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1 AUTOPHAGY AS AN EMERGING ARENA FOR PLANT-PATHOGEN

2 **INTERACTIONS**

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Abstract

Autophagy is a highly conserved degradation and recycling process that controls cellular homeostasis, stress adaptation, and programmed cell death in eukaryotes. Emerging evidence indicates that autophagy is a key regulator of plant innate immunity and contributes with both pro-death and pro-survival functions to antimicrobial defences, depending on the pathogenic lifestyle. In turn, several pathogens have co-opted and evolved strategies to manipulate host autophagy pathways to the benefit of infection, while some eukaryotic microbes require their own autophagy machinery for successful pathogenesis. In this review, we present and discuss recent advances that exemplify the important role of pro- and antimicrobial autophagy in plant-pathogen interactions.

Highlights

- Autophagy is an integral part of plant-pathogen interactions.
- A large variety of microbial pathogens target or are targeted by plant autophagy.
- 39 Autophagy in eukaryotic microbial pathogens is essential for pathogenesis.
- 40 Plant autophagy participates in defense responses against invading microbes.
- Successful pathogens have evolved strategies to manipulate plant autophagy.

Introduction

Autophagy is an evolutionary conserved process in eukaryotes that employs double-membrane vesicular structures, termed autophagosomes, to enclose and deliver cytoplasmic material for vacuolar/lysosomal degradation and recycling [1]. Depending on how the cellular cargo is recruited to the developing autophagosomes, autophagy can act as an unspecific (bulk) catabolic pathway for nutrient remobilization and energy supply, or as selective mechanism to eliminate superfluous and harmful compounds including aggregated proteins and damaged organelles [2]. While basal levels of autophagy serves mainly cellular homeostasis and quality control, increased autophagy activity allows adaptation to stressful conditions caused by a large variety of developmental and environmental cues [3]. Besides the significant contribution to cellular and organismal

survival, autophagy has been implicated in the regulation and execution of programmed cell death (PCD) in various eukaryotic organisms [4]. In plants, autophagy is increasingly recognized for its central importance in development, reproduction, metabolism, senescence and tolerance to abiotic and biotic stresses [5,6]. In this review, we focus on the role of autophagy during plant–pathogen interactions. In particular, we discuss the most recent evidence showing that plant autophagy may benefit either the host by participating in immune responses, or the invading agent, by contributing to infection.

The plant immune system has evolved several layers to fend off pathogenic organisms [7]. Perception of conserved microbial-associated molecular patterns (MAMPs) by surface receptors leads to activation of basal defenses known as MAMP-triggered immunity (MTI). Adapted pathogens interfere with MTI by secreting effectors that, in turn, can be recognized by resistance (R) genes to initiate effector-triggered immunity (ETI). ETI often culminates in a local PCD reaction at the site of pathogen attack, termed the hypersensitive response (HR) [8]. During the last years, it has become evident that autophagy is engaged in various aspects of plant immunity [9]. Most notably, autophagy was shown to regulate basal resistance as well as immunity- and disease-related cell death responses to microbial pathogens with different infection strategies. However, due to the concomitant involvement of plant autophagy in homeostatic, metabolic and developmental processes, the dissection of autophagic mechanisms underlying host immunity and microbial pathogenesis is still in its infancy.

 Most plant pathogens except viruses do not enter the cytoplasmic space, and there is limited evidence for direct autophagic targeting of pathogens or their individual components in in a process resembling xenophagy in metazoans Interestingly, similar to microbes in other host organisms [10,11], an increasing number of examples indicate that phytopathogens are able to manipulate plant autophagy to their own advantage. As detailed below, these include inhibition of autophagy mechanisms contributing to immunity [12-14] and the activation of autophagy pathways to target defense compounds or to potentially enhance nutrient acquisition [15-17].

The role of autophagy in eukaryotic plant pathogens

It is well established that autophagy components and pathways in eukaryotic microbes are important for pathogenesis and plant invasion. Several studies published in the last decade and summarized in [18] showed that microbial autophagy mediates the development of appressoria, specialized infection structures used by fungi and oomycetes to enter the plant tissue. More recently, new components mediating autophagy-dependent plant infection by fungi have been discovered (Figure 1). The conserved retromer complex is involved in protein trafficking from endosomes to the trans-Golgi network, and was shown to be essential for autophagy-dependent host penetration by the rice blast fungus Magnaporthe oryzae [19]. Interestingly, retromer also contributes to the regulation of autophagy-dependent immune cell death in plants [20]. Furthermore, the M. oryzae Rab GTPase MoYpt7 is required for fungal autophagy, appressoria development and pathogenicity [21]. Autophagy is also involved in hyphal fusion and positively regulates the virulence of Fusarium oxysporum [22]. In Botrytis cinerea the autophagy gene BcATG1 is essential for pathogenesis, besides playing a critical role in numerous developmental processes [23]. In several other phytopathogenic fungi, autophagic regulation of organelle quantity has been shown to play a major role in the metabolic switch responsible for the transition to virulence [24].

