STRAIN IMPROVEMENT OF *ASPERGILLUS ORYZAE* FOR ENHANCED BIOSYNTHESIS OF PHYTASE THROUGH CHEMICAL MUTAGENESIS

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Abstract

The present study deal with strain improvement through random chemical mutagenesis of phytase producing *Aspergillus oryzae* UJIIB-29. *Aspergillus oryzae* wild strain UJIIB-29 was exposed to three different chemical mutagens i.e. sodium azide, nitrous acid and ethyl methane sulfonate (EMS). Thirty isolates obtained after treating with chemical mutagens showing the large zone of phytic acid hydrolysis were picked and subjected to secondary screening for phytase production using submerged fermentation on phytase screening media. The mutant NA-4 obtained after treating wild strain with nitrous acid exhibited maximum activity of phytase (44 ± 0.01 U/ml/min). It was approximately 1.9 folds higher than enzyme activity (23.1 ± 0.01 U/ml/min) of parent strain. To further increase the phytase activity, fermentation conditions like fermentation media, time of incubation, incubation temperature, pH of the medium, carbon source and its different concentrations and size of inoculum were optimized. The maximum phytase units (58.98 ± 0.01 U/ml/min) were achieved in glucose phosphate broth (M) having 1% starch, 0.4% ammonium sulphate as a carbon and nitrogen sources, respectively, at 30° C, having pH 4.5 after 7days of incubation period. As a result of optimization, 1.3 times increased in phytase activity (58.98 ± 0.01 U/ml/min) was achieved. Nitrous acid proved to be a potent mutagen for strain improvement. Furthermore, optimized fermentation parameter can help in scaling up the phytsae production using *Aspergillus oryzae* NA-4.

Key words: Mutagenesis; Catalysis; Submerged fermentation, Enxyme, Optimization.

Introduction

Phytase (EC 3.1.3.8) also known as phosphomonoesterase that hydrolyze the phytic acid in a sequential manner. Phytase releases the myoinositol, inositol phosphate and soluble form of inorganic phosphate (P) (Zeller et al., 2015; Li et al., 2021). Based on position of phosphate group (PO₄³⁻) that hydrolyzed first, phytase enzymes are classified into two groups i.e. 3-phytase (E.C. 3.1.3.8) and 6-phytase (E.C. 3.1.3.26). When first phosphate molecule is released from phytic acid, others are also released by step wise action of phytases or acid phosphatases (Haefner et al., 2005; Da Silva et al., 2019).

Phytic acid, also called Ins P_6 (myo inositol 1, 2, 3, 4, 5, 6 hexakisphoaphate) is major source of organic phosphate in nutrition of animal and storage of phosphate in plants. In cereals, legumens and oilseed crops phytic acid commit approximately 60-90% of total phosphate content (Raboy, 2020). The complex of magnesium, potassium, magnesium and calcium with phytic acid is called phytate that is present in the plants.

Phytase are produced from various animals, microorganisms and plants sources. Because of substrate specificity, high thermostability, catalytic efficiency and low production cost phytases obtain by microbial sources have been targeted in research work (Sabu *et al.*, 2002; Dailin *et al.*, 2019). Several different filamentous fungal species such as *Aspergillus fumigatus, Aspergillus oryzae, Aspergillus carbonarius* and *Aspegillus niger* are reported for phytase production by fermentation (Casey & Walsh, 2004; Salmon *et al.*, 2012). The fungus belongs to *Aspergillus* (genus) yield most suitable and active extracellular enzyme. Therefore, at industrial level for the

production of phytase the *Aspergillus* spp. has been preferred (Jatuwong *et al.*, 2020 Tanruean *et al.*, 2021).

