

Upregulation of ATG3 contributes to autophagy induced by the detachment of intestinal epithelial cells from the extracellular matrix, but promotes autophagy-independent apoptosis of the attached cells

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Abbreviations: ATG, autophagy-related; BCL2L1, Bcl2-like1; CDK4, cyclin-dependent kinase 4; ECM, extracellular matrix; EGFR, epidermal growth factor receptor; GABARAP, GABA(A) receptor-associated protein; GFP, green fluorescent protein; LC3/ MAP1LC3, microtubule-associated protein 1 light chain 3; PE, phosphatidylethanolamine; qPCR, quantitative polymerase chain reaction; SD, standard deviation; SH3GLB1/Bif1, SH3-domain GRB2-like endophilin B1; *siRNA*, small interfering RNA; UVRAG, UV radiation resistance-associated.

Detachment of nonmalignant intestinal epithelial cells from the extracellular matrix (ECM) triggers their growth arrest and, ultimately, apoptosis. In contrast, colorectal cancer cells can grow without attachment to the ECM. This ability is critical for their malignant potential. We found previously that detachment-induced growth arrest of nonmalignant intestinal epithelial cells is driven by their detachment-triggered autophagy, and that *RAS*, a major oncogene, promotes growth of detached cells by blocking such autophagy. In an effort to identify the mechanisms of detachment-induced autophagy and growth arrest of nonmalignant cells we found here that detachment of these cells causes upregulation of ATG3 and that ATG3 upregulation contributes to autophagy and growth arrest of detached cells. We also observed that when ATG3 expression is artificially increased in the attached cells, ATG3 promotes neither autophagy nor growth arrest but triggers their apoptosis. ATG3 upregulation likely promotes autophagy of the detached but not that of the attached cells because detachment-dependent autophagy requires other detachment-induced events, such as the upregulation of ATG7. We further observed that those few adherent cells that do not die by apoptosis induced by ATG3 become resistant to apoptosis caused by cell detachment, a property that is critical for the ability of normal epithelial cells to become malignant. We conclude that cell-ECM adhesion can switch ATG3 functions: when upregulated in detached cells in the context of other autophagy-promoting events, ATG3 contributes to autophagy. However, when overexpressed in the adherent cells, in the circumstances not favoring autophagy, ATG3 triggers apoptosis.

Introduction

Many types of epithelial cells grow *in vivo* attached to a form of the extracellular matrix (ECM) called the basement membrane. Detachment of such cells from the ECM causes their growth arrest and ultimately, apoptosis.^{1–3} The latter form of cell death is called anoikis.³ It is now believed that these phenomena preclude growth of the indicated cells outside of their original

location. Unlike normal epithelial cells, cells composing carcinomas, cancers of epithelial origin, are well known to be able to survive and grow without being attached to the ECM.^{4–6} Such ability is thought to be critical for carcinoma progression.^{7,8} This notion is based on numerous experimental data. For example, the ability of cancer cells to survive and proliferate without being attached to the ECM as colonies in soft agar has served as a standard for malignant transformation for several decades.^{9,10} Also,

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the ability to grow anchorage-independently without adhesion to the ECM is conferred to cancer cells by activation of major oncoproteins, such as RAS, SRC, EGFR, and CTNNB1/ β -catenin, as well as by loss of tumor suppressor genes, such as PTEN.^{2,11-14} Furthermore, treatments that block the ability of cancer cells to grow without attachment to the ECM, suppress their in vivo tumorigenicity and the capacity for metastasis.^{4,5,15-21} Finally, acquisition by the nonmalignant cells of the ability to grow anchorage-independently results in a spontaneous attainment of in vivo tumorigenicity by these cells.^{22,23} Thus, the ability of tumor cells to grow without adhesion to the ECM represents a potential novel target for cancer treatment. The development of such treatments has been hampered by the fact that the mechanisms by which the nonmalignant cells stop growing and die after detachment from the ECM and those by which cancer cells acquire the ability to grow anchorage-independently are understood only in part.

In an effort to understand these mechanisms we found previously that anoikis of nonmalignant intestinal epithelial cells is driven by detachment-induced downregulation of the anti-apoptotic protein BCL2L1/BclX_L and detachment-induced upregulation of the pro-apoptotic molecule FASLG/Fas ligand as well as that of a cell-death promoting protein kinase CHEK2/Chk2.^{4,24,25} We also established that oncogenic RAS, an oncoprotein that often occurs in colorectal cancer, blocks anoikis of colon cancer cells via multiple mechanisms that involve the downregulation of pro-apoptotic proteins BAK1/Bak and CASP2/caspase-2 and the upregulation of BCL2L1 as well as that of apoptosis inhibitors BIRC3/cIAP2 and XIAP.^{4,5,18,26}

In searching for the mechanisms by which detachment of nonmalignant intestinal epithelial cells blocks their growth we found that growth arrest of detached cells is driven by their detachment-induced autophagy. Autophagy, degradation of cellular macromolecules and organelles, begins when a membrane of unknown origin surrounds a part of the cytoplasm or an organelle.²⁷ The resulting vesicle (autophagosome) fuses with the lysosome to form the autolysosome, and lysosomal enzymes degrade the vesicle content.²⁸ Autophagy, whose mechanisms are understood in part, is regulated by over 20 ATG proteins.²⁹ Autophagosome formation is begun in part by the action of class III phosphatidylinositol 3-kinase and BECN1/Beclin 1 and is then driven by a conjugate between the lipid phosphatidylethanolamine (PE) and proteins of the MAP1LC3/LC3 subfamily as well as that between PE and the GABARAP subfamily of proteins.²⁹ MAP1LC3 lipidation starts when its C-terminus binds the E1 ubiquitin-activating enzyme-like protein ATG7. ATG7-activated MAP1LC3 is then linked to the E2 ubiquitin-conjugating enzyme-like ATG3, and then to PE.²⁹ GABARAP is lipidated via a similar mechanism.^{30,31}

We found that ablation of BECN1 blocks detachment-induced autophagosome formation in the nonmalignant intestinal epithelial cells and detachment induced reduction of the fraction of these cells in the S-phase of the cell cycle without affecting their viability.¹ We further observed that oncogenic RAS triggers degradation of BECN1 protein in malignant intestinal epithelial cells by activating one or more members of the

calpain family of proteases.¹ We found that BECN1 loss results in the suppression of detachment-induced autophagy in the RAS-transformed cells and promotes proliferation of the indicated cells without adhesion to the ECM.¹

Our findings are consistent with observations indicating that autophagy can act as a tumor-suppressing mechanism. For example, loss of BECN1 and other mediators of autophagy, such as ATG5, ATG7, and SH3GLB1/Bif-1, is known to promote various types of cancer in mice.³²⁻³⁵ Furthermore, anti-cancer drugs, such as erlotinib, block EGFR-dependent tumor growth in vivo at least in part, by activating BECN1 and inducing autophagy of tumor cells.³⁶ Overall, the data emerging from the literature indicate that excessive autophagy can suppress tumor growth by causing growth arrest of cancer cells and perhaps via other mechanisms.^{37,38} This is presumably the reason why activation of major oncogenes, such as *RAS*, *EGFR*, and *AKT*, or loss of tumor suppressor genes, such as *TP53*, restrict (but do not completely block) autophagy in cancer cells.^{1,36,39,40} Conversely, given that such cells often exist in a state of metabolic stress (e.g. that caused by lack of properly organized vasculature),⁴¹ some degree of autophagy is likely required by these cells for the generation of essential cellular macromolecules (that are required for cell metabolism) out of the respective monomeric components that become available after the degradation of nonessential molecules via autophagic mechanisms. This is likely the reason why complete suppression of autophagy prevents growth of various types of tumors in vivo.^{42,43}

It has to be noted that many conclusions regarding the role of autophagy in tumor growth were made based on the susceptibility of mice lacking various mediators of autophagy to cancer.^{32,34,35,42,43} Furthermore, levels of some of these mediators are altered in diverse types of human tumors.^{44,45} Based on the changes in these levels, conclusions regarding the role of autophagy in the progression of respective cancers were made.^{44,45} However, some critical mediators of autophagy seem to be able to play autophagy-unrelated roles in cells. For example, ATG12, one of the important stimulators of autophagosome formation, was found to cause apoptosis via physical interactions with components of the cellular apoptotic machinery (such as the BCL2 family members) in a manner independent of its ability to trigger autophagy.⁴⁶ The capacity to kill cells regardless of its involvement in autophagy has also been demonstrated for ATG3, another protein critical for autophagosome formation.⁴⁷ Thus, the contribution of such proteins to cancer could, in principle, be explained by their autophagy-unrelated functions, such as the ability to control apoptosis.

Since the mechanisms by which detachment of intestinal epithelial cells from the ECM triggers autophagy (and such autophagy is critical for the suppression of anchorage-independent growth of cancer cells¹) are not known, we explored these mechanisms in the present study. We found that detachment-induced upregulation of ATG3 contributes to autophagy of detached cells and the reduction of the fraction of these cells in the S-phase of the cell cycle. We established that in addition to ATG3, detachment-induced upregulation of other autophagy mediators, such as ATG7, also contributes to detachment-induced autophagy.

We found that when ATG3 is overexpressed in the attached intestinal epithelial cells by itself, it does not cause autophagy but instead triggers their apoptotic death. Thus, cell-ECM interactions represent a regulator of the biological functions of ATG3: when ATG3 is upregulated in detached cells in the context of other autophagy-promoting events this protein contributes to the induction of autophagy. However, when overexpressed in the adherent cells, in the circumstances that do not favor autophagy induction, ATG3 triggers apoptosis. We also found that those (relatively few) adherent intestinal epithelial cells that adapt to the presence of increased ATG3 levels and survive despite ATG3 upregulation acquire anoikis resistance, a property that is typically displayed by the malignant cells.⁴⁸

Results

Detachment of intestinal epithelial cells from the ECM triggers autophagosome formation

Previously we used electron microscopy (EM) to demonstrate that detachment from the ECM triggers autophagosome formation in the nonmalignant spontaneously immortalized rat intestinal epithelial cells IEC-18.¹ Another approach for detecting this process that we used in the past was assaying the cells for autophagosome formation by counting the percentage of cells with puncta formed by green fluorescent protein (GFP)-tagged MAP1LC3B (GFP-LC3B) that was introduced into these cells by transient transfection.¹ Detecting the change from diffuse cytoplasmic to punctate (characteristic of autophagy) distribution of GFP-LC3B in cells is an often used assay of autophagosome formation.⁴⁹

We have now confirmed these findings by several additional approaches. First we assayed the cells before and after detachment for the number of puncta per cell formed by GFP-LC3B (that was introduced into these cells by transient transfection) which is now thought to represent a more appropriate method of detecting autophagosome formation than determining the percentage of the cells with such puncta in the total cell population.⁴⁹ As shown in **Figure 1A and B**, detachment resulted in a substantial increase in the number of GFP-LC3B puncta per cell. To demonstrate that detachment-induced increase in these puncta reflects increased autophagosome formation, rather than decreased fusion of the autophagosomes with the lysosomes, we treated attached and detached cells with bafilomycin A₁, an inhibitor of vacuolar H⁺-ATPases and a suppressor of autophagosome-lysosome fusion.⁴⁹ We found that bafilomycin A₁ treatment noticeably increased the number of GFP-LC3B puncta per cell in the attached cells (**Fig. 1A and B**). This indicates that IEC-18 cells undergo a significant degree of basal autophagy which can be disrupted by treatment with bafilomycin A₁. However, detached bafilomycin A₁-treated cells displayed a substantially higher number of GFP-LC3B puncta per cell than bafilomycin A₁-treated attached cells (**Fig. 1A, B**). These data are consistent with a scenario, according to which detachment increases the formation of autophagic vacuoles, rather than disrupts their fusion with the lysosomes in the indicated cells.

