Regulation of $Ca^{2+}/Calmodulin-Dependent Protein Kinase Kinase α by$ cAMP-Dependent Protein Kinase: II. Mutational Analysis¹

Takako Kitani,² Sachiko Okuno, and Hitoshi Fujisawa³

Department of Biochemistry, Asahikawa Medical College, Asahikawa 078-8510

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We previously reported that rat brain Ca²⁺/calmodulin-dependent protein kinase (CaM**kinase) IV is inactivated by cAMP-dependent protein kinase (PKA) [Kameshita, I. and Fujisawa, H. (1991)** *Biochem. Biophys. Res. Commun.* **180, 191-196]. In the preceding paper, we demonstrated that changes in the activity of CaM-kinase IV by PKA results from the phosphorylation of CaM-kinase kinase** a **by PKA and identified six phosphorylation sites, Ser²⁴ for autophosphorylation, and Ser⁵² , Ser⁷⁴ , Thr¹⁰⁸ , Ser⁴⁵⁸ , and Ser⁴⁷⁵ for phosphorylation by PKA. In the present study, a causal relationship between the phosphorylation and change in the activity toward PKlV peptide has been studied using mutant enzymes with amino acid substitutions at the six phosphorylation sites. The following conclusions can be drawn from the experimental results: (i) Phosphorylation of Ser74 and/or unidentified sites causes an increase in activity; (ii) phosphorylation of** Thr¹⁰⁸ or Ser⁴⁵⁸ causes a decrease in the activity; (iii) the inhibitory effect of the phospho**rylation of Thr¹⁰⁸ is canceled by the stimulatory effect of the phosphorylation, but that of Ser⁴⁸⁸ is not; and (iv) the inhibitory effects of Thr¹⁰⁸ and Ser⁴⁵⁸ are synergistic In contrast to the activity toward PKlV peptide, the activity toward CaM-kinase** *TV* **appears to be decreased by the phosphorylation of Thr¹⁰⁸ , but not significantly affected by the phosphorylation of Ser⁴ * 8 .**

Key words: cAMP-dependent protein kinase, Ca²⁺/calmodulin, CaM-kinase kinase, phos**phorylation site, protein phosphorylation.**

Our previous finding (1) that Ca²⁺/calmodulin-dependent protein kinase (CaM-kinase) IV is inactivated by incubation with cyclic AMP-dependent protein kinase (PKA) under the phosphorylation conditions provided a valuable example of cross-talk between $Ca²⁺$ -signaling and cyclic AMP-signaling pathways, because CaM-kinase IV is thought to be a key enzyme in the $Ca²⁺$ -signaling system in cells (2). In the preceding paper (3), the inactivation of CaM-kinase IV by PKA was demonstrated to result from the prevention of the activation of CaM-kinase IV through CaM-kinase kinase α by PKA, and several phosphorylation sites on CaM-kinase kinase α , Ser²⁴ for autophosphorylation and Ser⁵², Ser⁷⁴, Thr¹⁰⁸, Ser⁴⁵⁸, and Ser⁴⁷⁵ for PKA, were identified. However, biochemical studies did not reveal a clear relationship between phosphorylation at these phosphorylation sites and changes in enzyme activity. In the present paper, mutational analyses based on the biochemical data for the phosphorylation sites were performed in an

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attempt to reveal the molecular mechanism underlying the phosphorylation-activity relationship of CaM-kinase kinase. During the course of the studies, Wayman et al. reported mutational studies of the putative phosphorylation sites of CaM-kinase kinase α , suggesting the importance of Thr¹⁰⁸ and Ser⁴⁵⁸ in the inactivation by PKA *(4).*

EXPERIMENTAL PROCEDURES

Materials—[y-³²P]ATP (5,000 Ci/mmol) and the Thermo sequenase fluorescent-labelled primer cycle sequencing kit with 7-deaza-dGTP were from Amersham Pharmacia Biotech. Microbial protease inhibitors (pepstatin A, leupeptin, antipain, and chymostatin) were from the Peptide Institute (Osaka). DEAE-cellulose (DE52), phosphocellulose paper (P81), and 3MM paper were from Whatman. FluoroTrans, a polyvinylidene difluoride membrane, was from Pall Gelman Laboratory. NHS-LC-biotin and avidin conjugated with horseradish peroxidase were from Pierce. PKTV peptide (KKKKEHQVLMKTVCGTPGY) (5) was synthesized with a Shimadzu PSSM-8 automated peptide synthesizer. GeneEditor™ *in vitro* site-directed mutagenesis system was from Promega. pETlld was from Novagen. Restriction enzymes and other DNA modifying enzymes were purchased from Takara Shuzo, Toyobo, or New England Biolabs. All other reagents were of the highest grade commercially available.

Protein Preparations—Calmodulin was purified from *E. coli* cells transformed with expression vector pETlld carrying a cDNA encoding chicken calmodulin *(6),* essentially as described by Gopalakrishna and Anderson (7). The cDNA

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² Affiliated with the Laboratory for Radioactive Isotope Research, Asahikawa Medical College.

³ To whom correspondence should be addressed. E-mail: fujisawa@ asahikawa-med.ac.jp

Abbreviations: CaM-kinase, calmodulin-dependent protein kinase; PKA, cyclic AMP-dependent protein kinase; SDS-PAGE, SDS-polyacrylamide gel electrophoresis.

encoding chicken calmodulin was kindly donated by A.R. Means *(8).* Biotinylated calmodulin was prepared from the recombinant chicken calmodulin as described by Mangels and Gnegy (9). Recombinant rat CaM-kinase kinase *a* expressed in *E. coli (10)* was purified as described previously *(11).* PKA (catalytic subunit of cyclic AMP-dependent protein kinase) was purified from bovine heart as described previously *(12).* Recombinant rat CaM-kinase IV expressed in Sf9 cells was purified as described previously *(6).* Recombinant rat CaM-kinase $\Gamma V(K_{71}R)$, in which Lys⁷¹ (ATP-binding site) was replaced with arginine, was expressed in Sf9 cells and purified as described previously *(11).*

