

# Role of tumor and host autophagy in cancer metabolism

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**Macroautophagy (referred to here as autophagy) degrades and recycles cytoplasmic constituents to sustain cellular and mammalian metabolism and survival during starvation. Deregulation of autophagy is involved in numerous diseases, such as cancer. Cancers up-regulate autophagy and depend on it for survival, growth, and malignancy in a tumor cell-autonomous fashion. Recently, it has become apparent that autophagy in host tissues as well as the tumor cells themselves contribute to tumor growth. Understanding how autophagy regulates metabolism and tumor growth has revealed new essential tumor nutrients, where they come from, and how they are supplied and used, which can now be targeted for cancer therapy.**

Autophagy is a catabolic process that captures and degrades damaged proteins and organelles in lysosomes. Autophagy is regulated by >30 autophagy-related (ATG) proteins (Klionsky et al. 2011). During this multistep process, cytosolic components are sequestered in double-membrane vesicles called autophagosomes, which then fuse with lysosomes to form autolysosomes (Mizushima 2007; Klionsky and Codogno 2013; Feng et al. 2014). The contents of autolysosomes are broken down by the degradative enzymes supplied by lysosomes into products such as amino acids, nucleic acids, sugars, and fatty acids, which are recycled into central carbon metabolism (Mizushima and Klionsky 2007; Rabinowitz and White 2010; Guo et al. 2016; Zong et al. 2016). In normal cells and tissues, autophagy is active at a low basal level to sustain cellular homeostasis and protein and organelle quality control through the elimination of damaged organelles and protein aggregates. Autophagy can also be induced by different stresses such as nutrient starvation, hypoxia, oxidative stress, and infection to allow adaptation and survival (Komatsu et al. 2005). Autophagy can be selective or nonselective (Khaminets et al. 2016; Mancias and Kimmelman 2016; Pickles et al. 2018; Wyant et al. 2018). Selective autophagy identifies and targets specific cargo for degradation. Selective autophagic degradation pathways include mitophagy for mitochondria, xenophagy

for bacteria, ribophagy for ribosomes, and ferritinophagy for ferritin, ensuring that the appropriate substrates are degraded and recycled under specific conditions to maintain homeostasis.

Without autophagy, neonates deficient for *Atg5* or *Atg7* survive for only 12 h due to the neonatal starvation period, which can be extended to 24 h with force feeding. *Atg5*- or *Atg7*-deficient neonates also show reduced concentrations of essential amino acids and branched chain amino acids in the circulation. These results indicate the critical importance of autophagy to provide amino acids necessary to survive the neonatal starvation period (Kuma et al. 2004; Komatsu et al. 2005). Autophagy is also required for adult mice to survive fasting (Fig. 1A; Karsli-Uzunbas et al. 2014). Indeed, conditional (acute) systemic (whole-body) *Atg7* deficiency in adult mice leads to gradual depletion of dedicated nutrient stores of lipid in white adipose tissue (WAT), glycogen in liver, and protein in muscle (muscle wasting, also known as cachexia). Fasting conditionally autophagy-deficient adult mice causes rapid depletion of these dedicated nutrient stores and failure to maintain circulating glucose levels, which leads to death from hypoglycemia (Fig. 1A,B). Supplementation of *Atg7* conditionally deficient mice with glucose during fasting rescues muscle wasting and mouse survival (Karsli-Uzunbas et al. 2014). Thus, the loss of autophagy creates a systemic metabolic defect in mammals, possibly increasing dependency on and consumption of circulating nutrients, necessitating excessive catabolism of dedicated nutrient stores (lipid, glycogen, and muscle protein) (Karsli-Uzunbas et al. 2014). The metabolic imbalance created by conditional autophagy deficiency in adult mice may reflect a mechanism of cachexia seen in cancer patients that needs to be explored further.

