



Research Article

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Bacteria *Bacillus velezensis* BIM B-439 D as the Basis of Biopreparation Betaprotectin to Control Root Rots of Cucumber



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Abstract

Molecular-genetic and functional characterization of bacterial strain *B. velezensis* BIM B-439 D as the basis of biopesticide Betaprotectin was carried out. Bacterial genome contained 10 loci determining synthesis of polyfunctional enzyme complexes responsible for production of several antimicrobial compounds and lacking genes of drug resistance and pathogenicity. It was shown that the microbial culture suppressed development of a broad spectrum of fungal and bacterial pathogens infecting agricultural crops, raised germination power and sprouting rate of cucumber seeds, displayed high survival rate in natural soil microbiocenoses accounting for elevated phytoprotective and growth-stimulating potential and attractive application prospects as the active principle of biopesticide Betaprotectin. Prolonged application record of biopesticide throughout 10-year vegetation period to promote greenhouse cucumber culture provided for stable phytosanitary control of root rots, caused favorable effect on plant status and increased yields of vegetable products.

Keywords: Bacteria *Bacillus velezensis* BIM B-439 D; Greenhouse cucumber; Root rots; Biopesticide betaprotectin

Introduction

Root rots of greenhouse cucumber are widely distributed and deleterious diseases affecting this crop in all cultivation areas. Typical pathogens are represented by soil-borne fungi *Pythium debaryanum*, *Rhizoctonia solani*, *Ascohyta cucumis*, genus *Fusarium* [1-3]. Pathogenic fungi apart from deterioration of rhizosphere, release toxic metabolites inhibiting growth and development of cucumber plants [4]. Strains of microbial antagonists and the derived biopreparations are successfully used to control population density of many phytopathogenic species, including causal agents of cucumber root rots [5-7]. Of special interest are sporulating bacteria of genus *Bacillus* exerting antagonistic impact on microbial pathogens and displaying phyto-regulating activity, i.e. the ability to stimulate growth and development of cultivars, to induce their resistance to damaging factors, to initiate conducive influence on biological activity and make-up of soil microbiota. Such favorable effects

are determined by the capacity of numerous bacilli to produce antibiotics, phytohormones, hydrolytic enzymes contributing to utilization of diverse substrates and to generate agents involved in mineralization of phosphororganic compounds resulting in plant-digestible ion PO_4^{2-} . Growth-promoting activity of the bacilli may be also accounted for by the ability to synthesize vitamins and immunomodulators enhancing plant resistance potential [8-15]. Thus, the increased biological activity and phytostimulating function lay the basis for wide application prospects of *Bacillus* cultures in biotechnological sector as the promising sources of bioactive agents and the key ingredients of efficient biopesticides. Nowadays spore-forming bacteria of genus *Bacillus* serve as an active principle of a broad spectrum of biopreparations applied to curb root rots of cucumber crops around the world, e.g. Biosubtilin (Biotech International Ltd., India), RhizoVital (ABiTEP GmbH, Germany), Serenade (AgroGuest, USA), Sonata (AgroGuest, USA), etc [14-19]. To improve competitive edge of

phytoprotective biopreparations it is vital to incorporate into their formulas superactive microbial strains capable to carry out long-term stable control of phytopathogen development. Investigation of genetic and physiological properties of the cultures enables to evaluate and predict biotechnological potential thereof. Aim of this research was characterization of the studied strain *B. velezensis* BIM B-439 D and assessment of long-term application efficiency of the derived biopesticide Betaprotectin for control of root rots attacking greenhouse cucumber crop.

Objects and Methods

Object of study - bacterial strain *Bacillus velezensis* BIM B-439 D distinguished by high antagonistic activity toward a broad spectrum of phytopathogenic fungi and bacteria. Test cultures to evaluate antimicrobial activity: fungi *Penicillium expansum*, *Alternaria tenuis*, *Fusarium graminearum*, *Fusarium oxysporum*, *Fusarium culmorum*, *Fusarium avenaceum*, *Botrytis cinerea*, *Phoma betae* and phytopathogenic bacteria *Pseudomonas syringae* var. *lachrymans*, *Ps. fluorescens*, *Xanthomonas campestris* subsp. *campestris*, *Pectobacterium carotovorum* subsp. *carotovorum*, *Clavibacter michiganensis* subsp. *michiganensis*, *Agrobacterium tumefaciens*, phytopathogenic deposited in Belarusian collection of non-pathogenic microorganisms, Institute of Microbiology, National Academy of Sciences of Belarus. Fungi cultures were maintained on agar nutrient media (MPA, LB, HBA, 2 % agar), the fungi were grown on PDA medium (2 % agar). Submerged culture of bacterial strain *B. velezensis* BIM B-439 D was carried out in liquid modified Meynell medium [19] with molasses (30 g/l) in Erlenmeyer flasks on the shaker-incubator (200 rpm) at temperature 30±2 °C during 48h.

