

# Autophagy-Virus Interplay: From Cell Biology to Human Disease

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## Introduction

This review investigates the various functions cellular autophagy has in fighting off viral infection of the cell.

## Conclusions

Autophagy can be used in a classical way by binding viral proteins via p62 and SMURF through ubiquitin dependent and independent autophagy. It then allows antigens (proteins from the virus) to be presented via MHC to immune cells for antibody creation. Autophagy associated proteins can also inhibit pro-inflammatory signals by binding protein complexes responsible for exciting pro-inflammatory cytokines (communication proteins/signals). Autophagy can also destroy damaged mitochondria that may be increasing pro-inflammatory signaling, aberrantly, by releasing sizeable amounts of reactive oxygen species.

## Amendments



## Autophagy-Virus Interplay: From Cell Biology to Human Disease

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Autophagy is a highly conserved intracellular degradation process that targets protein aggregates and damaged organelles. Autophagy is also implicated in numerous viral infections, including human immunodeficiency virus-1 (HIV-1), influenza A (IAV) and herpes simplex virus-1 (HSV-1). Depending on the virus, autophagy can restrict or promote viral replication, and play key roles in modulating inflammation and cell survival. In this review, we consider examples of autophagy-virus interplay, highlighting the protective role of autophagy in human infections. We summarize recent discoveries and emerging themes illuminating autophagy's role in immunity and inflammation upon viral infection. Finally, we discuss future prospects and therapeutic implications, and potential caveats associated with using autophagy to control viral infections in humans.

**Keywords:** autophagy, inflammation, HIV-1, HSV-1, IAV, viral immunity

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### INTRODUCTION

Autophagy captures cytoplasmic contents, such as excess or defective proteins and organelles, for degradation by the lysosome. It is initiated in response to various stimuli, including nutritional state of the cell and environmental stresses such as starvation and hypoxia. As such, autophagy is an important process of regulating cellular homeostasis and survival. It is a well-studied process that is orchestrated by over 35 autophagy-related (ATG) proteins and can be organized into multiple steps: phagophore initiation, membrane elongation, autophagosome formation and autophagosome fusion with hydrolytic lysosomes (Figure 1A; Mizushima et al., 2011). Autophagy can be selective in terms of cargo capture via the recruitment of selective autophagy receptors.

Autophagy receptors can interact with ubiquitin tags that decorate the cargo [via its ubiquitin-binding domain (UBD)], and with LC3 proteins of nascent autophagosomes [via its LC3-interacting region (LIR) motif] (Stolz et al., 2014). Some autophagy receptors, particularly p62/SQSTM1 and optineurin, are regulated by Tank-binding kinase 1 (TBK1)-mediated phosphorylation, and are key players in the autophagic degradation of invasive pathogens (Wild et al., 2011; Pilli et al., 2012; Sparrer et al., 2017).

Autophagy plays a key role in cellular immunity to human infections (Randow et al., 2013; Paul and Münz, 2016). In the case of human viral infections, autophagy can be either pro-viral or antiviral (Levine et al., 2011; Jackson, 2015). Some viruses hijack the autophagy machinery for their intracellular survival, while others express specific proteins to evade autophagy and propagate in host cells. Antiviral autophagy can (1) selectively target pathogens for degradation, (2) promote pathogen recognition and inflammatory cytokine responses, (3) regulate inflammation, (4) control cell survival, (5) promote antigen presentation and/or (6) be regulated in a paracrine-mediated fashion, a departure from the classic cell autonomous route (Figure 1B; Levine, 2005;

1. Autophagy can be specific in its targeting by binding to particular ubiquitin tags (ubiquitin is a protein that attaches to proteins that are targeted for degradation), as well as LC3 proteins, which are found in the autophagosome (the vesicle that encircles the broken proteins).

Levine et al., 2011; Jackson, 2015; Paul and Münz, 2016). By focusing on these themes of antiviral autophagy, this review will highlight the protective nature of autophagy. We will draw from examples of human viral infections representing significant disease burden, whose interplay with autophagy has been supported by experimental and/or clinical evidence.

