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Bacterial strain *Pseudomonas avellanae* 6CH2 with anti-Fusarium activity in mitigation of herbicidal stress in wheat plants

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Abstract

The aim of this work was to study a new strain of microorganisms resistant to herbicides of different chemical structure and its impact on herbicidal stress in wheat (*Triticum aestivum* L.). Herbicides with synthetic auxins 2,4-D (2-ethylhexyl ether), dicamba (Octapon, Chistalan) and metsulfuron-methyl (Nanomet) had a phytotoxic (stress) effect on wheat plants, decreasing their weight up to 11%, reducing the amount of chlorophylls up to 10% and increasing the proline quantity by 2.5-5.7 times in leaves. The *Pseudomonas avellanae* strain 6CH2 was isolated from chemical factory soil and had a number of useful properties: suppression of phytopathogenic *Fusarium*, herbicide-resistance, synthesis auxins and molecular nitrogen fixation. Its special features did not disappear under the influence of herbicides. Because of these properties, *P. avellanae* strain 6CH2 had an anti-stress effect on wheat plants if they were jointly treated with herbicides and bacteria. Spraying with herbicides increased the concentration of proline in wheat leaves by 1.7-2.8 times, while bacterial addition made it at least the same as in the control group. Inoculation with bacteria *P. avellanae* strain 6CH2 increased the total chlorophyll amount by 1.19-1.26 times against the background of herbicidal stress. Thus, bacteria *P. avellanae* strain 6CH2 can be used as an anti-stress agent when spraying wheat crops with herbicides.

Keywords: Pseudomonas, Wheat, Anti-fungal activity, Anti-stress effect, Chlorophyll, Proline.

1. Introduction

When cultivating agricultural crops, traditionally, great attention is paid to combating weed flora. Along with soil tillage and crop rotations, chemical herbicides are the main methods for controlling unwanted vegetation. They are used more often than other plant protection products (Aktar et al., 2009). Such widespread and intensive use provokes the spread of resistant weeds, which in turn prompts agricultural producers to increase the doses of herbicides and combine several active substances in order to improve the efficacy of chemical herbicides (Zargar et al., 2019). The efforts to control perennial and persistent weeds can also inhibit crop plants (Light et al., 2005; Kumar and Singh, 2010) and results in residual effects next year (Su et al., 2018). The current situation will remain until the next generation of products aiming with a different mechanism of action compared to currently produced herbicides.

Therefore, the search for means to reduce herbicidal stress in agricultural crops becomes relevant. The use of specialized strains of microorganisms for this purpose began to be considered only recently and is reflected in a few publications. Bourahla *et al.* (2018) ascertained the ability of the *Pseudomonas putida* strain to improve physiological and biochemical parameters (chlorophyll, carotenoids, malondialdehyde, enzyme activity) and reduce the manifestations of oxidative stress in durum

wheat seedlings against the background of 10⁴ norfluazone in hydroponic culture. Burkholderia cepacia strain mitigated toxicity, increased the size, dry matter, the ability to form nodules, the content of nutrients in seeds of chickpea plants, reduced the levels of proline and malondialdehyde if the amount of glyphosate in the soil was 4.332 mg/kg (Shahid and Khan, 2018). Strain Mesorhizobium in the presence of cladinophope (up to 1.2 mg/kg of soil) increased biomass, nodule and leghemoglobin content, nutrient uptake, seed yield and grain protein in chickpeas (Ahemad and Khan, 2010b). Quizalafop-p-ethyl- and clodinafop-tolerant Rhizobium isolate increased biomass, symbiotic properties, nutrients uptake and seed yield of lentil (Ahemad and Khan, 2010a) and pendimethalin resistant cereals growth stimulating Azotobacter salinestris (Chennappa et al., 2018) were also described.