cDNAs for mutants of CaM-kinase kinase α in which threonine and/or serine residues identified as phosphorylation sites by autophosphorylation or PKA (3) were replaced with other amino acids, such as alanine, aspartate, or glutamate, were prepared by site-directed mutagenesis as follows. cDNAs for the CaM-kinase kinases $(S_{24}A)$, $(S_{24}D)$, $(S_{52}A)$, $(S_{52}D)$, $(S_{74}A)$, and $(S_{74}D)$ were prepared using the single-stranded DNA obtained from pUC118, into which the *Xbal-Xbal* fragment (288 bp) of pETCaMKKa *(10)* was inserted, as a template, and appropriate synthetic oligonucleotides as primers, essentially according to the method of Kunkel *et al. (13).* The mutations were confirmed by the dideoxynucleotide chain-termination method *(14)* using a DNA sequencer, LI-COR model 4000L. The pETlld carrying each cDNA containing the entire coding sequence of the mutant CaM-kinase kinase α was constructed by ligation of three fragments, the *Ncol—Xbal* fragment (250 bp) of the pUC118 containing the mutated segment, and the *Xbal-Sfil* fragment (702 bp) and *Sfil-Ncol* fragment (6.2 kbp) of $pETCaMKK\alpha$. cDNAs for the CaM-kinase kinases (S₄₅₉A), $(S₄₅₈D), (S₄₇₅A),$ and $(S₄₇₅D)$ were prepared similarly using the single-stranded DNA obtained from pUC118, into which the *Pstl-BamHl* fragment (282 bp) of pETCaMKKa was inserted, as a template, and appropriate synthetic oligonucleotides as primers. After the mutations were confirmed, the pETlld carrying the cDNA for the mutant enzyme was constructed by ligation of the *Pstl—BamHl* fragment (282 bp) of the pUC118 containing the mutated segment and the *BamHI-Sfil* (6.6 kbp) and *Sfil-Pstl* segments (318 bp) of $pETCaMKK\alpha$. cDNAs for the CaM-kinase kinases (T₁₀₈A), (T₁₀₈D), (T₁₀₈E), (S₄₅₈E), and (S₄₇₅E) were prepared using the GeneEditor™ *in vitro* site-directed mutagenesis system (Promega) with pETCaMKKa *(10)* as a template and appropriate synthetic oligonucleotides as primers. A pETlld carrying cDNA for the mutant enzyme $(S_{24}D/S_{62}D/S_{475}D)$ was constructed by ligating the *BamHI*-*Xmal* fragment (5.8 kbp) of the pETlld carrying the cDNA for the enzyme $(S_{24}D)$, the *Xmal–Sfil* fragment (814 bp) of pETlld carrying the enzyme *(SfJ)),* and the *Sfil-BamHl* fragment (600 bp) of pET11d carrying the enzyme $(S_{475}D)$. A pET11d carrying cDNA for the enzyme $(S_{52}D/S_{74}D)$ was constructed by ligating the *Ncol—Hgal* fragment (193 bp) of the pET11d carrying the enzyme $(S_{k2}D)$, the *Hgal-BamHI* fragment (1,359 bp) of pET11d carrying the enzyme $(S₁D)$, and the BamHI-Ncol fragment (5.6 kbp) of pETCaMKKa. pET11d carrying cDNAs for the enzymes $(S_{52}D/S_{74}D/S_{458}D)$ and $(S_{52}D/S_{74}D/S_{475}D)$ were constructed by ligating the *BamHI-Sfil* fragment (6.6 kbp) of the pETlld carrying the enzyme $(S_{52}DS_{74}D)$ and the *Sfil-BamHI* fragment (600 bp) of the pET11d carrying the enzymes $(S_{458}D)$ and $(S_{475}D)$, respectively. A pETlld carrying a cDNA for the enzyme $(S_{52}D/S_{14}D/S_{154}D/S_{175}D)$ was constructed by ligating the *BglR-Bgin* fragment (1,519 bp) of the pETlld carrying the enzyme $(S_{50}D/S_{74}D/S_{458}D)$ and the *BglII-BglII* fragment (5.7 kbp) of pET11d carrying the enzyme $(S_{475}D)$. pET11d carrying cDNA, for the mutant enzymes $(S_{62}D/S_{43}D)$ and $(T_{108}D/S_{458}D)$ were constructed by ligating the *BamHI-Sfil* fragment (6.6 kbp) of the pETlld carrying the enzymes $(S_{\kappa 0}D)$ and $(T_{108}D)$, respectively, and the *Sfil-BamHI* fragment (600 bp) of the pET11d carrying the enzyme $(S_{\text{ass}}D)$. pET11d carrying cDNA, for the enzymes $(S_{\kappa9}D/S_{47\kappa}D)$, $(S_{74}D/S_{475}D)$, and $(T_{106}D/S_{475}D)$ were constructed by ligating the *BamHI-Sfil* fragment (6.6 kbp) of the pETlld carrying the enzymes $(S_{62}D)$, $(S_{74}D)$, and $(T_{106}D)$, respectively, and the *Sfil-BamHl* fragment (600 bp) of the pETlld carrying the enzyme $(S_{476}D)$. All mutations were confirmed by the dideoxynucleotide chain-termination method *(14)* with a DNA sequencer (LI-COR model 4000L). *E. coli* strain $BL21(DE3)$ (15) was transfected with a pET11d carrying cDNA for a mutated enzyme and grown to an $A_{\rm em}$ value between 0.6 and 1.0 at 30°C in 50 ml of M9ZB medium containing 200 μ g/ml ampicillin. Isopropyl β -D-thiogalactoside (IPTG) was then added to a final concentration of 1 mM. After 2.5 h, the bacteria were harvested by centrifugation, washed in buffered saline, suspended in 1 ml of 20 mM Tris-HCl (pH 7.5 at 4°C) containing 1 mM dithiothreitol, 1 mM EGTA, 1 mM EDTA, 1 mM phenylmethylsulfonyl fluoride, and $10 \mu g/ml$ each of microbial protease inhibitors (pepstatin A, leupeptin, antipain, and chymostatin), and then disrupted by sonic oscillation. The supernatants were obtained by centrifugation, and the mutated enzymes were purified to apparent homogeneity by SDS-PAGE, as shown in Fig. 1, by streptomycin treatment, and by chromatographies on DE52 and calmodulin-Sepharose columns as described previously *(11).*