**PATHOGEN OR VIRAL ANTIGEN CLEARANCE BY AUTOPHAGY**

Selective autophagy has been reported to control several human viral infections *in vitro*, leading to the clearance of pathogens or viral antigens and host cell survival. One of the first viruses shown to engage the autophagy machinery was Sindbis virus (SV), a positive-stranded RNA alphavirus that typically causes mild disease in humans. Autophagy receptors p62 and SMURF1 were found to target SV capsid, in a ubiquitin-independent manner, to autophagosomes in human HeLa cells and mouse embryonic fibroblasts (MEFs) (Orvedahl et al., 2010, 2011). Depletion of p62 in HeLa cells or Atg5 in mouse neurons *in vivo* led to an accumulation of toxic SV capsid and higher virus-induced mortality, without altering SV replication (Orvedahl et al., 2010). Whether SV can express proteins to counteract autophagy is not yet known.

Similar to SV, p62 is involved in targeting toxic Chikungunya virus (CHIKV) capsid to autophagosomes in HeLa cells, but in a ubiquitin-dependent manner (Judith et al., 2013). CHIKV, also a positive-stranded RNA alphavirus, is a mosquito-borne virus causing severe pathologies in humans that range from febrile arthralgia, rash to encephalopathy (Couderec and Lecuit, 2015). siRNA-mediated depletion of p62 led to an increase in both cell mortality and CHIKV replication, while p62 overexpression promoted cell viability in CHIKV-infected cells (Judith et al., 2013). In contrast, another autophagy receptor NDP52 was shown to have a proviral role although overall autophagy played a cytoprotective role in CHIKV infection (Judith et al., 2013).

Human immunodeficiency virus-1 (HIV-1), a lentivirus that is transmitted sexually through infected body fluids, is targeted by autophagy in human CD4+ T cells *in vitro* (Sagnier et al., 2015). HIV-1 primarily infects CD4+ T cells, consequently compromising the individual's immune defense and leading to acquired immunodeficiency syndrome (AIDS) (Maartens et al., 2014). HIV-1 integrates its DNA into the host's genome, and recruits its transactivator protein Tat to activate viral transcription. Tat protein is therefore essential for HIV-1 replication. However, Tat protein is targeted by p62 in HEK293T cells, which directs it to autophagosomes in a ubiquitin-independent manner (Sagnier et al., 2015). Consistent with this, depleting p62 in HEK293T cells resulted in accumulation of Tat protein, and enhancing autophagy in a chronically HIV-1-infected T cell line led to reduction of Tat protein levels (Sagnier et al., 2015). Furthermore, peripheral blood mononuclear cells (PBMCs) from HIV-1-infected non-progressor individuals showed higher number of HIV-1 particle-containing autophagic vesicles compared with HIV-1-infected normal progressors, suggesting a role for autophagy in limiting

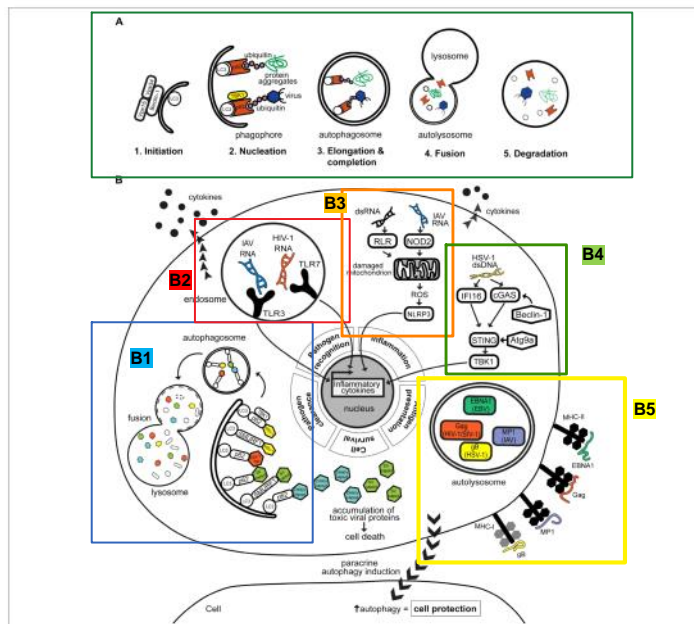
the pathogenesis of HIV-1 *in vivo* (Nardacci et al., 2014). However, HIV-1 can modulate autophagy and expresses multiple autophagy inhibitors, such as Vif, Nef, and Env, which operate in a cell type-specific manner (Liu et al., 2017). Hence, autophagy can be viewed to control HIV-1 replication by targeting viral components to degradation in specific cell types.