Liquid submerged cultures of phytopathogenic bacteria were incubated in Hottinger broth in Erlenmeyer flasks on the shaker (180-200 rpm) at temperature 30±2 °C for 24h, phytopathogenic fungi were cultured in potato-glucose broth in Erlenmeyer flasks on the shaker-incubator (140 rpm) at temperature 24±2°C during 36-48h. Cell and spore titers of liquid bacterial culture *B. velezensis* BIM B-439 D were determined by Koch method [20]. To facilitate spore counts prior to inoculation on agar medium thermal treatment of bacterial dilutions was conducted at 80 °C for 10 min. Antagonistic activity of cultural liquid of strain *B. velezensis* BIM B-439 D was estimated by method of delayed antagonism (wells technique). Sterile water was used in control variants [21]. To evaluate enzymatic activity of bacterial cultures standard qualitative methods were engaged, taking into account utilization degree of specific substrates-inducers: starch for amylolytic activity assays, casein for proteolytic activity, xylan for xylanase measurements, Na⁺-carboxymethylcellulose for cellulase assays, colloid chitin – for chitinolytic activity [22].

Growth-stimulating activity of cultural liquid of bacteria *B. velezensis* BIM B-439 D was investigated on seeds and seedlings of tomato (hybrid variety Torrero F1) and cucumber (hybrid variety Rodnichok). Assessment of seed quality and germination

was performed according to directions of State standard 12038-84. Sterile water was taken as the control. Survival of bacteria *B. velezensis* BIM B-439 D in sod-podzol soil (pH 6.0-6.2) was judged by the results of model experiments (upon addition of 48h bacterial cultural liquid bearing spores in minimal concentration 1x10⁷ CFU/g soil). The titer of bacterial spores in the samples was counted by Koch method after heating soil suspensions. Soil moisture content was kept throughout the experiment on the 50-60 % level. Concentrations of micromycetes and aerobic bacterial microbiota in soil were determined by method of finite dilutions [20]. DNA libraries for full-genome sequencing were prepared with Nextera XT («Illumina», USA). Nucleotide sequences were identified using MiSeq («Illumina», USA) equipped with MiSeq Reagent Kit v3 (USA). Data quality was verified with FastQC [23]. Low-resolution reads were filtered with Trimmomatic-0.36 [24-27]. Readings were assembled with SPAdes-3.11.1 [28]. Preliminary genome annotation was realized with the aid of RAST 2.0 and PROKKA 1.12 package [26,27] Individual genetic loci were compared using web-resource antiSMASH [28]. Mauve (v. 20150226) software was applied for comparison of genomes [29]. Average Nucleotide Identity (ANI) was computed with web-service JSpeciesWS [30]. The search for genes of antibiotic resistance and pathogenicity relied on resources ResFinder 4.1 and PathogenFinder 1.1 provided by the web-server of the Center of Genome Epidemiology, Technical University, Denmark [31].

Large-scale efficiency trials of biopesticide Betaprotectin in regard to cucumber pathogenic agents were performed in greenhouse complex of Grodno vegetable factory, Belarus growing hybrid variety Mirabella F1. The technology of cucumber cultivation is standard for this agricultural zone. Betaprotectin efficiency tests were conducted on model vegetable culture against natural infection background. The dosage of Betaprotectin expense for treatment of greenhouse cucumber equaled 50 l/ha. The vegetables were watered with 2 % working solution of biopesticide via installed irrigation system, 4-5 times during vegetation season with fortnight intervals. The unit dose of Betaprotectin working solution - 250 ml per plant. The treatment periods: for the first crop rotation cycle – February to May, for the second crop rotation cycle - July to October. Phytosanitary monitoring of greenhouse cucumber was accomplished throughout the whole period starting from initial supply of Betaprotectin until the end of vegetation season. Spread and severity rate of the diseases were controlled 10 days after plant treatment by checking 10 plants evenly distributed across field plot diameter in each sample repetition series. By the end of vegetation period root system and adjacent stem regions of plants were inspected. Economic efficiency of biopesticide was estimated as the amount of harvested cucumbers saved versus the control.

