Monoclonal Antibodies to a Membrane Glycoprotein Induce the Phosphorylation of Histone H1 in Sea Urchin Spermatozoa

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Abstract. Two groups of mAbs reacting with external domains of a major sea urchin sperm membrane glycoprotein of 210 kD were isolated. Previous studies have shown that group I mAbs inhibit the acrosome reaction induced by egg jelly and also cause large increases in intracellular Ca²⁺ ([Ca²⁺]i). Group II mAbs, at comparable levels of cell surface binding, neither inhibit the egg jelly-induced acrosome reaction nor cause increases in [Ca²⁺]. In this paper, we investigate the ability of these mAbs to induce the cAMP-dependent phosphorylation of sperm histone H1. Group I mAbs induce H1 phosphorylation to the same level and on the same peptide, as occurs upon treatment of sperm with egg jelly. These mAbs also activate adenylate cyclase to the same extent as egg jelly. Group II mAbs do not induce H1 phosphorylation and are only poor activators of adenylate cyclase. Group I mAbs compete with each other, but not with group II mAbs, for binding to the cell surface. These data indicate that the activation of adenylate cyclase is an initial event in the pathway leading from the binding of mAbs to a specific domain of the 210-kD protein at the cell surface, to the discrete phosphorylation of histone H1 in highly condensed sperm chromatin. The domain on the 210-kD protein recognized by group I mAbs plays a critical role in signal transduction during the early events of fertilization.

Sea urchin eggs are covered by a jelly layer that can be solubilized by brief exposure to pH 5 seawater. The major macromolecule of egg jelly is a fucose sulfate-rich glycoconjugate (FSG) that induces the acrosome reaction of sperm (Segall and Lennarz, 1979, 1981; Garbers et al., 1983; DeAngelis and Glaube, 1987). This morphological change consists of the exocytosis of the acrosome granule and the polymerization of actin to form the acrosomal process, which is covered by the membrane destined to fuse with the egg (reviewed in Lopo, 1983; Tilney, 1985). FSG has profound physiological effects on sperm, including the opening of ion channels, resulting in the net influx of Ca²⁺ and Na⁺, the net efflux of K⁺ and H⁺, and the depolarization of the K⁺-supported membrane potential (reviewed by Schackmann, 1986, 1988). In addition to these ionic events, FSG induces the activation of adenyly cyclase and protein kinase, the elevation of cAMP concentrations (reviewed in Garbers and Kopf, 1980), and the elevation of inositol trisphosphate (Domino and Garbers, 1988). These effects of FSG are all dependent on the presence of external Ca²⁺. The major substrate phosphorylated in vivo by the FSG-induced increases in cAMP-dependent protein kinase activity is sperm-specific histone H1 (Porter and Vacquier, 1986), which is phosphorylated on a single serine site in the peptide RKGS(PO₄)SNAR (Porter et al., 1988a). The phosphorylation of H1 on this single site in response to FSG may be a prerequisite for the hyperphosphorylation of H1 by egg kinases and its loss from sperm chromatin within 10 min after the sperm nucleus enters egg cytoplasm (Green and Poccia, 1985; Porter and Vacquier, 1988; Porter et al., 1988b).

These effects of FSG on sperm are complex, and the relationships between the changes in ionic permeability and the mobilization of the cyclic nucleotide cascade leading to H1 phosphorylation are not well understood. Although it has never been rigorously demonstrated, these effects of FSG on sperm must involve interaction with extracellular domains of cell surface proteins. In an attempt to identify sperm surface proteins involved in the recognition of FSG and the transduction of this signal across the plasma membrane, mAbs were isolated that react with extracellular domains of sperm membrane proteins (Trimmer, 1987). The major antigenic determinant on the sperm surface is a glycoprotein of 210–260 kD (the 210-kD protein). Two groups of mAbs reacting mono-
specifically with this protein were isolated. Group I mAbs inhibit the FSG-induced acrosome reaction (Trimmer et al., 1985; Trimmer and Vacquier, 1986), yet induce large increases in intracellular Ca²⁺ (Trimmer et al., 1986). Group II mAbs do not exhibit these effects on sperm. Here we show that the group I mAbs, but not the group II mAbs, are potent inducers of adenylyl cyclase and the phosphorylation of histone H1. This indicates that the 210-kD protein mediates the activation of sperm adenylyl cyclase. The domain of the 210-kD protein recognized by group I mAbs must be critical for the interaction of FSG with sperm during fertilization.