To verify these results by a complementary method we infected IEC-18 cells with a tandem mCherry-GFP-tagged LC3-encoding Moloney murine leukemia virus. GFP does not generate fluorescence at an acidic pH, and mCherry-GFP-LC3 can only yield green puncta outside of the lysosomes (the lumen of which is acidic).⁴⁹ In contrast, mCherry can give rise to red fluorescence both at the cytosolic and the lysosomal pH, and thus mCherry-GFP-LC3 can promote the formation of the red puncta both inside and outside of the lysosomes.⁴⁹ We found that mCherry-GFP-LC3-producing cells displayed an increased number of the yellow puncta (formed due to simultaneous fluorescence of both mCherry and GFP) after being detached for 7 h, and that the number of such puncta was significantly reduced after the cells were detached for 30 h (**Fig. 1C, D**). Conversely, the number of the red puncta was increased after the cells were detached for 7 h and remained increased after the cells were detached for 30h (**Fig. 1C, E**). These data are consistent with a scenario according to which detachment triggers increased formation of the autophagic vacuoles (signified by the presence of yellow dots) which later fuse with the lysosomes (and this results in the disappearance of the yellow dots due to GFP inactivation at the lysosomal pH). Since mCherry is stable both at the cytosolic and lysosomal pH, the number of autophagic vacuoles generating red fluorescence increased after cell detachment and remained high even when these vacuoles fused with the lysosomes.

We further found that detachment of IEC-18 cells resulted in a significant decrease in the amount of the nonlipidated MAP1LC3B (often called MAP1LC3B-I) in these cells and a strong increase in the amount of the lipidated MAP1LC3B (often referred to as MAP1LC3B-II), a transition that represents one of the hallmarks of autophagosome formation (**Fig. 1F**).⁴⁹ Detachment did not change the amount of the *MAP1LC3B* mRNA in these cells (**Fig. S1A**). To confirm that the increase in the amount of MAP1LC3B-II in detached cells reflects increased MAP1LC3B lipidation, rather than decreased autophagy-dependent degradation of MAP1LC3B-II (as occurs during autophagy)⁴⁹ we measured such lipidation before and after treatment of attached and detached cells with bafilomycin A₁ (**Fig. 1G, H**). Attached cells treated with this drug displayed a strong increase in the amount of MAP1LC3B-II compared to that of MAP1LC3B-I (under these conditions MAP1LC3B-I could only be detected after a long exposure of the respective western blot (**Fig. S1B, C**)). This increase is consistent with the possibility that bafilomycin A₁ treatment disrupted the basal autophagy in the attached cells which prevented MAP1LC3B-II degradation and caused its accumulation in these cells. Importantly, the amount of MAP1LC3B-II in bafilomycin A₁-treated detached cells was even higher than that in the attached cells treated with this drug (**Fig. 1G, H**). Hence, detachment likely promotes increased MAP1LC3B synthesis and lipidation, rather than blocks MAP1LC3B-II autophagy-dependent degradation.

When GFP-LC3B is delivered to the lysosome as a part of the autophagosome, the inner autophagosomal membrane (and the LC3 component of the fusion protein) is degraded, while the relatively more degradation-resistant GFP component remains intact and emerges as free GFP on a western blot. This free GFP

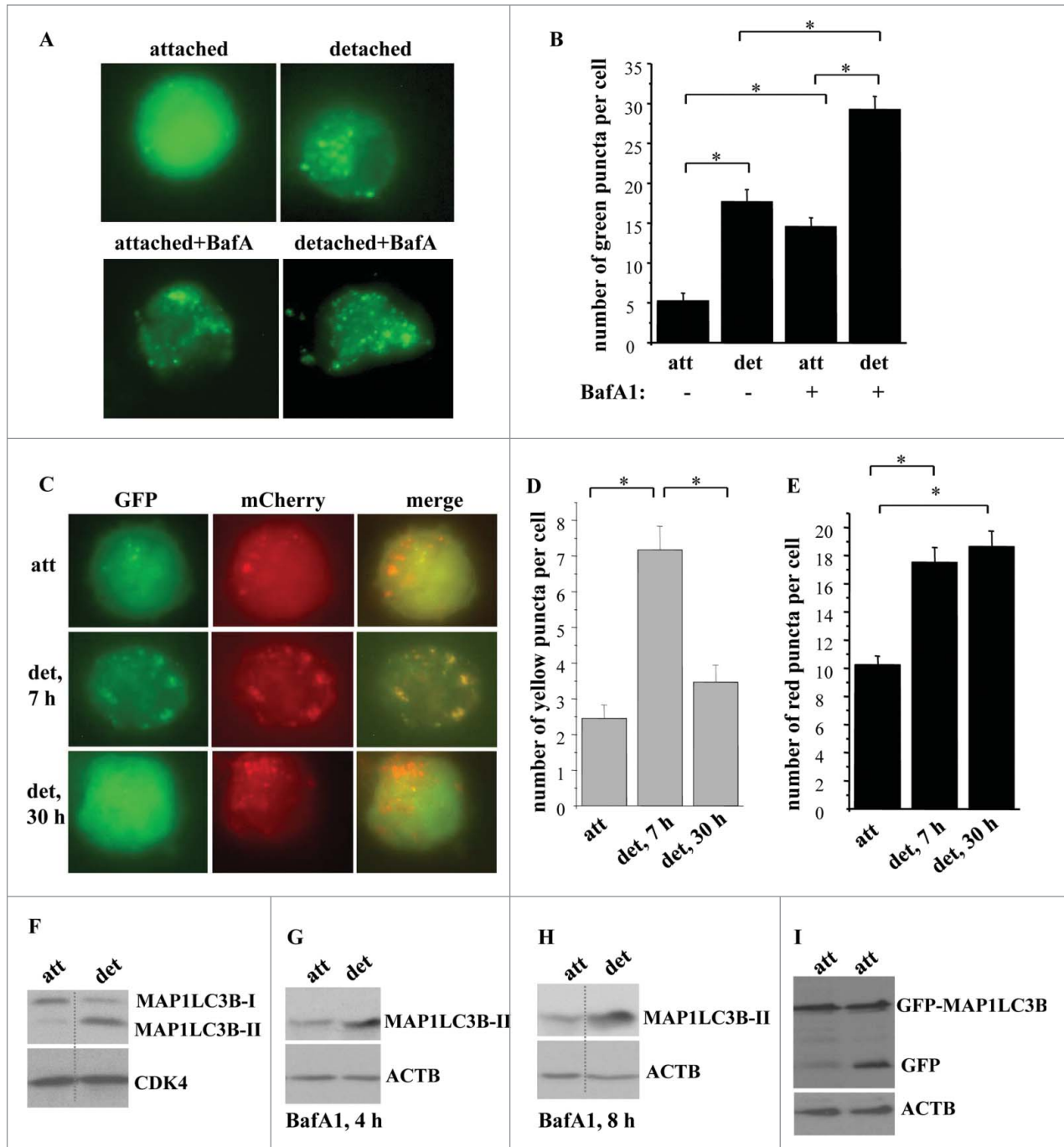


Figure 1. Detachment from the ECM triggers autophagosome formation in intestinal epithelial cells. **(A, B)** Nonmalignant intestinal epithelial cells IEC-18 cells were transfected with the GFP-LC3B expression vector, cultured attached to or detached from the ECM for 24 h in the absence (–) or in the presence (+) of 50 nM baflomycin A₁ (BafA1) and green puncta per cell were counted. **(A)** Representative fluorescence microscopy images of the attached and detached cells. **(B)** Quantification of the number of green puncta per cell. The numbers represent the average of the number of puncta per cell observed for 30–40 cells plus the SE. This experiment was repeated 3 times with similar results. **(C–E)** IEC-18 cells were transduced with the mCherry-GFP-LC3-encoding virus, cultured attached to (att) or detached from (det) the ECM for the indicated times and green, red, or yellow puncta were counted. **(C)** Representative fluorescence microscopy images of the attached and detached cells. **(D, E)** Quantification of the number of yellow **(D)** and red **(E)** puncta per cell performed within the same experiment. The numbers represent the average of the number of puncta per cell observed for 30–40 cells plus the SE. This experiment was repeated twice with similar results. **(F)** IEC-18 cells were cultured attached (att) to or detached from (det) the ECM for 20 h and assayed for MAP1LC3B expression by western blot. Positions of MAP1LC3B-I and MAP1LC3B-II on the gel are indicated. **(G, H)** IEC-18 cells were cultured attached (att) to or detached from the ECM (det) for the indicated times and in the presence of 50 nM baflomycin A₁ (BafA1) and assayed for MAP1LC3B expression by western blot. **(I)** IEC-18 cells were transfected with the GFP-LC3 expression vector, cultured attached to or detached from the ECM for 24 h and assayed by western blot by using an anti-GFP antibody. Positions of GFP-LC3B and free GFP on the gel are indicated. CDK4 was used as a loading control in **(F)** and ACTB in **(G–I)**. *, $p < 0.05$.

emergence represents another way to monitor autophagy.⁴⁹ We observed in this regard that detachment triggered a significant increase of free GFP in GFP-LC3B-transfected IEC-18 cells (Fig. 1I). Collectively, the data shown in Figure 1 indicate that detachment triggers increased autophagy of intestinal epithelial cells.

