

Sensor-based phenotyping of above-ground plant-pathogen interactions

Tanner, Florian; Tonn, Sebastian; de Wit, Jos; Van den Ackerveken, Guido; Berger, Bettina; Plett, Darren

DOI

[10.1186/s13007-022-00853-7](https://doi.org/10.1186/s13007-022-00853-7)

Publication date

2022

Document Version

Final published version

Published in

Plant Methods

Citation (APA)

Tanner, F., Tonn, S., de Wit, J., Van den Ackerveken, G., Berger, B., & Plett, D. (2022). Sensor-based phenotyping of above-ground plant-pathogen interactions. *Plant Methods*, 18(1), Article 35.
<https://doi.org/10.1186/s13007-022-00853-7>

Important note

To cite this publication, please use the final published version (if applicable).
Please check the document version above.

Copyright

Other than for strictly personal use, it is not permitted to download, forward or distribute the text or part of it, without the consent of the author(s) and/or copyright holder(s), unless the work is under an open content license such as Creative Commons.

Takedown policy

Please contact us and provide details if you believe this document breaches copyrights.
We will remove access to the work immediately and investigate your claim.

REVIEW

Open Access



Sensor-based phenotyping of above-ground plant-pathogen interactions

Florian Tanner^{1*}, Sebastian Tonn^{2†}, Jos de Wit³, Guido Van den Ackerveken², Bettina Berger¹ and Darren Plett¹

Abstract

Plant pathogens cause yield losses in crops worldwide. Breeding for improved disease resistance and management by precision agriculture are two approaches to limit such yield losses. Both rely on detecting and quantifying signs and symptoms of plant disease. To achieve this, the field of plant phenotyping makes use of non-invasive sensor technology. Compared to invasive methods, this can offer improved throughput and allow for repeated measurements on living plants. Abiotic stress responses and yield components have been successfully measured with phenotyping technologies, whereas phenotyping methods for biotic stresses are less developed, despite the relevance of plant disease in crop production. The interactions between plants and pathogens can lead to a variety of signs (when the pathogen itself can be detected) and diverse symptoms (detectable responses of the plant). Here, we review the strengths and weaknesses of a broad range of sensor technologies that are being used for sensing of signs and symptoms on plant shoots, including monochrome, RGB, hyperspectral, fluorescence, chlorophyll fluorescence and thermal sensors, as well as Raman spectroscopy, X-ray computed tomography, and optical coherence tomography. We argue that choosing and combining appropriate sensors for each plant-pathosystem and measuring with sufficient spatial resolution can enable specific and accurate measurements of above-ground signs and symptoms of plant disease.

Keywords: Plant disease, Phenotyping, Imaging sensors, Plant-pathogen interactions, Biotic stress, Signs and symptoms

Background

Worldwide yield losses in major crops due to pathogens and pests are estimated to be 17–30% [1]. In this review, we will focus on plant pathogens, i.e., organisms or biotic agents that can cause disease [2] and not on pests such as insects and nematodes. Plant pathogens belong to various taxa including viroids, viruses, phytoplasmas, bacteria, oomycetes and fungi [3]. When a pathogen interacts with a plant, structural, physical, and biochemical changes can occur in both the plant and the pathogen. Depending on plant genotype, pathogen strain, and environmental

conditions, the outcome of plant-pathogen interactions (PPI) may be disease, a physiological disturbance of the plant [3–5].

Disease resistance breeding and precision agriculture are key strategies to reduce yield losses due to plant disease in a sustainable way. Both rely on detection, identification and quantification of plant disease on various scales. In disease resistance breeding and pre-breeding, PPI are examined at the cell, tissue, whole plant and field plot level. Zooming in to the cell or tissue level can uncover the distinct mechanisms that determine plant resistance or susceptibility, and precise quantification of disease or resistance levels in whole plants or field plots aids the selection of the best genotypes. In precision agriculture, early and precise disease detection in the field enables efficient crop protection, e.g. by targeted

*Correspondence: florian.tanner@adelaide.edu.au

†Florian Tanner and Sebastian Tonn contributed equally to this work

¹ Australian Plant Phenomics Facility, School of Agriculture, Food and Wine, University of Adelaide, Urrbrae, SA, Australia

Full list of author information is available at the end of the article



© The Author(s) 2022. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

pesticide application or eradication of diseased plants. The challenge to detect and quantify plant disease in an unbiased and precise way initiated the field of plant disease phenotyping [6–8].

In general, “plant phenotyping” describes the study of the manifestation of a genotype under specific environmental conditions [9]. In the context of PPI the phenotype consists of changes that can be described as contrasting indications of disease: signs and symptoms. Whereas these terms were originally used for changes that are visible to the human eye, here we will use them also for changes that can be detected by non-invasive sensors.

Following the American Phytopathological Society (APS) Illustrated Glossary of Plant Pathology, a “symptom” is an indication of disease by reaction of the host [2]. These plant reactions include changes to pigmentation (e.g. necrosis, chlorosis), primary and secondary metabolism, and thermal energy dissipation (Fig. 1). A “sign” is an indication of disease from direct observation of a pathogen or its parts (e.g. sporulation, formation of fruiting bodies, mycelium, bacterial ooze) [2].

While the signs and symptoms are specific for each plant-pathosystem and influenced by environmental conditions, we classify them by their shared characteristics for this review. In practice, signs and symptoms most often do not appear in isolation but occur simultaneously. For example, chlorosis, necrosis, and sporulation may successively co-occur in the same area of an infected leaf.

Phenotyping of PPI can be addressed with invasive methods. For example, colonization of a plant leaf by a pathogen can be detected, classified and quantified by quantitative polymerase chain reaction (qPCR) or for bacterial pathogens by measuring colony-forming units in a homogenate. Such invasive methods can be precise and objective. However, they are necessarily destructive

and limited in speed and scalability, limitations that can be overcome by non-invasive sensors.

Non-invasive sensing offers the possibility of time-course measurements, higher throughput and lower costs [8, 10, 11]. The classic approach for non-invasive phenotyping is visual inspection by humans. This can yield precise and accurate estimates if raters are well trained and appropriate scales are used. However, visual estimates are prone to subjectivity, offer limited speed and scalability, are often qualitative rather than truly quantitative, and are innately limited to the visible spectrum of light [12]. Sensor-based non-invasive phenotyping has the potential to increase throughput and precision, and can detect disease signs and symptoms that are invisible to the human eye [7]. Essentially, non-invasive sensors capture the changes in interactions between electromagnetic radiation and matter (Fig. 2).

The most established sensors for non-invasively measuring PPI are red–green–blue (RGB), hyperspectral, thermal, and fluorescence sensors (Table 1). Less often, monochrome sensors, Raman spectroscopy, and tomographic sensors have been used [6, 13–16].

By focusing here on the biological aspects of disease phenotyping, namely the signs and symptoms, we highlight the sensor-based technologies that are most suitable for specific plant-pathosystems. We group the signs and symptoms into five generalised categories (Fig. 1 and Table 2) to illustrate the common biological processes that underlie sensor-detected signals, to highlight similarities between different plant-pathosystems and to

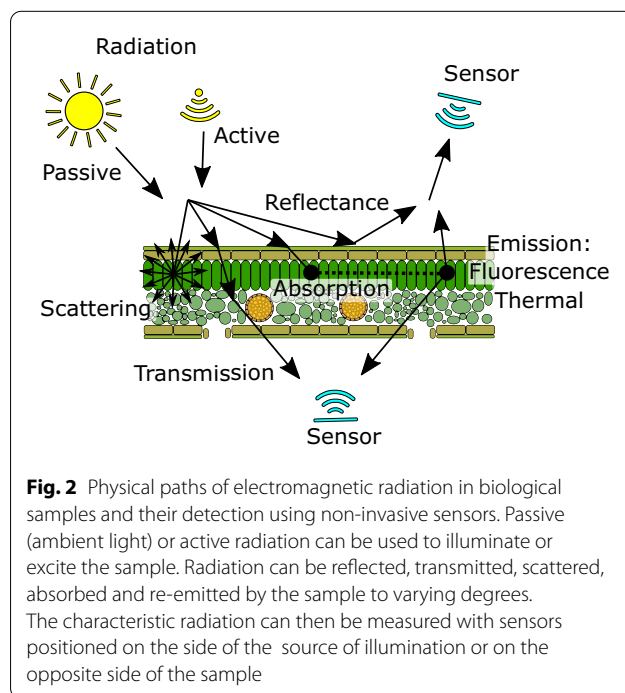
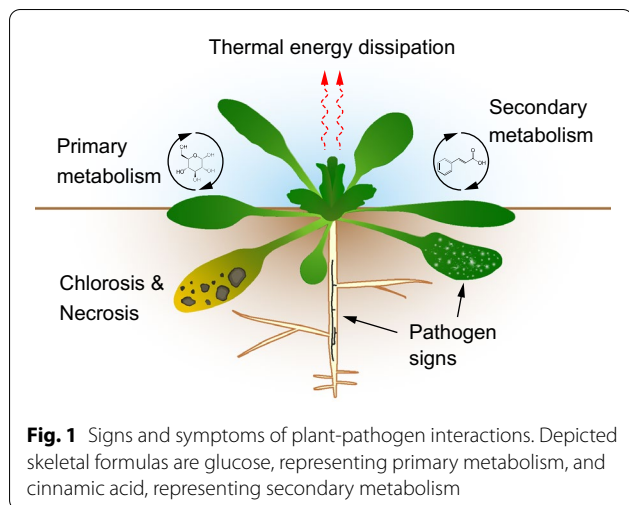


Table 1 Summary of sensors that have been used for phenotyping PPI

Sensor, technology	Imaging/non-imaging	Active/passive	Effect measured	Excitation/illumination wavelengths	Measured wavelengths
Monochrome	Imaging	Mainly active	Reflectance	Variable	Variable
RGB	Imaging	Mainly active, passive at large scale	Reflectance	Variable, usually visible spectrum	Range: ~400–700 nm R: ~600 nm G: ~530 nm B: ~460 nm
Hyperspectral	Both	Mainly active, passive at large scale	Reflectance, transmission	Variable	400–2500 nm
Thermal	Mainly imaging	Passive	Emission	NA	8–15 μ m
Chlorophyll fluorescence (kinetics)	Imaging/non-imaging	Active	Emission	400–700 nm	~650–800 nm
Fluorescence	Imaging/non-imaging	Active	Emission	Mostly 300–400 nm	Mainly 400–700 nm
Raman spectroscopy	Non-imaging	Active	Inelastic scattering of photons (Raman scattering)	Variable, often 785–830 nm [17]	Raman bands, 400–2133 cm^{-1} [18]
Optical coherence tomography	Imaging	Active	Reflectance of coherent light	800–1000 nm or 1200–1400 nm	800–1000 nm or 1200–1400 nm
X-ray computed tomography	Imaging	Active	Attenuation, phase shift	~0.01–0.1 nm	Visible light using scintillator

thereby point out possibilities to transfer and adapt phenotyping solutions (Table 3).

Signs, symptoms and sensors

Pathogen signs

After successful infection, plant pathogens propagate on or inside their host plant, either by rapid replication (e.g. bacteria, viruses) or mycelial growth (fungi, oomycetes) [4]. Quantifying the pathogen, based on signs like spores or mycelium, is a direct measure of plant resistance, defined in a strict sense as the ability of the host to restrict pathogen growth [19]. When a pathogen grows on the surface of the plant, sensors that directly capture the optical changes caused by its physical presence can be used for non-invasive measurement (Table 3).

Powdery mildews are surface-colonizing pathogens representing a variety of obligate biotrophic fungi that can cause disease on various host plants. Growing on the surface, these fungi only penetrate epidermal cells and use haustoria to acquire nutrients. RGB imaging was successfully applied to quantify pathogen signs both for grapevine powdery mildew (*Erysiphe necator*) and cereal powdery mildew infection (*Blumeria graminis* spp.) [20, 21]. In both studies, mycelial growth on detached leaf pieces was imaged in automated phenotyping systems. For grapevine powdery mildew, this system included a movable stage and a DSLR camera [20]. For cereal powdery mildew, a monochrome charge-coupled device (CCD) camera was combined with narrow-bandwidth illumination and a robotic arm system. The

best correlation to visual estimates of infected area was achieved with a simple segmentation algorithm that uses the minimum of the three RGB values [21]. While the throughput of imaging can be easily increased compared to visual scoring for both cereal powdery mildew and grapevine powdery mildew, the preparation of leaf samples remains a bottleneck for these plant-pathosystems.

