Effect of glucose-6-P on the catalytic and structural properties of glycogen phosphorylase a

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Kinetic studies of muscle phosphorylase a in cationic buffer (pH 6.8) demonstrate that glucose-6-P competitively inhibits the binding of the substrate, glucose-1-P, to the enzyme. The inhibitory effect of glucose-6-P is largely overcome by glycerol-2-P. AMP counteracts inhibition of the enzyme by glucose-6-P, while glucose and glucose-6-P can interact to produce a synergistic inhibition of phosphorylase a activity. Preincubation of phosphorylase a with glucose-6-P at 20°C results in ~3-fold increase in activity, while ultracentrifugation experiments carried out under the same conditions showed that phosphorylase a can be converted to dimers by glucose-6-P.

Phosphorylase a Inhibition

Activation Dimerization

Structure-function

1. INTRODUCTION

It has been reported that glucose-6-P does not significantly influence the enzymatic activity of skeletal muscle glycogen phosphorylase a (EC 2.4.1.1) [1]. However, it is known that glucose-6-P activates phosphorylase phosphatase in vitro, and this activation is attributed to glucose-6-P induced changes of the substrate conformational phosphorylase a [2,3]. Furthermore, this ligand causes marked changes in the fluorescence properties of acetamidosalicylate-phosphorylase a or in the electron-spin resonance spectrum of a spinlabelled phoshorylase a, and formally competes with glycerol-2-P [4].

Experiments presented here demonstrate that in the absence of glycerol-2-P, a common buffer salt used in kinetic studies of phosphorylase, glucose-6-P considerably affects phosphorylase a activity, while, for the first time, ultracentrifugation provides a direct evidence for the dissociation of phosphorylase a tetramer to dimers by glucose-6-P.

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2. MATERIALS AND METHODS

Phosphorylase b was isolated from rabbit skeletal muscles as in [5]. The 4-times crystallized enzyme was treated as in [6] to remove bound AMP. Phosphorylase a was prepared from phosphorylase b by phosphorylation with rabbit muscle phosphorylase kinase [7]. The enzyme was recrystallized at least 3 times in 50 mM glycerol-2-P, 50 mM 2-mercaptoethanol, 1 mM EDTA buffer (pH 6.8). The crystals were dissolved in a buffer containing 5 mM glycerol-2-P (pH 6.8). 5 mM 2-mercaptoethanol, and 1 mM EDTA, treated twice with acid-washed Norit (1 mg/mg protein), and purified on a column of Sephadex G-25 equilibrated with the same buffer. Possible contamination of enzyme by phospho-dephospho-phosphorylase hybrid was checked as in [8]. Enzyme assayed under standard conditions (16 mM glucose-1-P) could not be further activated with 75 mM glucose-1-P. Protein concendetermined from tration was absorbance measurements at 280 nm as in [9].

Initial reaction rates were determined in the direction of glycogen synthesis. Reaction mixtures

were 0.5 ml and contained 3-5 μ g/ml phosphorylase a, 1% glycogen, 0.5 mM dithiothreitol, 0.5 mM EDTA, and 5 mM triethanolamine/HCl buffer (pH 6.8). Enzyme and glycogen were preincubated for 15 min at 30°C before initiating the reaction with glucose-1-P, care being taken to convert $\leq 20\%$ of glucose-1-P during the assay. P_i released in the reaction was measured as in [10].

Sedimentation velocity experiments were carried out using an MSE Centriscan 75 preparative and analytical ultracentrifuge, with a 10-mm single-sector cell at 60000 rev./min and $20 \pm 0.5^{\circ}$ C. Sedimentation coefficients obtained from direct measurements of the scanner traces were corrected for viscosity and density of the buffer to water at 20° C. The percentage of components with different sedimentation coefficients was estimated as in [11].

Oyster glycogen (BDH Chemicals) was freed of AMP as in [12]. AMP, glucose-1-P (dipotassium salt), and glucose-6-P (monosodium salt) were products of Sigma Chemical Co. (St Louis MO). Caffeine and glucose were purchased from Serva, and glycerol-2-P was obtained from E. Merck.

3. RESULTS

Fig.1 shows the inhibitory effects of increasing concentrations of glucose-6-P on phosphorylase a activity at various concentrations of glucose-1-P. An apparent competitive inhibition is indicated, since extrapolation of these plots to the ordinate, yields the same value for V (66.6 μ mol.min⁻¹.mg⁻¹). The double-reciprocal plots also show concave upward curvature, suggesting a homotropic cooperativity between substrate binding sites [13]. Thus, the result of adding increasing concentrations of glucose-6-P is to increase the apparent K_m values for the substrate and to increase also the value of Hill coefficient (fig.1 inset, table 1).

