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Potential of bacterial chitinolytic, *Stenotrophomonas maltophilia*, in biological control of termites

Faiza Jabeen^{1*}, Ali Hussain², Maleeha Manzoor¹, Tahira Younis¹, Azhar Rasul¹ and Javed Iqbal Qazi³

Abstract

Termites are important pest of crops, trees, and household wooden installments. Two species *Coptotermes heimi* and *Heterotermes indicola* are the major species of termites that results in great economic loss in Asia including Pakistan. Chitinases have drawn interest because of their relevance as biological control of pests. The study was performed to screen chitinolytic bacteria from dead termites and to determine their chitinolytic activity in degrading chitin content of termites. Ten isolates were obtained forming clear zones on chitin-containing agar plates. One isolate (JF66) had the highest (3.3 mm) chitinolytic index. Based on sequence of 16S rRNA gene, the isolate was identified as *Stenotrophomonas maltophilia* with (99%) similarity under Accession number KC849451 (JF66), and DNA G + C content was found to be (54.17%). *S. maltophilia* (JF66) produces chitinases upto 1757.41 U/ml at 30 °C and pH 6.0 employing diammonium phosphate as a nitrogen source. Chitinase gene was also extracted and gets sequenced that confirmed its presence. Whole culture and different concentrations of crude enzyme of the isolate were tested on the chitin covering of termites. Mortalities showed that crude enzyme of isolate could degrade chitin of both species of the termites *C. heimi* and *H. indicola*. Chitinase produced by *S. maltophilia* had potential application as biocontrol agent for termites, but it is assumed that purification of chitinases may produce more prominent results.

Keywords: Termite, *Stenotrophomonas maltophilia*, Chitinases, Biological control

Background

Men have always required control measures for destructive insects like mosquitoes, bed bugs, crop pests, and many those competing with their assets such as termites. The most commonly used pest control measures are synthetic pesticides (i.e., malathion, methyl parathion, DDT, and lindane) which are the most widely used pest control measure but their frequent utilization has created a distressed ecological balance among pests resulting in resistance development in the pests as well as persistence in environment with deteriorative potential. Natural enemies are also affected when pesticides are applied hastily, which resulted in a resistant pest population, damaging crops and other products (Pohanka 2006; Thomas and Read 2007; Kumar et al. 2008). More than (99.9%) of pesticides

move into the environment rather than to the target which adversely influences public health and other valuable biota and pollutes the soil and water as well as the atmosphere (Pimentel 1995; El-Heneidy et al. 2015). The negative impact on environment and human health has educated scientist to track other substitutes. In this regard, biological control is one of the alternatives which proved to be safe and help in lowering pest density (Kumar et al. 2008 and Rakshiya et al. 2016).

Among pests, the most damaging to wood, household wooden installments, and crops are termites (Akhtar and Rashid 2001; Debelo and Degaga 2017). Pakistan is an agricultural country, and its economy depends on cash crops that are severely damaged by termite's infestation. In this region, two termite species *Heterotermes* and *Coptotermes* belong to Rhinotermitidae family are of immense economical importance. They build shelter tubes with soil, fecal material, and saliva but feed on vital plant parts and wood (Jenkins et al. 2002; Qasim et

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al. 2015). In Pakistan, especially the two termite species *Coptotermes heimi* and *Heterotermes indicola* attack wood and wooden material (Saliha et al. 2012).

Biological control has been acknowledged as more environment friendly and a safer method than reliance on chemical pesticides (Eilenberg and Hokkamen 2006; Kumar et al. 2008). In this scenario, microbial control agents are harmless as well as efficient because of their efficacy and undisruptive nature for non-target organisms (Lacey et al. 2001; Boisvert 2005). Bacterial chitinases, especially those from thermophilic bacteria showed a high potential against termites' attack in an environmentally safe manner. Chitinases are the enzymes capable of chitin hydrolysis to low molecular weight end products that are of great interest as a potential biological control agent (Huang and Chen 2004; Makino et al. 2006; Wang et al. 2006). So, chitinolytic bacteria have a great potential in controlling insects under selected conditions (Zhu et al. 2008; Osman et al. 2015). Termites like other insects bear chitin for polymerizing and strengthening their exoskeleton and other tissue/organs like trachea and gut epithelium (Merzendorfer and Zimoch 2003; Al-Sawalmih 2007). The gut of termites is indicative to design control strategy involving chitin degradation leaving insect incapable of digestion and absorption (Wiwat et al. 2000; Shternshis et al. 2002). Thus, application of bacterial chitinases for eradicating termite damages might protect humans from side effects of toxic synthetic chemicals. This is the first attempt regarding the application of bacterial chitinases in biological control of termites.