Microorganisms do not produce the enzyme in a sufficient quantity to cater the needs of industries. Hence, by changing the microorganism's genome, better rate of enzymes production and their characteristics are achieved. To change the genetic material different techniques include recombinant protoplast fusion, recombinant DNA technology and mutation are used (Ushasree et al., 2017). At industrial scale, random mutagenesis for stain improvement plays a very significant role in developing the economical process of microbial fermentation. Two various kinds of mutagenesis such as site direct mutagenesis and random mutagenesis are usually preferred. In the random mutagenesis, physical mutagens and chemical mutagens are favored. Physical mutagens include gamma irradiations, X-rays and ultraviolet irradiations. But, Chemical mutagens includes ethidium bromide, N-methyl N-nitrosoguanidine (MNNG), ethyl methanesulfonate (EMS), nitrous acid, sodium azide and nitrosotrimethyl guanidine (NTG) (Iftikhar et al., 2010; Goraya et al., 2017; Hassaan et al., 2021).

Phytase plays a very significant part in industries belong to animals feed as phytase increases the absorption and digestion of phosphorous contents and several other organic or inorganic nutrients that are insufficiently accessible in diet supplement of several monogastric animals (Vasudevan *et al.*, 2017). Phytase has very important role in the biomedical and biotechnological applications because of its anti-inflammatory, antioxidant and anti-cancerous properties. Therefore, in current study random chemical mutagenesis of *A. oryzae* UJIIB-29 was carried out to enhance phytase production.

Material and Methods

Fungal strains: The fungal strain of *A. oryzae* UJIIB-29 that produce the phytase was obtained from IIB (Institute of Industrial Biotechnology), Government College University Lahore. Fungal culture was kept on culture slant of Potato dextrose agar (PDA) and stored in refrigerator at 4°C for further use.

Mutagenesis

Spore suspension: Spore suspension was prepared from 3 days old culture from potato dextrose agar (PDA) slant in 5ml sterile dH₂O. The volume of the suspension was raised up to 50ml with 0.01% solution of tween-20. Glass beads was added to break the hyphal mycelia in the spore suspension and placed in shaking incubator for 30min at 30°C. The spore suspension was filtered to remove the fungal mycelia. By serial dilution spore suspension have 10^6 spores/ml was prepared. Homocytometer was used to count the spores under light microscope (Haq *et al.*, 2014).

Sodium azide treatment: Took 9ml spore suspension having 10^6 spore/ml in different test tubes and different concentrations (0.25-2.5%) of sodium azide were mixed up. The reaction was stopped by adding 1ml glycerol (40%) after 30 min. The samples were added in different eppendorfs (Eco Biosciences). The tubes were centrifuged at 4°C and 6000rpm for 15min. Decanted the supernatant and pellet was washed with autoclaved dH₂O. Repeat the same procedure for 2-3 times and pellet was re-dissolved in phosphate buffer (Jayaraman & Ilyas, 2010).

Ethyl methane treatment: The spore suspension (900ml) containing 10^6 spores/ml was transferred in various eppendorfs (Eco Biosciences). Different concentrations (10ul- 40ul) of pure Ethyl methane sulfonate were added in eppendorfs (Eco Biosciences). Incubated the reaction mixture at room temperature for 30min. After incubation period, 40% glycerol (500ul) was mixed in the reaction mixture to stop reaction. The reaction mixture was centrifuged at 4°C and 6000 rpm for 15 min. Decanted the supernatant and pellet was washed with sterile phosphate buffer. Repeat the same procedure for 2-3 times (Haq *et al.*, 2014).

Nitrous acid treatment: Spore suspension (1ml) having 10^6 spores/ml was added in different test tubes. 1 ml of nitric acid (0.1N) and 1ml of different concentrations (24mM-40mM) of sodium nitrite were mixed up in them. Incubated the reaction mixture in the shaking incubator for 30 min at 30°C and 120 rpm. To stop the reaction 40% glycerol (1ml) was added in the reaction mixture. Reaction mixture was centrifuged at 4°C and 6000 rpm for 10 min. The pallet was washed with sterile d.H₂O after supernatant was removed. The same method was repeated thrice (Haq *et al.*, 2014).

Screening of mutant strains: After treating with chemical mutagens, the suspension of spores was poured and spread aseptically on phytase screening media (PSM) in petri plates. Incubated the plates at 30°C in static incubator for 96 h. Fungal growth pattern was analyzed frequently every 3-4 days. The mutant colonies were selected qualitatively, based upon the diameter of phytic acid hydrolytic zone. The mutant colonies were examined quantitatively by further subjecting them to submerged fermentation for phytase production.

