Dephosphorylation of phosphorylase kinase by a histone H1-stimulated phosphoprotein phosphatase

Ronald L. Mellgren, Susan E. Wilson and Keith K. Schlender

Department of Pharmacology and Therapeutics, Medical College of Ohio, C.S. 10008, Toledo, OH 43699, USA

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A histone H1-stimulated protein phosphatase isolated from rabbit skeletal muscle dephosphorylated [32 P]phosphorylase kinase. The rate of dephosphorylation was markedly increased by 5-50 μ g histone H1/ml. Only the α and α' subunits were dephosphorylated in the absence or presence of histone H1. This is consistent with previous results suggesting that the H1-stimulated phosphatase is a type-2 protein phosphatase. The present studies suggest that the histone H1-stimulation is the result of a direct interaction of the histone with the phosphatase.

Dephosphorylation

Phosphorylase kinase

Histone H1

Protein phosphatase

1. INTRODUCTION

Recently, we described the isolation of a heatstable activator of phosphorylase phosphatase [1], and demonstrated that it was histone H1 [2]. In subsequent studies, it was found that a histone H1-stimulated phosphorylase phosphatase from rabbit skeletal muscle could be inhibited by 0.1 mM ATP, but not by phosphatase inhibitor-1 [3]. A bovine aorta histone H1-stimulated phosphatase was also insensitive to heat-stable phosphatase inhibitor-1 and -2 [4]. Thus, these H1-stimulated phosphatases appeared to be type-2 phosphatases according protein nomenclature in [5,6]. To characterize further the H1-stimulated rabbit muscle phosphatase we have investigated the influence of histone H1 on phosphatase activity utilizing as substrate phosphorylase kinase phosphorylated on the α,α' and β subunits. These studies provide further evidence that the H1-stimulated phosphatase is a type-2 protein phosphatase, and suggest that stimulation by H1 is the result of its interaction with the phosphatase and not with substrates.

2. MATERIALS AND METHODS

The histone H1-stimulated phosphatase and phosphatase-1 catalytic subunit were purified from rabbit skeletal muscle as in [3]. Rabbit skeletal muscle [32P]phosphorylase a was prepared as in [7], and phosphatase inhibitor-1 as in [8]. Phosphorylase kinase was purified from rabbit skeletal muscle by the method in [9] through the Sepharose gel filtration step, and phosphorylated with $[\gamma^{-32}P]$ ATP (400–1000 cpm/pmol) and cyclic AMP-dependent protein kinase catalytic subunit essentially as in [10]. The phosphorylated phosphorylase kinases used in this study contained $0.92-1.24 \mu \text{mol}$ phosphate per μmol $(\alpha + \alpha')$ subunits and $0.55-0.58 \mu mol$ phosphate per μmol subunit. The standard conditions dephosphorylation were: $0.5-1.0 \,\mu\text{M}$ ³²P as phosphorylase a or phosphorylase kinase ($\alpha + \alpha'$)subunits in 100 µl of 50 mM imidazole-HCl, 1.8 mM EDTA, 0.5 mM EGTA, 2 mM dithiothreitol, 0.2 mg bovine serum albumin/ml, 2% glycerol (pH 7.2). Dephosphorylation was carried out at 30°C in the absence or presence of various concentrations of histone H1. The histone H1 was purified from calf thymus as in [11]. Phosphatase activity was determined by release of 10% tri-chloroacetic acid-soluble ³²P from substrates [7]. SDS-polyacrylamide gel electrophoresis was performed in 7.5% slab gels using the buffer system in [12].

3. RESULTS

3.1. Dephosphorylation of phosphorylase kinase by a histone H1-stimulated protein phosphatase

The skeletal muscle histone H1-stimulated protein phosphatase which was previously shown to dephosphorylate [32 P]phosphorylase a [3] also had activity on 32 P-labeled phosphorylase kinase. In the absence of histone, the ratio of phosphatase activities on phosphorylase kinase νs phosphorylase a was 0.33 when both substrates were assayed at the same concentration (1.0 μ M accessible 32 P),

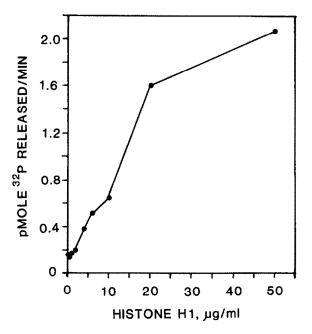


Fig. 1. The effect of histone H1 on phosphorylase kinase phosphatase activity. Phosphorylase kinase phosphatase activity was determined as described in section 2 in the presence of various concentrations of histone H1. The concentration of 32 P-labeled ($\alpha + \alpha'$)-subunits was 0.5 μ M. Each assay contained 39 munits of H1-stimulated phosphorylase phosphatase activity/ml measured in the presence of histone H1, using the unit definition and assay described in [3].