## Autophagy and cancer

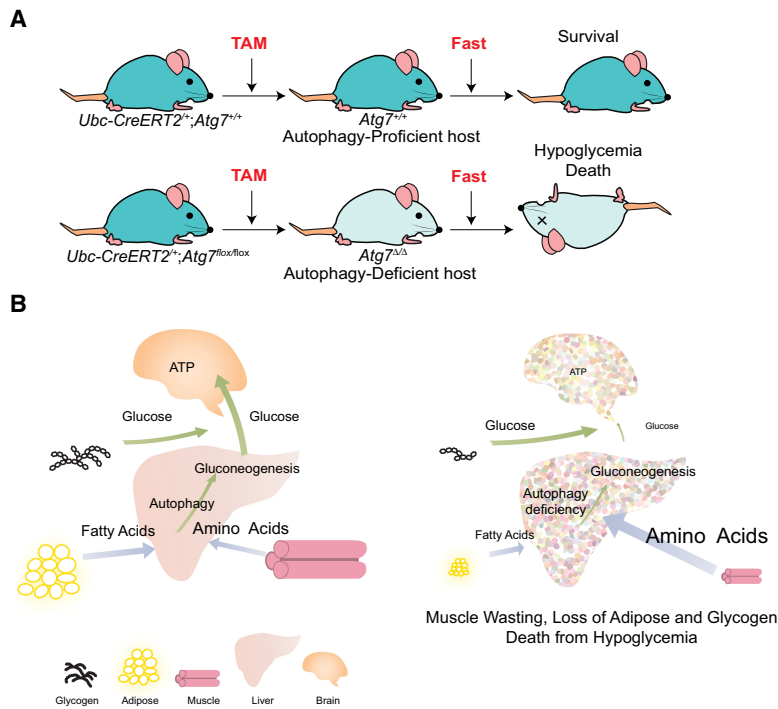
Given the role of autophagy in protein and organelle turnover, intracellular trafficking, and mammalian

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**Figure 1.** Autophagy is required to sustain circulating nutrients during fasting, critical for the survival of adult mice. (A) Mammalian survival during fasting requires autophagy. Treatment with tamoxifen (TAM) leads to conditional whole-body deletion of *Atg7* in adult *Ubc-CreERT2<sup>+/-</sup>;Atg7<sup>lox/lox</sup>* mice. While autophagy-proficient hosts can survive during fasting, autophagy-deficient hosts die from hypoglycemia when fasted (Karsli-Uzunbas et al. 2014). (B) Autophagy is essential to sustain circulating nutrients during fasting. Fasting autophagy-deficient hosts leads to the loss of WAT, glycogen in the liver, and proteins in muscle (cachexia) and the failure to maintain circulating glucose.

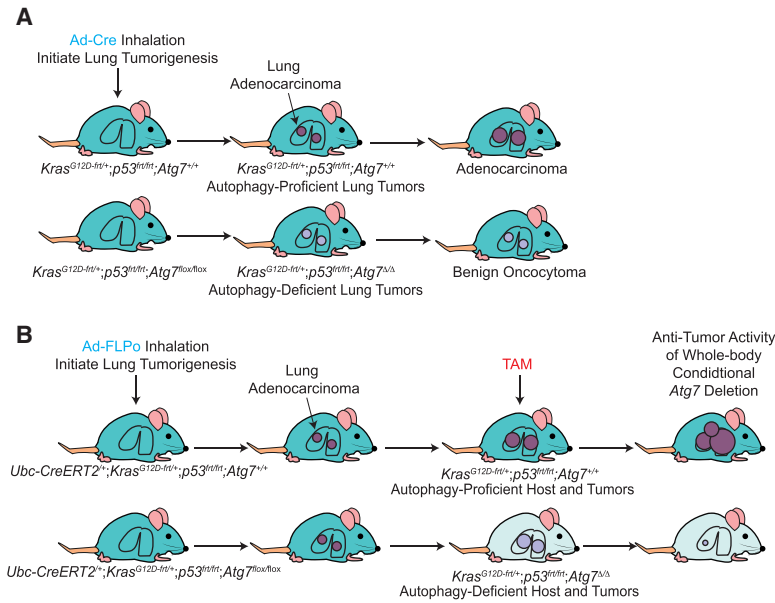
metabolism, it is not surprising that its deregulation has been implicated in numerous diseases such as metabolic diseases, neurodegenerative diseases, infectious diseases, and cancer (Mizushima et al. 2008). For example, conditional whole-body deletion of *Atg7* in adult mice increases susceptibility to *Streptococcus* infection, and most of these mice die at 2–3 mo of age due to neurodegeneration (Karsli-Uzunbas et al. 2014). Mutations of *Atg* genes have also been identified in various human diseases, such as Parkinson diseases and lysosomal storage disorders (Jiang and Mizushima 2014). It is now clear that autophagy is induced in many cancers and promotes their survival. In doing so, cancers have merely usurped the normal and critical survival mechanism of autophagy to enhance their growth and malignancy; hence, targeting autophagy is an important and novel approach to cancer therapy. Understanding the underlying mechanisms by which autophagy functions in cancer can guide its development as an anticancer approach.