Herpes simplex virus-1 (HSV-1) is a ubiquitous, neurotropic  $\alpha$ -herpesvirus with a global seroprevalence of 67% (Locker et al., 2015). It typically manifests as benign, self-limiting mucocutaneous ulcers, but in rare cases may cause life-threatening herpes simplex encephalitis (HSE) (Whitley and Bozeman, 2001). Reports have identified HSV-1-encoded ICP34.5 and Us11 as autophagy inhibitors, which exert their effects by targeting Beclin-1 and protein kinase R (PKR), respectively (Orvedahl et al., 2007; Lussignol et al., 2013). Studies in primary MEFs and mice described increased autophagy following infection with ICP34.5-deficient HSV-1, as suggested by an increased number of autophagosomes and virions in neuronal autophagosomes, respectively (Tallóczy et al., 2006; Alexander et al., 2007). During HSV-1 infection, autophagy appears to be operating in a cell type-specific manner (Yordy et al., 2012). For example, in contrast to mitotic cells such as MEFs and mouse keratinocytes, post-mitotic mouse primary neuronal cells predominantly use autophagy over interferon (IFN) as a viral control mechanism (Yordy et al., 2012). In MEFs, HSV-1 was shown to be selectively targeted by p62 and SMURF1. HSV-1-infected *smurf1*<sup>-/-</sup> MEFs failed to target HSV-1 virions to autolysosomes, resulting in more HSV-1 virions in autophagosomes (Orvedahl et al., 2011). In mouse primary trigeminal neurons *in vitro*, HSV-1 infection triggered the formation of p62-mediated autophagosomes (Katzenell and Leib, 2016). Finally, infection of HEK293T cells with ICP34.5-deficient HSV-1 led to the recruitment of TBK1- and p62-mediated autophagy, and viral restriction (Sparrer et al., 2017). However, the exact viral target of p62 and SMURF1, and the role of ubiquitin in mediating receptor-virus interactions, is not yet known. Reports using human HFFF2 fibroblasts and mouse dendritic cells (DCs) have shown that autophagy is triggered by HSV-1 double-stranded DNA (dsDNA), and is independent of viral replication but dependent on STING (McFarlane et al., 2011; Rasmussen et al., 2011).

**AUTOPHAGY AND CELLULAR IMMUNITY: PATHOGEN RECOGNITION AND CYTOKINE RESPONSES**

Cellular immunity requires the detection of viral pathogen-associated molecular patterns (PAMPs) by their cognate receptors to produce antiviral cytokines, such as type-I IFNs (Iwasaki, 2012). Autophagy can deliver viral PAMPs to their receptors, and help amplify the production of inflammatory cytokines. The capacity of autophagy to facilitate viral recognition and modulate downstream cytokine production has been demonstrated in the case of HIV-1 infections. During HIV-1 infection in human primary plasmacytoid DCs, autophagy plays a key role in presenting the HIV-1 RNA genome to its cognate immune

2. Proteins within the cell (p62 and SMURF1) attach to a viral capsid (the "shell" of the virus containing the viral RNA) and this targets the virus for autophagy (in a ubiquitin dependent manner, as explained in note 1),
3. P62, when repressed, leads to more cellular death (presumably, because autophagy is not functional to clear out virus); however, when artificially overexpressed, the cells survive better.