The described bacteria, which can increase plant resistance to herbicidal stress, belong to different genera: *Pseudomonas, Rhizobium, Mesorhizobium, Bacillus, Azotobacter.* All of them are rhizosphere or endophytic microorganisms that can actively interact with the plant. The authors of the research note that they have such properties as nitrogen-fixing and phosphatemobilizing activity, synthesis of auxins, exopolysaccharides, 1-aminocyclopropane-1-carboxylate deaminase, and siderophores.

It is believed that bacteria associated with plants can enhance the non-specific stress resistance (including

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resistance to herbicides), inducing universal protective reactions in the plants (Tétard-Jones and Edwards, 2015). Therefore, it is possible to obtain a single strain of microorganisms that favorably affects plants against a wide range of different herbicides. The publications we found do not contain data on whether a single bacterium can increase the plant's resistance to herbicides of different structures or to a combination of two herbicides. Therefore, experimental testing of this possibility is actual.

The aim of our work was to study a new strain of microorganisms resistant to herbicides of different chemical structure and its impact on herbicidal stress in wheat

Wheat was chosen because of its importance in human nutrition and a lot of publications about toxic reactivity of wheat plants to herbicides (Song *et al.*, 2007, Bezuglova *et al.*, 2019).

The presence of other useful qualities (fixing molecular nitrogen, fighting diseases) would increase the practical and commercial value of these bacteria.

2. Materials and methods

2.1. Microorganism

The organism of interest in this research was the bacterial strain 6CH2, isolated of the soil contaminated with petrochemicals from the territory of an industrial enterprise (Republic of Bashkortostan, Russia). The pure culture was characterized according to its cultural, morphological, physiological, and biochemical characteristics using the well-established procedures (Gerhardt *et al.*, 1981; Garrity *et al.*, 2005).

2.2. Molecular identification of bacteria

Isolation of total DNA was carried out according to the method described in (Wilson et al., 1995). Amplification of the 16S rRNA gene fragment was carried out using bacterial primers 27F (5` AGAGTTTGATC (A / C) TGGCTCAG 3') and 1492R (5' ACGG (C / T) TACCTTGTTACGACTT 3') on a My Cycler amplifier (Bio-Rad Laboratories, USA). Isolation and purification of PCR products was carried out from low-melting agarose using the Wizard PCR Preps reagent kit (Promega, United States) according to the manufacturer's recommendations. Sequencing of the obtained PCR fragments of the 16S rRNA gene was performed using the Big Dye Terminator v. 3.1 kit (Applied Biosystems Inc., USA) on an ABI PRIZM 3730 automated sequencer (Applied Biosystems Inc., USA) according to the manufacturer's instructions supplied.

The search for homologous sequences was carried out using the EzBioCloud databases (http://www.ezbiocloud.net/eztaxon). A dendrogram of phylogenetic similarity was constructed in the MEGA version 7 program (http://www.megasoftware.net) by the Neighbor-Joining method (Saitou and Nei, 1987) using the Kimura model (Kimura, 1980).

2.3. Antagonism against phytopathogens

Antagonistic potential against phytopathogens was determined during joint cultivation of bacteria and filamentous fungi in Petri dishes (Chetverikov and Loginov, 2009) on Czapek Dox agar medium. Test objects were *Bipolaris sorokiniana* (Sacc.) Shoemaker VKM F-

529, Fusarium culmorum (W.G. Smith) Sacc. VKM F-844, F. gibbosum Appel et Wollenw VKM F-848, F. graminearum Schwabe VKM F-1668, F. solani (Mart) Sacc. VKM F-142, F. oxysporum Schltdl VKM F-137, F. nivale (Fr.) Ces. Ex Sacc. VKM F-3106, F. semitectum VKM F - 1938, F. avenaceum VKM F - 132, Alternaria alternate (Fr.) Keissl. VKM F-3047, Rhizoctonia solani J.G. Kuehn VKM F-895. Cultures were obtained from the All-Russian Collection of Microorganisms. The fungi were maintained on Czapek Dox agar.

2.4. Measurement of bacterial nitrogenase activity

The acetylene reduction assay was used as an indicator of nitrogenase activity of strain 6CH2, with ethylene used as a label and measured by gas chromatography (Hardy *et al.*, 1973; Korshunova *et al.*, 2013).

