

AMP-activated protein kinase phosphorylates and inactivates liver glycogen synthase

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Recombinant muscle GYS1 (glycogen synthase 1) and recombinant liver GYS2 were phosphorylated by recombinant AMPK (AMP-activated protein kinase) in a time-dependent manner and to a similar stoichiometry. The phosphorylation site in GYS2 was identified as Ser⁷, which lies in a favourable consensus for phosphorylation by AMPK. Phosphorylation of GYS1 or GYS2 by AMPK led to enzyme inactivation by decreasing the affinity for both UDP-Glc (UDP-glucose) [assayed in the absence of Glc-6-P (glucose-6-phosphate)] and Glc-6-P (assayed at low UDP-Glc concentrations). Incubation of freshly isolated rat hepatocytes with the pharmacological AMPK activators AICA riboside (5-aminoimidazole-4-carboxamide-

1-β-D-ribofuranoside) or A769662 led to persistent GYS inactivation and Ser⁷ phosphorylation, whereas inactivation by glucagon treatment was transient. In hepatocytes from mice harbouring a liver-specific deletion of the AMPK catalytic α1/α2 subunits, GYS2 inactivation by AICA riboside and A769662 was blunted, whereas inactivation by glucagon was unaffected. The results suggest that GYS inactivation by AMPK activators in hepatocytes is due to GYS2 Ser⁷ phosphorylation.

Key words: A769662, 5-aminoimidazole-4-carboxamide-1-β-D-ribofuranoside (AICA riboside), AMP-activated protein kinase (AMPK), glucagon, glycogen synthase, hepatocyte.

INTRODUCTION

Glycogen is a branched homopolymer made of glucosyl units and is essential for whole-body glucose homeostasis. During hyperglycaemia, glucose is incorporated into glycogen by the combined action of GYS (glycogen synthase) and glycogen branching enzyme, which catalyse α-1,4- and α-1,6-glycosidic bond formation respectively [1]. In mammals, a muscle isoenzyme (GYS1) is ubiquitously expressed, but in liver the hepatic isoenzyme (GYS2) is predominantly expressed [1]. These isoenzymes are 70% identical and possess a highly conserved catalytic central region flanked by two variable extremities [2].

The rate of incorporation of glucose into liver glycogen directly correlates with GYS activity, which is regulated by allosteric and phosphorylation/dephosphorylation mechanisms [2–4]. Several protein kinases inactivate GYS by phosphorylation. However, GYS inactivation, at least in skeletal muscle, can be overcome to some extent by Glc-6-P (glucose 6-phosphate) an allosteric stimulator, whose intracellular concentration rises when glucose transport is increased [5,6]. A total of nine different phosphorylation sites are found in the three domains of GYS1, seven of which are conserved in GYS2 [2,7]. The N-terminal domain contains site 2 (Ser⁷) in GYS1 phosphorylated by different protein kinases, including PKA (cAMP-dependent

protein kinase), PhK (phosphorylase kinase) and AMPK (AMP-activated protein kinase), leading to GYS inactivation [8,9]. Site 2 phosphorylation allows site 2a (Ser¹⁰) phosphorylation by the constitutively active CK1 (casein kinase 1), resulting in further and substantial enzyme inactivation [10]. The central domain contains sites 3a (Ser⁶⁴⁰), 3b (Ser⁶⁴⁴), 3c (Ser⁶⁴⁸), 4 (Ser⁶⁵²) and 5 (Ser⁶⁵⁶), whose phosphorylation is hierarchical and also inactivates GYS [7]. Phosphorylation at site 5 by CK2 allows sequential phosphorylation of the other sites by GSK3 (glycogen synthase kinase 3). In addition, sites 3a and 3b can be phosphorylated directly by other kinases [11–13]. The C-terminal domain, which is absent in GYS2, contains site 1a (Ser⁶⁹⁷) phosphorylated by PKA, and site 1b (Ser⁷¹⁰) phosphorylated by Ca²⁺/calmodulin-dependent protein kinase II [2,14]. The relative importance of each site in the control of GYS1 activity has been studied by site-directed mutagenesis, revealing a crucial role for site 2, 2a, 3a and 3b phosphorylation for the control of GYS1 expressed in COS cells [15]. For GYS2, on the other hand, mutation of Ser⁷ to an alanine residue suffices to fully activate the enzyme expressed in FTO2B cells [16], suggesting that phosphorylation at Ser⁷ might be more important for GYS2 inactivation than phosphorylation at the other conserved GSK3/CK1/CK2 sites. One candidate kinase for phosphorylation of GYS2 at Ser⁷ is AMPK, known to phosphorylate GYS1 site 2

Abbreviations used: ACC, acetyl-CoA carboxylase; AICA riboside, 5-aminoimidazole-4-carboxamide-1-β-D-ribofuranoside; AMPK, AMP-activated protein kinase; CK, casein kinase; eEF2, eukaryotic elongation factor-2; Glc-6-P, glucose 6-phosphate; GP, glycogen phosphorylase; GSK3, glycogen synthase kinase 3; GST, glutathione transferase; GYS, glycogen synthase; HEK, human embryonic kidney; KO, knockout; LKB1, liver kinase B1; LS, liver-specific; PKA, cAMP-dependent protein kinase; PP1c, protein phosphatase 1γ catalytic subunit; PTG, protein-targeting glycogen; SRM, single reaction monitoring; UDP-Glc, UDP-glucose; WT, wild-type.

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in vitro and *in vivo* [8,17,18]. Moreover, this site lies in a favourable consensus for AMPK phosphorylation in GYS2.

AMPK is a heterotrimeric protein kinase composed of a catalytic α -subunit and two regulatory subunits (β and γ) [19–21]. Each subunit has different isoforms allowing the expression of 12 different complexes, excluding splice variants. AMPK activation occurs during metabolic stress and requires activation loop Thr¹⁷² phosphorylation of its α -catalytic subunits by upstream kinases. Once activated, AMPK stimulates ATP-producing pathways and inhibits anabolic processes to maintain intracellular energy status [19,20]. AMPK activation also influences whole-body energy homeostasis through appetite control and re-orientation of metabolism [19,21]. Important effects of AMPK activation in liver, the main site of action of the anti-diabetic drug metformin, are to reduce glucose production and lipid synthesis by decreasing the expression of key gluconeogenic and lipogenic enzymes [22,23]. However, metformin also inhibits glucose production by decreasing ATP levels independent of AMPK [24]. AMPK in liver is also activated under energy stress, such as fasting, caloric restriction and exercise [25]. Pharmacological and phytochemical compounds, such as AICA riboside (5-aminoimidazole-4-carboxamide-1- β -D-ribofuranoside), the thienopyridone A769662, and the herbal medicines berberin and resveratrol, have also been used to activate liver AMPK [25]. AMPK activation in liver restores energy status by switching from an anabolic to a catabolic state, i.e. by increasing lipid β -oxidation and/or decreasing synthesis of proteins, fatty acids, triacylglycerol and cholesterol. AMPK activation also has long-term effects by repressing lipogenic gene expression and enhancing mitochondrial biogenesis [26,27].

Despite physiological AMPK activation in liver and the presence of site 2 of GYS2 in a favourable consensus for AMPK, the role of AMPK in the regulation of liver GYS has not received attention. In the present study, we demonstrate that AMPK phosphorylates and inactivates recombinant GYS2 via Ser⁷ phosphorylation. We also report that AMPK activation by pharmacological activators leads to GYS inactivation and Ser⁷ phosphorylation in freshly isolated rat hepatocytes and that GYS inactivation by the compounds was blunted in incubated hepatocytes from mice whose two AMPK catalytic subunits had been genetically deleted.