A. Initiation by the congregation of several key autophagy proteins onto a vesicle/membrane is known as initiation - then, once p62 binds LC3, the ubiquitinated (tagged) viral proteins are linked to p62 and initiate nucleation into a structure known as the phagophore - once the vesicle closes, it forms the autophagosome, trapping the viral proteins within. Then, a lysosome binds/fuses the autophagosome to create the autolysosome - this leads to an increase in acidity and the introduction of degradation enzymes to destroy the viral proteins.

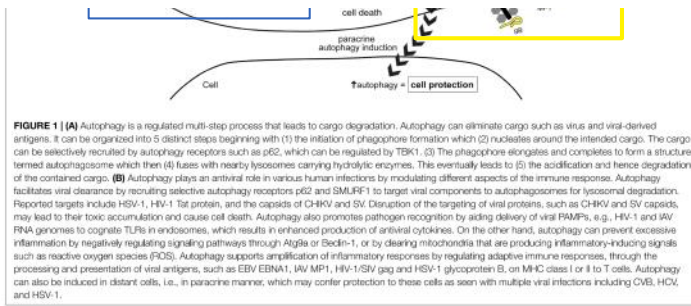
B1. Here, autophagy is occurring as described in A, but specifically binding p62 and SMURF with viral proteins to be degraded. This is the classic attack of autophagy.

B2. If the autophagosome does not undergo "maturation" wherein the lysosome binds to create the aforementioned autolysosome, this vesicle/endosome will simply trap the viral proteins, DNA, and RNA in with "Toll Like Receptors" (TLR), which will bind these antigens (viral DNA/RNA, etc.) and promote the production of cellular cytokines (inflammation signals) to raise the alarm to the rest of the cell, as well as other cells.

B3. If mitochondria are damaged from the introduction of viral proteins, DNA, and RNA, they will generate more reactive oxygen species, thereby stimulating more cytokine (inflammatory signals) production and release from the cell. To control this, autophagy will eliminate damaged mitochondria, known as mitophagy.

B4. Proteins implicated in the autophagy machinery (Beclin, ATG9) will bind particular subcomponents of the STING complex of proteins, which would normally translocate to the nucleus and bind genes that promote inflammatory cytokines (communication/signaling proteins). Since overdoing the inflammatory signaling can cause the cell to die or the over recruitment of immune cells, these autophagy proteins (Beclin and ATG9) binding the STING complex proteins inhibits their function (by inhibiting their ability to form the STING complex) and inhibiting their translocation to the nucleus thereby controlling/decreasing the inflammatory signaling from the cell.

B5. With the formation of the autolysosome (from 4,5 of autophagy), the viral proteins are degraded into



receptor Toll-like receptor-7 (TLR7) in endosomes, leading to the induction of IFN $\alpha$  (Lee et al., 2007; Zhou et al., 2012). Silencing the expression of ATG7 in plasmacytoid DCs *in vitro* leads to a significant decrease in IFN $\alpha$  production following HIV-1 infection, highlighting a crucial role for autophagy in mediating TLR7-IFN signaling (Zhou et al., 2012).

signaling can cause the cell to die or the over recruitment of immune cells, these autophagy proteins (Beclin and ATG9) binding the STING complex proteins inhibits their function (by inhibiting their ability to form the STING complex) and inhibiting their translocation to the nucleus thereby controlling/decreasing the inflammatory signaling from the cell.

B5. With the formation of the autolysosome (step 4-5 of autophagy), the viral proteins are degraded into smaller, less functional pieces; however, once degraded, the cell presents these pieces outside of itself, to immune cells, through major histocompatibility complexes (MHC) that take these pieces of viral protein (called antigens) and allows immune cells to take them up and create antibodies to these antigens.

