Regulation of CDK7 substrate specificity by MAT1 and TFIIH

Amgen Institute and Department of Medical Biophysics, University of CAK has been isolated from *Xenopus* and mammalian Toronto, 620 University Avenue, Suite 706, Toronto,

The cyclin-dependent kinase (CDK)-activating kinase

is a degradation product, although it can be reconsitted

CAK has been proposed to function in the constituted cycle progression, DNA repair and RNA polymerase II

cXK

the ring finger protein MAT1 (Nigg, 1996). It activates 1995). TFIIH phosphorylates CDC2, CDK2, TFIIF, TFIIE, other CDKs *in vitro* by phosphorylating a conserved TATA-binding protein (TBP) and the CTD (Lu *et al.*, threonine residue in their T-loop domain (Morgan, 1995). 1992; Serizawa *et al.*, 1992; Ohkuma and Roeder, 1994; threonine residue in their T-loop domain (Morgan, 1995). Recently, the *in vivo* significance of CDK phosphorylation Serizawa *et al.*, 1995; Shiekhattar *et al.*, 1995). It is not by CAK has been questioned since the enzyme with this known how efficiently the GTFs TFIIE, TFIIF and TBP function in budding yeast is not homologous to CAK are phosphorylated by free CAK. In one report, phos-(Kaldis *et al.*, 1996; Thuret *et al.*, 1996). CAK plays an phorylation of a CTD peptide by reconstituted CAK was important role in controlling transcription by phosphorylat- enhanced by MAT1 (Adamczewski *et al.*, 1996); however, ing the C-terminal domain (CTD) of the RNA polymerase it was not clear whether this effect was due to altered II (pol II) large subunit and has been implicated in substrate specificity or simply to stabilization of the repair of UV-damaged DNA (Adamczewski *et al.*, 1996; CDK7–CycH complex. Hoeijmakers *et al.*, 1996). The yeast CDK7 homologue, Here we have studied different forms of natural CAK Kin28, is essential, and its conditional inactivation blocks immunoprecipitated from cell extracts and recombinant pol II transcription and inhibits CTD phosphorylation CAK prepared from baculovirus-infected insect cells. Our

Krassimir Y.Yankulov and David L.Bentley¹ *in vivo* (Cismowski *et al.*, 1995; Valay *et al.*, 1995; Akhtar *et al.*, 1996).

Form of the Supervisor of the Supervisor CDK7-CycH-MAT1,
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et al., 1993; Devault *et al.*, 1995; Fisher *et al.*, 1995). It

cyclins A and B enhanced the phosphorylation of CDC2 by CAK (Fisher and Morgan, 1994), whereas CycD **Introduction Inhibited phosphorylation of CDK2** (Higashi *et al.*, 1996).
In addition, the association of CDK inhibitors with CDK– CAK is a cyclin-dependent kinase (CDK) which contains cyclin complexes inhibited their phosphorylation by CAK a catalytic CDK7 (p40^{MO15}) subunit, cyclin H (CycH) and *in vivo* (Kato *et al.*, 1994) and *in vitro* (Apreli in vivo (Kato et al., 1994) and in vitro (Aprelikova et al.,

results show that association with MAT1 and 'core' TFIIH modulates the activity of CDK7 at the level of substrate specificity.

Results

MAT1 alters the substrate specificity of recombinant CAK

The MAT1 subunit of CAK stabilizes the association of CDK7 with CycH and was proposed to serve as an assembly factor (Devault *et al.*, 1995; Fisher *et al.*, 1995; Tassan *et al.*, 1995). We investigated whether MAT1 might also affect the catalytic properties of CDK7. Human CAK subunits were co-expressed in Sf9 cells and immunoprecipitated by anti-CDK7 antibodies. Extensive precautions were taken to inhibit phosphatases during preparation
of the recombinant proteins (see Materials and methods)
as phosphorylation is important for stability of the bipartite
as phosphorylation is important for stabili CDK7–CycH complex (Fisher *et al.*, 1995). We found that precipitation. (**B**) Autoradiograph of kinase reactions with bipartite and immunoprecipitation was more effective than conventional tripartite recombinant CAK. + ind immunoprecipitation was more effective than conventional tripartite recombinant CAK. + indicates blocking of anti-