EXPERIMENTAL

Materials

AICA riboside (Toronto Research Chemicals) and glucagon were from Novo Nordisk. A769662 was kindly provided by Dr Anudharan Balendran (AstraZeneca, Mölndal, Sweden). The AMARA [28] and MR6 [29] peptides were kindly synthesized by Dr Vincent Stroobant (Ludwig Institute, Brussels, Belgium). Anti-(p-Thr¹⁷² AMPK α -subunit), anti-[p-Ser⁷⁹ ACC1 (acetyl-CoA carboxylase 1)] (Millipore) and anti-[total eEF2 (eukaryotic elongation factor-2)] (Santa Cruz Biotechnology) antibodies were from the sources indicated. Anti-(total AMPK $\alpha 1/\alpha 2$), anti-(total GYS1) and anti-(p-Ser⁷ GYS1) antibodies were kindly provided by Professor Grahame Hardie (College of Life Sciences, University of Dundee, Dundee, Scotland, U.K.). Anti-GYS2 and anti-[p-Ser¹⁴ GP (glycogen phosphorylase)] antibodies [5] were from the MRC Protein Phosphorylation Unit (University of Dundee, Dundee, Scotland, U.K.). Anti-(p-Ser⁷ GYS2) was raised in rabbit as described previously [16], and anti-(sheep IgG coupled to horseradish peroxidase), anti-(total GP) (Sigma) and anti-(rabbit IgG coupled to horseradish peroxidase) (GE Healthcare) antibodies were from the sources indicated.

Recombinant GYS1 and GYS2 were purified as described previously [5]. Briefly, HEK (human embryonic kidney)-293 cells were co-transfected with expression vectors for untagged GYS1 or GYS2 and GST (glutathione transferase)-tagged glycogenin. After 36 h, the cells were lysed and GYS–GST–glycogenin complexes were purified by chromatography on glutathione–Sepharose with elution by glutathione. Recombinant bacterially expressed $\alpha_1\beta_1\gamma_1$ AMPK was fully activated by incubation with recombinant bacterially expressed LKB1 (liver kinase B1)–Mo25–STRAD (STE20-related kinase adapter protein) complex, both kindly provided by Dr Dietbert Neumann (ETH Zurich, Switzerland), as described previously [30] except that LKB1 was used instead of calmodulin-dependent protein kinase kinase- β . PKA catalytic subunits were purified from bovine heart as described previously [31]. Bovine hearts were obtained from the local slaughterhouse. Recombinant CK1 was from New England Biolabs. PP1c–GST (PP1c is protein phosphatase 1 γ catalytic subunit) was purified using GSTrapTM HP columns (GE Healthcare) from a lysate of *Escherichia coli* BL21 bacteria transformed with pGEX-PP1c, and cultured in the presence of 1 mM MnCl₂.

Experiments involving the preparation of hepatocytes from anaesthetized rats and mice were approved by the local animal ethics committee.

Protein kinase assays

AMPK and PKA were assayed by measuring ³²P incorporation from 0.1 mM [γ -³²P]MgATP (specific radioactivity of 1000 c.p.m./pmol) into 200 μ M AMARA peptide (AMARAAS-AAALRRR) for AMPK and 200 μ M MR6 (PVRMRNSFT) peptide for PKA, as described previously [32]. A protein kinase activity of 1 unit corresponds to the amount of enzyme that catalyses the incorporation of 1 nmol of ³²P/min into the peptide substrate under the appropriate assay conditions.

Phosphorylation of recombinant GYS preparations

Recombinant GYS proteins (2.5 μ g) were first dephosphorylated by incubating with 0.7 μ g of purified recombinant PP1c–GST in 30 μ l of dephosphorylation buffer (50 mM Hepes, pH 7.5, 1 mM MnCl₂, 0.1 mM EGTA and 0.1% 2-mercaptoethanol) for 30 min at 30 °C and the reactions were stopped by adding 150 nM microcystin-LR (Sigma). The preparations were then phosphorylated with 200 m-units of AMPK or 200 m-units of PKA in 60 μ l of reaction mixture containing 25 mM Hepes, pH 7.5, 0.5 mM MnCl₂, 50 μ M EGTA, 0.05% 2-mercaptoethanol, 200 μ M AMP, 10 mM MgCl₂, 175 nM microcystin and 0.1 mM [γ -³²P]MgATP (specific radioactivity 1000 c.p.m./pmol) at 30 °C. Aliquots (10 μ l) were removed at the times indicated in the Figure legends and stopped in SDS/PAGE sample buffer [100 mM Tris/HCl, pH 6.8, 10% (v/v) glycerol, 5% (v/v) 2-mercaptoethanol, 8% (w/v) SDS and 0.025% Bromophenol Blue]. Proteins were separated by SDS/PAGE (10% gels) for staining with PAGEBlueTM (Thermo Scientific). GYS protein was quantified by gel infrared imaging (LI-COR OdysseyTM) along with known amounts of rabbit aldolase whose concentration had been determined from the calculated molar absorption coefficient and the measured absorbance at 280 nm [30]. For determination of ³²P incorporation, GYS bands were excised from the gel, dissolved in 500 μ l of 3% (w/v) H₂O₂ (Sigma) by heating for 2 h at 80 °C, mixed with 5 ml of scintillant (Ultima GoldTM; PerkinElmer) and radioactivity was counted. Stoichiometries of phosphorylation were expressed as mol of ³²P incorporated/mol of GYS using the calculated molecular mass of each isoenzyme.

Analysis of phosphorylated recombinant GYS preparations by MS

Recombinant GYS1 and GYS2 proteins (1.5 μg) were dephosphorylated then phosphorylated, using non-radioactive MgATP, by PKA and AMPK as described above. Following SDS/PAGE and in-gel trypsin digestion, peptides were analysed by static nano-electrospray ionization–tandem MS in an LTQ XL ion-trap mass spectrometer (ThermoFisher Scientific) and the phosphorylation sites were pinpointed using an MS²/MS³ neutral loss method. For quantification by SRM (single reaction monitoring), tryptic peptides were analysed by LC–tandem MS with the LTQ XL equipped with a microflow ESI source interfaced to a Dionex Ultimate Plus Dual gradient pump, a Switchos column switching device and a FAMOS Autosampler (Dionex). Separation was on a BioBasic C18 column (180 $\mu\text{m} \times 15 \text{ cm}$; ThermoFisher Scientific) equilibrated in solvent A [5% (v/v) acetonitrile and 0.05% formic acid in water] using a 90 min gradient from 0 to 70% solvent B [80% (v/v) acetonitrile and 0.05% formic acid in water] at a flow rate of 1.5 $\mu\text{l}/\text{min}$. The mass spectrometer was operated in a data-dependent manner to follow up four SRM transitions corresponding to mono-, di- and un-phosphorylated forms of the Ser⁷ and Ser¹⁰ GYS2 peptide and an internal tryptic peptide from GYS2 as a reference. Abundances of each molecular species were determined by peak area intensity integration.