2.5. Measurement of bacterial indolyl-3-acetic acid

The capacity of strain 6CH2 to synthesize indolyl-3-acetic acid (IAA) was measured by immunoenzyme assay was carried out as described (Bakaeva *et al.*, 2020).

2.6. Modeling herbicide stress

For modeling herbicide stress in plants, selective herbicides containing auxin-like substances 2,4-D (2-ethylhexyl ether) - Octapon, 2,4-D (2-ethylhexyl ether) and dicamba (sodium salt) - Chistalan (LLC AHK-AGRO, Russia) and metsulfuron-methyl - Nanomet (LLC Pesticides.ru, Russia) were used. They are designed to destroy perennial, annual dicotyledonous weeds that grow among spring barley, spring and winter wheat (State catalog..., 2020).

2.7. Wheat cultivation and treatments

Plants of soft spring wheat (Triticum aestivum L.) cultivar Kinelskaya Yubileinaya were grown in Climate Chamber MLR-352H-PE (PHC Europe BV, Netherlands) in one-liter containers filled with a mixture of sand and black soil in a ratio of 1: 9, at a PAR photon flux density of 190 µmol • m⁻² • s⁻¹, 14 -hour photoperiod and temperature 26 ° C. Soil humidity was maintained at 60-80% of the total humidity capacity. On the seventh day after germination, when the third sheet start forming, plants were sprayed with a herbicide, a suspension of bacteria Pseudomonas avellanae strain 6CH2, or their mixture: per vessel 0.9 μl Octapon, 0.9 μl Chistalan, 13 μg Nanomet, 5×10^7 CFU of the strain 6CH2 (based on the working concentrations of herbicides taking into account the application rates according to the regulations). According to the regulations for herbicides used, the second or third leaf stage is the earliest period when herbicide treatment is acceptable. Control plants were not sprayed.

Growth and weight characteristics of shoots and roots were determined 14 days after spraying. For each variant of the experiment, 30 plants were grown.

On the third day after the treatment of plants, the content of chlorophyll and proline in the leaves was determined. The content of proline and chlorophyll in the leaves, dry weight of roots and shoots, and shoot length were determined using freshly plucked plant parts.

2.8. Chlorophyll measurement

The chlorophyll content was determined spectrophotometrically on spectrophotometer Selecta UV-

2005 (Selecta, Spain) after extraction with 96% alcohol (Vernon, 1960) and was expressed in mg/g dry weight.

2.9. Proline measurement

The proline content was determined using a ninhydrin reagent, as described previously (Bates *et al.*, 1973), using a calibration curve constructed with standard L-proline («Sigma», United States) and expressed in μg / g dry weight.

2.10. Statistical analyses

Statistical analyses were performed with descriptive statistics (mean) and two-sample unpaired t-test (p=0.05) to determine statistically significant differences (P < 0.05) between treatments using MS Excel. Data were expressed as averages \pm confidence interval.

3. Results

The cells of the 6CH2 strain are Gram-negative non-spore-forming mobile rods with 1-4 polar flagella, grow slowly on nutrient agar. When they are cultivated on sucrose media, colonies with a diameter of 2-4 mm, round,

shiny, translucent are formed. The optimum growth temperature is 23-25 ° C. The maximum growth temperature is 35 ° C. The metabolism is respiratory. Bacteria liquefied gelatin, did not reduce nitrates, formed levan, did not hydrolyze starch, did not form indole, and curdled milk poorly. They produced acid during fermentation of dextrose, sucrose, glycerin and formed a blue-green fluorescent pigment, when cultured on King B medium (King *et al.*, 1954).

