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Evaluation of chitinase enzyme in fungal isolates obtained from golden potato cyst nematode (*Globodera rostochiensis*)

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Abstract

The structure of nematode eggshell indicates that proteases and chitinases are necessary for infection by nematophagous fungi, and most of them produce these enzymes. Therefore the aim of this study was to assay chitinase activity in 34 isolates of various species of fungi obtained from golden potato cyst nematode (*Globodera rostochiensis*), as the most important potato pest, over two days of fungal growth (24 and 96 h). Chitinase specific activity was determined by measuring the release of reducing saccharides from colloidal chitin by the *N*-acetylglucosamine-dinitrosalicylate method at 540 nm. Colorimetry based on image processing technique was used to discover colour changes in minimal synthetic medium for validation of chitinase activity. The 34 isolates were identified based on morphological and molecular features including internal transcribed spacer (ITS) regions of ribosomal DNA. Results of the chitinase specific activity measurement showed the chitinase specific activity in all of them for 96 h was higher than for 24 h. Among isolates, the maximum and minimum chitinase specific activity respectively belonged to isolate 154 (0.56 U mg⁻¹) and isolate 6 (0.15 U mg⁻¹) in 24 h, also isolate 113 (1.02 U mg⁻¹) and isolate 6 (0.40 U mg⁻¹) in 96 h. Colorimetric results confirmed that enzyme activity was associated with colour changes. The 34 fungal isolates were classified in 11 genera most of which belonged to *Fusarium*. Finally, two isolates, 113 (*Fusarium oxysporum*) and 154 (*Trichoderma atroviridae*), with the highest chitinase enzyme activity are introduced as potent isolates to control the golden potato cyst nematode that can be used for chitinase enzyme production which is supposed to be used in commercial formulation.

Key words: chitinase, fungi, *Globodera rostochiensis*, potato.

Introduction

The potato (*Solanum tuberosum* L.) is the fifth most valuable crop after wheat, rice, corn and barley in the world (Zarghani et al., 2014). Iran is the third-largest producer of potato in Asia, where the production rate in 2015 was estimated about 5 million tons (Ahmadi et al., 2015).

The golden cyst nematode, *Globodera rostochiensis* (Wollenweber, 1923; Behrens, 1975), is the most destructive pathogen of potato (Gitty et al., 2011). Until recent years the use of chemical pesticides has been one of the most reliable methods for the management of the nematodes (Saifullah, Khan, 2014). However, according to international agreements, the use of chemical nematicides in many countries has been banned because of the deleterious effects on humans, the environment, reduction of the C:N ratio of soil, persistence in soil, pollution of groundwater and the high price of the nematicides (Dong et al., 2004).

Biological control of plant parasitic nematodes is a good management strategy (Sharon et al., 2009). Many natural enemies of plant parasitic nematodes occur that may be useful for biological control. These include pathogens, predators, competitors and antagonists of which 76% are fungi (Khezri Nezhad, 2004). The fungal antagonists have been most extensively studied and are considered the most applicable for biological control of nematodes. The use of biological agents to control of plant-parasitic nematodes is less effective compared with chemical method, but positive result of fungal agents has been encouraging (Nguyen et al., 2007). Tobin et al. (2008) reported that *Pochonia chlamydosporia* was a good biocontrol agent for the potato cyst nematode. López-Lima et al. (2013) reported that the populations of the potato cyst nematode decreased up to 89% with *Paecilomyces*. *Trichoderma* also successfully controlled plant-parasitic nematodes, especially cyst nematodes.