We also confirmed that similar to what we observed before, detachment of intestinal epithelial cells triggers a significant reduction of the fraction of the cells in the S phase of the cell cycle (Fig. 2A) (and a concomitant increase in the G₁-phase, Fig. S2A) and apoptosis (detected by the ability to bind ANXA5, a characteristic property of apoptotic cells) (Fig. 2B; Fig. S2B).^{1,50}

Detachment of intestinal epithelial cells from the ECM triggers ATG3 and ATG7 upregulation

In an effort to understand the mechanisms of detachment-induced autophagy of intestinal epithelial cells we found that detachment of IEC-18 cells does not alter the expression of autophagy mediators, such as UVRAG, SH3GLB1/Bif-1, ATG16L1, and ATG10, in these cells (Fig. 3A–D). We also found that all of ATG5 is present in a complex that most likely represents ATG5 conjugated with ATG12 in both attached and detached cells (Fig. 3E). The fact that all cellular ATG5 can be conjugated with ATG12 even in the absence of autophagy has been described by several groups.⁴⁹

We further found that detachment of IEC-18 cells resulted in a significant upregulation of autophagy mediators ATG3 and ATG7 (Fig. 4A, B). These events were not unique to IEC-18 cells as we also observed upregulation of both ATG3 and ATG7 in the case of nonmalignant Hkh-2 human intestinal epithelial cells (Fig. 4C, D). These cells were derived from human colon carcinoma cells HCT-116 (that carry one oncogenic *KRAS* allele) by targeted deletion of the activated *KRAS* allele via homologous recombination.⁵¹ Hkh-2 cells are non-tumorigenic in mice and are incapable of growing without adhesion to the ECM.^{23,51} Furthermore, we found previously that detachment from the ECM significantly enhances autophagosome formation in these cells.¹ We also found that detachment of IEC-18 cells causes a significant upregulation of the *ATG3* mRNA (Fig. 4E) but not that of the *ATG7* mRNA (Fig. 4F). Hence, detachment-induced ATG3 upregulation could occur due to increased transcription of the *ATG3* gene or increased *ATG3* mRNA stability. Conversely, it is possible that detachment-dependent upregulation of ATG7 is the consequence of

increased ATG7 protein synthesis or enhanced ATG7 protein stability.

We further observed that detachment-induced ATG3 and ATG7 upregulation occurs at the same time as MAP1LC3B lipidation (Fig. 4G–I). These events were detected by us as early as at 2 h post-detachment. We published previously that detachment of IEC-18 cells increases the number of autophagic vacuoles per cell (detected by EM) from approximately 1 in the attached cells to 5 at 5 h post-detachment and further, to 12 at 17 h post-detachment.¹ Thus, as might have been expected, ATG3 and ATG7 upregulation and MAP1LC3B lipidation is followed by the formation of autophagic vacuoles.

ATG3 upregulation is required for autophagosome formation in detached intestinal epithelial cells

To test whether detachment-induced ATG3 upregulation is required for autophagy of detached intestinal epithelial cells we ablated ATG3 in *detached* IEC-18 cells by transfecting them with 2 separate *ATG3*-specific small interfering (si)RNAs (Fig. 5A). We observed that *siRNA*-transfected detached IEC-18 cells displayed ATG3 levels comparable to those observed by us in the attached IEC-18 cells (Fig. 5A). We then assayed attached and detached cells for autophagosome formation by counting the number of puncta formed per cell by GFP-LC3B (after transient transfection of these cells with the GFP-LC3B expression vector). We found that ATG3 ablation significantly reduced

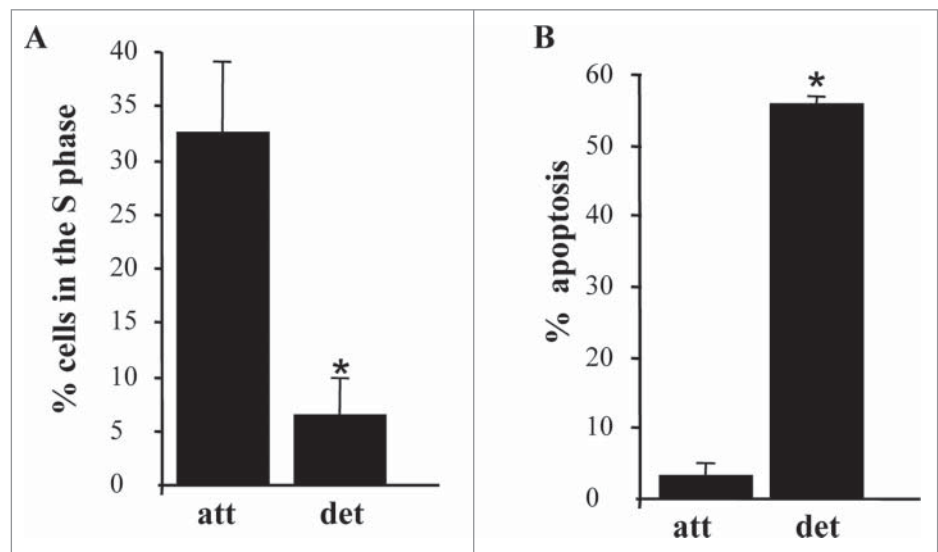


Figure 2. Detachment from the ECM triggers growth arrest and apoptosis of intestinal epithelial cells. (A) IEC-18 cells were cultured attached to or detached from the ECM for 15 h and assayed for the distribution of the cells in phases of the cell cycle by flow cytometry. Percentage of the cells in the S phase of the cell cycle is shown. The numbers represent the average of 2 independent experiments plus the SD. (B) The cells were cultured as in (A), stained with propidium iodide (PI) and ANXA5 and assayed for ANXA5 and PtdIns binding by flow cytometry. Percent apoptosis was calculated as the sum of the percentages of ANXA5-positive-PI-negative cells (undergoing early stages of apoptosis) and ANXA5-positive-PtdIns-positive cells (undergoing late stages of apoptosis). The numbers represent the average of 2 independent experiments plus the SD. * Indicates that the p-value was less than 0.05.

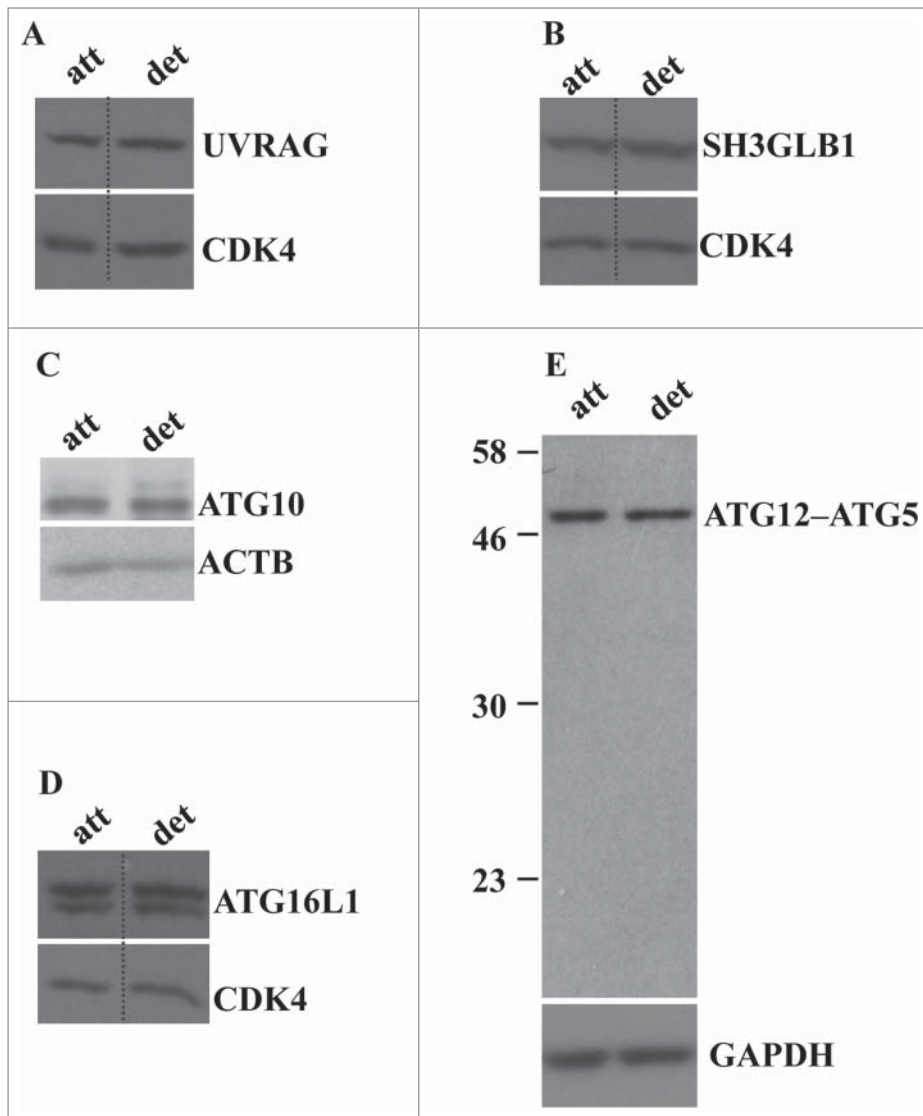


Figure 3. Detachment from the ECM does not alter the levels of UVRAG, SH3GLB1, ATG16L1, ATG10, and ATG12-ATG5 conjugate in intestinal epithelial cells. (A–D) IEC-18 cells were cultured attached to (att) or detached (det) from the ECM for 20 h and assayed for the expression of the indicated proteins by western blot. CDK4 was used as a loading control in (A, B, D), ACTB in (C) and GAPDH (glyceraldehyde 3-phosphate dehydrogenase) in (E).

autophagosome formation in the detached cells (Fig. 5B). We also observed that ATG3 loss noticeably reduced the levels of MAP1LC3B-II (Fig. 5C) in detached IEC-18 cells. We further established that ATG3 ablation blocked lipidation of GABARAP (Fig. 5E).⁵² Covalent GABARAP binding to PE represents another marker of autophagy.⁵² We found that attached IEC-18 cells carried almost exclusively unlipidated GABARAP (often referred to as GABARAP-I) whereas detached cells displayed a noticeable increase in the presence of the lipidated form (GABARAP-II) (Fig. 5D).⁵³ We also observed that detachment did not alter the level of *GABARAP* mRNA (Fig. S3A) in IEC-18 cells. We established that ATG3 loss significantly reduced the amount of GABARAP-II in the detached cells (Fig. 5E).

Collectively, our data indicate that the reversal of detachment-induced ATG3 upregulation blocks detachment-induced autophagy of intestinal epithelial cells.

We further found that enforced ATG3 downregulation significantly increased the fraction of detached cells in the S-phase of the cell cycle (Fig. 5F; Fig. S3B). Given that further to detachment and growth arrest, IEC-18 cells undergo apoptosis, we measured such apoptosis of these cells before and after ATG3 ablation.^{2,4} We found that ATG3 loss did not reduce apoptosis of the detached cells (Fig. 5G; Fig. S3C).