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The role of autophagy in plant immunity

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Despite some remaining controversy, both pro-death and pro-survival functions of autophagy are now generally recognized to contribute to anti-microbial defenses and disease resistance, depending on the pathosystem and pathogenic lifestyle.

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Autophagy can have a positive regulatory role during HR [25] (Figure 1). Several Arabidopsis mutants disrupted in core autophagy (*ATG*) genes or related pathway components displayed significantly reduced HR upon infection with avirulent strains of the bacterium *Pseudomonas syringae* pathovar (pv) *tomato* (*Pst* DC3000) harboring the effector proteins AvrRps4 or AvrRpm1 [20,26-28]. However, autophagy defects seemed to compromise *R* gene-mediated disease resistance only in case of *Pst* DC3000 AvrRps4 [20,29], supporting the earlier observed decoupling of HR from growth restriction for AvrRpm1-containing bacteria [30]. Knock-down of *ATG6* homologs in wheat further revealed the engagement of autophagy in broad-spectrum immunity conditioned by the *Pm21 R* gene towards the powdery mildew fungus *Blumeria graminis* f. sp. *tritici* (*Bgt*) [31].

Intriguingly, constitutive activation of autophagy in *Nicotiana benthamiana* due to silencing of the ATG3-interacting cytosolic glyceraldehyde-3-phosphate dehydrogenase (GAPC) enhanced *N* gene-mediated HR and resistance against *Tobacco mosaic virus* (TMV) [32]. This finding substantiates the death-promoting effect of enhanced autophagy during ETI [33], and explains the increased TMV accumulation previously noted in HR lesions of autophagy-deficient *N. benthamiana* leaves [34]. Furthermore, it adds to the emerging picture that the positive role of autophagy in immunity-related PCD is opposite to its function in preventing premature senescence and runaway cell death outside of the primary infection sites [28,35].

How autophagy exerts the dual roles during HR activation and containment is not well understood. The influence of autophagy on cellular survival is likely linked to homeostatic functions required to counterbalance infection-induced systemic responses such as ROS production, salicylic acid (SA) signaling, accumulation of misfolded/aggregated proteins, and endoplasmic reticulum stress [26,28,36]. In contrast, the pro-death mechanism of autophagy remains largely undefined, but may also involve the regulation of SA homeostasis and/or the level of NON-EXPRESSOR OF PATHOGENESIS-RELATED GENES 1 (NPR1), that negatively impacts HR [26,28,37]. Future work could further address the potential engagement of selective autophagic processes, e.g. in the removal of negative HR regulators [35].

There is compelling evidence and a broad consensus that autophagy positively controls plant resistance to necrotrophic pathogens (Figure 1). Autophagy deficiency in Arabidopsis mutants resulted in spreading necrotic lesions and enhanced fungal growth upon infection with *B. cinerea, Alternaria brassicicola, and Plectosphaerella cucumerina* [38-40], and restored susceptibility to a non-pathogenic mutant strain of *Sclerotinia sclerotiorum* [13]. Notably, autophagy-mediated disease resistance to *B. cinerea* engages the upstream regulator BAG6 (BCL2-ASSOCIATED ATHANOGENE FAMILY PROTEIN 6) [41]. While Arabidopsis *bag6* mutants were defective in autophagy induction and hypersusceptible to *B. cinerea*, ectopic expression of BAG6 in *N. benthamiana* leaves activated autophagy and cell death, which prevented fungal infection [41]. Hence, pathogen-induced necrotic cell death and disease development is restricted by autophagy and/or immunity-related (autophagic) PCD. This mechanism agrees with the inhibition of necrosis by autophagy during execution of vacuolar cell death in development [42]. The molecular basis of the

crosstalk remains largely unknown, although it is evident that protection from *B. cinerea* infection occurs independently of selective autophagy mediated by the cargo receptor NEXT TO BRCA1 GENE 1 (NBR1) [29]. Resistance to necrotrophs may be also mediated by autophagy via modulation of hormone homeostasis, e.g. to stimulate jasmonic acid (JA) defence signaling removal of plant- and removal of pathogen-derived toxic cellular constituents [39].