Results and Discussion

Sporulating bacterial strain *Bacillus velezensis* BIM B-439 D was isolated from sod-podzol soil sampled in Belarus and its

bacterial genome was sequenced and deposited in database of GenBank NCBI under registration number CP032144 [16-18]. Genome of strain *B. velezensis* BIM B-439 D is represented by circular chromosome sized 3 978 954 b.p. containing 46.50 % of G/C pairs (Figure 1). The deciphered chromosome includes 3969 predicted genes, including 3769 protein-encoding sequences, 27 rRNA genes, 86 tRNA genes, 82 pseudogenes, 9 genetic loci responsible for synthesis of 27 rRNA molecules and 86 genes determining synthesis of tRNA. Analysis of *B. velezensis* BIM

B-439 D genome revealed 10 loci common for genus *Bacillus* (the total length 336520 b.p.) defining synthesis of multifunctional enzyme complexes catalyzing production of several antimicrobial compounds. Using ResFinder 4.1 and PathogenFinder 1.1 resources, it was found that genome of strain *B. velezensis* BIM B-439 D lacked antibiotic resistance genes and islets of pathogenicity, accounting for their elevated phytoprotective potential and attractive application prospects as the basis of biopesticides.

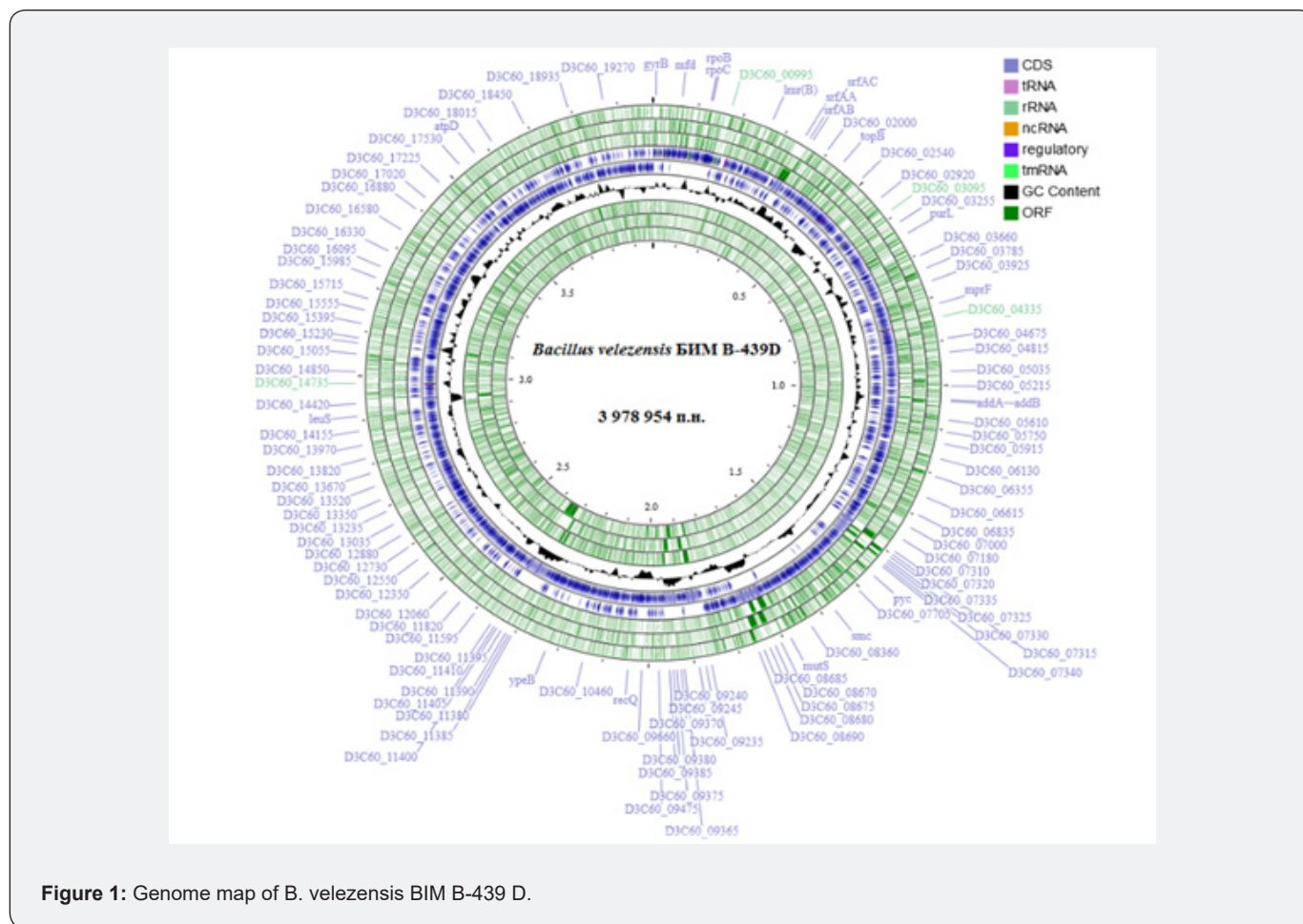


Figure 1: Genome map of *B. velezensis* BIM B-439 D.

Biotechnological potential of strain *B. velezensis* BIM B-439 D was analyzed in laboratory and large-scale production experiments. Wells technique demonstrated (Table 1) that bacteria *B. velezensis* BIM B-439 D displayed inhibitory action toward a broad range of phytopathogens causing most widespread diseases of agricultural crops, including root rots of cucumber. It was observed that swelling and deformation of hyphae occurred in phytopathogenic fungi exposed to active metabolites of tested bacteria, ultimately resulting in their complete lysis (Figure 2). Bacterial strain *B. velezensis* BIM B-439 D shows the ability to utilize casein, starch, cellulose, colloid chitin and xylan, evidencing generation by bacterial cells of variegated

