Regulation of mammalian spliceosome assembly by a protein phosphorylation mechanism

Jacqueline E.Mermoud, Patricia T.W.Cohen¹ and Angus I.Lamond2

European Molecular Biology Laboratory, Meyerhofstrasse 1, Postfach 102209, D-69012 Heidelberg, Germany and 'MRC Protein Phosphorylation Unit, Department of Biochemistry, University of Dundee, Dundee, DD1 4HN, UK

2Corresponding author

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Splicing of mRNA precursors (pre-mRNA) is preceded by assembly of the pre-mRNA with small nuclear ribonucleoprotein particles (snRNPs) and protein factors to form a spliceosome. Here we show that stimulat-
ing Ser/Thr-specific protein dephosphorylation ing Ser/Thr-specific protein dephosphorylation selectively inhibits an early step during mammalian spliceosome assembly. Treatment of HeLa nuclear splicing extracts with human protein phosphatase ¹ (PP1) expressed in Escherichia coli, or PP1 purified from rabbit skeletal muscle, prevents pre-spliceosome E complex (early complex) formation and stable binding of U2 and U4/U6.U5 snRNPs to the pre-mRNA. PP1 does not inhibit splicing catalysis if added after spliceosome assembly has taken place. Addition of purified SR protein splicing factors restores spliceosome formation and splicing to PP1-inhibited extracts, consistent with SR proteins being targets regulated by phosphorylation. These data extend earlier observations showing that splicing catalysis, but not spliceosome assembly, is blocked by inhibiting protein phosphatases. It therefore appears that pre-mRNA splicing, in common with other biological processes, can be regulated both positively and negatively by reversible protein phosphorylation.

Key words: protein phosphatase/protein phosphorylation/ RNA splicing/spliceosome assembly/SR proteins

Introduction

Splicing of nuclear mRNA precursor transcripts (premRNA) involves ^a two-step reaction mechanism and takes place in a dedicated complex termed a spliceosome (for reviews, see Green, 1991; Guthrie, 1991; Ruby and Abelson, 1991; Moore et al., 1993; Rymond and Rosbash, 1993). The function and composition of spliceosomes have been studied in both mammalian and yeast in vitro splicing systems. In both systems they comprise three major RNA-protein subunits, the Ul, U2 and U4/U6.U5 small nuclear ribonucleoprotein particles (snRNPs; reviewed by Guthrie and Patterson, 1988; Lührmann et al., 1990), and additional non-snRNP protein splicing factors (reviewed by Lamm and Lamond, 1993). Spliceosomes assemble on pre-mRNA substrates in a stepwise manner, involving the formation of distinct pre-splicing complexes that represent functional intermediates in the pathway. The order of assembly appears to be obligatory, with U1 snRNP binding to the ⁵' splice site, U2 snRNP interacting with the ³' end of the intron prior to U4/U6.U5 assembly (reviewed by Lamond, 1993; Moore et al., 1993). In yeast and mammals, the ⁵' and ³' splice sites are recognized and functionally associated with one another in one of the earliest characterized pre-splicing complexes containing U¹ snRNP, but not U2 or U4/U6.U5 snRNPs (Reed, 1990; Michaud and Reed, 1991, 1993; Rosbash and Seraphin, 1991; Jamison et al., 1992).

As well as the snRNP subunits, a number of nonsnRNP protein splicing factors play important roles during mammalian spliceosome assembly. These include a group of proteins that share a common motif consisting of repeating serine-arginine dipeptides (Bimey et al., 1993). There is evidence that most, if not all, proteins with the SR motif are phosphorylated in vivo (Wooley et al., 1988; Roth et al., 1990; Woppmann et al., 1993; Gui et al., 1994). A subset of SR-containing splicing factors, including SRp2O, ASF/SF2, SC-35, SRp4O, SRp55 or SRp75, are recognized through a shared phosphoepitope by the same monoclonal antibody, called mAb104 (Roth et al., 1990, 1991; Zahler et al., 1992). This group of proteins can be co-purified using a two-step salt precipitation method (Zahler et al., 1992). The characteristic serinearginine repeat motif is also found in several other splicing factors that are not detected by mAb104 and do not copurify with the other SR proteins. These include both subunits of the non-snRNP splicing factor U2AF (i.e. U2AF⁶⁵ and U2AF³⁵) and the U1 snRNP-specific 70 kDa protein (Theissen et al., 1986; Query and Keene, 1987; Spritz et al., 1987; Zamore et al., 1992; Zhang et al., 1992). To distinguish between these SR-containing splicing factors, we will refer to the proteins recognized by mAblO4 as SR104 proteins.

Although its detailed function in the splicing mechanism is not yet known, mutational studies have shown that the SR domain in U2AF⁶⁵ and in ASF/SF2 is required for their activity as essential splicing factors (Zamore et al., 1992; Caceres and Krainer, 1993; Zuo and Manley, 1993). U2AF, SR104 proteins and U1 snRNP are all required for spliceosome assembly or the stabilization of early presplicing complexes (Chabot and Steitz, 1987; Frendewey et al., 1987; Zillmann et al., 1987; Barabino et al., 1990; Krainer et al., 1990; Fu et al., 1992; Mayeda et al., 1992; Zahler et al., 1992, 1993; Zamore et al., 1992). For example, SR104 proteins can commit pre-mRNAs to the splicing pathway (Fu, 1993). ASF/SF2 has also been shown to cooperate with U¹ snRNP in binding pre-mRNA (Khotz et al., 1994), and protein-protein interaction between SC35 (ASF/SF2) and other SR proteins has been demonstrated (Wu and Maniatis, 1993).