Since at least some autophagy mediators are known to play autophagy-unrelated roles in cells, we decided to confirm the role of autophagy in detachment-induced reduction in the fraction of intestinal epithelial cells in the S phase of the cell cycle by ablating a different autophagy mediator in these cells.^{46,47} Since, according to our data (Fig. 5E), GABARAP likely mediates the effect of ATG3 on autophagy, we tested the effect of GABARAP ablation on the cell cycle progression of detached IEC-18 cells. Similar to what was observed in case of ATG3 loss, we found that GABARAP ablation by *siRNAs* (Fig. 6A) triggered a significant increase in the fraction of detached IEC-18 cells in the S phase of the cell cycle (Fig. 6B; Fig. S4). We conclude that detachment-induced ATG3 upregulation and subsequent autophagy is necessary for detachment-induced reduction in the fraction of intestinal epithelial cells in the S phase of the cell cycle.

ATG3 upregulation triggers apoptosis, rather than autophagosome formation and growth arrest, in the attached intestinal epithelial cells

We further tested whether ATG3 upregulation is sufficient for the induction of autophagy and subsequent growth arrest of attached intestinal epithelial cells. To achieve ATG3 upregulation in the attached IEC-18 cells we infected them with an ATG3-encoding Moloney murine leukemia virus carrying a puromycin resistance gene. We found that about 90% of control cells that were not transduced with the virus died within the first 24 h of puromycin treatment (not shown). We also observed that after the infection of IEC-18 cells with the control virus the ratio of the number of colonies formed by these cells in

monolayer culture after puromycin treatment to that formed by the cells infected with the same amount of the control virus but not treated with puromycin was approximately 0.7 (not shown). Thus, the efficiency of viral transduction under these conditions was approximately 70%. We further established that when infected with the *ATG3*-encoding virus and treated with puromycin for 24 h, the adherent IEC-18 cells produced *ATG3* at levels that were significantly higher than those in the cells infected with the same number of the control viral particles and comparable to those observed in the detached uninfected IEC-18 cells (Fig. 7A). Thus, we have found conditions under which *ATG3* could be produced in the attached cells at the level similar to that induced by cell detachment.

We subsequently assayed the control and *ATG3* expressing cells (generated after infection of IEC-18 cells with respective viruses and a 24 h-long selection with puromycin) for autophagy by detecting the conversion of MAP1LC3B from the nonlipidated form to the PE-conjugated form. Whereas, as expected, detachment of IEC-18 cells resulted in a quantitative conversion of MAP1LC3B-I to MAP1LC3B-II (Fig 7B, left), we observed no change in MAP1LC3B lipidation in the adherent IEC-18 cells in 20h after the introduction of *ATG3* in the cells (Fig. 5B, right), nor in 12 h or 48 h after generating *ATG3*-overproducing cells (not shown). Likewise, similar to what is shown above (Fig. 1B), we found that detachment of IEC-18 cells resulted in a substantial increase in the number of GFP-LC3B puncta per cell, whereas the introduction of *ATG3* in these cells has no impact on GFP-LC3B distribution in these cells when they were attached to the ECM (Fig. 7C). Furthermore, unlike what was observed after cell detachment (Fig. 2A), over-expression of *ATG3* in the adherent cells did not change the fraction of these cells in the S-phase of the cell cycle (Fig. 7D). Thus, it appears that *ATG3* upregulation is not sufficient for the

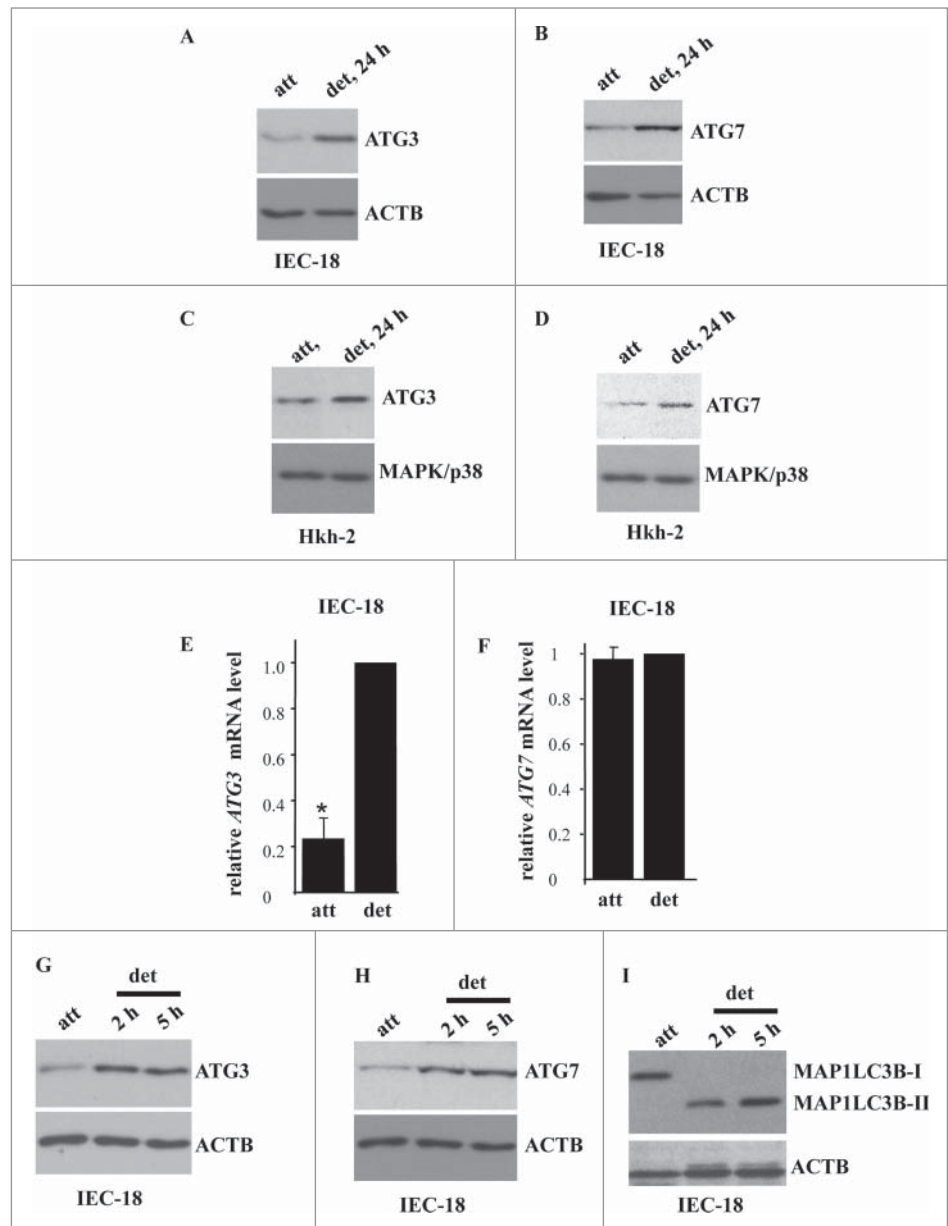


Figure 4. Detachment from the ECM triggers *ATG3* and *ATG7* upregulation in intestinal epithelial cells. (A, B) IEC-18 cells were cultured attached (att) to or detached from (det) the ECM for the indicated times and assayed for *ATG3* (A) or *ATG7* (B) protein expression by western blot. (C, D) Human nonmalignant intestinal epithelial Hkh-2 cells were cultured attached (att) to or detached (det) from the ECM for the indicated times and assayed for *ATG3* (C) or *ATG7* (D) protein expression by western blot. (E, F) IEC-18 cells were cultured attached to or detached from the ECM for 24 h and assayed for *ATG3* (E) or *ATG7* (F) mRNA levels by qPCR. The observed *ATG3* and *ATG7* mRNA levels were normalized by the levels of *RNA18S/18S* rRNA, which were also determined by qPCR. The resulting levels of *ATG3* (E) or *ATG7* (F) mRNA in the detached cells were arbitrarily designated as 1.0. The data represent the average of 4 independent experiments plus the SD in (E) and 2 independent experiments in (F). * indicates that the p-value was less than 0.05. (G-I) IEC-18 cells were cultured attached to or detached from the ECM for the indicated times and assayed for *ATG3* (G), *ATG7* (H) or MAP1LC3B (I) protein expression by western blot. Positions of MAP1LC3B-I and MAP1LC3B-II on the gel are indicated. ACTB was used as a loading control in (A, B, G-I) and MAPK/p38 in (C, D).

induction of autophagy and growth arrest in attached intestinal epithelial cells.

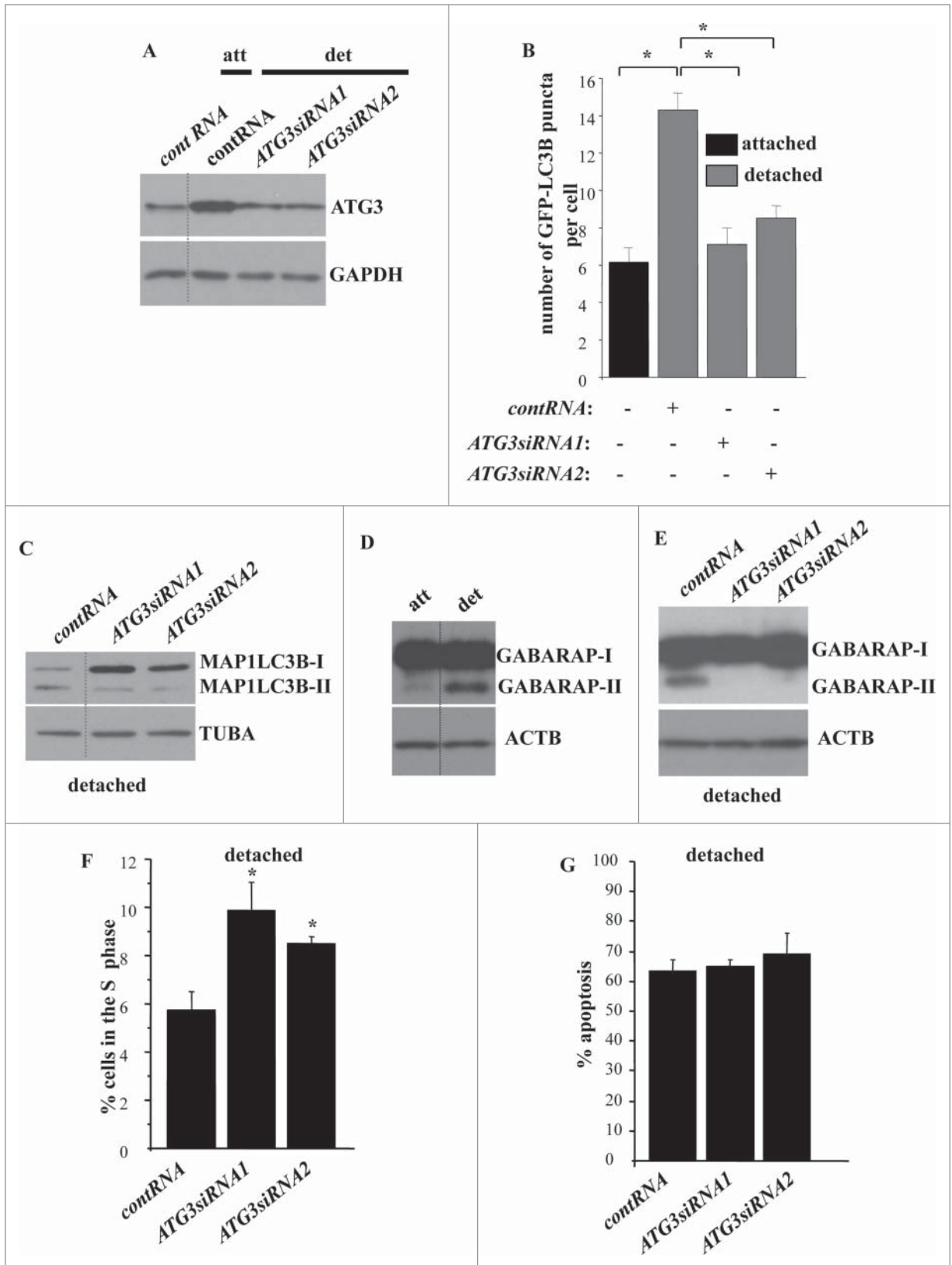


Figure 5. For figure legend, see page 1238.

