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Subunit composition of AMPK trimers present in the cytokinetic apparatus:

Implications for drug target identification

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Abstract

AMP-activated protein kinase has been shown to be a key regulator of energy homeostasis; it has also been identified as a tumor suppressor and is required for correct cell division and chromosome segregation during mitosis. The enzyme is a heterotrimer, with each subunit having more than one isoform, each encoded by a separate gene (two α, two β and three γ isoforms). In human endothelial cells, the activated kinase subunit of AMPK in the cytokinetic apparatus is α2, the minority α subunit, which co-localizes with β 2 and γ 2. This is the first demonstration of a trimeric complex of AMPK containing the γ 2 regulatory subunit becoming selectively activated and being linked to mitotic processes. We also show that α 1 and γ 1, the predominant AMPK subunits, are almost exclusively localized in the cytoskeleton, while α 2 and γ 2 are present in all subcellular fractions, including the nuclei. These data suggest that pharmacological interventions targeted to specific AMPK subunit isoforms have the potential to modify selective functions of AMPK.

Keywords

AMP-activated protein kinase; isoforms; cytokinesis; subcellular localization; phosphorylation

Introduction

AMP-activated protein kinase (AMPK) is an evolutionarily conserved sensor and regulator of cellular energy balance (reviewed in ref. 1). It is widely accepted that AMPK exists as an αβγ heterotrimer. Phosphorylation of the catalytic α subunit by LKB1 or Ca^{2+} -calmodulindependent protein kinase kinase β results in activation of AMPK. The β subunit binds glycogen and acts as a scaffold for the other subunits.² Upon binding AMP when the AMP/ATP ratio increases, the regulatory γ subunit allosterically activates the kinase further.

Each AMPK subunit has more than one isoform, and each is encoded by a separate gene (α1, α2; β 1, β 2; γ 1, γ 2, γ 3). Given the number of subunit isoforms, there are 12 different

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possible combinations of AMPK trimers. Evidence is emerging that particular trimeric combinations are preferentially activated and mediate specific roles. For example, α2/β2/γ3 is selectively activated during exercise in human skeletal muscle, and this complex is likely responsible for phosphorylation of acetyl-coA carboxylase.³ This may be the result of the distinct subcellular localization of this complex, and differential localization may also explain the other reported selective activations of α 2-AMPK or β 1-AMPK complexes.³⁻⁵ AMPK containing α2 was reported to have greater AMP dependence than α1-AMPK, and large proportion of $a2$, but not $a1$, was shown to be in the nucleus.⁶ There is some evidence that subcellular localization of AMPK may be dependent on the regulatory subunit, as γ 1containing complexes have been shown to be directed to the Z-line in differentiated myotubes via the binding of the cytoskeletal protein plectin.⁷ However, none of the other γ isoforms have yet been localized.

The expression profile of AMPK subunit isoforms has been investigated.⁸ Based on coimmunoprecipitation and subsequent activity measurements, the γ1 isoform seems to be the major regulatory subunit in all cells; γ 3 is largely specific to skeletal muscle. Although γ 2 is not the major regulatory subunit of AMPK, it is also widely expressed, and the allosteric activation of complexes containing γ 2 is higher than that of γ 1-AMPK.⁹ Mutations in *PRKAG2*, which encodes the γ 2 subunit, cause hypertrophic cardiomyopathy with conduction disease; 10 these mutations are exclusively found in the nucleotide-binding domains, and some of them directly involved in binding nucleotides, AMP or ATP.¹¹

The regulatory role of AMPK in cellular energy homeostasis is not just the direct modulation of catabolic and anabolic processes; it also functions as a tumor suppressor via the inhibition of cell division, and it is implicated in the maintenance of cell polarity.¹ AMPK activates myosin by phosphorylating the non-muscle myosin regulatory light chain,¹² and non-muscle myosin II has been shown to be directly involved in karyokinesis.¹³ Activated AMPK, the phosphorylated form of the α subunits (P- α), was shown to be in transient association with several mitotic structures, including centrosomes, spindle poles, the central spindle midzone and the midbody during mitosis and cytokinesis.¹⁴ Moreover, during the G_1/S -to-M-phase transition, AMPK activation resembles strikingly that of wellcharacterized "chromosomal passenger" proteins, such as Aurora B, INCENP or histone H3,¹⁵ and, therefore, the tumor suppressive activity of AMPK may manifest in the mitotic/ cytokinetic phase of the cell cycle.16 AMPK has also been shown to regulate meiotic maturation of mouse $\text{ocytes}^{17,18}$ and has been reported to activate directly stress-promoted transcription by phosphorylating histone H2B.¹⁹

In this paper, we have used isoform-specific antibodies to investigate AMPK subcellular localization by western blotting and immunofluorescence. We report that the activated AMPK complex linked to the mitotic apparatus is α 2/β2/γ2, and we provide further evidence for the compartmentalization of different AMPK trimers, most likely directed by the regulatory γ subunit.