B3

Autophagy also plays an important role in mediating cytokine production during infection by influenza A virus (IAV). IAV, an RNA virus, is a pandemic threat and global health concern. It targets epithelial cells of the respiratory tract, and in severe cases may cause pneumonia or pulmonary damage (Paules and Subbarao, 2017). In the case of infection with highly virulent IAV strains H1N1 and H5N1, high morbidity has been attributed to excessive host-induced inflammatory cytokine production (Peris et al., 2010). IAV infection was shown to induce autophagy in primary human blood macrophages, which regulates the production of CXCL10 and IFN $\alpha$ . When these cells were depleted of ATG5 or treated with the autophagy inhibitor 3-methyladenine (3-MA), they produced lower CXCL10 and/or IFN $\alpha$  levels (Law et al., 2010). The precise mechanism by which autophagy facilitates this aspect of IAV infection in human blood macrophages is unknown, but is thought to involve recognition of viral RNA by endosomal TLR3 (Law et al., 2010). In contrast, IAV-induced autophagy in MEFs can prevent IFN $\beta$  production, and enhance viral replication (Perot et al., 2018). The induction of autophagy during IAV infection is complex: it is initially cytoprotective but is later counteracted by the IAV matrix protein 2 (M2) which targets Beclin-1 to block lysosomal fusion with autophagosomes (Gannagé et al., 2009). Taken together, these data suggest an immunopathological role of autophagy in controlling cytokine production and IAV infection, in potentially a tissue-specific manner.

From the examples of HIV-1 and IAV, autophagy plays a fundamental role in modulating the primary antiviral response, by promoting viral recognition through TLR-dependent signaling and inflammatory cytokine production.

### ANTI-INFLAMMATORY ACTIONS OF AUTOPHAGY

In addition to promoting inflammation, autophagy is crucial for preventing prolonged and excessive inflammation detrimental to the host. Components of the autophagy machinery, such as Beclin-1 and Atg9a, interact with the cytoplasmic type-I-IFN-inducing STING-TBK1 pathway. A study found that Beclin-1 interacts with dsDNA sensor cGAS to dampen IFN $\beta$  production in HEK293T cells stimulated with dsDNA or infected with HSV-1 (Liang et al., 2014). This interaction also leads to clearance of dsDNA through autophagy, limiting the otherwise persistent IFN $\beta$ -mediated inflammation (Liang et al., 2014). The depletion of Beclin-1 was shown to increase cGAS-mediated IFN $\beta$  production while reducing HSV-1 replication in RAW264.7 mouse macrophages (Liang et al., 2014). In MEFs stimulated with dsDNA, Atg9a had a similar role and negatively regulated the STING-TBK1-IFN pathway by binding to STING and preventing its assembly with TBK1 in LC3-positive structures (Saitoh et al., 2009). Atg9a-knockout mice revealed an increase in IRE3 phosphorylation and IFN $\beta$  production following dsDNA stimulation (Saitoh et al., 2009). However, whether HSV-1 infection (or infection of other dsDNA viruses) is subject to Atg9a-mediated regulation remains unknown.

Autophagy may also negatively regulate inflammation indirectly by clearing host DAMPs, such as reactive oxygen species (ROS) released by mitochondria. This has been observed in mice BMDCs where IAV genomic RNA is detected by the NOD2-RIPK2 pathway, which activates ULK1 to induce RIPK2-mediated autophagic clearance of damaged mitochondria (Lupler et al., 2013). In RIPK2-deficient mice BMDCs infected with IAV, mitochondria accumulated in cells resulting in elevated production of superoxide. This led to the hyperactivation of the NLRP3 inflammasome and an increased secretion of the inflammatory cytokine interleukin (IL)-18 (Lupler et al., 2013). A similar observation was made in Atg5-deficient MEFs and mouse primary macrophages stimulated with dsRNA analog poly(I:C), which led to excessive RIG-I-like-receptor signaling (Tal et al., 2009). Furthermore, ectopic P-granules autophagy protein 5 homolog (EPG5; a protein that regulates autolysosomal formation) has been shown to control pulmonary inflammation. Lung macrophages from Epg5 $^{-/-}$  mice showed excessive production of inflammatory IL-1 $\beta$  and IL-13 cytokines, resulting in resistance to IAV infection (Lu et al., 2016).

