Ca²⁺ Uptake Coupled to Glycogen Phosphorolysis in the Glycogenolytic-Sarcoplasmic Reticulum Complex from Rat Skeletal Muscle

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The glycogenolytic-sarcoplasmic reticulum complex from rat skeletal muscle accumulates Ca^{2+} upon stimulation of glycogen phosphorolysis in the absence of added ATP. It is shown that an efficient Ca^{2+} uptake involves the sequential action of glycogen phosphorylase, phosphoglucomutase and hexokinase, which generate low concentrations of ATP (approximately $1-2~\mu M$) compartmentalized in the immediate vicinity of the sarcoplasmic reticulum Ca^{2+} , Mg^{2+} -ATPase (the Ca^{2+} pump). The Ca^{2+} uptake supported by glycogenolysis in this subcellular structure is strongly stimulated by micromolar concentrations of AMP, showing that the glycogen phosphorylase associated with this complex is in the dephosphorylated b form. The results point out that the flux through this compartmentalized metabolic pathway should be enhanced in physiological conditions leading to increased AMP concentrations in the sarcoplasm, such as long-lasting contractions and in ischemic muscle.

Introduction

Glycogen particles are intimatelly associated with sarcoplasmic reticulum (SR)¹ in rabbit skeletal muscle (Wanson and Drochmans, 1972, Entman *et al.*, 1980). Subcellular fragmentation allows for preparation of glycogenolytic sarcoplasmic reticulum complexes (Entman *et al.*, 1980). Moreover, SR membrane preparations from rabbit skeletal muscle contain large amounts of phosphorylase in its dephosphorylated form (**b** form), and also glycogen and several enzymes di-

Abbreviations: $K_{0.5}(AMP)$, concentration of AMP that produced 50% of maximum activation; Ap_5A , P^1P^5 -di-(adenosine-5') pentaphosphate; Ca^{2+},Mg^{2+} -ATPase, Ca^{2+} and Mg^{2+} dependent adenosine triphosphatase (EC 3.6.1.38); EGTA, ethylene glycol bis(β-aminoethyl ether)-N,N'-tetraacetic acid; GL-SR, glycogenolytic-sarcoplasmic reticulum complex; hexokinase, ATP: D-hexose 6-phosphotransferase (EC 2.7.1.1); IU, amount of enzyme that released 1 μmol product per min at saturating substrate concentration; PGM, phosphoglucomutase (α -D-glucose-1,6 phosphomutase, EC 5.4.2.2); phosphorylase, glycogen phosphorylase (EC 2.4.1.1); SR, sarcoplasmic reticulum; Tes, 2-[(2-hydroxy-1,1 bis-(hydroxymethyl) ethylamine]; Tris, tris (hydroxymethyl) aminomethane.

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rectly involved in energy metabolism in muscle, such as myokinase, creatine kinase and AMP deaminase (Cuenda *et al.* 1994). Early studies with GL-SR have shown that this association allows to trigger a faster glycogen phosphorolysis at the onset of skeletal muscle contraction (Entman *et al.*, 1980).

Since the phosphorylated phosphorylase (a form) has low affinity for binding to SR membranes (Cuenda et al., 1991), GL-SR acts as a phosphorylase b reservoir in skeletal muscle cells due to the large SR membrane network of these cells (Cuenda et al., 1995). The activity of phosphorylase b is dependent of AMP and, under physiological conditions which raise the AMP levels in skeletal muscle sarcoplasm such as anoxia and during contraction (Busby and Radda, 1976, Newsholme and Leech, 1983) glycogen phosphorolysis in the vicinity of SR membranes should be stimulated.

