IDENTIFICATION AND QUANTIFICATION OF WHEAT ROOTS FLAVONOIDS INOCULATED WITH NATIVE RHIZOBACTERIA

Hércules Tancredo Moreira 1, Luciana Grange 1, Isac George Rosset 1, Robson Fernando Missio *1

*1 Federal University of Paraná (UFPR), Palotina, Paraná, Brazil

DOI: https://doi.org/10.29121/granthaalayah.v8.i10.2020.2127

ABSTRACT

In this study, we identified the presence and quantity of flavonoids produced in wheat roots inoculated with rhizobacteria. Our goal is to confirm the efficiency of standard isolates and show new strains with biotechnological potential to promote plant growth. The experiment was carried out with different isolates under inoculation in the following situations: T1-Azospirillum brasilense; T2-Herbaspirillum seropedicae; T3-Azospirillum brasilense and Herbaspirillum seropedicae co-inoculation; T4-native Enterobacter sp. n° 203; T5- native Enterobacter sp. n° 208; T6-native Enterobacter sp. n° 493; T7-Control only under nitrogen fertilization (N+); T8-Control without nitrogen (N-) and bacterial inoculation. Agronomic characteristics were assessed after 42 days of inoculation. Identification and quantification of flavonoids were carried out through HPLC, using an analytical curve with four standards based on Coumarin, Quercetin, Isoflavone and Rutin. Regarding the production of total flavonoids, two (203 and 493) out of the three native strains we tested were statistically significant, exceeding the values obtained from the inoculation of standard strains, which presented association with grasses (Azopirilum e Herbaspirillum). Standard bacteria, when inoculated in isolation, presented, along with those that received N+ treatment, the highest values for length and root and aerial part dry mass. New studies need to be carried out in order to confirm the technological use of these native strains as inoculant, as these bacteria may contribute to Biological Nitrogen Fixation (BNF) in wheat culture, either by competition or synergism.

1. INTRODUCTION

Wheat (Triticum aestivum L., from the grasses group) is considered the second most produced grain crop in the world and is an essential constituent of human nutrition worldwide [1]. The Brazilian wheat production in 2019 was approximately 5.2 million tons [2], and the State of Paraná was responsible for more than half of the national production [3].

Several obstacles make production and dissemination of wheat crop throughout Brazil difficult. Such obstacles include technical and biological factors during farming, problems in diseases control [4], other species that are more easily cultivated such as maize and soybean (which are given priority), besides the lack of credit or credit problems and attractive prices [5]. Another limiting factor for the cultivation of many crops, especially wheat, is the cost of
fertilizers, particularly the ones containing nitrogen [6]. Therefore, technology development and studies have been carried out in order to make it easier for maize, rice and wheat crops to absorb nitrogen [7]. The lines of research most studied included genetic improvement and increase of associative and beneficial relationships between the plant and nitrogen-fixing microorganisms [8].

The first stage of BNF is extremely important since it is when the bacterium-plant biochemical signaling happens. On that stage, thousands of microorganisms are attracted by the exudates of the roots and quickly approach each other to assemble and/or colonize root tissues [9]. In this highly competitive moment, the most efficient bacteria need to be the first to start the symbiotic fixation process and/or to establish free-living assemblages before the plant starts to establish inefficient or phytopathogenic relationships [10].

Among the most studied plant exudates are those from the flavonoids group [11]. They are phenolic compounds identified as hydrosoluble secondary metabolites present in almost all plant organs and have differentiated functions in the metabolism of plants [12]. Many studies showed the participation of flavonoids in antioxidant and stimulant processes, being beneficial to the plant [13].

Research reported that flavonoids make up most of plants exudates; however, few studies have focused on their production in roots tissues [14]. Moreover, even fewer studies have been carried out to identify the different types of those compounds and their respective functions, especially of those involved in the communication between rhizobacteria and plants [15]. Therefore, we emphasize the importance of the study as it investigates those compounds in wheat culture, which presents low responsiveness to microbial communication and high nutritional demands [16].