Enzyme Production

Preparation of inoculum: The conidial suspension was prepared from 72h old fungal culture in sterile dH_2O . Autoclaved dH_2O (10ml) was added in each slants having profuse conidial growth on its surface. The fungal conidia were scratched aseptically with the aid of inoculating loop. The homogenous suspension of spores was formed by vigorously mixing the solution. The numbers of spores were counted with the help of homocytometer and adjusted as 2.6×10^6 spores/ml.

Submerged fermentation: Phytase production was carried out in the phytase screening media containing glucose (3g), Ammonium sulphate (1.0g), KCL (0.1g), MgSO4. 7H₂O (0.02g), NaCl (0.04g), CaCl₂ (0.002g), FeSO₄.7H₂O (0.002g) MnSO₄ (0.002g) and sodium phytate (0.25g) in 200ml d.H₂O. After sterilization, spore suspension (1ml) was added in the fermentation medium. The fermentation medium was incubated at 30°C and 150 rpm in shaking incubator.

Extraction of enzyme: After 7 days (168h) of incubation the cultural broth was filtered with filter paper. After filtration, the filtrate was centrifuged for 10-15 min at 6000rpm. Supernatant was used for estimation of enzyme activity and pellet was discarded.

Enzyme assay: Enzyme assay was performed at 50°C in four test tubes named as blank, control, control enzyme and experimental. The reaction mixture (Experimental) had 0.5ml of 0.1M MgSO₄ solution, 0.5 ml crude enzyme and 1 ml substrate solution. In control 1ml substrate solution and 0.1M MgSO₄ (0.5ml) was added. Control enzyme contained only 0.5ml crude enzyme. Blank contained only 2ml of d.H₂O. Test tubes were incubated at 50°C in water bath for 15min. After incubation time, 1ml Trichloro acetic acid (TCA) solution (10%) was added in each test tubes expect control enzyme. Tausky Shorr Color Reagent (1ml) was added in all test tubes expect blank and mixed the control enzyme with control. The absorbance was taken by using the spectrophotometer at 660nm.

One unit phytase activity means, the enzyme quantity requires to liberate one micromole of inorganic phosphate (1umol) in one minute in standard enzyme assay conditions (Racmawati *et al.*, 2017).

Units/ml Enzyme = $\frac{(micromoles of phosphate release) (df)(5)}{(T) (0.5)(2)}$

where:df is a dilution factor.5 is total volume of reaction mixture.2 volume taken from the reaction mixture.0.5 is the volume of enzyme.All the values are measured in ml.

Optimization of fermentation parameters: The cultural conditions i.e. fermentation medium, time of incubation, incubation temperature, medium pH, carbon source and its concentration, nitrogen source and its concentration and size of inoculum for phytase production from wild strain (UJIIB-29) and mutant strain (NA-4) of *Aspergillus oryzae* were optimized.

Statistical analysis

Three replicates of operational variables were statistically analyzed by using one way ANOVA.

Results and Discussion

In order to increase the phytase production from A. oryzae UJIIB-29, sixty-two mutant isolates were obtained after exposing fungal spores suspension with various concentrations of nitrous acid, sodium azide and Ethyl methane sulfonate (EMS) for 30min. After qualitative (primary) screening, out of these sixty-two mutant strains, only thirty were subjected to quantitative (secondary) screening because in comparison to wild strain of Aspergillus oryzae UJIIB-29 only these 30 mutant strains exhibited the larger zones of phytic acid hydrolysis. Submerged fermentation was used for secondary screening of these 30 mutant strains on PSM (Phytase screening media). Phytase activity by these mutant strains is illustrated in (Fig. 1) (mutagens obtained after sodium azide treatment), (Fig. 2) (mutagens obtained after nitrous acid treatment) and (Fig. 3) (mutagens obtained after EMS treatment). Mutant strain NA-4 obtained after treating with nitrous acid gave maximum phytase activity (44±0.01 U/ml/min), among these 30 mutant strains. 1.9 times increased in phytase activity was achieved in comparison to enzyme activity (23.1±0.01 U/ml/min) by wild strain. Nitrous acid might change the structure of deoxyribonucleic acid (DNA) by substituting the amino group (NH₂) with hydroxyl (OH) group at C-6 of the adenine, guanine and cytosine. As result adenine (A) changes into hypoxanthine, guanine (G) changes into xanthine (X) and cytosine (C) changes into uracil (U).