Phosphorylation Reaction—The phosphorylation of proteins or peptides by protein kinases was carried out at 30°C in the standard phosphorylation mixture comprising 50 mM Mops-NaOH (pH 7.0 at 30'C), 2 mM dithiothreitol, 5 mM Mg(CH₃COO)₂, 0.1 mM nonradioactive or $[\gamma$ -³²P]ATP, 1 μ M calmodulin, 0.1 mM EGTA, 0.2 mM CaCl₂, and the indicated amounts of proteins or peptides. After incubation for the indicated times, the incorporation of $[^{32}P]$ phosphate into the protein substrates was determined by the 3MM paper method of Corbin and Reimann *(16),* except that the filter papers were washed with ice-cold 10% trichloroacetic acid containing 2 mM ATP. The incorporation of $[{}^{32}P]$ phosphate into the peptide substrates was determined by the phosphocellulose paper method of Roskoski *(17).*

Other Procedures—The concentration of calmodulin was determined spectrophotometrically using an absorption coefficient at A_{280} (1 mg/ml) of 0.21 (18) and a molecular weight of 16,700 *(19, 20).* Other proteins were determined by the method of Lowry *et al (21),* as modified by Peterson, *(22)* using bovine serum albumin as a standard. The amount of CaM-kinase $IV(K_{71}R)$ was corrected for an overestimation by a factor of 1.15 made by the Lowry's method *(23).* SDS-PAGE was carried out according to the method of Laemmli *(24).* Gel overlay assay by biotinylated calmodulin was performed as described by Kincaid *et al. (25).*

RESULTS

As reported in the preceding paper (3), the activity of CaM-

kinase kinase α toward the PKIV peptide is increased by incubation with PKA in the presence of Ca2+/calmodulin under phosphorylation conditions, but decreased by incubation with PKA in the absence of Ca2+/calmodulin, and six phosphorylation sites, one for autophosphorylation (Ser²⁴) and five for phosphorylation by PKA (Ser⁵², Ser⁷⁴, Thr¹⁰⁸, Ser⁴⁵⁸, and Ser⁴⁷⁵) are present in CaM-kinase kinase α . The phosphorylation-activity relationship was investigated by the biochemical study, but a definite conclusion could not be drawn. To specify which of the six phosphorylation sites are involved in changes in enzyme activity, 25 mutant enzymes, in which the phosphorylatable serine and/or threonine residues were replaced with alanine to prevent phosphorylation or with aspartate or glutamate to mimic phosphorylation, were prepared by site-directed mutagenesis, and the respective mutant enzymes expressed in *E. coli* were purified to apparent homogeneity by SDS-PAGE, as shown in Fig. 1. Among the 25 mutant enzymes, 22 gave single protein bands at a position corresponding to the wild-type enzyme, and the other three, CaM-kinase kinases $(S_{24}D)$, $(S_{24}D/S_{458}D)$, and $(S_{24}D/S_{52}D/S_{475}D)$, in which Ser²⁴ was replaced with aspartate, gave single bands that migrated more slowly. SDS-PAGE analysis followed by autoradiography of the wild-type enzyme autophosphorylated by incubation with $[\gamma^{32}P]ATP$ in the presence of $Ca^{2+}/$ calmodulin under the phosphorylation conditions gave a similar slower migrating radioactive band (data not shown), indicating that mutant enzymes in which Ser^{24} is

Table I shows the specific activities of the wild-type and 25 mutant CaM-kinase kinases *a* toward PKTV peptide before and after preincubation for 30 min at 30'C with or without PKA in the presence or absence of Ca²⁺/calmodulin under phosphorylation conditions. None of the 26 CaMkinase kinases showed a significant change in activity following incubation in the absence of Ca²⁺ and PKA, indicating that the enzymes were stably maintained under the experimental conditions.

The activities of the 25 mutants of CaM-kinase kinase *a* toward PKTV peptide measured without preincubation are compared with that of the wild-type enzyme, as shown in Fig. 2. Among the 25 mutant enzymes, 9 showed significantly lower activities than the wild-type enzyme. Among them, 8 mutants contained aspartate or glutamate as a substitute for Thr¹⁰⁶, aspartate for Ser⁴⁵⁸, or both. The fact that all of mutants containing aspartate or glutamate for Thr¹⁰⁸, or aspartate for Ser⁴⁵⁸ showed low activities as compared with the wild-type enzyme suggests that the phosphorylation of Thr¹⁰⁸ or Ser⁴⁶⁸ causes a decrease in the enzyme activity toward PKIV peptide. The activity of the double mutant $(T_{108}D/S_{408}D)$, in which both Thr¹⁰⁶ and $Ser⁴⁵⁸$ were replaced with aspartate, was only 28% of that of

 $(S_{456}A)$, $(S_{456}D)$, $(S_{456}E)$, $(S_{475}A)$, $(S_{475}D)$, $(S_{475}E)$, $(S_{52}D/S_{74}D)$, $(S_{24}D/$ $(\mathbf{S}_{\mathbf{4}S}\mathbf{D})$, $(\mathbf{T}_{106}\mathbf{D}/\mathbf{S}_{\mathbf{4}S}\mathbf{D})$, $(\mathbf{S}_{\mathbf{5}4}\mathbf{D}/\mathbf{S}_{\mathbf{4}4}\mathbf{D})$, $(\mathbf{S}_{\mathbf{5}2}\mathbf{D}/\mathbf{S}_{\mathbf{4}75}\mathbf{D})$, $(\mathbf{S}_{\mathbf{7}4}\mathbf{D}/\mathbf{S}_{\mathbf{4}75}\mathbf{D})$, $(T_{100}D/S_{475}D)$, $(S_{24}D/S_{62}D/S_{475}D)$, $(S_{52}D/S_{74}D/S_{475}D)$, and $(S_{52}D/S_{74}D/$ S_{odd} D/S₄₇₀D), were purified as described under "EXPERIMENTAL PROCEDURES," subjected to SDS-PAGE on a 7.5% acrylamide gel, and then stained with Coomassie Brilliant Blue R-250.

the wild-type enzyme, much lower than the activities of the respective single mutants ($T_{108}D$) and (S₄₅₈D), whose activities were 73 and 69%, respectively, of the wild-type enzyme. These data indicate the synergistic effect of the phosphorylation of Thr¹⁰⁸ and Ser⁴⁵⁸. In contrast to the $S_{\text{ass}}D$ mutant, $S_{\text{avg}}E$ and $S_{\text{avg}}A$ showed almost the same activity as the wild-type enzyme. This suggests that the substitution of glutamate for Ser⁴⁶⁸ does not mimic the phosphorylation of Ser⁴⁵⁸, although the substitution of aspartate does. The