It was found that glucose-6-P inhibition of phosphorylase a except for triethanolamine—HCl, can be equally well demonstrated in Tris—HCl and also imidazole—HCl buffers (pH 6.8). On the contrary, this inhibition is largely reversed by glycerol-2-P and maleate. In the presence of 2 mM glucose-6-P, increasing concentrations of glycerol-2-P in the assay decrease both the apparent K_m value for glucose-1-P and the Hill coef-

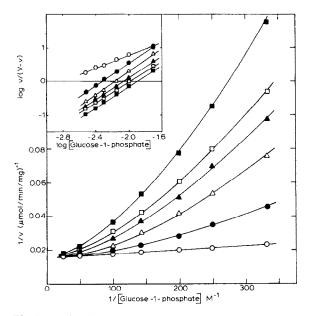


Fig.1. Kinetics of glucose-6-P inhibition of phosphorylase a with respect to glucose-1-P. Doublereciprocal plots for glucose-1-P binding in the absence (0) or presence of 1 (\bullet), 2 (Δ), 3 (\blacktriangle), 4 (\square) and 5 mM (**III**) glucose-6-P. The enzyme was assayed at 30°C in 5 mM triethanolamine-HCl buffer (pH 6.8), containing 0.5 mM dithiothreitol and 0.5 mM EDTA, with 1% glycogen. [Glucose-1-P] was varied from 3-40 mM. Inset: Hill plots for glucose-1-P. The V value with respect to glucose-1-P was 66.6 μ mol.min⁻¹.mg⁻¹. Log v/(V-v) was plotted against log(glucose-1-P) which yields Hill n (Hill coefficient) as the slope and the apparent $K_{\rm m}$ values [14].

ficient (table 1). However, glycerol-2-P alone in the concentrations tested has no effect on the enzyme activity. Practically, the enzyme is not inhibited by 2 mM glucose-6-P in the presence of 20 mM glycerol-2-P in agreement with [1].

The results of the kinetics of glucose inhibition of phosphorylase a obtained under the same conditions are summarized in table 1. Since it appears that glucose-6-P acts in a similar way as glucose which also increases the homotropic cooperativity between substrate molecules [14], we examined the joint action of these two inhibitors on the enzyme activity. A marked synergistic effect of the two in inhibiting phosphorylase a occurs as indicated in fig.2a. In fig.2b, the intercept on the x-axis gives $-\alpha K_i$ (the inhibition constant for glucose in the presence of glucose-6-P) [15]. Thus, from the plot,

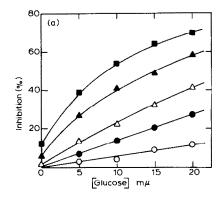
Table 1 Kinetic parameters for phosphorylase a

Additions	K _m (mM)	n	
None	1.6 ± 0.2	1.0	
Glucose-6-P (1 mM)	4.6	1.6	
Glucose-6-P (2 mM)	6.9	1.7	
Glucose-6-P (3 mM)	8.5	1.7	
Glucose-6-P (4 mM)	10.3	1.55	
Glucose-6-P (5 mM)	12.3	1.6	
Glucose-6-P (2 mM) +			
glycerol-2-P (2.5 mM)	6.0	1.5	
Glucose-6-P (2 mM) +			
glycerol-2-P (5 mM)	4.8	1.4	
Glucose-6-P (2 mM) +			
glycerol-2-P (10 mM)	3.5	1.25	
Glucose (10 mM)	4.2	1.3	
Glucose (20 mM)	7.0	1.35	
Glucose (30 mM)	9.6	1.45	

The apparent K_m and n (Hill coefficient) values with respect to glucose-1-P were calculated from the data of fig.1 as described in the legend. Kinetic experiments of glucose inhibition or glycerol-2-P effect on glucose-6-P inhibition of phosphorylase a were performed under the same conditions as in fig.1, and the data obtained were plotted as Hill plots [14] to calculate the kinetic parameters. In the absence of inhibitor the mean K_m is quoted together with the standard deviation of the mean (10 determinations)

 $\alpha K_i = 1.2$ mM, and if we take $K_i = 6$ mM (an approximate value obtained from the data of table 1, based on the assumption that K_m , app. = $K_m(1 + I/K_i)$), then α , an interaction factor, is 0.2. However, we cannot calculate the inhibition constant for glucose-6-P in the presence of glucose, because the replot of the slopes vs the corresponding glucose-6-P concentrations yields a curve (fig.2b, inset). It is worth noting that the synergism between glucose and glucose-6-P exists even in the presence of glycerol-2-P, but in this case higher concentrations of glucose-6-P are required.

