



A Study on High Throughput Phenotyping for Vegetable Crops

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Abstract

Conventional phenotyping approaches for vegetable crops such as Solanaceae, Bulb, and Root crops have contributed significantly to the development of numerous varieties. Despite this, traditional phenotyping procedures are insufficient because of the longer time required to produce a variety, poor genetic gain, environmental influences, and other externalities that impact phenotype-based selection. A novel recent approach of high throughput phenotyping (HTP) is regarded a potential tool for addressing the problems of traditional phenotyping. The advent of sensor, computer vision, automation, and sophisticated machine learning technologies sparked the creation of high-throughput phenotyping technology in the prior decade. HTP platforms are being used to conduct non-destructive evaluations of the whole plant system in a variety of crops. HTP provides precise measurements and suggests the collection of high-quality and accurate data, which is required for standardizing phenotyping for genetic dissection and genomic assisted breeding techniques such as genome-wide association studies (GWAS), linkage mapping, marker-assisted selection (MAS), and genomic selection (GS). The rest of this chapter examines the application of high-throughput phenotyping tools in genomic-assisted breeding for vegetable crops.

Keywords; Vegetables, High throughput phenotyping, Genomic assisted breeding

Introduction

The availability of resources for farmers is minimal, making crop management strategies that maximize crop output difficult to implement. Agriculture systems that are over managed may be detrimental to a sustainable agricultural system. While improved genetics, management, and environmental adaptations contributed to the increased production of major commodity crops, quantifying their relative contributions is difficult due to the environment's complex interactions and dynamic nature and management practices. Crop managers face significant challenges in maintaining a consistent and high-yielding crop production level in an unexpected climate, as crop management tactics rely heavily on prior practices. The discrepancy between potential and actual agricultural yields may be significant for certain crops. It is suggested in studies that yield may be enhanced when both best-adapted variety and agronomic practices are applied in the field.

However, the improvement of the genetic structure of plants increased the complex trait like yield. To maximize agricultural productivity, crop management practices must address a variety of practical constraints. Moreover, the advancements in breeding technology continue to promote yield gains in staple crops globally.

Conventional phenotyping techniques are prohibitively expensive, time-consuming, slow, and frequently harmful, and they only allow for the analysis of a few variables at a time. However, traditional breeding operations are being transformed into more efficient contemporary breeding programs by incorporating emerging technologies, most notably high-throughput phenotyping.

A new technology, non-destructive phenotyping, adds a new dimension to the data collection process by increasing the precision, speed, and analysis of captured data. According to scientists, agricultural productivity is expected to increase significantly shortly due to genetic enhancement enabled by high-throughput phenotyping technologies. Sensing technology, data processing, and analysis advancements have significantly improved field and crop management tactics. Apart from focusing on a variety of traits that indicate the water content, chlorophyll content, biomass, and growth potential of a plant.

However, the development of novel vegetable varieties and target environments poses significant challenges in terms of high-throughput and precision phenotyping, modeling, and collaboration with vegetable breeders. Accurate phenotyping was required for several aspects like physiological, morphological, structural, biochemical and molecular characteristics to develop high-yielding vegetable cultivars that were more resistant to biotic and abiotic stresses. As a result, breeders can conduct multiple trials under various growth conditions and with a variety of lines to map populations, breed populations, mutant populations, and the germplasm pool. The remainder of this chapter discusses how high-throughput phenotyping technologies can be used to optimize breeding operations in genomic assisted breeding for vegetable yield gains.

What is High-Throughput Vegetable Phenotyping?

One hundred years ago (Johannsen 1903, 1911), The term "phenotype" was coined as a counterpoint to the concept of "genotypes", and refers to a collection of methodologies and processes for accurately assessing plant growth, architecture, and composition at various sizes. Historically, plant breeders have analyzed hundreds to thousands of plant phenotypes using visual observations, manual tools and among other techniques. Information on the tools for HTP in vegetables is provided in Figure 1.

A fully functional HTP system is composed of supporting hardware (sensors and platforms) and a computing component that communicates with one another (data process and analytics). The research will analyze and integrate a variety of advanced imaging techniques commonly used in computed tomography (CT), into HTP systems in this rapidly growing market. While the industrial sector is driving sensor technology advancements, efforts are being made to incorporate them into agricultural high-throughput systems (AHTP).

The data processing and the analytic system is the most critical component of an HTP system. The current generation of HTP systems, particularly those with high-resolution imaging capabilities, can collect multidimensional data on crops from a large number of people. On the other hand, researchers will quickly discover that they are capable of being overwhelmed by massive amounts of data. In conjunction with ongoing community initiatives, HTP technology has the potential to play a critical role in resolving the breeder's dilemma and expediting the development of new crop varieties with advanced traits.

A uniform set of criteria for assessing agricultural qualities in multiple dimensions can be achieved using equipment such as spectrum reflectance, photogrammetry, and computer vision. Timely and accurate measurements of agricultural characteristics High-throughput phenotyping devices enable breeding programs to increase their capacity to manage a larger breeding population while maintaining the same level of selection intensity. For example, HTP platforms based on unmanned aerial systems (UAS) could be used to rapidly scan breeding grounds. Advanced sensors capture information about the crop that the human eye or senses are unable to see or perceive.

Advanced data analytics and artificial intelligence models extract previously undiscovered information from human and sensor data, and they hold enormous promise for identifying novel agricultural characteristics. The additional elements can be used to characterize plant performance during a particular developmental stage (for example, emerging, blooming, or harvesting) or to assess crop dynamic responses to environmental changes over the course of a growing season. Along with increasing the amount of data available for assessing minute genetic differences between genotypes, unique crop characteristics have the potential to increase genetic diversity within the crop population.