 $S_{74}A$, $S_{74}D$, and $S_{74}D/S_{475}D$ mutants showed significantly higher activities than the wild-type enzyme, suggesting that the phosphorylation of Ser⁷⁴ causes an increase in enzyme activity, and that the activation is probably due to the conversion of this serine to another residue because replacing Ser⁷⁴ not only with aspartate, but also with alanine caused an increase in the enzyme activity. The fact that the activity of the enzyme $(S_{52}DS_{74}DS_{458}D)$ was almost the same as that of the enzyme $(S_{\mu s}D)$ suggests that the inhibi-

TABLE I. Specific activities of CaM-kinase kinase α mutants toward PKIV peptide. Approximately 20 μ g/ml of purified wild-type and mutated CaM-kinase kinase α , in which alanine (A), aspartate (D), and glutamate (E) residues were substituted for the phosphorylatable serine (S) or threonine (T) residues, were preincubated at 30'C in the standard phosphorylation mixture containing nonradioactive 0.1 mM ATP with or without 0.5 μ g/ml PKA in the presence or absence of Ca²⁺/calmodulin, as indicated. Before and after preincubation for 30 min, a 10- μ l aliquot was incubated at 30°C for 1 min in a final volume of 50 μ l of phosphorylation mixture containing 0.2 mM PKIV peptide and 0.1 mM [y-²²P]ATP (about 200 cpm/pmol). The incorporation of ^{[32}P]phosphate into the peptide was determined by the phosphocellulose paper method. The results are expressed as the specific activities (nmol/min/mg) and each value represents the mean \pm SD of four or five independent determinations.

tory effect of the phosphorylation of Ser⁴⁵⁸ is dominant over the stimulatory effect of the phosphorylation of Ser⁷⁴. Mutant enzymes in which two other phosphorylatable sites, Ser²⁴ and Ser⁴⁷⁵, were each replaced with aspartate, showed almost the same activity as the wild-type enzyme, suggesting that the phosphorylation of Ser²⁴ or Ser⁴⁷⁵ causes no significant change in the enzyme activity. The activity of the mutant $(S_{\alpha}D)$, in which another of the six residues identified as phosphorylation sites (3) , Ser⁵², was replaced with aspartate, was somewhat lower than that of the wild-type enzyme, but the activities of the double and triple mutants $(S_{52}DS_{74}D), (S_{52}DS_{475}D), (S_{24}DS_{52}DS_{475}D),$ and $(S_{52}DS_{74}DS_{475}D)$ containing aspartate in place of Ser⁵² were almost the same as that of the wild-type enzyme, suggesting that the phosphorylation of Ser^{52} does not always result in a decrease in enzyme activity.

Figure 3 shows the effects of the substitutions of alanine, aspartate, or glutamate residues for the phosphorylatable serine and/or threonine residues on the increase in en-zyme

Fig. **2. The effects of substitutions of Ala, Asp, or Glu for the phosphorylatable Ser or Thr residues of CaM-kinase kinase a on the activity toward P&IV peptide.** The ratios of the specific activities toward PKTV peptide of the indicated enzymes to that of the wild-type enzyme (283 \pm 22 nmol/min/mg) were calculated from the results shown in Table **I.**

Fig. 3. **Changes in the activities of the CaM-kinase kinase a mutants by autophosphorylation.** The ratios of the specific activities toward PKTV peptide of the indicated wild-type and mutant enzymes preincubated in the presence of Ca²⁺/calmodulin $[fourth column (+Ca²⁺)$ of Table I] to those in its absence [second column $(-Ca²⁺)$ of Table I] (solid bars), and the ratios of the activities of the indicated enzymes $(+Ca²⁺$ column of Table I) to that (365 nmol/min/mg) of the wildtype enzyme preincubated in the presence of Ca²⁺/calmodulin (dotted bars) were calculated from the results shown in Table I.

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phorylation. Thus, the effect of the phoshorylation of Ser⁷⁴ on the activation of the enzyme appeared to be additive to that of the autophosphorylation. The activities of the mutated enzymes containing aspartate in substitution for Ser⁴⁶⁸, such as the enzymes $(T_{108}DS_{458}D)$, $(S_{52}DS_{74}DS_{458}D)$ S_{475} D), (S₂₄D/S₄₅₈D), and (S4₅₈D), after autophosphorylation were much lower than that of the wild-type enzyme (Table I), indicating that the decrease in enzyme activity caused by the substitution of aspartate for Ser⁴⁵⁸ continues after autophosphorylation. Thus, the phosphorylations of Thr¹⁰⁸ and Ser⁴⁵⁸ appear to cause a decrease in the enzyme activity toward PKTV peptide, and that the decrease caused by the phosphorylation of Thr^{108} is canceled out by autophosphorylation of the enzyme while that of $Ser⁴⁵⁸$ is not.

Figure 4 shows the effects of the substitutions on the increase in enzyme activity caused by incubation with PKA in the presence of Ca2+/calmodulin under the phosphorylation conditions, where the enzymes should be phosphorylated by itself as well as by PKA. The wild-type enzyme was activated about 2-fold by incubation with PKA under these conditions. Among the 25 mutated enzymes, all four mutants that contained aspartate or glutamate in substitution for Thr¹⁰⁸, $(T_{108}D)$, $(T_{108}E)$, $(T_{108}D/S_{458}D)$, and $(T_{108}D/$ S_{475} D), were activated to a greater extent than the wildtype enzyme, and the three mutants not containing aspartate for Ser⁴⁶⁸ showed almost the same activity as the wildtype enzyme after incubation with PKA. Another mutants, $(\dot{T}_{108}D/S_{458}D)$, containing aspartate for Ser⁴⁶⁸, showed much lower activity than the wild-type enzyme. This, together with the above results (Fig. 3), indicates that the effect of the phosphorylation of Thr¹⁰⁸ is also canceled out by incubation in the presence of Ca2+/calmodulin both will and without PKA. The other 21 mutants without aspartate or glu t amate substitutions for Thr 108 were activated to a similar or lesser extent than the wild-type enzyme, and among them, mutants containing alanine or aspartate substitutions for Ser⁷⁴, (S_n, A) , (S_n, D) , $(S_n, D/S_n, D)$, $(S_{74}DS_{475}D), (S_{52}DS_{74}DS_{475}D),$ and $(S_{52}DS_{74}DS_{455}D/S_{475}D),$ showed a tendency to be less activated by incubation with PKA in the presence of $Ca^{2+}/calmodulin$ than the wild-type enzyme. This result, together with the previous observation