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Chitinases, glucanase and protease are considered the most important mechanism for successful biological control of nematodes (Sharon et al., 2001; Safari Motlagh, Samimi, 2013). Saifullah and Khan (2014) explained that *Trichoderma harzianum* enzymatically penetrated cysts and eggs resulting in death of *G. rostochiensis*. Also Saifullah and Khan (2014) utilizing low-temperature scanning electron microscopy (LTSEM) observed initial infection and the penetration by *T. harzianum* to cyst and egg wall. Manzanilla Lopez et al. (2013) stated that *P. chlamydosporia* is one of the most important fungi for the biocontrol of plant parasitic nematodes. Santos et al. (2013) demonstrated that chitinase and protease of *P. chlamydosporia* was involved in control of the potato cyst and root-knot nematodes.

Plant parasitic nematodes cause important economic losses in agricultural crops. Activity of the fungal egg-parasites leads to immobility and death of the embryos and results in a reduction of nematode population density. The disintegration of nematode eggs can only be caused by enzymatic action. Knowledge of various aspects of chitinolytic enzymatic systems allows the development of a new generation of chitinases and the design of better strategies of biological control (Gortari, Hours, 2008). Fungal chitinases are produced by nematophagous and entomopathogenic fungi to aid infection (Tikhonov et al., 2002). Because chitin is a dominant component of the eggshell in nematodes, the use of chitinase producing fungi is a good strategy for biological control of the golden potato cyst nematode (Morton et al., 2004). In the present study, we evaluated the chitinase production by various nematophagous fungi isolated from the golden potato cyst nematode, as the most important potato pest.

Materials and methods

Fungal isolates. Thirty four fungal isolates were recovered from infected eggs of the golden potato cyst nematode (*Globodera rostochiensis*) in Hamedan, Iran, in 2015. The fungi were grown on PDA (potato dextrose agar) at 25°C.

Morphological and molecular identification. All of the isolates were identified based on morphological (Nagami et al., 2006) and molecular features including internal transcribed spacer (ITS) regions (ITS1, ITS2 and 5.8S gene) of ribosomal DNA.

DNA extraction. For DNA extraction from 34 isolates, fungal mycelia were grown on potato dextrose broth (PDB) for 7 days, then were lyophilized by freeze-drying. Total genomic DNA was extracted from lyophilized mycelia with a QIAGEN DNeasy Plant Mini Kit (Germany). The extracted genomic DNA was diluted in 50 µl distilled water and checked in terms of quality and quantity by means of both 0.8% agarose gel electrophoresis and spectrophotometer technique based on absorption at 260 nm.

Polymerase chain reaction (PCR) amplification. ITS regions (ITS1, ITS2 and 5.8S gene) of ribosomal DNA were amplified with the ITS1 (forward primer) and ITS4 (reverse primer) primers. PCR amplifications were performed using a thermocycler (Corbett Research, Australia) with total reaction volume of 25 µl. In this study, Takara EmeraldAmp GT PCR Master Mix to amplify the ITS region for sequencing was used. Final component concentrations per reaction containing 12.5 µl Takara

EAGT Master Mix (optimized mix of *Taq* polymerase, MgCl₂ and dNTPs), 2.5 µl forward primer (5 µM), 2.5 µl reverse primer (5 µM), 6.5 µl sterile distilled DNA-grade water, 1 µl template DNA (10–500 ng µl⁻¹). Amplification reactions were carried out using the following cycle profile: initial denaturation at 94°C for 1 min followed by 35 cycles including; denaturation at 94°C for 1 min, annealing of primer 51°C for 1 min, extension at 72°C for 1 min and a final extension at 72°C for 8 min. PCR products were separated by electrophoresis technique using 1% agarose gel in 1X Tris-Borate-EDTA (TBE) buffer by adding 12 µl SYBR-safe 10,000X concentrate DNA stain to melted agarose before running of the gel and finally visualized under ultraviolet illuminator. SYBR-safe is a non-toxic chemical that is used to visualize DNA under UV light in agarose gel electrophoresis. The gel image was recorded using BioDoc gel Documentation System (UVP, USA).

