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Addendum

Delivery of cytoplasmic proteins to autophagosomes

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Autophagy represents a signaling-dependent regulated process that allows the degradation of some cellular proteins in autophagosomes, and plays a critical role in the management of cellular homeostasis under various stress conditions. In recent years, selective degradation of cytoplasmic proteins during stress has attracted considerable scientific interest. Here we examined the ability of resveratrol to induce autophagy in a variety of human cancer cell lines. We found that resveratrol-induced autophagy is accompanied by colocalization of proline-, glutamic acid-, and leucine-rich protein-1 (PELP1) with the green fluorescent protein-microtubule-associated protein 1 light chain 3 (GFP-LC3) in autophagosomes. In addition, we found that hepatocyte growth factor-regulated tyrosine kinase substrate (HRS), a previously shown PELP1-interacting protein, is co-recruited to autophagosomes in the presence of resveratrol. Although autophagy has been assumed to be a bulk and non-selective degradation process, in recent years, evidence of selective degradation of cytosolic proteins and organelles by autophagy is mounting. These observations suggest that the interaction of the target protein(s) with the delivery protein or proteins such as HRS facilitates the transport of certain cytoplasmic proteins to autophagosomes for their selective degradation, and thus, could influence the cytoplasmic as well as nuclear functions of nuclear receptor coregulators. Since PELP1 and, perhaps, other nuclear receptor coregulators are widely dysregulated in human cancers, these findings highlight the significance of the autophagic selective degradation of PELP1 following resveratrol (or other phytoestrogens) treatment in developing future strategies to use resveratrol under cancer prevention and therapeutic settings.

The formation of autophagosomes is initiated by the formation of double-membrane-bound structures inside an intact cell, upon encounter of environmental stress such as nutrient starvation, pathogen infection or chemical and radiation treatment. In mammalian cells,

microtubule-associated protein 1 light chain3 (LC3) is recruited to the membranous structure during the mature process, which includes envelopment and sequestration of cytosolic proteins and organelles such as the Golgi apparatus, mitochondria and endoplasmic reticulum (ER). Subsequently, matured autophagosomes fuse with endosomes and lysosomes, converting into autolysosomes to degrade the captured cytoplasmic components. Although such a degradation process, commonly known as macroautophagy, has been assumed to be bulk and non-selective, accumulating evidence suggests a potential selective degradation of organelles by autophagy.^{1,2} For example, peroxisomes that participate in hydrogen peroxide and fatty acid metabolism are proliferated in lower eukaryotes such as *Saccharomyces cerevisiae*, *Hansenula polymorpha* and *Pichia pastoris* and in mammalian cells during growth on fatty acids.³⁻⁵ Alterations in nutrient conditions induce rapid and extensive peroxisome degradation whereas other organelles such as mitochondria, Golgi and cytosolic proteins show much lower levels of degradation, suggesting that the elimination of peroxisomes is selective.³ The ER is selectively sequestered in autophagosomal structures induced when the unfolded protein response is overwhelmed.⁶ Under ER stress conditions, ER clearance is controlled by autophagy, and autophagy-related proteins such as Atg1 and Atg8 are involved in the onset of autophagy and the survival of yeast cells.^{7,8} In addition, autophagy may also be involved in the selective degradation of mitochondria^{9,10} as suggested by the autophagic degradation of the mitochondrial protein Uth1 in yeast.⁹ However, data from electron microscopy studies suggest that mitochondrial sequestration during autophagy is regulated in two temporally distinct manners in both Uth1-dependent and -independent manners.¹⁰ In brief, selective sequestration and degradation of ER, peroxisomes and mitochondrial components plays a pivotal role in the processes important in cellular stress management.^{5,8,10}

In recent years, selective degradation of cytoplasmic proteins has attracted considerable scientific interest.¹¹⁻¹⁴ For example, Onodera and Ohsumi reported¹² the loss of acetaldehyde dehydrogenase in yeast cells and its sequestration into autophagosomes under nitrogen starvation conditions; Yu et al. showed selective degradation of catalase under caspase 8 inhibitor-induced autophagy in a mouse cell line;¹³ and Qing et al have linked the induction of autophagy to the suppression of the 90 kDa heat shock protein (Hsp90) by geldanamycin, which in turn allows the proteasome-independent degradation of I κ B kinase (IKK).¹⁴

Proline-, glutamic acid- and leucine-rich protein 1 (PELP1/ also known as MNAR) is a novel nuclear receptor coregulator that