The sequence (1413 bp) of the 16S rRNA gene from the isolated strain corresponding to 21-1433 positions of the *E. coli* nomenclature was determined and subsequently deposited in GenBank as number MT703877. The bacterial species *Pseudomonas avellanae*, *P. syringae*, *P. cannabina*, and *P. mandelii* were closest to the studied sample. The sequence similarity between strains 6CH2 and *P. avellanae* BPIC 631 was 99.15%, and with *P. syringae* KCTC 12500, *P. cannabina* CFBP 2341, and *P. mandelii* NBRC 103147 - 98.94%. To clarify the phylogenetic position of the strain, we compared its nucleotide sequence of the 16S rRNA gene and that of nearby *Pseudomonas* secies and constructed a dendrogram (Fig. 1).

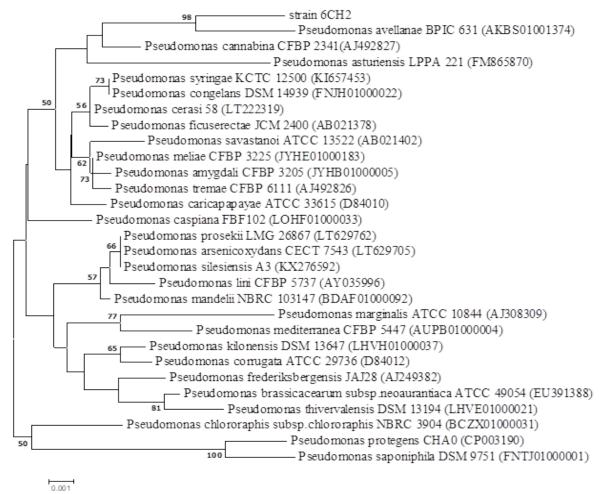


Figure 1. Phylogenetic position of the 6CH2 strain according to the analysis of the nucleotide sequence of the 16S rRNA gene. The scale shows the evolutionary distance corresponding to 1 nucleotide change in every 1000 nucleotides. The numbers show the statistical significance of the branching order determined using the "bootstrap" analysis (the values of the "bootstrap" analysis are shown above 50%). The data obtained was presented in this way to make it possible to identify the studied strain as Pseudomonas avellanae 6CH2.

The investigated strain showed antagonism against the spectrum of phytopathogenic *Fusarium* and other phytopathagenic micromycetes (table 1), without exerting much growth suppression of micromycetes from other genera. It was found that the direct interaction of test fungi of the genus *Fusarium* and the strain *P. avellanae* 6CH 2 significantly slowed the formation of mycelium compared to the control without bacteria (the spore germination delay was 24-96 h), and the morphology of the pathogens hyphae was strongly changed.

 Table 1. Antagonistic activity of the bacterial strain Pseudomonas

 avellanae 6CH2

Phytopathagenic micromycetes	Diameter of the fungal growth inhibition zone, mm
Fusarium culmorum VKM F - 844	12.0±1.5
F. gibbosum VKM F – 848	16.8±2.0
F .graminearum VKM F – 1668	14.6±1.4
F. nivale VKM F – 3106	10.4±0.5
F. semitectum VKM F – 1938	14.2±1.2
F. solani VKM F – 142	16.2±1.5
F. avenaceum VKM F – 132	20.6±2.2
F. oxysporum VKM F-137	14.4±1.3
Bipolaris sorokiniana	≤5.0
Alternaria alternate VKM F-3047	≤5.0
Rhizoctonia solani VKM F-895	≤5.0

The nitrogen-fixing activity of *P. avellanae* 6CH2 was 19.8 nmol $C_2H_4 \cdot h^{-1} \cdot ml^{-1}$. The nitrogenase activity did not decrease under the influence of the studied herbicides in the above-mentioned concentrations.

The herbicide-resistant strain 6CH2 was able to synthesize IAA both when grown in the presence and in the absence of the herbicides. The maximum IAA production was 189 ± 12 ng / ml in the culture liquid without herbicides and did not decrease by more than 10% in their presence.