Using purified SR membranes and purified gly-cogen phosphorylase *b* from rabbit skeletal muscle we have shown that this association can form *in vitro* a metabolic shuttle by coupling glycogenolysis to support Ca²⁺ uptake through production of ATP in the vicinity of the SR Ca²⁺,Mg²⁺-ATPase (Cuenda *et al.*, 1993). This metabolic pathway can be written as follows.

$$(Glycogen)_{n-1}$$

$$(Glycogen)_n + Pi \xrightarrow{1} Glc \ 1-P \xrightarrow{2}$$

$$ADP \quad Glucose$$

$$Glc \ 6-P \xrightarrow{3} ATP \xrightarrow{4} Ca^{2+} uptake$$

where Glc 1-P and Glc 6-P are glucose-1-phosphate and glucose-6-phosphate, respectively; and 1-4 are the enzymes phosphorylase, PGM, hexokinase and Ca²⁺,Mg²⁺-ATPase, respectively. As noted in (Cuenda et al., 1993), a major bioenergetic advantage of this compartmentalized metabolic pathway is the efficient use in the cell of low concentrations of high energy phosphate in the form of ATP generated from low energy phosphate compounds, such as Glc 6-P. The flux through this metabolic pathway will be blocked upon activation of the glycogen phosphorylase b kinase at the onset of muscle contraction, because the phosphorylated a form of glycogen phosphorylase has low affinity for binding to SR membranes (Cuenda et al., 1991).

In this communication we report experimental data showing that our previous results in rabbit skeletal muscle can be extended to rat skeletal muscle where a GL-SR subcellular structure is also present. Using the isolated GL-SR from rat skeletal muscle we show that this metabolic shuttle should operate more efficiently in physiological conditions where the AMP concentration raises in the micromolar range, such as ischemic muscle and in tetanic or long lasting contractions.

Materials and Methods

GL-SR has been prepared from skeletal muscle of male Fischer rats, following the protocol of Entman *et al.* (1980). Phosphorylase of rat skeletal muscle was prepared as described by Gutierrez-Merino *et al.* (1980). The protein concentration was determined using the method of Lowry (Lowry *et al.*, 1951), with bovine serum albumin as standard.

The glycogen content of GL-SR preparations has been measured following the phenol sulfuric

method (Dubois *et al.*, 1956), using glycogen as standard, and was found to be on average 25 ± 5 µg glycogen per mg of protein (n > 10).

Ca²⁺ uptake by GL-SR membranes was measured with the use of ⁴⁵Ca²⁺ and Millipore HA filters as indicated elsewhere (Cuenda *et al.*, 1993). All the data reported in this paper are the average of triplicate experiments carried out with, at least, two different GL-SR preparations.

Enzyme assays

The ATPase activity was measured using the coupled enzyme system pyruvate kinase/lactate dehydrogenase as indicated in Cuenda *et al.* (1990), with the following assay medium: 100 mm Tes (pH 7.0)/ 100 mm KCl/ 20% v/v glycerol/ 5 mm MgCl₂/ 2.5 mm ATP/ 0.1 mm CaCl₂/ 0.25 mm NADH/ 0.42 mm phospho*enol*pyruvate/ 7.5 IU pyruvate kinase/ 18 IU lactate dehydrogenase/ 5 mm NaN₃/ 0.4 μg/ml calcimycin and 10 μg GL-SR protein·ml⁻¹. The Ca²⁺-dependent ATPase activity of different preparations (*n*>10) of the rat GL-SR complex at 25 °C ranged from 0.85 to 1.45 μmol ATP hydrolyzed per min per mg protein.

Unless stated otherwise, the phosphorylase activity had been measured in the direction of glycogen degradation as indicated in Cuenda *et al.* (1991), with the following assay medium: 50 mm imidazole (pH 6.9)/ 10 mm magnesium acetate/ 12 mm KH₂PO₄/ 0.63 mm NADP+/ 0.45 g·l⁻¹ glycogen/ 50 µm glucose-1,6-diphosphate/ 5 IU PGM/ 5 IU glucose-6-phosphate dehydrogenase and 100 µg GL-SR protein·ml⁻¹. The phosphorylase activity in the presence of 1 mm AMP of different preparations of the rat GL-SR (n > 10) at 30 °C ranged from 0.12 to 0.72 µmol glucose-1-phosphate produced per min per mg protein.