In table 2, the effects of glucose-6-P, glucose and caffeine on phosphorylase a activity are presented. Caffeine, which is known to competitively inhibit the binding of glucose-1-P to the enzyme and also to act synergistically with glucose [15], can also interact with glucose-6-P to produce an apparent synergistic inhibition of phosphorylase a activity. Furthermore, in the presence of the



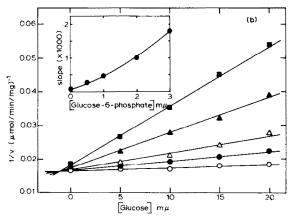


Fig. 2. (a) Inhibition of phosphorylase a activity by glucose, glucose-6-P and a combination of both inhibitors. The enzyme was assayed with 20 mM glucose-1-P and 1% glycogen in 5 mM triethanolamine—HCl buffer (pH 6.8), containing 0.5 mM dithiothreitol and 0.5 mM EDTA, at 30°C. Glucose concentrations were as indicated in the figure. Glucose-6-P concentrations were 0 (\bigcirc), 0.5 (\bullet), 1 (\triangle), 2 (\triangle) and 3 mM (\blacksquare). (b) Dixon plots of $1/\nu \nu s$ [glucose] at different fixed [glucose-6-P] (the above mentioned).

Inset: The replot of slopes vs [glucose-6-P].

3 ligands together, enzyme activity is nearly eliminated. However, AMP is able to completely reverse these inhibitory actions.

Fig. 3 shows that preincubation of phosphorylase a (5 mg/ml) with 5 mM glucose-6-P at 20°C results in a marked increase of enzymatic activity. At 30 min, activation with glucose-6-P is ~3-times greater than that observed without glucose-6-P. The data in fig. 3 also show that stimulation of enzymatic activity can also be demonstrated if enzyme is preincubated with 50 mM glucose in agreement with [16], while the addition of 50 μ M AMP

Table 2

Synergistic inhibition of phosphorylase *a* activity by glucose-6-P, glucose, and caffeine

Additions	Activities (%)			
	- AMP		+ 50 μM AMP	
	100	(53)	100 (73)	
Glucose-6-P (1 mM)	67		98	
Glucose (5 mM)	86		96	
Caffeine (0.5 mM)	57.:	5	97	
Glucose-6-P (1 mM) + glucose (5 mM)	27		99	
Glucose-6-P (1 mM) +				
caffeine (0.5 mM)	16		96	
Glucose (5 mM) +				
caffeine (0.5 mM)	16		93	
Glucose-6-P (1 mM) + glucose (5 mM) +				
caffeine (0.5 mM)	2.	5	93	

The enzyme was assayed with 5 mM glucose-1-P and 1% glycogen in the absence or presence of AMP as in section 2. Activities indicated are the percentage remaining when compared to controls taken as 100%. Values in parentheses are the control activities in μ mol.min⁻¹.mg⁻¹

to phosphorylase a which had been preincubated for 44 min with glucose-6-P results in an instantaneous decrease of activity. To relate alterations of enzymatic activity to quaternary structure of phosphorylase a, sedimentation velocity experiments were carried out under the same conditions of stimulation of activity (20°C). The following results were obtained (table 3):

- (i) Phosphorylase a in the presence of 5 mM glucose-6-P sediments mainly as a dimer $(s_{20,w} = 8.8)$, while a minor fast-sedimenting component is formed with an $s_{20,w}$ of value of 13.4 corresponding to the tetramer [16];
- (ii) The identified dimeric form of phosphorylase a in the presence of glucose-6-P is able to be associated to the tetrameric form with low concentrations of AMP;
- (iii) Glucose (10 mM) and glucose-6-P (2 mM) when added independently to the enzyme cause partial dissociation to dimers, while a cooperative promotion of dimer formation when the two ligands are added together can be observed.