Materials and Methods

mAbs

The production of mouse mAbs reacting with external domains of sea urchin sperm plasma membrane proteins has been described in detail (Trimmer et al., 1985, 1986, 1987; Trimmer, 1987; and Trimmer and Vacquier, 1988). This description includes preparation of the immunogen, screening by ELISA, labeling sperm with ³²P, detergent extraction of labeled sperm, immunoprecipitation of labeled protein, gel electrophoresis of immunoprecipitates and autoradiography, production of ascites fluid, isotyping of immunoglobulins, Fab fragment production, ¹²⁵I labeling of antibodies, and competitive binding experiments. In the experiments reported in this paper, mAb preparations used were IgG partially purified from ascites fluid in borosilicate tubes, 5 Ixl stock sperm suspension was added, and slow rotary centrifugation at 1,000 g for 5 min and washed twice in seawater containing 0.1 M glycine and 1 mg/ml BSA. A stock suspension of fixed cells was pre-

Miscellaneous Procedures

0.1 M glycine and 1 mg/ml BSA. A stock suspension of fixed cells was pre-

Adenylyl Cyclase

The method described by Schultz and Jakobs (1983) was modified as follows. Buffer A contained 90 mM Hepes, pH 7.5, 6.6 mM cAMP, 3 mM ATP, 15 mM MγG, and 60 mM NaN₄. Buffer B contained 10 mM NaCl, 10 mM KCl, 2 mM Hepes (pH 7.5), 20 mM NaN₄, 300 µg/ml creatine kinase, and 45 mM creatine phosphate (pH 7.5). Buffer C contained 10 mM NaCl, 10 mM KCl, 2 mM Hepes, 5 µM EDTA, 0.1 mM DTT, 20 mM NaN₄, and 12 mg/ml BSA. 1-ml aliquots of all three buffers were stored at −70°C. One part buffer A was mixed with one part buffer B and the isotope ([³²P]ATP, [²⁵I]Biochemicals) was added (1 µCi/84 µl A + B). Usually, 750 µl of this mixture was placed in a 1.5-ml Eppendorf tube in a 23°C water bath. The cell lysate was then prepared exactly as follows. A 1-ml portion of seawater, containing egg jelly (185 µg fucose/ml) or mAb was placed in Eppendorf tubes at 23°C. A fresh 1:10 dilution of semen (dry sperm) was made, 60 µl (600 µg protein) was added, and the tube was capped and inverted. At 120 s, the tube was microfuged for 45 s, the supernatant was immediately removed by aspiration and 1 ml 0°C seawater was added, the cell pellet was resuspended with a 1-ml disposable pipette tip, the cells were pelleted by 45 s of centrifugation, the supernatant was carefully removed, and 500 µl buffer C (0°C) was added to lyse the cells. The lysate was piled up several times in the pipette tip, and at exactly 8 min after exposure of the cells to the test medium, 375 µl of lysate was added to 750 µl of reaction mixture. Every 3 min, 120 µl of reaction mixture (~36 µg sperm protein) was removed and added to 480 µl of 0.2 M zinc acetate, followed by 600 µl 0.2 M NaClO₃. The tubes were capped, vortexed, and either stored frozen or centrifuged 5 min at top speed in a microfuge. 1 ml of supernatant was put over 3 ml columns of neutral aluma equilibrated in 100 mM Tris pH 7.5. The columns were washed into scintillation vials with two portions of 3 ml Tris buffer and the Cerenkov radiation was determined in a scintillation counter. [³²P]cAMP was used to monitor recovery of [³²P]cAMP formed; recovery was virtually 100%. No [³²P]ATP substrate was detectable in the column eluates. Data points were linear from 3 to 18 min and activity was proportional to added sperm protein. Phosphodiesterase inhibitors were not included in the assay.