In our study, treatment of wheat plants with herbicides Nanomet and Chistalan at the stage of emergence of the third leaf resulted in suppression of their growth; shoot (up to 14%) and root weight (up to 18%) (Fig. 2), shoot length (up to 12%) (Fig. 3) were significantly less than in the control group of plants not exposed to the herbicide. The variant with the herbicide Octapon, which slightly stimulated the mass accumulation and shoots growth, stood out from the general paradigm. This synthetic auxin, whose effect on sensitive plants is due to the excessive accumulation of auxins in them and a violation of their distribution between organs (Grossmann, 2007), worked in our case as a weak growth stimulant.

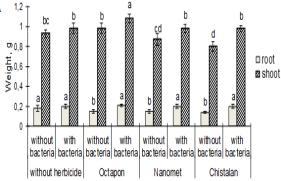


Figure 2. Effect of herbicides and spraying with strain P. avellanae 6CH2 on root and shoot weight of wheat measured 14 days after. Mean values \pm confidence interval are presented (n=30). Significantly different means of each parameter are marked with different letters (p \le 0.05)

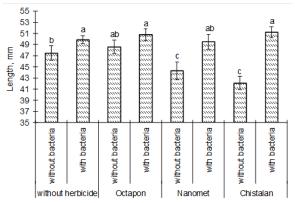


Figure 3: Effect of herbicides and spraying with strain P. *avellanae* 6CH2 on shoot length of wheat measured 17 days after. Mean values \pm confidence interval are presented (n=30). Values sharing same letters differ non-significantly (P>0.05)

Another manifestation of the negative effect of all tested herbicides on plants was a decrease in the total content of chlorophylls a and b (up to 10%) (Fig. 4).

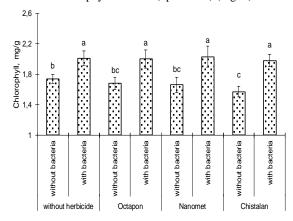


Figure 4. Effect of herbicides and spraying with strain P. avellanae 6CH2 on chlorophyll amount in wheat leafs measured 3 days after. Mean values \pm confidence interval are presented (n=30). Values sharing same letters differ non-significantly (P>0.05)

Bacterial treatment without herbicides had a positive effect on the length of the shoots and, accordingly, on the chlorophyll amount (an increase of 16%). Taking into account that the chlorophyll content is an important indicator of the state of plants under stress (Ashraf and Harris, 2013), it can be unambiguously asserted that bacterial treatment with the strain 6CH2 was not stressful for the tested plants. The positive effect of bacteria was expressed in mitigation the negative influence of the herbicide on the photosynthetic apparatus. It was reflected in the pigments quantity in plants. In all variants, inoculation with bacteria led to an increase in the total chlorophyll amount against the background of herbicidal stress.

When treating wheat plants with mixtures of bacteria and herbicides, a significant increase in all growth characteristics was observed relative to plants treated with herbicides only. The mass of plant roots reached the control parameters, and the mass and length of shoots exceeded the values in the control group by 8-16%. The increase in the mass of plant can be attributed to the bacterial indolylacetic acid responsible for root system extension and improvement of absorption of mineral elements and water. Bacteria can also increase the nitrogen and phosphorus available to plants, as shown above. The assumption of improved nutrition seems plausible because the increase in plant mass occurred in all variants of the experiment with the bacterium regardless of the herbicide type.

In our experiment, spraying with herbicides increased the concentration of proline in plants (Fig. 5). If the spraying with synthetic auxins increased the concentration of proline by 3 times relative to the control, then in the case of treatment with Nanomet, the increase was more than 500%. The treatment of plants with a mixture of herbicides and bacteria did not initiate the accumulation of proline; its concentration in them practically did not differ from the plants of the control group.

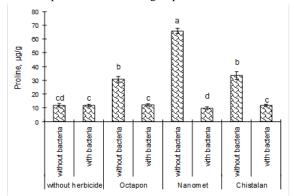


Figure 5. Effect of herbicides and spraying with strain *P. avellanae* 6CH2 on proline amount in wheat leafs measured 3 days after. Mean values \pm confidence interval are presented (n=30). Values sharing same letters differ non-significantly (P>0.05)

4. Discussion

The main problem of using herbicides is their negative effect on basic agricultural crops. Herbicide treatment can cause oxidative stress in plants, which is manifested in a slowdown in growth processes and a decrease in productivity in the final (Light *et al.*, 2005). At the same time, herbicidal treatment is the only real way to control weeds on an industrial scale, even taking into account the non-absolute selectivity of herbicides. And reducing the toxic load of herbicides on cultivated plants is a real problem.