### Autophagy in tumor cells promotes cancer

The role of autophagy in cancer has been explored extensively in genetically engineered mouse models (GEMMs). GEMMs for cancer in which autophagy was specifically ablated in tumor cells demonstrated that autophagy loss can promote formation of benign lesions associated with tissue damage and inflammation but that many aggressive cancers require autophagy for growth, survival, and malignancy (White 2012; Kimmelman and White 2017; Onorati et al. 2018). In several of these models, autophagy can act as a tumor-suppressive mechanism during the early stage of tumorigenesis by suppressing reactive oxygen species

(ROS), DNA damage, tissue damage, inflammation, and genome instability, which are known inducers of tumor initiation (Karantza-Wadsworth et al. 2007; Komatsu et al. 2007; Mathew et al. 2007, 2009, 2014; Yang et al. 2011; Deretic et al. 2013; Rosenfeldt et al. 2013; Strohecker et al. 2013). In tissues such as the pancreas and liver, where tumor initiation is caused by chronic tissue damage and inflammation, ablation of *Atg7* or *Atg5* leads to induction of benign pancreatic intraepithelial neoplasia (PANIN) in *Kras<sup>G12D</sup>/+* mice and spontaneous liver adenomas, respectively, that fail to progress to malignancy (Takamura et al. 2011; Rosenfeldt et al. 2013; Yang et al. 2014). As these tumors remain benign, this indicates that even though depletion of autophagy can increase tumor initiation in the pancreases and livers of mice, autophagy is required for tumors to progress to a malignant stage. While these findings from GEMMs are interesting, the general infrequency of mutations in essential autophagy genes in human cancers has indicated that this does not represent a mechanism of cancer causation in humans and that the vast majority of human cancers preserve autophagy function (Laddha et al. 2014; Lebovitz et al. 2015).

Genetic ablation of essential autophagy genes in numerous GEMMs for cancer has revealed an important role for autophagy in promoting tumor growth, survival, and malignancy. In these GEMMs, essential autophagy genes are deleted in tumor cells that arise spontaneously in the context of a normal tumor microenvironment and functional immune system. Deletion of *Atg5* or *Atg7* in *Kras<sup>G12D</sup>*- or *Braf<sup>V600E</sup>*-driven lung cancer (Fig. 2A; Guo et al. 2013; Strohecker et al. 2013; Karsli-Uzunbas et al. 2014), *Braf<sup>V600E</sup>;Pten<sup>-/-</sup>*-driven melanoma (Xie et al. 2015), *Kras<sup>G12D</sup>*-driven pancreatic ductal adenocarcinoma



**Figure 2.** Tumor cell-autonomous autophagy as well as systemic autophagy promote tumor growth. (A) Tumor cell-autonomous autophagy promotes tumor growth. AdenoCre virus inhalation initiates lung tumorigenesis by activation of *Kras*<sup>G12D</sup> and deletion of *Trp53*, and tumor-specific autophagy deficiency is induced by deletion of *Atg7*. Loss of tumor cell-autonomous autophagy inhibits tumor growth and leads to the development of benign oncocytomas (Guo et al. 2013). (B) Pronounced antitumor activity of systemic autophagy ablation in tumor-bearing mice. AdenoFLPo virus inhalation initiates lung tumorigenesis by activation of *Kras*<sup>G12D</sup> and deletion of *Trp53*. When lung cancer is established, conditional whole-body autophagy deficiency is achieved by tamoxifen (TAM) injection and deletion of *Atg7*. Loss of host autophagy and tumor cell-autonomous autophagy inhibits tumor growth to a greater extent than tumor cell-autonomous autophagy alone (Karsli-Uzunbas et al. 2014).

(PDAC) (Rosenfeldt et al. 2013; Yang et al. 2014), or *Pten*<sup>-/-</sup>-driven prostate cancer (Santanam et al. 2016) with or without *Trp53* deletion decreases tumor progression. Similar results were obtained with the deletion of *Becn1* or FAK family-interacting protein of 200 kDa (*Fip200*) in *Palb2*<sup>-/-</sup> or polyoma middle T (*PyMT*) oncogene-driven breast cancer GEMMs (Wei et al. 2011; Huo et al. 2013) or the deletion of *Atg13* or Unc-51-like autophagy-activating kinase (*Ulk1*) in a *Kras*<sup>G12D</sup>-driven glioblastoma model (Gammoh et al. 2016). Deletion of *Atg7* in the intestine epithelium in adenomatous polyposis coli (*APC*)<sup>+/-</sup> mice suppresses tumor initiation by increasing the immune response and perhaps also by increasing metabolic stress (Levy et al. 2015). Finally, autophagy is particularly critical for the survival and growth of *Kras*<sup>G12D</sup>; *Lkb1*<sup>-/-</sup> (liver kinase B1)-driven lung cancer, as these tumors lack LKB1 and the ability to induce AMP kinase and adapt to metabolic stress (Bhatt et al. 2019). Thus, many cancers benefit from and require functional autophagy for their growth and progression.