In the present study, bacterial chitinases production by *Stenotrophomonas maltophilia* and their termiticidal activity was quantified.

Materials and methods

Chitin extraction of crab

Blue crab (*Callinectes sapidus*) was purchased from market and washed with water, and shells were separated. The shells were oven dried at 105 °C and crushed to powder. Chitin polymer was extracted according to the method of Rhazi et al. (2000). Finally, extracted chitin powder was dried and saved in bottles.

Processing for colloidal chitin

Chitin was processed according to the method of Jabeen and Qazi (2014). The extracted chitin in filter paper was washed several times by autoclaved distilled water till the spent wash water's pH became 7. The colloidal chitin was removed from the filter paper, weighed, and stored in dark bottle at 4 °C.

Sampling and isolation of the chitinolytic bacterium

Termites were sampled from trees of Quaid-e-Azam campus area, University of the Punjab, Lahore, Pakistan.

The samples were saved in tightly capped sterilized bottles and transported to laboratory for further processing. The dead termites were enriched in selective medium prepared after Furukawa et al. (1978). The medium contained 1% prepared colloidal chitin as sole carbon source; NH_4SO_4 , 1.0; KH_2PO_4 , 0.2; K_2HPO_4 , 1.6; NaCl, 0.1; $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$; $\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$, 0.01; $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$, 0.02 (g/L) dissolved in distilled water. After 5 days of incubation at 30 °C temperature, dilution was made and spread on chitin agar plates, incubated at 30 °C for 3 days. Zones of clearance appeared around colonies and then pure cultured through nutrient agar.

Chitinase assay

Chitinolytic activity of the isolate was determined by the estimation of released reducing sugars from the chitin as described by Sadafi et al. (2001). The standard curve was plotted with *N*-acetylglucosamine (NAG) in the range of 100 to 600 µg/ml. One unit of chitinolytic activity was described as 1 µmol of liberation of NAG per milligram of protein per minute.

Protein test

The protein content was estimated by the method described by Bradford (1976). Bovine serum albumin (BSA) prepared from 2 to 10 µg/ml range with two class intervals was used as standard. Calibration curve was then plotted by performing regression analysis of A^{595} absorbance versus corresponding concentrations of the standards.

16S rRNA gene sequencing

Freshly grow bacterial colony was suspended in 5 ml sterilized nutrient broth and grown for overnight. The culture was centrifuged at 10,000 for 10 min, and pellet was processed for DNA extraction. Bacterial 16S rRNA gene was amplified by using the universal primers 27F(5'-AGAG TTGATCMTGGCTCAG-3') and 1492R(3'-TACG G{Y}TACCTTGTTACG-5') (Oligo, USA). Extraction and amplification was performed according to the method described in Jabeen and Qazi (2014). Sequencing was done from Korea and matched with the nucleotide database available at Gene Bank, using BLAST tool in NCBI (<http://www.ncbi.nlm.nih.gov>) for recognition of the highest percentage similarity with the described species.

Phylogenetic analysis

BLAST sequences were imported into the clustalW program package. The sequence was aligned with the closest relative and phylogenetic tree was plotted by using the neighbor joining program in clustalW package (Fig. 1).

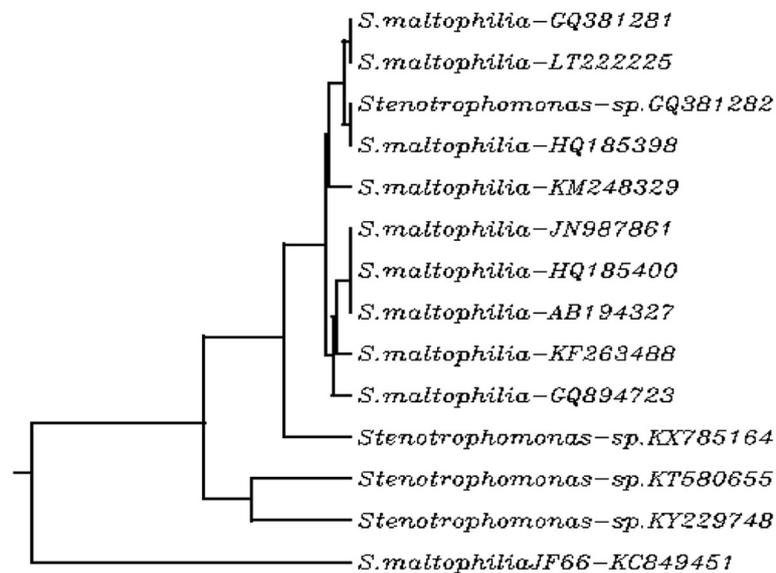


Fig. 1 Phylogenetic tree of the isolate showing highest homology with *Stenotrophomonas* sp. consensus neighbor-joining phylogenetic rooted tree from sequences of the 16S rRNA gene, encompassing *S. maltophilia*

Optimization of chitinase production

Effects of temperature, pH, and nitrogen source on chitinase production were studied by growing the isolate in chitin broth at different temperature ranging from 20 to 60 °C, pH ranging from 4 to 11 for 5 days. Different nitrogen sources such as ammonium chloride, trypton, gelatin, peptone, ammonium oxalate, ammonium dihydrogen phosphate, yeast extract, and urea were supplemented to the chitin medium to study their influence on chitinase production.