enzyme complex: protease, amylase, β -endoglucanase, chitinase, xylanase. Production of a wide spectrum of hydrolytic enzymes may act as an additional antagonistic factor of bacteria *B. velezensis* BIM B-439 D augmenting their competitive efficiency in rivalry with pathogens from soil microbiota. To assess competitive and adaptive potential of strain *B. velezensis* BIM B-439 D, survival of the bacteria was traced in natural microbial cenoses. In the course of laboratory experiments bacterial cultural liquid was supplied into soil containing native strains of pathogenic and saprophytic micromycetes (cell titer at least $6.0 \cdot 10^4$ CFU/g) and bacteria (the minimal titer $3.0 \cdot 10^3$ CFU/g). Experimental results established that upon 40 days $\geq 90\%$ of viable spores of *B. velezensis* BIM

B-439 D survived in soil samples (Figure 3), evidencing enhanced adaptation rate and competitive ability of the culture introduced into natural soil microbiocenoses. Phytostimulating activity of bacteria *B. velezensis* BIM B-439 D was also evaluated. The effect of liquid bacterial culture on seed germination and development of cucumber seedlings was tested. The results have shown that seed

treatment with bacterial cultural liquid significantly upgraded germination efficiency. The best results were achieved under the 20-min impact of 5 % solution of cultural liquid of *B. velezensis* BIM B-439 D: germination power of cucumber seeds soaked in 5% solution of bacterial liquid culture rose by 16 % on the average, sprouting rate by 14 % (Figure 4).

Table 1: Antagonistic activity of bacteria *B. velezensis* BIM B-439 D against pathogens of vegetable and forage crops (estimated by wells technique).