There is increasing evidence that selective autophagy, such as mitophagy or ferritinophagy, also has a role in cancer (Chang et al. 2017; Um and Yun 2017; Santana-Codina and Mancias 2018). For mitophagy, several studies have suggested a tumor suppressor role for PARKIN (Fujiwara et al. 2008; Poulgiannis et al. 2010; Zhang et al. 2011), whereas BCL-2-interacting protein 3 (BNIP3) seems to display protumorigenic and antitumorigenic roles (Maes et al. 2014; Chourasia and Macleod 2015; Chourasia et al. 2015). The different isoforms of the selective cargo receptor for ferritinophagy, nuclear receptor coactivator 4 (NCOA4;  $\alpha$  and  $\beta$ ), may have a differential role by either inhibiting or promoting cancer cell proliferation (Peng et al. 2008; Wu et al. 2011). The exact role of selective autophagy in cancer still needs further investigation, as its involvement seems to be cell type- and context-dependent.

**Autophagy sustains tumor cell metabolism**

Whereas basal autophagy functions at a low level in normal cells and tissues, numerous cancer cell lines have a high level of basal autophagy, which is necessary to meet elevated metabolic demand and allow cell survival in vitro and tumorigenesis in vivo (Degenhardt et al. 2006; Guo et al. 2011, 2013; Lock et al. 2011; Yang et al. 2011; Viale et al. 2014). Comprehensive metabolic analysis of RAS-driven tumor cells with and without genetic ablation of autophagy revealed that autophagy is required to prevent energy crisis and maintain nucleotide pools during starvation. Autophagy accomplishes this by recycling macromolecules, thereby providing bioenergetic and biosynthetic substrates to the TCA cycle, which maintains energy homeostasis and nucleotide levels (Guo et al. 2011, 2016; Strohecker et al. 2013). In *Kras*<sup>G12D</sup>; *Lkb1*<sup>-/-</sup>-driven lung cancer models, deficiency in *Atg7* reduces the amino acid substrate supply to mitochondria, causing excessive fatty acid oxidation, which depletes lipid stores and promotes energy crisis (Bhatt et al. 2019). Therefore, an important mechanism by which autophagy promotes tumor growth, survival, and malignancy is through its ability to sustain essential metabolic functions of tumor cells. One means by which RAS-driven cancers achieve this is by up-regulating basal autophagy by directly activating the MiT/TFE-regulated transcription program for autophagy and lysosomal biogenesis (Perera et al. 2015).

**Autophagy deficiency selectively kills tumor cells, thereby providing a therapeutic window**

Knowing that many tumors are autophagy-dependent, as autophagy deficiency compromises growth, survival, and malignancy, raises the question of whether tumors are more sensitive to loss of autophagy than normal tissues. Differential sensitivity between tumor and normal

tissues is essential for a therapeutic window in cancer therapy. To test this genetically, conditional whole-body deletion of *Atg7* in adult mice with established KRAS-driven lung cancer was performed to simulate the consequences in cancer patients following autophagy inhibitor treatment. Conditional whole-body deletion of *Atg7* in mice with lung cancer produced dramatic antitumor activity prior to significant damage to normal tissues (Karsli-Uzunbas et al. 2014). Similar findings were obtained with systemic induction of expression of a dominant-negative ATG4b in mice with KRAS-driven PDAC (Yang et al. 2018). The preferential sensitivity of some tumors compared with normal tissues upon systemic loss of autophagy indicated the existence of a therapeutic window for cancer therapy.