chromatography (Ni²⁺ NTA and S Sepharose) in separat antibody with excess antigenic peptide. The positi chromatography (Ni^{2+} –NTA and S-Sepharose) in separat-
ing recombinant CAK from the endogenous CTD kinase
are indicated. (C) MAT1 + CycH, CycH + CDK7 and MAT1 + activity in Sf9 lysates. Co-immunoprecipitation of CycH CycH + CDK7 samples were prepared as in (A). Parallel kinase
and MAT1 with CDK7 was confirmed by Western blot assays were performed with pol II and GST-CDK2(K33R). and MAT1 with CDK7 was confirmed by Western blot analysis (data not shown). We did not succeed in reconstituting an immunoprecipitable ERCC2–CAK complex by mixing extract from Sf9 cells expressing ERCC2 with extract from cells co-expressing $CDK7 + CycH +$ ratios for the CDK7–CycH complex varied between 0.23 MAT1 or by co-expressing all four subunits in Sf9 cells and 2.4 in different co-infections, whereas the ratio for (data not shown). MAT1–CDK7–CycH complexes varied between 2.9 and

reactions were performed with a mixture of two substrates: however, we always found that in parallel infections the GST–CTD and GST–CDK2(K33R). The GST–CDK2- tripartite complex had a higher CTD/CDK2 phosphoryl- (K33R) substrate is inactive as a kinase due to a mutation ation ratio by a factor of 4.5 on average. Side-by-side in its catalytic centre. It is phosphorylated on residue kinase reactions with immunopurified calf thymus RNA T160 of CDK2 to equal extents in the presence or absence pol II (Thompson *et al.*, 1990) as substrate showed of CycA (Poon *et al.*, 1993). The GST–CTD substrate that phosphorylation by CDK7–CycH–MAT1 was also contains full-length mouse CTD with 52 heptad repeats enhanced relative to CDK7–CycH by 2.2-fold (Figure 1C, (consensus YSPTSPS). In its hyperphosphorylated state lanes 2 and 3). The smaller effect observed with this (CTD-0), it migrates more slowly than in its hypophos- substrate may result from the fact that it is purified as a phorylated CTD-A state. Kinase reactions were linear for mixture of hypo- and hyperphosphorylated forms. at least 2 h, and phosphorylation of the two substrates We considered the possibility that the higher CTD/ was measured after 30 min. Under these conditions, the CDK2 phosphorylation ratio in the presence of MAT1 relative phosphorylation rates for the two substrates are a could be a non-specific effect of stabilizing CDK7–CycH measure of substrate specificity, i.e. the ratio of k_{car}/K_M which results in greater kinase activity per reaction. To for the two substrates. Note that this ratio is independent test this possibility, we titrated tri for the two substrates. Note that this ratio is independent of substrate concentration (Fersht, 1985). 1–3) and bipartite (Figure 2A, lanes 4–6) CAK while

expressed in Sf9 cells. Western blotting showed equal strated that the enhancement of CTD versus CDK2 levels of CDK7 and CycH respectively in the two lysates phosphorylation by MAT1 was independent of the total (data not shown). Immunoprecipitates of both lysates with amount of kinase activity in the reaction. anti-CDK7 antibody had significant kinase activity towards To confirm the effect of MAT1 on substrate preference, GST–CTD and GST–CDK2(K33R). This activity was we prepared CAK complexes by a different method. dependent on infection with both CDK7 and CycH viruses CDK7, CycH and MAT1 were expressed individually in (Figure 1B, lanes 1 and 2 and data not shown) and Sf9 cells. Equal amounts of CDK7 and CycH extracts was abolished by blocking the anti-CDK7 antibody with plus either mock-infected or MAT1-infected extract were antigenic peptide (Figure 1B, lanes 4 and 6). Tripartite mixed for 1 h and then immunoprecipitated with anti-CAK consistently phosphorylated GST–CTD more extens- CDK7 antibodies. The CAK complexes prepared in this ively than the bipartite complex relative to GST–CDK2 way had approximately equal activity toward GST– (Figure 1B, lanes 3 and 5). The average ratio of CTD to CDK2(K33R) but the MAT1-containing complex had CDK2 signals was 4.06 ($n = 11$) for CDK7–CycH–MAT1 much higher activity toward GST–CTD (Figure 2B, lanes and 0.88 $(n = 11)$ for CDK7–CycH co-expressed in Sf9 2 and 3). We consistently observed that the bipartite cells (see Figure 6). The CTD/CDK2 phosphorylation complex assembled from individually expressed subunits

In order to measure substrate preference directly, kinase 5.4. We do not understand the basis for this variation;

 $CDK7 + CycH$ or $CDK7 + CycH + MAT1$ were co-
keeping the substrates constant. This experiment demon-