The PGM and hexokinase activities were measured at 35 °C using the coupled enzyme glucose 6-phosphate dehydrogenase with the following assay mixture: 100 mm Tes-Tris (pH 7.0)/ 50 mm KCl/ 10 mm MgCl₂/ 20 mm H₂PO₄K/ 50 μm CaCl₂/ 0.48 mm NADP+/ 1 IU glucose 6-phosphate dehydrogenase, and 10 mm glucose 1-phosphate/ 50 μm glucose 1,6-diphosphate for PGM activity measurements or 3 mm glucose/ 2 mm ATP for the hexokinase activity assay medium. Variations in NADP+ absorbance were recorded at 340 nm.

The AMP content of ADP solutions had been determined using the coupled enzyme AMP-deaminase as indicated in Tovmasian *et al.* (1990), with a value for the differential extinction coefficient ($\in_{AMP} - \in_{IMP}$) at 265 nm of 8.2×10^3 M⁻¹ cm⁻¹.

Chemicals

Bovine serum albumin, ADP, AMP, ATP, phosphoenolpyruvate, EGTA, glucose-1,6-diphosphate, glucose 6-phosphate, Tris (TRIZMA base) and TES were obtained from Sigma. Glycogen, Ap₅A, NADH, NADP⁺, calcimycin, pyruvate kinase, lactate dehydrogenase, PGM, hexokinase and glucose 6-phosphate dehydrogenase were purchased from Boehringer Mannheim. Hexokinase and PGM were sulfate-free by centrifugation. All the other chemicals used in this study were obtained from Merck.

Results and Discussion

Our GL-SR preparations from Fischer rats skeletal muscle had a phosphorylase activity which is dependent upon the presence of AMP and it is inhibited by caffeine (Fig. 1A), therefore, indicating that the phosphorylase bound to these complexes is mainly in its b (dephosphorylated) form. This is a property shared with the GL-SR complex from rabbit skeletal muscle (Entman et al., 1980). The GL-SR and the SR from rabbit skeletal muscle also had myokinase activity (Entman et al., 1980; Cuenda et al., 1994), which produces ATP from AMP and ADP, and this would interfere with the assays of Ca²⁺ uptake supported by glycogen phosphorolysis in the absence of added ATP. We, thus, confirmed the presence of myokinase activity of the GL-SR complex from rat skeletal muscle, which on average was found to be $0.14 \pm 0.05 \,\mu\text{mol}$ ADP per min per mg GL-SR protein. Because the SR Ca²⁺ pump is highly efficient even in the pres-

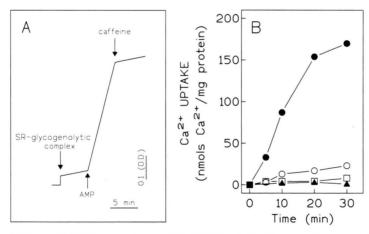


Fig. 1. Phosphorylase activity and Ca^{2+} uptake by rat GL-SR. Panel A. Phosphorylase activity of the rat GL-SR, monitored by the production of NADPH with the coupled enzyme system PGM/glucose-6-phosphate dehydrogenase. The phosphorylase activity was measured at 35 °C in the following assay medium: 50 mm Tes Tris (pH 7.0)/ 10 mM MgCl₂/ 20 mm Pi/ 60 mm KCl/ 40 μ M CaCl₂/ 5 mm NaN₃/ 0.5 mm NADP+/ 3.6 g· $^{-1}$ glycogen/ 0.1 mm ADP/ 50 μ M glucose-1,6-diphosphate/ 32 IU PGM/ 10 IU glucose-6 phosphate dehydrogenase. The different arrows show the addition of 0.1 mg GL-SR protein/ml, 1 mm AMP and 10 mm caffeine. The absorbance (O.D.) was measured at 340 nm, in quartz cuvettes of 1 cm light-path. The calibration bar of 0.1 O.D. corresponds to 16.08 nmols/ml of glucose-1-phosphate, as one mol of NADPH is produced per mol of glucose-1-phosphate and the standard molar absorption coefficient of NADPH is 6220 M $^{-1}$ cm $^{-1}$.