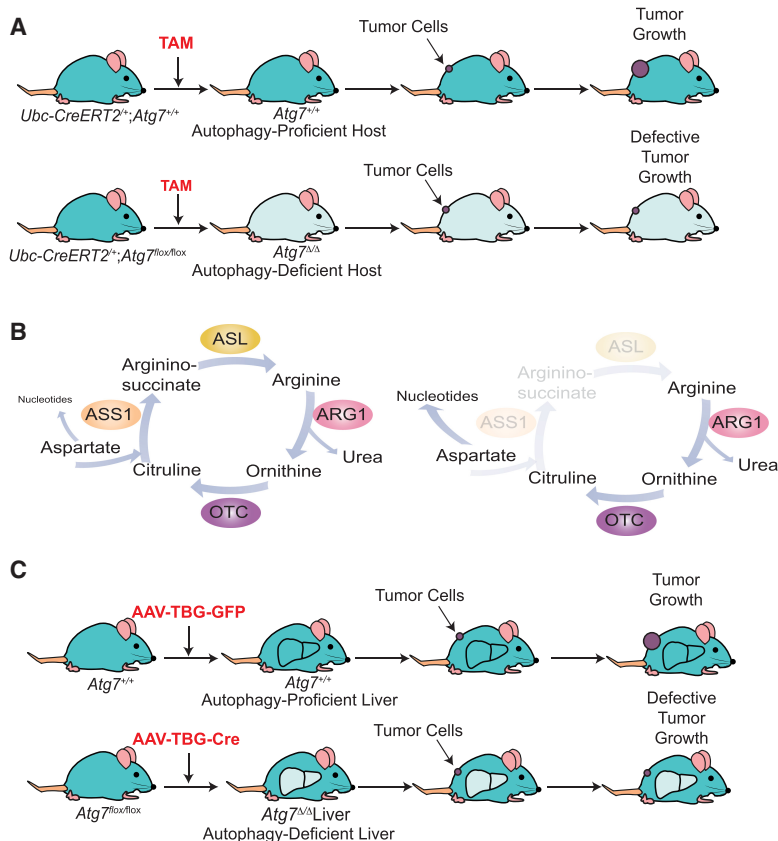
### Host autophagy promotes tumor growth

A remarkable finding from the conditional whole-body deficiency in autophagy in mice with cancer was the substantial tumor regression, which was far greater than that which occurred with tumor-specific autophagy ablation. In one example, conditional deletion of *Atg7* throughout mice with *Kras*<sup>G12D</sup>- and *Trp53*<sup>-/-</sup>-driven lung cancers caused greater tumor regression than tumor-specific *Atg7* deletion, suggesting that host autophagy as well as tumor cell-autonomous autophagy contributed to tumor growth (Fig. 2A,B; Karsli-Uzunbas et al. 2014). In a second example, conditional induction of systemic expression of

a dominant-negative ATG4b in mice with *Kras*<sup>G12D</sup>- and *Trp53*<sup>-/-</sup>-driven PDAC produced similar results (Yang et al. 2018). In a third example, the important role of host autophagy in tumor growth was demonstrated in a *Drosophila melanogaster* malignant tumor model in which deletion of *Atg13* and *Atg14* in different compartments demonstrated that autophagy in the tumor microenvironment and in distant tissues was required for tumor growth (Katheder et al. 2017). To directly test whether host autophagy as well as tumor cell-autonomous autophagy promote tumor growth in mammals, autophagy-competent cancer cell lines were allografted onto autophagy wild-type and autophagy-deficient (conditional whole-body *Atg7* deleted) host mice. Remarkably, deletion of host-specific autophagy impaired the growth of multiple cancer cell lines, including melanoma, carcinogen-induced urothelial carcinoma, and nonsmall cell lung cancer (NSCLC) cell lines (Fig. 3A, Poillet-Perez et al. 2018). Similarly, autophagy in stromal cells in the local tumor microenvironment is necessary for the efficient growth of PDAC (Sousa et al. 2016). Thus, autophagy in both the tumor cells themselves and the host promotes tumorigenesis.

### Host autophagy sustains tumor metabolism

As there is clear evidence that host autophagy promotes tumor growth, the next question is as follows: How does host autophagy promote tumor growth? Tumors



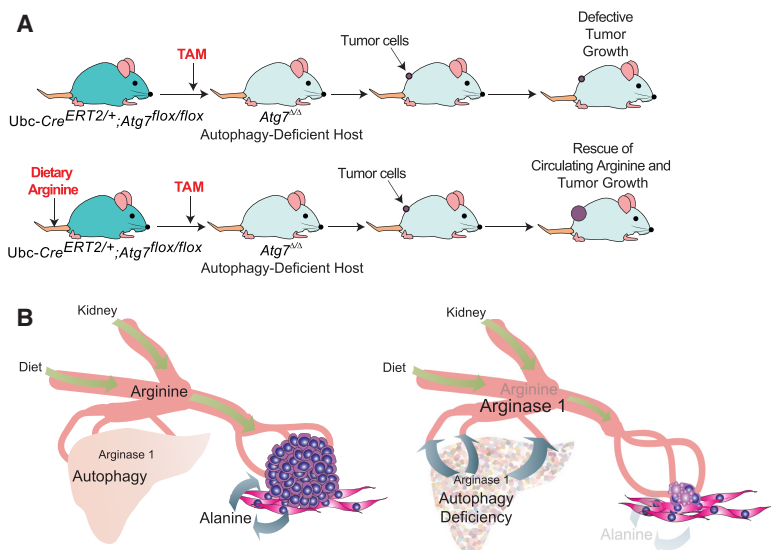
**Figure 3.** Whole-body autophagy and liver autophagy are essential for the growth of arginine-auxotrophic tumors. (A) Host autophagy promotes tumor growth. Treatment with TAM leads to conditional whole-body deletion of *Atg7*. Loss of host autophagy dramatically decreases the growth of autophagy-competent tumor cells, demonstrating the role of nontumor cell-autonomous autophagy in tumor growth (Poillet-Perez et al. 2018). (B) Arginine auxotrophy in cancer. Many cancer cells are auxotrophs for arginine, as they do not express argininosuccinate synthase 1 (ASS1) or argininosuccinate lyase (ASL), two enzymes required for de novo arginine biosynthesis. Tumors downregulate expression of these enzymes in order to use aspartate for nucleotide biosynthesis instead of the urea cycle (Rabinovich et al. 2015). (ARG1) Arginase 1; (OTC) ornithine transcarbamylase. (C) Liver autophagy promotes tumor growth. Tail vein injection of AAV-TBG-cre leads to the specific deletion of *Atg7* in the liver. Loss of autophagy in the liver mimics the effect of host autophagy loss on tumor growth (Poillet-Perez et al. 2018).