Phytopathogen	Diameter of pathogen growth inhibition zone, mm
<i>Pseudomonas syringae</i> var. <i>lachrymans</i>	37±0.8
<i>Pectobacterium carotovorum</i>	24±0.9
<i>Xanthomonas campestris</i> subsp. <i>campestris</i>	24±0.8
<i>Clavibacter michiganensis</i> subsp. <i>michiganensis</i>	22±0.9
<i>Agrobacterium tumefaciens</i>	23±0.6
<i>Fusarium graminearum</i>	30±1.0
<i>Fusarium oxysporum</i>	29±0.5
<i>Fusarium culmorum</i>	24±0.7
<i>Fusarium avenaceum</i>	23±1.0
<i>Botrytis cinerea</i>	27±0.9
<i>Penicillium expansum</i>	23±0.8
<i>Phoma betae</i>	31±0.6
<i>Phytium aphanidermatum</i>	20±0.9

Owing to the pronounced phytoprotective and growth-stimulating activity bacterial strain *B. velezensis* BIM B-439 D was applied as the basis of biopesticide Betaprotectin intended for control of microbial infections in cucumber and tomato cultivars. Efficiency trials of biopesticide Betaprotectin (expense 50 l/ha) on cucumber seedlings grown in nurseries in small-scale hydroponic culture during the first crop rotation cycle diminished distribution of the pathologies by 25.6%, whereas the intensity of root rot damage fell by 20.9%. The level of biological efficiency of applied biopesticide reached 40.3% (Table 2). According to the Table 3, the similar relationship was recorded likewise during the second rotation cycle of cucumber culture. Application of biopesticide Betaprotectin curtailed the average spread of root rots by 14.8%, and the extent of root damage was 25.8% lower than in the non-treated control variant. Biological efficiency of Betaprotectin reached the average mark 41.4%. Betaprotectin supply enabled

to raise harvests of cucumbers during the first crop rotation cycle from 28.5 kg/m² to 31.0 kg/m². The saved vegetable yield equaled 2.5 kg/m², or 8.3 % up the control (Table 4). Application of Betaprotectin during the second crop rotation cycle resulted in extra 0.93 kg/m² yield of cucumber, gaining additional 7.1 % above the control harvest (Table 5). During 2012-2021 regular supply of biopesticide Betaprotectin 4-5 times in the course of cucumber vegetation season, with fortnight intervals in each crop rotation cycle provided for stable phytosanitary control of root rot diseases (Table 6). Extent of cucumber root damage within the framework of Betaprotectin testing program ranged from 36.5 % in 2012 to 27.5 % in 2015. In contrast, intensity of control plant lesions constituted 53.8-60.4 %. The improved phytosanitary status considerably upgraded cucumber yields varying from 44.0 kg/m² in 2013 to 47.6 kg/m² in 2015 as compared with the control values 40.4-42.5 kg/m².

Table 2: Effect of liquid Betaprotectin on spread and pathological damage of root rots attacking greenhouse cucumber (the first crop rotation cycle, 2012-2013).

Experimental variant	Spread of the disease, %	Plant infection, %	Biological efficiency %
Control (untreated)	76.3	51.9	-
liquid Betaprotectin	50.7	31	40.3

Table 3: Effect of liquid Betaprotectin on spread and pathological damage induced by root rot pathogens of greenhouse cucumber (the second crop rotation, 2012-2013).

Experimental variant	Spread of the disease, %	Plant infection, %	Biological efficiency %
Control (untreated)	86.9	62.4	-
liquid Betaprotectin	72.1	36.6	41.4

Table 4: Effect of biopesticide Betaprotectin on productivity of greenhouse cucumber (the first crop rotation cycle 2012-2013).

Experimental variant	Harvest collected from 1 m ² area, kg	Harvest saved on the date of checking:	
		kg/m ²	% of the control
Control (untreated)	28.5	-	-
Liquid Betaprotectin	31	2.5	8.3

Table 5: Effect of biopreparation Betaprotectin on productivity of greenhouse cucumber (the second crop rotation cycle 2012-2013).

Experimental variant	Harvest collected from 1 m ² area, kg	Harvest saved on the date of checking:	
		kg/m ²	% of the control
Control (untreated)	13	-	-
Liquid Betaprotectin	13.92	0.93	7.1

Table 6: Application efficiency of biopesticide Betaprotectin against root rots of cucumber at production facilities of Grodno vegetable factory (the mean of two crop rotation cycles).