Then ExoSAP-IT for PCR product cleanup was done. ExoSAP-IT reagent treats PCR products ranging in size from less than 100 bp over 20 kb with absolutely no sample loss by removing unused primers and nucleotides. To purify the samples, 5 µl post-PCR product was mixed with 2 µl ExoSAP-IT reagent for a combined 7 µl reaction volume, and incubated at 37°C for 15 min to degrade the remaining primers and nucleotides. Finally the samples were incubated at 80°C for 15 min to inactivate the ExoSAP-IT reagent.

The ITS1, ITS2 and 5.8S sequences were obtained by sequencing both strands in opposite directions using the PCR amplification primers, ITS1 and ITS4 in Genomics Resource Laboratory at Massachusetts University, USA.

Enzyme assay. Colloidal chitin was prepared following the procedure of Tikhonov et al. (2002). For enzyme assay in liquid media, the 34 isolates were grown in minimal synthetic medium (MSM) g l⁻¹: (NH₄)₂SO₄, 2.8, urea, 0.6, KH₂PO₄, 4, CaCl₂·2H₂O, 0.6, MgSO₄, 0.2, FeSO₄·7H₂O, 0.01, ZnSO₄·H₂O, 0.0028 and CoCl₂·6H₂O, 0.0032 containing colloidal chitin (1 g l⁻¹) (Zeilinger et al., 1999) at 25°C. The culture medium was filtered through Whatman paper No. 3 filter followed by filtration through 0.2-mm Millipore polydifluoropropylene membranes. The filtrate obtained was analyzed for chitinolytic activity.

Chitinase activity was determined by measuring the release of reducing saccharides from colloidal chitin by the *N*-acetyl-glucosamine-dinitrosalicylate method according to the method described by Monreal and Reese (1969). A reaction mixture containing 200 µl 0.5% chitin provided in citrate phosphate buffer (0.05 M, pH = 6.6) and 200 µl enzyme solution was incubated at 37°C for 1 h. Then 1 ml of dinitrosalicylic acid reagent (DNS) was added to reaction. The reaction was heated in boiling water for 5 min and was centrifuged at 6,000 rpm for 5 min and absorbance at 540 nm was measured. Protein concentration was determined according to Bradford (1976) with bovine serum albumin (Sigma) as the standard. Chitinase specific activity was calculated from the rate of enzyme activity divided per the mass of total protein. Enzyme assay was done to determine the most promising isolates for biological control of *G. rostochiensis* 24 and 96 h after fungal growth.

Data of the chitinase specific activity was subjected to analysis of variance (ANOVA) by software SAS, version 9.0 (Statistical Analysis System Institute Inc., USA) in a completely randomized design with three replicates.

Image processing. The field of digital image processing refers to processing digital images by means of a digital computer. Note that a digital image is composed of a finite number of elements, each of which has a particular location and value. Because digital images are displayed as a discrete set of intensities, the eye's ability to discriminate between different intensity levels is an important consideration in presenting image-processing results. Colour intensity of digital images has 256 levels that all of them could not be seen with human's eyes exactly (Gonzalez, Woods, 2002).

Chitinase enzyme activity causes the colour change in enzyme solution that is indistinguishable by human eyes. As a result, to analyze these phenomena, due to the ability of image processing to detect low colour change levels, this method was used. At first colour images were taken from isolates enzyme solution using a 12.1 MP digital camera Canon SX200is (Japan). Histogram equalization was done to reduce the noise of images using pre-processing algorithms (Gonzalez, Woods, 2002). Then by colour space transformation from RGB (red, green, blue) to HSV (hue, saturation and value) and $L^*a^*b^*$ using software *MATLAB*, version 2013b, the obtained colour components (R, G, B, H, S, V, L^* , a^* and b^*) were evaluated to find the best colour component that can confirm the chitinase activity colour changes.