that a significant amount of phosphate is incorporated into Ser⁷⁴ upon incubation with PKA in the presence of Ca²⁺/ calmodulin (3) , indicates that the phosphorylation of Ser⁷⁴ may be involved to some extent in the activation of CaMkinase kinase α by PKA in the presence of Ca²⁺/calmodulin. The fact that the activities after incubation with PKA in the presence of $Ca²⁺/calmodulin$ of all mutants containing aspartate for Ser⁴⁶⁸, $(S_{\text{ass}}D)$, $(T_{\text{obs}}DS_{\text{ass}}D)$, $(S_{\text{ss}}DS_{\text{res}}DS_{\text{ass}}D)$, and $(S_{52}DS_{74}DS_{458}DS_{475}D)$, were much lower than that of the wild-type enzyme, indicates that the decrease in enzyme activity that occurs upon the phosphorylation of Ser⁴⁶⁸ is not canceled by incubation with PKA in the presence of Ca2+/calmodulin, as well as without PKA as described above. The activations of mutant enzymes $(S_{\kappa\rho}D/\sqrt{S_{\kappa\rho}D})$ $S_{74}DS_{458}D$) and $(S_{52}DS_{74}DS_{458}DS_{475}D)$ by incubation without PKA in the presence of Ca^{2+}/cal calmodulin were significantly higher than that of the wild-type enzyme (Fig. 3), but the activation by incubation with PKA were almost the same as that of the wild-type enzyme (Fig. 4), supporting the above discussion that the phosphorylation of Ser^{74} is involved in the activation of the enzyme by PKA in the presence of Ca2+/calmodulin.

Figure 5 shows the effects of the mutations on the decrease in the enzyme activity caused by incubation with PKA in the absence of $Ca²⁺/calmoduli$ under the phosphorylation conditions, where autophosphorylation should not occur. The activity of the wild-type enzyme decreased to two-thirds of its original activity upon phosphorylation by PKA under the experimental conditions. Among the 25 mutants, 5 did not show significant decreases in enzyme activity after incubation: $(T_{108}DS_{468}D)$ showed an increase, $(S_{458}A)$ showed no change, and $(\overline{T_{106}}E)$, $(\overline{T_{108}}D)$, and $(\overline{T_{108}}A)$ showed little, if any, decrease. Four of the 5 mutants, $(T_{108}A)$, $(T_{108}D)$, $(T_{108}E)$, and $(T_{108}D/S_{458}D)$, contained substitutions of nonphosphorylatable amino acids for Thr¹⁰⁸. These results, together with the previous observation that a significant amount of phosphate is incorporated into Thr¹⁰⁸ upon incubation with PKA (3) , indicate the involvement of the phosphorylation of Thr¹⁰⁸ in the decrease in enzyme activity caused by incubation with PKA in the absence of $Ca^{2+}/calmodulin$. The other mutant, $(S₄₅₉A)$,

Fig. **4. Changes in the activities of the CaM-kinase kinase a mutants by PKA (plus Ca*7calmodulin).** The ratios of the specific activities toward PKTV peptide of the indicated enzymes preincubated in the presence of PKA and Ca²⁺/calmodulin flast col umn $(+Ca²⁺/+PKA)$ of Table I] to those in their absence [second column $(-Ca²⁺)$ of Table I] (solid bars), and the ratios of the activities of the indicated enzymes $(+Ca^{2+}/+PKA)$ column of Table I) to that (546 nmol/min/mg) of the wild-type enzyme preincubated in the presence of PKA and Ca²⁺/calmodulin (dotted bars) were calculated from the results shown in Table I.

which contained no Thr¹⁰⁶ substitution, not only showed no significant decrease in activity, but also the highest activity after incubation among the wild-type and 25 mutant enzymes. Another mutant enzyme containing a substitution for Ser⁴⁵⁸, $(S_{458}E)$, showed the second highest activity, although its activity decreased somewhat after incubation with PKA. In contrast to these two mutants, $(S_{468}D)$ was inactivated to the same extent as the wild-type enzyme by incubation with PKA- These results, together with the above finding that the substitution of glutamate for Ser⁴⁵⁸ does not mimick the phosphorylation of Ser⁴⁵⁸ while substitution by aspartate does, indicate that blocking the phosphorylation of Ser⁴⁵⁸ inhibits the inactivation of the enzyme by PKA in the absence of $Ca^{2+}/calmodulin$. Thus, the phos-

phorylation of Ser⁴⁵⁸ appears to promote either the phosphorylation of Thr¹⁰⁸ by PKA or the inactivation of the enzyme by the phosphorylation of Thr¹⁰⁸, or both. As shown in the preceding paper (3), CaM-kinase kinase α undergoes phosphorylation at Ser⁴⁵⁸ by PKA in the absence of Ca^{2+/} calmodulin, but little, if any, phosphorylation at Ser⁴⁵⁸ in its presence, and the rate of the phosphorylation of Thr¹⁰⁸ by PKA in the absence of Ca²⁺/calmodulin is one-fifth of that in its presence, supporting the concept that the phosphorylation of Ser⁴⁵⁸ promotes the phosphorylation of Thr¹⁰⁸ by PKA. On the other hand, the fact that the activity of the double mutant $(T_{108}D/S_{458}D)$, which is approximately 26% that of the wild-type enzyme, is much lower than the activities of the respective single mutants $(T_{108}D)$ and $(S_{168}D)$,

tion mixture containing $2 \mu M$ calmodulin, 0.1 mM [γ -²²P]ATP (about 1,000 cpm/pmol), and 50 µg/ml of CaM-kinase IV(K_nR), and the incorporation of [³²P]phosphate into the CaM-kinase IV was determined by the 3MM paper method. (A) The ratios of the activities of the indicated enzymes to that of the wild-type enzyme without preincubation. (B) The ratios of the activities of the indicated enzymes preincubated in the presence of PKA and Ca^{2+}/cal odulin to those in their absence (solid bars), and the ratios of the activities of the indicated enzymes to that of the wild-type enzyme preincuodulin to those in their absence (solid bars), and the ratios of the activities of the indicated enzymes to that of the wild-type enzyme preincu-
hated in the presence of PKA and Ca²⁺/calmodulin (dotted bars) (C) The rat presence of PKA to those of the enzymes preincubated in its absence (solid bars), and the ratios of the activities of the indicated enzymes to that of the wild-type enzyme preincubated in the presence of PKA (dotted bars).