Results

Characterization of mAbs

The group I mAbs, J10/14 (IgG₂), J18/2 (IgG₃), J16/5 (IgG₄), and J17/10 (IgG₂); and group II mAbs, J4/4 (IgG₂) and J18/27 (IgG₃), react monospecifically with the 210-kD protein, as shown by autoradiograms of SDS-PAGE analysis of immunoprecipitates of detergent extracts of ¹²⁵I-labeled sperm proteins (Fig. 1). The lane of total detergent extract of radiolabeled sperm used as the starting material for the
Figure 1. Autoradiogram of immunoprecipitates of mAbs. All six mAbs react exclusively with the 210-kD protein. The total 125I-labeled sperm membrane protein used as the starting material for the immunoprecipitates appears in the far right lane with approximate relative mass given in kilodaltons (kD). Group I mAbs are J18/2, J10/14, J16/5, and J17/10. Group II mAbs are J4/4 and J18/27. T8/40 serves as a negative control.

immunoprecipitates shows the specificity of the mAbs for the 210-kD protein.

**mAb-induced Phosphorylation of H1**

To investigate the effects of these mAbs on the in vivo phosphorylation of H1, the ATP pools of sperm were labeled with 32P and the cells were diluted into seawater containing egg jelly or mAb. Group I mAbs (J10/14, J16/5, and J17/10) at 100 µg/ml induced H1 phosphorylation to the same extent as did egg jelly (Fig. 2). Group II mAbs (J4/4 and J18/27), and the negative control mAb T8/40, were noninductive at concentrations from 5 to 400 µg/ml (Fig. 2). Both J10/14 IgG and Fab fragments were inductive at concentrations as low as 5 µg/ml. mAb J18/2 (group I) was unique in that at high concentrations (100–200 µg/ml, Fig. 2) it was noninductive, but at lower concentrations it induced H1 phosphorylation to the same extent as seen upon treatment of sperm with egg jelly (Fig. 3). In this same experiment, mAb J17/10 (group I) was inductive to 200 µg/ml, whereas J18/27 (group II) was noninductive at all concentrations tested.

**H1 Site Phosphorylated by mAb Treatment**

Previous chemical analysis for phosphate of purified H1 had

Figure 2. mAb-induced phosphorylation of histone H1. Total sperm protein (35µg) stained with Coomassie brilliant blue is shown in the left two lanes. T, tubulin; H1, histone H1; CH, core histones. The nine lanes to the right of the protein stain are autoradiograms of total sperm protein (35 µg). +Jelly, whole egg jelly at 150 µg fucose per ml. Sperm were exposed for 10 min to 100 µg mAb/ml before fixation. Note that J18/2 did not induce H1 phosphorylation.

Figure 3. An autoradiogram showing that high concentrations of J18/2 do not induce H1 phosphorylation, whereas lower concentrations are inductive. In this experiment, J18/2 at 25 µg/ml was as inductive as egg jelly, but with increasing concentration of this mAb, less phosphorylation of H1 occurred, whereas with J17/10, (and also J10/14 and J16/5) this decrease in H1 phosphorylation with increasing mAb concentration was not observed. Only the H1 (arrow) portion of the autoradiogram is presented. mAb concentrations in micrograms per milliliter are shown over each lane. J18/27 was noninductive from 25 to 400 µg/ml. +Jelly, positive control; -Jelly, negative control.
shown that before treatment with egg jelly there is 0.16 mol phosphate/mol protein, and that after egg jelly treatment this number increases to 0.46 (Porter and Vacquier, 1986). Two-dimensional, thin-layer separation of Endo Arg C digests of the NH₂-terminal CNBr-generated fragment of H1 yielded only one labeled peptide (Porter et al., 1988a). Mixing experiments showed that this same peptide was the only one labeled in vivo when the H1 kinase was activated by treating sperm with the phosphodiesterase inhibitor 3-isobutyl-1-methylxanthine, or when sperm were permeabilized with detergent in the presence of [γ-³²P]ATP and cAMP. The sperm H1 kinase is cAMP dependent and the sequence of the phosphorylated peptide (RKGS(PO₄)SNAR) is a preferential site for cAMP-dependent protein kinase (Porter et al., 1988a). To test whether this same site was phosphorylated in response to mAb treatment, the ³²P-labeled histone H1 was purified from mAb J10/14-treated sperm and the two-dimensional Endo Arg C maps were prepared. The same ³²P-labeled peptide previously reported as being labeled upon egg jelly treatment was the only major spot appearing on the autoradiogram of the peptide map (Fig. 4). Thus, in the highly condensed native sperm chromatin, the endogenous cAMP-dependent protein kinase phosphorylates H1 on the same peptide, whether activated by egg jelly, phosphodiesterase inhibitors, or mAbs.

