



Non-recombinant Mutagenesis of *Bacillus subtilis* MTCC 2414 for Hyper Production of Laccase

Venkatanagaraju Erumalla*, Chittaranjan Das, Akihito Choppy

Associate Professor, Therapeutic Enzymes (Division), Department of Life Sciences,
Garden City College of Science and Management Studies, 16th KM, Old Madras Road, Bangalore-560049, Karnataka, India.

ARTICLE HISTORY

Received: 06.11.2016

Accepted: 13.12.2016

Available online: 30.12.2016

Keywords:

Bacillus subtilis MTCC 2414, Laccase, Ultraviolet rays, Ethidium bromide, Ethyl methane sulfonate.

ABSTRACT

The advent of protein engineering and sophisticated molecular technologies has opened possibilities for screening variants of enzymes and tailor-made proteins from microorganisms with enhanced production yields which may be of interest for specific commercial applications. During this investigation the indigenous strain of *Bacillus subtilis* MTCC 2414 procured from Institute of Microbial Technology (IMTECH) Chandigarh, India was improved for enhanced laccase production by using physical mutagen (ultraviolet rays) and chemical mutagens viz., 0.5 mg/ml ethidium bromide and 0.5 mg/ml ethyl methane sulfonate. Mutant GCBR 4 with hyper laccase production (178.8 ± 3.67 U/ml) was obtained after treating wild strain for 80 min with ethidium bromide. The effectiveness of hyper laccase producing mutant GCBR4 indicates its possible applicability in various biotechnological and industrial processes.

*Corresponding author:

Email : venkatanagarajue@gmail.com

Tel.: +91 - 7676680687

INTRODUCTION

Enzymes are delicate protein molecules ubiquitous in occurrence, and are essential for cell growth and differentiation[1, 2]. The extracellular enzymes are of commercial value and find multiple applications in various industrial sectors[3]. Although there are many microbial sources available for producing extracellular enzymes, only a few are recognized as commercial producers[4]. Of these, strains of *Bacillus sp.* dominate the industrial sector[5]. Laccase (p-diphenol: oxygen oxidoreductase; EC 1.10.3.2; also known as p-diphenol oxidase; *p-DPO*; p-diphenolase) is a copper-containing hydrolase[6], which has an ability to catalyze the oxidation of a wide variety of organic and inorganic compounds by coupling it to the reduction of oxygen to water[7].

There is an increasing demand for laccase in the market for various applications such as biopulping[8], biobleaching[9], denim bleaching[10], organic synthesis[11], decolorization[12], dechlorination of xenobiotic compounds[13], bioremediation [14], plant fiber modification, ethanol production, wine

stabilization, baking[15], cosmetic and dermatological preparations [16], biofuel cells *etc.*[17]. Many microorganisms like species of *A. lipoferum*[18], *Aquifexaolicus*[19], *Azospirillum lipoferum*[20], *B. subtilis*[21], *B. sphaericus*[22], *B. halodurans* LBH-1 [23], *Escherichia coli* [24], *Marinomonas mediterranea* [25], *Oceanobacillus iheyensis*[26], *P. maltophilia*[27], *P. syringae*[28], *P. fluorescens* GB-1[29], *P. putida* GB-1[30], *P. desmolyticum* NCIM 2112[31], *P. aerophilum*[32], *Streptomyces sp.*[33], *Thermusthermophilus* TTC1370[34], *Xanthomonas campestris*[35] have been evaluated for the production of laccase.

Microbes serve as the preferred source of laccase because of their rapid growth, the limited space required for their cultivation, and the ease with which they can be genetically manipulated to generate new enzymes with altered properties[2]. However, high cost and low yields of laccase have been the main problems for its industrial production. Therefore, there is a great need to develop new strains with inexpensive mutagens that provides a high laccase yield.

For industrial use enzyme must be produced at low cost and should be reusable and reproducible[2]. To achieve this many techniques has been developed for strain improvement. Strain improvement is usually done by mutating the microorganism that produces the enzyme by techniques such as classical mutagenesis, which involves exposing the microbe to physical mutagens such as X-rays, γ -rays, UV rays, etc., and chemical mutagens such as NTG, EMS, EtBr, etc.[2,36].