Enzyme characterization

The enzyme was characterized by incubating enzyme to different pH levels from 4 to 11 by using different buffers acetate (4–5), phosphate (6–7), Tris HCl (8–9), and glycine NaOH buffers (10–11) similarly temperatures ranging from 20 to 60 °C for 30 min in shaking water bath. The crude enzyme was also incubated with different concentrations of chitin substrate at optimum pH and temperature for 30 min. The enzyme activity was estimated, as described before.

Amplification of chitinase gene

The extracted DNA was also used to amplify chitinase gene(s) employing the primer sequence used by Lee et al. (2007), i.e., F(5'-AATGGGGAATTCGCAAAGCCAGTCTGAC-3'), and R(5'CTCTCTCTTATC-CTCGAGTATCAAAGTATGAT-3') (Macrogen, Korea) using Pro-mega Go-Taq® Flexi DNA Polymerase (MGW Biotech, Germany). Following Lee et al. (2007), PCR was prepared and amplified. The gel was electrophoresed then along with 1 Kb plus ladder of Invitrogen™ (catalog number:

10787018) and PCR product. The PCR products were purified by using QIA quick PCR purification kit protocol, and purified product was got sequenced commercially from Macrogen, Korea.

Partial purification of chitinase enzyme

The extracellular chitinases was purified through ammonium sulphate precipitation and dialysis. Five hundred milliliter supernatant of selected strain was precipitated with ammonium sulphate at different saturation levels (20–80%) by increasing the concentration of salt by 10% each time following the method described by Jabeen (2011). Chitinase activities and protein concentrations were measured every time during addition of salt fractions to calculate the specific activities of enzyme.

Experiments on termites

Collection of termites

Termites' members of *Heterotermes* and *Coptotermes* were collected from the field and transported to the laboratory. Termites were allowed to remain within the piece of bark of tree for 24 h. Termites were recovered and placed in a cool place (28 ± 2 °C) in a glass small box covered with black cloth till further use.

Concentrating bacterial chitinases for termiticidal assessment

The bacterium was grown in 50 ml FJ medium at their corresponding enzyme optima. The growth was centrifuged in sterile centrifuge tubes, and three different concentrations were made as 1X, 5X, and 10X. Three test tubes were prepared with 10 ml of supernatant in each

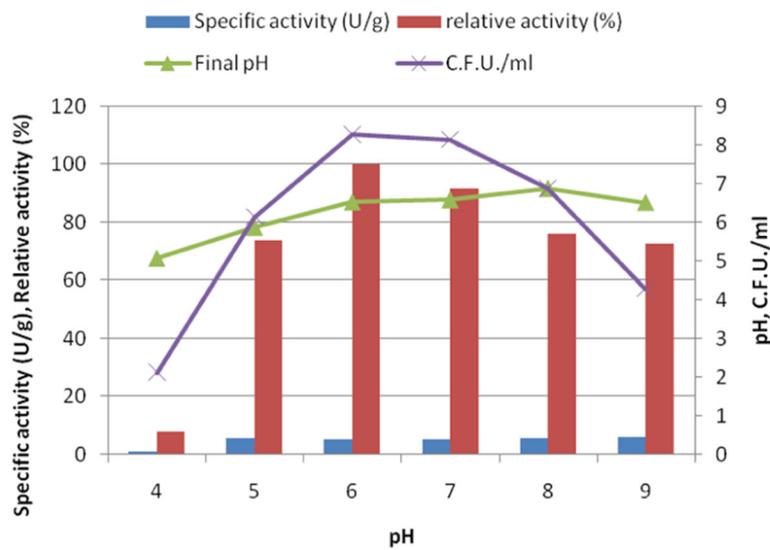


Fig. 2 Effect of pH on specific and relative activity of chitinase produced by *S. maltophilia* JF66 following 5 days of incubation at 37 °C and 140 rpm

sterile capped test tube. The 10 ml supernatant in two of the test tubes (5X, 10X) was dried by exposing to room temperature at sterile conditions till it becomes 5 and 1 ml to attain 5X and 10X concentrations, respectively. A third tube was remained as it is as 1X.