Thus, the present study identified the presence and quantity of flavonoids produced in wheat roots inoculated with rhizobacteria, aiming to confirm the efficiency of standard isolates as well as to show new strains with biotechnological potential to promote plant growth.

2. MATERIALS AND METHODS

For the experiments, it was chosen to grow CD 150 wheat, early cycle. It is defined as a short plant with medium to high fertility, high grain yield potential and high plant lodging resistance, which makes it tolerate drought periods, and today it represents the most grown cultivar in western Paraná [17].

The standard microorganisms Azospirillum brasilense and Herbaspirillum seropedicae were obtained from the EMBRAPA soybean crop bank, and the Enterobacter sp. (nº 203, 208 and 493) native strains, from the UFPR-Palotina crop collection. The experiment was carried out following a fully casualized design with 8 treatments (T) and 4 replicates. The plants were inoculated with different isolates and were submitted to the following treatments: T1 - Azospirillum brasilense; T2 - Herbaspirillum seropedicae; T3 - A. brasilense and H. seropedicae co-inoculation; T4 - native Enterobacter sp. nº 203; T5 - native Enterobacter sp. nº 208; T6 - native Enterobacter sp. nº 493; T7 - Control only under nitrogen fertilization (N+); T8 - Control without nitrogen (N-) and bacterial inoculation. All treatments were submitted to fertilization, being used two nutritious solutions previously prepared, considering nitrogen values of 0 mg L⁻¹ (Table 1) and 50 mg L⁻¹ (Table 2).

Sowing was carried out in vases containing medium-grain-sized sand sterilized in autoclave [18]. The seeds were disinfected with a 1% sodium hypochlorite solution, shaken for 3 minutes, followed by a disinfection with 70% alcohol, shaken for 1 minute. Those steps were followed by washing with deionized water and sterilization for 6 times straight [19]. After sowing, the vases were taken to a greenhouse at 25ºC under controlled humidity (around 60%).

<table>
<thead>
<tr>
<th>Reagente</th>
<th>mg L⁻¹</th>
<th>P</th>
<th>K</th>
<th>Ca</th>
<th>Mg</th>
<th>S</th>
<th>Cl</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sulfato de Magnésio</td>
<td>530</td>
<td>50.35</td>
<td>68.90</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fosfato de Potássio</td>
<td>250</td>
<td>57.5</td>
<td>72.50</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cloreto de Cálcio</td>
<td>550</td>
<td>149.65</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cloreto de Potássio</td>
<td>340</td>
<td>176.80</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Micronutrientes*</td>
<td>1 mL</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*100 mL⁻¹ (FeEDTA-4,21; ZnSO₄₀,32; H₃BO₃₀,12; CuSO₄₀,50; Na₂Mo 4-0,015; MnSO₄.H₂O-1,01)
Table 2: Nutritional solution used in the experiment with 50 mg L\(^{-1}\) of nitrogen.

<table>
<thead>
<tr>
<th>Reagente</th>
<th>mg L(^{-1})</th>
<th>N</th>
<th>P</th>
<th>K</th>
<th>Ca</th>
<th>Mg</th>
<th>S</th>
<th>Cl</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nitrato de Potássio</td>
<td>357</td>
<td>49.98</td>
<td>130.30</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sulfato de Magnésio</td>
<td>530</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fosfato de Potássio</td>
<td>250</td>
<td></td>
<td>57.50</td>
<td>72.50</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cloreto de cálcio di-hidratado</td>
<td>550</td>
<td></td>
<td></td>
<td></td>
<td>149.65</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cloreto de Potássio</td>
<td>10</td>
<td></td>
<td>52.00</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Micronutrientes*</td>
<td>1 mL</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*g 100 mL\(^{-1}\) (FeEDTA-4.21; ZnSO\(_4\)-0.32; H\(_3\)BO\(_3\)-0.12; CuSO\(_4\)-0.50; Na\(_2\)Mo 4-0.015; MnSO\(_4\).H\(_2\)O-1.01)

After 42 days, the plants were collected and separated the aerial parts and the roots from the portion next to soil. After that, the length of both vegetative parts was measured. The plant material was dried for 72 hours at 45 °C in an oven with air circulation and renewal in order to obtain the dry matter (DM).