Results

Cytosolic, membrane/particulate, nuclear and cytoskeletal fractions of HUVECs were prepared by sequential detergent extraction.²⁰ The overall protein composition was analyzed by SDS-PAGE, which revealed the expected presence of vimentin (54 kDa) as a clear marker in the cytoskeleton (Fig. 1A). Western blots of the separated proteins of each fraction were probed with antibodies specific to AMPK subunit isoforms (Fig. 1B). The α1 subunit, which accounts for more than 80% of the total α protein in HUVEC,²¹ was exclusively detected in the cytoskeleton fraction by western blotting. The less abundant α subunit, α2, was detectable as a doublet at low levels in western blots of each fraction. The activated kinase subunit (P-α) was present predominantly in the cytosolic and the membrane/particulate fractions, with a lesser amount detected in the nuclear fraction. The γ 1 subunit is the major regulatory subunit in $HUVEC^{21}$ and was detected in the cytoskeleton. The minority regulatory subunit, γ 2, was present in all fractions.

To gain information on the specific localization of AMPK subunits in HUVEC, we used confocal microscopy to examine cells stained with subunit-specific antibodies. Of particular interest was the localization of activated AMPK complexes during the cell cycle, as P-α has recently been shown to associate transiently in human cancer-derived epithelial cells with several mitotic structures including centrosomes, spindle poles, the central spindle midzone and the midbody during mitosis and cytokinesis.¹⁴ We found a similar pattern in dividing HUVEC and, in particular, detected strong staining with P-α antibody of the midbody and cleavage furrow (co-localization with actin) (Fig. 2A). As there is some evidence that the γ subunit may direct AMPK subcellular localization,⁷ we examined the distribution of the γ isoforms in these cells. In spite of the recent report of the nuclear presence of γ 1,²² we did not observe γ 1 in the nuclei, nor did we find co-localization with the activated AMPK during mitosis. In general, γ1 staining was dispersed throughout the cytoplasm (Fig. 2B). However, strong staining of the midbody, similar to P- α , was also shown for γ 2, suggesting that activated AMPK at that site contains this regulatory subunit (Fig. 2B). The staining of the midbody was achieved with two different γ 2 antibodies: one was raised against a short C-terminal peptide, γ2C (Fig. 2C); the other antibody was raised against a longer midsegment of the protein (Fig. 2B). Specific co-localization of P- α and γ 2 was not possible, as the P- α and both γ 2 antibodies were raised in rabbit. Both γ 2 antibodies also stain the nuclei (but not the nucleoli) and the condensed chromosomes (Figs. 2B, 3A and B), and we found intense staining of γ 2 around the separating chromosomes (Fig. 2B). The staining around the condensed and segregating chromosomes suggests that AMPK containing γ2 may be associated with the kinetochore. Interestingly, no P-α was detected around the segregating chromosomes during mitosis.

In order to determine the catalytic subunit present along with γ 2 at the midbody, HUVEC were stained with antibodies to the α isoforms. The α1 subunit was found to be dispersed throughout the cell with little apparent nuclear staining (no co-localization with nuclear γ 2) and did not concentrate at the midbody (Fig. 3A). In contrast, distinct midbody staining was shown for α 2, with distinct co-localization with γ 2 at this position (Fig. 3A); some α 2 nuclear staining was also found in non-dividing cells (data not shown). Attempts to use the

Cell Cycle. Author manuscript; available in PMC 2015 June 03.

α antibodies with the P-α antibody were unsuccessful, presumably due to strong binding of the P-α antibody hindering the other α antibody.

The distribution of the β subunits in dividing HUVEC contrasted markedly. β1 was shown to be clearly extranuclear, with typically a granular appearance; in some instances, the staining coincided with the midbody, although did not precisely colocalize with γ 2 (Fig. 3B). The β 2 antibody was found to give nuclear and midbody staining, with the latter corresponding to the P-α localization. We conclude that the subunit composition of activated AMPK at the midbody is α2/β2/γ2.

