

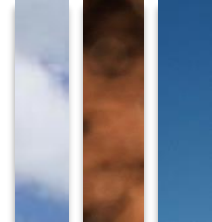
Phyto2Energy Project

**WP2: Plant-microbes
interactions:
improving biomass
production and
remediation**

PHYTO2ENERGY

**Phytoremediation driven energy crops
production on heavy metal degraded areas as
local energy carrier**





***The new bioinoculum: some aspects of
(1) development and
(2) evaluation***



WP2: Results

What has been done ?

Development of the inocula application method



C

Phyto2Energy
 4L: 4% Molasses + 1% inulin
 Date of preparation: 10.03.2016 r.

The product contains three components: A,B,C which had to be mixed before use.
 A,B,C components mixing time ___/___/___
 YYYY/MM/DD

INSTRUCTION

Component A: bacterial lyophilisate

↓

Component B: 0.85% NaCl

↓

Component C: 4% molasses + 1% inulin

↓ 24h incubation

Dilution and application of the product

1st step
 Add Component B (0.85% NaCl) to Component A (bacterial liophylisate) and mix.

2nd step
 Add mix Component A (bacterial liophylisate) + Component B (0.85% NaCl) to Component C (4% molasses + 1 % inulin). Then mix and leave for 24h at room temperature.

3rd step
 Use inocula obtained according to the step 2 in 10% solution (4L inoculum + 36L water).



The overall goal is:

To

Demonstrate the freeze-drying as a promising method for bioinoculum production on the example of *Pseudomonas putida* strains

The specific aim:

Evaluation of some biological activities on example of *P. putida* after freeze-drying

This aim will be reached by the comparing the activities of freeze-dried and not freeze-dried cells of *P. putida*

WP 2: Why the lyophilization process was chosen ?

Freeze dried cultures have some **advantages**, including:

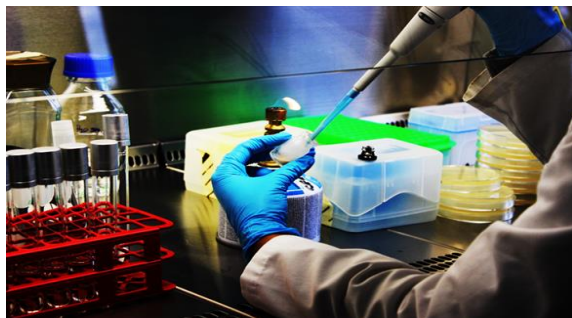
- their low volume,
- convenience for transportation and storage,
- ease of use.

Disadvantages:

- freeze-drying can cause many types of damage to cells, including a loss of viability,
- reduction of metabolic activity, and
- changes in cell morphology, etc.

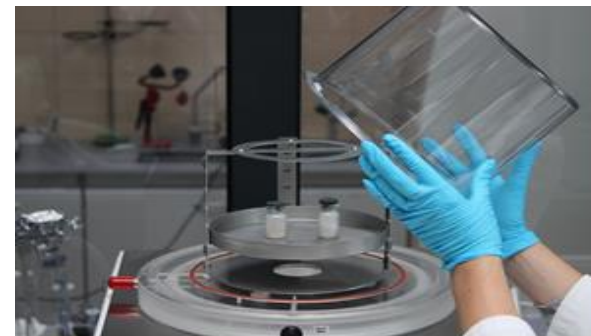


WP2: Evaluation of lyophilization efficiency – number of bacteria



| Time (days) | Number of bacteria (after lyophilization - AL) [cfu/ml] | Number of bacteria (before lyophilization - BL) [cfu/ml] | BSR = (logAL/logBL) x 100 |
|-------------|---------------------------------------------------------|----------------------------------------------------------|---------------------------|
| 0 | 2.4×10^{10} | 4.8×10^{11} | 88.87 |
| 3 | 1.3×10^{10} | 4.8×10^{11} | 86.59 |
| 14 | 5.4×10^9 | 4.8×10^{11} | 83.32 |
| 60 | 3.0×10^9 | 4.8×10^{11} | 81.14 |
| 180 | 2.1×10^8 | 4.8×10^{11} | 71.23 |
| 360 | 6.4×10^7 | 4.8×10^{11} | 58.77 |