Nowadays, genetic studies on QTL mapping and genome-wide association studies (GWAS) may be used to identify critical genetic variables underlying or associated with yield increase using HTP-based phenotypes. By utilizing marker-assisted selection (MAS) during breeding, it is possible to improve the incorporation of genetic characteristics associated with desirable agricultural characteristics into the existing vegetable germplasm. Integrating breeding populations enables more precise selection, shorter breeding cycles, and increased genetic gain. Large-scale phenotyping enables the collection of massive amounts of agricultural data with high spatiotemporal resolution and the identification of novel crop characteristics. This technique enables the integration of crop and environmental data, as well as management data. Prescriptive phenotyping allow the breeders to develop crop qualities in response to the breeder and consumer requests.

High-Throughput Phenotyping (HTP) Platforms for Vegetable Crops

High throughput phenotyping is a non-destructive technology that creates a good way for measuring the plant phenotype under laboratory and field circumstances. Using sophisticated automation and robotics, imaging (2D and 3D) methods, innovative sensors, hardware and software, these systems monitors a range of plant growth and development aspects. HTP is based on real-time monitoring of plant growth and development in smart glasshouses and physiological and biochemical reactions. Along with plant growth rates and biomass accumulation, the visible imaging system quantifies a range of characteristics, such as canopy architecture and phenology. On the other hand, a hyperspectral imaging system can identify internal properties such as sugar, starch, protein, and moisture content, as well as a range of factors associated with stress. Multiple pictures at varying time intervals and wavelengths are acquired by HTP devices to create data for software-based analysis.

High-Throughput Phenotyping in Genomic Assisted Breeding

The advanced technology of high throughput phenotyping has been successfully used for rapid evaluation of plants traits in glass houses and controlled conditions. Both the academic and industrial sectors have attempted to develop high-throughput screening (HTS) technologies to adapt various crops including vegetables for a variety of breeding purposes. Jansen and colleagues (2009) developed the GROWSCREEN FLUORO to assess stress

tolerance in rosette plants using leaf growth and chlorophyll fluorescence characteristics. Flood, et al. (2016) developed the Phenovator, which can screen over 1000 *Arabidopsis* plants for photosynthesis, growth, and multispectral reflectance multiple times per day. These studies aimed to decipher agriculture's chronological evolution and assess crop's genetic responses.

In vegetable studies, image features are frequently used to replace manual measurements and increase data collection efficiency (phenotyping), or in conjunction with genomic analyses such as quantitative trait loci (QTLs) and genome wide association studies (GWAS) mapping to evaluate genetic variation in crops or to predict crop performance. Several quantitative traits are identified in vegetables crops.

High-Throughput Phenotyping under Controlled Condition

A regulated environment is frequently defined in plant science as an enclosed enclosure in which certain environmental variables such as light condition, temperature, humidity, temperature and CO₂ level are controlled and monitored. Greenhouses, growth chambers, temperature chambers, and nursery rooms are only a few facilities often used in plant research to investigate plant responses to controlled environmental conditions. In controlled environments, plant phenotyping systems are made of sensors, automated control systems, data processing, management systems, and computer software that all work in concert to provide results. The controlled environment is smaller (diameters) and more equipped than natural habitats, which simplifies the deployment of automated phenotyping devices much more than in the wild. These systems collect data on agricultural attributes in a high-throughput manner through the use of sensors, automation, and control systems. The current state of high-throughput plant phenotyping systems in controlled environments was discussed and the sensors used to assess plant characteristics in such systems.

Root Phenotyping

Although the root system dictates the positioning of roots in the soil, little is known about the roots of plants when they are not in the soil. It is critical to understand the anatomical properties of roots in order to appreciate water transport, nutrient absorption, root carbon costs, and root interactions with microorganisms such as mycorrhizal fungi. Finally, the most unclear root phenes are those that are reliant on physiological and flux-related processes. When it comes to root research, they are seldom quantified and have received far less attention in "high-throughput" settings than in traditional ones. According to current thought, the physiological phenes of roots represent a vast and unexplored frontier in root research. Roots are notoriously difficult to analyze. As a result of this difficulty, the genetic and functional foundations of root phenes are less established than those of aboveground phenes.

To close this "phenotyping gap," a shift away from traditional phenotyping toward image-based phenotyping has occurred, which enables relatively high throughput while maintaining root measurement accuracy. Numerous platforms make use of two-dimensional imaging via cameras and propagate plants via seedlings on agar plates, germination paper, or fabric cloth in bins. Additionally, readers are encouraged to peruse this manual. Even though controlling environmental factors is advantageous for characterizing root phenotypes, this chapter focuses on strategies applicable to field-grown plants. The integration of root phenes and functional phenomics will need the phenotyping of several root phenes at the same time. It is anticipated that standard approaches will be tested in the field, which will address the root

cause of the "phenotyping gap." We must dig deeper and harder to fully realize the promise of roots for agricultural revolutionization.

Conclusion

In summary, agricultural HTP technology has the ability to solve the breeder's equation for maximum genetic gain by increasing the intensity and precision of selection, improving the detection of genetic variations, and decreasing breeding cycles. Crop HTP technology is a multidisciplinary and comprehensive approach that integrates research in agronomy, information science, engineering sciences, and biology. Additionally, it leverages cutting-edge computer and artificial intelligence technologies to provide a more comprehensive solution. Numerous advanced data analysis techniques (e.g., machine learning, deep learning) are being used to examine the different phenotypic data available for crops and develop predictive and prescriptive models for crop phenotyping in a highly automated, multi-dimensional, big-data environment. This section will provide the most current information on HTP technology and its applications in plant breeding, genetics, genomics assisted breeding, and some case studies to assist future researchers in developing and improving HTP technology.

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