and in the same assay buffer (see section 2). The addition of histone H1 greatly increased the rate of dephosphorylation of the phosphorylase kinase in a concentration-dependent manner (fig.1). Neither the basal activity nor the H1-stimulated activity

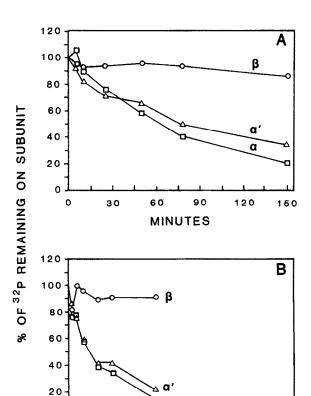


Fig.2. Specificity of the H1-stimulated phosphatase for the α and α' subunits of phosphorylase kinase. Samples of phosphorylase kinase (0.69 µM total 32P) were incubated in the standard dephosphorylation buffer in the absence or presence of 8 µg histone H1/ml (panels A and B, respectively). H1-stimulated phosphatase (78 munits/ml) was added to each mixture, and at various times aliquots were removed and heated at 100°C for 2 min in the presence of 1.5% SDS and 220 mM β mercaptoethanol. The samples were electrophoresed as described in section 2, stained with Coomassie blue, destained, and the α,α' and β subunits excised and counted in a liquid scintillation counter. The percentage of ³²P counts remaining in each subunit was determined by comparison to the label present in the subunits of a sample which was electrophoresed before the addition of phosphatase.

60

MINUTES

90

120

150

0

0

30

was inhibited by phosphatase inhibitor-1, but activity in the presence or absence of H1 was stimulated by 5 mM Mn²⁺ (not shown).

3.2. Specificity toward the phosphorylated subunits of phosphorylase kinase

Dephosphorylation of the α,α' and β subunits by the H1-stimulated phosphatase was determined by liquid scintillation counting of the Coomassiestained subunits excised from SDS-gels (fig.2). The phosphatase was highly specific for the α and α' subunits (fig.2, panel A). The addition of histone H1 greatly accelerated dephosphorylation of these subunits, but did not allow dephosphorylation of the β -subunit (fig.2, panel B). In a similar experiment, utilizing purified catalytic subunit of type-1 phosphatase it was possible to dephosphorylate nearly totally the β -subunit without dephosphorylating the α and α' subunits (not shown).

4. DISCUSSION

The rabbit skeletal muscle histone H1-stimulated protein phosphatase was capable of dephosphorylating [32P]phosphorylase kinase, and the rate of dephosphorylation was greatly increased by histone H1 (fig.1). Thus, the H1-stimulation of this phosphatase is not restricted to the dephosphorylation of phosphorylase a, the only other substrate previously studied [1-4]. The effect of histone H1 on both phosphorylase a and phosphorylase kinase dephosphorylation suggests, but does not prove, that the stimulation is not a substrate-directed effect. If phosphorylase a is used as substrate, the latter possibility must be considered since protamine, which activated the H1-stimulated phosphatase [3], has a direct inhibitory effect on phosphorylase a [13]. However, at a concentration of 15 µg/ml, histone H1, which was a better activator of the phosphatase than protamine [3], did not inhibit phosphorylase a activity (unpublished). This concentration of histone H1 dramatically activated the phosphatase activity with phosphorylase kinase (fig.1) or phosphorylase a [3] as the substrate. Thus, the activation of the phosphatase by histone H1 could be resolved from polycationinduced inhibition of phosphorylase a. Furthermore, maximum activation of phosphorylase phosphatase activity could be achieved in the presence of a 10-fold molar excess of phosphorylase a relative to histone H1 [1,3]. Since histone H1-stimulation appears to be phosphatase-directed, it is likely that histone H1 may influence the dephosphorylation of other phosphoprotein substrates in addition to phosphorylase a and phosphorylase kinase. For example, there may be nuclear phosphoproteins which serve as substrates for an H1-stimulated phosphatase. In recent experiments, H1-stimulated phosphorylase phosphatase activity has been detected in rat liver nuclear preparations (unpublished).

The specificity of the histone H1-stimulated phosphatase for the α and α' subunits of phosphorylase kinase (fig.2) confirms the previous suggestion that the enzyme is a type-2 phosphatase [3,4]. Among the type-2 phosphatases, only phosphatases 2B and 2C have known metabolic regulators, Ca^{2+} and Mg^{2+} respectively [6]. There is no known metabolic regulator of phosphatase 2A [6], the previously described type-2 protein phosphatase which most closely resembles the H1-activated phosphatase. It will be of interest to determine if a histone H1-like activator exists in the cytoplasm, and to establish if histone H1 is important in regulating the dephosphorylation of nuclear phosphoproteins.

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