In animals, autophagy is a key mechanism in the fight against invading intracellular bacterial and viral pathogens. In contrast, there is surprisingly little knowledge about the contribution of autophagy to basal resistance against viruses, the major intracellular pathogens in plants. Autophagy has been associated with plant antiviral RNA silencing by mediating the targeted degradation of viral silencing suppressors including the cucumovirus protein 2b and potyvirus protein HCpro [43]. Interestingly, potyviral challenge of Arabidopsis lines with reduced expression of the negative autophagy regulator TARGET OF RAPAMYCIN (TOR) revealed strongly decreased levels of Watermelon mosaic virus, whereas Turnip mosaic virus accumulation was only slightly affected [44]. Although the significance of these findings has yet to be verified under autophagy-deficient conditions, they imply an antiviral role of autophagy against some potyviruses, and potentially other unrelated viral species. In this context, it remains to be determined whether autophagy can directly eliminate viruses in a process similar to mammalian xenophagy [45].

Finally, the role of autophagy in basal resistance to (hemi)biotrophic pathogens is a matter of ongoing debate. So far, there is no evidence that autophagy is directly involved in the regulation of MTI. In addition, despite some conflicting results, autophagy deficiency seems to rather enhance resistance to the virulent bacterial strain *Pst* DC3000 and some powdery mildew fungal species [9]. These findings could be partly linked to the impact of autophagy on SA levels and signaling, which might be further tested in plant systems with enhanced autophagy levels.

Pathogen manipulation and pro-microbial role of autophagy

Considering the long-lasting co-evolutionary battle between plants and their pathogens, it is not surprising that successful microbes have evolved sophisticated strategies to modulate autophagy to their benefit (Figure 1).

The necrotroph *S. sclerotiorum* requires the phytotoxin oxalic acid (OA) to trigger unrestricted host cell death and establish successful infection. OA-deficient mutants are non-pathogenic and activate autophagy leading to restrictive HR-like cell death and resistance [13]. Autophagy deficiency restored pathogenicity, indicating that *S. sclerotiorum* secretes OA to suppress antimicrobial autophagy. A similar autophagy-mediated mechanism operates in the non-host *Ustilago maydis*-barley interacton. The biotrophic smut fungus *U. maydis* is recognized by barley, triggering a defense response that neutralizes the pathogen and prevents disease, but results in large necrotic areas and stunted leaf growth. In contrast, *U. maydis* mutants lacking the *Pep1* effector show hallmarks of autophagy at the attempted penetration site and remain restricted to the infected area, which might indicate that *Pep1* is an autophagy inhibitor [12]. These findings suggest that autophagy suppression might be a virulence strategy shared by pathogens with completely different lifestyles.

In line with this notion, binding and activation of TOR by the Cauliflower mosaic virus (CaMV) P6 protein has recently been proposed to inhibit autophagy and impact resistance responses to bacterial pathogens [14]. CaMV infection and transgenic expression of P6 increased the susceptibility to *Pst* DC3000 infection and facilitated growth of the effector-delivery deficient *Pst* mutant *hrc*⁻. This effect appears to be in agreement with P6-induced impairment of MTI responses including oxidative burst and SA accumulation. However, it would be surprising if P6 suppression of autophagy is causally linked to the observed phenotype, as *atg* mutants have been shown to display enhanced rather than reduced SA levels and bacterial resistance [38]. Hence, future efforts need to clarify the involvement of autophagy during CaMV infection and to reveal the potential role of TOR-binding of P6 to modulate this pathway for enhanced pathogenicity.

Other pathogens induce autophagy as part of their infection strategy. For example, the secreted effector AWR5 from the bacterium *Ralstonia solanacearum* inhibits TOR, which results in the activation of autophagy [17]. Although the mechanistic details of this host-pathogen interaction remain to be elucidated, a tantalizing scenario would be that autophagy induction in the host stimulates plant cell dismissal and metabolic re-routing. This would be beneficial for *R. solanacearum* during its transition to the necrotrophic phase by facilitating nutrient acquisition. Viral pathogens might also promote and hijack

autophagy pathways to invade host cells. For instance, the viral silencing suppressor P0 was shown to trigger autophagic degradation of ARGONAUTE1, an essential component of antiviral RNA-induced silencing complexes [16]. Given the frequent connections between viruses and autophagy in animals [46], future research will most likely provide more cases of virus-induced autophagic degradation of antiviral defense components in plants, perhaps even including small RNAs.

Another interesting example for the manipulation of the host autophagy machinery by a plant pathogen comes from the hemibiotrophic oomycete *Phytophthora infestans*. The RXLR effector protein PexRD54 was shown to bind to a specific host ATG8 protein, which prevented interaction of ATG8 with the autophagy cargo receptor Joka2/NBR1 [15]. Joka2-mediated selective autophagy was further reported to positively influence plant resistance to *P. infestans*; hence, depletion of Joka2 by PexRD54 enhances susceptibility of the host. Interestingly, both Joka2 and PexRD54 trigger the formation of autophagosomes and activate autophagy. This led the authors to speculate that Joka2 facilitates removal of plant or pathogen proteins that negatively impact immunity, whereas PexRD54 might co-opt the autophagy pathway to selectively eliminate defense-related compounds or to recycle and redistribute nutrients in favor of the pathogen.