GYS assay

The recombinant GYS1/2 preparations were dephosphorylated then rephosphorylated by AMPK or PKA, with or without CK1 (5 units to ensure maximal inactivation) as described above using non-radioactive MgATP. The reaction mixtures were placed on ice for the GYS assay using a modification of the method described in [33] by following the incorporation of UDP-[U-¹⁴C]-D-Glc (PerkinElmer) into glycogen. Briefly, to determine the K_m for UDP-Glc (UDP-glucose), 100 ng of GYS was incubated in 100 μl of assay buffer containing 25 mM Hepes, pH 7.8, 50 mM NaF, 1 mM EDTA, 0.7% shellfish glycogen and 0, 0.05, 0.1, 0.5, 1, 2.5 or 4.6 mM UDP-[U-¹⁴C]-D-Glc with or without 10 mM Glc-6-P. To determine the K_a for Glc-6-P, GYS2 was incubated as described above in the presence of 0.25 mM UDP-[U-¹⁴C]-D-Glc and 0, 0.01, 0.02, 0.05, 0.125 or 0.25 mM Glc-6-P. The K_a of GYS1 for Glc-6-P was measured with 0.1 mM UDP-[U-¹⁴C]-D-Glc and 0, 0.02, 0.05, 0.125, 0.25 or 1 mM Glc-6-P. The reactions were stopped by spotting 90 μl aliquots on to 2 cm \times 2 cm Whatman 31 ET CHR papers (GE Healthcare), which were plunged into ice-cold 66% (v/v) ethanol. After washing in ethanol and a final wash in acetone, the dried papers were transferred to vials for scintillation counting. The kinetic parameters V_{max} , K_m for UDP-Glc and K_a for Glc-6-P were obtained by non-linear curve fitting using GraphPad Prism.

AMPK $_{\alpha 2\text{LS}}$ ^{-/-} mice

LS (liver-specific) AMPK $_{\alpha 2}$ ^{-/-} mice were obtained by crossing floxed AMPK $_{\alpha 2}$ mice with mice expressing Cre recombinase under the control of both the mouse albumin regulatory elements and the α -fetoprotein enhancers. AMPK $_{\alpha 1\alpha 2\text{LS}}$ ^{-/-} mice were then bred by crossing these mice with AMPK $_{\alpha 1}$ ^{-/-} mice [34].

Preparation and incubation of freshly isolated rat and mouse hepatocytes

Freshly isolated hepatocytes from anaesthetized overnight-starved male Wistar rats (approximately 200 g of body weight) were prepared by perfusion with collagenase as described

previously [35]. A scaled-down procedure was also used to prepare isolated cells from starved WT (wild-type) and AMPK $_{\alpha 1\alpha 2\text{LS}}$ ^{-/-} mice (approximately 30 g of body weight). Hepatocytes (50 mg of wet weight/ml) were gassed with O₂/CO₂ (19:1) and pre-incubated for 40 min in Krebs–Henseleit buffer (118 mM NaCl, 4.7 mM KCl, 2.5 mM CaCl₂, 1.18 mM KH₂PO₄, 1.18 mM MgSO₄ and 25 mM NaHCO₃, pH 7.4) with 30 mM glucose, 5 mM glutamine and 5 mM leucine at 37°C to fully activate GYS [36]. Agonists were then added to the incubation medium at final concentrations of 1 mM AICA riboside, 30 μM A769662 (from a stock solution in DMSO) or 10⁻⁷ M glucagon (control incubations without A769662 received 0.1% DMSO). Duplicate samples were taken at 0, 5, 15 and 30 min. For GYS assay, 250 μl of cell suspension was removed, mixed with 50 μl of ice-cold buffer containing 120 mM glycylglycine, pH 7.4, 75 mM KF, 7 mM EDTA and 0.5% shellfish glycogen and immediately frozen in acetone/solid CO₂. Thawed extracts were centrifuged and aliquots (25 μl) were assayed for GYS as described above with 2.5 mM UDP-Glc, but in the absence of Glc-6-P and in the presence of 9.3 mM Na₂SO₄ [37]. In some experiments, GP was assayed as described previously [37]. For AMPK assays and immunoblotting, cells were harvested by rapidly centrifuging 250 μl of cell suspension (Beckman bench Microfuge B; 20 s at 9500 g). After removing supernatants, the cell pellets were immediately frozen in liquid nitrogen for storage at -80°C. Extracts were prepared and precipitated with 10% (w/v) poly(ethylene glycol) 6000 for AMPK assays and immunoblotting [32].

Immunoblotting

Extracts from rat and mouse hepatocytes (50 μg of protein) were subjected to SDS/PAGE. Anti-(rabbit IgG) conjugated to IR dye 800 (Rockland) and anti-(sheep IgG) conjugated to Alexa Fluor[®] 680 (Molecular Probes) were used to measure the phosphorylation state of AMPK and ACC by LI-COR Odyssey[™] infrared imaging as described previously [32]. The phosphorylation states of GYS2 Ser⁷ and GP Ser¹⁴ were measured by chemiluminescence as described previously [32], and equal sample loading was verified by blotting for full-length eEF2.

Other methods

Protein was estimated by the method of Bradford [38] with γ -globulin as a standard. The results are expressed as means \pm S.E.M. for the indicated number of individual experiments. The statistical significance of the results was assessed using a paired two-sided Student's *t* test.

RESULTS

AMPK phosphorylates Ser⁷ of GYS2

GYS1 site 2 (Ser⁷) is phosphorylated by AMPK *in vitro* [8] and in AICA riboside-treated and contracting muscles [18,39,40]. The corresponding site in GYS2 (Ser⁷) also lies in a favourable consensus for phosphorylation by AMPK (Figure 1A). We first compared the time course of phosphorylation of recombinant GYS1 by equal amounts of AMPK or PKA catalytic subunits in terms of activity units measured towards substrate peptides. After 40 min of incubation, when the stoichiometry of ³²P incorporation was maximal, phosphorylation by PKA was approximately 50% higher than that observed with AMPK, reaching 0.46 ± 0.03 compared with 0.31 ± 0.02 mol of ³²P incorporated/mol of GYS1 for AMPK (Figure 1B). This result is consistent with the fact that