Fig. 2. MAT1 shifts CAK substrate preference. (**A**) CAK substrate preference is independent of total kinase activity. Kinase assays with baculoviral CAK complexes were as in Figure 1B. Serial 1:3 dilutions were used in lanes 1–3 and 4–6. The gel was quantified by Phosphorimager (PI) (Molecular Dynamics) and arbitrary units were plotted as well as CTD/CDK2 ratios. (**B**) Substrate preferences of bipartite and tripartite CAK reconstituted from individually expressed subunits. Extracts from 10⁵ infected Sf9 cells were mixed for 1 h on ice, immunoprecipitated and assayed for kinase activity. Extract from 10⁵ mock-infected Sf9 cells was added to samples 1 and 2. (**C**) Dose-dependent shift in CAK substrate specificity by MAT1. Sf9 cells were infected with CycH and CDK7 viruses at an m.o.i. of 3–5 (lanes 1–9) and MAT1 virus at an m.o.i. from 0.5 (lane 2) to 18 (lane 9). Kinase assays were performed as in Figure 1B.

assembled by mixing of individual components behaved

had especially low reactivity toward the CTD. A control We also tested whether the effect of MAT1 on the mix of MAT1- and CycH-infected extracts had no immuno- substrate specificity of CDK7–CycH was dose dependent. precipitable kinase activity (Figure 2B, lane 1). In sum-
Sf9 cells were infected with the CDK7 and CycH viruses mary, the substrate preferences of CAK complexes at a constant m.o.i. of 3–5 and the MAT1 virus was titrated assembled by mixing of individual components behaved from m.o.i. 0 to 18. CAK was precipitated from each in the same way as those made by co-expression. infected cell extract with anti-CDK7 antibody and assayed

Fig. 3. Natural free CAK and TFIIH differ in their substrate specificities. (**A**) Scheme for preparation of TFIIH and free CAK from HeLa cell extract by sequential immunoprecipitation (IP) with rabbit anti-CDK7 and anti-p62 peptide antibodies. The final TFIIH and free CAK preparations were analysed by Western blotting with monoclonal anti-p62 (3C9), anti-CDK7(2F8) and anti-ERCC2 (2F6). A 2-fold greater fraction of the TFIIH was loaded relative to free CAK. (B) Kinase and Western blot analysis of HeLa TFIIH (lane 4) and free CAK (lane 5) prepared as in (A). Lanes 1-3: controls immunoprecipitated with anti-GST, anti-p62 or anti-CDK7 with 2-fold greater loading of lane 2. (**C**) Comparison of the kinase activities of TFIIH with total CAK from HeLa cells and *Xenopus* oocytes. TFIIH (lanes 3 and 7) and total CAK (lanes 2, 4 and 6) were prepared by single or double IP as indicated; 2.5-fold larger fractions of TFIIH (lanes 3 and 7) were analysed relative to total CAK (lanes 2 and 6). (**D**) Immunopurified HeLa TFIIH is activated by TFIIE. TFIIH was purified as in (A) and kinase activity toward RNA pol II was assayed as in Figure 1C. Recombinant TFIIE was added at 5 µg /ml.

(Figure 2C). This experiment showed that enhancement assay. Western blotting showed that the anti-p62 precipitate of the CTD phosphorylation relative to CDK2 phosphoryl- contained both p62 and CDK7, whereas the supernatant ation was approximately linearly related to the amount of was effectively cleared of 'core' TFIIH as demonstrated MAT1 virus used to infect the culture. We conclude that by the absence of p62 (Figures 3A and B and 5). The the MAT1 subunit confers on CDK7–CycH a preference presence of ERCC2 in the supernatant from the antifor RNA pol II CTD as a substrate over CDK2. p62 immunoprecipitation indicates that our 'free' CAK

ing isoforms of CAK, we purified them by immuno- of this complex in HeLa cells (Fisher *et al.*, 1995). precipitation from HeLa whole cell extract according to Free CAK and TFIIH immunopurified from HeLa cells

with GST–CTD and GST–CDK2(K33R) substrates form of CAK were treated identically before the kinase preparation also contains ERCC2–CAK (Figure 3A). **TFIIH and free CAK differ in their substrate** Attempts to purify bipartite CAK from anti-CDK7 eluates **preference** by depletion with anti-MAT1 antibodies were unsuccess-To compare the substrate specificities of naturally occurr- ful, probably because of the low abundance or instability

the scheme in Figure 3A. Free CAK and TFIIH-associated both had abundant kinase activity whereas anti-GST CAK were immunoprecipitated with anti-CDK7 peptide control immunoprecipitates had none (Figure 3B, lane 1; antibody, and then eluted from the precipitate with excess C, lanes 1 and 5). There were substantial differences peptide. The eluted material was re-immunoprecipitated between the kinase activities of the CAK and TFIIH with antibodies against either MAT1 or the p62 subunit samples. First, TFIIH had a stronger preference for the of TFIIH. The anti-p62 precipitate contains TFIIH while GST–CTD substrate over GST–CDK2(K33R) than did the supernatant contains free CAK. The anti-MAT1 pre- free CAK (Figure 3B, lanes 4 and 5). The average ratios cipitate is a mixture of TFIIH and free CAK and serves of total CTD/CDK2 phosphorylation signals were 14.4 as a control for total CAK activity after the second round and 4.7 for TFIIH $(n = 7)$ and CAK $(n = 5)$, respectively of immunoprecipitation. This purification strategy has the (see also Figure 6). Secondly, the TFIIH-associated kinase advantage that both the TFIIH-associated and the free caused a pronounced mobility shift of a fraction of the