Results and discussion

The 34 fungal isolates were identified according to morphological and molecular features and were classified in 11 genera (*Alternaria*, *Beauveria*, *Candida*, *Cylindrocarpon*, *Fusarium*, *Humicola*, *Lecanicillium*, *Paecilomyces*, *Plectosphaerella*, *Trichoderma* and *Uloclidium*) most of which belonged to *Fusarium* (Table 1). As a producer of a variety of chitinase enzymes, the filamentous fungus, *Fusarium* spp. can be an important means of biological control for *G. rostochiensis*. Ibrahim et al. (2009) isolated some fungi including *F. oxysporum*, *Aspergillus* spp., *Rhizoctonia solani*, *Paecilomyces lilacinus* and *Verticillium chlamydosporium* from *Meloidogyne incognita* eggs that *F. oxysporum* had the most biocontrol effect on the nematode. Also this result is similar to report of Ruanpanun et al. (2010). They isolated various fungal species from *M. incognita* most of which were *Penicillium* and *Fusarium*.

The results of chitinase specific activity showed significant differences not only among isolates but also between two days of fungal growth at the 0.001 level (Table 2). All of the 34 isolates had chitinase activity and chitinase specific activity in all of the isolates was higher in 96 h than 24 h. Among these isolates, the maximum chitinase specific activity belonged to isolate 154: *T. atroviridae* (0.56 U mg⁻¹) and isolate 113: *F. oxysporum*

Table 1. Results of identification of the fungal isolates according to morphological and molecular features

Isolate No.	Genus and species	Isolate No.	Genus and species
6	<i>Candida parapsilosis</i>	83	<i>Fusarium oxysporum</i>
8	<i>Fusarium equiseti</i>	93	<i>Candida</i> sp.
11	<i>Plectosphaerella</i> sp.	97	<i>Fusarium oxysporum</i>
12	<i>Fusarium oxysporum</i>	109	<i>Fusarium solani</i>
14	<i>Fusarium solani</i>	111	<i>Candida parapsilosis</i>
18	<i>Fusarium solani</i>	113	<i>Fusarium oxysporum</i>
19	<i>Candida parapsilosis</i>	123	<i>Fusarium oxysporum</i>
27	<i>Alternaria alternata</i>	129	<i>Fusarium equiseti</i>
30	<i>Alternaria alternata</i>	140	<i>Candida parapsilosis</i>
40	<i>Plectosphaerella cucumerina</i>	141	<i>Humicola grisea</i>
49	<i>Cylindrocarpon olidum</i>	144	<i>Candida parapsilosis</i>
52	<i>Uloclidium dauci</i>	145	<i>Fusarium oxysporum</i>
56	<i>Fusarium equiseti</i>	147	<i>Fusarium oxysporum</i>
62	<i>Fusarium solani</i>	151	<i>Beauveria bassiana</i>
63	<i>Fusarium solani</i>	152	<i>Lecanicillium muscarium</i>
66	<i>Fusarium equiseti</i>	153	<i>Paecilomyces</i> sp.
76	<i>Candida parapsilosis</i>	154	<i>Trichoderma atroviridae</i>

Table 2. Analysis of variance (mean square) of the chitinase specific activity in 24 and 96 h after fungal growth among 34 isolates

Sources of variance	df	Specific activity (24 h) U mg ⁻¹	Specific activity (96 h) U mg ⁻¹
Isolate	33	0.03932***	0.06152***
Error	68	0.00036	0.00889
CV%		8.10	14.49

CV – coefficient of variation, df – degree of freedom; *** – significant at 0.001 level

(1.02 U mg⁻¹) in 24 h and 96 h, respectively. Also isolate 6: *C. parapsilosis* (0.15 U mg⁻¹ and 0.40 U mg⁻¹) in 24 h and 96 h had the minimum chitinase specific activity (Figs. 1 and 2).