### Binding of mAbs to the Sperm Surface

To test if the failure of mAbs J4/4 and J18/27 to induce the phosphorylation of H1 was due to a lack of sperm surface binding, binding analyses were performed under the conditions used for the phosphorylation experiments. We previously reported that J4/4 saturates all binding sites at ~125 μg/ml when the sperm concentration is 4 × 10⁸ cells/ml (Trimmer et al., 1985). We observed no phosphorylation of H1 in J4/4 at 200 μg/ml and sperm concentrations of 1 × 10⁴ cells/ml. We also reported that J10/14 IgG saturates cell surface binding sites at ~125 μg/ml, and that there is no difference in number of binding sites in living cells as compared with paraformaldehyde/glutaraldehyde fixed cells (Trimmer et al., 1985; Trimmer, 1987).

In Table I, we present data on the relative binding of these six anti–210-kD mAbs to sperm under incubation conditions equivalent to those used for the in vivo phosphorylation of H1 by mAbs (0–200 μg/ml). Fixed cells were used in these experiments, as we have never observed a quantitative difference between the binding of mAb to fixed as opposed to living sperm (the hybridomas were selected in an ELISA using fixed cells [Trimmer et al., 1985]). Also, at subsaturating concentrations, mAbs J18/2 and J18/27 cause the 210-kD protein to be shed from the surfaces of living cells (Trimmer and Vacquier, 1988). The surface binding data are presented as the relative amount of mAb bound at 100 μg/ml as a percentage of the amount bound at 200 μg/ml (Table I). Using this cell surface binding assay, it is evident that at 100 μg/ml, mAbs J18/27 and J10/14 are present at saturating concentrations, whereas J4/4, J18/2, J16/5, and J17/10 are slightly below saturation. The relative amount of fluorescent second antibody bound at 200 μg mAb/ml was lowest for J4/4, being about one third to one fourth of the group I mAbs. However, the other group II mAbs, J18/27, bound roughly the same amount of second antibody as the group I-treated cells (Table I). The differences in inductive ability exhibited by group I and group II mAbs thus are not due simply to differences in cell surface binding.

To study in more detail the binding of these mAbs and how it relates to the striking differences in their ability to induce H1 phosphorylation, we performed competitive binding analysis using ¹²⁵I-J10/14. The group I mAbs J16/5 and J18/2 compete as well as J10/14 itself for ¹²⁵I-J10/14 binding to cells, whereas the group II mAbs (J4/4 and J18/27) are non-competitive at all concentrations tested, to 250 μg/ml (Fig. 5). It therefore appears as if the group I mAbs bind at or near the same site on the 210-kD protein, whereas the group II mAbs bind a different region(s) of the protein.

From the data presented thus far, it is clear that mAb J18/2 is unique among the group I mAbs in that it displays a narrow concentration dependence in its ability to induce H1 phosphorylation, even though it clearly binds to the 210-kD protein at high levels at the noninductive concentrations (Table

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### Table I. Relative Binding of Anti–210-kD mAbs