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dithiothreitol, and 0.1 mM EDTA. Aliquots of 2- μ l were incubated at 30'C for 1 min in a final volume of 50 *yl* of the standard phosphorylawhich are 73 and 69%, respectively, of the wild-type enzyme (Fig. 2). This supports the concept that the phosphorylation of Ser⁴⁵⁸ promotes the inactivation of the enzyme when Thr¹⁰⁸ is phosphorylated by PKA. Thus, the phosphorylation of Ser⁴⁵⁸ appears not only to promote the rate of the Thr¹⁰⁸ phosphorylation by PKA, but also to enhance the extent of the inactivation of the enzyme by the Thr¹⁰⁸ phosphorylation by PKA. The preceding paper (3) demonstrates that the rate of phosphorylation of Ser⁴⁶⁸ in CaM-kinase kinase α by PKA (0.008 mol phosphate/mol of CaM-kinase kinase α /min) in the presence of Ca²⁺/calmodulin is much slower (about 36-fold) than that in its absence (0.29 mol phosphate/mol of enzyme/min). Thus, a primary factor in determining whether the activity of CaM-kinase kinase α increases or decreases after phosphorylation by PKA appears to be the phosphorylation of Ser⁴⁶⁸ by PKA, which occurs in the absence but not the presence of Ca2+/calmodulin.

Since the incubation of CaM-kinase kinase α with PKA in the presence of Ca2+/calmodulin results in an increase in activity toward PKTV peptdde, but a decrease in activity toward CaM-kinase IV (3) , the effects of the substitutions on the activity toward CaM-kinase IV were examined as shown in Fig. 6. The enzymes $(S_{74}A)$ showed a higher activity and $(T_{108}D)$ a lower activity toward CaM-kinase IV (Fig. 6A) as well as PKIV peptide (Fig. 2) than did the wild-type enzyme. On the other hand, $(S_{458}D)$, which showed much lower activity toward PKTV peptide than the wild-type enzyme (Fig. 2), had almost the same activity toward CaMkinase IV as the wild-type enzyme, suggesting that the phosphorylation of Ser⁴⁵⁸ does not play an important role in controlling the activity toward CaM-kinase IV, in contrast to the activity toward PKTV peptide. In contrast to the activity toward PKTV peptide, the activity of CaM-kinase kinase α toward CaM-kinase IV is not significantly affected by incubation in the presence of Ca^{2+}/cal calmodulin (autophosphorylation), and decreased by incubation with PKA in both the presence and absence of Ca^{2+}/cal calmodulin (3). Hence, the effects of substitutions for the phosphorylatable serine or threonine residues on changes in the enzyme activity toward CaM-kinase TV through phosphorylation by PKA in the presence (Fig. $6B$) and absence (Fig. $6C$) of Ca²⁺/ calmodulin were examined. The activity of the wild-type enzyme decreased to approximately one-half of its original activity after incubation with PKA in both the presence and absence of $Ca^{2+}/calmodulin$, and the changes in the activities of the wild-type and mutant enzymes after incubation with PKA in the presence of Ca²⁺/calmodulin were almost identical to those in its absence. The activities of $(T_{108}A)$ and $(T_{108}D)$ did not decrease significantly after incubation with PKA in the presence or absence of Ca²⁺/calmodbadon what I first in the presence of absence of 0a 7 cannou-
ulin, indicating that the phosphorylation of Thr¹⁰⁸ is involved in the decrease in the activity toward CaM-kinase TV by the incubation with PKA in the presence or absence of Ca²⁺/calmodulin. The fact that the activity of $(T_{108}D)$ after incubation with PKA was much higher than that the wildtype enzyme, although lower than that of $(T_{\text{tot}}A)$, indicates type enzyme, annough lower man mat or $(1_{106}A)$, multales wiat aspartate in place of the comp partly minites phos-
phorylated Thr¹⁰⁸ in enzyme action catalyzing the phosphorylation of CaM-kinase IV. The mutant $(S_{\text{avg}}D)$ showed essentially the same activity, even after incubation with PKA in the presence or absence of $Ca²⁺/calom$ as the r KA in the presence or absence or Ca /Camioutum, as the
wild-type enzyme. Thus, the phosphorylation of Ser⁴⁶⁸ by

PKA, which is important for controlling the PKIV peptidephosphorylating activity as described above, appears not to be important for controlling the CaM-kinase TV-phosphorylating activity. An alternative possibility is that aspartate in place of Ser⁴⁵⁸ does not mimic phosphorylated Ser⁴⁵⁸ in the phosphorylation of CaM-kinase IV. None of the 10 mutated enzymes in which each of the five serine or threonine residues identified as PKA phosphorylation sites (3) was replaced by asparate or alanine was inactivated more by incubation with PKA, and all showed significantly lower activities after incubation with PKA than did the wild-type enzyme.

To obtain a clue as to the mechanism for the changes in

Fig. 7. **The effects of the concentrations of peptide or protein substrate and ATP on the activity of CaM-kinase kinase a (T1MD/S4I8D).** (A and B) Approximately 200 ng of wild-type CaM-kinase kinase α (O) or mutant $(T_{100}D/S_{460}D)$ (\bullet) was incubated in a final volume of 50μ of the standard phosphorylation mixture containing 0.2 mM [γ -²²P]ATP (about 50 cpm/pmol) and various concentrations of PKIV peptide (A), or 0.2 mM PKIV peptide and various concentrations of $[\gamma^{22}P]$ ATP (about 50 cpm/pmol) (B) at 30°C for 1 min, and the incorporation of [³²P]phosphate into the peptide was determined by the phosphocellulose paper method. (C and D) Approximately 2.4 ng of wild-type CaM-kinase kinase α (o) or mutant $(T_{\text{top}}DS_{\text{max}}D)$ (\bullet) was incubated in a final volume of 50 μ l of the standard phosphorylation mixture, except that the concentration of calmodulin was 5 μ M, containing 0.1 mM [γ -²²P]ATP (about 1,000 cpm/pmol) and various concentrations of CaM-kinase IV(K₇,R) (C), or 50 μ g/ml (0.94 μ M) of CaM-kinase $IV(K_{71}R)$ and various concentrations of [y- ^{27}P]ATP (about 1,000 cpm/pmol) (D) at 30°C for 1 min, and the incorporation of [^{*}P]phosphate into CaM-kinase IV was determined by the 3MM paper method.