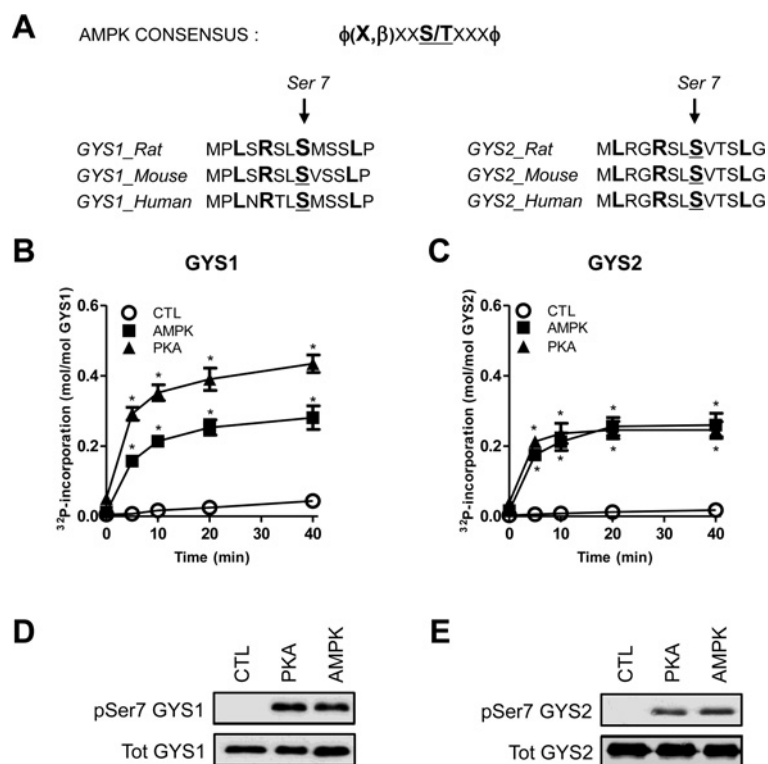


Figure 1 Phosphorylation of recombinant GYS1 and GYS2 by AMPK and PKA

(A) Upper panel: AMPK consensus sequence surrounding the phosphorylated serine/threonine residue, where ϕ represents a hydrophobic residue, β is a basic residue and X is a variant. The parentheses indicate that the order of the variable and basic residues at the $-4/-3$ positions is not critical. Lower panel: an alignment of the N-terminal sequences of rat, mouse and human GYS1 and GYS2 is shown. Time course of phosphorylation of GYS1 (B) or GYS2 (C) after dephosphorylation by PP1c-GST and incubation for 0, 5, 10, 20 and 40 min with 0.1 mM [γ - 32 P]MgATP (control, CTL) or in the presence of 200 m-units of recombinant activated $\alpha_1\beta_1\gamma_1$ AMPK or purified PKA catalytic subunits. Results are means \pm S.E.M. for three independent experiments. * $P < 0.05$ compared with the control values (paired two-sided Student's t test). Representative immunoblot from three incubations of GYS1 (D) and GYS2 (E) phosphorylated as described above for detection with the appropriate anti-pSer⁷ GYS1 or GYS2 antibodies.

there are two sites for PKA and one AMPK site described for the muscle GYS1 isoenzyme [7,8]. Like GYS1, GYS2 was also phosphorylated in a time-dependent manner by PKA and AMPK, but to a similar extent (maximal stoichiometry of 0.29 ± 0.02 and 0.25 ± 0.02 mol of 32 P incorporated/mol of GYS2 respectively; Figure 1C). For PKA, only one phosphorylation site in GYS2 has been reported [2] and the comparable stoichiometry of phosphorylation of GYS1 and GYS2 by AMPK (Figure 1C) would suggest the presence of a single AMPK phosphorylation site in GYS2. Overall the stoichiometries of phosphorylation of the two GYS preparations were low, possibly because only a proportion of recombinant protein was in the proper conformation for phosphorylation by protein kinases.

In order to identify the site phosphorylated by AMPK in GYS2, recombinant GYS1 and GYS2 preparations were phosphorylated with non-radioactive ATP and either AMPK or PKA under conditions required to reach maximal phosphorylation. After SDS/PAGE and in-gel trypsin digestion, the samples were analysed by tandem MS and SRM. Only the single-phosphorylated tryptic peptide containing Ser⁷ was detected in AMPK- or PKA-phosphorylated GYS1 and GYS2 by SRM analysis. Tandem MS analysis of the peptide identified Ser⁷ as the residue phosphorylated by AMPK and PKA in the GYS isoenzyme preparations. Moreover, by SRM, only the singly phosphorylated peptide was detected with no indication of phosphorylation at Ser¹⁰. Immunoblotting of recombinant GYS1 and GYS2 phosphorylated by PKA and AMPK with a phospho-specific antibody raised against phosphorylated Ser⁷ for each

isoenzyme confirmed that this site was phosphorylated by the two kinases (Figures 1D and 1E). Taken together, the results demonstrate that AMPK phosphorylates GYS2 at Ser⁷ *in vitro*.

Effect of phosphorylation by AMPK and PKA on the kinetic properties of GYS1 and GYS2

The recombinant GYS1 and GYS2 preparations were phosphorylated with non-radioactive ATP and either AMPK or PKA under conditions required to reach maximal phosphorylation. GYS was then assayed in the presence or absence of 10 mM Glc-6-P, a positive allosteric effector of GYS. The ratio of GYS activity measured with and without Glc-6-P provides an activity ratio to assess GYS inactivation induced by phosphorylation. After phosphorylation by AMPK or PKA, GYS1 was inactivated compared with non-phosphorylated control GYS1 assayed at low UDP-Glc concentrations (Figure 2A), but the effect was opposed at high UDP-Glc. AMPK- and PKA-induced GYS1 inactivation was also reflected by a decreased sensitivity towards stimulation of activity by Glc-6-P when assayed at low UDP-Glc (Figure 2B). In addition, at low UDP-Glc concentrations (0.05–0.5 mM) within the physiological range, a significant decrease (61 to 27% and 64 to 32%) in GYS2 activity ratio was observed after phosphorylation by AMPK or PKA respectively, which was counteracted by high UDP-Glc concentrations (2.5 and 4.6 mM) (Figure 2C). In addition, AMPK- and PKA-induced phosphorylation decreased the affinity of GYS2 for Glc-6-P when measured at low UDP-Glc (Figure 2D).

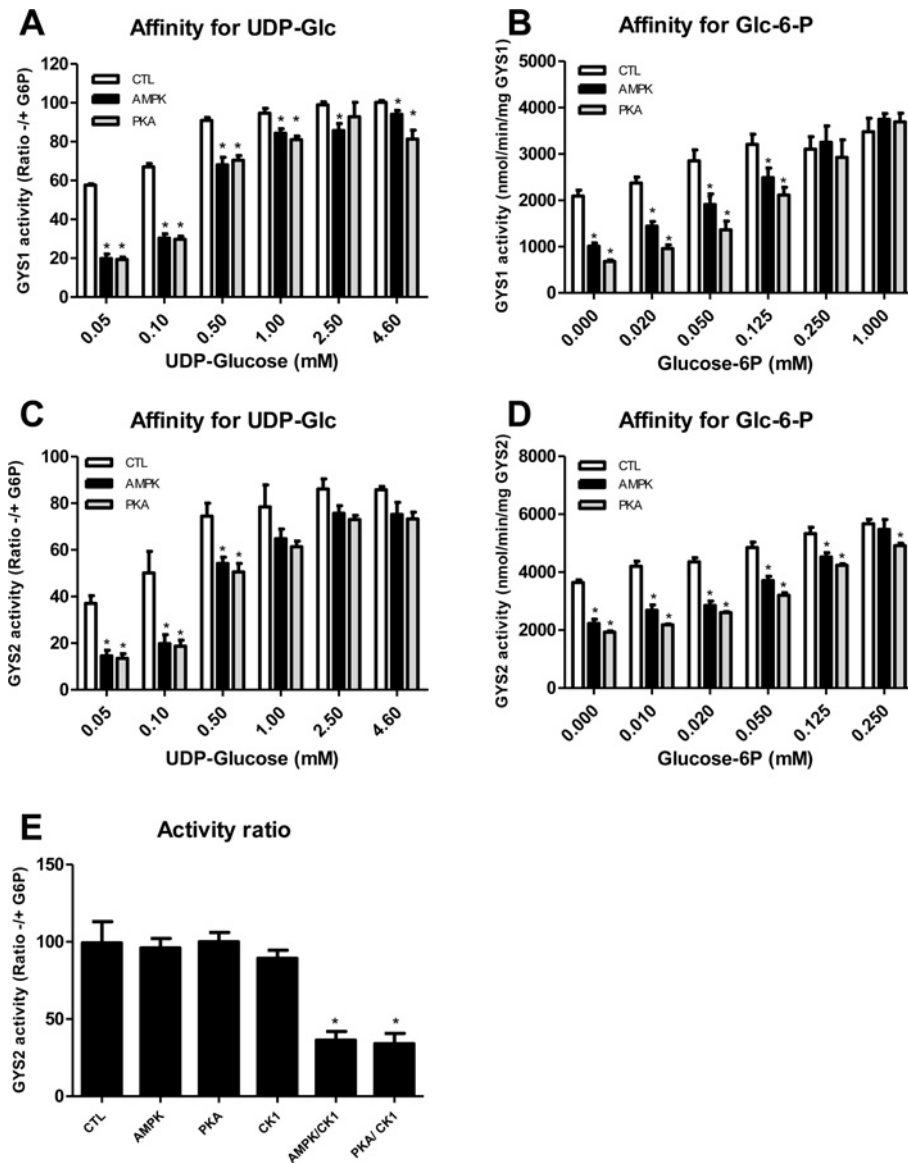


Figure 2 Effects of phosphorylation by AMPK and PKA on GYS1 and GYS2 activity as a function of UDP-Glc or Glc-6-P concentration