Fig. 1. Protein phosphatases inhibit pre-mRNA splicing and splicing complex formation. A ^{32}P -labelled Ad1 pre-mRNA substrate was incubated for 120 min at 30°C under standard splicing conditions in the absence or presence of recombinant or purified catalytic subunits of Ser/Thr specific protein phosphatases. (A) and (B) The lanes marked 'Ctrl' correspond to splicing reactions carried out in the absence of phosphatase (lanes 1 and 3). In lane 2, human PPIy expressed in E.coli (0.1 U/µl) was added to the assay from the beginning of the reaction. In lanes 4, PPIy (0.1 U/µl) was added 20 min after the start of the reaction. (C) Human PPly expressed in E.coli (0.15 U/ μ l; lane 2), purified PPI catalytic subunits from rabbit skeletal muscle (0.5 U/µl; lane 3) and PP2A (0.7 U/µl; lane 4) were added at the beginning of the reaction. 'Ctrl' (lanes 1 and 4), no phosphatase was added. (A) and (C) The splicing products of Adl pre-mRNA were resolved on ^a 10% polyacrylamide/8 M urea gel. The structures of the premRNA, splicing intermediates and products are indicated by cartoons with exons indicated as solid boxes and introns as lines. Markers are endlabelled MspI-digested pBR322 fragments ('Mrks'). (B) Spliceosome complexes were separated on an agarose-polyacrylamide composite gel. Prespliceosome (A complex) and spliceosome complexes (B and C complexes) are indicated with arrows. 'N' represents non-specific complexes.

Recent data have indicated that inhibiting protein dephosphorylation on serine or threonine residues blocks the catalytic steps of splicing, but not spliceosome assembly (Mermoud et al., 1992; Tazi et al., 1992, 1993). HeLa nuclear splicing extracts contain both protein phosphatase ¹ (PPI) and PP2A activities (Mermoud et al., 1992). When these activities are inhibited by treatment with the specific phosphatase inhibitors okadaic acid, tautomycin or microcystin-LR, inactive spliceosomes are formed containing U1, U2 and U4/U6.U5 snRNPs (Mermoud et al., 1992; Tazi et al., 1992). Thiophosphorylation of the U¹ snRNP-specific 70 kDa protein causes a similar effect (Tazi et al., 1993). Splicing can be reactivated in extracts treated with phosphatase inhibitors by adding exogenous catalytic subunits of PP1 or PP2A (Mermoud et al., 1992). There is also evidence that forms of PPI and PP2A may differentially affect the first and second catalytic steps of splicing, indicating that at least two, and possibly more, separate dephosphorylation events are required for catalysis.

Here we show that stimulating specific Ser/Thr protein dephosphorylation events inhibits spliceosome formation at an early step of the assembly pathway in HeLa splicing extracts. We also present evidence that the phosphorylation

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state of the SR104 family of protein splicing factors may be crucial for their function in spliceosome assembly.

Results

Protein phosphatases inhibit spliceosome assembly

The effects of Ser/Thr-specific protein phosphatases on pre-mRNA splicing were investigated in HeLa nuclear extracts (Figure 1). Treatment of the extract with human PPI, isoform y, expressed in Escherichia coli and added from the beginning of the assay, potently inhibits splicing (Figure IA, lane 2). A parallel analysis revealed that PPIy inhibited early steps in spliceosome assembly (Figure 1B). Lane ¹ (panel B) shows the typical pre-splicing complex $(A \text{ complex})$ and spliceosomes $(B + C \text{ complexes})$ formed in a control splicing assay which are abolished in extracts treated with PPly (Figure 1B, lane 2). All pre-mRNAs tested were similarly inhibited by PP1y in a concentrationdependent manner (data not shown). No spliced products were observed even after long incubation times, indicating that the inhibitory effect does not reflect a reduction in the kinetics of spliceosome assembly (data not shown). Inhibition of splicing was also observed using PP^I catalytic

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Fig. 2. The phosphatase-mediated block of spliceosome formation is sensitive to phosphatase inhibitors. The inhibition of splicing complex assembly caused by phosphatase treatment was prevented when PPl'y was pre-treated with microcystin-LR. (A) and (B) Lanes 1 and 5 show control splicing reactions with AdI pre-mRNA carried out in the absence of inhibitor. The splicing extract was treated with either microcystin-LR (0.6 μ M; lane 2) or with with PP1 γ (0.1 U/ μ l; lane 3). Lane 4 corresponds to the addition of pre-mixed PPly/microcystin-LR to the splicing reaction. (A) shows the separation of RNA intermediates and products on a denaturing gel, while (B) shows a parallel analysis of splicing complex assembly on a native agarose-polyacrylamide composite gel.

subunits purified from rabbit skeletal muscle (Figure IC, lane 3). Similarly, the catalytic subunit of PP2A, which has overlapping substrate specificity with PP1, also blocked splicing (Figure 1C, lane 4) and spliceosome assembly (data not shown).