<table>
<thead>
<tr>
<th>Group</th>
<th>mAb</th>
<th>A</th>
<th>B</th>
<th>Relative fluorescence</th>
</tr>
</thead>
<tbody>
<tr>
<td>II</td>
<td>J4/4</td>
<td>87</td>
<td>140</td>
<td></td>
</tr>
<tr>
<td>II</td>
<td>J18/27</td>
<td>100</td>
<td>620</td>
<td></td>
</tr>
<tr>
<td>I</td>
<td>J18/2</td>
<td>80</td>
<td>400</td>
<td></td>
</tr>
<tr>
<td>I</td>
<td>J10/14</td>
<td>100</td>
<td>460</td>
<td></td>
</tr>
<tr>
<td>I</td>
<td>J16/5</td>
<td>89</td>
<td>630</td>
<td></td>
</tr>
<tr>
<td>I</td>
<td>J17/10</td>
<td>96</td>
<td>590</td>
<td></td>
</tr>
</tbody>
</table>

Column A, relative amounts of mAb bound to fixed cells at 100 μg mAb/ml, expressed as a percentage of the amount bound at 200 μg mAb/ml. Column B, arbitrary units of fluorescent second antibody bound after treatment of cells with 200 μg mAb/ml.
Figure 5. Competitive binding using $^{125}$I-J10/14. $^{125}$I-J10/14 (20 µg) was mixed with the indicated micrograms of each unlabeled antibody and then sperm were added. Bound J10/14 was determined with a gamma counter (Trimmer et al., 1987). Each point is the average of three samples.

I and Fig. 5). An experiment was performed to determine if J10/14 or J16/5 could override J18/2 to yield the maximum mAb-inducible level of H1 phosphorylation. Cells were diluted into J18/2 at 100 µg/ml and after 5 min, either J10/14 or J16/5 was added to concentrations ranging from 0 to 100 µg/ml (5 µg/ml J10/14 is fully inductive). At 100 µg/ml, J18/2 does not induce the translocation and shedding of surface proteins (Trimmer and Vacquier, 1988). The results (Fig. 6) show that J18/2 at 100 µg/ml induced only a small amount of H1 phosphorylation that was not increased by later addition of J10/14 or J16/5 (as might be expected from Fig. 5). Treatment of sperm with either 100 µg/ml J10/14 or J16/5 alone gave high levels of H1 phosphorylation, as did egg jelly (Fig. 6). These data indicate that the blocking of the common binding site for these three group I mAbs by noninductive concentrations of mAb J18/2 disrupts the ability of both J10/14 and J16/5 to induce H1 phosphorylation.

An additional experiment was performed to determine if pretreatment of cells with J18/2 would alter the concentration dependence of FSG-induced H1 phosphorylation. As shown in Fig. 7, pretreatment with 100 µg/ml J18/2 completely inhibits the FSG-induced phosphorylation of H1. In contrast, the binding of the group II mAb J4/4 has no effect on the concentration dependence of the FSG-induced phosphorylation, relative to FSG alone. These data indicate that the site recognized by the group I mAb plays a role not only in the mAb-induced phosphorylation, but also in the phosphorylation of H1 induced by FSG.

**mAb Activation of Adenylate Cyclase**

To determine the relationship between the activation of adenylate cyclase and H1 phosphorylation, sperm were treated with these six mAbs or egg jelly, washed, lysed, and the adenylate cyclase activity was determined. Six such experiments are presented in Table II, the enzyme rates being calculated from the linear slope of six time points. The fold increase in enzyme activity induced by egg jelly alone ranged from 2.49 to 13.45, with a mean of 6.79. The group I mAbs stimulate adenylate cyclase activity to the same level as does egg jelly, whereas the group II mAbs (and wheat germ agglutinin [WGA]) failed to induce these large increases in adenylate cyclase activity. We have previously shown that the phosphorylation of H1 by the endogenous protein kinase is cAMP dependent (Porter et al., 1988a). The ability of group I mAbs, but not group II mAbs, to activate adenylate cyclase is consistent with the idea that increased levels of cAMP activate the kinase phosphorylating histone H1 (Porter and Vacquier, 1986; Porter et al., 1988a).