GYS1 (A) or GYS2 (C) activity ratio of non-phosphorylated GYS (control, CTL) or GYS phosphorylated by AMPK or PKA measured in the presence or absence of 10 mM Glc-6-P with 0.05, 0.1, 0.5, 1, 2.5 and 4.6 mM UDP-glucose. GYS1 (B) or GYS2 (D) activity of non-phosphorylated enzyme (CTL) or enzyme phosphorylated by AMPK or PKA was measured with variable Glc-6-P concentrations and 0.1 mM or 0.25 mM UDP-Glc respectively, as described in the Experimental section. (E) GYS activity ratio of non-phosphorylated GYS2 (CTL) or GYS2 phosphorylated by AMPK, PKA, CK1 or in combination as indicated and measured with 2.5 mM UDP-Glc. Results are means \pm S.E.M. for three independent experiments. * $P < 0.05$ compared with the control values (paired two-sided Student's t test).

GYS1 or GYS2 phosphorylation by AMPK or PKA had no significant effect on V_{max} in the presence or absence of Glc-6-P (Table 1). In addition, there was no significant effect of phosphorylation of the GYS isoenzymes by AMPK or PKA on the K_m for UDP-Glc measured in the presence of Glc-6-P (Table 1). However, when GYS2 was assayed in the absence of Glc-6-P, phosphorylation by AMPK or PKA increased the K_m for UDP-Glc from 0.39 mM to 0.70 and 0.88 mM respectively, and similar effects of phosphorylation were seen for GYS1. Phosphorylation of GYS1 and GYS2 by AMPK and PKA increased the K_a for Glc-6-P when measured at low UDP-Glc (Table 1). Thus phosphorylation of both GYS1 and GYS2 by AMPK or PKA leads to GYS inactivation by decreasing the

affinity for both UDP-Glc (in the absence of Glc-6-P) and Glc-6-P (at low UDP-Glc concentrations). After phosphorylation by combinations of PKA/CK1 and AMPK/CK1 for GYS assay under identical conditions with those used to measure GYS in intact cells with 2.5 mM UDP-Glc, a 60% decrease in GYS2 activity ratio was observed (Figure 2E).

AMPK activation leads to Ser⁷ phosphorylation and inactivation of GYS in freshly isolated rat hepatocytes

Freshly isolated hepatocytes from overnight-starved rats were incubated with two different pharmacological AMPK activators (A769662 or AICA riboside) and with glucagon, which

Table 1 Effects of phosphorylation by AMPK and PKA on the kinetic properties of recombinant GYS1 and GYS2

Summary of the values of K_m for UDP-Glc and V_{max} measured with or without 10 mM Glc-6-P and K_a for Glc-6-P after curve-fitting of the data shown in Figure 2. Results are means \pm S.E.M. for three independent experiments. * $P < 0.05$ compared with the control values (paired two-sided Student's t test).

(a) K_m for UDP-Glc

	Glc-6-P	K_m (mM)		V_{max} (μ mol/min per mg of protein)	
		GYS1	GYS2	GYS1	GYS2
Control	–	0.32 \pm 0.04	0.39 \pm 0.05	8.62 \pm 0.28	10.07 \pm 0.34
	+	0.15 \pm 0.02	0.15 \pm 0.02	8.04 \pm 0.26	10.76 \pm 0.32
AMPK	–	0.65 \pm 0.14*	0.70 \pm 0.10*	8.54 \pm 0.56	10.69 \pm 0.47
	+	0.13 \pm 0.03	0.18 \pm 0.02	7.77 \pm 0.34	12.10 \pm 0.26
PKA	–	0.66 \pm 0.09*	0.88 \pm 0.20*	8.75 \pm 0.34	10.15 \pm 0.76
	+	0.19 \pm 0.05*	0.20 \pm 0.04	8.57 \pm 0.46	11.24 \pm 0.53

(b) K_a for Glc-6-P

	K_a for Glc-6-P (mM)	
	GYS1	GYS2
Control	0.07 \pm 0.03	0.04 \pm 0.01
AMPK	0.13 \pm 0.03*	0.12 \pm 0.03*
PKA	0.19 \pm 0.03*	0.12 \pm 0.01*

inactivates liver GYS via PKA-induced phosphorylation. AMPK activation was assessed by measuring AMPK activity, AMPK Thr¹⁷² phosphorylation and phosphorylation of ACC Ser⁷⁹, its best known downstream target. Treatment with AICA riboside induced an increase in AMPK activity, AMPK Thr¹⁷² phosphorylation and ACC Ser⁷⁹ phosphorylation, which persisted for up to 30 min of incubation (Figures 3A, 3D and 3E). Treatment with A769662 caused no significant increase in AMPK activity (Figure 3A), but transiently increased AMPK Thr¹⁷² (Figure 3D) and downstream ACC (Figure 3E) phosphorylation. Although glucagon has been shown to activate the AMPK pathway in liver by lowering energy levels [41], in our acute/short-term incubations of hepatocytes the hormone had no significant effect either on AMPK activity or on AMPK and ACC phosphorylation (Figures 3A, 3D and 3E). Glucagon treatment, however, induced a rapid and transient decrease in GYS activity, which correlated with GYS2 Ser⁷ phosphorylation (Figures 3B and 3F). The transient effect of glucagon on GYS activity can be explained by the fact that in the presence of high glucose (30 mM in the present study), following PKA-induced GP phosphorylation/activation by glucagon, GP becomes progressively inactivated (Figures 3C and 3G), relieving inhibition of GYS phosphatase, which can then dephosphorylate GYS to return its activity to basal (Figure 3B). In sharp contrast with glucagon, pharmacological AMPK activation induced a slower, but persistent, inactivation of GYS (Figure 3B), reflected by an increase in Ser⁷ GYS2 phosphorylation (Figure 3F). Interestingly, incubation of hepatocytes with AICA riboside resulted in a slow increase in GP activity and Ser¹⁴ phosphorylation, but these effects were not seen on incubation with A769662 (Figures 3C and 3G). Taken together, the results suggest that pharmacological AMPK activation leads to GYS inactivation via GYS2 Ser⁷ phosphorylation in hepatocytes, as previously shown for GYS1 in skeletal muscle [18].