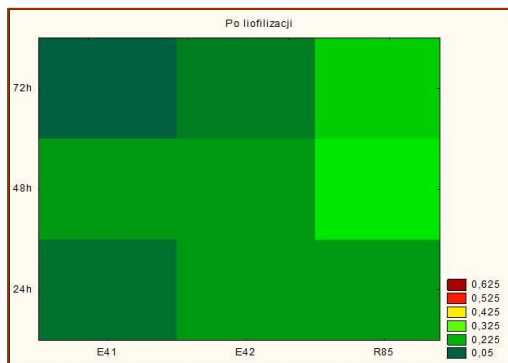
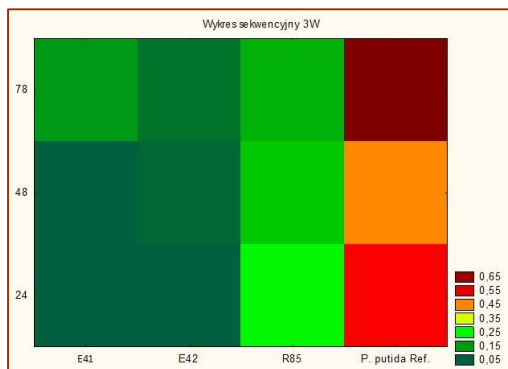
| Time (days) | Number of bacteria (after lyophilization - AL) [cfu/ml] | | | Number of bacteria (before lyophilization - BL) [cfu/ml] | | | BSR = (logAL/logBL) x 100 | | |
|-------------|---------------------------------------------------------|----------------------|----------------------|----------------------------------------------------------|----------------------|----------------------|---------------------------|--------------|--------------|
| | E41 | E42 | R85 | E41 | E42 | R85 | E41 | E42 | R85 |
| 0 | 1.9×10^{10} | 1.4×10^{10} | 3.4×10^{10} | 1.0×10^{11} | 4.9×10^{10} | 2.5×10^{11} | 93.28 | 94.85 | 92.44 |
| 180 | 2.5×10^8 | 8.2×10^8 | 2.2×10^8 | 1.0×10^{11} | 4.9×10^{10} | 2.5×10^{11} | 76.36 | 83.34 | 73.16 |



The following properties of *P. putida* strains were evaluated before and after freeze-drying:

- ❖ Enzymes activities – API-ZYM® (BioMerieux) test and the plate method with various media
- ❖ Biofilm formation - crystal violet (CV) method
- ❖ Biosurfactant production – blood agar, drop-collapse method, surface tension measurements
- ❖ Antibiotic sensitivity- disc diffusion method, PMs
- ❖ FAME analysis
- ❖ Antifungal activity
- ❖ Phenotype MicroArrays – metabolic activities (PM1, PM2, PM 3, PM4, PM9, PM10, PM11, PM12 and PM13 plates)

- Biofilm formation**



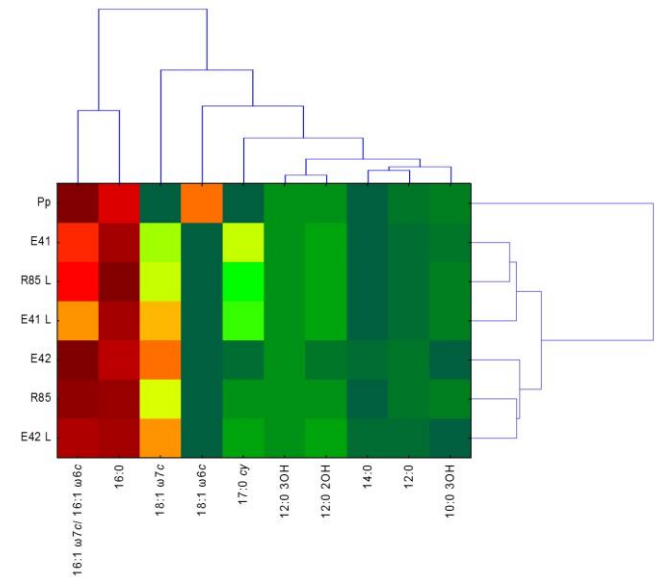
- Biosurfactant production**



| Strains | Surface tension (mN/m) | | |
|----------------------|------------------------|---------------------------------|----------------------------------|
| | Before lyophilization | After lyophilization (6 months) | After lyophilization (12 months) |
| <i>P. putida</i> E41 | 25.96 ± 0.49 | 35.18 ± 0.28 | 31.57 ± 0.37 |
| <i>P. putida</i> E42 | 52.23 ± 0.46 | 50.93 ± 0.63 | 51.27 ± 0.67 |
| <i>P. putida</i> R85 | 54.62 ± 0.46 | 45.52 ± 0.64 | 48.74 ± 0.59 |

FAME method proved its applicability for monitoring the changes in the cellular fatty acids composition of microorganisms under stress conditions like lyophilization

| FAMEs | Strain | | | | | | |
|------------------|---------|-----------------------|-------|-------|----------------------|-------|-------|
| | Control | Before lyophilization | | | After lyophilization | | |
| | Pp | E41 | E42 | R85 | E41 | E42 | R85 |
| saturated (s) | 43,51 | 61,68 | 41,32 | 51,30 | 60,02 | 49,15 | 61,03 |
| unsaturated (us) | 34,59 | 38,30 | 57,56 | 48,33 | 39,93 | 50,54 | 38,74 |
| s:us ratio | 1,26 | 1,61 | 0,72 | 1,06 | 1,50 | 0,97 | 1,58 |

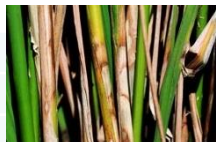


The FAME patterns showed that the freeze-drying was mostly influenced on the FAME profile of R85 strain.