reprogram their metabolism to support biosynthetic and energetic pathways necessary for growth, proliferation, and survival. Tumors obtain their entire nutrient supply from the host to fuel these pathways (DeBerardinis and Chandel 2016; Pavlova and Thompson 2016; Vander Heiden and DeBerardinis 2017). As autophagy is important for sustaining host metabolism at both the cellular and mammalian levels, particularly during nutrient limitation (Rabinowitz and White 2010; Goldsmith et al. 2014; Karsli-Uzunbas et al. 2014; Kimmelman and White 2017), the loss of host autophagy may create a systemic metabolic defect and unfavorable environment for tumor growth. Two main sources of tumor nutrients are the host circulation and the local tumor microenvironment; as such, they were the first places to look for a role for host autophagy in promoting cancer metabolism.

Using conditional whole-body deletion of *Atg7* or *Atg5* and autophagy-competent cancer cell lines, it was found that host autophagy maintains tumor growth by sustaining the levels of circulating arginine (Poillet-Perez et al. 2018). These autophagy-deficient mice have altered serum metabolite levels. Among these metabolites, arginine is notably reduced in serum from autophagy-deficient host mice (Karsli-Uzunbas et al. 2014; Poillet-Perez et al. 2018). Arginine is a semiessential amino acid that is derived from three different sources: the diet, de novo synthesis, and protein turnover. Arginine is involved in multiple biological pathways, such as the urea cycle, mechanistic target of rapamycin (mTOR) activation (Chantranupong et al. 2016), and synthesis of nitric oxide, creatine, polyamines, and proteins (Morris 2007). It has long been known that numerous human cancers are arginine auxotrophs due to the silencing of enzymes such as argininosuccinate synthase 1 (ASS1), which is responsible for de novo arginine synthesis (Fig. 3B; Dillon et al. 2004; Delage et al. 2010; Patil et al. 2016). Without expression of the enzymes necessary for de novo arginine biosynthesis, tumor cell growth thereby is completely dependent on a supply of exogenous arginine. Urea cycle dysregulation

(UCD) through loss of ASS1 allows cancer cells to use aspartate for pyrimidine synthesis to support proliferation rather than for arginine synthesis and the urea cycle (Fig. 3B; Rabinovich et al. 2015; Nagamani and Erez 2016; Keshet et al. 2018). In a similar way, the mitochondrial electron transport chain can enable aspartate-derived nucleotide synthesis, which is required for tumor growth (Birsoy et al. 2015; Garcia-Bermudez et al. 2018). Moreover, arginine deficiency in arginine-auxotrophic tumors leads to mitochondrial distress and exhausts aspartate by inducing asparagine synthetase, leading to tumor cell death (Cheng et al. 2018). UCD induces nucleotide imbalance that leads to an increase in transversion mutations, worse prognosis, and better response to immune checkpoint blockade (Lee et al. 2018). Given these well-known alterations in de novo arginine biosynthesis, usage, and dependency in cancer, the decreased circulating arginine caused by autophagy deficiency in mice was likely to have significant deleterious consequences for tumor growth. The next questions are as follows: Is this the case, and why does autophagy deficiency in mice cause depletion of circulating arginine?