Different media like glucose phosphate broth modified (GPBM), glucose phosphate broth (GPB), glucose peptone media (GPM), peptone potato dextrose broth (PPDB), corn starch media (CSM), Vogel's media (VM), malt extract broth (MEB), malt yeast broth (MYB), czapek dox media (CDM) and phytase screening media (PSM) were used as a fermentation medium for maximum phytase production from both wild strain (UJIIB-29) and mutant strain (NA-4) as shown in (Fig. 4). For both mutant strain (NA-4) and wild strain (UJIIB-29) of *Aspergillus oryzae*, the maximum activity of phytase (57.97 \pm 0.01 U/ml/min and 22 ± 0.01 U/ml/min, prospectively) was obtained on glucose phosphate broth modified. Haq *et al.*, (2020) also used the same composition of glucose phosphate broth for biosynthesis of phytase by *Aspergillus oryzae* and maximum phytase activity (22.3 U/ml/min) was reported at (GPBM). Magnesium sulphate (MgSO₄. 7H₂O) in the cultural medium acts as a co-factor for enzyme and reduce the permeability of the plasma membrane. Potassium chloride (KCL) in the medium enhances the conidial head formation. Calcium chloride (CaCl₂) helps to enhance the branches of hyphae and enhance the uptake of phosphate in the medium. Potassium dihydrogen phosphate maintains the pH for production of phytase by using the K⁺/H⁺ ions channels.

Effect of various incubation period (24h-216h) was examined on yield of phytase by mutant strain (NA-4) and wild strain (UJIIB-29) of Aspergillus oryzae as shown in (Fig. 5). The maximum enzyme activity (58.1 ±0.01 U/ml/min, 21.89±0.01 U/ml/min) by mutant strain (NA-4) and wild strain (UJIIB-29), respectively was observed after 168h. Further increase in incubation period after 192h, the phytase activity (16.00 ± 0.01 U/ml/min, 54.68 ±0.01 U/ml/min) was decreased from wild strain (UJIIB-29) and mutant strain (NA-4), respectively. Bhavsar et al., (2008) and Haq et al., (2020) also confirmed our wild strain's (UJIIB-29) results that maximum activity of phytase enzyme (22.3 U/ml/min and 68 IU/ml) was obtained after 168h incubation period. In case of mutant strains, Shah et al., (2009), Sing (2013) and Hassaan et al., (2021) reported the maximum phytase activity (80 IU/ml and 2881.60 U/ml) after 14 days, 4 days and 12 days of incubation period, respectively. The differences in results might be because the use of different species of Aspergillus and fermentation media for phytase production. The cells of the fungus enter the log phase at very slow rate due to slow consumption of media components.

Different incubation temperatures i.e. 20°C, 30°C, 40°C and 50°C were examined to observe their effect on phytase activity from both wild stain (UJIIB-29) and mutant strain (NA-4) as describe in (Fig. 6). The maximum enzyme activity (22.57 \pm 0.01U/ml/min, 58.4 \pm 0.01 U/ml/min) was achieved by wild strain (UJIIB-29) and mutant strain (NA-4), respectively at 30°C. But in case of mutant a slight decline in activity of phytase enzyme (57.82± 0.01 U/ml/min) was recorded when incubation temperature was 40°C. In case of wild strain (UJIIB-29), Singh (2014), Sandhya et al., (2015) and Haq et al., (2020) published the similar results. But in mutant strain (NA-4), Sapna and sing (2013) and Mahmood et al., (2021) reported that maximum phytase production was achieved at 35°C. The contradictions in the result might be due to the effect of random mutagenesis on fungal strain of Aspergillus oryzae. It exhibited that n Aspergilus oryzae could be a mesophilic fungus. Incubation temperature effect the microorganism growth as well as the product formation. At very high incubation temperature, metabolic functions of fungus disturbed due to denaturation of proteins and fungal cells starting die.