Exposing termites to the bacterial isolates

Filter paper discs of 10 mm were made, sterilized by autoclaving, and loaded with prepared concentrations, i.e., 1X, 5X, and 10X, of the bacterial culture fluids. The discs loaded with 10 µl/disc of given preparation were placed in Petri plates and 10 termites free of extraneous

dust etc. were exposed to them. A control was also run having filter paper discs moistened with simple sterilized distilled water. The Petri plates were covered and kept under darkness at 26 ± 2 °C and were observed for the termite mortality after every hour up to 8 h and then at 24 h post-exposure.

Preparation of bagasse made chipboard

One gram of Sugarcane Bagasse (SCB) was sterilized in each Petri plate and was mixed with 10 ml of a chitinolytic bacterial cell free cultural fluid. The chitinolytic chips of the SCB thus prepared were exposed to the

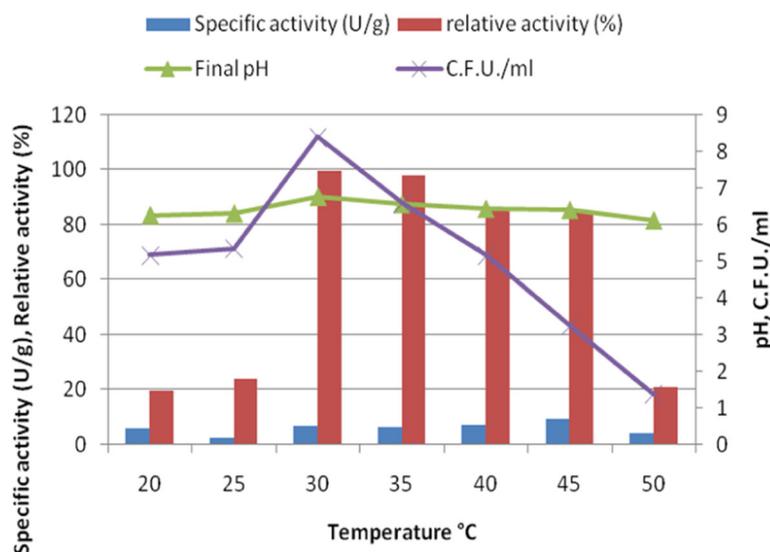


Fig. 3 Effect of different temperatures on specific and relative activities of chitinase produced by *Stenotrophomonas maltophilia* JF66 following 5 days of incubation at 140 rpm

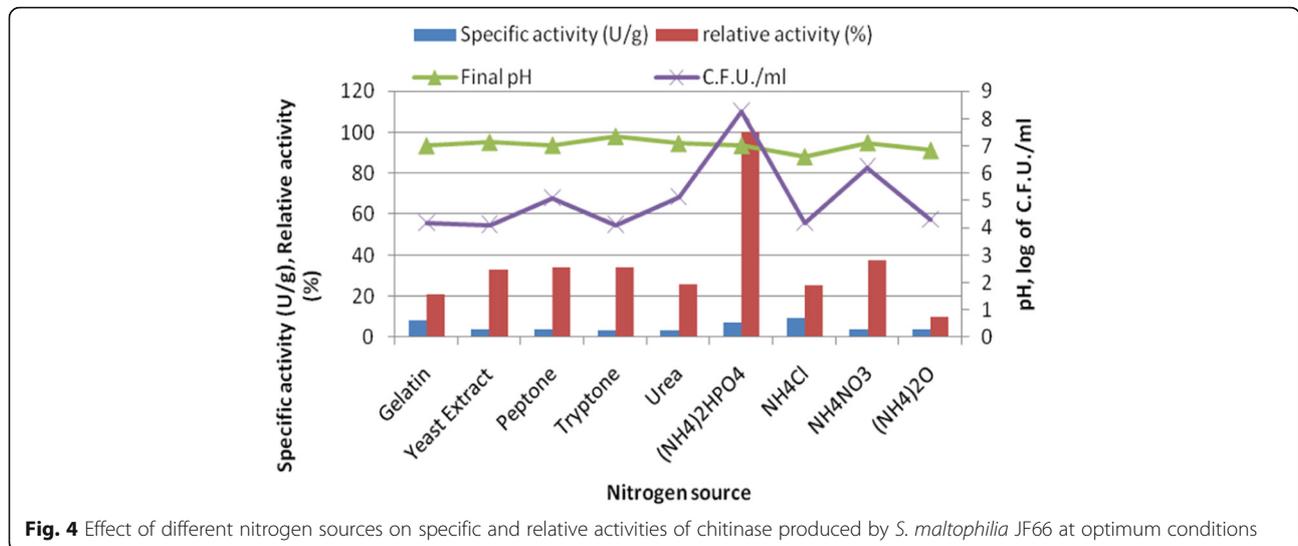


Fig. 4 Effect of different nitrogen sources on specific and relative activities of chitinase produced by *S. maltophilia* JF66 at optimum conditions

termites. Mortality rate of the termite was observed and recorded as mentioned before. The experiments were conducted in triplicates.