In this study, among the 34 isolates of various species of fungi, two well-known biocontrol genera *Trichoderma* and *Fusarium* had the highest enzymatic

activity. These fungi were previously reported by Sankaranarayanan et al. (2002), Safari Motlagh and Samimi (2013) and Saifullah and Khan (2014) as important biocontrol agents on *G. rostochiensis*. Also Safavi et al. (2010) have stated that several species of fungi including *Zoophthora radicans*, *Metarhizium anisopliae*, *Fusarium* sp. and *B. bassiana* are potent biocontrol agents of plant

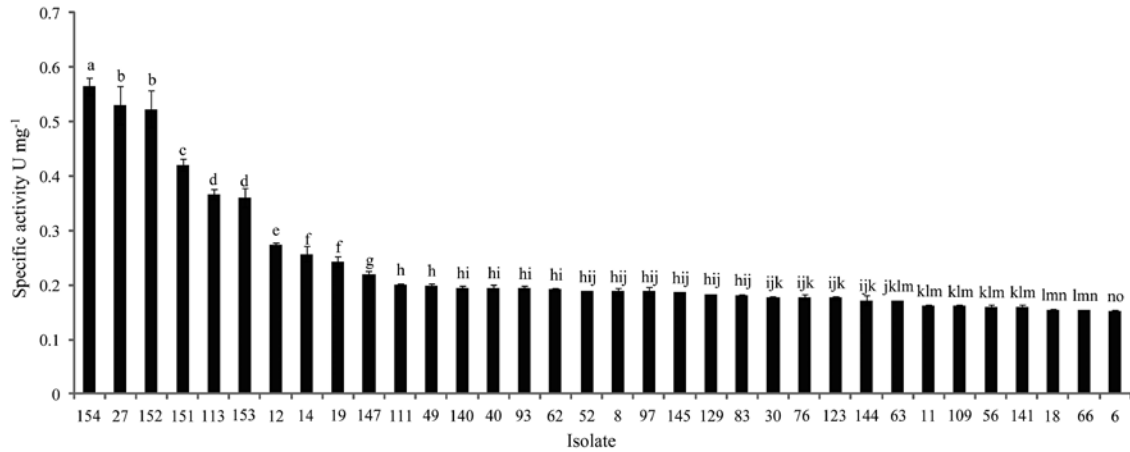


Figure 1. Chitinase specific activity of culture filtrates of 34 isolates in the minimal synthetic medium (MSM) in 24 h

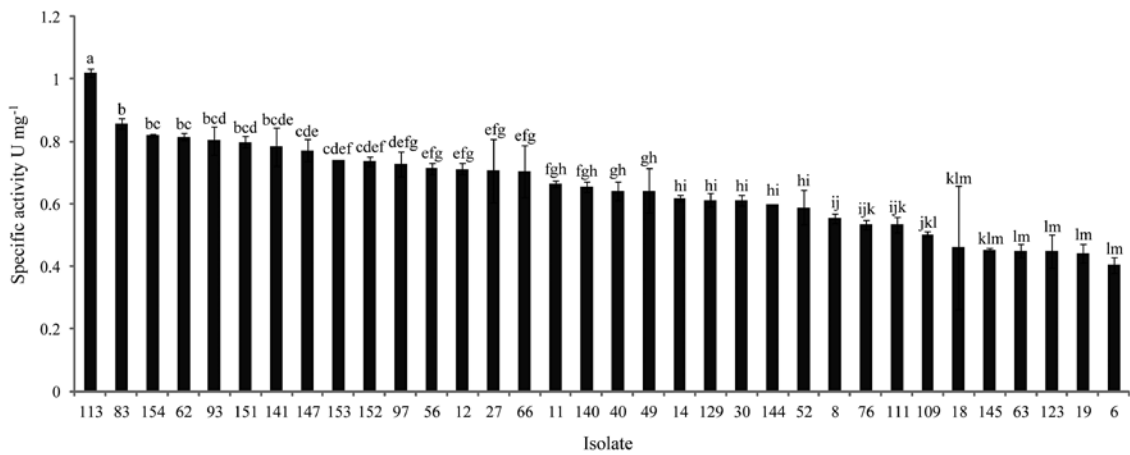


Figure 2. Chitinase specific activity of culture filtrates of 34 isolates in the minimal synthetic medium (MSM) in 96 h

pathogenic fungi and arthropods that can infect different insect species in screen house or field conditions.