5) although there was substantially less CDK7 in the TFIIH sample as determined by Western blotting. The **TFIIE and TFIIF are phosphorylated by TFIIH but** average ratios of CTD-0/CTD-A signals were 1.12 and **not by free CAK** 0.04 for TFIIH ($n = 7$) and free CAK ($n = 5$), respectively **In** addition to the 0.04 for TFIIH ($n = 7$) and free CAK ($n = 5$), respectively

(see also Figure 6). The accumulation of GST-CTD

molecules in the CTD-0 position when phosphorylated by

TFIIH (Figure 3B and C) was not an effect of a limitin

free CAK plus TFIIH, phosphorylated GST–CTD and
GST–CDK2(K33R) similarly to the original anti-CDK7 TFIIH phosphorylated the TFIIE p56 subunit and the eluate. As expected, the anti-p62 immunoprecipitate con-
taining only TFIIH preferentially phosphorylated the CTD
over CDK2 and hyperphosphorylated a significant fraction
ated either of these substrates to a detectable lev over CDK2, and hyperphosphorylated a significant fraction ated either of these substrates to a detectable level (Figure
of the CTD. This result therefore confirmed the enhanced 4, lanes 11, 12, 15 and 16). We consistently of the CTD. This result therefore confirmed the enhanced 4, lanes 11, 12, 15 and 16). We consistently observed that CTD/CDK2 phosphorylation ratios for TFIIH relative to phosphorylation of TFIIF by TFIIH was more efficient CTD/CDK2 phosphorylation ratios for TFIIH relative to free CAK. The contract of the than by TFIIE. In an experiment not shown, CAK also

with anti-*Xenopus* CDK7 peptide antibody and TFIIH was association with TFIIH.

prepared by CDK7 peptide elution and re-precipitation with cross-reactive anti-p62 antibody (Figure 3C, lanes 5–7). As we observed for the human enzymes, the ability to hyperphosphorylate the CTD co-purified with TFIIH (Figure 3C, compare lanes 6 and 7). In conclusion, the experiments in Figure 3 show that the ability to hyperphosphorylate the CTD is greatly enhanced in TFIIH relative to free CAK.

The increased phosphorylation of CTD by TFIIH relative to free CAK could result from contamination of immunoprecipitated TFIIH with TFIIE which stimulates the TFIIH kinase (Lu *et al.*, 1992). Two lines of evidence Fig. 4. TFIIH but not CAK can phosphorylate TFIIE and TFIIF. **Fig. 4.** TFIIH but not CAK can phosphorylate TFIIE and TFIIF. Kinase reactions were performed with: lanes 1–4, conventionally did not detect either the p34 or the p56 subunit of TFIIE purified TFIIH (HAP fraction) (Gérard et al., 1991); lanes 5–8, in immunoprecipitates with anti-CDK purified TFIIH (HAP fraction) (Gérard *et al.*, 1991); lanes 5–8,
immunoprecipitates with anti-CDK7, anti-p62 or anti-
immunoprecipitated TFIIH (as in Figure 3A); lanes 9–12, soluble
recombinant baculoviral tripartite CAK, prepared as in Figure 3A. Controls, C, contained no substrate. The still responsive to addition of TFIIE (Figure 3D). In this positions of phosphorylated GST-CDK2(K33R), GST-CTD-A, experiment. TFIIH purified by sequential positions of phosphorylated GST-CDK2(K33R), GST-CTD-A,

GST-CTD-0, the rap74 subunit of TFIIF and the p56 subunit of TFIIE

are indicated. The TFIIF preparation contains an ~50 kDa degradation

product of rap74 which is al recombinant TFIIE. The kinase activity of immunopurified GST–CTD substrate to the slowly migrating hyperphos-
phorylated CTD-0 form (Figure 3A, lane 4). In contrast,
free CAK did not convert GST–CTD to the CTD-0 form
(Figure 3B, lane 5). Note that total CTD phosphorylation
(CTD

staining (data not shown). TFIIH prepared by an alternative
protocol (anti-p62 immunoprecipitation followed by pep-
immobilization of our immunoprecipitated TFIIH, we also
tide elution and re-precipitation with anti-CDK7 a We also compared the abilities of total CAK and TFIIH failed to phosphorylate TBP. These results indicate that free purified from *Xenopus* oocytes to hyperphosphorylate the CAK does not recognize TFIIE and TFIIF as substrates; CTD. Total CAK was prepared by immunoprecipitation however, it acquires specificity for these substrates upon