Results and discussion

Processing of chitin containing waste

Chitin was used as a sole carbon source for the production of bacterial chitinases. Edible crabs, a waste product of food industries, were chosen for the present study. The shells were processed for removal of minerals by demineralization and proteins by deproteinization. From 4.0 g of the dried chitin wastes 0.984 g (24.6%), chitin was obtained. Following dehydration, the figure came down to 0.55 g and could represent only 13.8% of the total chitin waste. This crude chitin was further processed for attaining colloidal chitin for the production of chitinases. Crude chitin was processed with concentrated hydrochloric acid. The 5.0 g crude chitin used for

acid treatment yielded 54 g of colloidal chitin which was 1080% more colloidal chitin than the crude chitin.

Isolation and screening of chitinolytic bacteria

Eighty-four bacterial strains were isolated from different samples comprising dead termites, termite affected trees, termites' mounds, termite-affected gardens, and termite-affected fields. Samples were processed on chitin containing selective agar medium. Out of the 84 isolated bacterial strains, 35 exhibited vivid zones of hydrolysis on 1.0% chitin agar medium at pH 6.0. The maximum ratio of (ZS) and (CS) to the bacterial colony size was 5.0 mm of the isolate JF66. This isolate was pure cultured on the selective medium intervened by culturing on nutrient agar plates. Puspita et al. (2017) found a high chitinolytic index among various bacteria including *Stenotrophomonas maltophilia*. The pure culture was preserved on FJ medium agar slants. Glycerol stock was also prepared employing growth in FJ broth and was

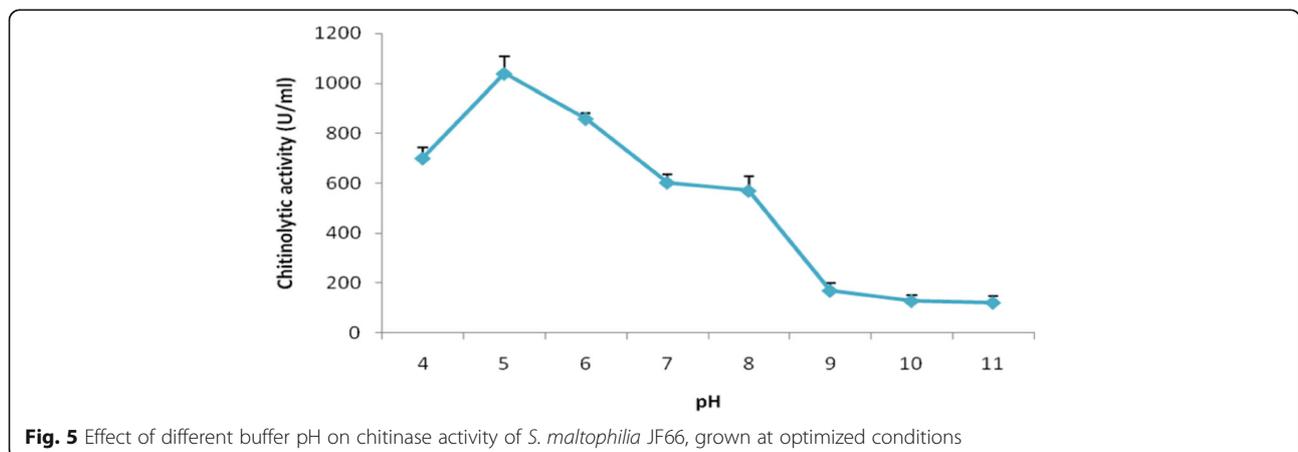


Fig. 5 Effect of different buffer pH on chitinase activity of *S. maltophilia* JF66, grown at optimized conditions

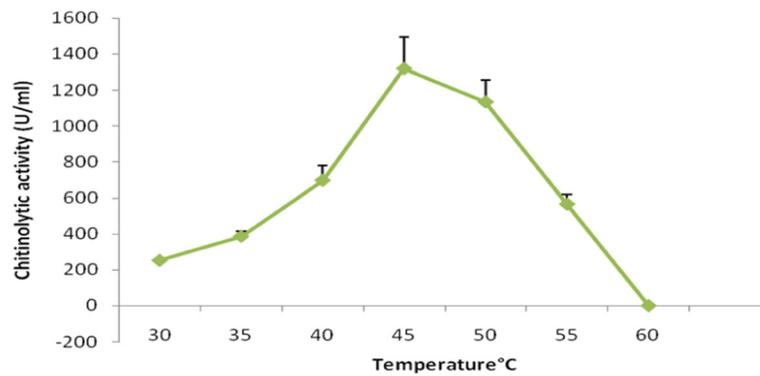


Fig. 6 Effect of incubation temperature on chitinase activity of *S. maltophilia* JF66, grown at optimized conditions

preserved at -20 °C. The isolate JF66 yielded maximum activity of 301.55 U of chitinase/ml.