The results of the linear regression of enzyme assay for two days, 24 h and 96 h, showed there is a good correlation between two conditions and the coefficient of determination was $R^2 = 0.7$ (Fig. 3). This coefficient showed that biological activity does not necessarily follow a linear relationship. But we can see relative changes between chitinase specific activities in 96 h than 24 h in 34 isolates had uptrend (Fig. 4).

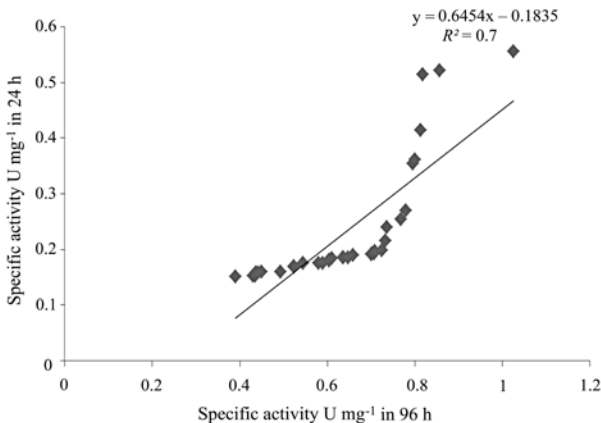


Figure 3. Regression relationship between chitinase specific activity in 24 h and 96 h in 34 isolates

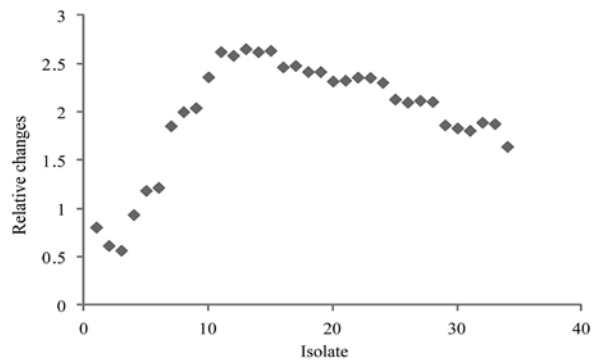


Figure 4. Relative changes between chitinase specific activity in 24 h and 96 h in 34 isolates

Image processing technique for colorimetry of the samples confirmed the results of chitinase activity. Results showed that among all 9 colour components (RGB, HSV and $L^*a^*b^*$), L^* had maximum correlation coefficient ($R^2 = 0.97$) with enzyme activity (Fig. 5). This component shows intensity of colour between 0 (black) to 100 (white) and confirms that intensity colour of isolates with high chitinase activity is low. This means that increase of chitinase activity causes darkening of enzyme solution colour.

Results showed that isolate 113 (*F. oxysporum*) with maximum chitinase activity equal to 1.02 U mg^{-1}

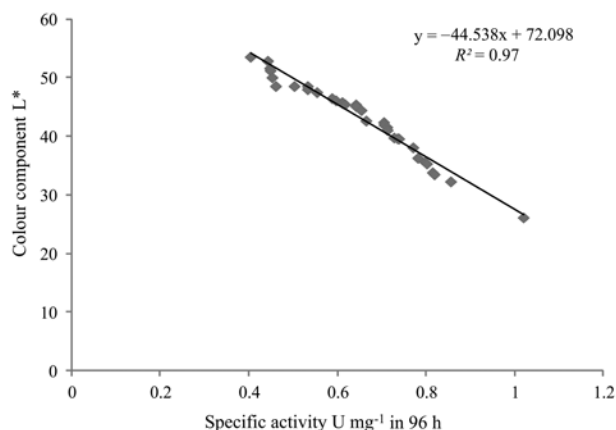


Figure 5. Regression relationship between chitinase specific activity in 96 h and colour component L* (image processing) in 34 isolates

had minimum L* component value and isolate 6 (*C. parapsilosis* with minimum chitinase activity equal to 0.40 U mg⁻¹) had maximum L* component value.