These reports highlight the important role of autophagy in attenuating inflammation. The autophagy machinery can limit inflammation by regulating cytosolic NLR- and STING-mediated signaling pathways through disposal of their ligands, inactivation of their cognate receptors or interaction with their downstream effector molecules.

### AUTOPHAGY AND CELL SURVIVAL

As shown from investigations of multiple human viral infections, autophagy plays a role in promoting cell survival and limiting pathogenesis. This has been demonstrated by the ability of mouse L cell mutant gro29 cells which have high basal autophagy to restrict HSV-1 replication (Le Sage and Banfield, 2012). In contrast, Atg161<sup>HM</sup> mice, which have reduced basal autophagy, showed high mortality following infection with CHIKV *in vivo* (Joubert et al., 2012). One mechanism by which autophagy may promote cell survival during viral infection is by degrading and preventing accumulation of toxic viral proteins, such as viral capsids, in the infected cells. This has been demonstrated in the case of CHIKV and SV infected cells, as discussed above (Orvedahl et al., 2010; Judith et al., 2013). Furthermore, reports have documented the cytoprotective effect of autophagy-enhancing drugs, such as vitamin D, MG132 and rapamycin, in viral infections. For example, primary human macrophages have shown benefit from vitamin D treatment, which limits HIV-1 replication *in vitro* (Campbell and Spector, 2012). Treating HSV-1-infected human corneal epithelial (HCE) cells with MG132 can reduce viral titres (Yakoub and Shukla, 2015). Moreover, pre-conditioning human fibroblasts *in vitro* with rapamycin has been shown to promote cell survival following HSV-1 infection (Ahmad et al., 2018). This cytoprotective role for autophagy that occurs early in HSV-1 infection appears to be a TBK1-dependent process (Ahmad et al., 2018). In agreement with this, TBK1

DAMPs: Damage Associated Molecular Patterns - created by the cell, like the mitochondria, to lead the cell to die.

**TABLE 1 |** Opportunities for autophagy-modifying therapeutic intervention in human viral diseases.

Virus	Available treatment for acute disease	Vaccine	Post-infectious inflammation/autoimmunity
CHIKV	None	No	Rheumatic inflammation
HIV-1	Antiretroviral therapy (ART)	No	HIV-associated neuroinflammation
HSV-1	Nucleoside analog (e.g., acyclovir)	No	Anti-N-methyl-D-aspartate receptor (NMDAR) encephalitis
IAV	Neuraminidase inhibitors (e.g., oseltamivir)	Yes	Acute infection-induced cytokine storm

deficiencies render human fibroblasts susceptible to HSV-1 infection and leads to increased cell mortality (Herman et al., 2012). These data support literature showing that TBK1 is a key player in protective autophagy against bacterial infections (Weidberg and Elazar, 2011), and extend its protective role to viral infection.

### AUTOPHAGY AND ADAPTIVE IMMUNITY: ANTIGEN PRESENTATION

Through its degradative function, autophagy is particularly useful for generating endogenous peptide antigens for major histocompatibility complex (MHC)-II presentation (Dengjel et al., 2005; Paul and Münz, 2016). In viral infections, autophagy generates viral antigens loaded onto MHC-I and MHC-II for presentation to T cells (Münz, 2017). Epstein-Barr virus (EBV) is an oncogenic  $\gamma$ -herpesvirus causing a spectrum of human diseases ranging from mononucleosis to lymphomas and carcinomas (Taylor et al., 2015). Historically, EBV nuclear antigen 1 (EBNA1) was one of the first viral antigens shown to be processed by autophagy and loaded on MHC-II molecules of EBV-transformed B cell lines (Paludan et al., 2005). Inhibition of autophagy leads to accumulation of EBNA1 in autophagosomes of EBV-transformed lymphoblastoid cell lines, and a decrease in EBNA1-specific CD4<sup>+</sup> T cell recognition via MHC-II (Paludan et al., 2005).