Unlike powdery mildews, many other filamentous pathogens like fungi and oomycetes form signs on the surface of the host only at the end of the disease cycle, in the form of spore bearing structures. Sporulation of *Cercospora beticola*, a polycyclic necrotrophic ascomycete fungus, occurs on the leaves of infected sugar beet in the area of the necrotic lesions. Hyperspectral microscopy was used to show that sporulation is correlated to an overall decrease in reflectance in the area of lesions in the spectral range of 400–900 nm [22]. However, the advantage of hyperspectral images over RGB images in this study is unclear since the proposed trait is the difference of the integral of reflectance over the entire spectral range of the camera. It would be interesting to know whether the difference in reflectivity over a narrower wavelength range (e.g. one of the RGB channels) could match or improve the quantification of fungal sporulation.

An RGB flatbed scanner was used to assess the interaction between a panel of 335 wheat cultivars and *Mycosphaerella graminicola* (Septoria tritici blotch) [23]. Leaves were collected from a field trial with natural infection and scanned. From the scans the density, size and

Table 2 Applications of sensor-based phenotyping of PPI

Sign/symptom	Plant	Pathogen	Sensor / Vector	Scale	Reference
Pathogen signs	<i>Arabidopsis thaliana</i> , <i>Hordeum vulgare</i>	<i>Fusarium graminearum</i> expressing GFP	Zeiss Stemi-C dissecting microscope with a 470 nm excitation filter and 535 nm emission filter	Whole plants (At), detached spikes (Hv)	[26]
	<i>Beta vulgaris</i> L.	<i>Cercospora beticola</i>	Hyperspectral microscope (PFD V10E), motorized stage	Individual lesions	[22]
	Cereals	<i>Blumeria graminis</i>	Monochrome CCD sensor, 4 channels captured	'Macrobot', robotic arm	[21]
	<i>Nicotiana benthamiana</i>	<i>Phytophthora infestans</i> expressing RFP	Monochrome camera with filter wheel and excitation lights (Patho-Screen imaging system, Phenovation, the Netherlands)	Detached leaves	[28]
Primary metabolism	<i>Nicotiana tabacum</i>	<i>Tobacco mosaic virus</i> expressing GFP	RGB camera under UV illumination (Blak-Ray Model B 100AP)	Detached leaves	[27]
	<i>Phaseolus vulgaris</i>	<i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> expressing <i>lux-eYFP</i> operon	nightOWL LB 983 in vivo imaging system (Berthold Technologies, Germany), confocal laser scanning microscope Zeiss LSM 880	Detached leaves	[34]
	<i>Triticum aestivum</i> L.	<i>Fusarium graminearum</i>	X-ray CT, Biomedical Imaging and Therapy beamline (BM11-BM, 05B1-1)	Spikelet	[114]
	<i>Vitis</i> spp.	<i>Erysiphe necator</i>	RGB (Nikon D850), automated motorized stage	Individual leaf discs, automated imaging	[20]
	<i>Arabidopsis thaliana</i>	<i>Pseudomonas syringae</i>	CF Imager (Technologica Ltd., UK), NPQ, fPSII, Fv/Fm	Individual plants	[149]
	<i>Cucumis melo</i>	<i>Podosphaera xanthii</i>	Open FluorCam 700 MF (Photon System Instruments), NPQ, fPSII	Individual leaves	[54]
	<i>Hordeum vulgare</i>	<i>Blumeria graminis</i> f. sp. <i>hordei</i>	Chlorophyll Fluorometer IMAGING-PAM M-series (Walz, Germany)	Individual leaves	[150]
	<i>Lactuca sativa</i>	<i>Biernia lactucae</i>	Open FluorCam 700 MF (Photon System Instruments), Fv/Fm	Leaf discs	[151]
	<i>Nicotiana benthamiana</i>	Pepper mild mottle virus	FluorCam (Photon System Instruments), NPQ, fPSII	Individual leaves	[152]
	<i>Olea europaea</i>	<i>Xylella fastidiosa</i>	Micro-hyperspectral imager (VNIR model, Headwall Photonics, USA), 400 – 885 nm, from aircraft 500 m above ground	Orchards	[48]
	<i>Oryza sativa</i>	<i>Rhizoctonia solani</i>	Non-imaging NeoSpectra micro handheld spectrometer (SiWare Systems, Canada), 1348–2551 nm	Individual leaf spots (non-imaging)	[61]
	<i>Solanum tuberosum</i>	<i>Phytophthora infestans</i> , <i>Alternaria solani</i>	Non-imaging field spectrometer SVC HR-1024i (350–2500 nm) (Spectra Vista Corporation, USA)	Individual leaf spots (non-imaging)	[153]

Table 2 (continued)

Sign/symptom	Plant	Pathogen	Sensor / Vector	Scale	Reference
Secondary metabolism	<i>Beta vulgaris</i>	<i>Cercospora beticola</i>	SWIR spectral camera, 970–2500 nm (HySpex SWIR-320 m-e line camera, Norsk Elektro Optikk A/S, Norway)	Detached leaves	[154]
	<i>Citrus sinensis</i> , <i>Citrus paradisi</i>	<i>Candidatus Liberibacter</i> spp.	Non-imaging handheld Raman spectrometer (Resolve spectrometer equipped with 831-nm laser source, Agilent, USA)	Detached leaves	[86, 87, 90]
	<i>Hordeum vulgare</i>	<i>Blumeria graminis</i> f.sp. <i>hordei</i>	UV line scanner, 250–500 nm (Head-wall Photonics)	Detached leaves	[81]
	<i>Nicotiana benthamiana</i>	Pepper mild mottle virus	Excitation with xenon-lamp + BP 340/75, imaging with CCD camera + BP 440/20 and BP 520/20	Individual leaves	[152]
	<i>Triticum aestivum</i>	<i>Puccinia tritici</i> , <i>Blumeria graminis</i> f. sp. <i>tritici</i>	Non-imaging fiber-optic fluorescence spectrometer (IOM GmbH, Germany) combined with 337 nm pulsed N2 laser	Individual leaf spots (non-imaging)	[73]
Necrosis and chlorosis	<i>Vitis vinifera</i>	<i>Plasmopara viticola</i>	Macroscopic (AZ100 multizoom, Nikon), ex. BP 340/26 and em. LP 371	Leaf parts	[69]
	<i>Arabidopsis thaliana</i>	<i>Pseudomonas syringae</i>	RGB (Nikon D5200 DSLR)	Seedlings growing in well-plates	[117]
	<i>Arabidopsis thaliana</i>	<i>Sclerotinia sclerotiorum</i>	RGB (USB camera, full HD 1080p) controlled by Raspberry Pi 3 Model B motherboards	Detached leaves	[115]
	<i>Beta vulgaris</i>	<i>Cercospora beticola</i>	RGB (camera Baumer HXG-40), multispectral camera (6 bands of 10 nm between 450 and 850 nm, AIRPHEN)	Phenobile 1,50 distance to canopy top (RGB), hexacopter (multispectral)	[119]
	<i>Triticum aestivum</i>	<i>Mycosphaerella graminicola</i>	RGB (flatbed scanner)	Detached leaves collected from field trial	[23]
	<i>Zea mays</i>	<i>Setosphaeria turcica</i>	RGB	Drone, 6 m above ground	[118, 120]

Table 2 (continued)

Sign/symptom	Plant	Pathogen	Sensor / Vector	Scale	Reference
Thermal energy dissipation	<i>Cucumis sativus</i> L.	<i>Fusarium oxysporum</i> f. sp. <i>cucumerinum</i>	FLIR SC620	Individual leaves	[123]
	<i>Ipomoea batatas</i> L.	Sweet potato feathery mottle virus (SPFMV) Sweet potato chlorotic stunt virus (SPCSV)	Top-view thermal camera (FLIR A615), PlantScreen conveyor system, NaPPI, Helsinki	Whole plant	[132]
	<i>Nicotiana tabacum</i> L.	Tobacco mosaic virus (TMV)	Infrared imager (Agema THV900LW), Cartesian positioning system in imaging chamber	Leaves	[131]
	<i>Olea europaea</i> L.	<i>Verticillium dahliae</i>	Temperature sensor (Apogee IRR-P), Fixed 1 m above canopy	Single tree canopy	[129]
Structural changes	<i>Olea europaea</i> L.	<i>Verticillium dahliae</i>	Broad-band thermal camera (FLIR SC655) on crewed aircraft	3000 ha, spatial resolution = 62 cm	[130]
	<i>Capsicum annuum</i>	<i>Stemphylium lycopersici</i>	Laboratory-OCT system, 4096-pixel line scan camera (spl4096-140 km, Basler)	Single leaves	[134]
	<i>Malus domestica</i>	<i>Marssonina coronaria</i>	Backpack-based OCT system, 2048-pixel line scan camera (spl2048-140 km, Basler, Germany)	Single leaves	[133]
	<i>Solanum tuberosum</i>	<i>Streptomyces scabies</i>	Medical X-ray CT scanner (Toshiba Xvision high-resolution CT scanner)	Single plants	[135]
	<i>Triticum aestivum</i>	<i>Fusarium graminearum</i>	Synchrotron-based phase contrast X-ray imaging with the Biomedical Imaging and Therapy beamline (BMIT-BM, 0581-1) at the Canadian Light Source	Single excised wheat spikes	[14]

Table 3 Suitability of sensors for phenotyping PPI

		RGB	Hyperspectral	Thermal	Fluorescence	Chlorophyll fluorescence (kinetics)	Raman spectroscopy	OCT	X-ray CT
Pathogen signs	Controlled	++	+	–	+	–	–	+	+
	Field	++	+	–	–	–	–	+	–
Primary metabolism	Controlled	–	+	–	–	++	+	–	–
	Field	–	+	–	–	++	+	–	–
Secondary metabolism	Controlled	–	+	–	+	–	+	–	–
	Field	–	+	–	+	–	+	–	–
Necrosis and chlorosis	Controlled	++	++	–	+	+	–	+	+
	Field	++	+	–	+	+	–	+	–
Thermal energy dissipation	Controlled	–	–	++	–	–	–	–	–
	Field	–	–	+	–	–	–	–	–

“Not used/unsuitable” (–), “Preliminary” (+) and “Widely used” (++)

melanisation of pycnidia (signs) as well as lesion size (a symptom) were measured. A Genome-Wide Association Study using the phenotypic data identified 26 chromosome intervals associated with *Septoria tritici* blotch resistance. Sixteen of these loci overlapped with intervals that had already been identified based on visual assessment in previous studies, while ten had not been described before, demonstrating the potential power of quantitative phenotyping [23]. This example also illustrates the benefit of a high resolution, e.g. being able to discriminate single pycnidia to determine density and size, thus overcoming the weakness of low spatial resolution of many field phenotyping methods.

Transgenic pathogen strains that express detectable markers such as fluorescent proteins or bioluminescence conferring enzymes are another alternative to track and quantify pathogen growth directly [24, 25]. Fluorescent proteins have mainly been used to study the infection process in vivo at the cell level using epi-fluorescence or fluorescence confocal laser scanning microscopy. But also at the level of whole plants or seedlings, pathogens expressing fluorescent proteins, including bacteria, fungi, oomycetes and viruses have been used to track, image and quantify infection and colonization [26–31]. Plant auto-fluorescence and low fluorescent protein signal intensity can hinder imaging at larger scales. Bioluminescence, so far mainly applied in bacteria, generates a light signal without prior excitation, therefore avoiding plant auto-fluorescence. But the low signal intensity requires imaging with sensitive cameras and extended exposure time in the dark (up to several minutes) [32, 33]. A recent study generated bioluminescent and fluorescent *Pseudomonas syringae* pv. *phaseolicola* [34]. The bioluminescence enabled detection of the bacteria at the leaf scale in a dedicated imaging chamber. Identified colonized plant parts

could then be sampled and further investigated under the fluorescence microscope making use of the expressed fluorescent proteins. Since enzymes producing bioluminescent compounds have also been identified in fungi, such a luminescence based approach might also be feasible to facilitate quantification and examination of fungal infection at the macro- and microscopic level [35]. But all these approaches are restricted by the requirement for both a protocol for genetic transformation of the pathogen of interest and for facilities authorized to carry out experiments with transgenic plant pathogens.