Fig. 1. Production of phytase by Sodium Azide treated mutant isolates. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 3. Production of phytase by EMS treated mutant isolates. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 5. Effect of incubation time on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 2. Production of phytase by Nitrous Acid treated mutant isolates. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 4. Effect of different cultural media on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 6. Effect of incubation temperature on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 7. Effect of pH of fermentation medium on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 9. Effect of different concentrations carbon source on activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 11. Effect of different concentrations of nitrogen source on phytase *activity*. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 8. Effect of different carbon sources on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 10. Effect of different nitrogen sources on phytase activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).



Fig. 12. Effect of inoculum size on phytase production activity. Among three replicates Y-bar showed the standard deviation (S.D. $\leq \pm 0.05$).

Different pH values of medium (4-8) were examined to check their effect on the phytase production by both wild strain (UJIIB-29) and mutant strain (NA-4) as illustrated in (Fig. 7). At pH 4.5, the maximum activity of phytase (58.7 ± 0.01 U/ml/min) was achieved from mutant strain (NA-4). Shivanna and Venkateswaran (2014) also recorded that maximum enzyme activity was achieved at pH 4.5 in submerged fermentation by Aspergillus niger. At 5.5 pH of fermentation medium, maximum activity of phytase enzyme (23.57 \pm 0.01 U/ml/min) was achieved by wild strain (UJIIB-29) of Aspergillus oryzae. Haq et al., (2020) and Odeniyi and Itaba (2020) observed that enzyme activity (22.3 U/ml/min and 1 U/ml) was maximum by Aspergillus oryzae and Aspergillus niger at medium pH 5.5. The presence of high concentration H⁺ ions effect the permeability of plasma membrane as well as nutrients mobility. The decrease or increase in pH change the amino acids charges and at enzyme active site that decrease the enzyme activity.

Various sources of carbon like glucose, starch, glycerol, xylose, mannitol, fructose, galactose, lactose, sucrose and starch were used to evaluate the production of phytase by both wild strain (UJIIB-29) and mutant strain (NA-4) in fermentation media as exhibited in (Fig. 8). Maximum enzyme activity $(23.7 \pm 0.01 \text{ U/ml/min})$ from wild strain was attained when glucose was used in the fermentation media. Soni and Khire (2007), Bhavsar et al., (2008) Qasim et al., (2017), Haq et al., (2020) and Mahmood et al., (2021) also recorded the similar results that activity of phytase was highest by A. niger, A. tubingensis and A. oryzae when glucose was used in the fermentation medium. The maximum phytase units (58.91 \pm 0.01 U/ml/min) in case of mutant strain (NA-4) were achieved, when starch (as a carbon source) used in the fermentation medium. The mutant strain (NA-4) showed a slight decrease in phytase activity (58.51±0.01 U/ml/min), when glucose used as a carbon source in fermentation media. Sapna & Singh (2013), Sing (2017) also observed the similar results that production of phytase was maximum from Aspergillus ssp. when starch used as source of in the fermentation media.

Different carbon source concentrations (0.5%-2%) were utilized in cultural medium to examine their effect on the phytase activity by mutant strain (NA-4) and wild strain (UJIIB-29) of Aspergillus oryzae as described in (Fig. 9). Phytase enzyme activity $(23.71 \pm 0.01 \text{ U/ml/min},$ 58.91 ± 0.01 U/ml/min) in case of wild strain (UJIIB-29) and mutant strain (NA-4) was maximum when carbon source (1%) was used in the cultural media. Shah et al., (2009) and Qasim et al., (2017) observed that phytase yield was maximum by suspending glucose as carbon source at concentration 3.5% and 1.5% in cultural media from A. niger and A. tubingensis respectively. The difference in the results might be because of different species of Aspergillus have different metabolical rate. But in case of mutant, Sing (2017) reports the similar results that activity of phytase was highest by Aspergillus oryzae when 1% starch was used in the fermentation media.