Panel B. Effects of caffeine, glucose and glucose-1,6-diphosphate on Ca^{2+} uptake by rat GL-SR in the absence of added ATP. The Ca^{2+} uptake by rat GL-SR was supported by glycogen phosphorolysis and measured in the absence of ATP. The assay medium contained 50 mm Tes-Tris (pH 7.0)/ 10 mm MgCl₂/ 20 mm Pi/ 60 mm KCl/ 40 μ 4 CaCl₂ (4000 cpm.nmol⁻¹)/ 5 mm NaN₃ and 0.1 mg GL-SR protein. ml⁻¹, supplemented with 3.6 g·⁻¹ glycogen/ 1 mm AMP/ 0.1 mm ADP/ 50 μ Ap₅A/ 50 μ glucose-1,6-diphosphate/ 0.14 mg.ml⁻¹ (28 IU) PGM, 0.13 mg.ml⁻¹ (18 IU) hexokinase. The different symbols correspond to: 1 mm AMP (\bullet), 1 mm AMP in the absence of glucose 1,6-diphosphate (\bigcirc), 1 mm AMP and 10 mm caffeine (\square), and 1 mm AMP and 50 mm glucose (\blacktriangle).

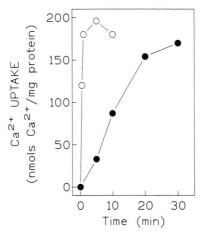


Fig. 2. Ca²⁺ uptake by rat GL-SR in the absence (\bullet) and presence (\bigcirc) of added ATP. In both experiments the assay medium contained 50 mm Tes-Tris (pH 7.0)/ 10 mm MgCl₂/ 20 mm Pi/ 60 mm KCl/ 40 μ m ⁴⁵CaCl₂ (4000 cpm·nmol⁻¹)/ 5 mm NaN₃ and 0.1 mg GL-SR protein·ml⁻¹. The Ca²⁺ uptake in control experiments was measured in the presence of 2.5 mm ATP (\bigcirc). In experiments performed in the absence of ATP (\bigcirc) the assay medium also contained: 3.6 g·l⁻¹ glycogen/ 1 mm AMP/ 0.1 mm ADP/ 50 μ m Ap₅A/ 50 μ m glucose-1,6 diphosphate/ 0.14 mg·ml⁻¹ (28 IU) PGM, 0.13 mg·ml⁻¹ (18 IU) hexokinase.

ence of micromolar ATP concentrations we added to all the assays 50 μ m of the myokinase inhibitor Ap₅A, a concentration which produced full inhibition of the myokinase activity of the rat GL-SR complex (data not shown).

When the assay medium is supplemented with the enzymes PGM, hexokinase and ADP plus the physiological PGM activator glucose 1,6-diphosphate, glycogen phosphorolysis by the phosphorylase associated with the rat GL-SR complex supports Ca²⁺ uptake in the absence of added ATP (Fig. 1B and 2). The overall activities of hexokinase and PGM added to the assay medium are much lower than those present in the sarcoplasm of rat skeletal muscle (Clark and Lardy, 1984). Inhibition of the phosphorylase activity of the GL-SR by caffeine blocks Ca²⁺ uptake (Fig. 1B). In separate control experiments, the effect of caffeine on PGM and hexokinase activities was measured. Caffeine (up to 10 mm) was found without significant effect on the activity of PGM and of hexokinase, as it produced less than 10% inhibition of their enzymatic activities when assayed under the same experimental conditions with varying concentrations of their substrates, glucose-1-phosphate (up to 10 mm) and glucose (up to 3 mm)/ ATP (up to 2 mm), respectively (results not shown). Ca²⁺ uptake shows an absolute requirement for PGM and hexokinase activities in the assay medium, such that removal of any of these enzymes results in blockade of Ca²⁺ uptake (data not shown). In the absence of the PGM activator glucose-1,6-diphosphate the rate of Ca²⁺ uptake is largely reduced (Fig. 1B). The inhibition of Ca²⁺ uptake by 50 mм glucose, which blocks the production of glucose and ATP from glucose-6-phosphate and ADP, see also Fig. 1B, demonstrated the need of the hexokinase activity to couple glycogen phosphorolysis to Ca²⁺ uptake by GL-SR. It is to be noticed that GL-SR preparations retain low amounts of hexokinase, as indicated by activity measurements (data not shown). However endogenous hexokinase activity is not able to sustain a measurable Ca²⁺ uptake. Similarly, the glycogen content of GL-SR preparations is too low to reliably measure Ca²⁺ uptake in the absence of added exogenous glycogen.

