

Transient Activation of Dihydropteridine Reductase by Ca2 -activated Proteolysis

Authors: Kawai, Kouji, Fujimoto, Kengo, Okamoto, Akira, Inaba, Akemi, Yamada, Hirotaka, et al.

Source: Zoological Science, 17(4): 437-443

Published By: Zoological Society of Japan

URL: https://doi.org/10.2108/0289-0003(2000)17[437:TAODRB]2.0.CO;2

BioOne Complete (complete.BioOne.org) is a full-text database of 200 subscribed and open-access titles in the biological, ecological, and environmental sciences published by nonprofit societies, associations, museums, institutions, and presses.

Your use of this PDF, the BioOne Complete website, and all posted and associated content indicates your acceptance of BioOne's Terms of Use, available at <u>www.bioone.org/terms-of-use</u>.

Usage of BioOne Complete content is strictly limited to personal, educational, and non - commercial use. Commercial inquiries or rights and permissions requests should be directed to the individual publisher as copyright holder.

BioOne sees sustainable scholarly publishing as an inherently collaborative enterprise connecting authors, nonprofit publishers, academic institutions, research libraries, and research funders in the common goal of maximizing access to critical research.

Transient Activation of Dihydropteridine Reductase by Ca²⁺-activated Proteolysis

Kouji Kawai, Kengo Fujimoto*, Akira Okamoto, Akemi Inaba, Hirotaka Yamada and Setsuko Katoh

Department of Biochemistry, Meikai University School of Dentistry, Sakado, Saitama 350-0283, Japan

ABSTRACT—A transient activation of dihydropteridine reductase (DHPR), which is the regenerating enzyme of tetrahydrobiopterin in the system of aromatic amino acid hydroxylases, was studied during the incubation of DHPR with Ca²⁺-activated protease, m-calpain. The DHPR subunit (29 k) was cleaved by m-calpain just before the 35th (Ser) and 48th (Val) residue from the N-terminus, generating two new fragments of 21 k and 19 k. By determining kinetic parameters, we found that 21 k and 19 k were more active than the native enzyme and that the activation of them was more remarkable and transient against the natural substrate of quinonoid dihydrobiopterin than against a synthesized substrate. Phosphorylation of DHPR by Ca²⁺/calmodulin-dependent protein kinase II controlled the sensitivity of the enzyme to the Ca²⁺ activated protease.

INTRODUCTION

Tetrahydrobiopterin (BH4) functions as a H-donor cofactor of aromatic amino acid hydroxylases (Nagatsu et al., 1972; Kaufman and Fisher, 1974), rate-limiting enzymes for producing monoamine neurotransmitters in the sympathetic nervous system, and becomes converted to guinonoid dihydrobioptrin (q-BH₂) through 4α -carbinolamine in the process. The regeneration of BH₄ from q-BH₂ by dihydropteridine reductase (DHPR; NADH: quinonoid dihydropteridine oxidoreductase [EC 1. 6. 99. 7]) with NADH allows this cofactor to function catalytically. Besides the biosynthesis of BH₄ from GTP by three enzymes (Katoh and Akino, 1986), GTP cyclohydrolase I (Hatakeyama et al., 1989), 6-pyruvoyltetrahydropterin synthase (Inoue et al., 1991), and sepiapterin reductase (SPR) (Sueoka and Katoh, 1982), the BH₄-recycling by DHPR is important to control the concentration of the cofactor in the cell. DHPR reduces with NADH various quinonoid dihydropterins derived from BH4 and other tetrahydropterins such as 6-methyl tetrahydropterin (6M-PH₄). DHPR has been purified to homogeneity from many sources (Craine et al., 1972; Hasegawa, 1977; Korri et al., 1977; Webber et al., 1978; Firgaira et al., 1981). The cDNAs of human (Dahl et al., 1987), mouse (Yang et al., 1996), and rat (Shahbaz et al., 1987) DHPR have been cloned, and their deduced amino acid sequences are almost identical.