Symptom: changes in primary metabolism

In plant-pathogen interactions, the plant primary metabolism is influenced both by manipulation of the pathogen and the immune response of the plant itself. Pathogen infection may modify source-sink relations in the plant or impair photosynthesis, while plant immune responses require additional resources from the pool of primary metabolites [36–38]. Together this may lead to detectable symptoms based on photosynthetic performance or altered accumulation and allocation of primary metabolites.

Photosynthetic performance can be probed by analyzing chlorophyll *a* fluorescence and the kinetics of chlorophyll *a* fluorescence (Chl-F) [39, 40]. According to the model of photosystem II (PSII) absorbed light energy can take three different paths: (i) drive photosynthesis (photochemical quenching); (ii) dissipate as heat (non-photochemical quenching); (iii) re-emit as fluorescence [39, 41].

Measuring the kinetics of Chl-F, the changes of Chl-F under different light conditions, e.g. before and after a saturating light pulse, allows separation of these components and calculation of diverse parameters that yield information about photosystem II (PSII) photochemistry, electron flux, and CO₂ assimilation [39, 40]. Commonly

used parameters are the maximum quantum efficiency of PSII photochemistry (F_v/F_m), the operating efficiency of PSII photochemistry (Φ_{PSII} , F_q'/F_m' or $\Delta F/F_m$), the level of photochemical quenching of PSII (qP or F_q'/F_v') or the level of non-photochemical quenching (NPQ) which estimates the rate constant for heat loss from PSII [40].

With commercially available Chl-F kinetics imaging systems these parameters can be mapped onto imaged leaves or whole plants, enabling identification of spatial heterogeneity that may be linked to localized pathogen infection [42]. Many of the Chl-F parameters (e.g. F_v/F_m , qP , NPQ) are measured on dark-adapted plants and commonly require light sources in close proximity to the plant to provide e.g. a saturating pulse to measure maximum fluorescence (F_m). Therefore, most Chl-F imaging systems are designed for growth chambers or greenhouses where LED panels can provide even illumination and plants can easily be dark adapted [42]. However, there are also field phenotyping systems that include Chl-F imaging with active illumination [43]. Dark adaptation in the field can be achieved by imaging at night or before dawn. But because the sensor needs to be close to the plants, Chl-F imaging with active illumination in the field is limited to ground-based phenotyping platforms which offer limited throughput compared to uncrewed aerial vehicles (UAV).

Analysis of spectral reflectance under sunlight does not require active illumination and is therefore an alternative for probing photosynthesis that is compatible with aerial vectors like UAVs or aircrafts. Spectral reflectance data can be used to build predictive models for photosynthetic parameters like maximum carboxylation rate of Rubisco or to estimate sun-induced chlorophyll fluorescence [44–47]. For example, sun-induced chlorophyll fluorescence, determined from spectral images taken from an aircraft, has successfully been used to estimate disease severity of olive trees infected with the bacterium *Xylella fastidiosa* [48]. But these approaches are technically challenging, both in data acquisition and data analysis, and interpretation is difficult because the relationship of reflectance, canopy geometrical structure, leaf physiology and variation in solar radiation is not fully understood [47, 49, 50]. So far, these challenges limit applications, despite the potential especially for large scale remote sensing of plant stress.

Chlorophyll fluorescence imaging systems with active illumination on the other hand have been used in numerous studies to monitor the effect of pathogen infection on plants [42]. A common response, in many cases prior to visual changes, is the decrease of Φ_{PSII} resulting from a decreased PSII electron transport as well as an increased heat dissipation rate (NPQ). This has also been observed for infections of many biotrophic pathogens like powdery

mildew of wheat and barley (*B. graminis*), powdery mildew of cucurbits (*Podosphaera xanthii*), downy mildew of lettuce (*Bremia lactucae*) and downy mildew of grapevine (*Plasmopara viticola*) [51–54]. These biotrophic pathogens often induce visible symptoms only at late infection stages, thus Chl-F imaging may be particularly useful to visualize and quantify early colonization.

A general drawback of Chl-F imaging is the lack of specificity as photosynthesis and Chl-F are influenced by many biotic and abiotic stress factors alike [55]. This could be partially overcome by taking into account the differences in spatial patterns of Chl-F changes. Patterns induced by localized pathogen infection might be distinguishable from patterns induced by abiotic stresses that affect the whole plant.

While Chl-F provides information about the current productivity of the plant, it does not allow for quantification of the actual concentration of primary metabolites. Changes in accumulation and allocation of sugars, starch, amino acids or proteins can, in principle, also be estimated directly via reflectance spectroscopy or imaging spectroscopy in the visible (VIS, 400–700 nm), near-infrared (NIR, 700–1000 nm) and shortwave infrared (SWIR, 1000–2500 nm) range [56–58]. Such approaches are based on combining non-invasive spectroscopic measurements with biochemical analysis of the same tissue to build predictive models. For example, Ely et al. [56] developed spectra-trait models for leaf starch, glucose, and protein content based on reflectance spectroscopy (500–2400 nm) and biochemical analysis of leaves of eight crop species. However, the usefulness of such models for linking spectral features to metabolic changes during PPI still requires validation.

So far, studies only indicate that reflectance spectroscopy may sense specific changes in primary metabolism during PPI. Gold et al. [59] collected reflectance spectra (400–2400 nm) with a portable non-imaging contact spectrometer from potato leaves at different infection stages of *Phytophthora infestans* or *Alternaria solani*. Using spectra-trait models they estimated pathogen-induced changes in leaf sugar, starch and nitrogen concentration and found increased sugar concentration during the biotrophic, necrotrophic and sporulation phase of *P. infestans*. But these estimates were not validated by chemical analysis and the applied spectra-trait models were originally developed on data from forests and grasslands [60]. Although the study shows clear differences in reflectance spectra between tissue infected with the different pathogens and tissue at different infection stages, the interpretation of these spectral differences remains unclear.

A similar study in rice with healthy and sheath blight (*Rhizoctonia solani*) affected plants found that differential

spectral regions could be linked to absorption features of starch, cellulose and protein content, although they were not chemically validated [57, 61].

Both of these studies used non-imaging spectrometers and found spectral features in the SWIR range to be important for detection of diseased plant tissue. Therefore, imaging SWIR sensors might be particularly useful to not only measure changes to primary metabolites but also to provide spatial information on these changes.

Symptom: changes in secondary metabolism

Plant secondary metabolites (PSM) are a large group of structurally and functionally diverse metabolites that are, as opposed to primary metabolites, considered non-essential for primary functions like photosynthesis, growth and reproduction [62, 63]. Those PSM that are involved in plant immunity are commonly classified into two groups, phytoanticipins and phytoalexins [64]. While phytoanticipins are constitutively produced and stored in plant tissue, phytoalexins are synthesized in response to a pathogen. Members of both groups show *in vitro* antimicrobial and insect-deterrent activity [65].

Independent of their function, PSM can be informative markers for preformed resistance (phytoanticipins), or a symptom of the infection progress, and magnitude or quality of the plant immune response (phytoalexins). Due to their specific absorption and, in some cases, fluorescence spectra they may be detected non-invasively, e.g. by reflectance and fluorescence spectroscopy or imaging [57, 66].

In grapevine, infection with the downy mildew pathogen *Plasmopara viticola* induces the accumulation of stilbenes, a group of phenolic compounds [67]. Pure stilbenes emit a violet-blue fluorescence around 400 nm when excited with UV light (335 nm) [68]. A fluorescence signal with a similar spectrum was imaged in downy mildew infected grapevine leaves at the cell level using confocal laser scanning microscopy, as well as at the tissue level using epifluorescence macroscopy [68–70]. Mass spectrometry imaging revealed co-localization of stilbenes with the violet-blue fluorescence signal, suggesting that stilbenes are indeed the source or at least contribute to the observed fluorescence [70]. Since stilbene synthase expression has also been linked to downy mildew resistance, the violet-blue fluorescence may not only enable detection and quantification of downy mildew infection but also allow for the assessment of differences in plant defense responses [71].

Blue (around 440 nm) and green (around 520 nm) fluorescence upon UV excitation has also been described in *Nicotiana benthamiana* infected with Pepper mild mottle virus, likely due to accumulation of the phenolic compound chlorogenic acid, and in wheat infected with leaf

rust (*Puccinia triticina*) or powdery mildew (*Blumeria graminis* f. sp. *Tritici*) [72, 73].

These examples indicate that changes in UV-excited blue and green fluorescence, induced by certain pathogens, is a conserved response across plant species. In fact, it has been described as a general conserved stress response, also to abiotic stresses including drought, nutrient deficiencies and increased UV irradiation [74–77]. Responsible fluorophores in most cases are likely stress-induced soluble and cell wall bound phenolic compounds that fluoresce in the blue-green spectrum [78]. For example, the fluorescent stilbenes accumulate in grapevine leaves also in response to prolonged UV-C irradiation [70]. Consequently, distinction between infections of different pathogens, or between biotic and abiotic stress, might not be possible. This is a critical limitation for phenotyping in field trials, where various stresses can occur simultaneously.

Plant secondary metabolite content of leaves may also be estimated via reflectance spectroscopy [79, 80]. Spectral indices or models for leaf traits like phenolic content have been mostly developed and validated for remote sensing in landscape ecology studies [60, 79]. For example, Kokaly and Skidmore [80] proposed that an absorption feature around 1660 nm is related to content of phenolic compounds in different plant species and showed that in fresh tea leaves (*Camellia sinensis*), this absorption feature correlates with total phenolic compound content.

Only a few studies have combined spectral measurements with biochemical analysis of diseased plants. This is required to link spectral features to physiological processes during PPI. Brugger et al. [81] explored spectral imaging in the UV range (250–400 nm), which is particularly interesting because many plant secondary metabolites involved in stress responses feature absorption maxima in that range. They found in barley infected with powdery mildew (*Blumeria graminis* f. sp. *hordei*) that changes in flavonoid content during the first 5 days of infection correlated with reflectance intensity around wavelengths that match flavonoid absorption spectra. But adverse interaction of the sensor with the required UV light source restrains interpretation of these results. Another study combined spectral imaging in the SWIR range (970–2500 nm) with untargeted metabolic fingerprinting of three different sugar beet genotypes infected with *Cercospora beticola* [82]. Although there were correlations between several secondary metabolites and spectral data, it remains unclear if this correlation is due to direct contribution of these metabolites to the reflectance spectrum. Combining imaging spectroscopy with mass spectrometry imaging could help to provide direct links between specific metabolite groups and spectral features.

Raman spectroscopy is another technology to measure changes in plant secondary metabolism. After excitation of the sample with a laser it measures the inelastic scattering of photons (also called Raman scattering), which can provide both qualitative and quantitative information about the chemical composition of the sample [15, 83]. Raman scattering can be collected with portable non-imaging handheld Raman spectrometers [84, 85] and this approach has been applied to detect viral, bacterial and fungal infections in plants [85–89]. For example, a handheld Raman spectrometer was used to detect *C. liberibacter* spp. infection in citrus trees. Using orthogonal partial least squares discriminant analysis, grapefruit leaves were classified into healthy, infected, and nutrient-deficient categories with 98% accuracy in the training set, but the authors did not validate the classification accuracy in a test set [86, 87]. Infection was associated with increased intensity of the Raman band assigned to lignin and phenolic compounds. Correspondingly, a follow-up study found increased *p*-coumaric acid content in infected leaves, a phenolic compound and lignin precursor whose Raman spectrum matches the disease associated bands [90]. Similar disease associated bands, likely corresponding to phenolic compounds, have also been described for virus infection in wheat, tomato and rose [15, 88, 91]. These studies used non-imaging Raman spectrometers, which only provide point measurements and do not yield any spatial information. But Raman spectroscopy can also be combined with digital imaging so that Raman spectra are recorded for each pixel [92]. This has been explored as a tool for quality and safety inspection in food, pharmaceutical, and biomedical sectors and, for example, to detect watermelon seeds infected by the bacteria *Acidovorax citrulli* [93, 94]. So far, the lengthy image acquisition has restricted the throughput and therefore applications of Raman imaging, but Lee et al. [94] report a relatively fast system that requires 250 s to image an area of five by twenty cm with a spatial resolution of 250 by 1024 pixels. Such systems may already be useful for certain phenotyping challenges, but further reducing the acquisition time would widen the range of possible applications.

