

## Addenda

# Topology of Double-Membraned Vesicles and the Opportunity for Non-Lytic Release of Cytoplasm

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### ABSTRACT

Infection of mammalian cells with several positive-strand RNA viruses induces double-membraned vesicles whose cytosolic surfaces serve as platforms for viral RNA replication. Our recent publication (Jackson et al. *PLoS Biol* 2005; 3:861-71) chronicled several similarities between poliovirus-induced membranes and autophagosomes, including induced co-localization of GFP-LC3 and LAMP1. Occasionally, the cytosolic lumen of these structures also contains viral particles; this likely results from wrapping of cytosol, which can contain high viral concentrations late in infection, by newly formed double membranes. Interestingly, RNAi treatment to reduce LC3 or Atg12p concentrations reduced yields of extracellular virus even more than intracellular virus. It is often assumed that exit of non-enveloped viruses such as poliovirus requires cell lysis. However, we hypothesize that autophagosome-like double-membranes, which can become single-membraned upon maturation, provide a long-sought mechanism for the observed non-lytic release of cytoplasmic viruses and possibly other cytoplasmic material resistant to the environment of maturing autophagosomes.

Autophagy is an important component of the innate immune response, limiting infections of mammalian cells with herpes virus<sup>1</sup> and with intracellular bacteria such as mycobacteria,<sup>2</sup> *Streptococcus*<sup>3</sup> and *Shigella*.<sup>4</sup> However, during infection with positive-strand RNA viruses such as poliovirus,<sup>5-7</sup> equine encephalitis virus,<sup>8</sup> murine hepatitis virus<sup>9</sup> and SARS virus,<sup>10</sup> mammalian cells accumulate double-membraned vesicles that are thought to serve as platforms for viral RNA replication. In a recent publication in *PLoS Biology*,<sup>11</sup> we and our co-authors chronicled many similarities between autophagosomes and the vesicles induced by both poliovirus and rhinovirus, including, in addition to their double-membraned structure, induced co-localization of GFP-LC3 and LAMP1, post-fixation staining with monodansylcadaverine<sup>12</sup> and the presence of cytosolic luminal contents. Of potential interest to the autophagy community is the existence of defined viral proteins such as poliovirus proteins 2BC and 3A that can, when expressed together but not separately, induce GFP-LC3 and LAMP1 co-localization and double-membraned vesicles.<sup>7,8,11</sup>

Co-localization of viral RNA replication complexes with components of the autophagosome did not demonstrate what the purpose of the co-localization was: do the host autophagosomes facilitate viral replication or help defend against microbial infection? For murine hepatitis virus, it was shown that elimination of Atg5p expression from murine ES cells caused a dramatic reduction in the production of extracellular, enveloped virions,<sup>13</sup> arguing that Atg5p was required in some step of viral production, maturation or cell exit. We showed that stimulation of the autophagy pathway with tamoxifen or rapamycin increased yields of poliovirus, a non-enveloped virus. Inhibition of the pathway with 3-methyladenine or with RNAi directed against either endogenous ATG12 or LC3 reduced the amount of intracellular virus,<sup>11</sup> consistent with a role for components of the autophagosome in RNA synthesis or packaging. However, these RNAi treatments caused an even greater reduction in the yield of extracellular virus.<sup>11</sup> One possible explanation of the observed preferential decrease in extracellular virions is that the reduction in autophagosome machinery decreased cell lysis early in infection. Alternatively, the reduced abundance of autophagosomal machinery may impair a pathway of non-lytic viral escape. We are intrigued by the latter possibility, because nonlytic delivery of cytosol to the extracellular milieu could be a unique characteristic of multilamellar vesicles. We and others have obtained ultrastructural images consistent with this interpretation, with viral particles,<sup>11,14</sup> and GFP-LC3<sup>11</sup> present in blebs of cytoplasm nearby or connected to the surface of infected cells.

All positive-strand RNA viruses of eukaryotes replicate their genomes on cytoplasmic membranes, yet the morphology and apparent origins of the targeted membranes differ

greatly from virus to virus. In a striking example, Flock House Virus, a nodavirus whose RNA replication cycle can be mimicked in *S. cerevisiae*, normally establishes RNA replication complexes on the outer mitochondrial membranes of infected cells.<sup>15</sup> However, when Flock House Virus RNA replication complexes were retargeted to the endoplasmic reticulum (ER), the efficiency of viral RNA replication actually increased.<sup>16</sup> These kinds of data suggest that different RNA viruses may co-opt different subcellular compartments for some reason other than the exigencies of replicating RNA, such as to alter cellular signal transduction in ways that are useful during infections of natural hosts.