Conclusions / Future directions

This review highlights the importance of autophagy in the field of plant-pathogen interactions. Autophagy has emerged as a central part of the plant weaponry against invading microbial pathogens. Its significance for plant defense is supported by the evolution of microbial strategies to manipulate the host autophagy machinery for enhanced virulence and disease establishment. In addition, autophagy in eukaryotic phytopathogens has evolved as an essential process in the development of functional infection structures. However, the examples illustrating the key roles of autophagy in plant-biotic interactions are still limited both in number and mechanistic detail. Current efforts in several laboratories around the world will certainly help to revert this situation in the coming years and further reveal the highly complex and multifaceted integration of autophagy into the plant immune system.

A key direction of future research will be the identification and characterization of selective autophagy receptors that drive plant defense responses and are still hidden in the gray shades of "bulk" autophagy. In a more refined interaction, we envisage that plants employ and pathogens manipulate particular selective autophagy pathways to benefit defense and disease, respectively. So far, very few autophagy cargo receptors and their substrates have been identified in plants, but the generally very complex outcome of disease in autophagy deficient plants may indicate that selective processes with distinct functions operate in parallel within the full autophagy response. To dissect these mechanisms in greater detail, we need to establish plant lines with increased "bulk" autophagy to support conclusions from knock-out mutants, and complement these general systems by targeting specifically the different selective autophagy pathways. In addition, due to concomitant, often overlapping roles of autophagy in cellular homeostasis and various developmental and environmental stress responses, it is essential to more precisely inhibit or activate autophagy by inducible and cell type-specific approaches.

Another important area of research relates to the largely unexplored crosstalk between autophagy and other cellular pathways that govern proteostasis, hormone signaling, and programmed cell death in plant-microbe interaction. Notably, the plant ubiquitin-proteasome system was recently found to be degraded by autophagy in response to nutrient starvation or chemical and genetic proteasome inhibition [47]. Whether a similar interplay occurs during immunity and disease is not known; however, recent evidence indicates that the 26S proteasome is central to plant immunity and targeted by multiple pathogen effectors to suppress SA-mediated host defenses [48].

 Overall, there are still only very few pathogens identified that directly modulate the plant autophagy machinery to the benefit of infection. Among these, suppression of autophagy seems to be most common strategy, whereas the potential subversion of bulk and selective pathways still remains merely speculative. However, the fundamental role of autophagy in host immunity and microbial pathogenesis anticipates that phytopathogens have evolved sophisticated capacities to evade and exploit autophagy as demonstrated for a multitude of metazoan pathogens, thus adding further complexity to this emerging arena of plant-microbe interactions.

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Figure Legend

Figure 1: Anti- and pro-microbial roles of autophagy during plant-pathogen interactions.

Autophagy is an integral part of plant immunity. Arabidopsis infection with avirulent *Pseudomonas syringae* pv. tomato DC3000 (*avrRps4*) induces autophagy, which contributes to the hypersensitive response (HR) and disease resistance. Infection of Arabidopsis with the necrotrophic fungus *Botrytis cinerea* triggers cleavage of the BAG6 protein, which results in autophagy activation and reduced disease development. Plant autophagy also participates in antiviral defense by targeted degradation of viral silencing suppressors such as the potyvirus protein HCpro and the cucumovirus protein 2b.

Plant pathogens manipulate the host autophagy machinery to counteract host defense and promote virulence. *Phytophthora infestans* effector PexRD54 binds ATG8 and outcompetes the plant selective autophagy receptor Joka2 from autophagosome association, thereby enhancing disease susceptibility of the host. The AWR5 effector from *Ralstonia solanacearum* inhibits TOR to activate autophagy, which is presumed to be beneficial for nutrient acquisition and successful infection. *Sclerotinia sclerotiorum* secretes the toxin oxalic acid to suppress autophagy and HR-like autophagic cell death as part of the host defense response against necrotrophic infection.

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Highlights (3-5 bullet points 85 characters each)

- Autophagy is an integral part of plant-pathogen interactions.
- A large variety of microbial pathogens target or are targeted by plant autophagy.
- Autophagy in eukaryotic microbial pathogens is essential for pathogenesis.
- Plant autophagy participates in defense responses against invading microbes.
- Successful pathogens have evolved strategies to manipulate plant autophagy.