There are a great number of literatures reported to use the strain improvement process for producing various industrial enzymes like lipase, chitinase, cellulase, glucoamylase, protease and fibrinolytic protease [2,37-39]. But there was no report available on mutation studies of *Bacillus subtilis* MTCC 2414 for laccase production. The present investigation was undertaken to improve the laccase quantitatively from *Bacillus subtilis* MTCC 2414 strain through exposure to physical and chemical mutagens.

MATERIALS AND METHODS

Experimental Chemicals

All chemicals and reagents of analytical grade were used in this research and are mostly purchased from sigma USA and Hi media Mumbai. All the experiments were conducted in triplicate and the mean values are considered.

Micro organism and Inoculum Preparation

The *Bacillus subtilis* MTCC 2414 strain that produces laccase was employed in the present study. The organism was procured from IMTECH, Chandigarh, India. Stock cultures were maintained in nutrient broth medium with 70% glycerol, cultures were preserved at -20°C [40]. The inoculum was prepared by transferring a loopful of stock culture (*Bacillus subtilis* MTCC 2414) to a certain volume (100 ml) of sterile nutrient broth, stock medium, then incubated it overnight at 37°C on a rotary shaker 200 rpm, before being used for inoculation[2]. A stock suspension was prepared and adjusted to 7×10^8 cells/ml.

Mutation and Selection

Mutagenesis by UV Irradiation

Four ml of the spore suspension containing 1×10^7 spores/ml

was pipetted aseptically into sterile petridish of 80 mm diameter having a flat bottom. The exposure of spore suspension to UV light was carried at a distance of 30 cm away from the UV lamp (15 W, 2537Å). The exposure times were 30 to 90 min [2,41-46]. Each UV exposed spore suspension was stored in the dark overnight to avoid photo reactivation, then was serially diluted in saline and plated in agar medium using 0.04% guaiacol. The plates were incubated for 24h at 37°C and the numbers of colonies on each plate was counted. Each colony was assumed to be formed from a single spore. Mutants for hyper production of laccase were detected visually by the intensity of zone sand were further selected based on their capacity of enzyme production in the liquid medium.

Mutagenesis by Ethidium bromide (EtBr)

Four ml of the UV mutated spore suspension containing 1×10^7 spores/ml was pipetted aseptically into 15 ml of (0.5 mg/ml) Ethidium bromide (EtBr) solution and incubated at 37°C . The sample (2 ml) of this solution was taken at intervals of 30 to 270 min and centrifuged immediately at 10,000 rpm for 5 min at 37°C [2,43,47-50]. The supernatant was decanted and the cell pellet obtained was resuspended in 5 ml saline to stop the reaction. The washed cell suspension was serially diluted in saline and plated on agar medium using 0.04% guaiacol. The plates were incubated for 24h at 37°C and the number of colonies on each plate was counted. Each colony was assumed to be formed from a single spore. Mutants for hyper production of laccase were detected visually by the intensity of zones and were further selected based on their capacity of enzyme production in the liquid medium.

Ethyl Methane Sulfonate (EMS) Mutagenesis

Four ml of the EtBr mutated spore suspension containing 1×10^7 spores/ml was pipetted aseptically into 15 ml of (0.5 mg/ml) ethyl methane sulfonate (EMS) solution and incubated at 37°C . The sample (2 ml) of this solution was taken at intervals of 30 to 270 min and centrifuged immediately at 10,000 rpm for 5 min at 37°C [2,43,44,49-51]. The supernatant was decanted and the cell pellet obtained was re suspended in 5 ml saline to stop the reaction. The washed cell suspension was serially diluted in saline and plated on agar medium using 0.04% guaiacol. The plates were incubated for 24h at 37°C and the number of colonies in each plate