The calcium ion plays a role as a second messenger with a regulatory involvement in many aspects of cellular signaling and regulates the activation of various protein kinases

* Corresponding author: Tel. +81-492-71-2503; FAX. +81-492-71-2503. E-mail: kengo@dent.meikai.ac.jp (Greengard, 1978) and proteases such as calpain. Recently, both DHPR and SPR, which are BH₄-recycling enzyme and BH₄-generating enzyme, respectively, were found to be phosphorylated by Ca²⁺-activated protein kinases (Katoh *et al.*, 1994). Calpain is a Ca²⁺-activated protease widely distributed in various organs such as skeletal muscle (Huston *et al.*, 1968; Dayton *et al.*, 1976), platelets (Phillips and Jakabova, 1997), axoplasm (Schlaepfer, 1974; Pant *et al.*, 1979), liver (Nishiura *et al.*, 1978), and heart (Waxman and Krebs, 1978; Drummond and Duncan, 1966). Calpain is inhibited by an endogenous inhibitor, calpastatin (Nishiura *et al.*, 1978). It is wellknown that calpain has a strict substrate specificity, and modulates functions of various physiologically active proteins by limited proteolysis (Suzuki, 1993).

We found, in this study, that DHPR is transiently activated by limited proteolysis by m-calpain.

MATERIALS AND METHODS

Chemicals

DHPR (sheep liver), Ca²⁺-activated protease (m-calpain), calpastatin, ferri-cytochrome c, 2,6-dichlorophenolindophenol (DCPIP), ATP, calmodulin, ethylene glycol-bis (β -aminoethyl ether)-N, N, N', N'-tetra acetic acid (EGTA) and protein markers were obtained from Sigma (U.S.A.). Nicotinamide adenine dinucleotide (reduced form; NADH) was obtained from Wako Pure Chem. (Japan), and polyvinylidene difluoride (PVDF) membrane (Immobilon-P^{SQ}), from Millipore (U.S.A.). Tetrahydrobiopterin (BH₄) and 6-methyl-5, 6, 7, 8-tetrahydropterin (6M-PH₄) came from Dr. Schricks Lab. (Switzerland). Sepiapterin reductase (SPR) and Ca²⁺/calmodulindependent protein kinase II were prepared from hemolysate (Sueoka and Katoh, 1982) and cerebral cortex (Yamauchi and Fujisawa, 1983), respectively, obtained from rats.

DHPR assay

The commercially obtained DHPR was further purified by HPLC using a Shodex DEAE-825 column (5×100 mm). DHPR was eluted by a linear gradient of 0-0.15 M KCl in 20 mM phosphate buffer, pH 7.5, at a flow rate of 0.5 ml/min. As the substrate of DHPR, 6-methyl quinonoid dihydropterin (q-MH₂) was generally prepared from 6M-PH4 by non-enzymatic oxidation with ferri-cytochrome c just before use. The amount of substrate reduced by DHPR in the presence of NADH was determined spectrophotometically as the amount of ferrocytochrome c (Katoh et al., 1970). The reaction mixture contained 50 mM Tris-HCl, pH 7.5, 0.05 mM ferri-cytochrome c, 0.05 mM NADH, 1 μ M 6M-PH₄, and 1 μ g DHPR in a final volume of 2 ml. The mixture without DHPR was incubated for 1 min after 6M-PH₄ had been added to form q-MH₂, and then the reaction was started by the addition of DHPR and monitored for 1 min at 550 nm (Hitachi Spectrophotometer U-3210). Assays were performed at 25°C. One unit (U) was defined as 1 µmole of cytochrome c reduced per min (Hasegawa, 1977). Concentration of tetrahydropterin was determined by spectrophotometric titration with DCPIP (Kaufman and Levenberg, 1959).