We further found that when the cells producing ectopic ATG3 were generated (in the process of a 24 h-long selection with puromycin) and placed in monolayer culture for an additional 20 h, a significant fraction of these cells displayed a typical feature of apoptosis, such as the ability to bind ANXA5 (Fig. 7E). When cultured in the presence of exogenous ATG3 for an extended period of time (7–10 d) a significant fraction of these cells (60–70%) lost viability as was evident from their inability to form colonies (Fig. 7F). Thus, upregulation of ATG3 in the adherent intestinal epithelial cells is insufficient for the induction of autophagy in these cells and triggers their apoptotic death.

Conjugation of ATG3 to ATG12 is not required for the ability of ATG3 to kill attached intestinal epithelial cells

Others found that ATG3 can cause apoptosis via covalent binding to ATG12 via lysine 243 (of ATG3).⁴⁷ It was proposed that the ATG12–ATG3 conjugate can increase the susceptibility of cells to various apoptotic stimuli by altering mitochondrial homeostasis. This apoptosis is thought to be mediated by mechanisms that are unrelated to autophagy.⁴⁷ It was demonstrated in this regard that an ATG3 variant in which lysine 243 is mutated to arginine (ATG3KR) is capable of causing autophagy but does not promote apoptosis.⁴⁷ We thus used this mutant to test the role of ATG12–ATG3 conjugation in ATG3-dependent death of adherent intestinal epithelial cells. We found that the ATG3KR mutant when expressed in IEC-18 cells (as described above for the wild-type ATG3) triggered loss of the adherent cells with the same efficiency as the wild-type ATG3 (Fig. 8).

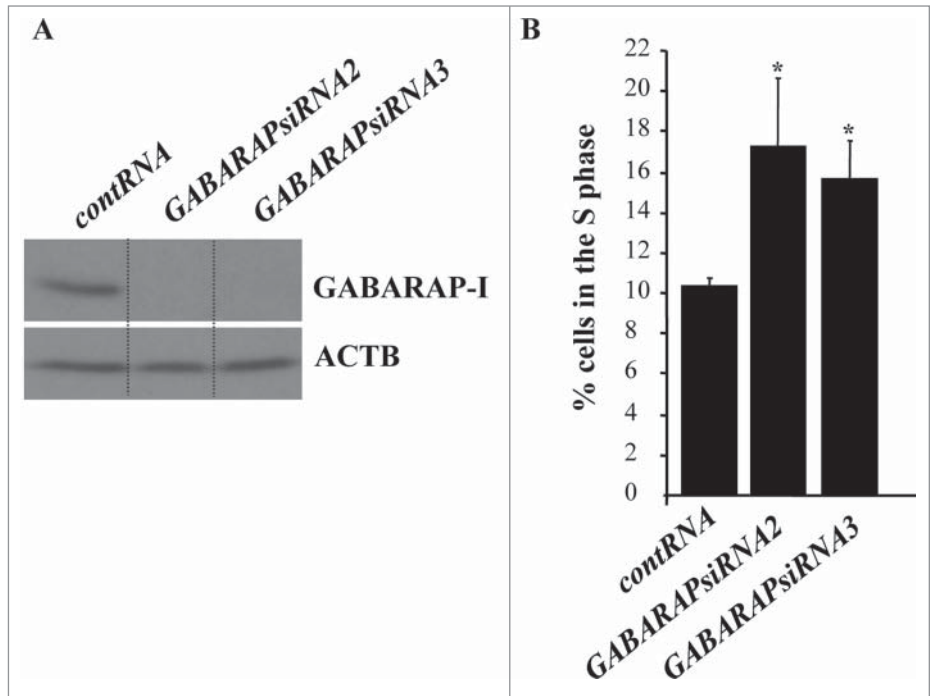


Figure 6. GABARAP ablation increases the fraction of detached intestinal epithelial cells in the S-phase of the cell cycle. (A) IEC-18 cells were transfected with 100 nM control RNA (contRNA) or GABARAP-specific small interfering RNA (GABARAPsiRNA)2 or 3 and assayed for GABARAP expression by western blot. ACTB was used as a loading control. (B) IEC-18 cells were treated as in (A), detached from the ECM for 14h and assayed for the distribution of the cells in phases of the cell cycle by flow cytometry. Percentage of the cells in the S phase of the cell cycle is shown. The numbers represent the average of 3 independent experiments plus the SD. * indicates that the p-value was less than 0.05.

Thus, ATG12 conjugation to ATG3 is unlikely required for the ability of ATG3 to kill attached intestinal epithelial cells.

Detachment-induced upregulation of ATG7 is required for detachment-induced autophagosome formation in intestinal epithelial cells

Since ATG3 upregulation cannot by itself induce autophagosome formation in intestinal epithelial cells we speculated that, in addition to such upregulation, other autophagy-promoting events occur in the detached cells. In addition to ATG3,

Figure 5 (See previous page). Enforced ATG3 downregulation blocks autophagy in detached intestinal epithelial cells and increases their fraction in the S phase of the cell cycle. (A) IEC-18 cells were transfected with 100 nM control RNA (contRNA) or ATG3-specific small interfering RNA (ATG3siRNA) 1 or 2, cultured attached to (att) or detached from (det) the ECM for 2 h and assayed for ATG3 expression by western blot. (B) IEC-18 cells were transiently transfected with a GFP-LC3B expression vector, then transfected as in (A) and cultured attached to or detached from the ECM for 14 h and assayed for the presence of GFP-LC3B puncta as in Figure 1. The results represent the average of the number of GFP-LC3B puncta per cell observed for 30–50 cells plus the SE. This experiment was repeated twice with similar results. (C) IEC-18 cells were transfected as in (A), cultured as in (B) and assayed for MAP1LC3B expression by western blot. Positions of MAP1LC3B-I and MAP1LC3B-II on the gel are indicated. (D) IEC-18 cells were cultured attached to or detached from the ECM for 20 h and assayed for GABARAP expression by western blot. Positions of GABARAP-I and GABARAP-II on the gel are indicated. (E) IEC-18 cells were transfected as in (A), cultured attached to or detached from the ECM for 16h and assayed as in (D). GAPDH was used as a loading control in (A), TUBA/α-tubulin in (C) and ACTB in (D, E). (F) IEC-18 cells were transfected as in (A), cultured detached from the ECM for 14 h and assayed for the distribution of the cells in phases of the cell cycle by flow cytometry. Percentage of the cells in the S phase of the cell cycle is shown. The numbers represent the average of 4 independent experiments plus the SD. * indicates that p-value was less than 0.05. (G) Cells were treated as in (C) and assayed for apoptosis as in Figure 2B. The numbers represent the average of 2 independent experiments plus the SD.