As a result of reduced MHC-II antigen presentation, mice with *Atg5*-deficient DCs intradermally injected with HSV-1 showed significantly lower IFN $\gamma$  production by CD4<sup>+</sup> T cells (Lee et al., 2010). In addition, autophagy can deliver viral antigens for MHC-I cross-presentation. Using a mouse BMA3.1A7 macrophage cell line for CD8<sup>+</sup> cell stimulation, HSV-1 glycoprotein B (gB) was presented on MHC-I in an autophagy-dependent manner (English et al., 2009; Radtke et al., 2013).

Autophagy is also vital for efficient stimulation of antiviral CD4<sup>+</sup> T cells in HIV-1/Simian immunodeficiency virus (SIV) and IAV infections. Knocking down LC3 protein or inhibiting autophagy using 3-MA in human DCs led to reduced antigen processing and MHC-II presentation, and a decrease in HIV-1-specific CD4<sup>+</sup> T cell response (Blanchet et al., 2010). On the other hand, enhancing autophagy in human DCs with rapamycin resulted in a more pronounced HIV-1-specific CD4<sup>+</sup>

T cell response (Blanchet et al., 2010). Fusing SIV gag protein to LC3 in mice BMDCs was also shown to improve antigen-specific CD4<sup>+</sup> T cell responses *in vitro* (Jin et al., 2014). Similar results were obtained *in vivo* where immunizing mice with SIV gag-LC3 resulted in a stronger humoral immune response, with CD4<sup>+</sup> T cells producing higher levels of IFN $\gamma$ , TNF $\alpha$  and IL-2 (Jin et al., 2014). Conjugating IAV matrix protein 1 (M1) to LC3 in HaCat human epithelial cells, B cells and DCs led to enhanced antigen-specific human CD4<sup>+</sup> T cell responses *in vitro*, as measured by IFN $\gamma$  (Schmid et al., 2007).

Taken together, autophagy can perpetuate the initial response to viral infection by priming and mediating T cell responses of the adaptive immune system to ensure effective viral clearance.

### BEYOND CELL AUTONOMOUS IMMUNITY: PARACRINE REGULATION OF AUTOPHAGY

Since its discovery, the primary focus of autophagy research has been to investigate its role on a cell autonomous level. Interestingly, two recent reports have demonstrated that autophagy can also be triggered at a cell population level (i.e., in a paracrine manner) to affect distant cells. A first report showed that autophagy could be triggered in distant and distinct cell types that can protect them from a variety of viral infections (Delorme-Axford et al., 2013). In this case, primary human placental trophoblasts can protect other cells from coxsackievirus B3 (CVB), hepatitis C virus (HCV), vesicular stomatitis virus (VSV) and vaccinia virus (VACV), by secreting signals that induce autophagy to resist infections. This concept is particularly relevant in the womb, allowing maternal trophoblasts to confer resistance to viral infections to the growing fetus.

A second report described the paracrine regulation of autophagy early in HSV-1 infection (Ahmad et al., 2018). In this case, HSV-1 infection of human fibroblasts was shown to induce autophagy in cells neighboring an infection site. Despite having functional basal autophagy, HSE patient-derived fibroblasts deficient in TBK1 specifically failed to mount paracrine-mediated autophagy during HSV-1 infection. The study further showed that autophagy induction early during infection may protect cells from death. The autophagic role of TBK1 has previously been associated with inflammation control in neurodegenerative amyotrophic lateral sclerosis (ALS) (Freischmidt et al., 2015). These observations highlight a potential involvement of TBK1 in controlling neuroinflammation through autophagy in HSE.