Conclusions

1. Based on morphological and molecular features, the 34 isolated fungi from *Globodera rostochiensis* were classified in 11 genera most of which belonged to *Fusarium*. As a producer of a variety of chitinase enzymes, the filamentous fungus, *Fusarium* spp., can be an important means of biological control for this nematode.

2. According to chitinase specific activity assay in 34 isolates of various species of fungi obtained from infected golden potato cyst nematode over two days of fungal growth (24 h and 96 h), isolates 154: *Trichoderma atroviridae* and 113: *Fusarium oxysporum* in 24 and 96 h had high activity, respectively. Also isolate 6: *Candida parapsilosis* in these two days had the minimum chitinase specific activity.

3. Use of image processing technique confirmed that there is a good relationship between enzyme activity and colour intensity of enzyme solution in 96 h. The colour of solution for isolates with high enzyme activity is darker than for isolates with low enzyme activity.

4. In general, the successful isolates, 154: *T. atroviridae* and 113: *F. oxysporum* in chitinase enzyme activity in the present study can be used for mass production of this enzyme which is supposed to be used in commercial formulation and also will be able to control golden potato cyst nematode.

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Fermento chitinazės aktyvumo įvertinimas grybų izoliatuose, gautuose iš auksinio bulvinio nematodo (*Globodera rostochiensis*)

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Santrauka

Nematodo kiaušinėlio apvalkalas rodo, kad proteazės ir chitinazės fermentai yra būtini užsikrėtimui nematofagiais grybais, kurių dauguma produkuoja šiuos fermentus. Tyrimo metu siekta iširti chitinazės veiklą grybų 34 rūšių izoliatuose, gautuose iš auksinio bulvinio nematodo (*Globodera rostochiensis*) per dvi grybo augimo dienas (24 ir 96 val.). Chitinazės specifinis aktyvumas nustatytas matuojant redukuojančių sacharidų išskyrimą iš koloidinio chitino, taikant *N*-acetil-gliucozamino-dinitrosalicilato (540 nm) metodą. Siekiant patvirtinti chitinazės aktyvumą taikytas kolorimetrijos metodas, paremtas vaizdo apdorojimu nustatant spalvų pokyčius minimalioje sintetinėje terpėje. Tirti 34 izoliatai buvo identifikuoti remiantis morfologiniais ir molekuliniais požymiais, taip pat ir ribosominės DNR vidiniais transkribuotais tarpiklio regionais. Chitinazės specifinio aktyvumo matavimo rezultatai parodė, kad visuose tirtuose izoliatuose jis buvo didesnis po 96 val. nei po 24 val. Maksimalus ir minimalus chitinazės specifinis aktyvumas po 24 val. nustatytas izoliatuose 154 (0,56 U mg⁻¹) ir 6 (0,15 U mg⁻¹), po 96 val. – izoliatuose 113 (1,02 U mg⁻¹) ir 6 (0,40 U mg⁻¹). Kolorimetrijos metodo rezultatai patvirtino, kad fermentų veikla susijusi su spalvos pokyčiais. Tirti grybų 34 izoliatai buvo suklasifikuoti į 11 genčių, iš jų dauguma priklausė *Fusarium* rūšiai. Du didžiausio chitinazės fermento aktyvumo izoliatai – 113 (*Fusarium oxysporum*) ir 154 (*Trichoderma atroviridae*) – gali kontroliuoti auksinį bulvinį nematodą ir būti panaudoti chitinazės fermento, skirto komerciniams preparatams, gamybai.

Reikšminiai žodžiai: bulvės, chitinazė, *Globodera rostochiensis*, grybai.