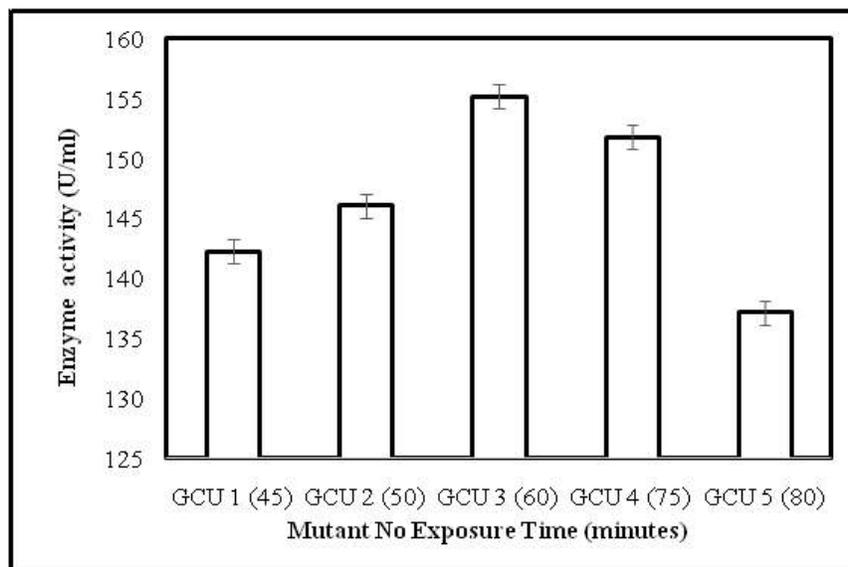


Fig 1. : Mutants after the treatment with UV radiations

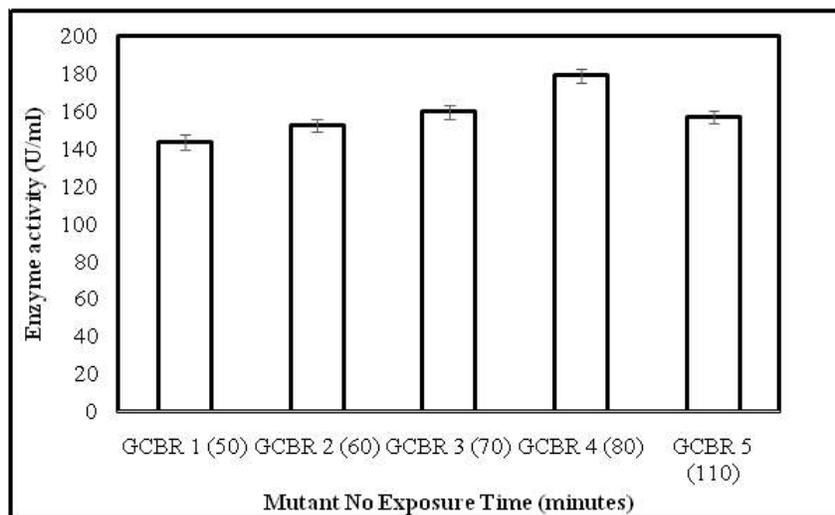


Fig 2. : Mutants after the treatment with EtBr

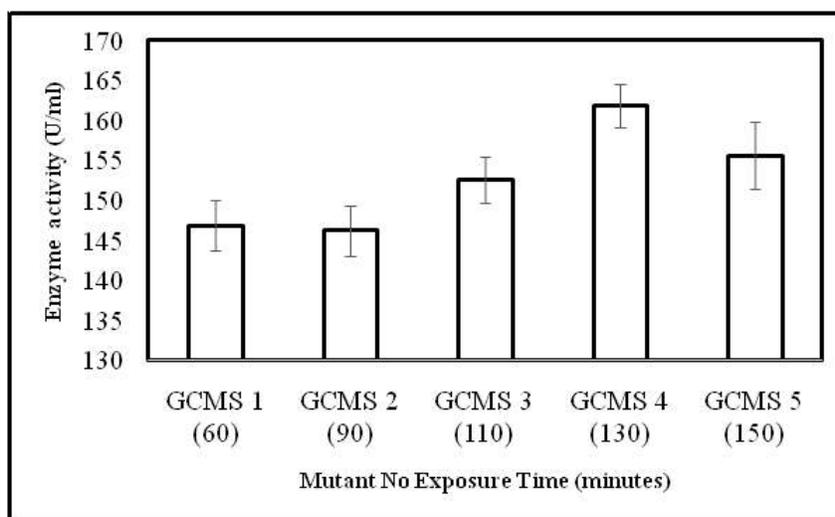


Fig 2. : Mutants after the treatment with EMS

were counted. Each colony was assumed to be formed from a single spore. Mutants for hyper production of laccase were detected visually by the intensity of zones and were further selected based on their capacity of enzyme production in the liquid medium.