Symptom: necrosis and chlorosis

Pathogen-induced chlorosis and necrosis are prominent symptoms of plant disease as they are visually evident and very common. Chlorosis results from changes in pigmentation, mainly the degradation of chlorophyll, and necrosis from the death of cells and tissue. Both may occur locally in lesions, with chlorosis often preceding or surrounding necrotic lesions. Viruses can cause chlorosis in diverse patterns that are often reflected in their name

(e.g. “chlorosis”, “mottle”, “mosaic”, “streak”, “vein clearing”, “yellowing”) [95].

Both chlorosis and necrosis can be induced by specific pathogen-produced metabolites or proteins, e.g. the chlorosis inducing coronatine from *Pseudomonas syringae* or the necrosis and ethylene-inducing peptide 1 (Nep1)-like proteins from *Botrytis cinerea* [96–101]. Particularly pathogens with a necrotrophic life style produce toxins that kill plant cells, by disrupting the plant cell membrane directly or by generating membrane damaging reactive oxygen species [102, 103]. In interactions with viruses and biotrophic pathogens like downy mildews, chlorosis is often induced in later stages of the infection, and chlorophyll degradation appears to be regulated by the same plant genes that control regular leaf senescence [104–106]. The area of chlorotic or necrotic tissue can serve as a good proxy to estimate spread of the pathogen, as well as the impact on yield, depending on plant developmental stage, type of the pathogen and stage of the infection.

While chlorotic tissue has reduced chlorophyll content, necrotic tissue lacks all pigments characteristic for healthy plant tissue. This results in changes of absorption and reflectance in the visible spectrum, evident as color ranging from yellow (chlorosis) to shades of brown to black (necrosis). Besides the lack of pigments, necrotic tissue also differs from healthy tissue in water content and three-dimensional structure due to the collapse of cells. A lower ratio of cell surface to intercellular air space due to a collapse of e.g. the spongy mesophyll leads to reduced reflectance of NIR radiation [107]. Changes in water content in necrotic tissue also impacts SWIR reflectance due to several water absorption peaks in the SWIR range [108, 109]. Consequently, sensors that detect reflectance in the VIS–NIR–SWIR range are useful to quantify chlorosis and necrosis. The contrast between healthy leaf tissue and lesions lacking chlorophyll may be enhanced by imaging the red steady-state chlorophyll fluorescence [110, 111]. The onset of cell death can also be visualised by imaging the increased chlorophyll fluorescence that results from the disassembly of the chloroplast thylakoid membrane in tissue undergoing cell death [112]. Additionally, the changes in plant internal tissue structure that precede and are associated with necrosis and tissue damage can be measured with different tomography methods, e.g. optical coherence tomography (OCT) or X-Ray computed tomography (CT) [113, 114].

In controlled experiments, particularly on samples that are easy to image such as detached leaves, RGB imaging is an established method to track and quantify chlorosis and necrosis. Barbacci et al. [115] combined a detached leaf assay with a setup for automated RGB

image acquisition and analysis to quantify necrotic lesion development on *Arabidopsis* inoculated with *S. sclerotiorum*. This enabled measurement of latency period and lesion doubling time (LDT) at high resolution (measurement every 10 min over 36 h) and of large sample sizes (120–270 leaves per imaging unit). Whereas latency period varied mostly between different *S. sclerotiorum* isolates, LDT was mainly determined by the plant genotype and independent of the inoculated isolate. Using the differences in LDT as a robust indicator of quantitative resistance led to the identification of the nucleotide-binding site leucine-rich repeat gene *LAZ5* as a negative regulator of quantitative resistance to *S. sclerotiorum*. On a similar scale, RGB imaging enabled quantification of chlorosis induced by *P. syringae* in *Arabidopsis* seedlings growing in 48-well plates [116, 117]. The assay was tested in a genome-wide association study to efficiently distinguish between presence and absence of effector-triggered immunity and confirmed the loci of known resistance genes.

In field experiments, necrosis in maize infected by *Setosphaeria turcica* (Northern corn leaf blight) or sugar beet infected by *C. beticola* (Cercospora leaf spot) has been assessed by sensors [118, 119]. Wiesner-Hanks et al. [118, 120] acquired images with a RGB camera mounted on a UAV flying 6 m above a maize field trial. The necrotic lesions captured in these images were manually annotated and used to train a CNN. Combining this CNN with a conditional random field method allowed automated segmentation of the aerial images to identify and quantify lesion area.

In a sugar beet field trial, Jay et al. [119] tested both RGB imaging from a ground-based vehicle and spectral imaging (six bands between 450 and 850 nm) from a UAV to assess *Cercospora* leaf spot severity. They determined the necrotic spot density from RGB images and green fraction from both RGB and spectral images. The image data was compared to visual severity scores given by an expert on a 1–9 scale. Spot density gave a better prediction for low (less severe) visual scores and green area was a better predictor for high visual scores. Consequently, combining these two features as input for a neural network enabled a good prediction of the visual disease scores. Because only the ground-based RGB sensor enabled measuring both of these features, it outperformed the aerial spectral sensor.

Symptom: thermal energy dissipation

Plant-pathogen interactions can result in a change of tissue temperature by affecting energy balance terms such as transpiration or light absorption [121]. These induced changes often precede other symptoms and are characterized by complex spatial and temporal

dynamics. This makes thermal energy dissipation an interesting candidate trait for early detection of disease. PPI can cause an increase in tissue temperature by inducing stomatal closure and vascular occlusion. Conversely, damage to cells and deregulation of stomatal opening can lead to decreased tissue temperature through uncontrolled transpiration [122, 123]. Photosynthetic performance has an influence on plant temperature as well because part of the absorbed light energy, if not emitted as fluorescence or converted in photochemistry, is dissipated as heat in the process of non-photochemical quenching [39].

Temperature can be remotely measured in the thermal infrared spectrum (TIR, 8000–15,000 nm) with radiometric sensors [124]. This technology has long been applied to measure abiotic stresses, particularly drought stress which is associated with an increased canopy temperature [125, 126]. Multiple plant-pathosystems have also been studied with thermal sensors [127].

Vascular pathogens that grow within the xylem of host plants can cause occlusion of the vascular system due to both their own growth and the plant responses to the pathogens. This can lead to symptoms similar to drought stress, as the hydraulic conductivity is inhibited by the occlusion [128]. In a field experiment, the crown temperature of olive trees under natural infection with *Verticillium dahliae*, a vascular fungal pathogen, was measured with infrared temperature sensors (Apogee IRR-P) mounted 1 m above the trees. The temperature was positively correlated with *Verticillium* wilt disease severity levels across multiple sites [129]. The same authors validated and vastly increased the throughput of their methods by mounting a broad-band thermal camera (FLIR SC655) on a crewed aircraft and flying it over 3000 ha of olive orchards [130].

The measurement of plant thermal energy dissipation is not easy to measure in the field as environmental effects and the influence of other stresses decrease specificity of the measurements. Therefore, many studies of plant thermal energy dissipation are performed in controlled environments.

An abscisic-acid-induced stomatal closure in leaves of cucumber infected with the vascular pathogen *Fusarium oxysporum* f. sp. *Cucumerinum* could be detected by an increase in temperature with a FLIR SC620 digital infrared camera in a controlled environment [123]. The maximum temperature of leaves of infected plants was reached nine days after infection. The authors also observed a fast decrease of temperature ten days after infection and attributed it to uncontrolled water loss due to cell damage. Eleven days after infection, leaf temperature rose again which was attributed to dehydration of the leaves.

Virus-plant interactions can also influence tissue temperature. Tobacco infected with tobacco mosaic virus shows a fast increase in leaf temperature at the initial infection site preceding a hypersensitive response. This increase in temperature is due to stomatal closure which is induced by salicylic acid accumulation. The same areas of the leaves later appear as necrotic lesions [131]. Changes in leaf temperature were also observed in sweet potato upon infection with two different viruses, Sweet potato feathery mottle virus (SPFMV) and Sweet potato chlorotic stunt virus (SPCSV) [132]. In this controlled environment experiment, leaf temperature was measured with a top-view thermal camera (FLIR A615) and differed between healthy plants, plants infected with SPFMV, and plants co-infected with both viruses. Higher temperatures were associated with higher disease severity scores [132].

Sensing structural changes

Preliminary studies using tomographic sensors for phenotyping PPI have been performed in various plant-pathosystems. As these sensors measure spatially resolved attenuation, refractive index variation and scattering strength inside tissue, they can be useful to non-invasively study internal structural changes which can allow for early detection and potentially identification of disease [113, 114, 133, 134]. Besides measuring internal plant and pathogen structure, X-ray CT scanning also provides the option to measure through substrate and was used to study changes in root morphology of potato affected by *Streptomyces scabies* [135].

Synchrotron-based phase contrast X-ray CT was used to measure differences in tissue degradation in wheat caused by *Fusarium graminearum* [114]. A known resistance mechanism to this fungus is the inhibition of fungal colonization from spikelet to spikelet inside the rachis internode. Traditional histological studies to quantify this type of resistance require destructive sampling. The tissue degradation leads to increased tissue porosity, which makes it possible to sense the pathogen spread by changes in X-ray attenuation.

Leaves of apple trees infected with *Marssonina coronaria* were measured with a custom-built backpack-based OCT sensor in the field [133]. Disease progression causes an enlarged gap between epidermis and palisade parenchyma that could be sensed by a reduction in backscattering. The presence of infection was confirmed using Loop-mediated isothermal amplification, a nucleic-acid based technique [133]. Authors from the same group applied similar techniques in leaves of *Capsicum annuum* infected with *Stemphylium lycopersici* [134].

These studies show the potential of tomographic sensors for early detection of disease as well as phenotyping of below ground structures.

Discussion

Phenotyping plant-pathogen interactions in the field is limited by specificity, canopy structure, and environmental conditions

Signs and symptoms are easier to measure in a controlled environment where plant material can be accessed from multiple angles at close proximity and under optimal illumination. Yet field phenotyping is a requirement for most disease resistance breeding programs and precision agriculture.

Traits identified in controlled environments at small scales such as lesion size or sporulation, may, in some cases, be transferrable to the field [7, 136, 137]. For example, Northern corn leaf blight causes large and obvious lesions on maize plants, a symptom that was measured on RGB images taken from a UAV at 6 m altitude [118, 120]. This way, the relative area of necrotic maize tissue could be measured with high throughput.

However, it is technically more demanding to measure signs and symptoms under field conditions. Multiple biotic and abiotic stresses can affect plants simultaneously and lead to a loss of specificity. For example, chlorosis may be caused both by a pathogen or abiotic stress. In a canopy, plants or plant parts can shade each other from the sensors, especially in tall growing crops where lower plant parts are covered. Applying sensors from a distance, e.g. when mounted on UAVs, can result in a lack of spatial resolution. Also, natural radiation influences sensor-based measurements and can confound the measurement of signs and symptoms.