TABLE II. **Summary of kinetic parameters of CaM-kinase kinase** α **(T₁₀₈D/S₄₅₈D). Kinetic parameters were obtained from the** double-reciprocal plots shown in Fig. 7. The V^{max} values were calculated from the apparent V_{max} values on the basis of the Michaelis equation $v = V/(1 + K/J[S])$.

	K for substrate (μM)	K_{m} for ATP (μM)	max (nmol/min/mg)
	Assayed with PKIV peptide		
Wild type	280	250	1,840
T_{108} D/S ₄₅₈ D	280	770	880
	Assayed with CaM-kinase $IV(K_nR)$		
Wild type	$0.46(24 \text{ µg/ml})$	40	1,910
T_{108} D/S ₄₅₈ D	1.2 $(63 \mu g/ml)$	87	2.270

the activity of CaM-kinase kinase α induced by PKA, the kinetic properties of $(T_{108}D/S_{458}D)$, which always showed the lowest activity toward PKTV peptide (Table I and Figs. 2- 5), were examined in comparison with the wild-type enzyme using PKTV peptide and CaM-kinase IV as substrates, as shown in Fig. 7, and the kinetic parameters are summarized in Table II. The substitutions of aspartate for both Thr¹⁰⁶ and Ser⁴⁵⁸ caused an approximately 3-fold increase in the K_m value for ATP but no change in the K_m for PKIV peptide, and a decrease in the V_{max} by about onehalf, when the enzyme was assayed with PKTV peptide as a substrate, and 2- to 3-fold increases in the K_m values for both ATP and CaM-kinase IV and an increase in the *Vmax,* when assayed with CaM-kinase $IV(K_{71}R)$ as a substrate. As reported in the preceding paper (3), incubation of CaMkinase kinase α with PKA in the absence of Ca²⁺/calmodulin results in a significant increase in the K_m for ATP but no increase in the K_m for the peptide substrate, and a slight decrease in the \hat{V}_{max} , when the enzyme is assayed with PKIV peptide, and $3-$ to 4-fold increases in both K_m values for ATP and the protein substrate and a 1.7-fold increase in the V_{max} , when assayed with CaM-kinase IV(K₇₁R). In contrast, incubation of the enzyme with PKA in the presence of Ca2+/cahnodulin results in a marked increase (approximately 3-fold increase) in *Vmia* when the enzyme is assayed with PKTV peptide (3). Thus, the kinetic properties of the mutant enzyme $(T_{108}DS_{458}D)$ are very similar to those of the enzyme phosphorylated by PKA in the absence of Ca^{2+} calmodulin, providing further support that the phosphorycannocum, providing further support that the phosphory-
lations of Ser⁴⁵⁸ and Thr¹⁰⁸ are important contributors to the change in activity caused by incubation with PKA in the absence of Ca2+/calmodulin.

DISCUSSION

The activity of CaM-kinase kinase α toward PKIV peptide is altered when the enzyme undergoes autophosphorylation or is phosphorylated by PKA, and six phosphorylation sites, one for autophosphorylation $(Ser²⁴)$ and five for phosphorylation by PKA (Ser⁵², Ser⁷⁴, Thr¹⁰⁸, Ser⁴⁵⁸, and Ser⁴⁷⁵) have been identified (3). Changes in activity produced by amino acid substitutions at the six phosphorylatable serine and threonine residues (Fig. 2) suggest that the phosphorylation of Ser⁷⁴ causes an increase and that the phosphorylations of Thr¹⁰⁸ and Ser⁴⁵⁸ synergistically cause a decrease in α activity. The replacement of Ser^{24} , which has been identified as the major autophosphorylation site (3), by amino acids such as alanine or aspartate had no significant affect either the enzyme activity (Fig. 2) or the extent of activation of the enzyme through autophosphorylation (Fig. 3), indicating that the phosphorylation of Ser²⁴ is not involved in the activation of the enzyme. These results, together with the previous observation that the amount of phosphate incorporated into Ser²⁴ upon autophosphorylation accounts for only 30% of the total phosphate incorporated into the enzyme (3), provide further support for the contention that the gradual activation of CaM-kinase kinase α upon autophosphorylation is due to a conformational change induced by phosphorylation at many unidentified sites on the enzyme (3). The enzyme $(S_{74}D)$, whose activity was 1.4-fold higher than that of the wild-type enzyme (Fig. 2), was further activated upon autophosphorylation to the same extent (about 1.3-fold) as the wild-type enzyme (Fig. 3 and Table I), indicating that the activation of the enzyme that occurs upon phosphorylation of Ser⁷⁴ is additive with the activation produced by autophosphorylation. The mutant enzymes containing the substitutions at Ser⁷⁴ were less activated upon phosphorylation by PKA in the presence of $Ca²⁺$ and calmodulin than the wild-type enzyme (Fig. 4). Therefore, the phosphorylation of Ser⁷⁴ appears to be involved in the activation of the enzyme by PKA, since a significant amount of phosphate has been demonstrated to be incorporated into $Ser⁷⁴$ upon phosphorylation by PKA in the presence of $Ca²⁺$ calmodulin (3).