Production of Laccase by Shake Flask Fermentation

Hyper laccase producing mutant strains of ultra-violet rays, ethyl methane sulfonate and ethidiumbromide were inoculated separately in the fermentation medium. The medium used for the production of laccase contains (g/l) 1.0 glucose, 1.0 peptone, 0.2 K_2HPO_4 , 0.04 $CaCl_2$, 0.02 $MgSO_4$, 0.0002 $FeSO_4$, 0.001 $ZnSO_4$ and 0.02 guaiacol (Hi media, Mumbai, India) as inducer for laccase production. The pH 6 of the medium was adjusted with 1N HCl/NaOH. Two percent (v/v) of 24h old inoculum suspension was transferred to 50 ml of growth medium in 250 ml Erlenmeyer flasks. These flasks were then placed in the rotary incubator shaker (Lab top) rotating at 200rpm for 24h at 37°C. After the completion of fermentation the whole fermentation broth was centrifuged at 10,000 rpm at 4°C for 10 min and the clear supernatant (crude enzyme) was used for the estimation of

laccase. All the experiments were run parallel in triplicate.

Determination of Laccase Activity

Laccase activity was measured by monitoring the oxidation of 1mM guaiacol (Hi media, Mumbai, India) buffered with 0.2 M sodium phosphate buffer (pH 6) at 420 nm for 1 min. The reaction mixture (900 μ l) contained 300 μ l of 1 mM guaiacol, culture filtrate, and 0.2 M sodium phosphate buffer (pH 6). One unit of the enzyme activity was defined as the amount of enzyme that oxidized 1 μ mol of guaiacol per minute. The enzyme activity was expressed in U/ml [52].

RESULTS

Improvement of Strain by Physical (UV Radiation) Mutagenesis

Data of Fig: 1 shows the production of laccase by UV treated strains of *Bacillus subtilis* MTCC 2414. The parental strains of *Bacillus subtilis* MTCC 2414 was subjected to UV treatment for different time intervals i.e. from 30 to 90 min. Of all the isolates investigated, maximum enzyme production (155.2 ± 1.19 U/ml/min) was obtained by GCU 3 which was selected after 60

min of UV treatment.

Improvement of Strain by Ethidium Bromide (EtBr) Mutagenesis

The mutant strains of *Bacillus subtilis* MTCC 2414 was selected after the treatment of the parental strain with EtBr (Fig: 2). The GCU 3 strain of *Bacillus subtilis* MTCC 2414 was subjected to EtBr treatment for different time intervals i.e. 30 to 270 min. Of all the isolates investigated, maximum enzyme production (178.8 ± 3.67 U/ml/min) was obtained by GCBR 4.

Improvement of Strain by Ethyl Methane Sulfonate (EMS) Mutagenesis

The strains of *Bacillus subtilis* MTCC 2414 were screened after the chemical treatment with EMS (Fig: 3). The GCBR 4 strains of *Bacillus subtilis* MTCC 2414 was subjected to EMS treatment for 30 to 270 min. Of all the isolates investigated, maximum enzyme production (161.8 ± 2.76 U/ml/min) was obtained by GCMS 4.

DISCUSSION

Five UV treated mutant strains of *Bacillus subtilis* MTCC 2414 were isolated on the basis of a bigger zone of guaiacol oxidation in the petri plates and named as GCU isolates. These strains were screened for laccase production under submerged fermentation. Of all the isolates investigated, maximum enzyme production (155.2 ± 1.19 U/ml/min) was obtained by GCU 3 which was selected after 60 min of UV treatment. The production of enzyme following the growth of the organism was found to be highly significant than other mutant derivatives. But this mutant was not stable. It may be due to the reason that the mutant produced by UV irradiations had undergone back mutations when they were exposed to light [41]. Five EtBr treated mutant strains of *Bacillus subtilis* MTCC 2414 was selected on the basis of a bigger zone of guaiacol oxidation in the petriplates and named as GCBR isolates. Further screening of the strains for laccase production under submerged fermentation. Of all the isolates investigated, maximum enzyme production (178.8 ± 3.67 U/ml/min) was obtained by GCBR 4. Five EMS treated mutant strains of *Bacillus subtilis* MTCC 2414 was isolated on the basis of a bigger zone of guaiacol oxidation in the petriplates and named as GCMS isolates. This mutant showed improvement in the production of the enzyme. Of all the isolates investigated, maximum enzyme production (161.8 ± 2.76 U/ml/min) was obtained by GCMS 4.