Discussion

The specific localization of activated AMPK with several mitotic structures in human cancer-derived epithelial cells has recently been reported in reference 14. We have confirmed this in HUVEC and identified the subunit composition of activated AMPK at the midbody to be α 2/β2/γ2. This is the first demonstration of AMPK containing the γ 2 regulatory subunit becoming selectively activated and being linked to a specific cellular process. As non-muscle myosin II has been identified as an AMPK substrate and is known to be involved in scission during telophase, we suggest that one of the functions of the α2/β2/γ2 AMPK complex may be to regulate non-muscle myosin-II during cytokinesis directly or indirectly.^{12,13} In a recent publication, it was shown that AMPK directly phosphorylates several proteins which coordinate different aspects of mitosis.23 By phosphorylating protein phosphatase 1 regulatory subunit 12C and p21-activated protein kinase, AMPK indirectly regulates myosin regulatory light chain phosphorylation and, therefore, cytoskeleton rearrangement. This study was performed by overexpressing the $a2$ catalytic subunit co-expressed with $\beta1/\gamma1$; however, the modified $\alpha2$ can form trimers with endogenous $\frac{\beta 2}{\gamma^2}$, which can be directed to the mitotic apparatus. On the other hand, NUAK2, a member of the AMPK family, also plays an important role in regulating the cytoskeleton and influencing cell proliferation and cell motility; aberration of both of these can result in tumorigenesis, and there may an interaction between the two pathways.²⁴

AMPK activation can occur selectively in the nucleus. Polo-like kinase 1, the key regulator of cell cycle events, has been implicated in the nuclear activation of AMPK during mitosis.²⁵ Furthermore, substantial P- α staining was detected first in the nuclei after radiation treatment in tumor cells, and the increase in cytoplasmic staining only occurred subsequently.²⁶ The nuclear activation of AMPK by ionizing radiation was independent of LKB1 but, instead, dependent on ATM, the DNA damage sensor.

Given the extranuclear localization of the γ 1 subunit, it is likely that other nuclear processes associated with AMPK (e.g., the activation stress-promoted transcription¹⁹) may also involve complexes containing γ 2. In addition, γ 2 being in the vicinity of the segregating chromosomes may indicate that γ 2-containing AMPK is required for the formation of kinetochores. AMPK involvement in faithful chromosome segregation has been reported in reference 14. The kinase subunit in this case might be a_1 , as this catalytic subunit was shown to associate with the condensed chromatin during meiotic maturation in mouse oocytes,¹⁷ and activated α 1 also appears to translocate into the nucleus during the

development of Huntington disease.²⁷ We only detected α 1 in the cytoskeletal fraction of HUVECs, while P- α and α 2 both were present in the nuclear fraction (Fig. 1B). An earlier study showed that a significant proportion of α2 but not α1 was found in the nucleus and also found that AMP activated the α 2-containing complexes to a greater extent than α 1-AMPK.⁶ Our current work suggests these α 2-containing complexes would have γ 2 as their regulatory subunit and, as AMPK containing γ 2 is more sensitive to changes of AMP $concentrations⁹$ this is the likely explanation of their greater activation.

It is unclear whether α2/β2/γ2 AMPK shuttles between the cytoplasm and the nucleus. Nuclear entry of α 2/β2/γ2 complexes may be directed by α 2, which has a nuclear localization signal,²⁸ and/or by γ 2, which, unlike the γ 1 isoform, has both a strong atypical bipartite nuclear localization signal (9-**KKK K**DV SSP GGS GG**K K**NA SQ**K RR**-31) and a nuclear exit signal (546-**L**SD **IL**Q A**L**I **L**-556). If some γ2 is permanently in the nucleus (Figs. 2 and 3), it could plausibly act by recruiting activated and translocated $\alpha/2/\beta$ 2 subunits into distinct positions at different stages of mitosis. Only a weak interaction was found between the β and the γ subunits, whereas the binding between α -β was reported to be very strong.²⁹ It has been reported that in human skeletal muscle biopsies, γ 2 is present but does not appear to participate in any AMPK complex.³⁰ Therefore the low contributions (less than 10%) of γ 2-containing complexes to the overall AMPK pool⁸ may be the result of γ 2's transient participation in an AMPK complex; hence, the majority of γ 2 may be not in complex and its cellular presence underestimated.

Based on our observations, we think that AMPK activation and function depends on its subcellular localization, and that it may be largely dependent on the regulatory γ subunit in the complex. We only detected γ 1 in the cytoskeleton, and the majority of α 1 is also in this fraction (Fig. 1B). Deletion of either α 1 or γ 1 causes premature death of erythrocytes in mouse, and that seems to be the only phenotype.³¹ AMPK has been shown to regulate cell structure,¹² and the loss of either α 1 or γ 1 is detrimental after the erythrocytes become enucleated. However, selective activation that depends on other AMPK subunits has also been reported. The α1-containing AMPK is preferentially phosphorylated in muscle cells upon metformin treatment, and fatty acid metabolism is inhibited, while AICAR preferentially activates α 2-containing complexes.³² These observations would suggest that AMPK activation is dependent on the kinase subunit. On the other hand, that may be simply the consequence of locating the α 1-AMPK or α 2-AMPK to certain positions in the cell and predisposing them for selective activation.