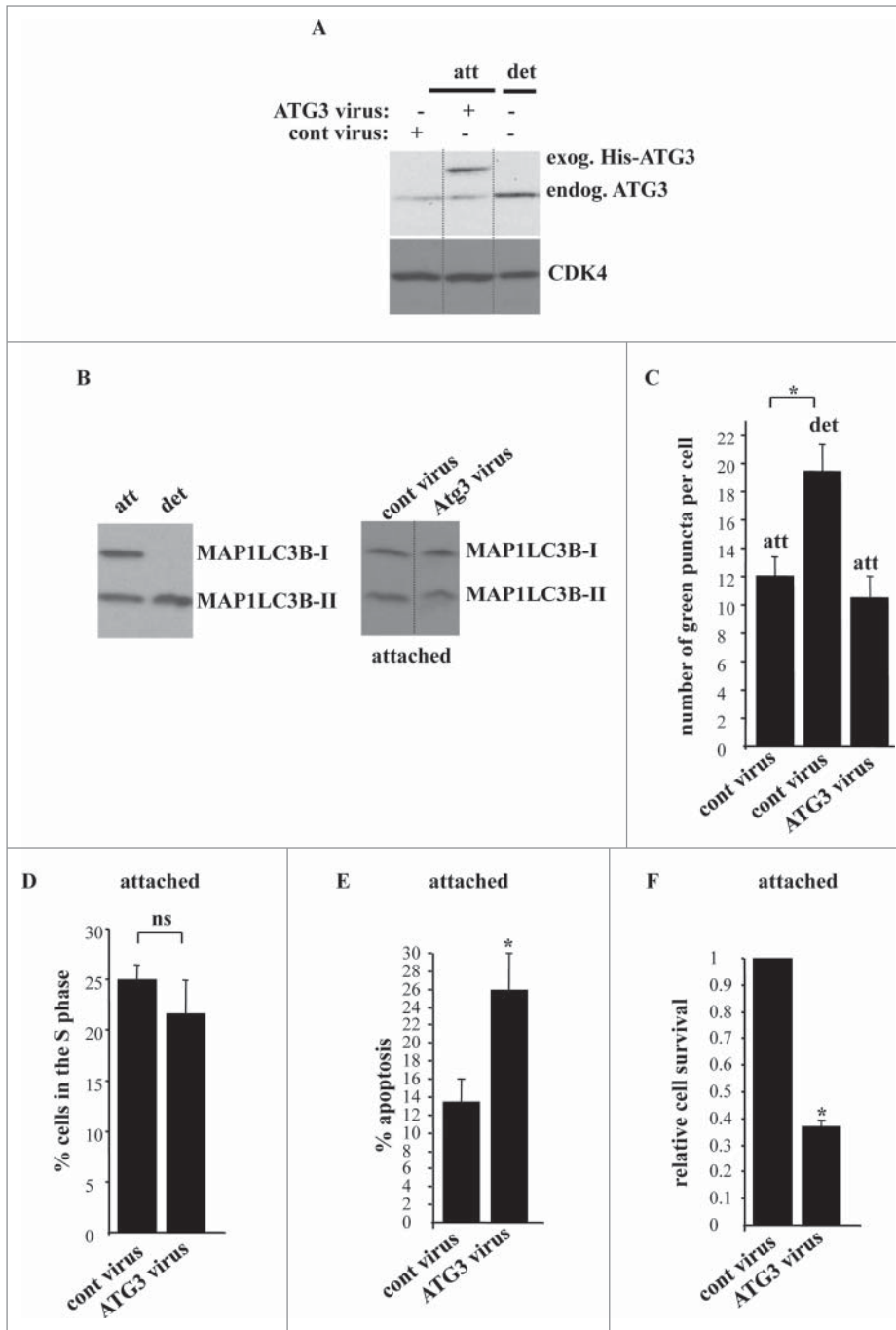


Figure 7. ATG3 upregulation in attached intestinal epithelial cells does not trigger their autophagy or cell cycle arrest but promotes apoptosis. **(A)** IEC-18 cells were infected with 4×10^4 viral particles per cell of the control retrovirus (cont virus) or ATG3-encoding retrovirus (ATG3 virus), treated with $2 \mu\text{g/ml}$ puromycin for 24 h and assayed for ATG3 expression by western blot along with IEC-18 cells that were detached for 20 h. The positions of endogenous ATG3 (endog. ATG3) and exogenous His-tagged ATG3 (exog. His-ATG3) on the gel are indicated. CDK4 was used as a loading control. **(B, left)** IEC-18 cells were cultured attached (att) to or detached (det) from the ECM for 20 h and assayed for MAP1LC3B expression by western blot. Positions of MAP1LC3B-I and MAP1LC3B-II on the gel are indicated. **(B, right)** IEC-18 cells infected with the indicated viruses as in **(A)** were cultured attached to the ECM for an additional 20 h and analyzed as in **(B, left)**. **(C)** IEC-18 cells infected with the indicated viruses as in **(A)** were transiently transfected with a GFP-LC3B expression vector and plated attached to or detached from the ECM for 20 h and analyzed for the presence of GFP-LC3B puncta as in **Figure 1**. The results represent the average of the number of GFP-LC3B puncta per cell observed for 30–45 cells plus the SE. **(D)** IEC-18 cells infected with the indicated viruses as in **(A)** were cultured attached to the ECM for an additional 24 h and assayed for the distribution of the cells in phases of the cell cycle by flow cytometry. Percentage of the cells in the S phase of the cell cycle is shown. The numbers represent the average of 3 independent experiments plus the SD. **(E)** IEC-18 cells infected with the indicated viruses as in **(A)** were cultured attached to the ECM for an additional 24 h and assayed for ANXA5 binding. The numbers represent the average of 2 independent experiments plus the SD. **(F)** IEC-18 cells infected with the indicated viruses as in **(A)** were cultured attached to the ECM for 7–10 d. Cell colonies were then stained and counted. The numbers represent the average of the triplicates plus the SD. This experiment was repeated twice with similar results. Relative cell survival was calculated as the number of colonies formed by the ATG3-expressing cells relative to that formed by the control cells which was arbitrarily designated as 1.0. * indicates that the *p*-value was less than 0.05.

detachment of intestinal epithelial cells triggered upregulation of ATG7 (Fig. 4B, D, H). We reasoned that such upregulation (possibly in concert with other yet unknown detachment-dependent events) could also contribute to detachment-induced autophagy. We therefore tested the role of ATG7 in autophagosome formation in detached cells.

We found that ablation of ATG7 by 2 different *siRNAs* (Fig. 9A, left, center), significantly reduced ATG7 levels in detached IEC-18 cells. ATG7 levels in detached cells transfected with one of these *siRNAs* were similar to those in the attached cells whereas transfection with the second *siRNA*

resulted in somewhat lower ATG7 levels in detached cells compared to the attached ones (Fig. 9A, right). However, transfection with each *siRNA* resulted in a similar (substantial) degree of reduction in the number of GFP-LC3B puncta per cell (the GFP-LC3B vector was introduced in cells via transient transfection) in detached IEC-18 cells (Fig. 9B). These data indicate that once detached cells lose the ability to upregulate ATG7 above a certain level (e.g., that observed in the attached cells), autophagosome formation cannot be induced in the detached cells. We did not observe consistent inhibition of MAP1LC3B lipidation in the detached cells

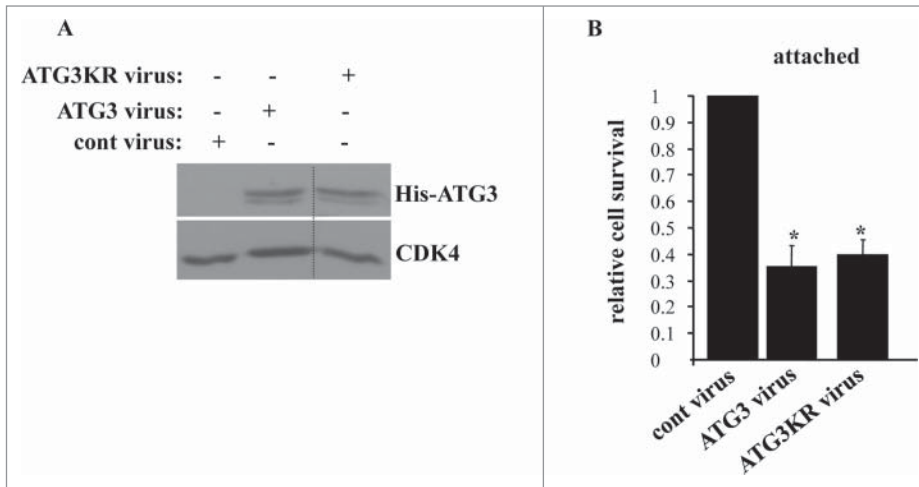
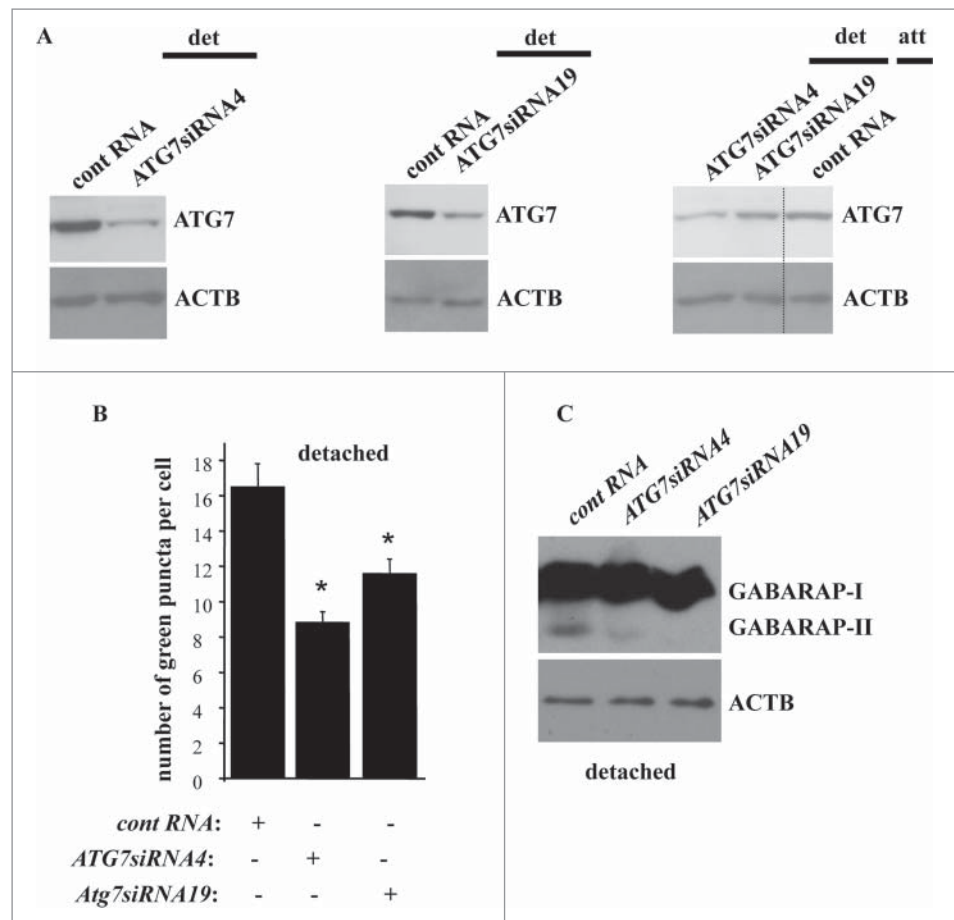


Figure 8. Lysine 243 of ATG3 is not required for the ability of ATG3 to kill attached intestinal epithelial cells. **(A)** IEC-18 cells were infected (+) or not infected (–) with 4×10^4 viral particles per cell of the control retrovirus (cont virus) or ATG3-encoding retrovirus (ATG3 virus), or the virus encoding ATG3 in which K243 was mutated to arginine (ATG3KR virus), treated with 2 $\mu\text{g}/\text{ml}$ puromycin for 24 h and assayed for ATG3 expression by western blot. CDK4 was used as a loading control. **(B)** IEC-18 cells treated as in **(A)** were assayed for survival as in **Figure 5F**. The numbers represent the average of triplicates plus the SD. This experiment was repeated twice with similar results. * indicates that the p-value was less than 0.05.

after enforced ATG7 downregulation, possibly due to the fact that the amount of the remaining ATG7 was sufficient to promote such lipidation. However, the amount of ATG7 remaining in the cells transfected with ATG7-specific siRNAs was clearly not sufficient for the lipidation of GABARAP since we found that ATG7 loss caused a significant inhibition of GABARAP lipidation in the detached cells (**Fig. 9C**).⁵² Hence, upregulation of ATG7 contributes to detachment-induced autophagy of intestinal epithelial cells. These data are consistent with a notion that detachment-induced upregulation of ATG3 stimulates autophagy of detached cells together with other detachment-induced autophagy-promoting events, such as, for example, detachment-induced upregulation of ATG7.