Although it is believed that monocotyledonous plants are insensitive to the herbicides based on the synthetic auxin 2,4-D, in the case of their application before the onset of the tillering stage, inhibition of wheat plant growth was noted (Kumar and Singh, 2010). It is also believed that cereals are relatively resistant to the action of sulfonylurea herbicides, but in some cases they do not cope with their toxic effects (Barrett, 1989), not to mention the problems of dicotyledons crops that follow them in the crop rotation.

In our experiment we observed a tendency to slow growth and reduce the amount of chlorophyll in plants treated with certain herbicides. But a more obvious marker of stress was the accumulation of proline in the leaves.

The formation and accumulation of the amino acid proline is a possible physiological reaction of wheat plants to stress caused by toxic substances including herbicides (Sharma and Dietz, 2006). To date, the osmoprotective, antioxidant, signal-regulatory and other functions of this multifunctional amino acid have been established (Szabados and Savoure, 2009; de Carvalho *et al.*, 2013).

Some authors associate the ability of cells to accumulate proline with a selective assessment of drought resistance of plant varieties and species (Chaves and Oliveira, 2004). Taking into account the fact that proline plays an important role in maintaining cellular metabolism and ensures the survival of plants in extreme conditions, we also determined its amount in wheat leaves.

Metsulfuron-methyl most strongly provoked the accumulation of proline in the leaves. We tend to associate this additional increase with the mechanism of action of sulfonylurea class herbicides on plants, the primary target of which is the acetolactate synthase (ALS) enzyme. ALS is the first enzyme on the biosynthetic pathway of branched-chain amino acids (valine, leucine and isoleucine) that functions in fungi, bacteria and higher plants (Brown and Cotterman, 1994). When it is inhibited, an excess of pyruvate and oxaloacetate are formed in plant cells, then they are transformed into α-ketoglutarate (Fig. 6). After that, the metabolic pathway leading to the synthesis of glutamate begins to function as much as possible. It is known that in plants proline can be synthesized in two ways - from glutamate or ornithine. However, it is believed that the synthesis of proline under the influence of stress occurs mainly along the glutamate pathway (Liang et al., 2013). In some plants, more than a hundredfold increase in proline content was noted in response to unfavorable factors.

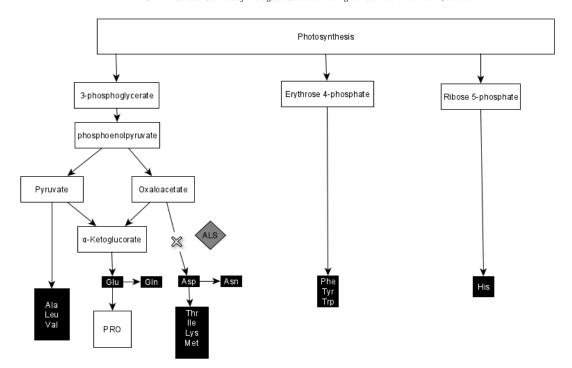


Figure 6. Biosynthesis of amino acids in plants under the influence of herbicides of the sulfonylurea class

We have previously shown that the investigated bacterial strain P. avellanae 6CH2 is stable and capable of growing at a high rate in media in the presence of herbicides based on synthetic auxins (Octapon, 10 ml / L; Chistalan, 5 ml / L) and sulfonylureas (Nanomet, 0.05 g / L) (Chetverikov, 2019).