Proteolysis of DHPR by Ca²⁺-activated protease (m-calpain)

DHPR was incubated with m-calpain (0.05-1 U) in the reaction mixture containing 50 mM Tris-HCl (pH 7.5), 5 mM CaCl₂, and 5 mM 2-mercaptoethanol for 5–180 min at 25°C. After the incubation, SDS for a 0.1% final concentration was added to the reaction mixture, which was then boiled for 3 min. The resultant solution was applied to SDS-PAGE (12.5% gel) performed according to Laemmli (Laemmli, 1970). Proteins on the gel were stained with Coomassie brilliant blue G-250.

Analysis of N-terminal amino acid sequence

After electrophoresis, separated proteins on the gel were transferred to a PVDF membrane at 400 mA for 3 hr in 100 mM sodium tetraborate. The transferred protein fragments on the PVDF membrane stained with Coomassie brilliant blue G-250 were cut out and directly applied to a protein sequencer. The N-terminal amino acid sequences of the fragments were determined by the Edman method (Edman and Begg, 1967) with a gas-phase protein sequencer (Applied Biosystems model 477A) with the chemicals and program supplied by the manufacturer.

Phosphorylation of DHPR

DHPR (5 μ g) was added to a reaction mixture (final 100 μ l) containing 50 mM Tris-HCl buffer (pH 7.5), 0.5 μ g Ca²⁺/calmodulindependent protein kinase II, 0.5 μ M calmodulin, 100 μ M ATP, 2 mM magnesium acetate, and 0.15 mM CaCl₂, and incubated at 25°C for 10 min, under which conditions DHPR is maximally phosphorylated (Katoh *et al.*, 1994).

Protein measurement

Protein concentration was determined by measuring its absorbance at 280 nm with bovine serum albumin as a standard (Hitachi Spectrophotometer U-3210).

RESULTS

Activation of DHPR by m-calpain

Effects of m-calpain and calcium ion on the activity of DHPR were observed *in vitro*. As shown in Fig. 1a, a transient activation of DHPR was observed during the initial period of the incubation of DHPR with m-calpain in the presence of calcium ions. DHPR activity increased to about 150% after the incubation for 5 through 30 min, and then it decreased gradually; however, about 120% activity was maintained until 180 min. This activation of DHPR was not observed when

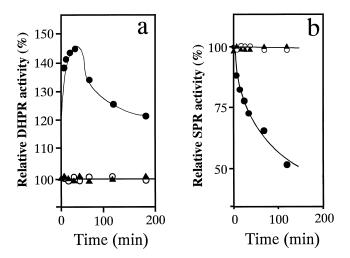


Fig. 1. Time-dependent activation of DHPR by m-calpain. (a) DHPR (10 μ g) was incubated with m-calpain (1 U) in the presence () or absence () of calpastatin (1 U) in the reaction mixture containing 5 mM CaCl₂, 5 mM 2-mercaptethanol, and 50 mM Tris-HCl buffer (pH 7.5) in a final volume of 100 μ l at 25°C. At various times, an appropriate volume containing 1 μ g DHPR was removed from the reaction mixture and measured for activity as described in the text. DHPR was incubated without m-calpain and calpastatin () as a control. (b) SPR was incubated in the presence or absence of m-calpain and calpastatin as described in (a), and assayed for activity by a published method (Katoh *et al.*, 1994).

Table 1. Effect of calcium ion on the activation of DHPR by m-calpain

Addition	DHPR activity (mU)	[%]
none	3.67	100
m-calpain	3.46	94
m-calpain+Ca ²⁺	5.21	142
m-calpain+Ca ²⁺ +EGTA	3.39	92
m-calpain+EGTA	3.39	92
m-calpain+Ca ²⁺ (5 min) +EGTA	3.78	103

DHPR (6 μ g) and m-calpain (1 U) were preincubated in the presence or absence of calcium ion and/or EGTA in the reaction mixture at 25°C for 5 min as described in Fig. 1a. In the last experiment in the table, the reaction mixture was treated with EGTA after preincubation of DHPR and m-calpain with calcium ions and further incubated for 5 min. DHPR equivalent to 1 μ g of the reaction mixture was assayed for activity by the method described in the text.