### CONCLUSION

Many open questions remain concerning the precise role of autophagy in human viral infections. Studies looking at human responses *in vivo* are rare, due to difficulty of conducting these studies. However, a wealth of studies using animal

*in vitro/ex vivo/in vivo* and human *in vitro/ex vivo* models have given remarkable insights into the role of autophagy in disease manifestation.

In this review, we discuss human viruses modulated by autophagy that represent a significant clinical burden. We highlight how autophagy is protective and may be used to enhance current treatment options (Table 1; Whitley and Roizman, 2001; Mescon, 2005; Maertens et al., 2014; Couderc and Lecuit, 2015). Several reports have shown the protective role of p62-mediated selective autophagy in various human pathogens (e.g., CHIKV, HIV-1, and HSV-1), making p62 an attractive therapeutic target. Enhancement of autophagy through p62 may provide an important therapeutic avenue for treatment of human viral diseases (Table 1). However, p62 also participates in other biological processes, such as cell proliferation and ubiquitin-proteasomal degradation (Liu et al., 2016b), and research focusing on employing p62 for therapeutic benefit should be aware of potential pleiotropic effects. The ability of p62 to interact with viruses independent of its conventional ubiquitin-binding domain also warrants further investigation. As shown in the case of HSV-1 and HIV-1 infections, augmenting autophagy using stimulants (such as rapamycin and vitamin D) can be beneficial to restrict viral replication and/or promote cell survival. Enhancing autophagy in vaccine therapies has also been beneficial, taking advantage of the role of autophagy in antigen priming. Promising results were observed in the case of IAV and HIV-1/SIV-1 infections, whereby increasing the autophagic targeting of viral protein gave rise to a heightened adaptive immune response (Schmid et al., 2007; Jin et al., 2014). Moreover, autophagy is important for both MHC-I and/or -II antigen presentation in HSV-1 and HIV-1 infections, as well as for regulating inflammation by facilitating antiviral inflammatory cytokine production. Autophagy's role in fine-tuning inflammation is also important during IAV infection, where it promotes inflammatory cytokine production and prevents excessive inflammatory responses. In addition to the effects on acute disease outcome, modulating autophagy may have a promising role in the prevention or treatment of various viral post-infectious inflammation/autoimmune disorders for which there are limited treatment options (Table 1; Kovalevich and Langford, 2012; Armangue et al., 2014; Couderc and Lecuit, 2015; Liu et al., 2016a). Harnessing autophagy's inflammation-reducing capacity can potentially prevent the development of these states, or help to resolve the inflammatory symptoms.

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Current research on autophagy is mostly focused on its role in cell autonomous immunity (Randow et al., 2013). However, recent studies have revealed a novel type of autophagy triggered in a paracrine manner in response to viral infections. Elucidating the role of paracrine-regulated autophagy may prove to be highly relevant in disease pathogenesis *in vivo*, and may be useful as a method of clinical intervention. It is tempting to speculate that similar mechanisms could also be extended to viral pathogenesis that disseminates to sensitive tissues, such as the central nervous system (CNS).

In conclusion, we have discussed here the protective nature of autophagy in light of important human viral infections, and highlighted potential therapeutic strategies that can be pursued through autophagy modulation. Most viral infections result in complex host-pathogen interplay, and therefore routes of intervention require careful consideration in terms of application. For example, certain studies of autophagy in HSV-1 infections have shown viral restriction whilst others have only demonstrated cytoprotective effects despite the presence of viral autophagy inhibitors, which may be partly due to cell type specificity. These studies reveal the intimate interactions of the virus and the host cell which will require further dissection if we wish to target the appropriate molecular pathways for antiviral therapies.

## AUTHOR CONTRIBUTIONS

LA, SM, and VS-S conceptualized and wrote this review. Figure was prepared by LA. All authors approved the final version of this review and agreed to be accountable for the content of the work.

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**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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