Different organic and inorganic sources of nitrogen i.e., peptone, sodium nitrite, ammonium chloride, ammonium nitrate, ammonium sulphate, potassium nitrite, urea, beef extract, yeast extract and tryptone were used to evaluate phytase production as shown in (Fig. 10). Both wild strain (UJIIB-29) and mutant strain (NA-4) showed maximum activity of enzyme (23.8 ±0.011 U/ml/min and 58.96± 0.023 U/ml/min, respectively), when ammonium sulphate was used in the cultural media. In case of wild strain (UJIIB-29), Qasim et al., (2017), Olubusola & Itaba (2020) and Kumari & Bansal (2021) supported our results that phytase production was maximum when ammonium sulphate was used as a nitrogen source. In case of mutant strain, Ajith et al., (2019) reported that phytase activity was maximum by using the sodium nitrate (NaNO₃) in fermentation media. The contradiction might be due to use of different Aspergillus spp. like Aspergillus foetidus and Aspergillus orvzae. Nitrogen is used for amino acid and nucleotides synthesis for microbes. These molecules are the basic units of large polymers such as proteins, deoxyribonucleic acid (DNA) and glucosamine.

Different nitrogen source (ammonium sulphate) concentrations (0.2%-0.8%) were used in fermentation medium to analyze their effect on phytase production by mutant strain (NA-4) and wild strain (UJIIB-29) of Aspergillus oryzae (Fig. 11). The mutant strain (NA-4) exhibited the maximum units of enzyme (58.96 \pm 0.01 U/ml/min), when 0.4% ammonium sulphate ((NH₄)₂ SO₄) concentration was used in fermentation medium. But wild strain (UJIIB-29) gave maximum units of enzyme (23.8± 0.01 U/ml/min), when 0.3% ammonium sulphate $((NH_4)_2 SO_4)$ concentration was used in the fermentation medium. Haq et al., (2020) also observed the similar results that activity of phytase enzyme was utilizing 0.3% maximum by nitrogen source concentration. But in mutant the maximum enzyme activity (58.96 \pm 0.01 U/ml/min) was attained when 0.4% concentration of ammonium sulphate was used in fermentation medium. Shah et al., (2009) recorded that maximum phytase units were attained by using 1% concentration of nitrogen source in the fermentation media. It might be because to use different Aspergillus species like Aspergillus niger and Aspergillus orvzae.

Size of inoculum is another significant factor that plays a significant role for enzyme production. The inoculum size was varied (1-8%) to evaluate its effect on phytase production by wild strain (UJIIB-29) and mutant strain (NA-4) as exhibited in (Fig. 12). In both wild strain (UJIIB-29) and mutant strain (NA-4), 4% inoculum size showed maximum yield of enzyme (24.6± 0.01U/ml/min and 58.98 \pm 0.01 U/ml/min, respectively). In wild strain (UJIIB-29), Vassilev et al., (2007) and Haq et al., (2020) also recorded the same results of phytase production from the Aspergillus niger and Aspergillus oryzae. But in mutant strain, shah et al., (2009) use the 1% inoculum size to achieve maximum production of phytase after mutagenesis of A oryzae. The contradiction in results might be because of effect of chemical mutagens during the random mutagenesis and use of different fermentation media.

Conclusion

Aspergillus oryzae mutant strain NA-4 was acquired as a promising microorganism for phytase biosynthesis that produced 1.9 times higher activity of phytase as compared to wild strain of Aspergillus oryzae. Optimization of various fermentation conditions highly influenced the phytase production. 1.3 times enhance in production of phytase was attained after optimization of various fermentation conditions. The recent study will aid to make an economical and effective biochemical process for biosynthesis of phytase at large scale. The process can be utilized in the poultry feed industry and have key significance for human ecologists and nutritionists.