Effects of pharmacological AMPK activators on GYS inactivation in freshly isolated hepatocytes prepared from liver-specific AMPK $\alpha 1/\alpha 2$ -deficient mice

As observed in rat hepatocytes, AMPK Thr¹⁷² phosphorylation increased in WT mouse hepatocytes incubated with AICA

riboside, whereas incubation with A769662 or glucagon had no significant effect (Figures 4A and 4E). In hepatocytes from WT mice, we also confirmed that treatment with AICA riboside or A769662 increased ACC Ser⁷⁹ phosphorylation, whereas incubation with glucagon had no effect on phosphorylation at this target site (Figures 4B and 4E). In hepatocytes from WT mice incubated with AICA riboside, A769662 or glucagon, GYS activity decreased rapidly, reaching ~40%, 50% and 90% inactivation after 15 min of incubation respectively (Figure 4F). Thus GYS inactivation by the various treatments in rat and WT mouse hepatocytes was greater than that seen *in vitro* following AMPK- and PKA-induced phosphorylation of recombinant GYS2 (Figures 2C and 2D), but was similar to the extent of recombinant GYS inactivation seen after phosphorylation by AMPK or PKA together with CK1 (Figure 2E). Interestingly, in contrast with glucagon treatment of rat hepatocytes, where transient changes in GYS activity and GYS2 Ser⁷ phosphorylation were observed (Figures 3B and 3F), GYS inactivation in response to glucagon treatment of WT mouse hepatocytes was maintained over the 30 min incubation period (Figure 4F). In addition, in WT mouse hepatocytes, incubation with both glucagon and AICA riboside increased GP Ser¹⁴ phosphorylation (Figure 4C), although the increase with glucagon was not transient, as observed in rat hepatocytes (compare Figures 4C and 3G).

In hepatocyte extracts from AMPK $\alpha 1\alpha 2$ LS^{-/-} mice, basal GYS activity was almost 2-fold higher (831 \pm 72 m-units/g of cells) than in extracts from WT mice (456 \pm 28 m-units/g of cells) (Figures 4F and 4G). In hepatocyte extracts from AMPK $\alpha 1\alpha 2$ LS^{-/-} mice incubated with AICA riboside or A769662, no signals for AMPK Thr¹⁷² phosphorylation, ACC Ser⁷⁹ phosphorylation or total $\alpha 1/\alpha 2$ AMPK by immunoblotting were detected (Figure 4E). In addition, AMPK activity in hepatocytes from $\alpha 1\alpha 2$ LS-KO (knockout) mice was substantially reduced and, compared with hepatocytes from WT mice, was not increased by AICAR treatment (Supplementary Figure S1 at <http://www.BiochemJ.org/bj/443/bj4430193add.htm>). Glucagon treatment of hepatocytes from AMPK $\alpha 1\alpha 2$ LS^{-/-} mice induced a rapid time-dependent decrease in GYS activity to an extent similar to that observed in hepatocytes from WT mice (Figures 4F and 4G). In hepatocytes from AMPK $\alpha 1\alpha 2$ LS^{-/-}

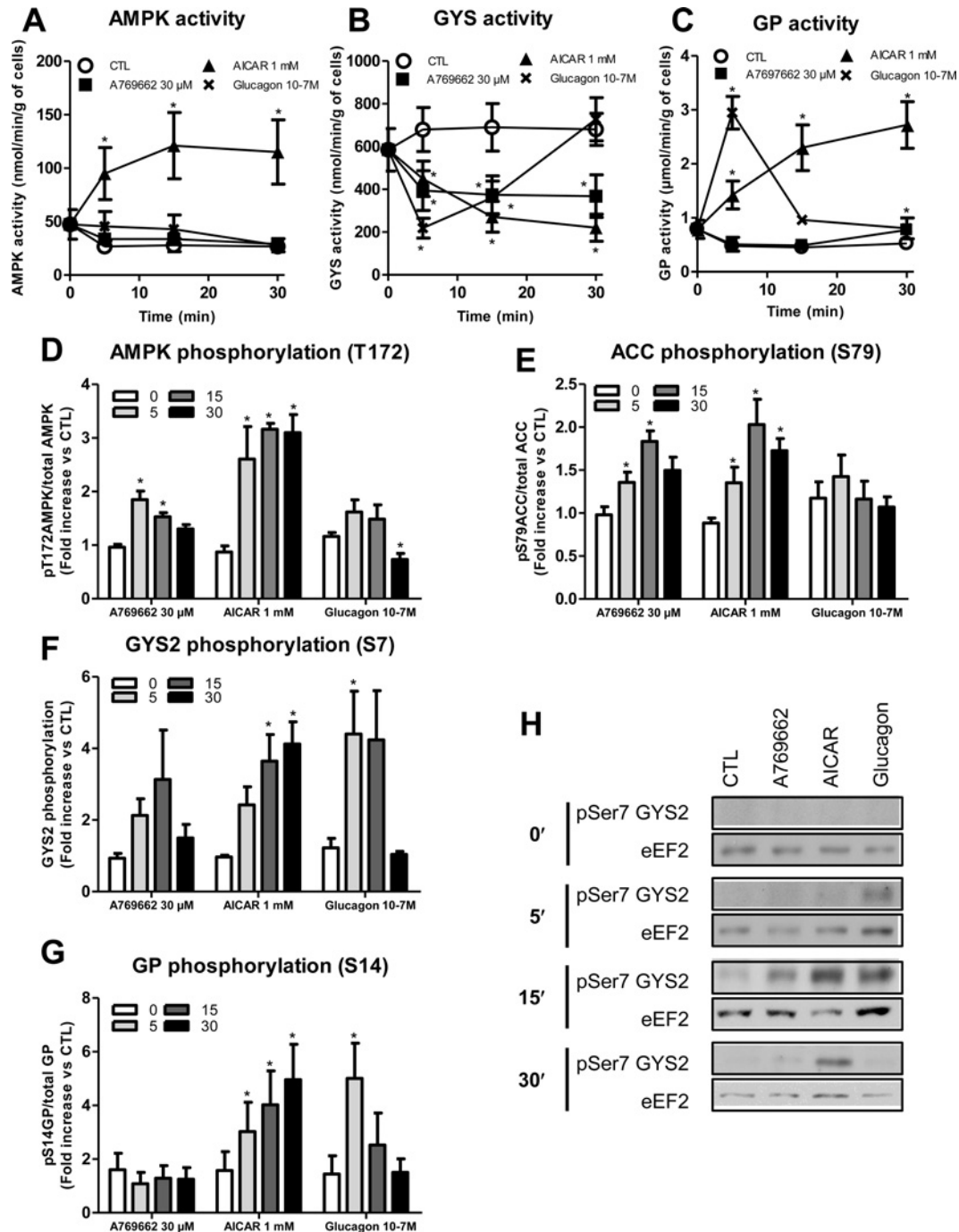


Figure 3 Time course of the effects of AICA riboside, A769662 and glucagon on AMPK activity, AMPK Thr¹⁷², ACC Ser⁷⁹ and GYS2 Ser⁷ phosphorylation in incubated rat hepatocytes

Freshly isolated rat hepatocytes were incubated for 30 min with 30 μ M A769662, 1 mM AICA riboside or 10⁻⁷ M glucagon compared with vehicle control (CTL). At the indicated times, the cells were freeze-stopped and extracts were prepared for the measurement of AMPK (A), GYS (B) and GP (C) activity and for immunoblotting AMPK Thr¹⁷² (D), ACC Ser⁷⁹ (E), GYS2 Ser⁷ (F) and GP Ser¹⁴ (G) phosphorylation. In (D and E) the histograms show the quantification of immunoblots by OdysseyTM imaging after ratios of the signals obtained with the anti-phospho compared with anti-(total protein) antibodies had been calculated. In (F and G) pGYS2 Ser⁷ and pGP Ser¹⁴ phosphorylation were expressed as fold increases after immunoblot detection by chemiluminescence. In (H) a representative immunoblot for pGYS2 Ser⁷ phosphorylation compared with eEF2 as a loading control is shown. Results are means \pm S.E.M. for four independent experiments. **P* < 0.05 compared with the control values (paired two-sided Student's *t* test).