Since the lowers activity produced by the substitution of Thr¹⁰⁸ with aspartate in $(T_{106}D)$ was increased much more greatly upon autophosphorylation (Fig. 3) and upon incubation with PKA in the presence of Ca^{2+}/cal calmodulin (Fig. 4), and more weakly decreased upon incubation with PKA in the absence of Ca^{2+}/cal calmodulin (Fig. 5), the decrease in the activity toward PKTV peptide caused by the phosphorylation of Thr¹⁰⁸ appears to be canceled by autophosphorylation as well as by phosphorylation by PKA in the presence or absence of Ca2+/calmodulin. In contrast, the lower activity produced by the substitution of Ser⁴⁵⁸ with appartate in (S_{avg})) increased upon autophosphorylation (Fig. 3) or upon incubation with PKA in the presence of $Ca²⁺/calmoduli$ n (Fig. 4), and decreased upon incubation with PKA in the absence of Ca^{2} /calmodulin (Fig. 5) to the same extent as the wild-type enzyme. Thus, the decrease in activity produced by the phosphorylation of Ser⁴⁶⁸, in contrast to Thr^{108} , appears to continue after autophosphorylation or phosphorylation by PKA. The double mutants $(T_{100}D/S_{458}D)$ almost always showed the lowest activities (Table I), which were significantly lower than the activities of $(S_{458}D)$, even after autophosphorylation (Fig. 3) or phosphorylations by PKA in the presence (Fig. 4) or absence (Fig. 5) of Ca^{2+}/cal calmodulin, suggesting that the inhibitory effect of the phosphorylation suggesung untuk internationally enter of the phosphorylation
of Thr¹⁰⁸ in an enzyme phosphorylated at Ser⁴⁵⁸ was not canceled by autophosphorylation or phosphorylation by canceled by admphosphorylation of phosphorylation by
PKA The rate of phosphorylation of Ser⁴⁶⁸ by PKA in the absence of Ca2+/calmodulin was more than 1 order of magnitude greater than that in the presence of $Ca²⁺/calom$ calmodulin (3) , indicating that binding of Ca²⁺/calmodulin to the enzyme inhibits the phosphorylation of Ser⁴⁶⁸ by PKA. Conversely, it has been reported that the phosphorylation of CaM-kinase kinase α by PKA in the absence of Ca²⁺/calmodulin inhibits the binding of calmodulin to the enzyme, odulin inhibits the binding of calinodulin to the enzyme,
and that this effect is lost when Ser⁴⁵⁶ is replaced by alaand that thus enect is lost when ser¹¹ is replaced by ala-
nine (d). Thus, the phosphorylation of Ser⁴⁶⁸ in CaM-kinase kinase α by PKA appears strongly to suppress the binding of Ca2+/calmodulin to the CaM-kinase kinase, and *vice*

versa, presumably because Ser⁴⁶⁸ is located in the calmodulin-binding domain (Fig. 1A). Our attempts to confirm that the phosphorylation of Ser⁴⁵⁸ in CaM-kinase kinase α by PKA abolishes the binding of calmodulin were unsuccessful, because it was difficult to phosphorylate only Ser⁴⁵⁸ sufficiently among the five PKA phosphorylation sites. On the other hand, the substitution of aspartate for Ser⁴⁵⁸, as in the case of the phosphorylation of Ser⁴⁵⁸, apparently abolishes the binding of $Ca^{2+}/calmodulin$ in the calmodulin overlay assay, as shown in Fig. 8. However, the mutant $(S_{458}D)$, as well as the wild-type or other mutant enzymes, was purified by affinity chromatography on calmodulin-Sepharose, and it showed no significant activity in the $\frac{1}{2}$ absence of Ca²⁺/calmodulin (data not shown) but significant activity in its presence as shown above, indicating that the $(S_{\mu\nu}$ D) can bind calmodulin. Furthermore, this indicates that the calmodulin overlay assay does not necessarily reflect the binding of calmodulin.

In contrast to the results obtained with PKTV peptide as a substrate, when the activity of CaM-kinase kinase α was measured with CaM-kinase IV as a substrate, only the substitution of aspartate for Thr¹⁰⁸ among the five phosphorylation sites for PKA resulted in a significant decrease in activity (Fig. $6A$), and only mutant enzymes in which Thr^{106} was replaced with other amino acids such as aspartate or alanine, underwent no significant changes in activity upon incubation with PKA in the presence (Fig. 6B) and absence (Fig. 6C) of Ca2+/calmodulin, while the wild-type and other mutant enzymes underwent significant decreases in activity. These results indicate that the phosphorylation of Thr¹⁰⁸ is most important for regulating the activity toward CaM-kinases IV and I of CaM-kinase kinase α by PKA, because the activities of CaM-kinase kinase α toward both CaM-kinases IV and I are regulated in a similar manner by

Fig. 8. Analysis of mutants of CaM-kinase kinase α by calmod**ulin overlay after SDS-PAGE.** Approximately 0.05 *\xg* samples of the wild type, mutant $(S_{\text{def}}D)$, and mutant $(S_{\text{def}}A)$ CaM-kinase kinase a were subjected to SDS-PAGE on a 7.5% acrylamide gel, and the separated proteins were transferred onto a polyvinylidene difluoride membrane (FluoroTrans, Pall Gelman Laboratory). The membrane was blocked with 5% nonfat milk for 30 min at 24'C, then incubated with $25 \mu g/ml$ biotinylated calmodulin for 60 min, followed by 2 μ g/ml avidin conjugated with peroxidase for 60 min, in 50 mM Tris-HCl (pH 7.5) containing 150 mM NaCl and 1 mM CaCl,. The positive bands that bound calmodulin were detected with diaminobenzidine tetrahydrochloride and H_2O_2 in the presence of $CoCl₂$

PKA (3). Thus, it appears that the phosphorylations of Thr¹⁰⁸ and Ser⁴⁵⁸ in CaM-kinase kinase α by PKA play a key role in the regulation of the activity toward PKTV peptide, but that the phosphorylation of only Thr¹⁰⁸ plays a key role in the regulation of the activity toward CaM-kinases IV and I.

A comparison of the kinetic properties of the wild-type enzyme and the mutant $(T_{108}D/S_{458}D)$, which almost always showed the lowest activity toward PKTV peptide, revealed that the substitutions of aspartate for both Thr¹⁰⁸ and Ser⁴⁶⁸ caused a large decrease in the V_{max} for PKIV peptide, but an increase in the V_{max} for CaM-kinase IV, and caused no change in the *Km* for PKTV peptide but a significant increase in the *Km* for CaM-kinase IV. These kinetic properties of $(T_{108}D/S_{458}\overline{D})$ are very similar to those of the enzyme phosphorylated by PKA in the absence of Ca²⁺/calmodulin (3). Thus, the mechanism of the changes in the activity of CaM-kinase kinase α caused by PKA appears to vary with the substrate used for assay.

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