Identification of isolate

16S rRNA gene sequencing revealed the isolate as *S. maltophilia* with 99% similarity, and their accession number obtained was KC 849451 with GC content 54.172. Alshehri et al. (2016) and Puspita et al. (2017) also identified the bacterium with the same method.

Optimization of growth and chitinase production

Maximum growth and chitinase production was recorded as 116×10^2 C.F.U./ml and 107.72, respectively, at initial pH 6. However, significant amount of enzyme was also produced at pH 7 and could attain (91.97%) of the level obtained at pH 6. Different pH profiles of chitinase produced by a variety of bacterial sources have been reported. Zarei et al. (2011) also found maximum chitinase production at pH employing *Serratia marcescens*. Narayana and Vijayalakshmi (2009) reported similar results with *Streptomyces* sp. Maximum specific

activity observed was 6.163 U/g at pH 9. The bacterial growth could approach 116×10^2 C.F.U./ml, following inoculation in the FJ medium with initial pH 4. While the highest growth up to 293×10^8 C.F.U./ml was recorded in case of pH 6. Growth of the bacterium remained within the two levels mentioned (Fig. 2). By changing the temperature, 35 °C was found to be the most suitable as the chitinase activity/production rises to 512.31 U/ml while raising the temperature up to 55 °C, 78.97% reduction in yield was recorded. In *Streptomyces* sp. optimum temperature for chitinase activity was reported to be 35 °C by Narayana and Vijayalakshmi (2009). At high temperature, decrease in enzyme production indicated its intolerance of elevated temperature (Fig. 3). Chitinase production is considerably increased while changing the nitrogen sources. Among various organic and inorganic nitrogen sources, diammonium hydrogen phosphate was found to be the more suitable supporting enzyme yield, rise to 1757.41 U/ml releasing 0.253 mg/ml protein content. Supplementing diammonium hydrogen phosphate 3.43-fold more chitinase

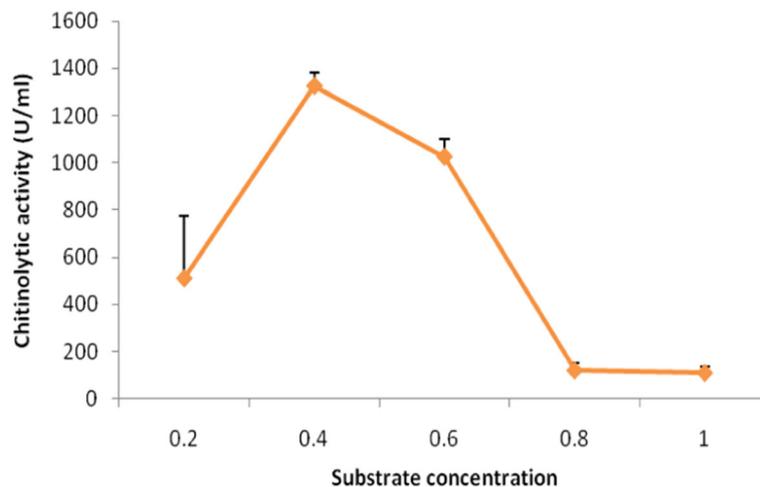


Fig. 7 Effect of substrate concentration on chitinase activity of *S. maltophilia* JF66, grown at optimized conditions

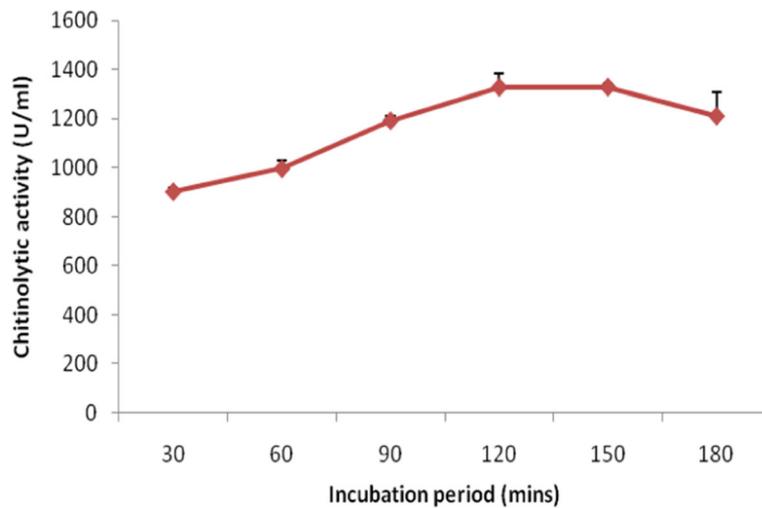


Fig. 8 Effect of incubation time on chitinase activity of *S. maltophilia* JF66 at optimized conditions