References

- Ajith, S., D. Ghosh, S. Shet, B. ShreeVidhya, D. Punith and A.V. Elanovan. 2019. Partial purification and characterization of synthetic bacteria phytase in *Piichiapastoris* using surface response methodology. *Judishapur J. Microbiol.*, 8(12):
- Bhavsar, K., P. Gujar, P. Shah, V.R. Kumar and J.M. Khire. 2008. Influence of pretreatment of agriculture residues on phytase production by *Aspergillus niger* NCIM 563 under submerged fermentation conditions. *Afri. J. Biotechnol.*, 7(8):
- Casey, A. and G. Walsh. 2004. Identification and characterization of a phytase of potential commercial interest. *J. Biotechnol*, 110: 313-322.
- da Silva, C.A., M.A. Callegari, C.P. Dias, A.M. Bridi, C.R. Pierozan, L. Foppa and R. Hermes. 2019. Increasing doses of phytase from Citrobacter braakii in diets with reduced inorganic phosphorus and calcium improve growth performance and lean meat of growing and finishing pigs. *Plos One*, 14(5): 0217-490.
- Dailin, D.J., S.Z. Hanapi, E. Elsayed, D. Sukmawati, N.I. Wan Azelee and J. Eyahmalay. 2019. Fungal phytases: biotechnological applications in food and feed industries, in recent advancement in white biotechnology through Fungi. *Fung. Biol.*, 1(9): 65-99.
- Goraya, A., F. Asghar, I. Javaid and S. Ali. 2017. Microbial strain improvement for overproduction of industrial products. J. Biol. Sci., 3(5): 10-29.
- Haefner, S., A. Knietsch and E. Scholten. 2005. Biotechnological production and applications of phytases. *Appl. Microbiol. Biotechnol.*, 68: 588-597.
- Haq, I.U., A. Nawaz, H. Mukhtar, Z. Mansoor, M. Riaz, M. Ahmed and S.M. Ameer. 2014. Random mutagenesis of Aspergillus niger and process optimization for enhanced production of glucose oxidase. *Pak. J. Bot.*, 46(3): 1109-1114.
- Haq, I.U., M.U. Javed, A. Nawaz, Y. Arshad and H. Mukhtar. 2020. Biosynthesis, purification, kinetic and thermodynamic analysis of phytase from Aspergillus oryzae. *Rev. Mex. Ing. Quim*, 19(Sup. 1): 227-242.
- Hassan, N., M.M. Khorshed, N.E. El-Bordeny, A.A. Abedo and M.M. Shoukry. 2021. Improvement of fungal phytase production and its in vitro application in ruminant nutrition. *Arab Uni. J. Agri. Sci.*, 29(1): 315-327.
- Iftikhar, T., M. Niaz, Y. Hussain, S.Q. Abbas, I. Ashraf and M.A. Zia. 2010. Improvement of selected strains through gamma irradiation for enhanced lipolytic potential. *Pak. J. Bot.*, 42(4): 2257-2267.
- Jatuwong, K., N. Suwannarach, J. Kumla, W. Penkhrue, P. Kakumyan and S. Lumyong. 2020. Bioprocess for production, characteristics, and biotechnological applications of fungal phytases. *Front. Microbiol.*, 11: 188.