Year of trials	Progression of root rots by the end of vegetation season (the mean value from 2 crop rotation cycles), %	Productivity of cucumbers, kg/m ²
2012	36.5	45.7
2013	31.1	44
2014	30.1	45.3
2015	27.5	47.6
2016	28.1	47.1
2017	31.4	44.3
2018	32.3	44.7
2021	31.2	45.3

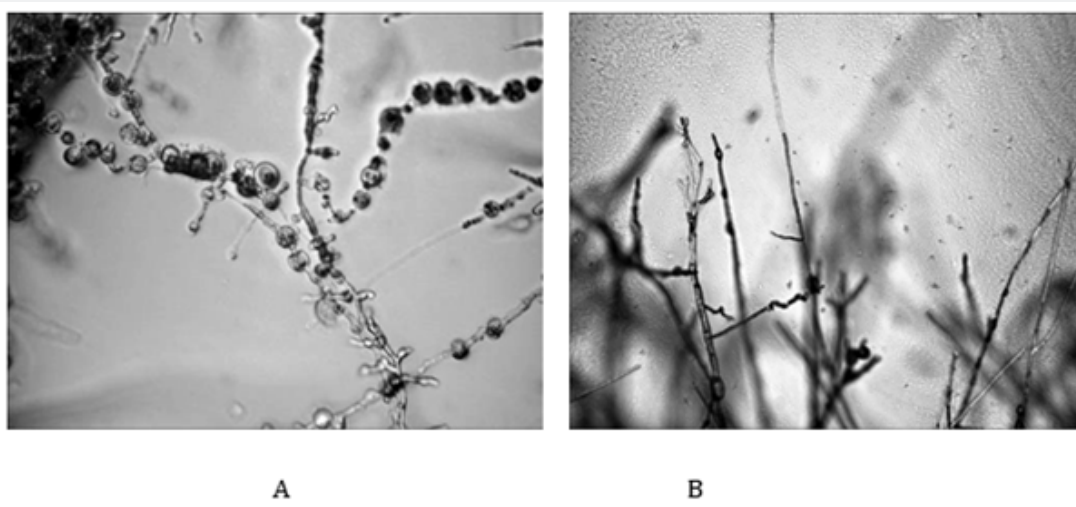


Figure 2: Effect of cultural liquid of bacteria *B. velezensis* BIM B-439 D on mycelial hyphae of *F. oxysporum* (microphoto): A - experiment, B – control.

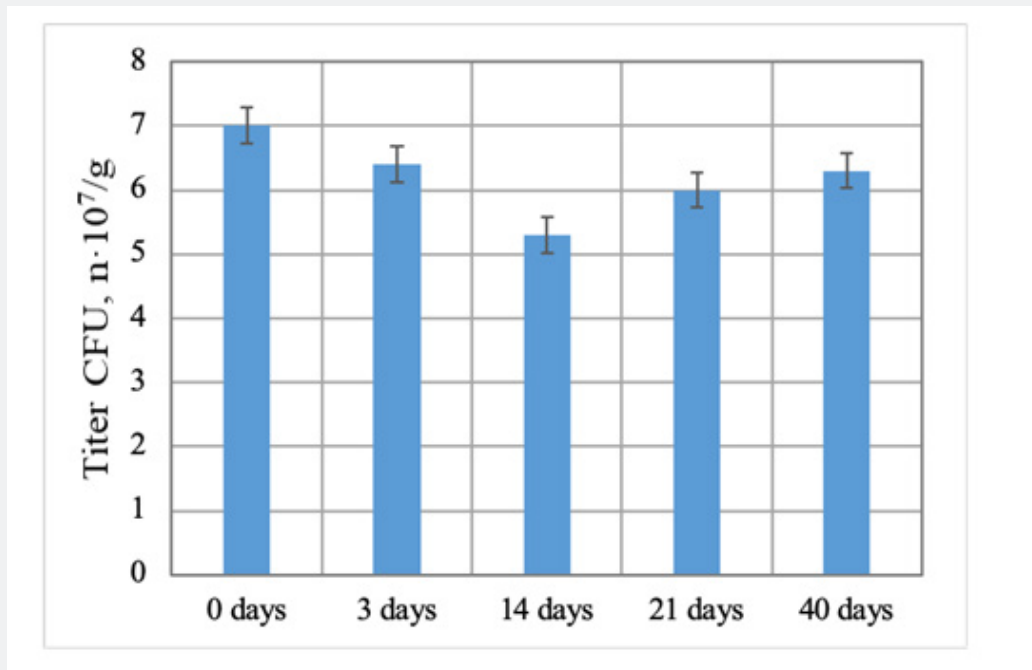


Figure 3: Survival rate of bacteria *B. velezensis* BIM B-439 D in soil.