DHPR was incubated with m-calpain in the presence of calpastatin, an endogenous specific inhibitor of m-calpain (Nishiura *et al.*, 1978). However, activation of DHPR by m-calpain was not observed when calcium ions were omitted before or after the incubation (Table 1). These results indicate that DHPR was activated by m-calpain that was stimulated by calcium ion and that calcium ions were also necessary for activation of DHPR after the proteolytic function of m-calpain. On the other hand, although SPR was also a substrate of m-calpain, the activity of SPR was decreased by the incubation with m-calpain (Fig. 1b).

Limited proteolysis of DHPR by m-calpain

DHPR is a homodimeric enzyme (Webber *et al.*, 1978), and the molecular weight of the subunit was estimated by SDS-PAGE as 29 k in this experiment. Proteolytic effect of m-calpain on DHPR molecule was analyzed by SDS-PAGE. When DHPR was incubated with m-calpain in the presence of calcium ion, two new fragments A (21 k) and B (19 k) were initially produced from the DHPR subunit (29 k); but only fragment B was observed after 120 min (Fig. 2). The latter finally disappeared, too, by further incubation. These findings indicate that the appearance of fragments A and B were dependent on limited proteolysis with m-calpain and that fragment A seemed to decompose faster than fragment B.

Hydrolyzed sites of DHPR by m-calpain

After the incubation of DHPR with m-calpain for 5 min as shown in Fig. 2, fragments A and B were isolated by SDS-PAGE for analysis with a protein sequencer. The amino acid

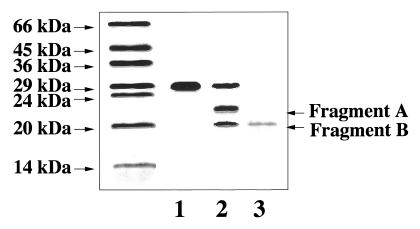


Fig. 2. Time-dependent proteolysis of DHPR by m-calpain. DHPR was incubated with m-calpain (1 U) at 25° C for 5 or 120 min as described in Fig. 1, and an appropriate volume containing 10 µg DHPR was applied to SDS-PAGE as described in the text. The leftmost lane indicates protein markers. Lanes 1, 2, and 3 indicate reaction mixtures incubated for 0, 5, and 120 min, respectively. Proteins on the gel were stained with Coomassie brilliant blue G-250.

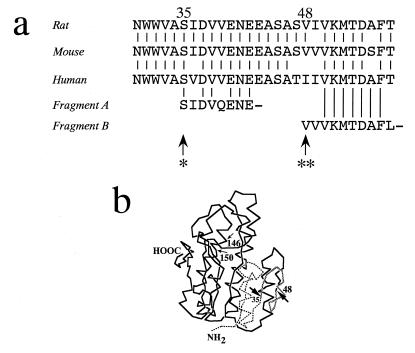


Fig. 3. Determination of cleavage sites of DHPR by m-calpain. (**a**) N-terminal amino acid sequences of fragment A and B determined by Edmen method (Edman and Begg, 1967) as described in the text with reference to the sequences of DHPR from various origins (Dahl *et al.*, 1987; Yang *et al.*, 1996; Shahbaz *et al.*, 1987). Asterisks indicate N-terminal sites of fragment A (*) and fragment B (**), respectively. (**b**) The position of cleavage sites in DHPR subunit. The X-ray data for rat DHPR were cited from the Protein Data Bank (PDB code 1DHR). Numbers 146 and 150 indicate the catalytic motif of DHPR (Tyr¹⁴⁶-X-X-Lys¹⁵⁰) for the pteridine substrate. The nicotinamide moiety of NADH is also located around the motif according to the data.