As indicated above, the activity of phosphorylase associated with GL-SR preparations is dependent upon AMP. Fig. 3 shows that Ca²⁺ uptake is stimulated by AMP, in the concentration range that stimulates the phosphorylase activity of the GL-SR (Fig. 3B), and in the range of AMP concentrations present in the cytoplasm of skeletal muscle cells, which has been reported to range between 2 and 10 µm in resting muscle (Newsholme and Leech, 1983; Yamada and Sugi, 1987), raising up to 100-200 µm during long-lasting exercise or in ischemic muscle (Newsholme and Leech, 1983). The results shown in the Fig. 3B also point out that the $K_{0.5}$ of AMP and the shape of the dependence upon AMP of Ca2+ uptake is different to that of the phosphorylase activity of GL SR. This result suggests that above a given concentration of AMP (> 0.1 mm) glycogen phosphorolysis is not the only rate-limiting step of this process. This is probably due to enhanced Ca2+ release from SR fragments by submillimolar concentrations of AMP (Meissner, 1984), that can counterbalance Ca²⁺ uptake. If this hypothesis is correct, addition of phosphorylase purified from rat skeletal muscle should produce a steeper dependence of Ca2+ uptake upon AMP concentration and higher levels of Ca²⁺ uptake at saturation by AMP. The results

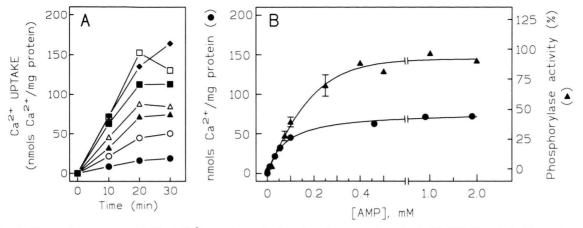


Fig. 3. Dependence upon AMP of Ca^{2+} uptake and phosphorylase activity of rat GL-SR. Panel A. Dependence upon AMP of Ca^{2+} uptake as a function of time. Ca^{2+} uptake was measured in the absence of ATP as indicated in the legend of the Fig. 2. The different concentrations of AMP (in μ M) were: 10 (\bullet), 32.5 (\circ), 55 (\blacktriangle), 100 (\Box) and 2000 (\bullet).

Panel B. Dependence upon AMP concentration of Ca^{2+} uptake and phosphorylase **b** activity. Ca^{2+} uptake as a function of AMP concentrations with 10 min of reaction (\bullet). The different values of Ca^{2+} uptake were obtained from the data of Panel A. Phosphorylase activity (\blacktriangle) had been measured as indicated in the Materials and Methods.

shown in the Fig. 4 are in good agreement with this prediction.