mice, GYS inactivation by AICA riboside was considerably reduced compared with hepatocytes from WT mice, whereas in response to A769662, GYS2 was inactivated by approximately 60% in spite of the loss of the AMPK catalytic subunits (Figures 4F and 4G). In hepatocytes from both WT and

AMPK $_{\alpha 1\alpha 2LS}^{-/-}$ mice incubated with glucagon or AICA riboside, GP Ser¹⁴ phosphorylation increased, but did not change in response to A769662 (Figures 4C and 4D). Unfortunately, immunoblotting of hepatocyte extracts from AMPK $_{\alpha 1\alpha 2LS}^{-/-}$ mice for GYS2 Ser⁷ phosphorylation was inconclusive because of

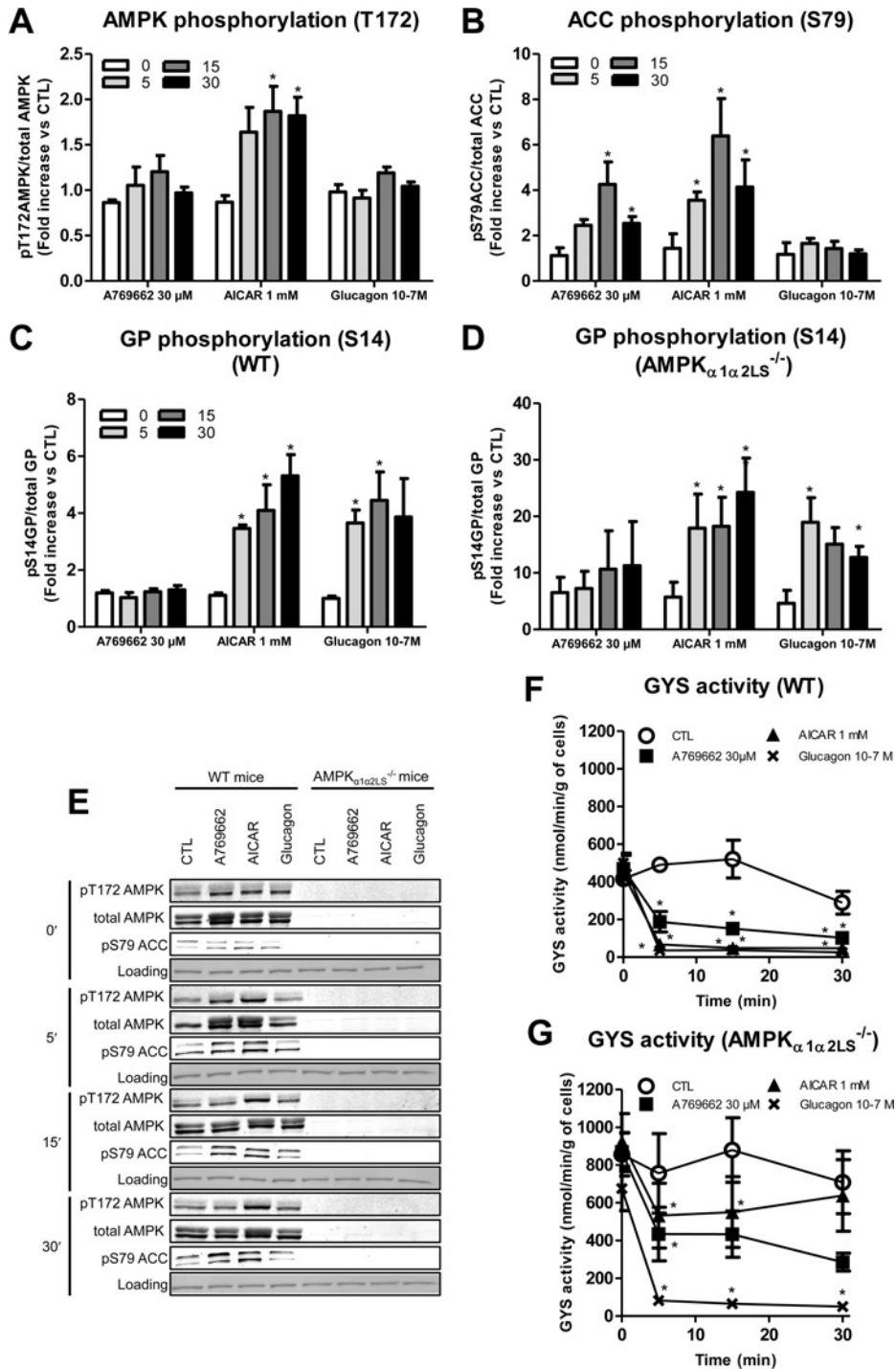


Figure 4 Time-course of the effects of AICA riboside, A769662 and glucagon on AMPK Thr¹⁷², ACC Ser⁷⁹ phosphorylation and GYS activity in incubations of hepatocytes from WT and $AMPK_{\alpha 1\alpha 2LS}^{-/-}$ mice

Freshly isolated hepatocytes from WT and $AMPK_{\alpha 1\alpha 2LS}^{-/-}$ mice were incubated for 30 min with 30 μ M A769662, 1 mM AICA riboside or 10⁻⁷ M glucagon compared with vehicle control (CTL). At the indicated times, the cells were freeze-stopped and extracts were prepared for immunoblotting AMPK Thr¹⁷² (A), ACC Ser⁷⁹ (B) and GP Ser¹⁴ (C) phosphorylation from WT hepatocyte incubations and GP Ser¹⁴ (D) phosphorylation from $AMPK_{\alpha 1\alpha 2LS}^{-/-}$ hepatocyte incubations, as described in the legend to Figure 3. (E) A representative immunoblot of AMPK Thr¹⁷², ACC Ser⁷⁹ phosphorylation and total AMPK ($\alpha 1/\alpha 2$) with pyruvate carboxylase revealed by the use of streptavidin-coupled horseradish peroxidase as a loading control. GYS activities in extracts from WT (F) and $AMPK_{\alpha 1\alpha 2LS}^{-/-}$ (G) hepatocyte incubations are also shown. Results are means \pm S.E.M. for three (F and G) or four (A–D) independent experiments. * $P < 0.05$ compared with the control values (paired two-sided Student's t test).

limited amounts of material available from only three separate experiments.

DISCUSSION

In the present study we show that both AMPK and PKA phosphorylate recombinant GYS2 to the same maximal stoichiometry and at a single site identified by MS as Ser⁷. GYS2 phosphorylation by AMPK or PKA led to enzyme inactivation by decreasing the affinity of the enzyme for both UDP-Glc (in the absence of Glc-6-P) and Glc-6-P (at low UDP-Glc concentrations) without affecting V_{\max} . Very similar effects of AMPK- and PKA-induced phosphorylation on the kinetic properties of GYS1 were observed. In skeletal muscle, AMPK activation via AICA riboside treatment results in GYS inactivation, which can largely be overcome by the rise in Glc-6-P due to the stimulation of glucose transport [6], explaining why glycogen synthesis can increase as a result of acute or repeated AICA riboside injection in rodents [42–44]. Phosphorylation of GYS1 at site 2 is a priming event for phosphorylation at site 2a by CK1, which inactivates the enzyme to a greater extent than seen after phosphorylation by PKA alone [10]. For GYS2, we observed that phosphorylation by both AMPK and CK1 led to inactivation, even when assayed with high concentrations of UDP-Glc (Figure 2E). This explains why GYS activity was substantially decreased in hepatocytes subjected to the various treatments, presumably due to phosphorylation at both sites 2 and 2a.