Unbiased proteomic profiling of serum revealed that in conditional whole-body autophagy-deficient host mice (either *Atg7* or *Atg5* deleted), low-circulating arginine is associated with increased levels and activity of a major enzyme that degrades arginine: arginase 1 (ARG1) (Poillet-Perez et al. 2018). ARG1 is localized in the liver and is involved in the degradation of arginine to ornithine. ARG1 can be released into the circulation following liver damage (Morris 2012). Indeed, liver-specific deletion as well as whole-body deletion of *Atg7* or *Atg5* cause the release of ARG1 into the circulation and decrease serum arginine and tumor growth (Fig. 3C). These findings indicate that ARG1 is released from hepatocytes following *Atg7* or *Atg5* deletion in the liver. Importantly, dietary arginine supplementation partially restores circulating arginine and tumor growth, confirming the importance of arginine for tumor growth (Fig. 4A). Thus, autophagy in the



**Figure 4.** Host autophagy promotes tumor growth through circulating arginine. (A) Dietary arginine supplementation partially rescues tumor growth on autophagy-deficient host mice. Treatment with TAM leads to conditional whole-body deletion of *Atg7*. Supplementation of the mice with arginine partially rescues tumor growth in autophagy-deficient host mice (Poillet-Perez et al. 2018). (B) Nontumor cell-autonomous autophagy promotes tumor growth by sustaining the supply of amino acids that are essential tumor nutrients. When autophagy is active in the liver, the release of ARG1 from hepatocytes is prevented, thereby maintaining circulating arginine that is important for the growth of arginine-auxotrophic tumors. Loss of autophagy in the liver causes the release of ARG1 from hepatocytes into the circulation, leading to the depletion of circulating arginine and inhibition of the growth of tumors auxotrophic for arginine (Poillet-Perez et al. 2018). Similarly, loss of autophagy in the stroma cells inhibits the secretion of alanine necessary for PDAC growth (Sousa et al. 2016).

liver prevents ARG1 release from hepatocytes and the degradation of circulating arginine that is important for the growth of arginine-auxotrophic tumors, highlighting a new metabolic vulnerability of cancer (Fig. 4B; Poillet-Perez et al. 2018; Venida and Perera 2019). Similarly, autophagy in pancreatic tumor stromal cells facilitates secretion of alanine, which is taken up by PDAC cells and used to support their metabolism and growth (Fig. 4B; Sousa et al. 2016). Thus, autophagy has an important role in controlling essential tumor nutrients by regulating both the local and systemic amino acid supply. Therapeutic inhibition of autophagy in cancer patients should limit the supply of these essential tumor nutrients, thereby impairing tumor growth in addition to inhibiting tumor promotion through tumor cell-autonomous autophagy.

### Conclusions and future directions

Recent studies have characterized the role of autophagy in tumor metabolism and highlighted the importance of both host and tumor autophagy in promoting metabolism and tumorigenesis. However, many additional questions remain. Although host autophagy sustains circulating arginine, allowing arginine-auxotrophic tumor cells to grow, not all arginine-auxotrophic cancer cell lines are defective for growth on an autophagy-deficient host (Poillet-Perez et al. 2018). This suggests the existence of tumor-specific adaptation mechanisms that need to be further studied.

Liver-specific deletion of *Atg7* or *Atg5* leads to decreased circulating arginine, but the tumor growth defect is less than that observed with conditional whole-body deletion of *Atg7* (Poillet-Perez et al. 2018). These data suggest that in liver-specific deleted hosts, autophagy in the microenvironment may locally feed the tumor with arginine, partially compensating for the loss of circulating arginine. Indeed, autophagy in the tumor microenvironment can provide amino acids as well as extracellular matrix molecules and interleukin-6, which promote PDAC growth and tumors in *Drosophila* (Sousa et al. 2016; Endo et al. 2017; Katheder et al. 2017). Note that macrophages in the tumor microenvironment can also influence tumor growth through ARG1-dependent depletion of arginine (Zea et al. 2005; Ellyard et al. 2010). While this was not observed in autophagy-deficient hosts, there are likely multiple roles for arginine in tumorigenesis. Moreover, LC3-associated phagocytosis (LAP) in the myeloid compartment, which uses essential autophagy components such as ATG7, also promotes tumor growth by engulfing dying cells and suppressing inflammatory polarization of the tumor-associated macrophages and an antitumor T-cell response (Cunha et al. 2018). Thus, inhibition of autophagy by targeting ATG7 may not only compromise tumor nutrition and metabolism but also induce an antitumor T-cell response through inhibition of LAP or by other mechanisms. Tumors might also be getting arginine through macropinocytosis, a nonselective form of endocytosis that provides amino acid supply (Recouvreux and Commisso 2017; Finicle et al. 2018). In fact, it has been

shown that RAS-driven PDAC and PTEN-deficient prostate cancer cells use macropinocytosis to overcome amino acid deprivation and support their growth (Commisso et al. 2013; Kamphorst et al. 2015; Davidson et al. 2017; Kim et al. 2018).