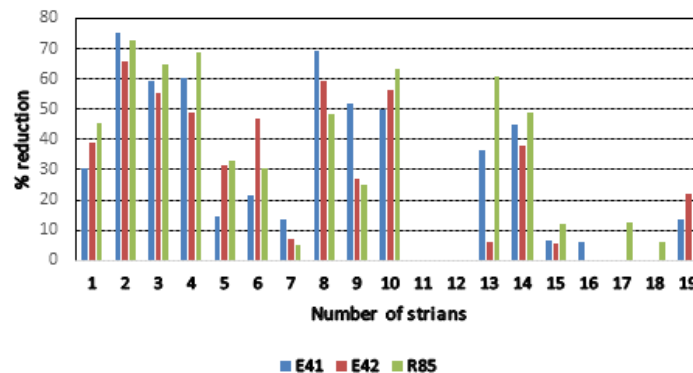
For this strain the highest ratio saturated:unsaturated FAMEs and decrease the membrane permeability were observed after lyophilization.

The analysis of individual FAMEs showed that the two fatty acids 17:0 cy and 16:1 ω7c/16:1 ω6c were characterized for R85 strain among all studied bacteria.

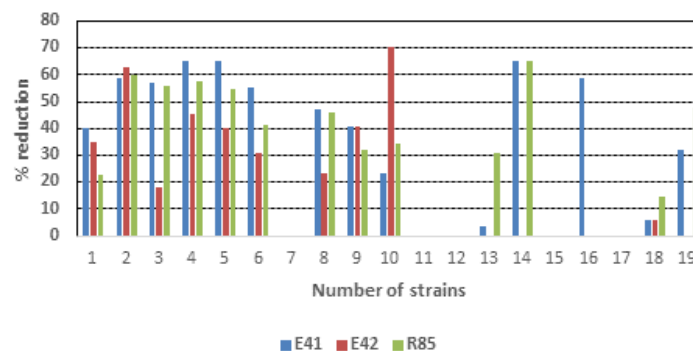
- 1 *Alternaria alternata*
- 2 *Boeremia strassesi*
- 3 *Colletotrichum dematium*
- 4 *Colletotrichum fuscum*
- 5 *Cylindrocarpon destructans*
- 6 *Diaporthe eres*
- 7 *Diplocereus hypericinum*
- 8 *Fusarium equiseti*
- 9 *Fusarium oxysporum*
- 10 *Phylloticta plantagnus*
- 11 *Rhizoctonia solani*
- 12 *Sclerotinia sclerotiorum*
- 13 *Fusarium avenaceum* - 2
- 14 *Fusarium oxysporum*
- 15 *Fusarium graminearum*
- 16 *Fusarium culmorum* - 2
- 17 *Fusarium graminearum*



Before lyophilization



After lyophilization

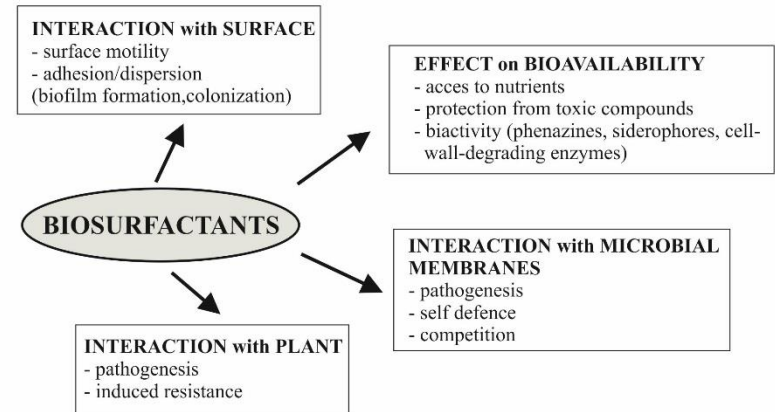
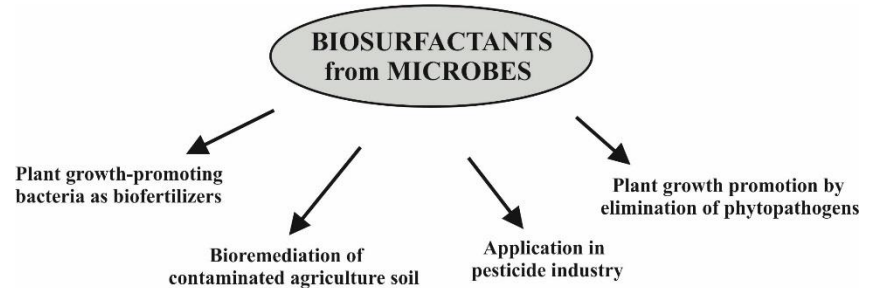


BIOSURFACTANTS are synthesized by environmental isolates, and has promising role in the agricultural industry.

Many rhizosphere and plant associated microbes produce biosurfactant; these biomolecules probably play also vital role in plant–microbe interactions.

In agriculture, biosurfactants can be used for plant pathogen elimination and for increasing the bioavailability of nutrient for beneficial plant associated microbes.

Biosurfactants can widely be applied for improving the agricultural soil quality by soil remediation.