The identification of a specific AMPK trimer that is specifically localized and activated during cytokinesis raises the question whether this complex may be selectively targeted pharmacologically to specifically modulate mitosis. While thienopyridone compounds target β1-containing complexes,33 most of the commonly used AMPK activators or inhibitors are not selective. Metformin, a potent activator of AMPK originally used to treat hyperglycemia and type 2 diabetes, also has anticancer properties; the latter is partly due to impaired cytokinesis.16 The other axis of cancer treatment is the AMPK-FoxO3A pathway, also activated by metformin; AMPK via the transcription factor FoxO3A modulates not only cell metabolism, but also stress resistance, autophagy and cell death.³⁴ There are a number of possible targets of AMPK. If the γ subunits are the chief mediator of AMPK localization,

Cell Cycle. Author manuscript; available in PMC 2015 June 03.

and, as a consequence, the compartmentalized complex is assigned to a specific signaling pathway, targeting the γ subunits could be a profitable direction for new, selective drug design for the manipulation of specific AMPK functions.

Experimental Procedures

Cell lines

Primary HUVEC (Invitrogen) were cultured in Medium 200 (Invitrogen) supplemented with Low-Serum Growth Supplement (Invitrogen). Cells were used between passages 2–5.

Reagents

The FractionPREP™ cell fractionation kit was purchased from BioViosion and used according to manufacturer's instructions.

Primary antibodies used were as follows: Rabbit mAB anti-P-AMPK α^{Thr172} (clone 40H9) and polyclonal rabbit AMPK γ 1 antibody (Cell Signaling Technology); rabbit polyclonal γ 2 anti-body—anti-PRKAG2 antibody (Atlas Antibodies); rabbit polyclonal C-terminal γ 2 antibody (gift from D. Carling); goat polyclonal AMPKγ1 (T-20) (sc-19138), goat polyclonal AMPKα1 (C-20) (sc-19128), polyclonal AMPKα2 (A-20) (sc-19129), polyclonal AMPKβ1 (N-18) (sc-19132) and polyclonal AMPKβ2 (E-20) (sc-19136) were all purchased from Santa Cruz Biotechnology, Inc.

For fluorescence detection of immunostaining, donkey anti-goat Alexa Fluor 488 and donkey anti-Rabbit Alexa Fluor 594 (both Invitrogen) were used. The actin network was visualized by Phalloidin-Tetramethylrhodamine B isothiocyanate (Phalloidin-TRITC) (Sigma) staining.

Immunofluorescence staining and confocal imaging

HUVEC were grown on poly-D-Lysine-coated coverslips in 12-well plates. Cells were fixed in 4% PFA, permeabilized in 0.2% Triton® X-100, blocked with 3% BSA in PBS then incubated with the appropriate primary and secondary antibodies.

Cells were mounted with *SlowFade®* Gold antifade reagent with DAPI (Invitrogen) and imaging was performed with a Leica TCS SP5 confocal laser-scanning microscope with a 63×, 1.4NA objective.

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Cell Cycle. Author manuscript; available in PMC 2015 June 03.

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Figure 1.

Subcellular distribution of AMPK subunit isoforms. (A) SDS-PAGE was stained with Coomassie brilliant blue. (B) Western blots with antibodies α1 and α2 (Santa Cruz); p-α and γ1 (Cell Signaling); γ2 (Atlas antibodies). Approximately $10⁷$ cells used for cell fractionation and equal volumes of the fractions were loaded on the SDS-PAGE. T, total cell extract; M, molecular weight markers; CS, cytosol; MP, membrane/particulate; N, nuclear; CSk, cytoskeletal fractions.

Figure 2.

Co-localization of activated AMPK with γ2 in dividing HUVEC. Immunofluorescence localization was performed with antibodies raised against p- α , γ 2 and a γ 2 C-terminal peptide. Actin was stained with phalloidin-TRITC. The different subunits are labeled in appropriately colored text in each part.

Figure 3.

Localization of AMPK subunits in HUVECs. (A) Localization of α 1 and α 2 along with γ 2. (B) Localization of β1 and β2 along with γ2 and phospho-α. Immunofluorescence localization was performed using antibodies raised against p-α, γ2 and α1/α2, β1/β2. The different subunits are labeled in appropriately colored text in each part.