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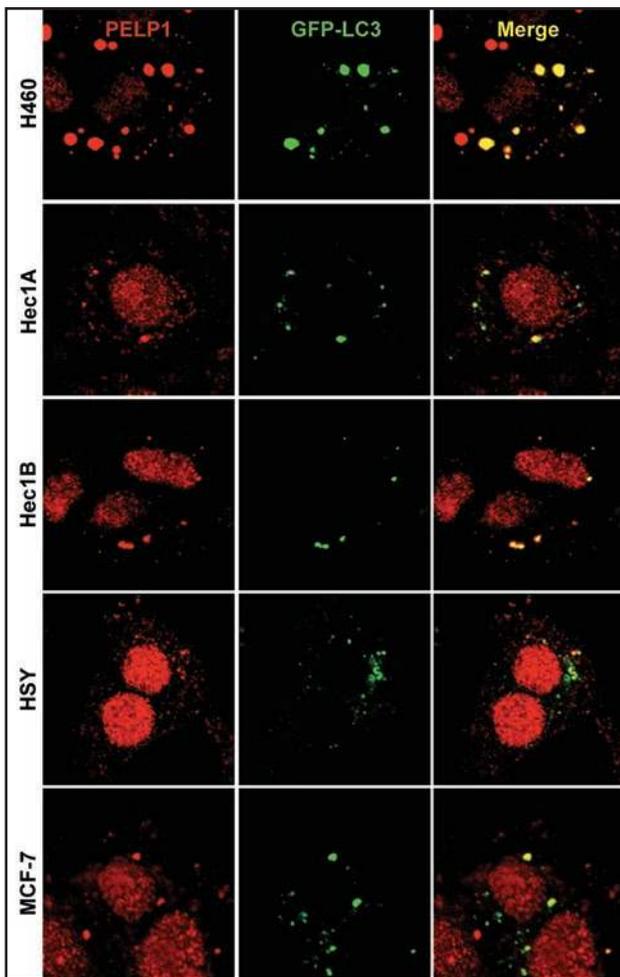


Figure 1. Colocalization of PELP1 and GFP-LC3 in resveratrol-induced autophagosomes in cancer cells. Lung cancer H460 cells, endometrial cancer Hec1A and Hec1B cells, salivary gland carcinoma HSY cells, and breast cancer MCF7 cells were transfected with GFP-LC3, treated with 100 μ M resveratrol for 24 hrs, fixed, and immunolabeled with anti-PELP1 antibody. Colocalization of PELP1 and GFP-LC3 was assayed by confocal microscopy.

modulates genomic and nongenomic signaling of the estrogen receptor. Earlier work from Kumar's lab has reported a mechanistic role of an early endosomal protein, the hepatocyte growth factor-regulated tyrosine kinase substrate (HRS), in the focal sequestration of PELP1 in the cytoplasm.¹⁵ To define the precise nature of such a sequestration focal point, Ohshiro et al.¹⁶ use a green fluorescent protein-LC3 (GFP-LC3) chimera and show that PELP1 and GFP-LC3 are co-recruited and degraded in autophagosomes in a variety of cancer cell types including lung, salivary gland, breast, and endometrial cancer cells upon treatment with the phytoestrogen resveratrol (Fig. 1). Since resveratrol also induces the arrest of the cell cycle at the G₁-phase,¹⁶ we have now shown that resveratrol-induced autophagy of the lung cancer H460 cells was accompanied by inhibition of cyclin D1, a key regulator of the G₁ to S-phase transition (Ohshiro K, Kumar R, unpublished observations). Since PELP1 interacts with the retinoblastoma protein (pRb) and its overexpression leads to accumulation of hyperphosphorylated pRb,¹⁷ we also found a reduction in the status of pRb phosphorylation

upon resveratrol treatment of H460 cells, suggesting the blockade of pRb inactivation by resveratrol and that PELP1 sequestration into autophagosomes further disrupts the interaction between PELP1 and pRb and causes the inhibition of pRb phosphorylation. Collectively, these findings illustrate an example of a modifying role of autophagy upon the status and functions of the cell cycle machinery.