Fig. 5. CDK7 is phosphorylated in vivo equally in free CAK and
TFIIH. HeLa cells were metabolically labelled with ³²PO₄, and TFIIH ratios of Phosphorimager values. (lane 4) and free CAK (lane 5) were prepared by double IP as in Figure 3A. As controls, single IPs with anti-GST, anti-p62 and anti-CDK7 are shown in lanes 1–3. Samples were transferred to PVDF TFIIE, TFIIF and TBP, and the CTD of the largest subunit membrane and $[^{32}P]CDK7$ was detected by Phosphorimager. of RNA pol II (Morgan, 1995; Hoeijmakers *et*

formation of a stable bipartite complex with CycH but is recombinant and natural CAK were prepared and assayed not essential in the presence of MAT1 (Devault *et al.*, with mixtures of substrates to determine their substrate 1995; Fisher *et al.*, 1995). We tested the hypothesis preferences. that a difference in phosphorylation of CDK7 in TFIIH We observed that phosphorylation of RNA pol II and compared with free CAK contributes to the difference in a full-length CTD fusion protein by recombinant CAK substrate specificity between these two forms of the reconstituted in insect cells is significantly enhanced by enzyme. HeLa cells were 3^2P -labelled and TFIIH and free MAT1 (Figures 1 and 2). The CTD/CDK2 phosphorylation CAK were purified by sequential immunoprecipitation ratio increased from 0.88 $(n = 11)$ for recombinant CDK7– with anti-CDK7 and anti-p62 antibodies (see Figure 3A). CycH to 4.06 $(n = 11)$ for CDK7–CycH–MAT1. Our Samples were resolved on an SDS-gel, transferred to data did not reveal a significant effect of MAT1 on PVDF membrane and ${}^{32}PO_4$ incorporation into CDK7 was phosphorylation of GST–CDK2, but we cannot exclude quantified by Phosphorimager. The membrane was then the possibility of an effect in the presence of partner quantified by Phosphorimager. The membrane was then reacted with anti-CDK7, anti-p62 and anti-MAT1 anti- cyclins. The catalytic activity of recombinant tripartite bodies. The anti-p62 Western blot showed that the free CAK closely resembled that of free cellular CAK in the CAK (Figure 5, lane 5) was not substantially contaminated ratio of both CTD/CDK2 and CTD-0/CTD-A phosphorylwith TFIIH (Figure 5, lane 4). Comparison of the ^{32}P and ation (see Figure 6). We think it unlikely, therefore, that Western blot signals did not reveal a significant difference MAT1 enhanced CTD phosphorylation by recruiting insect in relative phosphate incorporation into the CDK7 subunits 'core' TFIIH to the human CDK7–CycH complex. This of TFIIH versus free CAK. These results, therefore, do possibility is also unlikely because TFIIH is not an not indicate that a change in the overall level of CDK7 abundant factor and because MAT1 functions in a speciesphosphorylation causes the difference in substrate speci- specific way (Devault *et al.*, 1995).

CAK exists in at least two distinct states in the cell; as a far higher proportion of the CTD substrate was hyperphosfree complex and as a component of the pol II general phorylated by TFIIH than by free CAK (Figures 3 and transcription factor TFIIH. CAK phosphorylates many 4). The average ratio of CTD-0/CTD-A signals was 15 substrates *in vitro*, including several CDKs, the GTFs fold higher for TFIIH than for free CAK (Figure 6).

Fig. 6. Substrate specificities of different CAK complexes. CycH–CDK7 and MAT1–CycH–CDK7 values are averages from 11 experiments with baculoviral complexes reconstituted by co-infection of Sf9 cells. CAK and TFIIH indicate average ratios from five and seven experiments, respectively, with free CAK and TFIIH prepared from HeLa cells as described in Figure 3A. The CTD value is the sum of CTD-0 + CTD-A. CTD/CDK2 and CTD-0/CTD-A values represent

