

Optimization issues in phenotyping plateforms

Marian Brestic, André Rossi, Oksana Sytar, Marek Zivcak

▶ To cite this version:

Marian Brestic, André Rossi, Oksana Sytar, Marek Zivcak. Optimization issues in phenotyping plateforms. 20ème congrès annuel de la société Française de Recherche Opérationnelle et d'Aide à la Décision (ROADEF 2019), Feb 2019, Le Havre, France. hal-02308114

HAL Id: hal-02308114

https://hal.science/hal-02308114

Submitted on 8 Oct 2019

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Optimization issues in phenotyping plateforms

Marián Brestič¹, André Rossi², Oksana Sytar¹ and Marek Živčák¹

¹ Slovak Agricultural University in Nitra, 949 01 Nitra, Slovak Republic marian.brestic@uniag.sk, oksana.sytar@gmail.com, marek.zivcak@uniag.sk
² Université Paris-Dauphine, PSL, LAMSADE, F-75016 Paris, France andre.rossi@dauphine.psl.eu

Mots-clés: Biology, Phenotyping plateforms, Scheduling, Optimization

1 Introduction

The combined effects of climate change and population growth have made agriculture appear as one of the most important challenge of the century. In order to be able to produce food for everyone in the coming decades, there is an urgent need to build more knowledge about plants, and how to grow them in a sustainable way, i.e., how to produce enough biomass without resorting to pollutant chemicals inputs. Hence, plant biologists need to understand what makes plants resistant to diseases, parasites, insects and hydric stress, among other environment effects. The functional plant body, called the phenotype, is the result of the combination of the plan genotype with the environment: two plants with the same DNA will not have the same development under different environment conditions. As the so-called *plant* performance is the result of phenotype, a lot of effort has recently been devoted to study plants phenotype. More specifically, biologists run large scale experiments on hundreds or thousands of plants, for example to determine how resistant is a particular plant to a wide range of environment conditions. To this end, the plants are settled in pots, and these pots are moved by a conveyor belt from a cultivation chamber to different quantitative automated measurement stations, where RGB cameras, spectrography and other techniques are employed to monitor plants growth on a regular basis. The European project EPPN2020 [1] aims at gathering a large number of such facilities to share experience and accelerate the research progresses in plant biology. It currently involves 31 plant phenotyping platforms and 21 partners across Europe from 12 countries (including the Slovak University of Agriculture in Nitra), under the coordination of INRA, with a budget of €10 million for four years. It can be observed that the phenotyping platforms available in EPPN2020 and everywhere else in the world have a very similar structure (see Figure 1).

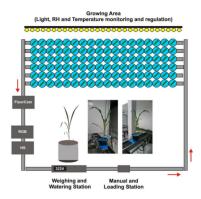


FIG. 1 – A synoptic view of the phenotyping platform in Nitra

2 Optimization problems

The exploitation of these phenotyping platforms poses some challenges that are analogous to the issues raised by traditional manufacturing or production systems, along with some specific features. We present below a list of problems that have been identified in the phenotyping platforms.

2.1 Maximizing the throughput

Phenotyping platforms are sometimes called *high-throughput* phenotyping platforms, for conveying the idea that they should be able to perform a many measurements in a short amount of time. Hence, as a first approach, a phenotyping platform can be modeled as a permutation flowshop. Indeed, the conveyor belt system does not allow a pot to overpass the pot that is located ahead of it on the conveyor belt, and since no by-passes are available, all the plants visit all the stations in the same order. As the biologists sometimes do not require all the plants to undergo all the measurements, a possible problem variant is to decide which plants to send to which measurement stations while fulfilling the minimum measurement requirements.

2.2 Ensuring an equal lighting to all plants

Plants development is very sensitive to light, so when comparing different kinds of environmental conditions, it is important to ensure that all the plants have received the same amount of light. Even if the phenotyping platforms in greenhouses like the one in Nitra are equipped with controlled agriculture-specific LEDs light systems, all the plants standing in the cultivation chamber do not receive the same amount of light, with differences that can reach 30%, which is sufficiently large to bias the results of the experiments. This is due to the fact that upgrading the lighting system would be very expensive, and would increase the energy consumption of the phenotyping platform. This problem is usually addressed by moving the plants randomly in the cultivation chamber during the night [2], which is considered a satisfactory approach only if applied over a sufficiently long period of time. Here also, optimization methods can be used to minimize the maximum gap between the amount of light received by any pair of plants in the system.

3 Conclusions et perspectives

The scheduling problems identified in this work will be presented in more details during the conference, along with some solutions applied to real instances.

4 Acknowledgment

This work was supported by a grant from Campus France, under the scheme of a Partenariat Hubert Curien Stefanik number 40491QC and corresponding Slovak project APVV-SK-FR-2017-0007.

Références

- [1] EPPN2020. https://eppn2020.plant-phenotyping.eu/
- [2] C. Brien, B. Berger, H. Rabie and M. Tester. Accounting for variation in designing green-house experiments with special reference to greenhouses containing plants on conveyor systems. *Plant Methods*, 9(5220), 2013.