Why do poliovirus and rhinovirus induce the formation of autophagosome-like membranes on which to replicate their RNA? Perhaps the subversion of components of the autophagy pathway by poliovirus and rhinovirus inactivates the innate immune response that autophagy can provide. In addition, the decrease in extracellular virus upon RNAi-mediated reduction of LC3 or Atg12p suggests that the topology of double-membraned vesicles provides an answer to a long-standing puzzle in the biology of non-enveloped viruses. Nominally lytic viruses are often assumed to spread exclusively via cell lysis. However, the possibility of non-lytic release of poliovirus and other picornaviruses has been suggested from numerous reports of persistently infected cell lines that continuously secrete infectious particles.<sup>17,18,14,19</sup> Even more convincingly, when polarized Caco-2 cultures, growing as intact monolayers, were infected with poliovirus, newly synthesized virus was shown to emerge only from the apical surface.<sup>14</sup> Assays showed that the monolayer remained intact, arguing that a non-lytic, and polarized, exit route of unknown origin had been utilized. We speculate that this non-lytic exit route could be provided by the unusual topology of organelles that begin as double-membraned vesicles with cytosolic contents in their lumen (Fig. 1). Early in picornavirus infection, double-membraned structures would entrap cytosol, but this cytosol would be free of virions. However, at later stages of infection, the cytosol trapped by newly generated double-membraned structures would often contain viral particles. Poliovirions and related enteroviruses are relatively resistant to the low pH and active proteolysis that would prevail within the lumen of these vesicles, should they mature. Maturation of autophagosomes results in the complete or partial degradation of the inner membrane: in this case, the fusion of the autophagosome with the plasma membrane would then allow the release of formerly cytosolic material (Fig. 1). If the autophagosome-like structure remains double-membraned, then fusion of the outer membrane with the plasma membrane would release packets of membrane-enclosed cytosol,<sup>11,14</sup> which may or may not remain intact.

The non-lytic release of cytosolic proteins is also a potential explanation for the apparently non-lytic, cell-to-cell spread of aggregated protein conglomerates.<sup>20-22</sup> It has long been known that autophagosomal structures cluster around aggregated proteins in cells<sup>23,24</sup> and recently a positive role for autophagy has been suggested in clearance of aggregates of huntingtin,<sup>25,26</sup> the protein that, when mutant, can cause Huntington's disease. It is possible that the unique topology of double-membraned vesicles is involved in the non-lytic escape of other cytosolic constituents besides viruses. It will be fascinating to determine the potential of the autophagy pathway to provide a non-lytic release mechanism for a variety of cytosolic constituents, although it will be difficult to devise assays as

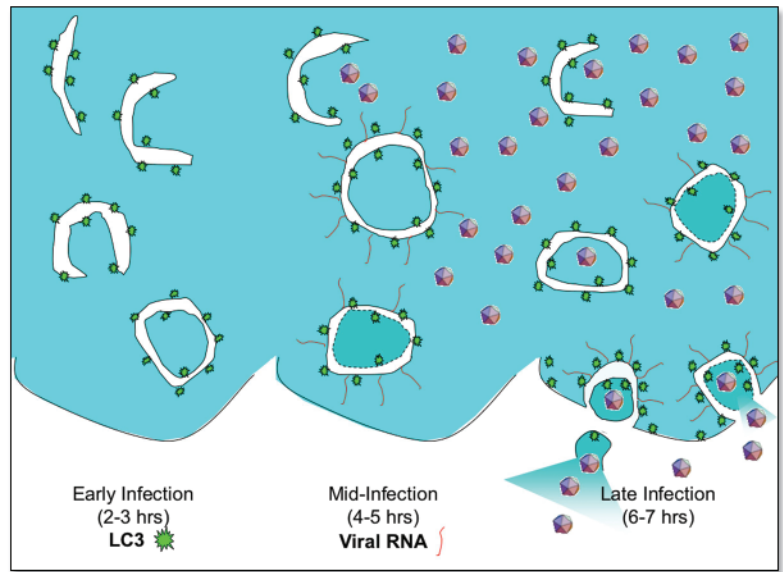


Figure 1. A potential role for double-membraned vesicles in non-lytic release of cytoplasm. Although poliovirus and other non-lytic viruses primarily exit cells via lysis, an alternative, non-lytic route of cell exit is consistent with many experimental observations.<sup>14,17-19</sup> While the surface of these membranes is the known site of viral RNA replication, virions can be observed within the vesicle lumen late in infection.<sup>5,6,11</sup> Extracellular cytoplasmic blebs containing both virus and GFP-LC3 have also been observed.<sup>11,14</sup>

sensitive as a viral plaque assay. This is, of course, one of the reasons that viruses have so often identified novel mechanisms in cell biology, and we speculate that the unique topology of a double-membraned compartment, which can transform into a single-membraned compartment to facilitate the direct release of cytosolic contents to the extracellular milieu, may be one such mechanism.

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