In freshly isolated rat hepatocytes, AMPK activation was seen on incubation with both AICA riboside and A769662, as reflected by increases in both AMPK Thr¹⁷² and ACC Ser⁷⁹ phosphorylation (Figures 3D and 3E). However, A769662 treatment did not increase AMPK activity as measured using a peptide-based assay (Figure 3A), which is in contrast with previous reports [45–47]. A769662 can both allosterically stimulate AMPK and promote AMPK α -subunit Thr¹⁷² phosphorylation [45,48], but the allosteric effect is lost during AMPK assay of poly(ethylene glycol) fractions from cell extracts. However the increase in downstream ACC phosphorylation is testimony to allosteric stimulation of AMPK by A769662 treatment in the present study (Figure 3E). Although A769662 caused a very modest, but significant, increase in Thr¹⁷² phosphorylation, our AMPK assay might not have been sensitive enough to detect this small increase. There are other reports in the literature showing that, in hepatocytes incubated with A769662, ACC phosphorylation was increased without a rise in AMPK α -subunit Thr¹⁷² phosphorylation [24,48]. The differences between these results and those previously reported [45–47] might be due to the use of different cell models (hepatocytes compared with HEK-293, CCL13 cells and mouse embryonic fibroblasts), A769662 concentrations or duration of treatment.

AMPK activation by treatment with A769662 or AICA riboside induced persistent GYS2 inactivation and Ser⁷ phosphorylation. Incubation with glucagon induced transient GYS2 inactivation and Ser⁷ phosphorylation due to activation of GYS phosphatase in the presence of glucose following inactivation of GP. This inverse relationship between the active forms of GP and GYS in liver was elegantly demonstrated many years ago following glucose administration to mice [49]. The fact that pharmacological AMPK activators induced a persistent inactivation of GYS2 indicates that AMPK might rapidly negatively regulate the activity of PP1c complexes involved in the control of glycogen metabolism. It is noteworthy that, in the long term, AMPK phosphorylates the R5/PTG (protein-targeting glycogen) subunit of the R5/PTG–

PP1c complex to accelerate its degradation via the malin–laforin complex [50].

Treatment of freshly isolated hepatocytes from WT mice with A769662, AICA riboside or glucagon led to GYS2 inactivation. However, in mouse hepatocytes incubated with glucagon, GYS inactivation was rapid, but not transient, and was sustained during the course of incubation, presumably because PP1 regulation is different in mouse and rat. In hepatocytes from AMPK $_{\alpha1\alpha2LS}^{-/-}$ mice, GYS activities were increased ~2-fold compared with activities in WT mice, suggesting that AMPK may have a negative effect on GYS2 gene transcription or translation. Indeed increased levels in GYS2 protein in hepatocyte extracts from AMPK $_{\alpha1\alpha2LS}^{-/-}$ mice were seen by immunoblotting (results not shown). It is noteworthy that in peroxisome-deficient hepatocytes where AMPK became activated, GYS expression decreased along with glycogen synthesis [51]. GYS inactivation by AICA riboside treatment was severely blunted in hepatocytes from AMPK $_{\alpha1\alpha2LS}^{-/-}$ mice (Figure 4G) compared with WT mice (Figure 4F). However, following treatment with A769662, there was still 50–60% inactivation of GYS2 in hepatocytes from AMPK $_{\alpha1\alpha2LS}^{-/-}$ mice (Figure 4G). It is now becoming clear, though, that A769662 has AMPK-independent effects, such as the inhibition of glucose production in hepatocytes from $\beta1$ -KO [46] and $\alpha1\alpha2$ LS-KO mice [24].

On the basis of our findings, we propose that AMPK activation in hepatocytes leads to GYS inactivation mediated by Ser⁷ phosphorylation. AMPK activation in liver during exercise would thus be expected to decrease glycogen synthesis and favour glycolysis to maintain ATP levels and reduce gluconeogenesis. AMPK has been shown to bind glycogen via the glycogen-binding domain of its β -subunits [52]. Moreover, branched oligosaccharides and limit dextrans, characteristic of highly degraded glycogen, inhibit AMPK activity [53] such that GYS2 would be active to replenish glycogen reserves following liver glycogen depletion during exercise. In addition, AMPK would bind glycogen via its β -subunit glycogen-binding domain to target it to GYS2 for control by phosphorylation under the appropriate conditions. Transgenic mouse models with knock-in mutations (GYS2 S7E and S7E/S10E, and mutations to abolish allosteric stimulation by Glc-6-P) would be useful to further delineate the role of control of GYS2 by AMPK in liver glycogen metabolism.

AUTHOR CONTRIBUTION

Laurent Bultot carried out most of the experimental work in the Brussels and Dundee laboratories. Bruno Guigas and Liliane Maisin performed mouse and rat hepatocyte incubations. Monique Beullens provided reagents and advice for dephosphorylation by recombinant PP1 catalytic subunits. Alexander Von Wilamowitz-Moellendorff expressed and purified recombinant GYS1/GYS2 and also conducted initial dephosphorylation/phosphorylation trials on recombinant GYS1/GYS2 *in vitro* for GYS assay. Nusrat Hussain carried out immunoblotting experiments and GYS kinetic studies. Didier Vertommen analysed GYS phosphorylation by MS. Joan Guinovart, Marc Foretz, Benoît Viollet and Kei Sakamoto provided essential reagents and advice. Louis Hue participated in the conception and design of the study, and analysis and interpretation of the data, together with Mark Rider who wrote the paper.

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SUPPLEMENTARY ONLINE DATA

AMP-activated protein kinase phosphorylates and inactivates liver glycogen synthase

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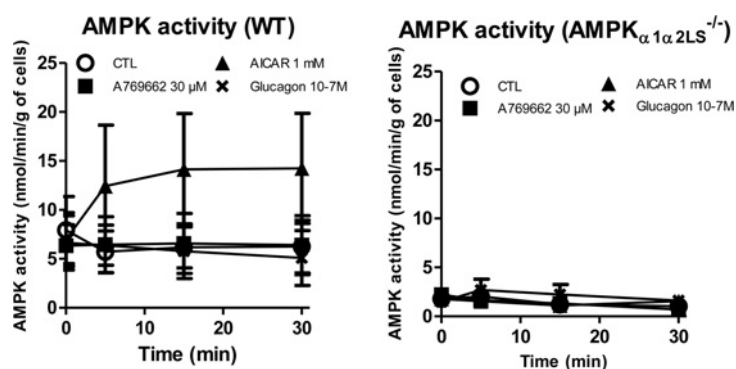


Figure S1 Time course of the effects of AICA riboside, A769662 and glucagon on AMPK activity in incubations of hepatocytes from WT and AMPK $\alpha_{1\alpha 2LS}^{-/-}$ mice

Freshly isolated hepatocytes from WT and AMPK $\alpha_{1\alpha 2LS}^{-/-}$ mice were incubated for 30 min with 30 μ M A769662, 1 mM AICA riboside or 10⁻⁷ M glucagon compared with vehicle control (CTL). At the indicated times, the cells were freeze-stopped and extracts were prepared for the measurement of AMPK activity. Results are means \pm S.E.M. for three independent experiments. Compared with hepatocytes from WT mice at zero time, AMPK activity in hepatocytes from AMPK $\alpha_{1\alpha 2LS}^{-/-}$ mice was significantly reduced [6.65 \pm 1.28 compared with 1.82 \pm 0.17 nmol/min per g of cells, $P < 0.05$ (paired two-sided Student's t test)].

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