**Discussion**

Sea urchin spermatozoa are ideal for studying the mechanism of signal transduction. They possess a large membrane...
Table II. Activation of Adenylate Cyclase with Egg Jelly and mAb

<table>
<thead>
<tr>
<th>Exp. No.</th>
<th>Group</th>
<th>mAb* or Jelly‡</th>
<th>Fold increase in rate over no jelly§</th>
<th>Percent of rate of jelly-treated sample</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Jelly</td>
<td>I J18/2</td>
<td>3.67</td>
<td>100</td>
</tr>
<tr>
<td></td>
<td></td>
<td>II J4/4</td>
<td>7.46</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>Jelly</td>
<td>I J10/14</td>
<td>1.45</td>
<td>19</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J10/14</td>
<td>7.58</td>
<td>102</td>
</tr>
<tr>
<td>3</td>
<td>Jelly</td>
<td>I J10/14</td>
<td>2.49</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>II J4/4</td>
<td>0.92</td>
<td>37</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J10/14</td>
<td>2.66</td>
<td>107</td>
</tr>
<tr>
<td>4</td>
<td>Jelly</td>
<td>I J16/5</td>
<td>13.45</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J17/10</td>
<td>11.57</td>
<td>118</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J18/27</td>
<td>7.30</td>
<td>74</td>
</tr>
<tr>
<td>5</td>
<td>Jelly</td>
<td>WGA1</td>
<td>1.75</td>
<td>13</td>
</tr>
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<td></td>
<td>I J18/2</td>
<td>13.36</td>
<td>99</td>
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<td></td>
<td>I J18/2*</td>
<td>11.57</td>
<td>86</td>
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<tr>
<td>6*</td>
<td>Jelly</td>
<td>I J4/4</td>
<td>3.84</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td>II J18/27</td>
<td>1.79</td>
<td>47</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J18/2</td>
<td>0.67</td>
<td>17</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J10/14</td>
<td>4.07</td>
<td>106</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J10/14</td>
<td>4.47</td>
<td>116</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J16/5</td>
<td>3.94</td>
<td>102</td>
</tr>
<tr>
<td></td>
<td></td>
<td>I J17/10</td>
<td>3.94</td>
<td>102</td>
</tr>
</tbody>
</table>

*mAb concentrations were 100 µg/ml, except in one J18/2 sample in exp. 5 and for all samples of exp. 6, in which the concentrations were 200 µg/ml.
‡ Egg jelly was 100 µg fucose/ml.
§ Basal adenylate cyclase specific activity in lysates of sperm not treated with egg jelly was 0.67 ± 0.38 nmol cAMP formed/min per mg protein (n = 7).
¶ Rates were calculated from the linear slopes of six time points over 18 min.
‖ WGA was 60 µg/ml, which yields 100% inhibition of the FSG-induced acrosome reaction (Podell and Vacquier, 1984).

Surface area (31–35 µm²; Cross, 1983) relative to their volume (8 µm³; Christen et al., 1982; Lee et al., 1983), they can be obtained as pure cells with ease and in vast quantity, and several methods exist to isolate the plasma membrane (reviewed in Vacquier, 1987). Egg jelly and its FSG component can be obtained from eggs with ease and its effects on sperm occur synchronously in a time span of seconds. In addition to FSG, egg jelly contains such small peptides as speract (Bentley et al., 1987) and speract (Schackmann and Chock, 1986; reviewed in Garbers et al., 1987) that bind to specific sperm surface receptors and alter cyclic nucleotide metabolism and ion channel activity. G proteins have also been reported in sperm membranes (Kopf et al., 1986; Bentley et al., 1986), and most recently GTP has been shown to control K⁺ channels in the flagellar membrane of sea urchin sperm (Lee, 1988). The FSG component of egg jelly is a very large, complex molecule (Segall and Lennarz, 1979, 1981; Garbers et al., 1983; DeAngelis and Glabe, 1987), and the mechanism by which it interacts with the sperm surface remains unknown. Because of the complexity of FSG, classical methods to identify the membrane proteins with which it interacts have been untenable. One approach to the analysis of such complex systems is to use antibodies as probes to identify proteins mediating signal transduction. Using this approach, we have shown that the 210-kD protein plays a critical role in transduction of the FSG response (Trimmer et al., 1985, 1986, 1987).