The reason why tumors need arginine for their growth is not totally clear. Arginine could promote tumor growth in different ways: by enabling protein synthesis or by sustaining mTOR activity, nitric oxide, or polyamine synthesis, all of which could be involved in promoting tumor growth (Chantranupong et al. 2016; Morris 2016; Saxton et al. 2016). In autophagy-deficient hosts, circulating arginine is degraded by ARG1 to produce ornithine (Fig. 3B; Poillet-Perez et al. 2018). It will be interesting to use in vivo isotope tracing to determine how arginine is used in tumors grown on wild type and autophagy-deficient hosts.

Circulating nutrients are important for tumor growth, and depleting essential tumor nutrients is an established therapeutic approach for cancer. For example, L-asparaginase is a frontline component in the treatment of acute lymphoblastic leukemia (Koprivnikar et al. 2017). As numerous tumors are auxotrophs for arginine, arginine deprivation therapy for cancer patients using enzymes that catabolize arginine, such as pegylated arginine deiminase (ADI-PEG20) or pegylated ARG1 (PEG-ARG1), are in development (Feun et al. 2015; Patil et al. 2016). Arginine deprivation with ADI-PEG20 in vitro leads to metabolism alteration and up-regulation of glutamine anaplerosis and cell death when used with glutaminase inhibitor (Kremer et al. 2017). The use of ADI-PEG20 alone or in combination with temozolomide in a mouse model of glioblastoma multiforme (GBM) leads to a decrease in tumor growth and extends mouse survival without obvious toxicity (Przystal et al. 2018). The impact of ADI-PEG20 on patients with ASS1-deficient tumors needs to be investigated further, and, encouragingly, a clinical trial demonstrated at least 30% stable disease in numerous cancer types, such as hepatocellular carcinoma (HCC) and melanoma (Patil et al. 2016). However, ADI-PEG20 is isolated from *Mycoplasma* and is immunogenic in humans; therefore, the use of other arginine-degrading enzymes, such as human PEG-ARG1, might be a more effective alternative. In vitro, PEG-ARG1 induces nonapoptotic cell death in GBM (Khoury et al. 2015). PEG-ARG1 also suppresses the tumor growth of malignant pleural mesothelioma xenografts and inhibits T-cell leukemia cell proliferation by inducing apoptosis through eIF2 $\alpha$  phosphorylation (Morrow et al. 2013; Lam et al. 2017), suggesting promising activity of PEG-ARG1 in human cancer. There are only a few clinical trials on PEG-ARG1: A stable disease rate of 26.7% was observed in HCC, and one patient with immunotherapy-resistant melanoma showed a sustained complete remission following treatment with PEG-ARG1 (Yau et al. 2013; De Santo et al. 2018). However, ARG1 is also a mediator of immune suppression. Inhibition of ARG1 with a small molecule inhibitor induced a proinflammatory environment and reduced tumor growth in a preclinical model (Steggerda et al. 2017). The effect of PEG-ARG1 needs to be further investigated in order to better understand the protumorigenic or antitumorigenic

role of ARG1. In order to further deprive tumors of arginine, it would be interesting to assess the combination of PEG-ARG1 with autophagy inhibitor. The use of chloroquine (CQ) or LY294002 to inhibit autophagy enhanced the toxicity and apoptosis induced by recombinant human arginase (rhARG) in NSCLC cells in vitro and potentiated the antitumor effect of rhARG in vivo (Shen et al. 2017).

Work on arginine identifies a new metabolic vulnerability of cancer and highlights the importance of better understanding the needs of arginine-auxotrophic tumors in order to best implement this therapy in the right patient population (Savaraj et al. 2010; Shen et al. 2017). In addition to tumors that have silenced enzymes required for de novo arginine biosynthesis, RAS- and BRAF-driven tumors that are highly autophagy-dependent are likely to be responsive to arginine deprivation therapy. Recent insights from understanding the nature of the autophagy addiction of RAS-driven cancers are providing further guidance. For example, coordinate inhibition of CRAF, BRAF, and ATG7 is selectively synthetically lethal in RAS-driven tumor cells in vitro (Lee et al. 2019; White 2019). Moreover, in RAS-driven cancers, inhibition of MAP kinase signaling with MEK or ERK inhibitors further induces autophagy, and their anticancer activity is significantly enhanced by genetic or pharmacologic (CQ) inhibition of autophagy (Bryant et al. 2019; Kinsey et al. 2019). It will be of interest to test whether this autophagy addiction of RAS-driven cancers is all or in part due to the need to maintain circulating arginine essential for tumor growth.

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