CONCLUSION

The search for promising strains of laccase producers is a continuous process. In this study the wild strain of *Bacillus subtilis* MTCC 2414 was improved for enhanced laccase production by using physical and chemical mutagens. GCBR 4 with hyper laccase production (178.8 ± 3.67 U/ml/min) was obtained after 80 min treating wild strain with Ethidium bromide. Finally, from the above results it was concluded that *Bacillus subtilis* MTCC 2414 mutant strain was developed by using EtBr treatment remained as a stable mutant after multiple culture cycles. GCBR 4 mutant showed 2 fold higher laccase production than the wild strain in flask fermentation. Thus the selected mutant has potential in minimizing the cost of enzyme for its biotechnological applications.

ACKNOWLEDGEMENTS

The authors would like to thank the chairman Garden City

Group of Institutions Dr. Joseph VG for providing laboratory facilities and supporting this work.

REFERENCES

- Sharma S, Aneja MK, Mayer J, Scholter M, Munch JC. RNA fingerprinting of microbial community in the rhizosphere soil of grain legumes. *Fems Microbiol Lett.* 2004; 240: 181-186.
- Venkatanagaraju E, Divakar G. *Bacillus Cereus* GD 55 Strain Improvement by Physical and Chemical Mutagenesis for Enhanced Production of Fibrinolytic Protease. *International Journal of Pharma Sciences and Research.* 2013a; 4(5): 81-93.
- Venkatanagaraju E, Divakar G. Optimization and Production of Fibrinolytic Protease (GD kinase) from Different Agro Industrial Wastes in Solid State Fermentation. *Current Trends in Biotechnology and Pharmacy.* 2013b; 7(3): 763-772.
- Gupta R, Beeg Q, Khan S, Chauhan B. An overview on fermentation, downstream processing and properties of microbial alkaline proteases. *Appl Microbiol Biotechnol.* 2002a; 60(4): 381-395.
- Gupta R, Beeg Q, Loran P. Bacterial alkaline protease: molecular approaches and industrial applications. *Appl Microbiol Biotechnol.* 2002b; 59(1): 15-32.
- Yoshida H. Chemistry of lacquer (Urushi). Part 1. *Journal of the Chemical Society of London.* 1883; 43: 472-486.
- Dwivedi UN, Singh P, Pandey VP, Kumar A. Laccases production. *Journal of Mol Catal.* 2011; 68: 117-128.
- Call HP, Mucke I. History, overview and applications of mediated lignolytic systems, especially laccase-mediator systems (Lignozym® process). *J Biotechnol.* 1997; 53: 163-165.
- Balakshin M, Capanema E, Chen CL. Biobleaching of pulp with dioxygen in the laccase mediator system reaction mechanisms for degradation of residual lignin. *J Mol Catal.* 2001; 13: 1-3.
- Pazarlıoğlu NK, Sarıik M, Telefoncu A. Laccase: production by *Trametes versicolor* and application to denim washing. *Process Biochem.* 2005; 40: 1673-1675.
- Aktas N, Tanyolaç A. Reaction conditions for laccase catalyzed polymerization of catechol. *Bioresour Technol.* 2003; 87: 209-211.
- Soares GMB, Amorim MTP, Hrdina R. Studies on the biotransformation of novel disazo dyes by laccase. *Process Biochem.* 2002; 37: 581-583.
- Ullah MA, Bedford CT, Evans CS. Reactions of pentachlorophenol with laccase from *Coriolus versicolor*. *Appl Microbiol Biotechnol.* 2000; 53: 230-235.
- Pointing SB. Feasibility of bioremediation by white-rot fungi. *Appl Microbiol Biotechnol.* 2001; 57: 20-23.
- Selinheimo E, Kruus K, Buchert J. Effects of laccase, xylanase and their combination on the rheological properties of wheat doughs. *J Cereal Sci.* 2006; 43: 152-156.
- Golzberner K, Walzel B, Zastrow L. Cosmetic and dermatological preparation containing copper binding