When PP1y was added to the splicing assay after 20 min, catalysis (Figure 1A, lane 4) and complex assembly (Figure lB, lane 4) were still observed. Parallel analysis confirmed that at this time spliceosome assembly, but not the first catalytic step of splicing, has occurred (data not shown). The simplest interpretation of this result is that dephosphorylation events inhibit spliceosome assembly, but do not block splicing if complex formation has already occurred.

If inhibition of spliceosome assembly is caused by protein dephosphorylation, it should be alleviated by pretreatment of the phosphatase with a specific phosphatase inhibitor. To test this, control reactions (Figure 2A and B, lanes ¹ and 5) were compared with splicing reactions in the presence of the phosphatase inhibitor microcystin-LR (Figure 2A and B, lanes 2) or $PPI\gamma$ (Figure 2A and B, lanes 3) and with reactions containing pre-mixed PP1 γ / microcystin-LR (Figure 2A and B, lanes 4). Microcystin-LR is ^a specific phosphatase inhibitor that binds irreversibly to the catalytic subunits of PP1 and PP2A (MacKintosh et al., 1990). It inhibits splicing, but allows assembly of splicing complexes (Mermoud et al., 1992; Figure 2A and B, lanes 2). In contrast, addition of PPl y abolishes splicing (Figure 2A, lane 3) and splicing complex formation

Fig. 3. ATP levels are stable in extracts treated with PPly or microcystin-LR. TLC analysis of α -³²P-labelled ATP incubated in HeLa nuclear extract under splicing conditions either in the absence or presence of splicing inhibitors. Lanes 1 and 5, $[\alpha^{-32}P]ATP$; lanes 2-4 correspond to different time points of a splicing reaction incubated without exogenous cold ATP or creatine phosphate (CP). ATP levels are stable in a standard control splicing reaction (lane 6) and in extracts where splicing was blocked by treatment with either human PP1 γ expressed in *E.coli* (0.3 U/µl; lane 7) or the phosphatase inhibitor microcystin-LR $(1 \mu M; \text{lane } 8)$, but not in the presence of alkaline phosphatase (0.05 U/ μ l; lanes 9 and 10). The samples of the standard splicing reactions including CP were analysed after ¹ h of incubation, the sample treated with alkaline phosphatase was also analysed at 10 min (lane 10). Parallel analysis confirmed that splicing was blocked in the extracts treated with PPly (lane 7), microcystin-LR (lane 8) and alkaline phosphatase (lanes 9 and 10), but not in the untreated control sample (lane 6; data not shown). The position of ATP and ADP is indicated with arrows; the identities of the other hydrolysis products were not investigated further.

(Figure 2B, lane 3). When PPly was pre-incubated with microcystin-LR, spliceosome formation is detected but catalysis is severely reduced (Figure 2A and B, lanes 4). This confirms that splicing inhibition is a specific effect of the phosphatase.

In summary, the data show that Ser/Thr protein dephosphorylation events block pre-mRNA splicing in vitro by specifically preventing spliceosome assembly.

PP1 does not affect ATP levels

ATP is required for spliceosome assembly and pre-mRNA splicing. To clarify that the inhibitory effects of protein phosphatases and phosphatase inhibitors do not arise indirectly through depletion of ATP from the extract, we analysed the levels of $[\alpha^{-32}P]ATP$ in splicing extracts by TLC (Figure 3). ATP levels are stable during ^a ^I h incubation under standard splicing conditions (Figure 3, lane 6) and in extracts treated with either $PP1\gamma$ (Figure 3, lane 7) or microcystin-LR (Figure 3, lane 8) at concentrations that inhibit splicing. In contrast, when alkaline phosphatase was added to the splicing assay, all the labelled ATP was hydrolysed within ¹ h (Figure 3, lane 9) and $\sim 50\%$ in the first 10 min (Figure 3, lane 10). Lanes 2-4 correspond to HeLa splicing extracts incubated without creatine phosphate (CP). These conditions do not support splicing and the data show that $>50\%$ of the ATP

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Fig. 4. An early step of splicing complex formation is affected by PP1 treatment. (A) In the presence of PP1, no pre-splicing E complex can be detected by gel filtration. Splicing reactions in the absence (untreated control) or presence of protein phosphatase (PP1; 0.1 U/µl) or in the presence of pre-mixed phosphatase with phosphatase inhibitor (PP1 + microcystin-LR; 0.1 U/ μ l + 1 μ M) were fractionated by gel filtration (see Materials and methods). The peaks containing E and H complexes are indicated. (B) Only U1 snRNP forms a stable complex with pre-mRNA after PP1 γ treatment. Affinity selection of snRNAs from splicing complexes assembled in the absence (lanes 2-4 and 6) or presence (lane 5) of PPIy (0.6 U/ gl). Nuclear extract was pre-incubated at 30'C for ¹⁰ min to hydrolyse the endogenous ATP. Splicing reactions were carried out for 40 min. The affinity selection was performed using biotinylated antisense 2'-O-Me RNA oligonucleotides targeted to the Ad1 pre-mRNA (Ryder et al., 1990). Affinity-selected RNAs were separated on ^a 10% denaturing polyacrylamide gel and analysed by Northern hybidization using probes complementary to each of the five spliceosomal snRNAs. Marker lane corresponds to unselected total HeLa nuclear RNA (lane I). Lane ² corresponds to incubation of pre-mRNA in splicing reactions without an antisense oligonucleotide, lane ³ shows the level of RNA selected in the absence of precursor, and lanes 4-6 correspond to reactions with both pre-mRNA and oligonucleotide present. In sample 6 no ATP/CP was added to the reaction.

is hydrolysed within 2 min (Figure 3, lane 2) and all of it within 20 min (Figure 3, lane 3). Additional experiments show that exogenously added creatine phosphokinase does not restore splicing or splicing complex formation to PP1-blocked extracts (data not shown). Increasing ATP concentration in the assay also failed to restore splicing to a PP1-treated extract (data not shown).