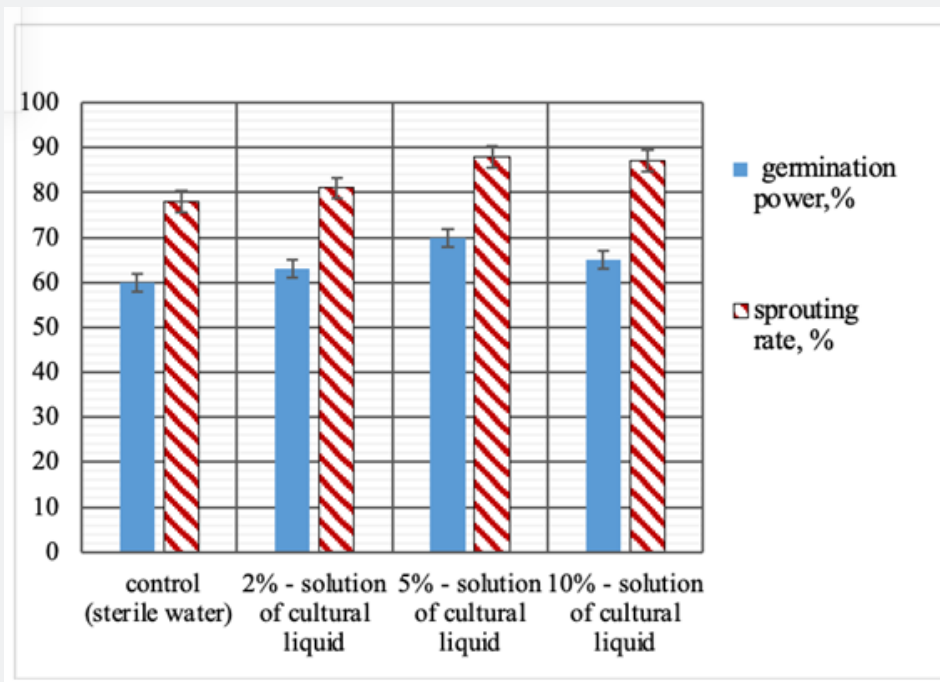


Figure 4: Efficiency of cucumber seed germination following treatment with solutions of bacterial cultural liquid.

Conclusion

Summing up, the analysis of genetic and functional characteristics of bacterial strain *B. velezensis* BIM B-439

D indicates that its promising biotechnological potential is appropriate for use in microbiological industry as the active principle of biopesticide Betaprotectin. Prolonged 10-year

application record of biopesticide Betaprotectin in greenhouse practice secures stable phytosanitary control of root rot diseases, causes beneficial effect on cucumber seedlings and promotes output of top-grade vegetable products.