membrane and ^{[32}P]CDK7 was detected by Phosphorimager. of RNA pol II (Morgan, 1995; Hoeijmakers *et al.*, 1996; Subsequently. Western blot analysis was performed with monoclonal Nigg 1006). This broad range of terrorts c Subsequently, Western blot analysis was performed with monoclonal
anti-p62 (3C9), anti-CDK7 (2F8) and rabbit anti-MAT1 antibodies.
that the enzyme has a relaxed substrate specificity, at least *in vitro*. Alternatively, the different isoforms of CAK could **Phosphorylation of CDK7 in TFIIH and free CAK be restricted in their specificity to different substrates.** Phosphorylation of CDK7 on residue T170 is required for In order to investigate this question, various forms of

ficity between TFIIH and free CAK. CAK phosphorylation of the CTD was stimulated significantly by association with TFIIH. The average ratios **Discussion** of CTD/CDK2 phosphorylation signals were 4.06 (*n* = 11) and 4.67 (*n* = 5) for recombinant tripartite CAK and **CAK** is regulated at the level of substrate free cellular CAK, respectively, whereas the ratio for **specificity** TFIIH was 14.4 (*n* = 7) (see Figure 6). Furthermore, a Hence, our data imply that TFIIH phosphorylates many CTD hyperphosphorylation is greatly enhanced in serine residues each time it binds a substrate molecule, TFIIH-associated CAK relative to free CAK (see Figure whereas free CAK phosphorylates only one or a few 6). GST–CTD hyperphosphorylation by free CAK was residues. observed previously when the enzyme was immuno-

CAK phosphorylated TFIIE (p56) or TFIIF (rap74) (Figure (Tassan *et al.*, 1995). In this case, cross-reaction of 4). In contrast, conventionally purified and immuno- the antibody with the GST–CTD substrate presumably precipitated TFIIH phosphorylated both these substrates stabilized the kinase–substrate interaction and promoted (Figure 4) (Ohkuma and Roeder, 1994; Yankulov *et al.*, processive phosphorylation (Tassan *et al.*, 1995). An 1995). This observation suggests that the ability to phos-

in a similar manner to enhance hyperphosphorylation.

In a similar manner to enhance hyperphosphorylation. phorylate TFIIE and TFIIF is acquired by CAK upon

to immunoprecipitate RNA pol II holoenzyme containing phosphorylations on a single GST–CTD molecule. Interall the GTFs (Ossipow et al., 1995). Under our immuno-
actions between TFIIE and TFIIF with 'core' TFIIH precipitation conditions, we did not detect RNA pol II or (Maxon *et al.*, 1994) may be responsible for the specificity TFIIE in the anti-CDK7 or anti-p62 precipitates by Western of TFIIH-associated CAK for these substrate TFIIE in the anti-CDK7 or anti-p62 precipitates by Western blotting (data not shown). In addition, TFIIH which had been purified extensively over seven columns (Ge´rard **Implications for the regulation of the activity of** *et al.*, 1991) phosphorylated TFIIF and TFIIE and hyper- **other CDKs** phosphorylated CTD similarly to the immunoprecipitated Here we demonstrate that the substrate preference of CAK TFIIH (Figure 4). Hence, we believe that these are is modulated by its association with MAT1 and TFIIH. properties of TFIIH *per se* rather than of a higher order The control of other CDKs at the level of substrate

Recently, a complex containing CAK (CDK7–CycH– observed, however, that CycA–CDC2 and CycA–CDK2 MAT1) and ERCC2 was separated from TFIIH (Drapkin kinases phosphorylate the Rb-related protein p107. *et al.*, 1996; Reardon *et al.*, 1996). Consistent with these whereas the corresponding CycB complexes do not (Peeper observations, we found appreciable amounts of ERCC2 *et al.*, 1993). In addition, interaction of p107 w observations, we found appreciable amounts of ERCC2 *et al.*, 1993). In addition, interaction of p107 with CycA–
but no p62 in our free CAK preparations (Figure 3A and CDK2 or CycE–CDK2 apparently restricted the substrate but no p62 in our free CAK preparations (Figure 3A and CDK2 or CycE–CDK2 apparently restricted the substrate B). The kinase activity of free CAK containing ERCC2 specificity of these kinases to p107 and not pRB (Zhu B). The kinase activity of free CAK containing ERCC2 specificity of these kinases to p107 and not pRB (Zhu toward CDK2 and CTD substrates closely resembled that $et al.$ 1995). The mechanism of regulating kinase activity toward CDK2 and CTD substrates closely resembled that *et al.*, 1995). The mechanism of regulating kinase activity of recombinant tripartite CAK rather than TFIIH (see by modulation of substrate specificity may therefore, of recombinant tripartite CAK rather than TFIIH (see by modulation of substrate specificity may, therefore, be Figure 6). In addition, free CAK did not phosphorylate of significance to other members of the CDK family in TFIIE or TFIIF (Figure 4). These data argue that association of CAK with ERCC2 is not sufficient to confer the characteristic substrate specificity of the TFIIH kinase.