Biosurfactant production:

- blood agar
- methylene blue agar
- oil spreading method
- drop-collapse method
- surface tension measurements



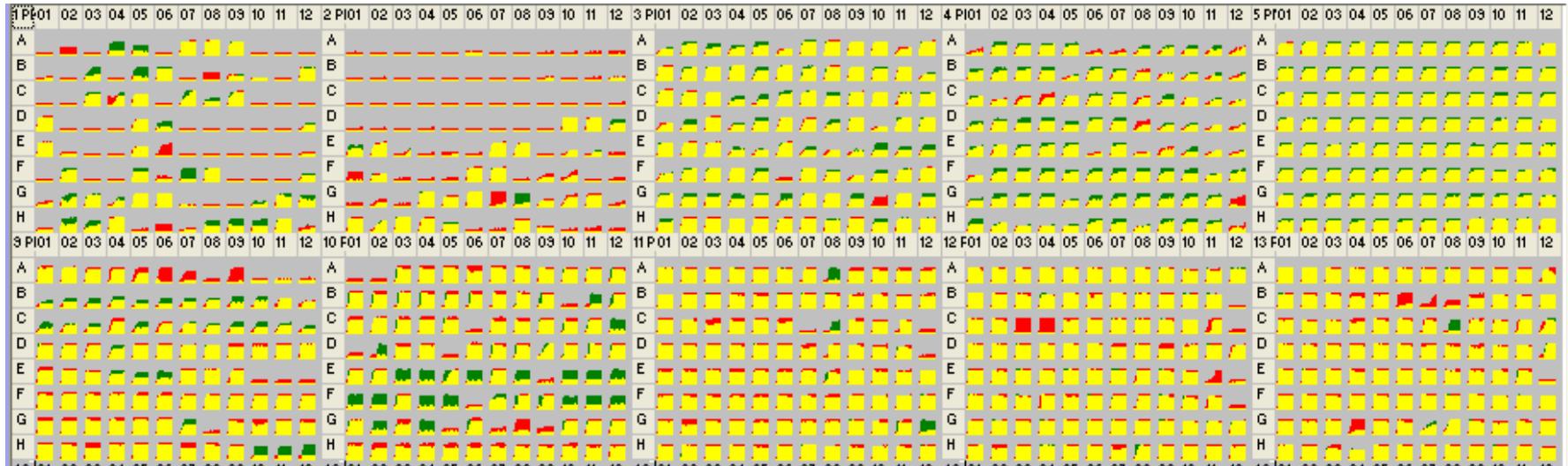
E 41

Phenotype MicroArrays:

PM1 and PM2 – carbon sources, PM3 – nitrogen sources, PM4 – phosphorus & sulfur sources, PM 5 – nutrient supplements, PM9 – osmolytes, PM10 – pH, PM11 – antibiotics, PM12 – antibiotics, PM13 – antibiotics +chemicals

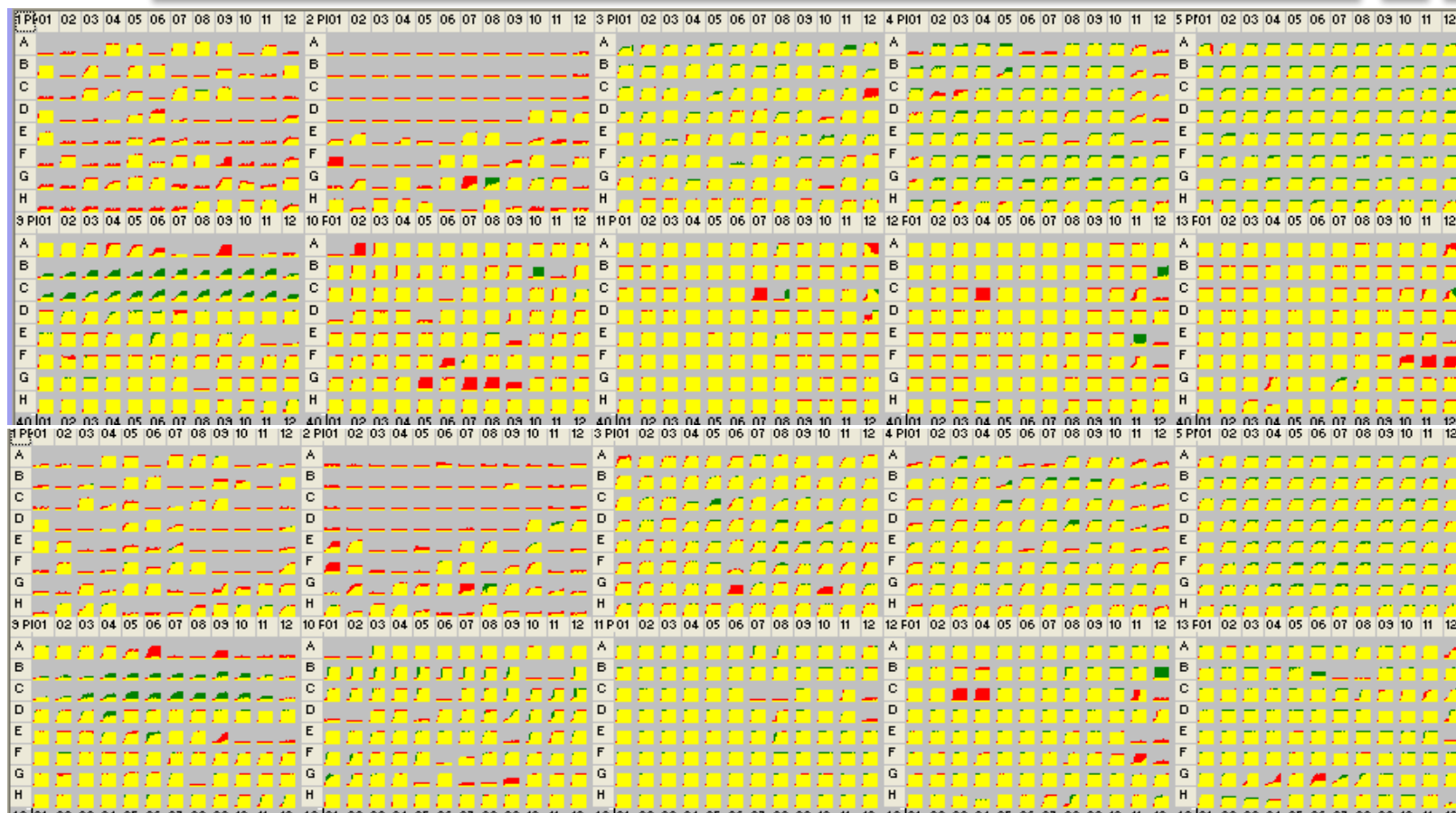
Green - before lyophilization; **red** – after lyophilization; **yellow** – common part (no changes)