In addition, the results of Fig. 4 demonstrated that the higher the phosphorylase concentration, the lower the concentration of AMP required to produce significant Ca^{2+} uptake by GL-SR preparations, e.g. the lower the $K_{0.5}(AMP)$ values. This effect is not due to an altered Ca^{2+} dependence of the Ca^{2+} pump of the SR membrane in the pres-

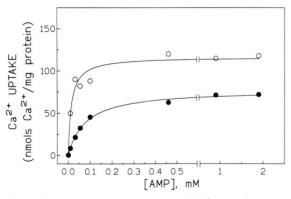


Fig. 4. Dependence upon AMP of Ca^{2+} uptake by rat GL-SR after 10 min reaction in the absence (\bullet) or presence (\bigcirc) of 0.1 mg·ml⁻¹ of phosphorylase *b* purified from rat skeletal muscle. The assay medium is indicated in the legend of Fig. 2 for Ca^{2+} uptake measurements in the absence of added ATP.

ence of phosphorylase, because the Ca²⁺ dependence of the Ca²⁺,Mg²⁺-ATPase of rat SR (devoid of phosphorylase) is not significantly different from that of the rat GL-SR (Fig. 5A). It is to be

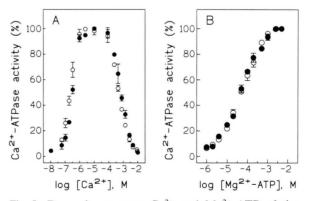


Fig. 5. Dependence upon Ca^{2+} and Mg^{2+} -ATP of the Ca^{2+} -ATPase activity of rat GL-SR. Panel A. Dependence of the Ca^{2+} -ATPase activity of rat GL-SR (\bullet) and SR (\bigcirc) upon Ca^{2+} concentration. Panel B. Dependence of the Ca^{2+} -ATPase activity of rat GL-SR upon Mg^{2+} -ATP. The activity measurements were done at 25 °C in the assay medium indicated in the Materials and methods, with varying Ca^{2+} (Panel A) or Mg^{2+} -ATP (Panel B) concentrations. The data are the average of n > 6 separate measurements at each Ca^{2+} and Mg^{2+} ATP concentrations. The 100% value of the Ca^{2+} -ATPase activity were 3.2 and 1.15 µmol ATP per min per mg protein for SR and GL SR, respectively.

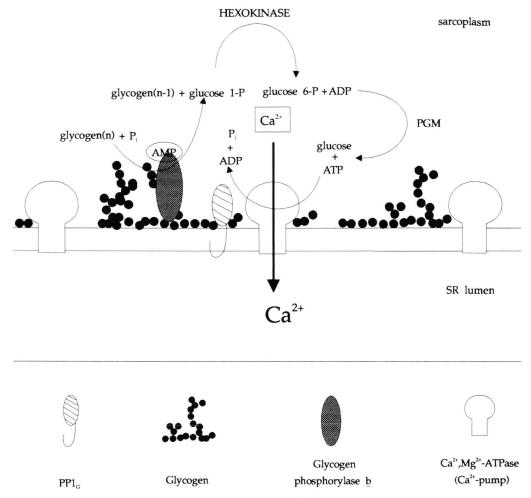


Fig. 6. Schematic view of the proposed metabolic shuttle for ATP production in GL-SR from rat skeletal muscle. Phosphorylase *b* binds to GL-SR through glycogen fragments as non-covalent anchoring sites (Wanson and Drochmans, 1972; Cuenda *et al.*, 1995), which in turn are anchored to SR membranes by the hydrophobic tail of the glycogen-associated form of protein phosphatase 1 (PP1_G) (Hubbard and Cohen, 1993). Since the location of phosphorylase **b** kinase of GL-SR relative to the SR membrane components is unknown and also because phosphorylase *b* kinase is not directly involved in this metabolic pathway, this protein has not been included in the diagram.

noted here that previous studies carried out in our laboratory with rabbit SR preparations showed that phosphorylase b binding to SR membranes does not alter Ca²⁺ binding to the Ca²⁺,Mg²⁺-AT-Pase, nor the Ca²⁺ dependence of this activity (Cuenda *et al.*, 1995). Since the concentration of phosphorylase in skeletal muscle sarcoplasm is ≥ 10 mg/ml (Fischer *et al.*, 1978), i.e. more than 100-fold higher than that used in the experiments reported in this study, these results indicate that this process takes place more efficiently at the concentrations of AMP reached in anoxia or dur-

ing skeletal muscle contraction, $100-200~\mu\text{M}$ (Busby and Radda, 1976; Newsholme and Leech, 1983).