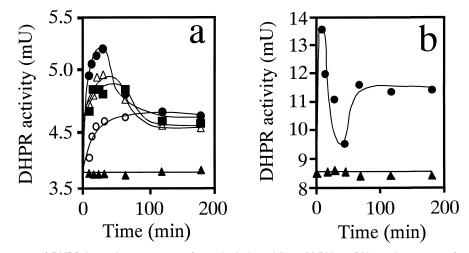


Fig. 4. Activation patterns of DHPR by various amounts of m-calpain by adding $6M-PH_4$ or BH_4 as the source of quinonoid dihydropterin substrate. DHPR (10 µg) was incubated with various amounts of m-calpain for various times at 25°C in the system (final 100 µl) of m-calpain as described in the text (1 U [], 0.5 U [], 0.25 U [], 0.05 U [], none []). After various times of incubation, the activity of DHPR (1 µg equivalent) was measured for 1 min in the system described in the text against q-MH₂ (**a**) and q-BH₂ (**b**).

sequence of sheep DHPR was previously analyzed except for about 100 amino acids in its N-terminal region (Lockyer *et al.*, 1987). The N-terminal sequences of fragments A and B analyzed in this study were determined as Ser-IIe-Asp-Val-Gln-Glu-Asn-Glu and Val-Val-Val-Lys-Met-Thr-Asp-Ala-Phe-Leu, respectively (Fig. 3a). These sequences correspond to the N-terminal regions around about 30 - 60th of DHPR subunits of rat (Shahbaz *et al.*, 1987), mouse (Yang *et al.*, 1996), and human (Dahl *et al.*, 1987) enzymes, as shown in Fig. 3a. These results indicate that the DHPR subunit was cleavaged by m-calpain between Ala-34 and Ser-35 and between Ser-47 and Val-48 to form fragments A and B, respectively (Fig. 3b).

Activities of fragments A and B

Fig. 4a shows time courses of the activities of DHPR against q-MH₂ during the incubation with various amounts of m-calpain. When DHPR was incubated for 5 min with 1 U mcalpain, the activity increased to about 150% of that of the native DHPR. The activity gradually decreased to about the 120% level, and this level remained until 180 min after the start of incubation. When a much smaller amount of m-calpain (0.05 U) was added, DHPR activity increased to about the 120% level, and it remained there until 180 min. These data (Fig. 4a) and the results by SDS-PAGE (Fig. 2) show that fragment A was more easily decomposed than fragment B and had higher activity than native DHPR or fragment B and that fragment B had higher activity than native DHPR. If the natural substrate (q-BH₂) of DHPR was used by the addition of BH₄, in the assay system of DHPR, a more rapid increase and decrease in DHPR activity was observed than with the chemically synthesized substrate (q-MH₂) during the initial time of incubation of DHPR and m-calpain, as shown in Fig. 4b.

Modulation of catalytic properties of DHPR by limited proteolysis

Fig. 4b also shows that native DHPR has higher activity toward q-BH₂ than toward q-MH₂. To know the differences in the catalytic properties between the native and modified DHPR, we analyzed the activities in the reaction mixture with m-calpain after the incubations for 5 min as the mixture of fragments A and B, and native DHPR, and for 120 min as fragment B. Fragments A and B and native enzyme could not be separated from each other by HPLC in suficient amount for the experiment because of further proteolysis during application of the mixture onto the column and because of similarity in molecular sizes of these fragments. As shown in Table 2, the sample incubated for 5 min had smaller Kms for both substrates than that containing fragment B (120 min-incubation), and fragment B had smaller ones than the native DHPR (0 min-incubation). All fragments show lower values in Km and larger values in Vmax/Km when the natural substrate q-BH₂ was used than when the chemically synthesized q-MH₂ was the substrate. Especially, the Vmax/Km value for q-BH₂ of the 5-min sample was about 2 fold that of the native DHPR (Table 2). These results indicate that fragment A has higher affinity and utilization for the natural substrate q-BH₂ than fragment B or native DHPR.