The extracts for flavonoids quantification were obtained from 300 mg of macerated dry root and submitted to reflux for 15 minutes. Were used 50 mL of 70% ethanol as the solvent. A qualitative filter paper was used to obtain the results, storing them in amber bottles in the refrigerator [20].

In order to quantify total flavonoids through spectrophotometry, it was built a 10-point standard curve with Quercetin and added 70% ethanol as the solvent (0.01; 0.05; 0.1; 0.3; 0.5; 1.0; 3.0; 5.0; 8.0 e 10 mg mL\(^{-1}\)). The equation of the line presented an R\(^2\) over 0.999. For each 2 mL of the samples we added 0.8 mL of 95% ethyl alcohol, 0.1 mL of 10% aluminum chloride, 0.1 mL of 1 mol L\(^{-1}\) sodium acetate, and 2 mL of distilled water. The solutions reacted for 40 minutes in the absence of light. The same procedures were performed for the negative control samples, and the 2 mL specimen was replaced by 2 mL of 70% alcohol. Then, was read the spectrophotometer (Spectrum SP 2000UV BellPotonics®) at a wavelength of 415 nm [21].

For HPLC analysis, it was built an analytical curve consisting of 4 standards based on Cumarin, Quercetin, Isoflavone and Rutin [22] diluted in methanol:water (70:30). In order to make a stock standard solution, were weighted 25 mg and diluted it in 25 mL of the solvent aforementioned. The curve was composed of 8 distinct concentration points diluted in methanol:water (70:30). These points were: 0.2 mg L\(^{-1}\); 0.4 mg L\(^{-1}\); 1 mg L\(^{-1}\); 5 mg L\(^{-1}\); 10 mg L\(^{-1}\); 50 mg L\(^{-1}\); 70 mg L\(^{-1}\); and 100 mg L\(^{-1}\). The linear equation presented a value of R\(^2\) above 0.998, at all events. The samples from all the treatments were filtered through PVDF (Polyvinylidene fluoride) filter and introduced into sterile vials for HPLC analysis (with UltiMate 3000 by Thermo Scientific®, using chromeleon 7.1).

For HPLC analysis, were used the following definitions, which were adapted from [23]: gradient system ACN:H\(_3\)PO\(_4\) (M:V) (Ph 2.75) 0-5 min ACN 22%, 5-6 min ACN 29%, 6-15 min ACN 29%, 15-16 min ACN 22%, 16-20 min ACN 22%; Phenomenex Column c18 25 cm x 4.6 mm x 5 μm; sample injection of 20 μL in mobile phase of 1.0 mL min\(^{-1}\), with system pressure at 1800 PSI and run time of 20 min at 30 °C, using UV-VIS detector at 260 nm.

All data were submitted to variance analysis and a comparison between the obtained averages was made using the t-test to 5% probability.

3. RESULTS AND DISCUSSIONS

Figure 1 shows the amount of total flavonoids as well as the amount of Quercetin and Rutin detected after HPLC quantification. The average value of total flavonoids found was 0.4455 µg g\(^{-1}\) root. The treatments with the highest average values were those inoculated with Enterobacter sp. number 203 and number 493, which produced 0.5044 and 0.4935 µg g\(^{-1}\) root respectively. These treatments differed statistically from Control N+ (T7), which had the lowest value (0.3829 μg g\(^{-1}\) of root), followed by inoculation with H. seropedicae (T2), with a production of 0.3964 µg g\(^{-1}\) of root.

The presence of nitrogen in the form of ammonia, in large quantities has significantly reduced the production of chemotactic flavonoids in roots [24]. That is partly because root tissues cease to trigger metabolic cascades (that are regulated by nutritional stress, i.e. nitrogen requirement) under excessive presence of nitrogen [25]. That demand triggers flavonoids production to communicate with beneficial microorganisms, in order to obtain the most assimilable form for plants (NO\(_2\)) and only the required amount of such nutrient [26]. If there is no biochemical communication between roots and microbial community, rhizosphere is weakened on enzymatic functions. That
may occur because of lack of nitrogen fixers and their nitrogenases, lack of growth promoters, plant hormones production, antibiotics, phosphate solubilization and other processes [27].