References

1. Karbozova RD (1993) Root rot of greenhouse cucumber and methods to diminish its infectivity. Synopsis of Ph.D thesis in agrosiences, specialty 06.01.11-plant protection. Almata, p. 21.
2. Grishechkina LD (2003) Diagnostics of vegetable crop diseases in greenhouse culture. *Journal of Phytoprotection and quarantine* 3: 45-50.
3. Rudakov OL, Rudakov VO (2016) Modification of methods to diagnose root rots of cucumber. *Journal of Plant protection* 4: 21-23.
4. Woloshuk CP (2013) Aflatoxins, fumonisins, and trichothecenes: a convergence of knowledge. *Journal of FEMS Microbiol. Rev.* 37(1): 94-109.
5. Gravel V, Martinez C, Antoun H, Tweddell RJ (2006) Control of greenhouse tomato root rot [*Pythium ultimum*] in hydroponic systems, using plant-growth-promoting microorganisms. *Can. Journal Plant Pathol* 28(3): 475-483.
6. Bosmans L, Buriijn ID, Gerards S, Rob M, Lore V, et al. (2017) Potential for biocontrol of hairy root disease by a *Paenibacillus* clade 8: 447.
7. Punja ZK, Yip R (2003) Biological control of damping-off and root rot caused by *Pythium aphanidermatum* on greenhouse cucumbers. *Can. Journal Plant Pathol* 25(4): 411-417.
8. Harwood CR, Mouillon JM, Pohl S, Arnau J (2018) Secondary metabolite production and the safety of industrially important members of the *Bacillus subtilis* group. *FEMS Microbiol. Rev* 42(6): 721-738.
9. Saharan BS, Nehra V (2011) Plant Growth Promoting Rhizobacteria: A Critical Review. *Journal of Life Sci Med Res* 21: 1-30.
10. Dragovoz VI (2014) Exometabolites of strain *Bacillus amyloliquefaciens* IMB B-7100 defining its phyto-stimulating activity. *Journal of Plant physiology and genetics* 46(6): 516-524.
11. Morgun VV, Kots YS, Kyrychenko EV (2009) Growth promoting rhizobacteria and their use in practice. *Journal of Fiziol. Biokhim. Kul't Rast* 41(3): 187-207.
12. Zabokritskiy NA (2015) The biologically active substances produced by probiotic microorganisms of the genera *Bacillus* and *Lactobacillus*. *Journal of Scient Art* 17(3): 8-19.
13. Bhushan BJ (2000) Production and characterization of a ther mostable chitinase from a new alkalophilic *Bacillus* sp. BG-11. *Journal of Appl Microbiol* 88(5): 800-808.
14. Abbas A, Khan SU, Wasim UK, Saleh TA, Hafeez M, et al. (2019) Antagonist effects of strains of *Bacillus* spp. against *Rhizoctonia solani* for their protection against several plant diseases: Alternatives to chemical pesticide. *C R Biologies*: 342(5-6): 124-135.
15. Aloo BN, Makumba BA, Mbega ER (2019) The potential of *Bacilli* rhizobacteria for sustainable crop production and environmental sustainability *Journal of Microbiological Research* 219: 26-39.
16. Berezhnaya AV, Evdokimova OV, Valentovich LN, Sverchkova NV (2019) Molecular-genetic analysis of chromosome loci determining antimicrobial properties of bacteria *B. velezensis* BIM B-439 D. *Journal of Prikl. biochem. Microbial* 55(4): 366-377.
17. Berezhnaya AV (2015) Molecular-genetic analysis of determinants encoding synthesis of antimicrobial metabolites by bacteria of genus *Bacillus* abstr. IX Int Sci conf, Belarus.
18. Titok MA, Valentovich LN, Berezhnaya AV, Kolomiets EI (2018) Genome analysis of bacterial strain *Bacillus amyloliquefaciens* BIM B-439 D. *Doklady Nat Acad Nauk Belarusi* 62(5): 592-600.
19. Meynell D (1967) Experimental microbiology. *Journal of Mir Publ*, pp.347.
20. Aseyeva IV (1984) Methods of soil microbiology and biochemistry. pp. 304.
21. Segy J (1983) Methods of soil microbiology Moscow. *Kolos Publ*, pp. 296.
22. Rukhlyadeva AP (1981) Methods to assay activity of hydrolytic enzymes. *Legpishcheprom Publ*, pp. 288.
23. Babraham Bioinformatics - FastQC a quality control tool for high throughput sequence data.
24. Bolger AM, Lohse M, Usadel B (2014) Trimmomatic: A flexible trimmer for Illumina sequence data. *Journal of Bioinforma. Oxf Engl* 30(15): 2114-2120.
25. SPAdes 3.11.1 Manual.
26. Aziz RK, Bartels D, Best AA, DeJongh M, Disz T (2008) The RAST Server: rapid annotations using subsystems technology. *BMC Genomics* 9: 75.
27. Seemann T (2014) Prokka: Rapid prokaryotic genome annotation. *Journal of Bioinformatics* 30(14): 2068-2069.
28. antiSMASH.
29. Darling AE, Mau B, Perna NT (2010) Progressive Mauve: Multiple Genome Alignment with Gene Gain, Loss and Rearrangement. *PLOS ONE* 5(6): e111147.
30. Richter M, Móra RR, Glöckner FO, Peplies J (2016) SpeciesWS: a web server for prokaryotic species circumscription based on pairwise genome comparison. *Journal of Bioinformatics* 32(6): 929-931.
31. CGE Server.



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