Table 2. Effect of m-calpain treatment on the kinetics of DHPR

Incubation time	Km (μM)		Vrel/Km	
	q-MH ₂	q-BH ₂	q-MH ₂	q-BH ₂
0	3.2	2.5	0.15 [100]	0.19 [100]
5 min	2.7	1.1	0.18 [120]	0.35 [184]
120 min	3.0	1.7	0.16 [107]	0.25 [132]

DHPR was preincubated with m-calpain and calcium ion for 5 or 120 min and determined in kinetic parameters as described in Fig. 5. DHPR equivalent to 1 μ g of the preincubation mixture was assayed for activity against various amounts of q-MH₂ and q-BH₂ as described in Fig. 5. Vrel; relative Vmax.

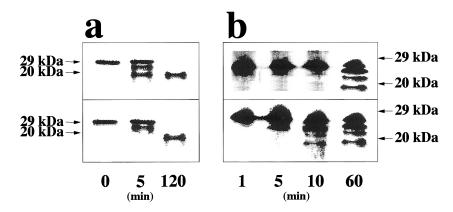


Fig. 5. Effect of phosphorylation of DHPR on the sensitivity to m-calpain. SDS-PAGE of phosphorylated and non-phosphorylated enzymes after being incubated with m-calpain. DHPR or SPR was phosphorylated by Ca²⁺/calmodulin-dependent protein kinase II as described in the text. The reaction mixture was further incubated with 1 U of m-calpain for various times, and an appropriate volume containing 10 μg DHPR and 30 μg SPR was applied to SDS-PAGE. (a) DHPR phosphorylated (lower) and non-phosphorylated (upper). (b) SPR phosphorylated (lower) and non-phosphorylated (upper).

Sensitivity of DHPR to m-calpain

DHPR and SPR can be phosphorylated by Ca²⁺/ calmodulin-dependent protein kinase II, as described in our previous report, although these enzymes are hardly affected in terms of their catalytic properties (Katoh et al., 1994). During the incubation of DHPR with m-calpain, besides the native DHPR, fragments A and B were observed after 5 min; whereas only fragment A was seen at that time when DHPR was incubated after having been phosphorylated with Ca2+/ calmodulin-dependent protein kinase II according to the method described previously (Katoh et al., 1994) (Fig. 5a [lower panel]). This indicates that DHPR became resistant to Ca2+activated protease by Ca2+-dependent phosphorylation. SPR was also found to be cleaved by m-calpain (Fic. 5b [upper panel]), although its activity decreased with incubation time, as shown in Fig. 1b. Three visible fragments of 26 k, 24 k, and 19 k were formed when the native SPR (28 k) was incubated with m-calpain for 60 min (Fig. 5b [upper]). If SPR was incubated with m-calpain after the former had been phosphorylated by Ca2+/calmodulin-dependent protein kinase II, these three fragments were observed even after 10 min (Fig. 5b [lower]). Accordingly, in the case of SPR, sensitivity to mcalpain was accelerated by Ca²⁺-dependent phosphorylation. Sensitivity of DHPR or SPR to V8 protease, which does not require calcium ions for its activity, however, was not affected by phosphorylation with Ca2+/calmodulin-dependent protein kinase II (data not shown).

DISCUSSION

In this study, we found a transient activation of DHPR by m-calpain, a widely distributed Ca²⁺-activated protease, *in vitro*. The activation of DHPR was observed during the initial period of the incubation of DHPR with m-calpain in the presence of calcium ions (Fig. 1a). By SDS-PAGE of the reaction mixture, two new fragments of 21 k and 19 k appeared from the native DHPR subunit of 29 k during the incubation (Fig. 2). These fragments on the gel were isolated, and their N-terminal amino

acid sequences were analyzed by the Edman method. The result indicated that the DHPR subunit was cleaved between Ala-34 and Ser-35 and between Ser-47 and Val-48 by m-calpain, and converted to 21 k and 19 k fragments, respectively (Fig. 3a). The positions of the hydrolysis by m-calpain in DHPR subunit were indicated on the three-dimensional structure obtained from X-ray data cited from the Data Bank (Su *et al.*, 1993) (Fig. 3b)