E41



E42

R85

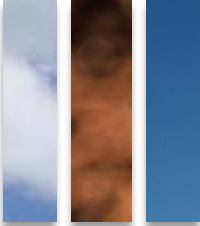


Assessment of the ecophysiology in soil using in vitro substrate degradation tests
(Establish various biodiversity indices and functional activities of communities (CLPPs) by the EcoPlates Biolog)

What is the biodiversity response to the different treatments implemented over time ?

- **T0:** 2014
- **T1:** 2015
- **T2:** 2016
- **T3:** 2017

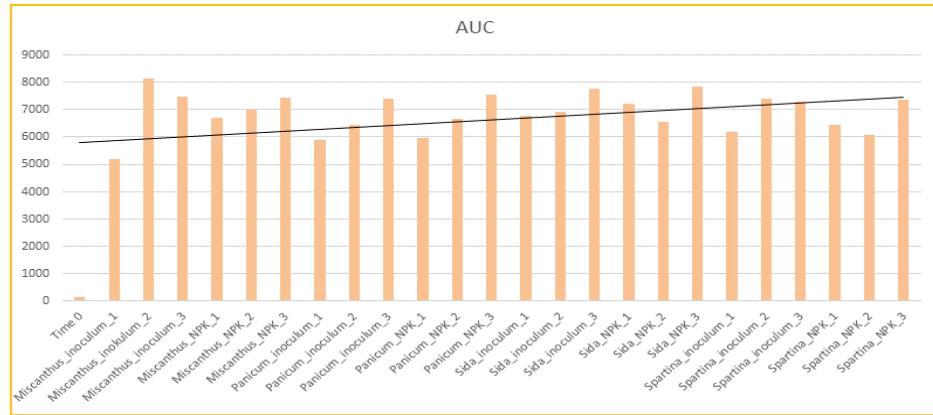
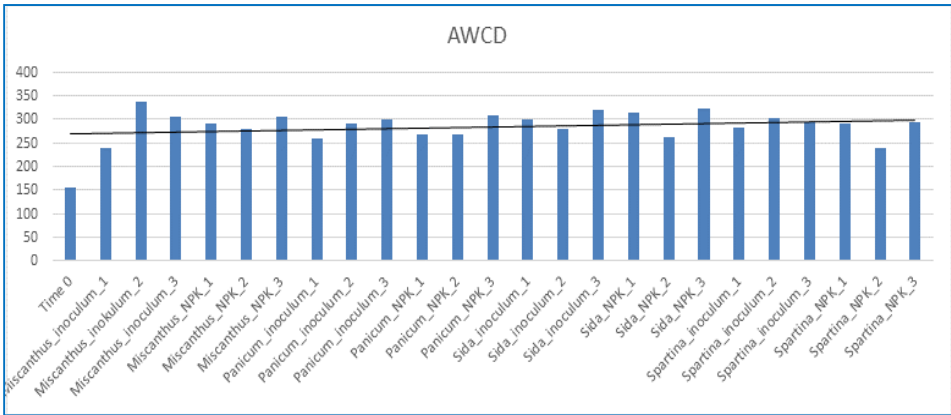
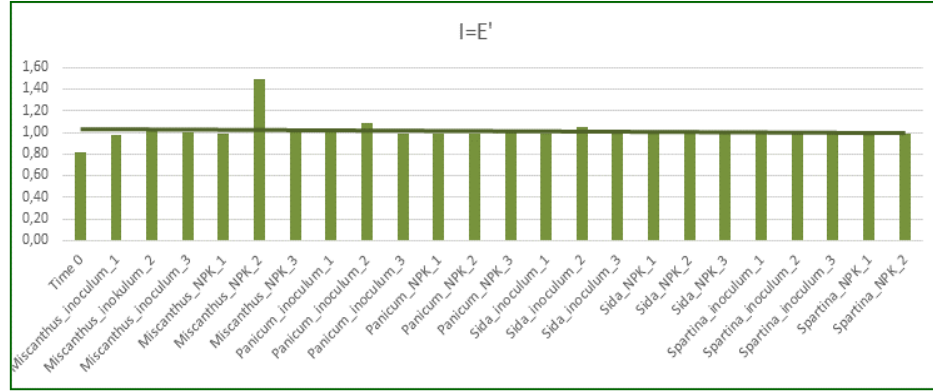
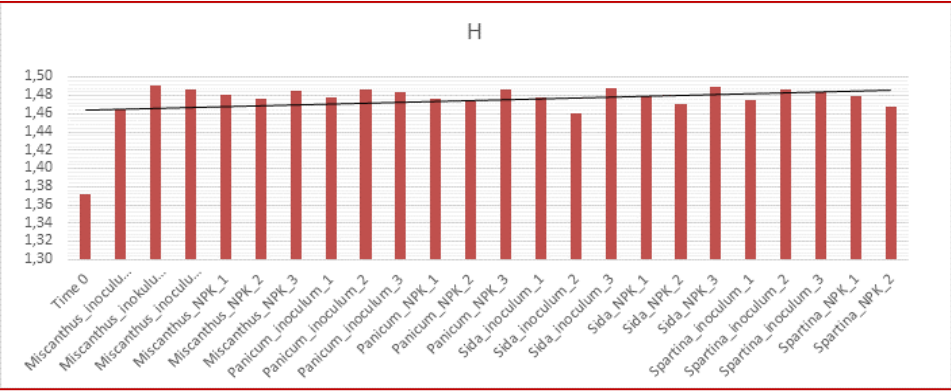
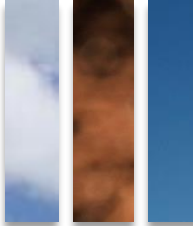