These results show that the studied bacteria exhibited other properties characteristic of PGP - bacteria: antagonism to phytopathogens, synthesis of a nitrogenase complex and phytohormones, including in the presence of herbicides. Inhibition of mycelium growth of Fusarium fungi and changes in hyphae morphology are the result of an action of metabolites of the antagonistic bacterial strain. Metabolites of Bacillus (Melentyev and Galimzyanova, 1999) and Azotobacter (Chetverikov and Loginov, 2009) had a similar impact on Fusarium. The nitrogen-fixing activity of P. avellanae 6CH2 correlated well with the values characteristic of other known nitrogen fixers (Bakaeva et al., 2020). According to the literature, the amount and activity of nitrogenase in bacteria decreases due to various stresses (Tripathi et al., 2002; Choi and Gal, 1998). Apparently, P. avellanae 6CH2 did not experience stress in the presence of herbicides and therefore the activity of nitrogenase did not decrease, while herbicides had almost no effect on nitrogen fixation and inhibition of fungal growth, their presence led to a slight decrease in the indolylacetic acid produced by bacteria. A similar decrease in IAA secretion was observed for the Burkholderia cepacia PSBB1 strain resistant to the herbicide glyphosate (Shahid and Khan, 2018).

Auxin synthesis by bacteria may be the main reason for the stimulating effect of PGPB strains on plants. Herbicide-resistant PGPB are likely to secrete auxin sufficiently even whey are applied to herbicide-contaminated soil. High production of auxins, on the contrary, can inhibit plants, as in the case of *Enterobacter sp.* I-3 (Park *et al.*, 2015).

It was shown that PGPBs can change the concentration of auxins not only by their synthesis, but also by their degradation. For example, *P. putida* 1290 can use auxins as a nutrient substrate, thereby eliminating the inhibitory effect of high concentrations of exogenous auxins on the plant (Leveau and Lindow, 2005). Therefore, for achievement of a stimulating effect, the final amount of auxin must correspond to the optimum for the given species under the given environmental conditions.

Thus, will the strain 6CH2, which is tolerant to 2,4-D and sulfonylureas and capable of synthesizing IAA in their presence, mitigate herbicidal stress in plants? Will the amount of phytohormones synthesized by them be sufficient?

The treatment of plants with bacteria 6CH2 in addition to herbicides initiated some positive consequences: stimulation of growth, production of chlorophyll, no need to accumulate a lot of proline. Similar effects from exposure to bacteria were observed under stresses induced by herbicides paraquat (Agafonova *et al.*, 2016), fusilad (Osman *et al.*, 2016), and glyphosate (Shahid and Khan, 2018). In the case of herbicides of the synthetic auxins group, it should be noted that the ability to degrade 2,4-D may be one of the factors providing a favourable effect of PGPR bacteria on plants (Jacobsen, 1997, Han *et al.*, 2015).

In the case of metsulfuron-methyl, an active substance of the herbicide Nanomet, bacteria can mitigate its negative effect on a cultivated plant, taking the impact on itself. First, detoxification can occur by accelerating the biodegradation process. Secondly, the toxicity of the herbicide can be reduced due to its primary binding to the bacterial enzyme ALS, since it has a more convenient conformation.

5. Conclusion

Herbicides with synthetic auxins (Octapon, Chistalan) and sulfonylureas (Nanomet) have a phytotoxic (stress) effect on wheat plants, significantly influencing their growth, reducing the amount of chlorophylls and increasing the proline quantity. The strain *P. avellanae* 6CH2 isolated in this study was capable of suppressing phytopathogenic micromycetes from the genus *Fusarium*, resistant to the herbicides, and also exhibited properties characteristic of PGPB: the synthesis of a nitrogenase complex and phytohormones, including in the presence of herbicides. These properties allowed the *P. avellanae* strain 6CH2 to have an anti-stress effect (to make studied parameters at least the same as in the control group) if wheat plants were jointly treated with herbicides and bacteria.

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Authors' contributions

SC, DS, MB designed study, performed the statistical analysis, wrote and edited the manuscript. DC, MT, TR, DS carried out the experiments and analyzed the samples.

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