The comparison of these data with those obtained in the presence of 2.5 mm ATP points out that the Ca²⁺ uptake saturates at approximately 200 nmol Ca²⁺/mg protein in both sets of data, and that the rate of Ca²⁺ uptake supported by glycogen phosphorolysis is about 5% of that obtained in the presence of 2.5 mm ATP. Attempts to directly measure the steady state concentration of ATP during Ca²⁺ uptake coupled to glycogen phospho-

rolysis failed, indicating that the ATP is kinetically compartmentalized in the GL-SR, i.e., the synthesized ATP has a restricted diffusion to the bulk medium. In several other cellular and subcellular systems, glucose metabolism in membrane associated compartments has been demonstrated to be functionally related to the activity of membrane ion pumps or channels through localized ATP synthesis (Paul et al., 1988; Weiss and Lamp, 1989; de Meis et al., 1992; Montero-Lomeli and de Meis, 1992). Compartimentalized ATP synthesis has also been demonstrated in skeletal muscle triads, which consist of transverse tubules connected to adjacent terminal cisternae of the SR membrane (Han et al., 1992). The local concentration of ATP in the GL-SR can be estimated from the dependence of the Ca²⁺,Mg²⁺-ATPase activity of GL-SR upon Mg²⁺-ATP (Fig. 5B). From the data shown in this Figure we obtain a steady state local concentration of $1-2 \mu M M g^{2+}$ -ATP.

In conclusion, in rat skeletal muscle the association of glycogenolysis in GL-SR membranes produces ATP nearby the SR Ca²⁺,Mg²⁺-ATPase

which is employed to drive the Ca²⁺ pump of the SR, as schematically drawn in the Fig. 6. This could lead to an efficient and faster decrease of cytosolic Ca²⁺ during the relaxation of the muscle, at a metabolic energy cost clearly lower than production of ATP elsewhere in the cell by glycolysis or by mitochondrial respiration. The compartmentalized production of ATP in GL-SR subcellular structures should increase in physiological situations in which the AMP concentration of the sarcoplasm is raised, such as in ischemia and in longlasting contractions (Busby and Radda, 1976; Newsholme and Leech, 1983), and might serve to protect muscle cells against the toxic effects of sustained high Ca²⁺ levels in the sarcoplasm.

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- Busby S.J.W. and Radda G.K. (1976), in Current Topics on Cellular Regulation, B.L. Horecker and E.R. Stadtman (eds.), (Vol. 10). Academic Press, New York, pp. 89–160.
- Clark M.G. and Lardy H.A. (1975), in Biochemistry of Carbohydrates (Biochemistry, Series One, Vol. 5), W.J. Whelan (ed.), MTP International Review of Science. Butterworths and University Park Press, London and Baltimore, pp. 223–266.
- Cuenda A., Henao F. and Gutiérrez-Merino C. (1990), Distances between functional sites of the Ca²⁺ + Mg²⁺-ATPase from sarcoplasmic reticulum using Co²⁺ as a spectroscopic ruler. Eur. J. Biochem. **194,** 663 667.
- Cuenda A., Centeno F. and Gutiérrez-Merino C. (1991), Modulation by phosphorylation of glycogen phosphorylase sarcoplasmic reticulum interaction. FEBS Lett. **283**, 273–276.
- Cuenda A., Nogues M., Gutiérrez-Merino C. and de Meis L. (1993), Glycogen phosphorolysis can form a metabolic shuttle to support Ca²⁺ uptake by sarcoplasmic reticulum membranes in skeletal muscle. Biochim. Biophys. Res. Commun. **196**, 1127–1132.
- Cuenda A., Henao F., Nogues M. and Gutiérrez-Merino C. (1994), Quantification and removal of glycogen phosphorylase and other enzymes associated with sarcoplasmic reticulum membrane preparations. Biochim. Biophys. Acta **1194**, 35–43.

