorans* NP1: biochemical characterization and its potential industrial applications, *Preparative Biochemistry & Biotechnology*, **49(2)**, 127–135 (2019)
4. Asha R., Niyonzima F.N. and Sunil S.M., Purification and properties of pullulanase from *Bacillus halodurans*, *Int. Res. J. Biol. Sci.*, **2(3)**, 35–43 (2013)
5. Bi J., Jing X., Wu L., Zhou X., Gu J., Nie Y. and Xu Y., Computational design of noncanonical amino acid-based thioether staples at N/C-terminal domains of multi-modular pullulanase for thermostabilization in enzyme catalysis, *Computational and Structural Biotechnology Journal*, **19**, 577–585 (2021)
6. Borchert M., Gjermansen M., Clark S., Henrissat B., Silow M.B. and Hallin F., Pullulanase variants and uses, United states patent, **54**, 1-67 (2014)
7. Catley B.J., Pullulan, a relationship between molecular weight and fine structure, *FEBS Letters*, **10(3)**, 190–193 (1970)
8. Davaeifar S., Shariati P., Tabandeh F. and Yakhchali B., Isolation and Identification of a New *Bacillus cereus* Strain and Characterization of its Neopullulanase, *Applied Food Biotechnology*, **2(2)**, 39–45 (2015)
9. Domań-Pytka M. and Bardowski J., Pullulan Degrading Enzymes of Bacterial Origin, *Critical Reviews in Microbiology*, **30(2)**, 107–121 (2004)
10. Duffner F., Bertoldo C., Andersen J.T., Wagner K. and Antranikian G., A New Thermoactive Pullulanase from *Desulfurococcus mucosus*: Cloning, Sequencing, Purification and Characterization of the Recombinant Enzyme after Expression in *Bacillus subtilis*, *J Bacteriol.*, **182(22)**, 6331–6338 (2000)
11. Gao H. and Gu W.Y., Optimization of polysaccharide and ergosterol production from *Agaricus brasiliensis* by fermentation process, *Biochemical Engineering Journal*, **33(3)**, 202–210 (2007)
12. Goh K.M., Mahadi N.M., Hassan O., Rahman R.N.Z.R.A. and Ilias R.M., A predominant β -CGTase G1 engineered to elucidate the relationship between protein structure and product specificity, *Journal of Molecular Catalysis B: Enzymatic*, **57**, 270–277 (2009)
13. Hii S.L., Ling T.C., Mohamad R. and Ariff A.B., Enhancement of Extracellular Pullulanase Production by *Raoultellaplasticola* DSMZ 4617 Using Optimized Medium Based on Sago Starch, *TOBIOTJ*, **3**, 1–8 (2009)
14. Hope G.C. and Dean A.C.R., Pullulanase Synthesis in *Klebsiella (Aerobacter) aerogenes* Strains Growing in Continuous Culture, *Biochem. J.*, **144**, 403–411 (1974)
15. Jyothi J., Abbulu K., Nanda G. and Kadimpati K.K., Enrichment of Extracellular Pullulanase Production by Isolated *Bacillus Cereus* Kksj 1981 through Optimization of Fermentation Conditions, *JGTPS*, **11(2)**, 7541–7548 (2020)
16. Malakar R., Yadav M., Malviya S.N. and Tiwari A., Effect of Media Composition and Culture Condition on Pullulanase Enzyme from Extremophile Bacterial Species, *International Journal of Biomedical and Advance Research*, **3**, 692–699 (2012)
17. Miller G.L., Use of Dinitrosalicylic Acid Reagent for Determination of Reducing Sugar, *Analytical Chemistry*, **3**, 426–428 (1958)
18. Muslim S., Mohammed A.A. and Fayyad R., A Novel Culture Medium Using Lettuce Plant (*Lactuca Sativa*) For Detection of Pullulanase Production by *Paenibacillus Macerans* Isolated from Agricultural Wastes, *Journal of Physics: Conference Series*, **1818**, 012034 (2021)
19. Nair S.U., Singhal R.S. and Kamat M.Y., Enhanced Production of Thermostable Pullulanase Type 1 Using *Bacillus cereus* FDTA 13 and Its Mutant, *Food Technol. Biotechnol.*, **44(2)**, 275–282 (2006)
20. Noorwez S.M., Ezhilvannan M. and Satyanarayana T., Production of a high maltose-forming, hyperthermostable and Ca^{2+} - independent amylopullulanase by an extreme thermophile *Geobacillus thermoleovorans* in submerged fermentation, *Indian J Biotechnol.*, **5**, 337–345 (2006)
21. Oladipo O., Afolayan D., Olusola L. and Festus I., Properties of a neutral, thermally stable and surfactant-tolerant pullulanase from worker termite gut-dwelling *Bacillus safensis* as potential for industrial applications, *Heliyon*, **8**, e10617 (2022)
22. Simair A.A., Qureshi A.S., Khushk I., Ali C.H., Lashari S., Bhutto M.A., Mangrio G.S. and Lu C., Production and Partial Characterization of α -Amylase Enzyme from *Bacillus* sp. BCC 01-50 and Potential Applications, *Bio Med Res. Int.*, **2017**, 1–9 (2017)
23. Singh R.S., Saini G.K. and Kennedy J.F., Maltotriose syrup preparation from pullulan using pullulanase, *Carbohydrate Polymers*, **80**, 401–407 (2010)
24. Swamy M.V. and Seenayya G., Thermostable Pullulanase and α -Amylase Activity from *Clostridium Thermosulfurogenes*SV9 Optmization of Culture Conditions for Enzyme Production, *Process Biochemistry*, **31**, 157–162 (1996)
25. Waleed M., Faiza A.A. and Nizar I., Production optimization of pullulanase enzyme produced by *Bacillus cereus* isolated from Syrian sources, *Int. Food Res. J.*, **22(5)**, 1824–1830 (2015)

26. Wei W., Ma J., Guo S. and Wei D., A type I pullulanase of *Bacillus cereus* Nws-bc5 screening from stinky tofu brine: Functional expression in *Escherichia coli* and *Bacillus subtilis* and enzyme characterization, *Process Biochemistry*, **49**, 1893–1902 (2014)

27. Zeikus J.G., Lowe S.E., Podkovyrov S.M. and Ramesh M.V., Cloning and sequencing of the *Thermoanaerobacterium*

saccharolyticum B6A-RI *apu* gene and purification and characterization of the amylopullulanase from *Escherichia coli*, *Applied and Environmental Microbiology*, **60**(1), 94-101 (1994).

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