- proteins for skin lightening. Int Pat Appl. 2004: WO2004017931.
17. Alcalde M. Lacases: Biological Functions, Molecular Structure and Industrial Applications. J Mac Cabe. 2007: 5: 461-476.
 18. Diamantidis G, Effosse A, Potier P, Bally R. Purification and characterization of the first bacterial laccase in the rhizospheric bacterium *Azospirillum lipoferum*. Soil Biol Biochem. 2000: 32: 919-927.
 19. Deckert G, Warren PV, Gaasterland WG, Young AL. The complete genome of the hyperthermophilic bacterium *Aquifexaelicus*. Nature. 1998: 393: 353-358.
 20. Givaudan A, Effosse A, Faure D, Potier P, Bouillant ML, Bally R. Polyphenol oxidase in *Azospirillum lipoferum* isolated from rice rhizosphere: Evidence for laccase activity in nonmotile strains of *Azospirillum lipoferum*. FEMS Microbiol Lett. 1993: 108: 205-210.
 21. Narayanan PM, Beenie J, Antony J, Murugan S. Production of Extracellular Laccase from *Bacillus subtilis* MTCC 2414 Using Agroresidues as a Potential Substrate. Biochemistry Research International. 2015: doi.org/10.1155/2015/765190.
 22. Claus H, Filip Z. The evidence of a laccase like activity in a *Bacillus sphaericus* strain. Microbiol Res. 1997: 152: 209-215.
 23. Ruijssenaars HJ, Hartmans S. A cloned *Bacillus halodurans* multicopper oxidase exhibiting alkaline laccase activity. Applied Microbiol Biotechnol. 2004: 65: 177-182.
 24. Kim C, Lorentz WW, Hoopes JT, Dean FF. Oxidation of phenolate siderophores by the multicopper oxidase encoded by the *Escherichia coli* *yacK* gene. J Bacteriol. 2001: 183: 4866-4875.
 25. Kalme S, Parshetti G, Jadhav S, Govindwar S. Biodegradation of benzidine based dye Direct Blue-6 by *Pseudomonas desmolyticum* NCIM 2112. Bioresour Technol. 2007: 98: 1405-1410.
 26. Takami H, Takaki Y, Uchiyama I. Genome sequence of *Oceanobacillus iheyensis* isolated from the Iheya Ridge and its unexpected adaptive capabilities to extreme environments. Nucl Acids Res. 2002: 30: 3927-3935.
 27. Isono Y, Hoshino M. Laccase like activity of nucleoside oxidase in the presence of nucleosides. Agri Biol Chem. 1989: 53: 2197-2203.
 28. Cha JS, Cooksey DA. Copper resistance in *Pseudomonas syringae* by periplasmic and outer membrane proteins. Proc Natl Acad Sci USA. 1991: 88: 8915-8919.
 29. Okazaki M, Sugita T, Shimizu M, Ohode Y, Iwamoto K. Partial purification and characterization of manganese oxidizing factors of *Pseudomonas fluorescens* GB-1. Applied Environ Microbiol. 1997: 63: 4793-4799.
 30. Brouwers GJ, Devring JPM, Cornelis P. *CumA*, a gene encoding a multicopper oxidase, is involved in Mn²⁺ oxidation in *Pseudomonas putida* GB-1. Applied Environ Microbiol. 1999: 65: 1762-1768.
 31. Solano F, Lucaselio P, Lopez serrano D, Fernandez E, Sanchezamat A. Dimethoxyphenol oxidase activity of different microbial blue multicopper proteins. FEMS Microbiol Lett. 2001: 204: 175-181.
 32. Fitzgibbon ST, Ladner H, Kim UJ, Stetter KO, Simon MI, Miller JH. Genome sequence of the hyper thermophilic archaeon *Pyrobaculum aerophilum*. Proc Natl Acad Sci. 2002: 99: 984-989.
 33. Arias ME, Arenas M, Rodriguez J, Soliveri J, Ball AS, Hernandez M. Kraft pulp biobleaching and mediated oxidation of a nonphenolic substrate by laccase from *Streptomyces cyaneus* CECT 3335. Applied Environ Microbiol. 2003: 69: 1953-1958.
 34. Miyazaki K. A hyperthermophilic laccase from *Thermus thermophilus* HB27. Extremophiles. 2005: 9: 415-425.
 35. Lee YA, Hendson M, Panopoulos NJ, Schroth MN. Molecular cloning, chromosomal mapping and sequence analysis of copper resistance genes from *Xanthomonas campestris* sp. Juglandis: Homology with small copper proteins and multicopper oxidases. J Bacteriol. 1994: 176: 173-188.
 36. Parekh S, Vinci VA, Strobel RJ. Improvement of microbial strains and fermentation processes. App Microbiol Biotechnol. 2004: 54: 287-301.
 37. Haki G, Rakshit SK. Development in industrially important thermostable enzymes a review. Biores Technol. 2003: 89: 17-34.
 38. Bapiraju KVSN, Sujatha P, Ellaiah P, Ramana T. Mutation induced enhanced biosynthesis of lipase. Afr J Biotechnol. 2004: 3(11): 618-624.
 39. Koncerova H, Vachova L, Chaloupka J. Mutants of *Bacillus megaterium* with altered synthesis of an exocellular neutral proteinase. Folia Microbiol. 1984: 29: 99-103.
 40. Richard RB, Murray PD. Guide to protein purification. 2nd ed Academic Press is an Imprint of Elsevier. 2009: 463: 14-15.
 41. Haq I, Khurshid S, Ali S, Ashraf MA, Rajoka MI. Mutation of *Aspergillus niger* for hyper production of citric acid from black strap molasses. World J Microbiol Biotechnol. 2001: 17: 35-37.
 42. Munazzah M, Ibraheem M, Sadia J. *Bacillus subtilis* improvement through UV and chemical mutagenesis for indigenously hyper produced urate oxidase. Pak J Life Soc Sci. 2012: 10: 123-129.
 43. Ambika V, Karuna D, Poonam S. Hyper-Production of Laccase by *Pseudomonas putida* LUA15.1 through Mutagenesis. J Microbiol Exp. 2016: 3(1): 3-8.
 44. Mahmoud MG, Rifaat HM, El Sayed OH, El Beih FM, Selim MS. Effect of inducers and process parameters on laccase production by locally isolated marine *Streptomyces lydicus* from Red Sea, Egypt. International Journal of Chem Tech Research. 2013: 5(1): 15-23.
 45. Rasheeda K, Gyana R. Strain improvement of white rot fungi *Pycnoporus cinnabarinus* with the influence of physical and chemical mutagens for enhancing laccase production. Journal of Scientific and Industrial Research. 2014: 73: 331-337.
 46. Adebayo E, Adegoke O, Julius K, Achana Y, Tarun B.