Figure 9. Detachment-induced upregulation of ATG7 contributes to autophagosome formation in detached intestinal epithelial cells. **(A)** IEC-18 cells were transfected with 100 nM control RNA (cont RNA) or ATG7-specific small interfering RNA (ATG7siRNA)4 (left) or 19 (center) and cultured detached from the ECM for 20 h (left, center). IEC-18 cells transfected as above were cultured attached to the ECM for 20 h in case of the control RNA or detached from the ECM for 20 h in case of ATG7siRNAs 4 and 19 and assayed for ATG7 expression by western blot. **(B)** IEC-18 cells were transfected (+) or not transfected (–) as in **(A)**, then transiently transfected with a GFP-LC3B expression vector, detached from the ECM for 12 h and assayed for the presence of GFP-LC3B puncta as in **Figure 1**. The results represent the average of the number of GFP-LC3B puncta per cell observed for 60–120 cells plus the SE. This experiment was repeated twice with similar results. * indicates that the p-value was less than 0.05. **(C)** IEC-18 cells were transfected as in **(A)**, cultured for 16 h detached from the ECM and assayed for GABARAP expression by western blot. Positions of GABARAP-I and GABARAP-II on the gel are indicated. ACTB was used as a loading control in **(A and C)**.



Adherent intestinal epithelial cells that survive after ATG3 upregulation display anoikis resistance

In the course of our studies we noticed that even though the majority of adherent intestinal epithelial cells die in response to the transduction with ATG3-encoding retrovirus, some of these cells survive (Fig. 7F). Furthermore, when expanded in the presence of puromycin (resistance to which is encoded by the respective viral vector) these cells displayed substantial levels of constitutively produced exogenous ATG3 (Fig. 10A). Thus, it appears that a fraction of IEC-18 cells are capable of resisting ATG3-induced apoptosis, and the total cell population is replaced with these apoptosis-resistant cells after ATG3 upregulation. Interestingly, when detached from the ECM, the cells overproducing ATG3 in a stable manner turned out to be significantly more viable than the control cells (Fig. 10B). Hence, these cells are resistant to detachment-induced apoptosis (a phenomenon called anoikis), a property that is thought to be essential for the malignant potential of cancer cells.^{3-5,7,8,17,18,23,55} We further found that the cells stably producing ectopic ATG3 did not display increased autophagosome formation compared to the control cells neither when attached to the ECM nor when detached for 12 h (Fig. 10C) or 17 h (not shown). Thus, their anoikis resistance does not seem to be related to the ability of ATG3 to promote autophagy.

In summary, we have identified in this study a novel mechanism of autophagosome formation in intestinal epithelial cells induced by their detachment from the ECM. We found before that this mechanism contributes to the blockade of anchorage-independent proliferation of nonmalignant intestinal epithelial cells. We also found in the past that *RAS* oncogene promotes growth of detached cell by blocking their autophagy (and the ability to grow without adhesion to the ECM is thought to be critical for the malignant potential of tumor cells).^{1,7,8} We have found now that detachment-induced autophagosome formation in the nonmalignant intestinal epithelial cells is mediated by more than one signal, including those triggered by detachment-induced upregulation of ATG3 and ATG7. We also observed

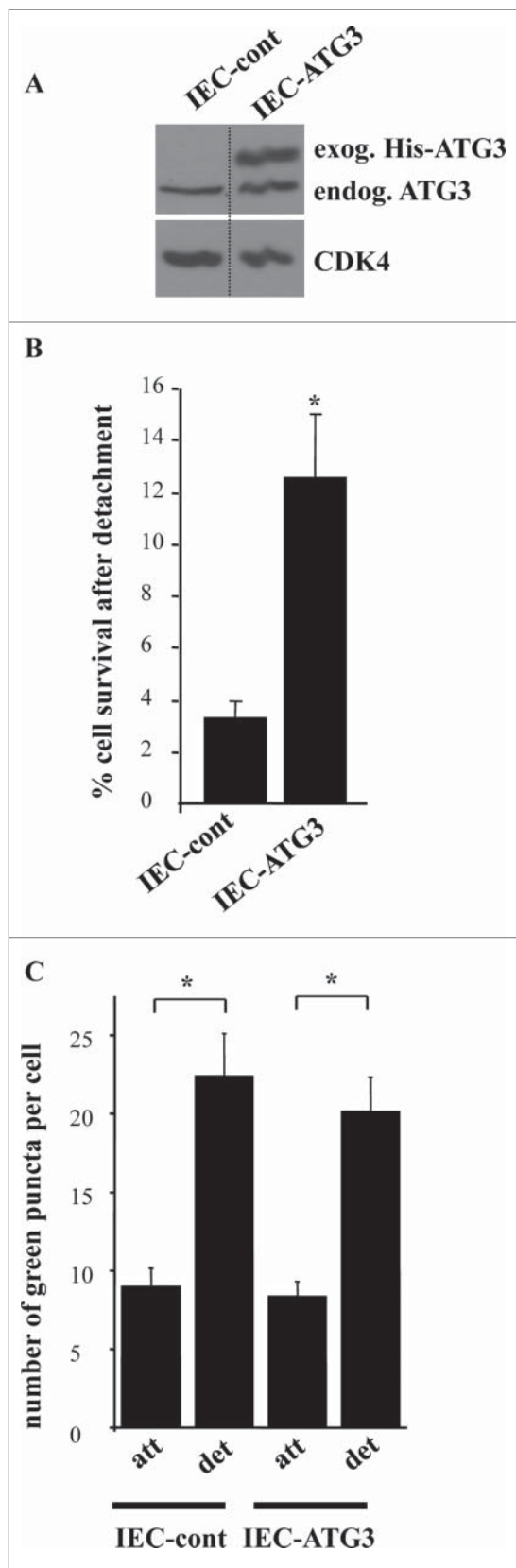


Figure 10. Intestinal epithelial cells that are resistant to apoptosis induced by ATG3 are also resistant to apoptosis triggered by their detachment from the ECM. **(A)** IEC-18 cells were infected with 4×10^4 viral particles per cell of the control retrovirus or ATG3-encoding retrovirus and cultured in the presence of 2 $\mu\text{g/ml}$ puromycin until the respective control cell line (IEC-cont) and ATG3-overexpressing cell line (IEC-ATG3) were established. Cells were then assayed for ATG3 expression by western blot as in **Figure 7A**. **(B)** The indicated cell lines were plated in monolayer culture either immediately or after being cultured without adhesion to the ECM for 16 h. Cell colonies were allowed to form for 7–10 d. The colonies were then stained and counted. “% cell survival after detachment” was calculated as the percentage of the colonies formed by the cells re-plated after detachment relative to that formed by the attached cells. The results represent the average of triplicates plus the SD. This experiment was repeated twice with similar results. * indicates that the p-value was less than 0.05. **(C)** The indicated cell lines were transiently transfected with a GFP-LC3B expression vector, cultured attached to (att) or detached from (det) the ECM for 12 h and assayed for the presence of GFP-LC3B puncta as in **Figure 1**. The results represent the average of the number of GFP-LC3B puncta per cell observed for 24–40 cells plus the SE.

that cell-ECM interactions can determine the biological functions of ATG3: when ATG3 is upregulated in detached cells in the context of other autophagy-promoting events (e.g. detachment-induced ATG7 upregulation) ATG3 contributes to the induction of autophagy. However, when overexpressed in the adherent cells, in circumstances that do not favor autophagy induction, ATG3 triggers apoptosis. Those cells that can adapt to the presence of ATG3 and survive become anoikis-resistant, i.e. they display features of malignant cells.

Discussion

In this study we have identified novel mechanisms of autophagy of intestinal epithelial cells induced by their detachment from the ECM. The importance of identification of these mechanisms stems from our earlier findings indicating that detachment-induced autophagy triggers growth arrest of such cells.¹ The ability of cancer cells to grow without adhesion to the ECM is thought to represent a critical prerequisite for tumor progression.^{4,5,10-14} We found that oncogenic RRAS promotes proliferation of intestinal epithelial cells in the absence of adhesion to the ECM by blocking their autophagy.¹ The mechanisms by which RAS exerts this effect were examined by us in the past. We found that RAS blocks detachment-induced autophagy by triggering calpain-dependent degradation of BECN1.¹ In agreement with our findings, others observed that inhibition of detachment-dependent autophagy enhances proliferation of breast epithelial cells carrying an oncogenic mutant of the PI3-kinase when these cells are placed in a 3-dimensional culture.⁵⁶ So far, the mechanisms by which detachment of the nonmalignant intestinal epithelial cells from the ECM promotes autophagy have remained unknown. We have shown now that such autophagy is triggered by detachment-induced upregulation of ATG3 and ATG7.

Our results indicate that detachment-induced upregulation of ATG3 stimulates autophagy of detached cells together with other detachment-induced autophagy-promoting events, such as, for example detachment-induced upregulation of ATG7. When upregulated in the adherent cells by itself, ATG3 does not trigger autophagy but induces apoptosis instead. These data are consistent with findings made by others that ATG3 and other mediators of autophagy, such as ATG12, can trigger apoptosis via mechanisms unrelated to their ability to promote autophagosome formation.^{46,47} Others proposed that in some cases the ability of ATG3 to trigger apoptosis requires the ability of this protein to conjugate with ATG12.⁴⁷ According to our data, ATG3-induced apoptotic death of attached intestinal epithelial cells occurs regardless of whether or not ATG3 can conjugate with ATG12. Understanding the mechanisms of ATG3-induced apoptosis represents the subject of our ongoing research.

Whether or not detachment-induced upregulation of ATG3 and ATG7 are the only events that contribute to detachment-induced autophagy of intestinal epithelial cells is presently not known. It was demonstrated that detachment of other types of epithelial cells (e.g., those of the breast) blocks several major

metabolic pathways in such cells.⁵⁷ Such metabolic stress could contribute to the induction of autophagy. In addition, it cannot be excluded that other yet unidentified autophagy-promoting changes in the expression of the regulators of autophagy are triggered by detachment of intestinal epithelial cells. Such events, together with detachment-dependent upregulation of ATG3 and ATG7 could in principle contribute to autophagy of detached intestinal epithelial cells. Hence, which combination of detachment-induced autophagy-promoting signals would be sufficient for the induction of autophagy in the adherent cells remains to be investigated. Studies aimed at addressing these questions are currently underway in our lab.

One potentially important physiological implication of our studies is that inhibition of autophagy due to inactivation or loss of ATG3 partners involved in this process can create a situation when ATG3 is no longer needed for autophagosome formation and instead of participating in this process triggers apoptosis. It is now thought in this regard that when autophagy is increased beyond a certain level it blocks tumor growth, and this is presumably the reason why oncogenes, such as *RAS*, *ERBB2*, *AKT*, and others have the potential to reduce autophagy in cancer cells.^{1,36,39} However, some level of autophagy in malignant cells is thought to be required for tumor growth as complete inhibition of autophagy achieved by, for example deletion of genes coding for autophagy mediators in mice (e.g. that of the *Atg7* gene) prevents tumor growth.⁴² One possible interpretation of these data is that cancer cells often experience disruption of metabolism due to their inadequate supply with blood and require basal levels of autophagy for the degradation of the nonessential cellular macromolecules and utilization of the resulting monomeric components of these molecules for the restoration of defective metabolic pathways.⁴² However, it is conceivable that loss of ATG7 or other autophagy-related proteins creates a situation when ATG3 which normally acts as an ATG7 partner in the induction of autophagy can no longer be involved in facilitating autophagy (since autophagosome formation is blocked due to the absence of ATG7) and promotes apoptosis of tumor cells instead.²⁹ Thus, lack of basal autophagy and ATG3-induced apoptosis could in principle cooperate with each other in suppressing tumor growth in mice lacking ATG7 or other autophagy-promoting proteins.