The effect of natural radiation can be controlled either by numerical correction during the data analysis, or by shading during the data acquisition, but both approaches are difficult due to the spatial and temporal variation of natural radiation. The technical limitations involving spatial resolution and natural radiation could be overcome partly by ground-based phenotyping platforms that carry sensors close to the canopy and offer a compromise between throughput and accuracy. For example, ground-based RGB imaging outperformed aerial RGB imaging for assessing *Cercospora* leaf spot in a sugar beet field trial [119]. The ground-based platform offered higher spatial resolution and artificial illumination which enabled measuring necrotic spot size and density, traits that could not be measured with the aerial platform. The specificity of measurements could potentially be further improved by sensor fusion, the integration of data from multiple sources [138, 139]. A combination of functional

plant traits derived from hyperspectral imaging (400–885 nm) and a thermal imaging sensor mounted on an aircraft enabled early detection of *Xylella fastidiosa*, a xylem-bound bacterial pathogen of olive trees [48].

These examples illustrate that it is feasible to increase throughput while maintaining accuracy also under field conditions. But it is critical that the challenges of specificity, canopy architecture, spatial resolution and natural radiation are considered and addressed with new solutions like improved sensor and vector technology or sensor fusion.

Non-invasive phenotyping of below-ground plant-pathogen interactions remains a challenge

Phenotyping of PPI below ground is still in an early stage of development as in-soil non-invasive phenotyping is difficult. Using invasive sampling and RGB imaging, changes of morphological root characteristics could be detected on soybean infected with *Fusarium* species and on alfalfa affected by *Phymatotrichopsis* root rot [140, 141]. Tomographic sensors are an option to measure PPI on roots growing in substrate in pots non-invasively [135]. With new X-ray CT scanners that are installed in automated phenotyping facilities, this method may become more accessible [142–144].

Sensor data helps to understand plant-pathogen interactions in more detail

Sensor-based phenotyping is commonly deployed to substitute traditional visual disease scores, for example to rank a genotype on a spectrum from resistant to susceptible in comparison with other genotypes [12]. Yet sensor-based phenotyping can capture the sum of processes that underlie PPI in more detail than what is reflected in traditional disease severity scores. To understand which specific PPI related changes influence the detected signals, sensor data needs to be linked with in situ measurements, particularly in the case of advanced sensors. This has been shown successfully in two studies that linked disease induced accumulation of phenolic compounds to specific Raman bands in Raman spectroscopy [90] and flavonoids to UV absorption features in spectral imaging [81]. Establishing and confirming such links will allow to non-invasively measure diverse aspects of PPI simultaneously and to transfer those findings between pathosystems and environments.

Advanced sensors expand the range of perceivable signs and symptoms but require complementary technologies

Implementation of sensor technology for phenotyping of PPI allows for measurement of a wide range of signs and symptoms (Table 2). Plant metabolites may be detected

with spectroscopic methods, and internal plant and pathogen structures can be detected with tomographic methods [15, 18]. Fluorescence imaging, possibly with tagged pathogens, and chlorophyll fluorescence imaging are other promising approaches for measuring pathogen growth and photosynthetic parameters. To build useful phenotyping systems, any improvements and innovation in sensor technology need to be matched with appropriate facilities, vector technology, data management and data analysis methods. In those fields, constant improvement is pivotal, such as increased payload of UAVs, addition of active illumination to ground-based phenotyping platforms, automation of indoor phenotyping systems, implementation of FAIR data standards and new machine learning methods for analysis [11, 145–148]. Integration of the resulting phenotypes with other-omics data can enable a more comprehensive interpretation of sensor data and will eventually lead to a deeper understanding of plant-pathogen interactions.

Abbreviations

CCD: Charge-coupled device; Chl-F: Chlorophyll *a* fluorescence; CNN: Convolutional neural network; CT: Computed tomography; FAIR: Findable, accessible, interoperable, reusable; F_m : Maximum fluorescence; F_v/F_m : Maximum quantum efficiency of PSII; IR: Infrared; LDT: Lesion doubling time; NIR: Near-infrared; NPQ: Non-photochemical quenching; OCT: Optical coherence tomography; PAM: Pulse-amplitude-modulated; PMMoV: Pepper mild mottle virus; PPI: Plant-pathogen interaction; PSII: Photosystem II; PSM: Plant secondary metabolite; qP: Photochemical quenching; RGB: Red–green–blue; SPCSV: Sweet potato chlorotic stunt virus; SPFMV: Sweet potato feathery mottle virus; SWIR: Shortwave-infrared; TIR: Thermal infrared; TMV: Tobacco mosaic virus; UAV: Uncrewed aerial vehicle; UV: Ultraviolet; UV-C: Ultraviolet C; VIS: Visible.

Authors' contributions

FT and ST analyzed references and drafted the manuscript; DP, BB, GA, and JW revised manuscript. All authors read and approved the final manuscript.

Funding

FT is supported by the University of Adelaide with a Scholarship International, and by The Plant Accelerator with a PhD stipend top-up. BB, DP and FT acknowledge the use of the facilities, and scientific and technical assistance of the Australian Plant Phenomics Facility, which is supported by the Australian Government's National Collaborative Research Infrastructure Strategy (NCRIS). ST and JW are supported by the PathoView project (16293) granted to GA, a TTW partnership project that is funded by Rijk Zwaan (de Lier, Netherlands) and the Dutch Research Council (NWO).

Availability of data and materials

Data sharing not applicable to this article as no datasets were generated or analysed during the current study.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

Author details

¹Australian Plant Phenomics Facility, School of Agriculture, Food and Wine, University of Adelaide, Urrbrae, SA, Australia. ²Department of Biology, Plant-Microbe Interactions, Utrecht University, 3584CH Utrecht, The Netherlands. ³Department of Imaging Physics, Delft University of Technology, Lorentzweg 1, 2628 CJ Delft, The Netherlands.

Received: 29 June 2021 Accepted: 8 February 2022

Published online: 21 March 2022

References

- Savary S, Willocquet L, Pethybridge SJ, Esker P, McRoberts N, Nelson A. The global burden of pathogens and pests on major food crops. *Nat Ecol Evol*. 2019;3:430–9. <https://doi.org/10.1038/s41559-018-0793-y>.
- D'Arcy CJ. Illustrated glossary of plant pathology. *Plant Heal Instr*. 2001. <https://doi.org/10.1094/PHI-I-2001-0219-01>.
- Niks RE, Parlevliet JE, Lindhout P, Bai Y. Breeding crops with resistance to diseases and pests. The Netherlands: Wageningen Academic Publishers; 2019.
- Agrios GN. How pathogens attack plants. In: *Plant pathology*. Amsterdam: Elsevier; 2005. p. 175–205.
- Agrios GN. Environmental factors that cause plant diseases. *Plant Pathol*. 2005. <https://doi.org/10.1016/b978-0-08-047378-9.50016-6>.
- Mutka AM, Bart RS. Image-based phenotyping of plant disease symptoms. *Front Plant Sci*. 2015;5:1–8. <https://doi.org/10.3389/fpls.2014.00734>.
- Simko I, Jimenez-Berni JA, Sirault XRR. Phenomic approaches and tools for phytopathologists. *Phytopathology*. 2017;107:6–17. <https://doi.org/10.1094/PHYTO-02-16-0082-RVW>.
- Mahlein AK, Oerke EC, Steiner U, Dehne HW. Recent advances in sensing plant diseases for precision crop protection. *Eur J Plant Pathol*. 2012;133:197–209. <https://doi.org/10.1007/s10658-011-9878-z>.
- Furbank RT, Tester M. Phenomics—technologies to relieve the phenotyping bottleneck. *Trends Plant Sci*. 2011;16:635–44. <https://doi.org/10.1016/j.tplants.2011.09.005>.
- Singh A, Ganapathysubramanian B, Singh AK, Sarkar S. Machine learning for high-throughput stress phenotyping in plants. *Trends Plant Sci*. 2016;21:110–24. <https://doi.org/10.1016/j.tplants.2015.10.015>.
- Sperschneider J. Machine learning in plant–pathogen interactions: empowering biological predictions from field scale to genome scale. *New Phytol*. 2019. <https://doi.org/10.1111/nph.15771>.
- Bock CH, Barbedo JGA, Del Ponte EM, Bohnenkamp D, Mahlein A-K. From visual estimates to fully automated sensor-based measurements of plant disease severity: status and challenges for improving accuracy. *Phytopathol Res*. 2020;2:9. <https://doi.org/10.1186/s42483-020-00049-8>.
- Sankaran S, Mishra A, Ehsani R, Davis C. A review of advanced techniques for detecting plant diseases. *Comput Electron Agric*. 2010;72:1–13. <https://doi.org/10.1016/j.compag.2010.02.007>.
- Mahlein A-K. Present and future trends in plant disease detection. *Plant Dis*. 2016;100:1–11. <https://doi.org/10.1007/s13398-014-0173-7.2>.
- Farber C, Mahnke M, Sanchez L, Kurouski D. Advanced spectroscopic techniques for plant disease diagnostics. A review. *TrAC Trends Anal Chem*. 2019;118:43–9. <https://doi.org/10.1016/j.trac.2019.05.022>.
- Oerke EC. Precision agriculture for sustainability. In: Stafford J, editor. *Precision agriculture for sustainability*. Routledge: Taylor & Francis; 2019.
- Payne WZ, Kurouski D. Raman-based diagnostics of biotic and abiotic stresses in plants. A review. *Front Plant Sci*. 2021. <https://doi.org/10.3389/fpls.2020.616672>.
- Sylvain T, Cecile LG. Disease identification: a review of vibrational spectroscopy applications. *Compr Anal Chem*. 2018;80:195–225. <https://doi.org/10.1016/bs.coac.2018.03.005>.
- Restif O, Koella JC. Concurrent evolution of resistance and tolerance to pathogens. *Am Nat*. 2004. <https://doi.org/10.1086/423713>.
- Bierman A, LaPlum M, Cadle-Davidson L, Gadoury D, Martinez D, Sapkota S, et al. A high-throughput phenotyping system using machine vision to quantify severity of grapevine powdery mildew. *Plant Phenomics*. 2019;2019:1–13. <https://doi.org/10.34133/2019/9209727>.
- Lück S, Strickert M, Lorbeer M, Melchert F, Backhaus A, Kilias D, et al. 'MacroBot'—an automated segmentation-based system for powdery mildew disease quantification. *Plant Phenomics*. 2020. <https://doi.org/10.1101/2020.03.16.993451>.
- Oerke EC, Leucker M, Steiner U. Sensory assessment of *Cercospora beticola* sporulation for phenotyping the partial disease resistance of sugar beet genotypes. *Plant Methods*. 2019;15:1–12. <https://doi.org/10.1186/s13007-019-0521-x>.
- Yates S, Mikaberidze A, Krattinger SG, Abrouk M, Hund A, Yu K, et al. Precision phenotyping reveals novel Loci for quantitative resistance to *Septoria tritici* Blotch. *Plant Phenomics*. 2019;2019:1–11. <https://doi.org/10.34133/2019/3285904>.
- Lorang JM, Tuori RP, Martinez JP, Sawyer TL, Redman RS, Rollins JA, et al. Green fluorescent protein is lighting up fungal biology. *Appl Environ Microbiol*. 2001;67:1987–94. <https://doi.org/10.1128/aem.67.5.1987-1994.2001>.
- Kassem II, Splitter GA, Miller S, Rajashekara G. Let there be light! Bioluminescent imaging to study bacterial pathogenesis in live animals and plants. *Biolumin Fundam Appl Biotechnol*. 2014;3:119–45.
- Skadsen RW, Hohn TM. Use of *Fusarium graminearum* transformed with gfp to follow infection patterns in barley and *Arabidopsis*. *Physiol Mol Plant Pathol*. 2004;64:45–53. <https://doi.org/10.1016/j.pmpp.2004.04.003>.
- Chapman S, Faulkner C, Kaiserli E, Garcia-Mata C, Savenkov EI, Roberts AG, et al. The photoreversible fluorescent protein iLOV outperforms GFP as a reporter of plant virus infection. *Proc Natl Acad Sci*. 2008;105:20038–43. <https://doi.org/10.1073/pnas.0807551105>.
- Domazakis E, Wouters D, Visser RGF, Kamoun S, Joosten MHJA, Vleeshouwers VGAA. The ELR-SOBIR1 complex functions as a two-component receptor-like kinase to mount defense against *Phytophthora infestans*. *Mol Plant Microbe Interact*. 2018;31:795–802. <https://doi.org/10.1094/mpmi-09-17-0217-r>.
- Wang K, Kang L, Anand A, Lazarovits G, Mysore KS. Monitoring in planta bacterial infection at both cellular and whole-plant levels using the green fluorescent protein variant GFPuv. *New Phytol*. 2007;174:212–23. <https://doi.org/10.1111/j.1469-8137.2007.01999.x>.
- Fujie M, Takamoto H, Kawasaki T, Fujiwara A, Yamada T. Monitoring growth and movement of *Ralstonia solanacearum* cells harboring plasmid pRSS12 derived from bacteriophage φRSS1. *J Biosci Bioeng*. 2010;109:153–8. <https://doi.org/10.1016/j.jbiosc.2009.07.012>.
- Hupp S, Rosenkranz M, Bonfig K, Pandey C, Roitsch T. Noninvasive phenotyping of plant-pathogen interaction: consecutive in situ imaging of fluorescing *Pseudomonas syringae*, plant phenolic fluorescence, and chlorophyll fluorescence in *Arabidopsis* leaves. *Front Plant Sci*. 2019;10:1239. <https://doi.org/10.3389/fpls.2019.01239PMID-31681362>.
- Du H, Wen C, Zhang X, Xu X, Yang J, Chen B, et al. Identification of a Major QTL (qRRs-10.1) that confers resistance to *Ralstonia solanacearum* in pepper (*Capsicum annuum*) using SLAF-BSA and QTL mapping. *Int J Mol Sci*. 2019;20:5887. <https://doi.org/10.3390/ijms20235887>.
- Xu X, Rajashekara G, Paul PA, Miller SA. Colonization of tomato seedlings by bioluminescent *Clavibacter michiganensis* subsp. *michiganensis* under different humidity regimes. *Phytopathology*. 2012;102:177–84. <https://doi.org/10.1094/phyto-03-11-0090>.
- Soldan R, Sanguankiatichai N, Bach-Pages M, Bervoets I, Huang WE, Preston GM. From macro to micro: a combined bioluminescence-fluorescence approach to monitor bacterial localization. *Environ Microbiol*. 2020. <https://doi.org/10.1111/1462-2920.15296>.
- Kotlobay AA, Sarkisyan KS, Mokrushina YA, Marcet-Houben M, Serbroykaya EO, Markina NM, et al. Genetically encodable bioluminescent system from fungi. *Proc Natl Acad Sci*. 2018;115:12728–32. <https://doi.org/10.1073/pnas.1803615115>.
- Berger S, Sinha AK, Roitsch T. Plant physiology meets phytopathology: plant primary metabolism and plant pathogen interactions. *J Exp Bot*. 2007;58:4019–26. <https://doi.org/10.1093/jxb/erm298>.
- Bolton MD. Primary metabolism and plant defense—fuel for the fire. *Mol Plant Microbe Interact*. 2009;22:487–97. <https://doi.org/10.1094/mpmi-22-5-0487>.
- Rojas CM, Senthil-Kumar M, Tzin V, Mysore KS. Regulation of primary plant metabolism during plant-pathogen interactions and its contribution to plant defense. *Front Plant Sci*. 2014. <https://doi.org/10.3389/fpls.2014.00017>.