The two fragments were more active than the native DHPR subunit. Fragment A was the most active type, since the activity in the 5-min incubation sample, in which fragments A and B and native DHPR were present (Fig. 2), was higher than that in the sample taken at 0 time (native DHPR) or 120 min (fragment B) (Fig. 2, Table 2). Vrel/Km values of these fragments were larger than those of native DHPR (Table 2). It is physiologically significant that the rate of activation of these fragments was more remarkable when the natural cofactor, BH₄ was used as the source of the substrate than when the synthesized cofactor, 6M-PH₄ was used (Table 2). Like SPR (Katoh and Sueoka, 1984; Sueoka and Katoh, 1985), DHPR was demonstrated to be a member of the short-chain dehydrogenase/reductase superfamily, based on its NAD(P)H requirement and primary structure (Jornvall et al., 1995). The enzymes of this family generally have a coenzyme-binding domain in their N-terminal region and a unique segment in their C-terminal region, for individual function, containing conserved residues of the essential motif Tyr-X-X-X-Lys. As well as SPR (Tyr¹⁷¹-X-X-Lys¹⁷⁵), DHPR contains a specific Tyr¹⁴⁶-X-X-X-Lys¹⁵⁰ motif in its sequence, and the pteridine substrate and nicotinamide ribose moiety of NADH were located near this sequence by crystallographic analysis (Su et al., 1993). These data and the results in this study indicate that cleavages of DHPR in the N-terminal region (34-35 and 47-48 in the sequence) by m-calpain in the presence of calcium ions modified the conformational structure around the active site of the enzyme and may spread out the domain around the pteridine site to increase the affinity strictly for the natural pteridine substrate (Fig 3b and Fig 4b). Small fragments of the N-terminus cut off by m-calpain, however, might still have been connected to fragment A or B in the presence of calcium ions since the activation was not observed when EGTA was added after the proteolytic reaction (Table 1).

m-Calpain has a strict substrate specificity, and bring about functional modulation of various proteins (Suzuki, 1993). DHPR and SPR were found to be good substrates for mcalpain in this study, and limitedly proteolyzed by the protease. DHPR showed a transient increase in its activity due to the proteolysis, whereas, SPR, in contrast, displayed a decreased activity, as was shown in Fig. 1a, b. The SPR subunit (28 k) was cleaved to at least three fragments of 26 k, 24 k, and 19 k during the initial incubation with m-calpain, as indicated in Fig. 5b; however, only one site of Val187 was detected by analysis of the N-terminal amino acid (data not shown). This result suggests that SPR was cleaved by m-calpain at its Cterminal region, which is located near the catalytic site of Tyr¹⁷¹-X-X-X-Lys¹⁷⁵, the motif contributing to the enzyme activity (Fujimoto et al., 1999), and then lost its activity. The sites of hydrolysis of DHPR and SPR by m-calpain found in this study however, were located in the sequences indicating negative scores for PEST sequences for the calpain reaction, as obtained by PESTfind Analysis (Rogers et al., 1986; Rechsteiner and Rogers, 1996). The finding shown in Fig. 5a, b, in which the cleavage of DHPR by Ca2+-activated protease is controlled by phosphorylation by Ca²⁺-dependent protein kinase, seems to be significant in vivo. Tyrosine hydroxylase, a BH₄-requiring enzyme for formation of catecholamines, is also limitedly proteolyzed and increased in its activity by m-calpain (Kiuchi et al., 1991). As well as SPR and DHPR (Katoh et al., 1994), tyrosine hydroxylase (Campbell et al., 1986) and nitric oxide synthase (Bredt et al., 1992) are also phosphorylated by Ca²⁺/ calmodulin-dependent protein kinase II and/or protei kinase C. It is physiologically understandable that enzymes involved in the biosynthesis and recycling of BH₄, and those requiring it might all be controlled by calcium ions through phosphorylation and proteolysis to regulate the amounts of neurotransmitters such as catecholamines, indoleamines, and nitric oxide in the cell.