production was attained while maximum specific activity 9.367 U/g was observed with ammonium chloride. In our findings, ammonium oxide, gelatine, ammonium chloride, and urea did not support enzyme production actively (Fig. 4). Saima et al. (2013) found malt for maximum chitinase production by *Aeromonas hydrophilia* and *A. punctata* while Jha et al. (2016) reported peptone as suitable nitrogen source.

Optimization and stability of chitinase activity

From the obtained results, it can be reported that pH maintenance plays an important role in chitinase activity. Maximum chitinase activity was found at pH 5 but decreased on raising the pH that reached null at pH 11

(Fig. 5). Thimoteo et al. (2017) determined 5 to 7.5 pH range optimum for chitinase activity. Temperature also affects the enzyme activity. Optimum chitinase activity observed was 1320.40 U/ml found at 45 °C. Sonel et al. (2014) also reported 45 °C temperature for the maximum stability of chitinases produced by *Bacillus subtilis*. The chitinase activity gradually decreases below and above this temperature. No chitinase activity was found at 60 °C, which showed (100%) decrease from optimum chitinase activity (Fig. 6). Different concentration of colloidal chitin was checked to conclude the best concentration for highest chitinase production that can be used at large scale. The enzyme activity increased to 1327.25 U/ml by applying 0.4% substrate concentration

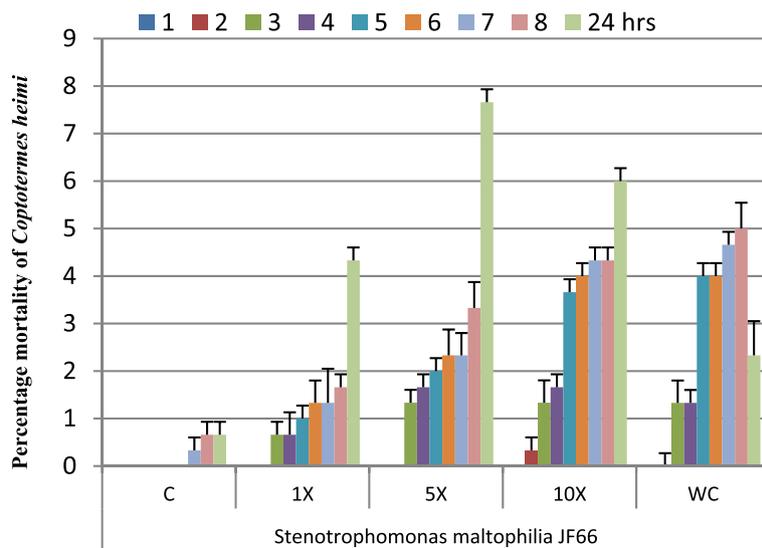


Fig. 9 Percentage mortality of *Coptotermes heimi* exposed to the cultures (WC) and cell-free culture fluids of varying strength of *S. maltophilia* JF66

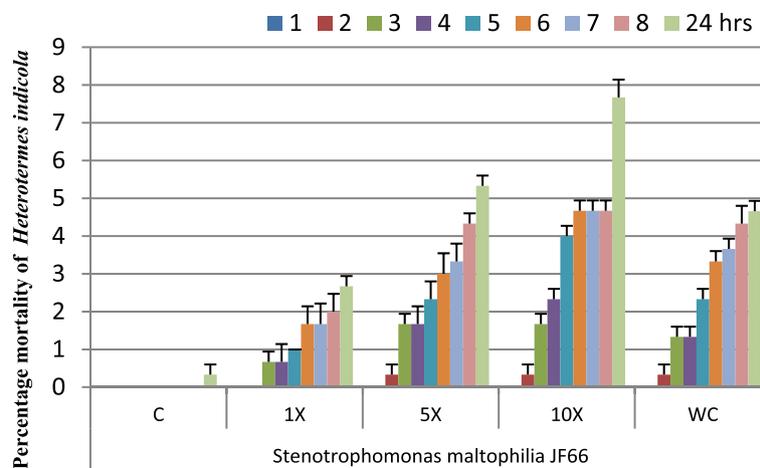


Fig. 10 Percentage mortality of *Heterotermes indicola* exposed to the cultures (WC) and cell-free culture fluids of varying strength of *Stenotrophomonas maltophilia* JF66