An early step of splicing complex formation is blocked in PPI-treated extracts

As judged by native gel electrophoresis, no pre-splicing A complex (which contains U1 and U2 snRNPs) can be detected in PPl -treated extracts (Figure 1B; see lanes ¹ and 2). Therefore, we addressed whether the pre-spliceosome E (early) complex (Reed, 1990; Michaud and Reed, 1991, 1993), which forms before A complex, could be detected after PPI treatment. E complex can be assayed by gel filtration as an additional peak eluting prior to hnRNP containing H complexes. We observed E complex in both the untreated control extract and in a control reaction where $PP1\gamma$ was inhibited by microcystin-LR (Figure 4A). In contrast, only H complexes form in PPly-treated extracts (Figure 4A).

to determine whether splicing snRNPs could still bind stably to pre-mRNA (Figure 4B). SnRNP binding to premRNA in either ^a control extract, or in extracts treated with PP1y, was studied by affinity selection using a biotinylated anti-pre-mRNA 2'-O-alkyl oligoribonucleotide (Ryder et al., 1990). Each of the U1, U2, U4, U5 and U6 snRNAs are detected in splicing complexes formed within 40 min in the control extract (Figure 4B, lane 4). The specificity of binding is confirmed by control experiments carried out in the absence of either antisense probe (Figure 4B, lane 2) or in the absence of pre-mRNA (Figure 4B, lane 3). When splicing complex assembly was inhibited by treatment with PPIy, Ul snRNP showed a higher level of stable complex formation with the premRNA (Figure 4B, lane 5), while the levels of the other four spliceosomal snRNAs were specifically reduced, relative to the untreated control extract (Figure 4B, lane 4). A similar pattern is observed in ^a control assay where ATP was omitted (Figure 4B, lane 6). Additional analysis showed that the unselected snRNAs were not degraded (data not shown).

Splicing extracts treated with PPly were analysed further

We conclude that the inhibitory effect of PP1 γ on

Restoration of splicing in PP1-inhibited extract

Having established that PP1y treatment inhibits an early step in the spliceosome assembly pathway, we next sought to identify conditions that could restore splicing activity to inhibited extracts. Splicing was assayed either in the absence of protein phosphatase (Figure 5, lanes 1-5) or in the presence of PP1 γ (Figure 5, lanes 6-9). Addition of untreated nuclear extract to the PPly-blocked extract restored splicing (Figure 5, see lanes 6 and 7) when added in low amounts which alone could not support splicing (Figure 5, lane 5). To address whether an RNA moiety was important for complementation, nuclear extract was pre-treated with micrococcal nuclease (MCN). The MCNtreated extract showed no splicing activity (Figure 5, lane 3), while ^a control extract treated with MCN under conditions where the nuclease is inactive does splice the pre-mRNA (Figure 5, lane 2). Addition of MCN extract, or MCN control extract, to the PP1y-blocked extract restored splicing (Figure 5, lanes ⁸ and 9). The MCN control extract restored splicing even when used in an amount that was too low to splice on its own (Figure 5, lane 4). We conclude that one or more MCN-resistant factors in nuclear extract can activate splicing in PP1 treated extracts.

SR104 proteins restore splicing in PP1-inhibited extract

Since SR104 splicing factors are known to be phosphoproteins that have essential functions early in splicing complex assembly, we tested whether their addition to PP1y-treated extracts would restore splicing. SR104 proteins were prepared from HeLa cells as described by Zahler et al. (1992), their purity confirmed by SDS-PAGE analysis and their function demonstrated by complementation of splicing in a HeLa S1OO extract (data not shown). Addition of purified SR104 proteins to a PPlyinhibited splicing extract restored spliceosome assembly (Figure 6B, see lanes ³ and 4) and mRNA formation (Figure 6A, see lanes 3 and 4) to a level comparable with the untreated control (Figure 6A and B, see lanes 4 and 1). The level of splicing/splicing inhibition could be manipulated by varying the concentration of exogenous $SR104$ proteins and PP1 γ , respectively. Thus, increasing the amount of PPly requires a corresponding increase in the amount of SR104 proteins to achieve a comparable level of complementation (data not shown). Furthermore, SR104 proteins enhanced splicing of adeno pre-mRNA and complex formation in untreated HeLa nuclear extract (Figure 6A and B, lanes 2). These conditions of increased splicing activity (Figure 6C, see lane ¹ with lanes 2 and 7) are still sensitive to phosphatase treatment (Figure 6C, lanes 3-6). Differences were observed between separate HeLa extract preparations, both in their splicing efficiency and in the level of splicing enhancement obtained upon