In this study, we show that the domain of the 210-kD protein defined by the group I mAbs activates adenylate cyclase and induces the cAMP-dependent phosphorylation of H1. We propose that these mAbs use the same pathway as does FSG in inducing these changes when sperm contact eggs. The precise mechanism by which binding of mAb to the group I domain activates adenylate cyclase is of general interest, and most likely involves a conformational change in the 210-kD protein that activates Ca²⁺ channels (Trimmer et al., 1986). The activation of the cyclase is strictly dependent on the presence of external Ca²⁺ (Garbers and Kopf, 1980). However, ionophore A23187 (2–40 µM) does not induce HI phosphorylation in sperm (Porter and Vacquier, 1986), but such negative data are inconclusive because ionophores act nonspecifically on cellular compartments and thus alter intracellular Ca²⁺ concentrations unphysiologically.

The three group I mAbs tested compete equally well for binding to cells (Fig. 5), and also show the same concentration dependencies for inhibition of the FSG-induced acrosome reaction (Fig. 4 in Trimmer and Vacquier, 1986). These mAbs activate adenylate cyclase but do not induce the acrosome reaction indicates that the cAMP-dependent protein kinase activity phosphorylating HI is not directly involved in the acrosome reaction. The group II mAbs (and WGA), although they bind to cells as avidly as group I mAbs, neither activate adenylate cyclase nor induce HI phosphorylation; they provide the proof that only certain domains of the 210-kD protein mediate signal transduction across the cell membrane. Additional understanding of the inductive and noninductive domains of the 210-kD protein will come from molecular cloning.

A separate topic of discussion is the possible biological significance of the phosphorylation of HI by the endogenous kinase on only one seryl residue in a single peptide (Porter et al., 1988a). Sea urchin sperm-specific histone HI is a unique type of HI possessing an extended NH₂-terminal tail with an unusual amino acid sequence (reviewed in Poccia, 1987). Within 10 min after fertilization, sperm HI becomes hyperphosphorylated (~10 phosphates/mol, Green and Poccia, 1985; Porter and Vacquier, 1988; Porter et al., 1988b), and lost from the sperm pronuclear chromatin, having been replaced by cleavage stage HI present in the egg cytoplasm (Green and Poccia, 1985; Poccia, 1987). In vitro labeling of HI in detergent permeabilized sperm shows that the endogenous kinase is completely cAMP dependent. Reactions with purified sperm HI and rabbit muscle cAMP-dependent protein kinase show a maximum stoichiometry of 1 mol phosphate incorporated per mol HI (Porter et al., 1988b). We can conclude from these data that there is only one site for phosphorylation by cAMP-dependent protein kinase in native sea urchin sperm HI. It is possible that this single phosphorylation event of HI, occurring by action of the FSG-induced kinase, may cause a change in chromatin structure that could expose other sites on HI to be phosphorylated by cAMP-independent kinases of the egg (Porter et al., 1988b), which would result ultimately in loss of sperm HI from the male pronucleus (Green and Poccia, 1985). Regardless of the rea-
son for its existence, this is the only system in which the phosphorylation of H1 in vivo occurs on a single defined site. Whether it has structural consequences to the chromatin remains to be determined.

Regarding the sperm histone kinase, it could be located outside the nucleus and then move through the nuclear envelope after activation, or it could be resident in the nucleus being packaged with the highly condensed chromatin. One surprising fact about its action is that it is so specific for H1. One would assume that of the many score of different proteins in the nucleus, the preferred sequence (Kemp et al., 1977) for phosphorylation of R/K-X-S (in which X is any amino acid) would be present in many other proteins that would label if the kinase were freely diffusing in the nucleus. The H1 kinase may be resident in the nucleus, tightly associated with H1, and thus inaccessible to other substrates. Regardless of the above speculation, this system is of value in studying how signals at the cell surface are transduced across the cell membrane and through the nuclear envelope to be ultimately expressed by the action of a protein kinase on a protein involved in chromatin structure. As recently discussed (Rauscher et al., 1988), little is yet known about this fundamental process that is central to our understanding of cellular activation.

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