39. Baker NR. Chlorophyll fluorescence: a probe of photosynthesis in vivo. *Annu Rev Plant Biol.* 2008;59:89–113. <https://doi.org/10.1146/annurev.arplant.59.032607.092759>.
40. Murchie EH, Lawson T. Chlorophyll fluorescence analysis: a guide to good practice and understanding some new applications. *J Exp Bot.* 2013;64:3983–98. <https://doi.org/10.1093/jxb/ert088PMID-23913954>.
41. Butler WL. Energy distribution in the photochemical apparatus of photosynthesis. *Annu Rev Plant Physiol.* 1978;29:345–78. <https://doi.org/10.1146/annurev.pp.29.060178.002021>.
42. Pérez-Bueno ML, Pineda M, Barón M. Phenotyping plant responses to biotic stress by chlorophyll fluorescence imaging. *Front Plant Sci.* 2019;10:1135. <https://doi.org/10.3389/fpls.2019.01135>.
43. Virlet N, Sabermanesh K, Sadeghi-Tehrani P, Hawkesford MJ. Field scanner: an automated robotic field phenotyping platform for detailed crop monitoring. *Funct Plant Biol.* 2017;44:143. <https://doi.org/10.1071/fp16163>.
44. Rascher U, Alonso L, Burkart A, Cilia C, Cogliati S, Colombo R, et al. Sun-induced fluorescence—a new probe of photosynthesis: first maps from the imaging spectrometer HyPlant. *Glob Chang Biol.* 2015;21:4673–84. <https://doi.org/10.1111/gcb.13017>.
45. Silva-Perez V, Molero G, Serbin SP, Condon AG, Reynolds MP, Furbank RT, et al. Hyperspectral reflectance as a tool to measure biochemical and physiological traits in wheat. *J Exp Bot.* 2018;69:483–96. <https://doi.org/10.1093/jxb/erx421>.
46. Aasen H, Wittenberghe V, Medina S, Damm A, Goulas Y, Wieneke S, et al. Sun-induced chlorophyll fluorescence II: review of passive measurement setups, protocols, and their application at the leaf to canopy level. *Remote Sens.* 2019;11:927. <https://doi.org/10.3390/rs11080927>.
47. Meacham-Hensold K, Fu P, Wu J, Serbin S, Montes CM, Ainsworth E, et al. Plot-level rapid screening for photosynthetic parameters using proximal hyperspectral imaging. *J Exp Bot.* 2020;71:2312–28. <https://doi.org/10.1093/jxb/eraa068>.
48. Zarco-Tejada PJ, Camino C, Beck PSA, Calderon R, Hornero A, Hernández-Clemente R, et al. Previsual symptoms of *Xylella fastidiosa* infection revealed in spectral plant-trait alterations. *Nat Plants.* 2018;4:432–9. <https://doi.org/10.1038/s41477-018-0189-7>.
49. Pinto F, Celesti M, Acebron K, Alberti G, Cogliati S, Colombo R, et al. Dynamics of sun-induced chlorophyll fluorescence and reflectance to detect stress-induced variations in canopy photosynthesis. *Plant Cell Environ.* 2020;43:1637–54. <https://doi.org/10.1111/pce.13754>.
50. Dechant B, Ryu Y, Badgley G, Zeng Y, Berry JA, Zhang Y, et al. Canopy structure explains the relationship between photosynthesis and sun-induced chlorophyll fluorescence in crops. *Remote Sens Environ.* 2020;241: 111733. <https://doi.org/10.1016/j.rse.2020.111733>.
51. Cséfalvay L, Di Gaspero G, Matouš K, Bellin D, Ruperti B, Olejníčková J. Pre-symptomatic detection of *Plasmopara viticola* infection in grapevine leaves using chlorophyll fluorescence imaging. *Eur J Plant Pathol.* 2009;125:291–302. <https://doi.org/10.1007/s10658-009-9482-7>.
52. Kuckenberg J, Tartachnyk I, Noga G. Temporal and spatial changes of chlorophyll fluorescence as a basis for early and precise detection of leaf rust and powdery mildew infections in wheat leaves. *Precis Agric.* 2009;10:34–44. <https://doi.org/10.1007/s11119-008-9082-0>.
53. Brugger A, Behmann J, Paulus S, Luigs HG, Kuska MT, Schramowski P, et al. Extending hyperspectral imaging for plant phenotyping to the UV-range. *Remote Sens.* 2019;11:1–11. <https://doi.org/10.3390/rs11121401>.
54. Polonio Á, Pineda M, Bautista R, Martínez-Cruz J, Pérez-Bueno ML, Barón M, et al. RNA-seq analysis and fluorescence imaging of melon powdery mildew disease reveal an orchestrated reprogramming of host physiology. *Sci Rep.* 2019;9:1–16. <https://doi.org/10.1038/s41598-019-44443-5>.
55. Lichtenthaler HK, Rinderle U. The role of chlorophyll fluorescence in the detection of stress conditions in plants. *C R C Crit Rev Anal Chem.* 1988;19:529–85. <https://doi.org/10.1080/15476510.1988.10401466>.
56. Ely KS, Burnett AC, Lieberman-Cribbin W, Serbin SP, Rogers A. Spectroscopy can predict key leaf traits associated with source–sink balance and carbon–nitrogen status. *J Exp Bot.* 2019;70:1789–99. <https://doi.org/10.1093/jxb/erz061>.
57. Curran PJ. Remote sensing of foliar chemistry. *Remote Sens Environ.* 1989;30:271–8. [https://doi.org/10.1016/0034-4257\(89\)90069-2](https://doi.org/10.1016/0034-4257(89)90069-2).
58. Serbin SP, Singh A, Mcneil BE, Kingdon CC, Townsend PA. Spectroscopic determination of leaf morphological and biochemical traits for northern temperate and boreal tree species. *Ecol Appl.* 2014;24:1651–69. <https://doi.org/10.1890/13-2110.1>.
59. Gold KM, Townsend PA, Chlus A, Herrmann I, Couture JJ, Larson ER, et al. Hyperspectral measurements enable pre-symptomatic detection and differentiation of contrasting physiological effects of late blight and early blight in potato. *Remote Sens.* 2020;12:286. <https://doi.org/10.3390/rs12020286>.
60. Wang Z, Chlus A, Geygan R, Ye Z, Zheng T, Singh A, et al. Foliar functional traits from imaging spectroscopy across biomes in eastern North America. *New Phytol.* 2020;228:494–511. <https://doi.org/10.1111/nph.16711>.
61. Conrad AO, Li W, Lee D-Y, Wang G-L, Rodriguez-Saona L, Bonello P. Machine learning-based presymptomatic detection of rice sheath blight using spectral profiles. *Plant Phenomics.* 2020;2020:1–10. <https://doi.org/10.34133/2020/8954085>.
62. Theis N, Lerdau M. The evolution of function in plant secondary metabolites. *Int J Plant Sci.* 2003;164:S93–102. <https://doi.org/10.1086/374190>.
63. Kessler A, Kalske A. Plant secondary metabolite diversity and species interactions. *Annu Rev Ecol Evol Syst.* 2018;49:115–38. <https://doi.org/10.1146/annurev-ecolsys-110617-062406>.
64. Bednarek P. Chemical warfare or modulators of defence responses—the function of secondary metabolites in plant immunity. *Curr Opin Plant Biol.* 2012;15:407–14. <https://doi.org/10.1016/j.pbi.2012.03.002>.
65. Piasecka A, Jedrzejczak-Rey N, Bednarek P. Secondary metabolites in plant innate immunity: conserved function of divergent chemicals. *New Phytol.* 2015;206:948–64. <https://doi.org/10.1111/nph.13325>.
66. García-Plazaola JL, Fernández-Marín B, Duke SO, Hernández A, López-Arbeloa F, Becerril JM. Autofluorescence: biological functions and technical applications. *Plant Sci.* 2015;236:136–45. <https://doi.org/10.1016/j.plantsci.2015.03.010>.
67. Pezet R, Gindro K, Viret O, Spring J-L. Glycosylation and oxidative dimerization of resveratrol are respectively associated to sensitivity and resistance of grapevine cultivars to downy mildew. *Physiol Mol Plant Pathol.* 2004;65:297–303. <https://doi.org/10.1016/j.pmpp.2005.03.002>.
68. Bellow S, Latouche G, Brown SC, Poutaraud A, Cervic ZG. In vivo localization at the cellular level of stilbene fluorescence induced by *Plasmopara viticola* in grapevine leaves. *J Exp Bot.* 2012;63:3697–707. <https://doi.org/10.1093/jxb/ers060PMID-22412183>.
69. Bellow S, Latouche G, Brown SC, Poutaraud A, Cervic ZG. Optical detection of downy mildew in grapevine leaves: daily kinetics of autofluorescence upon infection. *J Exp Bot.* 2012;64:333–41. <https://doi.org/10.1093/jxb/ers338PMID-23213137>.
70. Becker L, Bellow S, Carré V, Latouche G, Poutaraud A, Merdinoglu D, et al. Correlative analysis of fluorescent phytoalexins by mass spectrometry imaging and fluorescence microscopy in grapevine leaves. *Anal Chem.* 2017;89:7099–106. <https://doi.org/10.1021/acs.analchem.7b01002>.
71. Ciaffi M, Paolacci AR, Paolacci M, Alicandri E, Bigini V, Badiani M, et al. Transcriptional regulation of stilbene synthases in grapevine germplasm differentially susceptible to downy mildew. *BMC Plant Biol.* 2019. <https://doi.org/10.1186/s12870-019-2014-5>.
72. Pineda M, Gáspár L, Morales F, Szigeti Z, Barón M. Multicolor fluorescence imaging of leaves—a useful tool for visualizing systemic viral infections in plants. *Photochem Photobiol.* 2008;84:1048–60. <https://doi.org/10.1111/j.1751-1097.2008.00357.x>.
73. Bürling K, Hunsche M, Noga G. Use of blue–green and chlorophyll fluorescence measurements for differentiation between nitrogen deficiency and pathogen infection in winter wheat. *J Plant Physiol.* 2011;168:1641–8. <https://doi.org/10.1016/j.jplph.2011.03.016>.
74. Lichtenthaler HK, Miehe JA. Fluorescence imaging as a diagnostic tool for plant stress. *Trends Plant Sci.* 1997;2:316–20. [https://doi.org/10.1016/S1360-1385\(97\)89954-2](https://doi.org/10.1016/S1360-1385(97)89954-2).
75. Lang M, Lichtenthaler HK, Sowinska M, Heisel F, Miehe JA. Fluorescence imaging of water and temperature stress in plant leaves. *J Plant Physiol.* 1996;148:613–21. [https://doi.org/10.1016/S0176-1617\(96\)80082-4](https://doi.org/10.1016/S0176-1617(96)80082-4).
76. Hideg É, Juhász M, Bornman JF, Asada K. The distribution and possible origin of blue–green fluorescence in control and stressed barley leaves. *Photochem Photobiol Sci.* 2002;1:934–41. <https://doi.org/10.1039/b201916g>.