while decreased on adding more substrate and found to be reached 111.69 U/ml at 1.0% substrate which was 91.58% less than optimum chitinase activity (Fig. 7). Ajayi et al. (2015) reported 0.15% most suitable for two *Bacillus* species to produce maximum chitinase activity. Obtained results are near to findings of Karunya et al. (2011) who found 0.3% colloidal chitin for the maximum growth. Further chitinase activity was improved by increasing the incubation time that get higher by incubating the enzyme-substrate complex at 150 min (Fig. 8). Shirazi et al. (2007) revealed that considerable chitinase activity was found while increasing incubation time to 120 min.

Extraction of chitinase gene

The amplified chitinase gene showed about 1.7-kb distinct band. The purified amplicon was got sequenced commercially from Korea. Lee et al. (2007) isolated the chitinase gene from *Bacillus* sp. using the same primers and conditions.

Termiticidal effects of the bacterial cultures and their cell-free cultural fluids

The experiments were performed on two termite species, i.e., *Coptotermes heimi* and *Heterotermes indicola*. These species are threat in Punjab (Pakistan) for destruction of house hold wooden structures. Qureshi et al. (2014) also worked on the same species to control the termites by using antflagellates drugs for protozoan living in termites. In this regard, *Bacillus thuringiensis* was reported as insecticidal against many pest including termites by many workers.

This is the first report on effectiveness of bacterial chitinases on termites species frequently found in Pakistan destroying wood and crops. The termiticidal

activity was a function of the bacterial chitinases, as the highest mortalities up to 73% was recorded for 10X cell-free cultural fluids of *S. maltophilia* JF66 within 24 h of exposure in case of *Coptotermes heimi* (Fig. 9). Approaching levels of mortality was recorded for the whole *S. maltophilia* JF66. These observations are tempting to provoke into the bacterial viability following the cultural ingestion in the termites' gut. In case of *Heterotermes indicola* upto 77% mortalities was found within 24 h of exposure. Whole culture of *S. maltophilia* JF66 could cause only 47% mortality (Fig. 10). Husen et al. (2015) worked on Pentoxifylline that enhanced the chitinolytic activity to check termiticidal effect and inhibited the growth of termites.

It was found that more concentrated enzyme caused high termite mortality. Ten times cell-free cultural fluid (containing chitinases) exhibited the highest termite mortality. The whole culture also proved to be termiticidal in general.

Termiticidal effects of the bacterial cell-free cultural fluids with 10% sugarcane bagasse preparation

The highest mortality of 53% was obtained for the *Coptotermes heimi* by the *S. maltophilia* JF66 in contrast to 3.3% deaths of control animals, following 24 h of the exposure. More or less comparable results were obtained for *Heterotermes indicola*, where *S. maltophilia* JF66 caused 57% deaths following 24 h of exposure, in contrast to 6.7% deaths of the control animals. Different microbes were used with varying formulation by many authors without using sugarcane bagasse. Wright and Cornelius (2012) found 72.5% mortality of *Coptotermes formosanus* by applying entopathogenic fungi *Isaria fumosorosea* at 10^8 spores/ml on day 7. Similarly Devi et al. (2007) also revealed that 100% mortality of termite

species *Odontotermes obesus* was achieved by using Rhizobacterial isolates *R. radiobacter* and *Al. latus* following 1-h incubation. Treatment indicated that the strategy hypothesized for controlling the termite's damage of indoor artificial wooden structure need further attention of the researchers for materializing the application of the biological control.

Conclusion

Findings of this work showed that chitin can be extracted from edible crab shell chemically and further purification made it potent carbon and nitrogen source for isolation of chitinolytic bacteria. *S. maltophilia* produced chitinases by utilizing extracted crab chitin. Production of chitinases by *S. maltophilia* increased when their conditions were optimized. Presence of chitinase gene confirmed its chitinolytic activity. Percentage mortality employing chitinases to control the termites proved that chitinolytic *S. maltophilia* is an effective biological control agent of some insect pests. Further purification of chitinases and optimization of other supporting factors may improve the enzyme activity and mortality rate of termites.

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Availability of data and materials

The authors declare that they have no objection to the availability of data and materials.

Authors' contributions

Faiza Jabeen performed the experiments. Ali Hussain helped in statistical analysis. Maleeha Manzoor and Tahira Younis collected the samples. Azhar Rasul worked on collecting and arranging the data. Javed Iqbal Qazi supervised and facilitated the whole project. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Consent for publication

All authors are willing for publications.

Competing interests

The authors declare that they have no competing of interests.

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