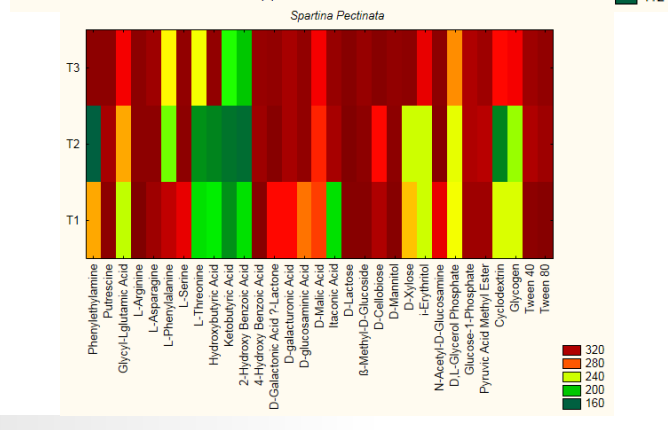
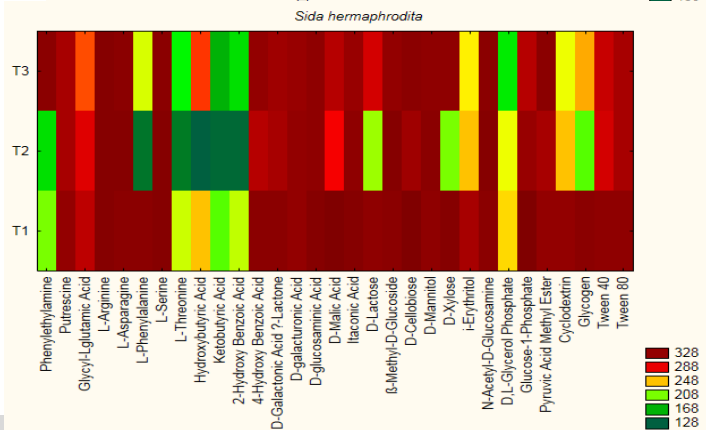
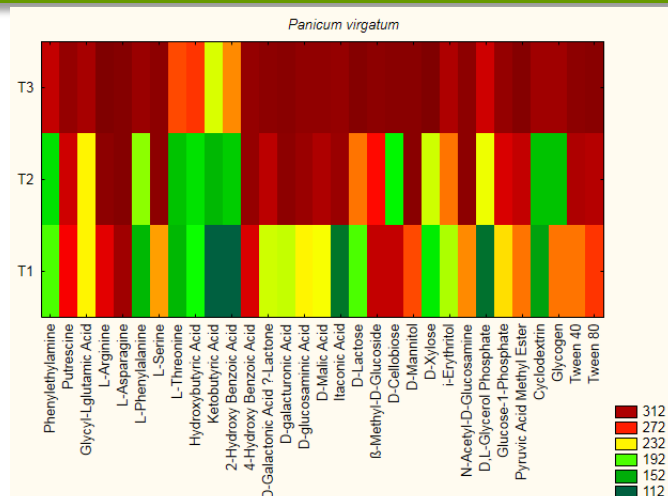
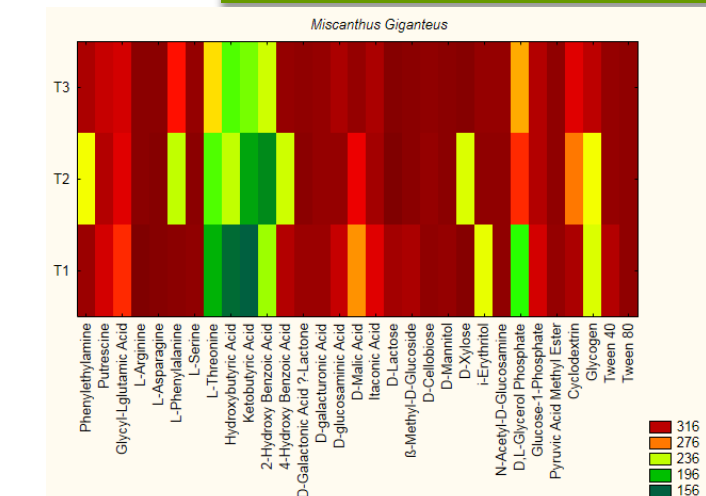