77. Heisel F, Sowinska M, Miehe JA, Lang M, Lichtenthaler HK. Detection of nutrient deficiencies of maize by laser induced fluorescence imaging. *J Plant Physiol.* 1996;148:622–31. [https://doi.org/10.1016/S0176-1617\(96\)80083-6](https://doi.org/10.1016/S0176-1617(96)80083-6).
78. Lang M, Stober F, Lichtenthaler HK. Fluorescence emission spectra of plant leaves and plant constituents. *Radiat Environ Biophys.* 1991;30:333–47. <https://doi.org/10.1007/bf01210517>.
79. Couture JJ, Singh A, Rubert-Nason KF, Serbin SP, Lindroth RL, Townsend PA. Spectroscopic determination of ecologically relevant plant secondary metabolites. *Methods Ecol Evol.* 2016;7:1402–12. <https://doi.org/10.1111/2041-210x.12596>.
80. Kokaly RF, Skidmore AK. Plant phenolics and absorption features in vegetation reflectance spectra near 1.66 μm . *Int J Appl Earth Obs Geoinf.* 2015;43:55–83. <https://doi.org/10.1016/j.jag.2015.01.010>.
81. Brügger A, Schramowski P, Paulus S, Steiner U, Kersting K, Mahlein AK. Spectral signatures in the UV range can be combined with secondary plant metabolites by deep learning to characterize barley–powdery mildew interaction. *Plant Pathol.* 2021;70:1572–82. <https://doi.org/10.1111/ppa.13411>.
82. Arens N, Backhaus A, Döll S, Fischer S, Seiffert U, Mock HP. Non-invasive presymptomatic detection of *Cercospora beticola* infection and identification of early metabolic responses in sugar beet. *Front Plant Sci.* 2016;7:1–14. <https://doi.org/10.3389/fpls.2016.01377>.
83. Butler HJ, Ashton L, Bird B, Cinque G, Curtis K, Dorney J, et al. Using Raman spectroscopy to characterize biological materials. *Nat Protoc.* 2016;11:664–87. <https://doi.org/10.1038/nprot.2016.036>.
84. Gupta S, Huang CH, Singh GP, Park BS, Chua N-H, Ram RJ. Portable Raman leaf-clip sensor for rapid detection of plant stress. *Sci Rep.* 2020. <https://doi.org/10.1038/s41598-020-76485-5>.
85. Farber C, Bryan R, Paetzold L, Rush C, Kuroski D. Non-invasive characterization of single-, double- and triple-viral diseases of wheat with a hand-held Raman spectrometer. *Front Plant Sci.* 2020;11:1–7.
86. Sanchez L, Pant S, Xing Z, Mandadi K, Kuroski D. Rapid and noninvasive diagnostics of Huanglongbing and nutrient deficits on citrus trees with a handheld Raman spectrometer. *Anal Bioanal Chem.* 2019;411:3125–33. <https://doi.org/10.1007/s00216-019-01776-4>.
87. Sanchez L, Pant S, Mandadi K, Kuroski D. Raman spectroscopy vs quantitative polymerase chain reaction in early stage Huanglongbing diagnostics. *Sci Rep.* 2020;10:10101. <https://doi.org/10.1038/s41598-020-67148-6>.
88. Mandrile L, Rotunno S, Miozzi L, Vaira AM, Giovannozzi AM, Rossi AM, et al. Nondestructive Raman spectroscopy as a tool for early detection and discrimination of the infection of tomato plants by two economically important viruses. *Anal Chem.* 2019;91:9025–31. <https://doi.org/10.1021/acs.analchem.9b01323>.
89. Egging V, Nguyen J, Kuroski D. Detection and identification of fungal infections in intact wheat and sorghum grain using a hand-held Raman spectrometer. *Anal Chem.* 2018;90:8616–21. <https://doi.org/10.1021/acs.analchem.8b01863>.
90. Dou T, Sanchez L, Irigoyen S, Goff N, Niraula P, Mandadi K, et al. Biochemical origin of Raman-based diagnostics of Huanglongbing in grapefruit trees. *Front Plant Sci.* 2021. <https://doi.org/10.3389/fpls.2021.680991>.
91. Farber C, Bryan R, Paetzold L, Rush C, Kuroski D. Non-invasive characterization of single-, double- and triple-viral diseases of wheat with a hand-held Raman spectrometer. *Front Plant Sci.* 2020. <https://doi.org/10.3389/fpls.2020.01300>.
92. Stewart S, Priore RJ, Nelson MP, Treado PJ. Raman imaging. *Annu Rev Anal Chem.* 2012;5:337–60. <https://doi.org/10.1146/annurev-anchem-062011-143152>.
93. Lohumi S, Kim MS, Qin J, Cho B-K. Raman imaging from microscopy to macroscopy: quality and safety control of biological materials. *TrAC Trends Anal Chem.* 2017;93:183–98. <https://doi.org/10.1016/j.trac.2017.06.002>.
94. Lee H, Kim M, Qin J, Park E, Song Y-R, Oh C-S, et al. Raman hyperspectral imaging for detection of watermelon seeds infected with *Acidovorax citrulli*. *Sensors.* 2017;17:2188. <https://doi.org/10.3390/s17102188>.
95. Agrios GN. Plant diseases caused by viruses. *Plant Pathol.* 2005. <https://doi.org/10.1016/b978-0-08-047378-9.50020-8>.
96. Bender CL. Chlorosis-inducing phytotoxins produced by *Pseudomonas syringae*. *Eur J Plant Pathol.* 1999;105:1–12. <https://doi.org/10.1023/A:1008692227307>.
97. Ciuffetti LM, Manning VA, Pandelova I, Betts MF, Martinez JP. Host-selective toxins, Ptr ToxA and Ptr ToxB, as necrotrophic effectors in the *Pyrenophora tritici-repentis*-wheat interaction. *New Phytol.* 2010;187:911–9. <https://doi.org/10.1111/j.1469-8137.2010.03362.x>.
98. Chakravarthy S, Worley JN, Montes-Rodriguez A, Collmer A. *Pseudomonas syringae* pv. tomato DC3000 polymutants deploying coronatine and two type III effectors produce quantifiable chlorotic spots from individual bacterial colonies in *Nicotiana benthamiana* leaves. *Mol Plant Pathol.* 2018;19:935–47. <https://doi.org/10.1111/mpp.12579>.
99. Van Kan JAL. Licensed to kill: the lifestyle of a necrotrophic plant pathogen. *Trends Plant Sci.* 2006;11:247–53. <https://doi.org/10.1016/j.tplants.2006.03.005>.
100. Oome S, Raaymakers TM, Cabral A, Samwel S, Böhm H, Albert I, et al. Nep1-like proteins from three kingdoms of life act as a microbe-associated molecular pattern in *Arabidopsis*. *Proc Natl Acad Sci.* 2014;111:16955–60. <https://doi.org/10.1073/pnas.1410031111>.
101. Hutchison ML, Tester MA, Gross DC. Role of biosurfactant and ion channel-forming activities of syringomycin in transmembrane ion flux: a model for the mechanism of action in the plant-pathogen interaction. *Mol Plant Microbe Interact.* 1995. <https://doi.org/10.1094/mpmi-8-0610>.
102. Seidl MF, Van Den Ackerveken G. Activity and phylogenetics of the broadly occurring family of microbial Nep1-like proteins. *Annu Rev Phytopathol.* 2019;57:367–86. <https://doi.org/10.1146/annurev-phyto-082718-100054>.
103. Möbius N, Hertweck C. Fungal phytotoxins as mediators of virulence. *Curr Opin Plant Biol.* 2009;12:390–8. <https://doi.org/10.1016/j.pbi.2009.06.004>.
104. Espinoza C, Medina C, Somerville S, Arce-Johnson P. Senescence-associated genes induced during compatible viral interactions with grapevine and *Arabidopsis*. *J Exp Bot.* 2007;58:3197–212. <https://doi.org/10.1093/jxb/erm165>.
105. Seo J-K, Kim M-K, Kwak H-R, Choi H-S, Nam M, Choe J, et al. Molecular dissection of distinct symptoms induced by tomato chlorosis virus and tomato yellow leaf curl virus based on comparative transcriptome analysis. *Virology.* 2018;516:1–20. <https://doi.org/10.1016/j.virol.2018.01.001>.
106. Wang Y, Tan J, Wu Z, Vandenlangenberg K, Wehner CT, Wen C, et al. STAYGREEN, STAY HEALTHY: a loss-of-susceptibility mutation in the STAYGREEN gene provides durable, broad-spectrum disease resistances for over 50 years of US cucumber production. *New Phytol.* 2019;221:415–30. <https://doi.org/10.1111/nph.15353>.
107. Slaton MR, Raymond Hunt E, Smith WK. Estimating near-infrared leaf reflectance from leaf structural characteristics. *Am J Bot.* 2001;88:278–84. <https://doi.org/10.2307/2657019>.
108. Wilson RH, Nadeau KP, Jaworski FB, Tromberg BJ, Durkin AJ. Review of short-wave infrared spectroscopy and imaging methods for biological tissue characterization. *J Biomed Opt.* 2015;20:30901. <https://doi.org/10.1117/1.JBO.20.3.030901>.
109. Kim MD, Zhang H, Zhou H, Du T, Wu Q, Mockler CT, et al. Highly sensitive image-derived indices of water-stressed plants using hyperspectral imaging in SWIR and histogram analysis. *Sci Rep.* 2015;5:15919. <https://doi.org/10.1038/srep15919>.
110. Chaerle L, Hagenbeek D, Bruyere ED, Der Straeten DV. Chlorophyll fluorescence imaging for disease-resistance screening of sugar beet. *Plant Cell Tissue Organ Cult.* 2007;91:97–106. <https://doi.org/10.1007/s11240-007-9282-8>.
111. Macioszek KV, Wielanek M, Morkunas I, Ciereszko I, Kononowicz KA. Leaf position-dependent effect of *Alternaria brassicicola* development on host cell death, photosynthesis and secondary metabolites in *Brassica juncea*. *Physiol Plant.* 2020;168:601–16. <https://doi.org/10.1111/ppl.12998>.
112. Landeo Villanueva S, Malvestiti MC, van Ieperen W, Joosten MHJ, van Kan JAL. Red light imaging for programmed cell death visualization and quantification in plant–pathogen interactions. *Mol Plant Pathol.* 2021;22:361–72. <https://doi.org/10.1111/mpp.13027>.