![Flavonoids, Quercetin, and Rutin levels](image)

**Figure 1:** Concentration of total Flavonoids, Quercetin, and Rutin in relation to total in wheat roots (CD150).

With the HPLC analysis (Figure 2), it was possible to quantify Quercetin and Rutin levels within the total flavonoid concentration of plant root, with a percentage that varies depending on treatment: from 2.2 to 2.9% for Quercetin and from 2.7 to 7.0% for Rutin. The values of Quercetin in all treatments correspond to an average of 2.55% of total flavonoids. Whereas, the values of Rutin were quite different. That is the case of the treatments with *A. brasiliense* and *A. brasiliense* in association with *H. seropedicae*, which reached, respectively, 2.7% and 7% of total flavonoids of the wheat root.

![HPLC analysis results](image)

**Figure 2:** Results of HPLC analysis in wheat roots (CD150).

Flavonoids, such as Quercetin and Rutin, are related to auxin transport, which affects the plant development [28]. Thus, such flavonoids can be considered endogenous regulators of auxin transport [29], protecting the plant from oxidative stress, since they are recognized as structures that sequester free radicals - not only in plants [30].

It is possible that potting conditions, under the lack of nitrogen as a nutrient within the recommendations of Cultivar (CD150), induced the formation of reactive O₂ species, causing increase in Quercetin and Rutin levels, as a protective effect against oxidative damage [25]. Different metabolic pathways of responding to biotic and abiotic stresses are normally triggered by genetically-engineered cascades, such as tissue dehydration, reduction of mitotic divisions in meristematic tissue, tissue apoptosis, reproductive tissues abortion, ethylene production and plant and crop death [31].
Was observed agronomic differences between root length and root dry matter and aerial dry matter treatments (Table 3). As for root length and root dry matter, the enhance is the treatments T1 - *A. brasiliense*, T2 - *H. seropedicae* and T7 - Control N+, which resulted in an average growth of 39.31 cm. The smallest root length was obtained by isolated inoculation with the native bacteria No. 208 (T5) (29.87 cm), and it did not differ statistically from T3, T4 and T6. On the other hand, studies carried out by our team (not published yet) show that the same strain, co-inoculated with standard *Azospirillum* in maize, was able to increase the vegetative growth by more than 50% compared to the plants obtained from the positive control (inoculated only with the standard bacteria).

T7 treatment (N+ control) provided the highest root dry matter (0.8282 g), followed by treatments T1 - *A. brasiliense* (0.5835 g), T2 - *H. seropedicae* (0.5274 g) and T5 - *Enterobacter* sp. 208 (0.5610 g), these three presented no statistical difference between each other. *Enterobacter* sp. 493 treatment provided the lowest root dry matter (0.3695 g). Those data (Table 3) confirm the efficiency of the standard strain, which were selected, evaluated and exhaustively tested through FBN study with non-legumes. Works from [32] and [33] on wheat and bean crops show that it is possible to verify significant agronomically efficiency of symbiosis and or of associative plants relationships cultivated with BPCVs, properly identified, registered and or recommended by the Ministry of Agriculture as promoters of plant growth.