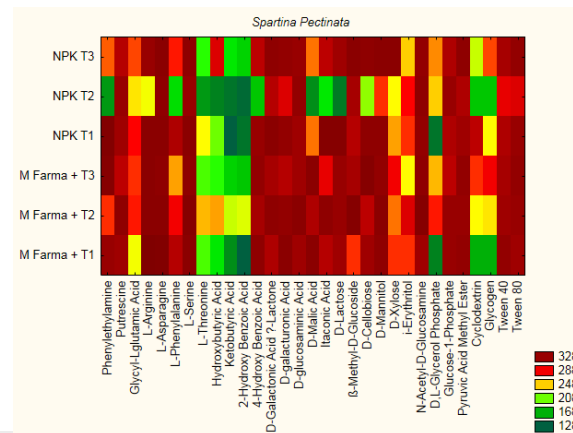
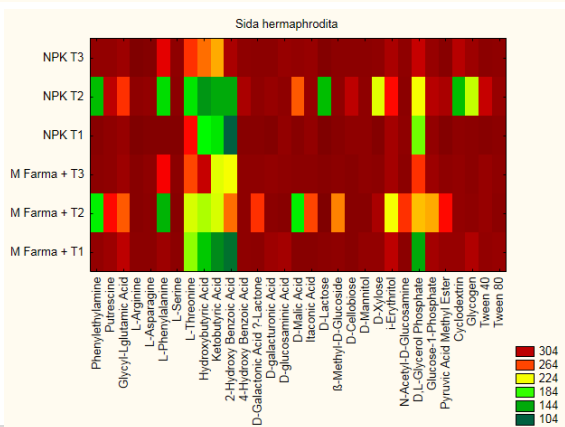
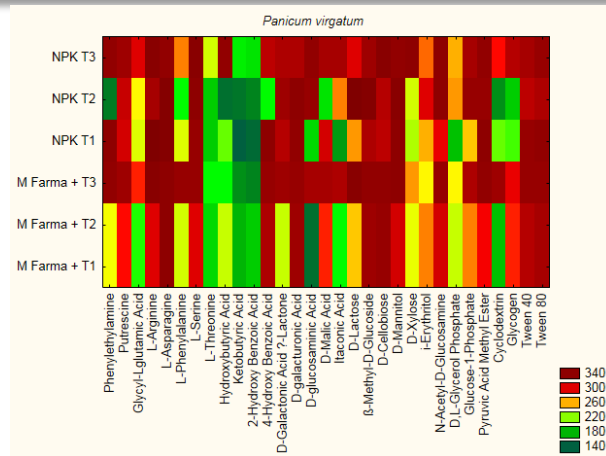
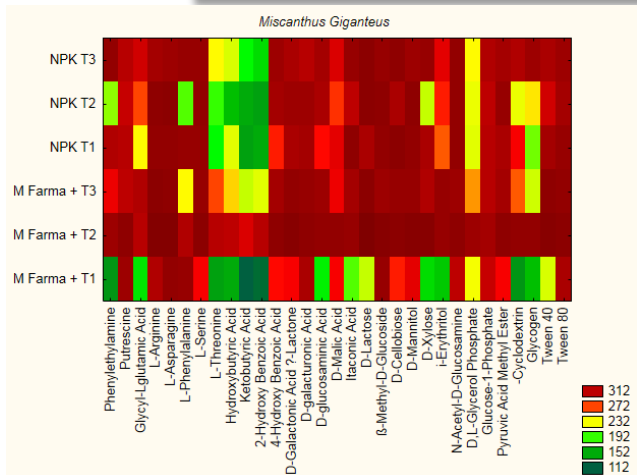
INDICES of biodiversity and biological activity :