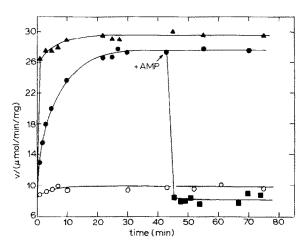


Fig. 3. Effect of glucose-6-P on the activity of phosphorylase a. Phosphorylase a (5 mg/ml) was preincubated at 20°C in 15 mM triethanolamine—HCl, 0.5 mM dithiothreitol, 0.5 mM EDTA buffer (pH 6.8) in the absence (○) or presence of 5 mM glucose-6-P (●) or 50 mM glucose (▲). Aliquots were removed at various intervals for measurement of activity at 20°C. The assay mixture (0.2 ml) contained 0.3 mg/ml of phosphorylase a, 1% glycogen, 20 mM glucose-1-P in 15 mM triethanolamine—HCl, 0.5 mM dithiothreitol, 0.5 mM EDTA buffer (pH 6.8) (30 s assay). AMP (50 µM) was added to enzyme which had been incubated for 44 min with glucose-6-P (■).

4. DISCUSSION

These results show that in the presence of the inhibitor, glucose-6-P, the previously cooperative phosphorylase a now displays cooperative glucose-1-P binding (fig.1). The kinetics of phosphorylase b with respect to glucose-1-P becomes also cooperative in the presence of glucose-6-P [13], while glucose-6-P is essentially without effect on phosphorylase b' activity [17], indicating that the first 19 residues in phosphorylase are important in the expression of homotropic and heterotropic interactions [18]. Xray diffraction studies on phosphorylase a crystals have shown that the AMP/ATP binding site is remarkably indiscriminate in binding phosphatecontaining compounds. There is also evidence that the glucose moiety of glucose-6-P is firmly bound to the crystals [19]. Glucose-6-P inhibition of the enzyme is competitively reversed by increasing substrate concentration (fig. 1) and this may be also due to some extent to the secondary binding of

Table 3

Effect of glucose, glucose-6-P, and AMP on the sedimentation properties of phosphorylase a

Additions	\$20,w	% of total
None	13.7	100
Glucose (50 mM)	8.3	100
Glucose (10 mM)	12.4	74
•	8.5	26
Glucose-6-P (2 mM)	13.2	63
	8.6	37
Glucose-6-P (5 mM)	8.8	75
	13.4	25
Glucose-6-P (20 mM)	8.9	83
	13.6	17
Glucose-6-P (2 mM) +		
glucose (10 mM)	8.9	100
Glucose-6-P (5 mM) +		
AMP (25 μM)	13.7	73
	8.8	27
Glucose-6-P (5 mM) +		
AMP (100 μM)	13.9	100

All ultracentrifugal sedimentation velocity measurements were performed with 5 mg enzyme/ml in 15 mM triethanolamine—HCl, 0.5 mM dithiothreitol, 0.5 mM EDTA buffer (pH 6.8) at 20°C. Other components were added as indicated. Other conditions were as in section 2

glucose-1-P in the same location as AMP [20]. The inhibitory action of glucose-6-P is also reversed by glycerol-2-P, suggesting that glycerol-2-P possibly binds in the AMP/ATP binding site.

Glucose-6-P in inhibiting the enzymatic reaction acts in a similar way as does glucose. Low concentrations of these ligands alone are only slightly inhibitory in saturating concentrations glucose-1-P. However, in combination they markedly inhibit the activity of the enzyme (fig.2). This synergistic effect indicates different binding sites for the two inhibitors [21] and the determined α -value of 0.2 for the glucose/glucose-6-P pair indicates that positive interactions exist between these two ligands in the enzyme-glucose-glucose-6-P complex [22]. It has also been reported that the inhibitors glucose and glucose-6-P are synergistic in their action on enzymatic activity and tryptic digestion of phosphorylase b [23,24].

Glucose and glycogen induce an increase of en-

zymatic activity of phosphorylase a at low temperature (15–23°C) promoting the dissociation of the tetrameric form to dimers [25]. Since phosphorylase a activation by glucose-6-P may be correlated with the formation of a dimeric species (fig.3, table 3), this suggests that stimulation is directly related to the dimerization of phosphorylase a. It has been reported that tetrameric phosphorylase a cannot be dissociated to the dimeric form by glucose-6-P [8,26]. The contradiction between our results and those in [8,26] might be due to the presence of anionic salts (glycerol-2-P and maleate) in their ultracentrifugation experiments.

This report presents evidence that the well known stimulatory effect of glucose-6-P in the dephosphorylation of phosphorylase a by phosphorylase phosphatase observed at low temperature [3], could be attributed to a dissociation phenomenon.

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