We have demonstrated here that those adherent intestinal epithelial cells that do not succumb to ATG3-induced apoptosis become anoikis-resistant. What are the possible mechanisms of such resistance? We found previously that anoikis of intestinal epithelial cells is driven by multiple mechanisms mediated by detachment-induced changes in the expression of various regulators of apoptosis.^{4,18,24,25} Importantly for this study, we and others observed previously that a small fraction of nonmalignant intestinal epithelial cells, such as IEC-18 and Hkh-2 used in this study, remains viable after being cultured without adhesion to the ECM and if expanded displays pronounced and well reproducible resistance to anoikis^{22,23} (the nature of this cellular heterogeneity is presently not known). These data indicate that intestinal epithelial cells contain a cell population in which at least some of the

anoikis-promoting mechanisms are blocked. It is possible that apoptosis induced by ATG3 is driven by mechanisms that overlap with those triggered by cell detachment. Thus, conceivably, unlike the majority of the cells which succumb to ATG3-induced apoptosis after ATG3 upregulation, the cells composing the anoikis-resistant fraction and hence, lacking the indicated apoptosis-promoting mechanisms, do not die in response to ATG3 upregulation.

What is the possible physiological relevance of the ability of the cells to survive and become anoikis-resistance in response to the upregulation of ATG3? It was found in this regard that ATG3 is upregulated, at least in some types of cells, under ischemia, a condition that frequently occurs in tumors due to the irregularities of the tumor vasculature.⁵⁸⁻⁶⁰ Ischemia induces apoptosis, and it is possible that such apoptosis is mediated at least in part by ATG3 upregulation.⁶⁰ Our data indicate that cells that remain viable despite ATG3 upregulation become anoikis-resistant. On the other hand, the fact that cancer cells can acquire resistance to ischemia-induced apoptosis is well documented.⁶¹ Thus, it is conceivable that cancer cells that do not die by ischemia-induced apoptosis and are hence selected for resistance to ATG3-induced apoptosis are at the same time selected for increased anoikis resistance. Anoikis resistance of cancer cells is thought to substantially contribute to their tumorigenicity.^{4,5,10-14} Hence, the ischemia-resistant cells would be expected to have a higher malignant potential than those before the onset of ischemia. Perhaps not by coincidence, tumors formed by cells that survive during hypoxia (the major consequence of ischemia) are particularly aggressive.⁶¹

ATG3 is upregulated in colorectal cancer cells during photodynamic therapy (PDT), a treatment that kills cancer cells by promoting their apoptosis.⁶² It was previously established that malignant cells that survive after PDT can form more aggressive tumors than those formed by the untreated cells.⁶³ It cannot be excluded that ATG3 contributes to PDT-induced apoptosis of cancer cells and that those cells that survive after such treatment (and are hence selected for resistance to ATG3-dependent apoptosis) possess increased anoikis resistance. Therefore tumors formed by such cells can be expected to display increased aggressiveness.

Levels of expression of some of the autophagy mediators in various human cancers were compared to those in respective normal cells in several studies, and used to assess the possible role of autophagy in the progression of these malignancies.^{37,44,45} To our knowledge, whether or not ATG3 levels change in the course of progression of any cancers has not yet been explored. Our data indicate that if changes in ATG3 expression are found in the future in human cancers, these changes could reflect the contribution of not only autophagy but also that of ATG3-dependent apoptosis in the course of the respective diseases.

In summary, we have found in this study that intestinal epithelial cell-ECM adhesion can switch the biological function of ATG3: ATG3 contributes to autophagy of detached cells but kills those attached to the ECM.

Materials and Methods

Cell culture

Cells were cultured as described previously.⁴ For suspension cultures cells were plated above a layer of 1% sea plaque agarose (IONZA, 50100) polymerized in α -minimal essential medium (MEM; Invitrogen, 11900) in case of IEC-18 cells and Dulbecco's modified essential medium (DMEM; Invitrogen, 12800) in case of Hkh-2 cells.

Western blot

This assay was performed as described previously.⁴ The following antibodies were used in the study. Anti-ATG3 (Cell Signaling Technology, 3415), anti-ATG7 (Cell Signaling Technology, 2631), anti-ATG16L1 (ProSci, 4425), anti-MAP1LC3B (Cell Signaling Technology, 3868), anti-GABARAP (Sigma, SAB2100873), anti-SH3GLB1 (ProSci, 45-143), anti-UVRAG (Abcam, ab72073), anti-ATG5 (Proteintech, 10181-2-AP), anti-ATG10 (Prosci 43990), anti-GFP (Clontech, 632381), anti-CDK4 (Santa Cruz Biotechnology, sc-601) and anti-AKTB (Santa Cruz Biotechnology, sc-130656).

RNA interference

RNAi was performed as described previously.²⁶ The sequences of the *siRNAs* were as follows. *ATG3 siRNA1*: UGUACAUCACUUACGACAA; *ATG3 siRNA2*: CGGAUGAAUUG GAGG CUAU; *ATG7 siRNA4*: GGCACGAACUGACCCAGAA; *ATG7 siRNA19*: AGUGAAUGCCA GCGGGUUC; *GABARAP siRNA2*: AUGAAAGCGUCUACGGUCU; *GABARAP siRNA3*: ACAAGAGGAGCAUCCGUU.

Quantitative polymerase chain reaction (qPCR)

This procedure was performed as described previously.¹⁸ Sequences of the primers used for the amplification of *ATG3* mRNA were GCAAACAAGAACCTATGACCTG and GTCTTCATACATGTGCTCAACTG, sequences of the primers used for the amplification of *ATG7* mRNA were GCTGGGAGAAGAACCAGAAA and GAGATTCAGATC-CACGGATGAC, sequences of the primers used for the amplification of *MAP1LC3B* mRNA were TCTTTGTAAGGGCG GTTCTG and TCACAAGCATGGCTCTCTTC, and sequences of the primers used for the amplification of *GABARAP* mRNA were TCAGGCTCAAACACCATCTC and GCGGCTGC ATCTACCATTA.

Expression vectors

pBabe puroWTATG3, and pBabe puroKRATG3 retroviral vectors were from Addgene (26926 and 26927, respectively). pBabe mCherry-GFP-LC3 vector was provided by Dr. N. Ridgway (Dalhousie University, Canada). pBabe as well as pHIT and pVSVG vectors (components of the retroviral packaging system) were provided by Dr. P. Lee (Dalhousie University, Canada). GFP-LC3B expression vector was provided by Dr. T. Yoshimori (Osaka University, Japan).

Production of the retroviruses

3×10^6 293T cells were plated on a 100 mm tissue culture dish, grown overnight and transfected with 10 μ g of either pBabe or pBabepuroWTAtg3 or pBabepuroKR Atg3 DNA, 5 μ g of pHIT and 5 μ g of pVSVG in the presence of 12 μ l of Lipofectamine 2000 (Invitrogen, 11668–019) in 6 ml of OPTI-MEM medium (Gibco, 31985–070). The medium was changed 4 h later to DMEM containing 10% fetal calf serum. The medium was collected 48 h later and filtered through a 0.45-micron filter unit. Viral concentration in the resulting solution was measured by use of the Retrovirus Quantitation Kit (Cell Biolabs Inc., VPK-120) according to the manufacturer's instructions.

Retroviral transduction

5×10^5 IEC-18 cells were plated in a 100 mm tissue culture dish and grown overnight. 4×10^4 viral particles per cell were then added to the cells in the presence of 8 μ g/ml polybrene (Sigma, 107689) and the cells were cultured for 24 h. Medium was then changed to fresh medium and the cells were cultured for 24 h in the presence of 2 μ g/ml puromycin. In the case of transduction of IEC-18 cells with the mCherry-GFP-LC3B virus, 6 ml of the virus-containing culture medium prepared as described above was added to 5×10^5 IEC-18 cells in the presence of 8 μ g/ml polybrene and the cells were further cultured for 24 h. Medium was then changed to fresh medium and the cells were cultured for an additional 24 h.

Detection of GFP-LC3B puncta in cells

In the case of the cells infected with retroviruses the cells were first transduced with the indicated viruses as described above and then transfected with 3 μ g GFP-LC3B expression vector in the presence of 6 μ l Lipofectamine 2000 in 6 ml of OPTI-MEM medium. Four h later the medium was changed to regular growth medium not containing puromycin. Twenty-four h later the cells were harvested by trypsinization, washed with phosphate-buffered saline (137 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 2 mM KH₂PO₄), and fixed with 4% paraformaldehyde for 20 min at room temperature. Images of the cells were captured by use of a Carl Zeiss Axiovert 200M microscope. In the case of the ATG3-deficient cells the cells were first transfected with the GFP-LC3B vector¹ and then with the control RNA and *ATG3*-specific *siRNAs*²⁶ and assayed as described above. In the case of the ATG7-deficient cells, the cells were initially transfected with respective RNAs²⁶ and then with the GFP-LC3B vector.¹ 8 h after the transfection ATG7-deficient cells were plated in monolayer or suspension.

Detection of mCherry-GFP-LC3 puncta in cells

Cells transduced with the mCherry-GFP-LC3-encoding virus as described above were analyzed by the use of a Carl Zeiss Axiovert 200M microscope for the presence of green, red, or yellow puncta.

Flow cytometry

This assay was performed as described previously.¹ Cells infected with retroviruses were cultured for 24 h in the absence of puromycin prior to being harvested for the assay.

Detection of apoptosis

Apoptosis was detected by use of ANXA5 as described previously.²⁵ Cells infected with retroviruses were cultured for 24 h in the absence of puromycin prior to being harvested for the assay.

Detection of clonogenic cell survival

Cell colonies were stained with Crystal Violet.²⁴ Cells infected with retroviruses were maintained in the presence of 2 μ g/ml puromycin for 7–10 d while the cell colonies were allowed to form.

Statistical analysis

Statistical analysis of the data in **Figures 1A, 2A, 3C, 4C, 5C, F, 6B, 7H, 8B, and C** was performed by the unpaired Student's *t*-test. Statistical analysis of the data in **Figures 1C, D and 5E**, was performed by the chi-square test for goodness-of-fit.

Western blot images

Western blot images were generated by Photoshop. When lanes were removed from these images, and separate parts of the same image were joined together, a vertical dotted line was used to indicate where the image was cut.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

Acknowledgments

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Supplemental Material

Supplemental data for this article can be accessed on the publisher's website.

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