- Jayaraman, R. and M.M. Ilyas. 2010. Strain improvement of Phosphate solubilizing fungal strains. J. Ecobiotech. J. Phytol., 2(5): 65-70.
- Kumari, N. and S. Bansal. 2021. Statistical modeling and optimization of microbial phytase production towards utilization as a feed supplement. *Biomass Convers & Biorefin.* https://doi.org/10.1007/s13399-021-01672-x
- Li, S., B. Wang, Y. Wang, X. Ma, L. Su, Y. Wang and L. Ren. 2021. Soluble organic nutrients induce ClaPhys expression to enhance phytase activity of watermelon roots. *Ann. App. Biol.*, https://doi.org/10.1111/aab.12756
- Mahmood, S., M.G. Shahid and M. Nadeem. 2021. Screening of phytate degrading fungi and optimization of culture conditions for phytase synthesis using agro-industrial byproducts. *Pak. J. Bot.*, 53(2): 763-770.
- Odeniyi, O. and L. Itaba. 2020. Production and characterization of an *Aspergillus niger* GS1S6 phytase cultivation in phytic acid rich agricultural substrates. *Fungal Territ.*, 3(3): 9-15.
- Qasim, S.S., K.A. Shakir, A.B. Al-Shaibani and M.K. Walsh. 2017. Optimization of culture conditions to produce phytase from Aspergillus tubingensis SKA. *Food Sci. Nutr.*, 8(07): 733.
- Raboy, V. 2020. Low phytic acid Crops: observations based on four decades of research. *Plants*, 9(2): 140.
- Rachmawati, D., S. Istiyanto and M. Maizirwan. 2017. Effect of phytase on growth performance, diet utilization efficiency and nutrient digestibility in fingerlings of Chanos chanos (Forsskal 1775). *Philipp J. Sci.*, 146(3): 237-245.
- Sabu, A., S. Sarita, A. Pandey, B. Bogar, G. Szakacs and C.R. Soccol. 2002. Solid-state fermentation for production of phytase by Rhizopus oligosporus. *Appl. Biochem. Biotech.*, 102: 251-260.
- Salmon, D.N.X., L.C. Piva, L.R. Binati, C. Rodrigues, V. de Souza and L.P. Enberghe, C.R. Soccol. 2012. A bioprocess for the production of phytase from Schizophyllum commune: studies of its optimization, profile of fermentation parameters, characterization and stability. *Bioproc. Biosyst. Eng.*, 35: 1067-1079.
- Sandhya, A., A. Sridevi, D. Suvarnalatha and G. Narasimha. 2015. Production and optimization of phytase by *Aspergillus niger. Pharm. Lett.*, 7: 148-153.
- Sapna and B. Singh. 2013. Improved production of proteaseresistance phytase by *Aspergillus oryzae* and its applicability in the hydrolysis of insoluble phytase. *J. Ind. Microbiol. Biotechnol.*, 40(8): 891-899.
- Shah P., K. Bhavsar, S.K. Soni and J.M. Khire. 2009. Strain improvement and up scaling of phytase production by *Aspergillus niger* NCIM 563 under submerged fermentation conditions. J. Ind. Microbiol. & Biotechnol., 36(3): 373.
- Shivanna G.B. and G. Venkateswaran. 2014. Phytase production by Aspergillus niger CFR 335 and Aspergillus ficuum SGA 01 through submerged and solid-state fermentation. The Scientific World J., 2014.
- Singh, B. 2013. Improved production of protease-resistant phytase by Aspergillus oryzae and its applicability in the hydrolysis of insoluble phytates. J. Indus. Microb. & Biotech., 40(8): 891-899.
- Singh, B. 2014. Phytase production by Aspergillus oryzae in solid-state fermentation and its applicability in dephytinization of wheat bran. *Appl. Biochem. & Biotechnol.*, 173(7): 1885-1895.
- Singh, B. 2017. Free and immobilized *Aspergillus oryzae* producing the protease resistant and thermostable phytase. *Biotech.*, 7: 213.
- Soni S.K. and J.M. Khire. 2007. Production and partial characterization of two types of phytase from *Aspergillus niger* NCIM 563 under submerged fermentation conditions. *World J. Microbiol & Biotech.*, 23(11): 1585-1593.

- Tanruean, K., W. Penkhrue, J. Kumla, N. Suwannarach and S. Lumyong. 2021. Valorization of lignocellulosic wastes to produce phytase and cellulolytic enzymes from athermophilic fungus,' thermoascus aurantiacus sl16w, under semi-solid state fermentation. J. Fungi, 7(4): 286.
- Ushasree, M.V., K. Shyam, J. Vidy and A. Pandey. 2017. Microbial phytase: impact of advance in genetic engineering in revolutionizing its properties and applications. *Bioresour. Technol.*, 245: 1790-1799.
- Vasudevan U.M., K. Shyam, J. Vidya and A. Pandey. 2017. Microbial phytase: impact of advances in genetic

engineering in revolutionizing its properties and applications. *Bioresour. Technol.*, 245: 1790-1799.

- Vassilev N., M. Vassileva, V. Bravo, M. Fernández-Serrano and I. Nikolaeva. 2007. Simultaneous phytase production and rock phosphate solubilization by *Aspergillus niger* grown on dry olive wastes. *Ind. Crops. Prod.*, 26(3): 332-336.
- Zeller, E., M. Schollenberger, I. Kühn and M. Rodehutscord. 2015. Hydrolysis of phytate and formation of inositol phosphate isomers without or with supplemented phytases in different segments of the digestive tract of broilers. J. Nutr. Sci., 4(1): 1-12.

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