- Improvement of Laccase Production in *Pleurotus pulmonarius* - LAU 09 by Mutation. Journal of Microbiology Research. 2012; 2(1): 11-17.
47. Michaelis G, Douglass S, Tsai M, Criddle RS. Mitochondrial DNA and suppressiveness of petite mutants in *Saccharomyces*. Biochem Genetics. 1971; 5: 487-495.
 48. Mohsinjaved MD, Ikram UL, Irfana MR. Multistep mutagenesis for the over expression of cellulose in *Humicola insolens*. Pak J Bot. 2011; 43: 669-677.
 49. Malik SH, Tehreema I, Ikramul HAQ. Enhanced Amyloglucosidase biosynthesis through mutagenesis using *Aspergillus niger*. Pak J Bot. 2011; 43: 111-119.
 50. Munazzah M, Ibraheem M, Sadia J. *Bacillus subtilis* improvement through UV and chemical mutagenesis for indigenously hyper produced urate oxidase. Pak J Life Soc Sci. 2012; 10: 123-129.
 51. Ferron VMA, Lopez CJL, Perez SJA, Sevilla FJM, Chisti Y. Rapid screening of *Aspergillus terreus* mutants for overproduction of lovastatin. World J Microbiol Biotechnol. 2005; 21: 123-125.
 52. Das N, Sengupta S, Mukherjee M. Importance of laccase in vegetative growth of *Pleurotus florida*. Applied and Environmental Microbiology. 1997; 63(10): 4120-4122.