**Table 3: Average values of agronomic variables studied in wheat (CD150).**

<table>
<thead>
<tr>
<th>Treatments</th>
<th>Root length* (cm)</th>
<th>Root dry matter (g)</th>
<th>Aerial length (cm)</th>
<th>Aerial dry matter (g)</th>
</tr>
</thead>
<tbody>
<tr>
<td>T1 - <em>Azospirillum</em></td>
<td>39.67 ab</td>
<td>0.5835 b</td>
<td>29.57 c</td>
<td>54.75 ab</td>
</tr>
<tr>
<td>T2 - <em>Herbaspirillum</em></td>
<td>38.92 ab</td>
<td>0.5274 bc</td>
<td>29.90 bc</td>
<td>49.00 b</td>
</tr>
<tr>
<td>T3 - <em>Herbas.+ Azospirillum</em></td>
<td>34.50 b</td>
<td>0.4071 cd</td>
<td>32.50 ab</td>
<td>46.67 bc</td>
</tr>
<tr>
<td>T4 - <em>Enterobacter</em> sp. n203</td>
<td>32.50 b</td>
<td>0.4178 cd</td>
<td>32.75 a</td>
<td>51.25 ab</td>
</tr>
<tr>
<td>T5 - <em>Enterobacter</em> sp. n208</td>
<td>29.87 b</td>
<td>0.5610 b</td>
<td>29.90 bc</td>
<td>54.75 ab</td>
</tr>
<tr>
<td>T6 - <em>Enterobacter</em> sp. n493</td>
<td>31.37 b</td>
<td>0.3695 d</td>
<td>34.25 a</td>
<td>50.50 b</td>
</tr>
<tr>
<td>T7 - Control N+</td>
<td>39.35 ab</td>
<td>0.8282 a</td>
<td>29.37 c</td>
<td>60.25 a</td>
</tr>
<tr>
<td>T8 - Control N-</td>
<td>48.87 a</td>
<td>0.3944 cd</td>
<td>29.00 c</td>
<td>37.50 c</td>
</tr>
<tr>
<td>CV (%)</td>
<td>20.65</td>
<td>17.67</td>
<td>6.05</td>
<td>12.52</td>
</tr>
<tr>
<td>Average</td>
<td>36.96</td>
<td>0.5154</td>
<td>30.85</td>
<td>50.71</td>
</tr>
</tbody>
</table>

*Averages followed by equal letters do not differ statistically by the t-test (p> 0.05).*

Although not statistically different from other treatments (Table 3), N-control showed the highest average root length (48.87 cm). This was probably because of direct nitrogen deficiency, which led to metabolism changes in the root cells, causing tissue stretching to search the nutrient in the environment with sandy and poor soil [34]. On the other hand, the weight value of root dry matter for this treatment (T8) was among the lowest: this is a reflection of stretching, which occurs at the expense of the promotion of vegetative growth itself [35].

The highest aerial dry matter values were obtained from treatments T1 (inoculated only with *A. brasiliense*), T5 (*Enterobacter* sp., Native 208) and T7 (Control N+), with an average of 56.58 g plant⁻¹. Similar results were observed in researches on wheat and inoculation with *Azospirillum* sp., correlating the amount of dry matter with high nitrogen availability [35].

4. **CONCLUSIONS AND RECOMMENDATIONS**

The next experimental step to test the agronomic potential of the native strains is to co-inoculate them with standard isolates for wheat crop; as well as to evaluate from agronomic and environmental aspects, whether the native strains can be used as collaborators of Biological Nitrogen Fixation (BNF); and or to indicate the need to preserve the native strains in agricultural environment aiming to improve through synergism and or competition the efficacy of strains commercially used as inoculants.

Treatments inoculated with the native strains Enterobacter sp. 203 and 493 (T4 and T6) were the ones with the highest concentration of total flavonoids in the root. Control N+ (T7) delivered a significant reduction of the presence of flavonoids in wheat roots.
Identification and Quantification of Wheat Roots Flavonoids Inoculated with Native Rhizobacteria

The highest values for length, root dry matter, and aerial dry matter were obtained from treatments inoculated with the isolates type *A. brasiliense* (T1) and *H. seropedicae* (T2), in addition to Control N+ (T7).

In order to continue exploring the agronomic potential of the native strains, it will be necessary to co-inoculate them with standard isolates for wheat crop and to evaluate if they can be used as collaborators of the BNF.

**SOURCES OF FUNDING**

This research received no specific grant from any funding agency in the public, commercial, or not-for-profit sectors.

**CONFLICT OF INTEREST**

The author have declared that no competing interests exist.

**ACKNOWLEDGMENT**

We would like to thank the Conselho Nacional de Desenvolvimento Científico e Tecnológico – CNPq and Universidade Federal do Paraná (UFPR) in special of Laboratório de Análise Instrumental.

**REFERENCES**