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Fig. 5. MCN-treated nuclear extract restores splicing to a PP1yblocked extract. Standard splicing assays in the absence (lanes 1-5) or presence (lanes 6-9) of PP1 γ . Unless otherwise stated, 7 μ l of extract are included in the reactions. Lane ¹ corresponds to an untreated control splicing reaction, lane ² to ^a MCN control extract (see Materials and methods). Lane 3 shows a standard splicing reaction with MCN-treated extract, lanes 4 and 5 show splicing assays performed using $2.5 \mu l$ of MCN control extract and $2 \mu l$ of untreated control extract, respectively. PP1 γ treatment (0.03 U/µl) shown in lanes 6-9 was performed by pre-incubation of the extract (42 μ l/ 3.7 U enzyme) for ¹⁰ min on ice. Extracts tested for complementation were added at the beginning of the splicing assay: water (lane 6); $2 \mu l$ of standard HeLa nuclear extract (lane 7); 2.5 pl of MCN control extract (lane 8) and 3 μ l of MCN-treated extract (lane 9). Low amounts of extract that alone were not sufficient to support splicing are marked with an asterisk.

adding exogenous SR 104 proteins. However, in all cases treatment with PPIy inhibited splicing.

Phosphorylation state of SR104 proteins is affected by PP1

If endogenous SR104 proteins in HeLa extracts are direct targets for the PPl-mediated inhibition of spliceosome assembly, their phosphorylation state should be altered upon PP1 treatment. To test this, SR104 proteins were detected in HeLa nuclear extracts incubated under splicing conditions using an SR104-specific monoclonal antibody (mAblO4; Roth et al., 1990, 1991; Zahler et al., 1992) that recognizes a phosphoepitope (Figure 7). In an immunoblot, the untreated control extract shows the characteristic pattem of SR104 proteins (Figure 7, lane 1; see Zahler et al., 1992). After PP1y treatment, only a band at \sim 30 kDa is detected, running with a slightly altered mobility to the doublet of \sim 30 kDa in the untreated control (Figure 7, see lanes ¹ and 2). This pattern was observed at early time points and persisted until the end of the reaction with no evidence of enhanced protein degradation

Fig. 6. Purified SR104 proteins restore splicing activity to ^a PP1y-blocked extract. The inhibition of splicing complex assembly caused by PPIy treatment was relieved by the addition of purified HeLa SR104 proteins in ^a concentration-dependent manner. (A) and (B) Splicing reactions carried out in the absence (lanes 1 and 2) or presence (lanes 3 and 4) of PP1 γ (0.03 U/µl). One microgram of purified SR 104 proteins/20 µl reaction was added at the start of the assay in the case of samples 2 and 4. (C) Standard splicing in an untreated control extract (lane 1) and in the presence of exogenously added SR104 proteins (1 μ g SR104/20 μ l, lanes 2-7). An increasing amount of PP1 γ was added to the SR104-treated extracts: 0.0125 U/µl (lane 3), 0.025 U/µl (lane 4), 0.05 U/µl (lane 5) and 0.1 U/µl (lane 6). (A) and (C) Analysis of splicing products by denaturing gel electrophoresis, cartoons indicating the identity of splicing intermediates and products. (B) Analysis of splicing complexes by native gel electrophoresis.

Fig. 7. Endogenous SR 104 proteins in HeLa nuclear extract are dephosphorylated upon PPIy treatment. The phosphorylation state of SR104 proteins was analysed by probing Western blots with mAblO4. An untreated splicing reaction control (lane 1) is compared with reactions containing extracts that were treated with PP1 γ (0.03 U/µl; lanes 2 and 3) and, in the case of sample 3, where splicing was restored by addition of exogenous HeLa SR104 proteins $(1.1 \mu g)$ SR104/20 µl reaction). Phosphatase treatment was performed by preincubation of the extract with PPIy for ¹⁰ min on ice. Purified SR 104 proteins were added after 5 min of incubation under standard splicing conditions. All samples shown were incubated for 70 min at 30°C.

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(data not shown). In contrast, mAblO4 could detect all the SR104 proteins in control extracts without exogenous PPl throughout the splicing reaction (data not shown). Parallel analysis demonstrated that the pre-mRNA was processed in the control extract, but not in the PPlytreated extract (data not shown). The usual pattem of SR104 proteins revealed by mAblO4 is detected again when splicing is restored to a PPly-treated extract by addition of exogenous SR104 proteins (Figure 7, lane 3). Similarly, additional experiments showed that PP1 γ also dephosphorylates purified SR 104 proteins (Mermoud, 1994; and data not shown). We conclude that PPly treatment removes the mAb ¹⁰⁴ phosphoepitope from most SR104 proteins and that this correlates with the inhibition of spliceosome assembly. There may also be changes in phosphorylation at other sites in SR104 proteins which are not recognized by mAbIO4.