113. Wijesinghe RE, Lee S-Y, Ravichandran NK, Shirazi MF, Kim P, Jung H-Y, et al. Biophotonic approach for the characterization of initial bitter-rot progression on apple specimens using optical coherence tomography assessments. *Sci Rep*. 2018. <https://doi.org/10.1038/s41598-018-33791-3>.
114. Brar GS, Karunakaran C, Bond T, Stobbs J, Liu N, Hucl PJ, et al. Showcasing the application of synchrotron-based X-ray computed tomography in host–pathogen interactions: the role of wheat rachilla and rachis nodes in Type-II resistance to *Fusarium graminearum*. *Plant Cell Environ*. 2019;42:509–26. <https://doi.org/10.1111/pce.13431>.
115. Barbacci A, Navaud O, Mbengue M, Barascud M, Godiard L, Khafif M, et al. Rapid identification of an *Arabidopsis* NLR gene as a candidate conferring susceptibility to *Sclerotinia sclerotiorum* using time-resolved automated phenotyping. *Plant J*. 2020. <https://doi.org/10.1111/tjpi.14747>.
116. Laflamme B, Middleton M, Lo T, Desveaux D, Guttman DS. Image-based quantification of plant immunity and disease. *Mol Plant Microbe Interact*. 2016;29:919–24. <https://doi.org/10.1094/mpmi-07-16-0129-ta>.
117. Martel A, Lo T, Desveaux D, Guttman DS. A high-throughput, seedling screen for plant immunity. *Mol Plant Microbe Interact*. 2020;33:394–401. <https://doi.org/10.1094/MPMI-10-19-0295-TA>.
118. Wiesner-Hanks T, Wu H, Stewart E, DeChant C, Kaczmar N, Lipson H, et al. Millimeter-level plant disease detection from aerial photographs via deep learning and crowdsourced data. *Front Plant Sci*. 2019;10:1–11. <https://doi.org/10.3389/fpls.2019.01550>.
119. Jay S, Comar A, Benicio R, Beauvois J, Dutartre D, Daubige G, et al. Scoring *Cercospora* leaf spot on sugar beet: comparison of UGV and UAV phenotyping systems. *Plant Phenomics*. 2020;2020:1–18. <https://doi.org/10.34133/2020/9452123>.
120. Wiesner-Hanks T, Stewart EL, Kaczmar N, DeChant C, Wu H, Nelson RJ, et al. Image set for deep learning: field images of maize annotated with disease symptoms. *BMC Res Notes*. 2018;11:440. <https://doi.org/10.1186/s13104-018-3548-6>.
121. Jones HG, Rotenberg E. Energy, radiation and temperature regulation in plants. In: eLS. Hoboken: Wiley; 2011. <https://doi.org/10.1002/9780470015902.a0003199.pub2>.
122. Chaerle L, Leinonen I, Jones HG, Van Der Straeten D. Monitoring and screening plant populations with combined thermal and chlorophyll fluorescence imaging. *J Exp Bot*. 2007;58:773–84. <https://doi.org/10.1093/jxb/erl257>.
123. Wang M, Ling N, Dong X, Zhu Y, Shen Q, Guo S. Thermographic visualization of leaf response in cucumber plants infected with the soil-borne pathogen *Fusarium oxysporum* f. sp. *cucumerinum*. *Plant Physiol Biochem*. 2012;61:153–61. <https://doi.org/10.1016/j.plaphy.2012.09.015>.
124. Brooker G. Introduction to sensors for ranging and imaging. London: Institution of Engineering and Technology; 2009.
125. Jackson RD, Idso SB, Reginato RJ, Pinter PJ. Canopy temperature as a crop water stress indicator. *Water Resour Res*. 1981;17:1133–8. <https://doi.org/10.1029/WR017i004p01133>.
126. Pinter PJ, Fry KE, Guinn G, Mauney JR. Infrared thermometry: a remote sensing technique for predicting yield in water-stressed cotton. *Agric Water Manag*. 1983;6:385–95. [https://doi.org/10.1016/0378-3774\(83\)90057-4](https://doi.org/10.1016/0378-3774(83)90057-4).
127. Nilsson HE. Remote sensing and image analysis in plant pathology. *Annu Rev Phytopathol*. 1995;33:489–527. <https://doi.org/10.1146/annurev.phyto.33.1.489>.
128. Yadeta KA, Thomma BPHJ. The xylem as battleground for plant hosts and vascular wilt pathogens. *Front Plant Sci*. 2013;4:1–12. <https://doi.org/10.3389/fpls.2013.00097>.
129. Calderón R, Navas-Cortés JA, Lucena C, Zarco-Tejada PJ. High-resolution airborne hyperspectral and thermal imagery for early detection of Verticillium wilt of olive using fluorescence, temperature and narrow-band spectral indices. *Remote Sens Environ*. 2013;139:231–45. <https://doi.org/10.1016/j.rse.2013.07.031>.
130. Calderón R, Navas-Cortés JA, Zarco-Tejada PJ. Early detection and quantification of Verticillium wilt in olive using hyperspectral and thermal imagery over large areas. *Remote Sens*. 2015;7:5584–610. <https://doi.org/10.3390/rs70505584>.
131. Chaerle L, Van CW, Messens E, Lambers H, Van Montagu M, Van Der Straeten D. Presymptomatic visualization of plant–virus interactions by thermography. *Nat Biotechnol*. 1999;17:813–6. <https://doi.org/10.1038/11765>.
132. Wang L, Poque S, Valkonen JPT. Phenotyping viral infection in sweet-potato using a high-throughput chlorophyll fluorescence and thermal imaging platform. *Plant Methods*. 2019;15:1–14. <https://doi.org/10.1186/s13007-019-0501-1>.
133. Lee J, Lee SY, Wijesinghe RE, Ravichandran NK, Han S, Kim P, et al. On-field in situ inspection for Marssonina coronaria infected apple blotch based on non-invasive bio-photonic imaging module. *IEEE Access*. 2019;7:148684–91. <https://doi.org/10.1109/ACCESS.2019.2946388>.
134. Ravichandran NK, Wijesinghe RE, Shirazi MF, Park K, Lee SY, Jung HY, et al. In vivo monitoring on growth and spread of gray leaf spot disease in capsicum annum leaf using spectral domain optical coherence tomography. *J Spectrosc*. 2016. <https://doi.org/10.1155/2016/1093734>.
135. Han L, Dutilleul P, Prasher SO, Beaulieu C, Smith DL. Assessment of common scab-inducing pathogen effects on potato underground organs via computed tomography scanning. *Phytopathology*. 2008;98:1118–25. <https://doi.org/10.1094/PHYTO-98-10-1118>.
136. Rebetzke GJ, Jimenez-Berni J, Fischer RA, Deery DM, Smith DJ. Review: high-throughput phenotyping to enhance the use of crop genetic resources. *Plant Sci*. 2019;282:40–8. <https://doi.org/10.1016/j.plantsci.2018.06.017>.
137. van Eeuwijk FA, Bustos-Korts D, Millet EJ, Boer MP, Kruijer W, Thompson A, et al. Modelling strategies for assessing and increasing the effectiveness of new phenotyping techniques in plant breeding. *Plant Sci*. 2019;282:23–39. <https://doi.org/10.1016/j.plantsci.2018.06.018>.
138. Pauli D, Chapman SC, Bart R, Topp CN, Lawrence-Dill CJ, Poland J, et al. The quest for understanding phenotypic variation via integrated approaches in the field environment. *Plant Physiol*. 2016;172:622–34. <https://doi.org/10.1104/pp.16.00592>.
139. Weiss M, Jacob F, Duveiller G. Remote sensing for agricultural applications: a meta-review. *Remote Sens Environ*. 2020;236: 111402. <https://doi.org/10.1016/j.rse.2019.111402>.
140. Mattupalli C, Seethepalli A, York LM, Young CA. Digital imaging to evaluate root system architectural changes associated with soil biotic factors. *Phytobiomes J*. 2019;3:102–11. <https://doi.org/10.1094/PBIOMES-12-18-0062-R>.
141. Arias MMD, Leandro LF, Munkvold GP. Aggressiveness of *Fusarium* species and impact of root infection on growth and yield of soybeans. *Phytopathology*. 2013;103:822–32. <https://doi.org/10.1094/PHYTO-08-12-0207-R>.
142. APPF Automated Phenotyping System. <https://www.plantphenomics.org.au/technologies/#greenhouses>. Accessed 28 Apr 2021.
143. Controlled Environment Phenotyping Facility (CEPF). <https://ag.purdue.edu/cepf/>. Accessed 31 Mar 2021.
144. PhenoConveyor-Plant Phenomics Research Center (PPRC). http://pprcen.njau.edu.cn/Research/R_D_Platforms.htm. Accessed 31 Mar 2021.
145. Wilkinson MD, Dumontier M, Aalbersberg IJJ, Appleton G, Axton M, Baak A, et al. Comment: the FAIR guiding principles for scientific data management and stewardship. *Sci Data*. 2016;3:1–9. <https://doi.org/10.1038/sdata.2016.18>.
146. Ferentinos KP. Deep learning models for plant disease detection and diagnosis. *Comput Electron Agric*. 2018;145:311–8. <https://doi.org/10.1016/j.compag.2018.01.009>.
147. Singh AK, Ganapathysubramanian B, Sarkar S, Singh A. Deep learning for plant stress phenotyping: trends and future perspectives. *Trends Plant Sci*. 2018;23:883–98. <https://doi.org/10.1016/j.tplants.2018.07.004>.
148. Papoutsoglou EA, Faria D, Arend D, Arnaud E, Athanasiadis IN, Chaves I, et al. Enabling reusability of plant phenomic datasets with MIAPPE 1.1. *New Phytol*. 2020. <https://doi.org/10.1111/nph.16544>.
149. De Torres ZM, Littlejohn G, Jayaraman S, Studholme D, Bailey T, Lawson T, et al. Chloroplasts play a central role in plant defence and are targeted by pathogen effectors. *Nat Plants*. 2015;1:15074. <https://doi.org/10.1038/nplants.2015.74>.
150. Brugger A, Kuska MT, Mahlein A-K. Impact of compatible and incompatible barley—*Blumeria graminis* f. sp. *hordei* interactions on chlorophyll fluorescence parameters. *J Plant Dis Prot*. 2017. <https://doi.org/10.1007/s41348-017-0129-1>.

151. Bauriegel E, Brabandt H, Gärber U, Herppich WB. Chlorophyll fluorescence imaging to facilitate breeding of *Bremia lactucae*-resistant lettuce cultivars. *Comput Electron Agric.* 2014;105:74–82. <https://doi.org/10.1016/j.compag.2014.04.010>.
152. Pineda M, Olejníčková J, Cséfalvay L, Barón M. Tracking viral movement in plants by means of chlorophyll fluorescence imaging. *J Plant Physiol.* 2011;168:2035–40. <https://doi.org/10.1016/j.jplph.2011.06.013>.
153. Gold KM, Townsend PA, Herrmann I, Gevens AJ. Investigating potato late blight physiological differences across potato cultivars with spectroscopy and machine learning. *Plant Sci.* 2020;295: 110316. <https://doi.org/10.1016/j.plantsci.2019.110316>.
154. Arens N, Backhaus A, Döll S, Fischer S, Seiffert U, Mock H-P. Non-invasive Presymptomatic Detection of *Cercospora beticola* Infection and Identification of Early Metabolic Responses in Sugar Beet. *Front Plant Sci.* 2016. <https://doi.org/10.3389/fpls.2016.01377>.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