In summary, we suggest that the MAT1 subunit of **Materials and methods** CAK, previously shown to facilitate assembly of CDK7 **Antibodies** in favour of the CTD; and that a high stoichiometry of CTD phosphorylation requires 'core' TFIIH. residues 131-148 (DVIQKNKLKLTREQEELE), human CycH

Modulating the conformation of the catalytic site may be antibody (CDQKDIAKKLSF) was purchased from Upstate Biotechno-
important in control of protein kinase activity (Morgan logy Inc. Rabbit anti-GST antibodies were affin important in control of protein kinase activity (Morgan
and De Bondt, 1994). For example, the extensive con-
formational changes which occur when the conserved
The Decomes phosphorylated (Russo *et al.*, 1996) might
and ER Thr160 becomes phosphorylated (Russo *et al.*, 1996) might raised against a comp
affect substrate recognition in some CDKs. However, our provided by M.Pandes. affect substrate recognition in some CDKs. However, our experiments revealed no evidence for involvement of
phosphorylation in modulating CDK7 substrate specificity.
There was no correlation between altered substrate recog-
nition by CDK7 in free CAK and TFIIH and altered
 10 nition by CDK7 in free CAK and TFIIH and altered overall phosphorylation of the enzyme (Figure 5). This $[\gamma^{32}P]$ ATP per reaction; 1 μg/ml aprotonin, 1 μg/ml leupeptin, 1 μg/ml overall phosphorylation of the enzyme (Figure 5). This $[\gamma^{32}P]ATP$ per reaction; 1 $\mu g/m$ aprotonin, 1 $\mu g/m$ leupeptin, 1 $\mu g/m$ experiment does not exclude the possibility that CDK7 is phosphorylated to equal extents o different complexes. Addition of MAT1 to CycH–CDK7 for 30 min at 30°C and terminated by addition of 5 μ l of 5× SDS dramatically alters its behaviour in gel filtration experi-
ments suggesting a significant conformational change SDS gels which were fixed and stained with Coomassie Brilliant Blue ments, suggesting a significant conformational change SDS gels which were fixed and stained with Coomassie Brilliant Blue
(Gighen at al. 1005). Such an effect of MAT1 gould be to verify equal loading. Dried gels were quant (Fisher *et al.*, 1995). Such an effect of MAT1 could be
involved both in promoting binding to CycH and in
phosphorylation of the CTD.
 $\frac{1}{2}$ (Melecular Dynamics).
The following protein substrates were used: a mix of 5

Neither recombinant tripartite CAK nor free cellular precipitated with antibodies raised against GST–MAT1 association with 'core' TFIIH. Such an interaction could, for example, decrease the Previously, a monoclonal anti-CDK7 antibody was used dissociation rate of the substrate thereby allowing multiple

complex of transcription factors or a contaminating kinase. Specificity has not been explored extensively. It has been
Recently, a complex containing CAK (CDK7–CycH–
observed, however, that CycA–CDC2 and CycA–CDK2 kinases phosphorylate the Rb-related protein p107, of significance to other members of the CDK family in addition to CAK.

Affinity-purified rabbit anti-peptide antibodies were prepared against
human CDK7 residues 329-343 (KRTEALEQGGLPKK), human MAT1 C-terminus (SKKSKHEEEEWTDDDLVESL) and human TFIIH p62 **Mechanisms of CDK7 substrate specificity**
 C-terminus (YNKLHTWQSRRLMKKT). The anti-p62 antibody cross-

regulation

not shown). Affinity-purified anti-X.laevis CDK7 C-terminal peptide **regulation** not shown). Affinity-purified anti-*X.laevis* CDK7 C-terminal peptide

of GST–CTD (Peterson *et al.*, 1992) and GST–CDK2(K33R) (Poon

RNA pol II. Expression and purification of recombinant TFIIF, TFIIE, GST-CTD and GST-CDK2(K33R) were described previously (Yankulov *et al.*, 1995). Immunoaffinity-purified calf thymus pol II (Thompson *Chem.*, **270**, 18195–18197. *et al.*, 1990) (a gift of Dr J.Greenblatt) contained both the hypophosphory-
lated (pol II A) and hyperphosphorylated (pol II 0) forms of the enzyme. a C-terminal domain kinase that controls mRNA transcription in lated (pol II A) and hyperphosphorylated (pol II 0) forms of the enzyme.