The data are consistent with SR 104 proteins being direct targets for regulation mediated by PPl. This predicts that addition of the specific phosphatase inhibitor microcystin-LR to PPly-treated extract should, if the extract contains the appropriate kinase activities: (i) change the phosphorylation state of the endogenous SR104 proteins and (ii) restore spliceosome assembly to the PP1 γ -treated extract. Parallel Westem blot and native gel analyses confirm this to be the case (Figure 8). Thus, $PP1\gamma$ treatment

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Fig. 8. Addition of microcystin-LR to PPly-treated extract results in rephosphorylation of SR104 proteins and restoration of spliceosome assembly. (A) Detection of endogenous SR proteins by probing Western blots with mAblO4. Samples were analysed after either 25 min (lanes 1-5), 55 min (lane 9), 85 min (lanes 6-8, 10, 12) or 120 min (lane 11) after starting the splicing assay. An untreated splicing control (lanes ¹ and 6) was compared with a sample where PPIy was added either after 25 min of incubation (lanes 2 and 7); or from the beginning of the reaction (lanes 3, 4, 8-11), or was pre-mixed with microcystin-LR and then added at the beginning of the reaction (lanes ⁵ and 12). We also tested whether SR proteins become rephosphorylated in extracts treated with PPly if the phosphatase is subsequently inactivated (samples marked 'D'). In this case, the extract was incubated for 25 min with PP1 γ (lane 4), then microcystin-LR was added to the extract to inhibit the phosphatase and the samples analysed at the times shown (lanes 9-11). (B) Native gel analysis of splicing complexes formed in samples treated with PPly and microcystin-LR as described in (A). The labels at the top of each lane indicate the corresponding samples analysed in (A) and (B). Samples were analysed after either 25 and 45 min (lanes 1-3, 7 and other data not shown), 55 min (lane 4), 85 min (lane 5) or 120 min (lane 6) incubation at 30°C.

removes phosphoepitopes recognized by mAb104 (Figure 8A, lanes 3, 4, 7 and 8) and inhibits splicing complex formation (Figure 8B, lane 3). Subsequent addition of microcystin-LR restores the phosphoepitope (Figure 8A, lanes 9-11) and spliceosome formation (Figure 8B, lanes 4-6). These data also demonstrate that the HeLa nuclear splicing extracts contain kinase activity(ies) that can phosphorylate the SR104 proteins at the epitope recognized

Fig. 9. The cartoon illustrates a model showing distinct steps where splicing may be regulated by ^a protein phosphorylation mechanism. Phosphorylation is required for early steps in spliceosome assembly (this study). Possible targets for this effect include SR104 protein splicing factors. Once a spliceosome containing Ul, U2, U4/U6.U5 snRNPs is assembled, at least two separate dephosphorylation events are required for catalysis, with PP1 needed for the first catalytic step and PP2A (and possibly also PPI) for the second catalytic step (Mermoud et al., 1992; Tazi et al., 1992). Dephosphorylation of the U1 70 kDa protein may be required for the first catalytic step since its thiophosphorylation blocks splicing, but not spliceosome assembly (Tazi et al., 1993).

by mAbl04. Addition of PP1y 25 min after starting the assay does not block spliceosome assembly (Figure 8B, lane 2) or catalysis (see Figure 1), but removes the phosphoepitope recognized by mAbl04 (Figure 8A, see lanes 2 and 7).

Discussion

In this study, we have shown that treatment of HeLa nuclear splicing extracts with either human PPl expressed in E.coli, or PP1 purified from rabbit skeletal muscle, selectively inhibits an early step during spliceosome assembly. After PP1 treatment, no pre-splicing E complex can be detected and while Ul snRNP can still form a stable complex with the pre-mRNA, stable binding of U2 and U4/U6.U5 snRNPs is prevented. This contrasts with the effect of adding specific phosphatase inhibitors to the extract, which block splicing catalysis but allow the formation of inactive spliceosomes containing U1, U2 and U4/U6.U5 snRNPs (Mermoud et al., 1992; Tazi et al., 1992). Therefore, as illustrated in Figure 9, protein phosphorylation can affect pre-mRNA splicing either positively or negatively by influencing separate steps in the splicing pathway. These findings are reminiscent of the complex phosphorylation mechanisms that have been shown to regulate many biological processes, including metabolism and the cell division cycle.

Inhibition of splicing by PP1 is a clear effect of protein phosphatase activity since (i) it is sensitive to the specific phosphatase inhibitor microcystin-LR and (ii) inhibition was observed with both purified rabbit muscle PPl and human PP1y purified to homogeneity from a bacterial expression system that is free of contaminating eukaryotic proteins. That PPl does not block catalysis when added after spliceosome assembly underlines the specificity of inhibition. This specificity is further supported by studies on alternative splicing substrates with competing ⁵' splice sites (Cardinali et al., 1994). In conditions where both ⁵' splice sites are active, PP1y treatment can block splicing to the upstream ⁵' splice site without inhibiting splicing to the downstream ⁵' splice site.

A major question that emerges is how protein phosphatases can affect the activity of specific components of the splicing machinery. The catalytic subunits of protein phosphatases are known to have inherent substrate specificities (Cohen, 1989). However, studies on cytoplasmic and nuclear forms of PPl and PP2A have shown that additional regulation of substrate specificity can be conferred upon native enzymes through the presence of targeting subunits (Cohen, 1989; Bollen et al., 1993; Hubbard and Cohen, 1993). It is possible, therefore, that the inhibition of spliceosome assembly is enhanced by the interaction of exogenous PP1 with free targeting subunits in the HeLa nuclear extract. PP1 activity is present in the HeLa nuclear splicing extracts (Mermoud et al., 1992). It will be important now to isolate native PPI from the HeLa nuclear extract and to characterize any potential regulatory subunits that can enhance its activity towards components of the splicing apparatus. Another important task is to establish whether PPl itself, or another protein phosphatase with overlapping substrate specificity, acts upon splicing factors in vivo.