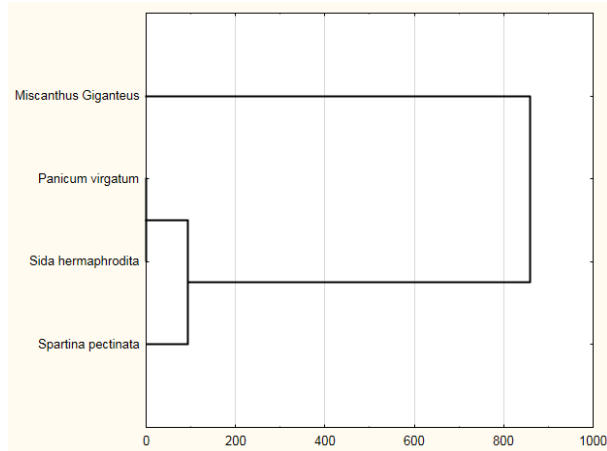
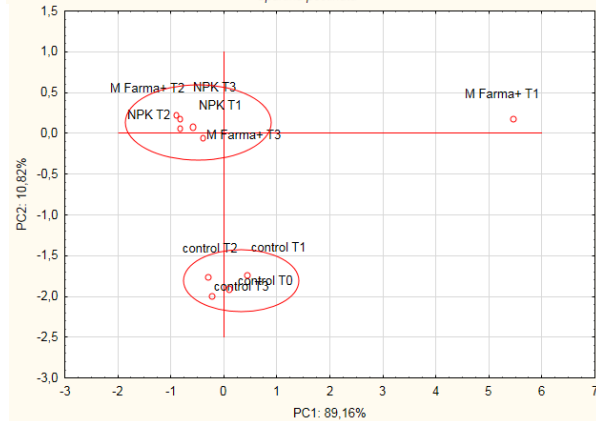
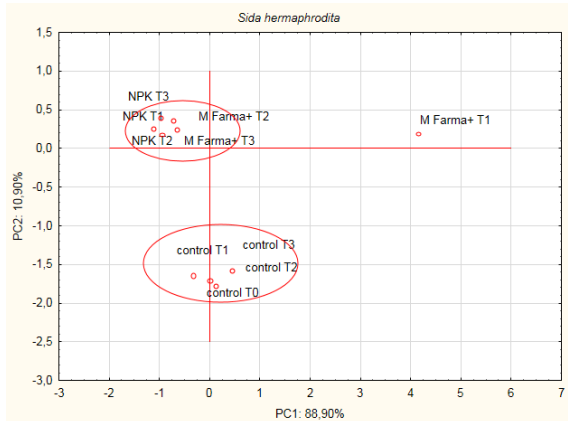
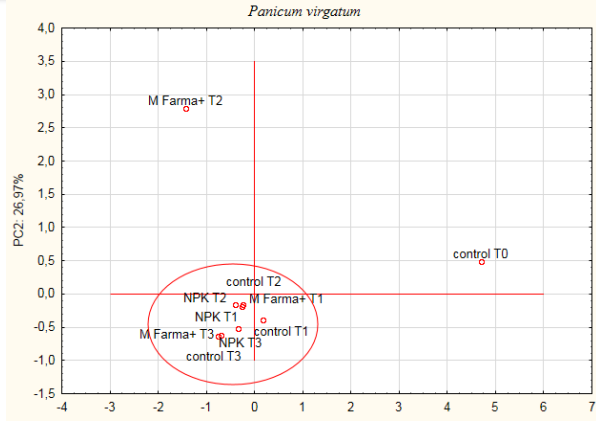
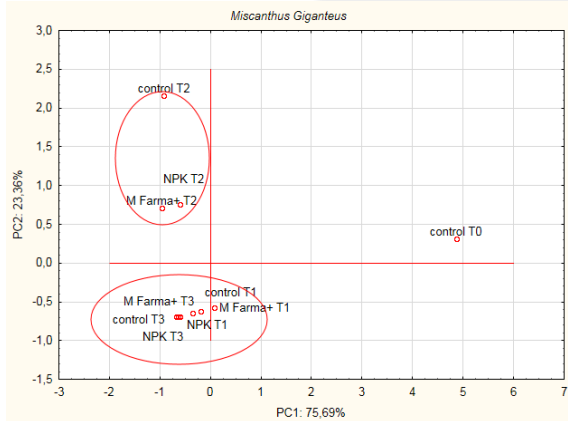
- AWCD (average well-color development) - $\text{AWCD} = \sum \text{OD}_i / 31$
- Shannon-Weiner functional diversity index - $H = - \sum p_i (\ln p_i)$, where p_i is the ratio of the activity on each substrate (OD_i) to the sum of activities on all substrates $\sum \text{OD}_i$
- Shannon Evenness (E) index - was calculated from Shannon-Weiner diversity index (H) and substrate richness (S) index as follows: $E = H / \ln S$
- AUC (Area Under the Curve) - $\text{AUC} = \sum (A_n + A_{n+1}) / 2 \times (t_{n+1} - t_n)$

Statistical analysis - principal components analysis (PCA) and cluster analysis (nearest neighbor method with Euclidian distance)









Thank you for your attention