Expression and purification of CAK from insect cells kinase (CAK) activity. *Mol. Cell. Biol.*, 15, 2983–2992.
Baculoviruses, expressing human CDK7, CycH, MAT1 and ERCC2, Devault.A., Martinez.A.M., Fesquet.D., Labbe.J.C. Baculoviruses, expressing human CDK7, CycH, MAT1 and ERCC2, Devault,A., Martinez,A.M., Fesquet,D., Labbe,J.C., Morin,N., were gifts of Drs D.Morgan, R.Fisher and D.Siderovski. Sf9 cells Tassan.J.P.. Nigg.E.A., Cavadore.J.C were gifts of Drs D.Morgan, R.Fisher and D.Siderovski. Sf9 cells Tassan,J.P., Nigg,E.A., Cavadore,J.C. and Doree,M. (1995) Mat1 (7×10⁵/ml) were infected at an m.o.i. of 3–5 for each virus (except in ('ménage-à-trois') a $(7 \times 10^5/\text{ml})$ were infected at an m.o.i. of 3–5 for each virus (except in ('ménage-à-trois') a new RING finger protein subunit stabilizing Figure 2C), harvested 28–36 h post-infection, washed twice in phosphate-
cyclin buffered saline (PBS) and lysed in ice-cold IPL buffer [10 mM Na **14**, 5027–5036.
HEPES pH 7.5, 10 mM KCl, 0.4 mM DTT, 0.5 mM EDTA, 0.5 mM Drapkin.R., Leroy EGTA, 10 mM 2-glycerophosphate, 0.5 mM Na₃VO₄, 1 mM NaF, 0.2% Human cyclin-dependent kinase-activating kinase exists in 3 distinct NP-40, 1 µM microcystin, 1 µg/ml aprotonin, 1 µg/ml leupeptin, 1 µg/ complexes. *Proc.* NP-40, 1 μM microcystin, 1 μg/ml aprotonin, 1 μg/ml leupeptin, 1 μg/ml pepstatin, 50 μg/ml phenylmethylsulfonyl fluoride (PMSF)]. KCl was ml pepstatin, 50 μg/ml phenylmethylsulfonyl fluoride (PMSF)]. KCl was Fersht,A.R. (1985) *Enzyme Structure and Function*. W.H.Freeman. added to 0.4 M, the homogenate was rocked for 30 min at 4°C and Fisher.R.P. and Morgan clarified by centrifuging for 15 min at 13 000 *g*. The supernatant was frozen at -70° C after adding glycerol to 20%. CAK activity from extract frozen at –70°C after adding glycerol to 20%. CAK activity from extract Fischer,L., Gérard,M., Chalut,C., Lutz,Y., Humbert,S., Kanno,M., corresponding to 10⁵ cells was immunoprecipitated with ~1 µg of anti-
Chambon P an were washed four times for 10 min in IPL buffer containing 0.3 M KCl and 0.2% NP-40, washed once more in kinase buffer without ATP and

cells, except that KCl was added to 0.35 M. The extract was diluted

immediately with an equal volume of lysis buffer without KCl, clarified

at 45 000 r.p.m. in the SW55 Beckman rotor for 30 min and filtered

through a 0. tor 2 h with 1/10 vol of protein A-Sepharose beads. Between 5 and 10 associated with cyclin D1 is not phosphorylated by cdk7-cyclin-H.

ug of affinity-purified anti-peptide antibodies were used to immunopre-

cipitate p62peptide for 2–18 h. The eluates were filtered through Bio-Rad spin
columns to remove any beads, and immunoprecipitated with a second
antibody. The resulting supernatant and beads (after three washes of 10
min) were used fo

Metabolic labelling of HeLa cells

HeLa cells (5×10⁷) were washed in phosphate-free media and incubated

with 0.9 mCi of ³²PO₄/ml for 3-4 h. Extraction was as described above

except that clarification was for 15

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and MAT baculoviruses and to D.Siderovski for ERCC2 virus. We also *Nature*, **368**, 160–163.

thank J.-M.Egly for gifts of TFIIH and antibodies and for communication Ossipow V. Tassan J.P. thank J.-M.Egly for gifts of TFIIH and antibodies and for communication Ossipow, V., Tassan, J.P., Nigg, E.A. and Schibler, U. (1995) A mammalian of results before publication. W.Dynan, T.Hunt, M.Pandes and J.Green-
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BSA, and used for kinase assays. Under these conditions, the kinase
reaction was linear for $>$ 2 h.
activating kinase. Cell, 83, 47–58.
activating kinase.
- **TFIIH**

Gérard,M., Fischer,L., Moncollin,V., Chipoulet,J.M., Chambon,P. and

HeLa cells were washed twice in PBS and lysed as described for Sf9

cells, except that KCl was added to 0.35 M. The extract was diluted

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