Which protein splicing factors may be the key targets whose activity is modulated by PP1? Leaving aside as yet unidentified splicing factors, there are several candidate phosphoproteins whose activities are required at early steps during spliceosome assembly. These include the SR family of non-snRNP protein splicing factors, the Ul snRNP-specific 70 kDa protein and both subunits of the non-snRNP splicing factor U2AF (i.e. U2AF⁶⁵ and U2A F^{35}). Although it is important to recognize that when dealing with a crude system, such as the HeLa splicing extract, it is difficult to exclude indirect effects, we have presented here several lines of evidence indicating that one or more of the SR104 proteins may be direct targets whose activity is affected by PPI. Not only are they required at an early stage of spliceosome assembly (Krainer et al., 1990, 1991; Ge et al., 1991; Fu and Maniatis, 1992a,b; Zahler et al., 1992, 1993; Fu, 1993), which correlates with the stage at which PPI inhibits, addition of purified SRI 04 proteins is sufficient to restore spliceosome assembly in a PP1-treated extract. SR104 proteins are reported to be reversibly phosphorylated in vivo (Roth et al., 1991; Gui et al., 1994). We show in this study that the phosphorylation state of endogenous SR104 proteins in HeLa nuclear extracts changes when splicing is inhibited by PPI. An interesting comparison can be drawn between the PPI-treated HeLa nuclear extract and HeLa S1OO extracts. In both cases, splicing and spliceosome assembly do not take place and the assembly block appears to occur

at a similar early step. Both extracts can be activated by addition of exogenous, purified SR 104 proteins (this study; Zahler et al., 1992). In S100 extracts, the splicing defect is caused by limiting amounts of endogenous SR104 proteins (Krainer et al., 1990; Fu et al., 1992; Zahler et al., 1993). In the PPl-treated HeLa nuclear extract, we propose that it may instead be the loss of function of SR104 proteins that is responsible for an analogous inhibition.

The phosphorylation state of at least some proteins in the HeLa splicing extract is dynamic, since phosphate groups are continuously added and removed by competing kinase and phosphatase activities. As a consequence, the steady-state level of protein phosphorylation will change rapidly in response to any treatment that affects the relative level of these competing activities. We show here that PPl can dephosphorylate SR104 proteins. Recently, several studies have reported the characterization of kinase activities that phosphorylate proteins with SR domains (Woppmann et al., 1993; Gui et al., 1994). One activity is associated with snRNP particles and phosphorylates serine residues in the serine-arginine repeat motif (Woppmann *et al.*, 1993). This activity phosphorylates the U1 snRNP 70 kDa protein at ^a subset of the sites of phosphorylation detected in vivo. Gui et al. (1994) identified a cell cycle-regulated serine kinase that can phosphorylate SR splicing factors, SRPKI (SR protein specific kinase 1). Purified SRPK1 can induce disassembly of speckled intranuclear snRNP structures in interphase nuclei. There is also evidence that the phosphorylation state of SR104 proteins is dynamic in vivo, since they were reported to be hyperphosphorylated in metaphase cells (Roth et al., 1990; Gui et al., 1994).

Apart from SR104 proteins, the other major candidates for regulated target proteins are U2AF and the U1 70 kDa protein, which both have an SR domain and are required at an early stage of spliceosome assembly (Ruby and Abelson, 1988; Séraphin and Rosbash, 1989; Barabino et al., 1990; Zamore et al., 1992; Zhang et al., 1992). Although we have no evidence at present whether U2AF or Ul 70 kDa is affected, either directly or indirectly, by treatment of HeLa nuclear extracts with PP1, it is possible that their activity could be regulated by phosphorylation mechanisms. In the case of U1 snRNP 70 kDa protein, Tazi et al. (1993) have shown that its thiophosphorylation inhibits splicing *in vitro*, but not spliceosome assembly. This is consistent with the dephosphorylation of Ul 70 kDa protein being required for catalysis after spliceosome formation has occurred (see Figure 9). It is not yet known whether the phosphorylation state of U1 70 kDa might also influence spliceosome assembly.

The E complex, i.e. the earliest functional mammalian pre-splicing complex detected, contains Ul snRNP, both subunits of U2AF and at least five additional spliceosomeassociated proteins. In E complex the ⁵' and ³' splice sites are functionally associated (Michaud and Reed, 1991, 1993). That no E complex forms in PP1-treated nuclear extracts suggests that either stable binding of factors to the respective ⁵' and ³' splice sites, or stable interactions between these components, are prevented by dephosphorylation. Recently, SC35 and ASF/SF2 were found to interact specifically with both the U1 70 kDa protein and with the small subunit of U2AF (Wu and Maniatis, 1993).