Selective capture of organelles during autophagy could be promoted by projections of the vacuole membrane, followed by contact and fusion, or engulfment by the vacuole of mitochondria¹⁰ and by the enclosure of ER by autophagosomes in an envelope derived from the ER itself.⁶ In addition, the actin cytoskeleton is reported to play an essential role in the elimination of excess peroxisomes by autophagy and the transport of precursor aminopeptidase I in the cytoplasm to vacuole targeting pathway.¹⁸ However, we do not have sufficient understanding of the processes involved in the selective sequestration of non-organelle cytosolic proteins in autophagosomes. Although literature exists about the recruitment to, and roles of autophagy-related proteins at, the autophagosome,¹⁹ it is still unknown exactly how target proteins selectively degraded or are delivered to autophagosomes. In this context, Ohshiro et al.¹⁶ found that resveratrol stimulation of cancer cells triggers a distinct colocalization of HRS with PELP1 in autophagosomes. Since HRS marks the early endosomes in the cytosol and plays a pivotal role in vesicular trafficking and signaling,¹⁵ the above findings implicate a potential role of HRS in transporting PELP1 to the autophagosomes. Interestingly, in *Drosophila*, HRS controls signaling by sorting and degradation of receptors in a ligand-independent manner.²⁰ A recent study has found that an ortholog of HRS, CeVPS-27 is required in the endosomal and autophagic pathways and play a crucial role in endocytic trafficking of low-density lipoprotein receptor-related protein 1 in *Caenorhabditis elegans*.²¹ Furthermore, at the same time of the publication of our current study,¹⁶ HRS was found to play a crucial role in the maturation of autophagosomes in mammalian cells.²² This study stems from the idea that phosphatidylinositol 3-phosphate (PI3P) plays crucial roles in endocytic and autophagic membrane trafficking and also binds to HRS via a FYVE domain.^{23,24} Further, since class I and class III PI3-kinases inhibit and activate the autophagic pathway, respectively, these kinases are expected to be involved at the initial stages of the pathway.²⁵ It is possible that PI3P might also interact with HRS during resveratrol-induced autophagy and might participate in the delivery of its interacting protein, PELP1 to endosomes/autophagosomes. However, this has not yet been formally demonstrated. Emerging results illustrate that in addition to cancer, autophagy is likely to be involved in many human diseases and could be exploited for therapeutic gains.^{26,27} For example, diseases caused by the accumulation or the aggregation of cytosolic proteins are likely to involve autophagy; α -synuclein accumulates and aggregates in neurons of patients with Parkinson's disease,¹¹ whereas mutant alpha-1-antitrypsin accumulates in liver cells of patients with liver inflammation and carcinoma.²⁸ And thus, targeting the activation of the step(s) responsible for selective degradation of target proteins by autophagy may be of therapeutic value. Further, as described above, IKK, which is essential for the activation of the NF κ B pathway, is selectively degraded in an autophagic process induced by geldanamycin.¹⁴ It is possible that the combination of geldanamycin and agents targeting the steps involved in the IKK-selective degradation pathway are likely to be of

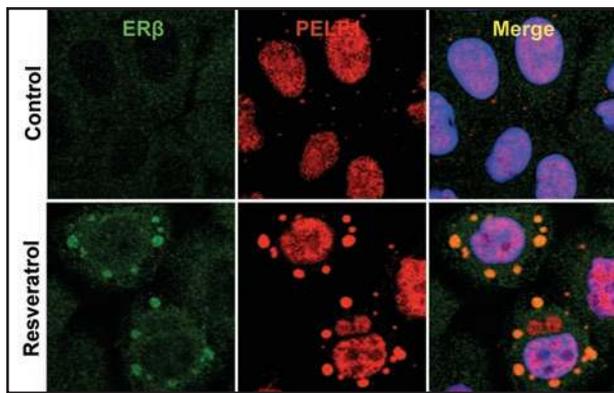


Figure 2. Localization of ER β to vacuole-like compartments containing PELP1. H460 cells were treated with 100 μ M resveratrol for 24 hrs, fixed, and immunolabeled with anti-PELP1 or anti-ER β antibodies. Colocalization of PELP1 and ER β was assayed by confocal microscopy.

therapeutic value in tumors driven by the NF κ B pathway. In the same vein, since PELP1 is overexpressed in a variety of human tumors,²⁹ one potential way to inhibit the growth of PELP1-overexpressing estrogen receptor (ER) α -positive cancer cells is to use resveratrol to trigger autophagy in such cells. Furthermore, we have also found that resveratrol also promotes the accumulation of ER β to vacuole-like structure in H460 cells (Fig. 2). Since PELP1 is known to interact with both ER α and ER β ,³⁰ it is tempting to speculate that the ER β might be also recruited by its interaction with PELP1 and degraded in autophagosomes under autophagic conditions induced by resveratrol. In brief, the reported autophagic selective degradation of PELP1 following treatment with resveratrol may offer new strategies to use resveratrol for cancer prevention in a therapeutic setting.

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