- Cuenda A., Nogues M., Henao F. and Gutierrez-Merino C. (1995), Interaction between glycogen phosphorylase and sarcoplasmic reticulum membranes and its functional implications. J. Biol. Chem. **270**, 11998–12004.
- de Meis L., Grieco M.A.B. and Galina A. (1992), Reversal of oxidative phosphorylation in submitochondrial particles using glucose-6-phosphate and hexokinase as an ATP regenerating system. FEBS Lett. **3,** 197–01.
- Dubois M., Giles K.A., Hamilton J.K., Rebers P.A., and Smith F. (1956), Colorimetric methods for determination of sugar and related substances. Anal. Chem. **28**, 350–356.
- Entman M.L., Keslensky S.S., Chu A. and Van Winkle W.B. (1980), The sarcoplasmic reticulum-glycogenolytic complex in mammalian fast twitch skeletal muscle. Proposed *in vitro* counterpart of the contractionactivated glycogenolytic pool. J. Biol. Chem. **255**, 6245–6252.
- Fischer E.G., Alaba J.O., Brautiga D.L., Kerrick W.G.L., Malencik D.A., Moeschler H.J., Picton C. and Pocinwong S. (1978), in: Versatility of proteins. C.H. Li (ed.), Academic Press, New York.
- Gutiérrez-Merino C., García-Blanco F., Pocovi M., Menéndez M. and Laynez, J. (1980), Regulation of glycogen phosphorylase b by AMP. J. Biochem. **87**, 1483–1490.

- Han J.-W., Thieleczek R., Varsányi M. and Heilmeyer Jr., L.M.G. (1992), Compartmentalized ATP synthesis in skeletal muscle triads. Biochemistry **31**, 377–384.
- Hubbard M.J. and Cohen P. (1993), On target with a new mechanism for the regulation of protein phosphorylation. Trends. Biochem. Sci. **18**, 172–177.
- Lowry O.H., Rosebrough N.J., Farr A.L. and Randall R.J. (1951), Protein measurements with the phenol reagent. J. Biol. Chem. **193**, 265–275.
- Meissner G. (1984), Adenosine nucleotide stimulation of Ca²⁺ induced Ca²⁺ release in sarcoplasmic reticulum. J. Biol. Chem. **259**, 2365–2374.
- Montero-Lomelí M. and de Meis L. (1992), Glucose 6-Phosphate and hexokinase can be used as an ATPregenerating system by the Ca²⁺ ATPase of sarcoplasmic reticulum. J. Biol. Chem. 267, 1829–1833.
- Newsholme E.A. and Leech A.R. (1983), in: Biochemistry for the Medical Sciences. John Wiley and Sons, Ltd., London, U.K., Chapters 7 and 9.

- Paul R.J., Bauer M. and Pease W. (1979), Vascular smooth muscle: aerobic glycolysis linked to sodium and potassium transport processes. Science 206, 1414–1416.
- Tovmasian E.K., Hairapetian R.L., Bykova E.V., Severin S.E.Jr. and Haroutunian A.V. (1990), Phosphorylation of the skeletal muscle AMP-deaminase by protein kinase C. FEBS Lett. **259**, 321–323.
- Wanson J.C. and Drochmans P. J. (1972), Role of the sarcoplasmic reticulum in glycogen metabolism. Binding of phosphorylase, phosphorylase kinase, and primer complexes to the sarcovesicles of rabbit skeletal muscle. J. Cell Biol. **54**, 206–224.
- Weiss J.N. and Lamp S.T. (1989), Cardiac ATP-sensitive potassium channels. Evidence for preferential regulation by glycolysis. J. Gen. Physiol. **94**, 911–935.
- Yamada T. and Sugi H. (1987), ³¹P-NMR study of the regulation of glycogenolysis in living skeletal muscle. Biochim. Biophys. Acta **931**, 170–174.