Therefore it was proposed that SR proteins play ^a role in promoting or stabilizing interactions between the separate ends of the intron, e.g. the Ul and U2 snRNP, at an early stage of spliceosome assembly. These 'bridging' functions may be affected by changes in phosphorylation mediated by PPI. The complex we detect accumulating after PPI treatment may correspond to an otherwise transient intermediate between the H and E complexes which has not been previously reported. Therefore, an important task will be to characterize in detail this arrested complex, and determine its composition and its relationship to previously described complexes that form at early stages of spliceosome assembly (Legrain et al., 1988; Séraphin and Rosbash, 1989; Michaud and Reed, 1991; Jamison et al., 1992; Fu, 1993). It will also be important to relate the present in vitro findings with possible in vivo roles of reversible phosphorylation in regulating pre-mRNA splicing. In this regard it is interesting that one essential pre-mRNA splicing factor in Schizosaccharomyces pombe, prp4+, codes for a predicted Ser/Thr kinase (Alahari et al., 1993). Based on the network of phosphorylation mechanisms involved in regulating a wide range of biological processes, we may anticipate that complex effects of phosphorylation on pre-mRNA splicing remain to be uncovered.

Materials and methods

Protein phosphatases and phosphatase inhibitors

Human PP1 γ was expressed in *E.coli* and purified as described by Alessi et al. (1993). The isolation and sequence of the PP1 γ clone was described by Barker et al. (1993). The catalytic subunits of PPI and PP2A (Cohen et al., 1988) were purified to homogeneity from rabbit skeletal muscle and kindly provided by Prof. Philip Cohen (University of Dundee). One unit of activity catalyses the dephosphorylation of ^I nmol of glycogen phosphorylase in ¹ min. Microcystin-LR was purchased from Sigma.

In vitro splicing assays

HeLa cell nuclear extracts were prepared as described by Mermoud et al. (1992). Splicing assays were performed using uniformly labelled, capped pre-mRNAs incubated with nuclear extracts using the in vitro splicing conditions described by Lamond et al. (1987). Adenovirus major late precursor (adeno pre-mRNA) was transcribed from Sau3A-digested plasmid pBSAdl (Konarska and Sharp, 1987). Splicing products were separated on 10% polyacrylamide/8 M urea denaturing gels, run in $1 \times$ TBE. Splicing complexes were separated on non-denaturing agarose-polyacrylamide composite gels, as described by Lamond et al. (1989).

Micrococcal nuclease treatment of nuclear extracts

MCN was purchased from Sigma and further purified by FPLC chromatography on MonoS, following the protocol of Young et al. (1991). MCN digestion was performed essentially as described by Krainer and Maniatis (1985). Sixty microlitres of HeLa nuclear extract were mixed with $CaCl₂$ to give ^a final concentration of ¹ mM and then MCN added. After incubation for 25 min at 30°C, the nuclease was inactivated by adding EGTA to ^a final concentration of ¹ mM. The MCN control extract (MCN Ctrl) corresponds to an otherwise identical sample where EGTA was present from the beginning of the incubation at 30°C. Aliquots of the MCN and MCN Ctrl extracts were deproteinized and analysed by Northern blotting to ensure that snRNAs were degraded in the MCN extract, but not in the MCN control.

Streptavidin agarose affinity selection assay and Northern hybridization analyses

Antisense affinity selection assays and Northern hybridization analyses were performed as described by Ryder et al. (1990). The sequence of the biotinylated, anti-adeno intron 2'-0-Me RNA oligo was as follows: $5'$ -IACCAIAUIIACICIICC-3' (I = inosine) with four additional biotinylated deoxycytidines at the ⁵' terminus (Sproat et al., 1989).

Gel filtration of pre-splicing E complexes

Splicing reactions (300 μ l) were set up with RNasin and dithiothreitol (DTT), but without the addition of exogenous ATP, magnesium and CP. After pre-incubation in the absence or presence of protein phosphatase for 5 min at room temperature, pre-mRNA was added and reactions were incubated for 14 min at 30^oC, and subsequently loaded directly onto ^a 1.5X50 cm Sephacryl S-500 column equilibrated in FPS buffer [20 mM Tris (pH 7.8), 0.1% Triton X-100, 60 mM KCl, 2.5 mM EDTA]. Fractionation by gel filtration was performed as described by Reed (1990); the column flow rate was increased to 0.15 ml/min..

TLC

للمرابط المعالمي

Aliquots (1/20th of each sample) of standard splicing reactions containing α -³²P-labelled ATP were separated by TLC on Polygram PEI cellulose MN ³⁰⁰ paper, developed using ³⁰⁰ mM aqueous trisodium citrate at pH 7.0. Standards of unlabelled ATP and ADP were located, after drying, under ^a 254 nm UV lamp.

SDS-PAGE and immunoblotting

For Western blots, 40 μ I splicing reactions containing 14 μ I of nuclear extract (9 mg/ml) were used. Reactions were stopped by the addition of Laemmli sample buffer and proteins were separated on 12.5% SDS-polyacrylamide gels using the method of Lehmeier et al. (1990). After transfer of proteins on nitrocellulose membranes using semi-dry electrotransfer, Western blot analysis was performed as described by Blencowe et al. (1993). The immunoblots shown were developed using ECL chemiluminescence (Amersham, UK), in combination with horseradish peroxidase secondary antibodies, as per the manufacturer's instructions.

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