ACKNOWLEDGMENTS

The authors wish to thank Dr. K. Haino (Jochi University, Japan) for her helpful advice concerning the Edman analysis. This work was supported in part by a grant from the Ministry of Education, Science, Sports and Culture of Japan (No. 100771018) and by a grant from the Miyata Foundation, Meikai University.

REFERENCES

- Bredt DS, Ferris CD, Snyder SH (1992) Nitric oxide synthase regulatory sites. Phosphorylation by cyclic AMP-dependent protein kinase, protein kinase C, and calcium/calmodulin protein kinase; identification of flavin and calmodulin binding sites. J Biol Chem 267: 10976–10981
- Campbell DG, Hardie DG, Vulliet PR (1986) Identification of four phosphorylation sites in the N-terminal region of tyrosine hydroxylase. J Biol Chem 261: 10489–10492

- Craine JE, Hall ES, Kaufman S (1972) The isolation and characterization of dihydropteridine reductase from sheep liver. J Biol Chem 247: 6082–6091
- Dahl HH, Hutchison W, McAdam W, Wake S, Morgan FJ, Cotton RG (1987) Human dihydropteridine reductase: characterisation of a cDNA clone and its use in analysis of patients with dihydropteridine reductase deficiency. Nucleic Acids Res 15: 1921–1932
- Dayton WR, Goll DE, Zeece MG, Robson RM, Reville WJ (1976) A Ca²⁺-activated protease possibly involved in myofibrillar protein turnover. Purification from porcine muscle. Biochemistry 15: 2150–2158
- Drummond GI, Duncan L (1966) The action of calcium ion on cardiac phosphorylase b kinase. J Biol Chem 241: 3097–3103
- Edman P, Begg G (1967) A protein sequenator. Eur J Biochem 1: 80-91
- Firgaira FA, Cotton RG, Danks DM (1981) Isolation and characterization of dihydropteridine reductase from human liver. Biochem J 197: 31–43
- Fujimoto K, Ichinose H, Nagatsu T, Nonaka T, Mitsui Y, Katoh S (1999) Functionally important residues tyrosine-171 and serine-158 in sepiapterin reductase. Biochim Biophys Acta 1431: 306–314
- Greengard P (1978) Phosphorylated proteins as physiological effectors. Science 199: 146–52
- Hasegawa H (1977) Dihydropteridine reductase from bovine liver. Purification, crystallization, and isolation of a binary complex with NADH. J Biochem 81: 167–177
- Hatakeyama K, Harada T, Suzuki S, Watanabe Y, Kagamiyama H (1989) Purification and characterization of rat liver GTP cyclohydrolase I. Cooperative binding of GTP to the enzyme. J Biol Chem 264: 21660–21664
- Huston RB, Krebs EG (1968) Activation of skeletal muscle phosphorylase kinase by Ca²⁺. II. Identification of the kinase activating factor as a proteolytic enzyme. Biochemistry 7: 2116–2122
- Inoue Y, Kawasaki Y, Harada T, Hatakeyama K, Kagamiyama H (1991) Purification and cDNA cloning of rat 6-pyruvoyltetrahydropterin synthase. J Biol Chem 266: 20791–20796
- Jornvall H, Persson B, Krook M, Atrian S, Gonzalez-Duarte R, Jeffery J, Ghosh D (1995) Short-chain dehydrogenases/reductases (SDR). Biochemistry 34: 6003–6013
- Katoh S, Akino M (1986) Biosynthesis of tetrahydrobiopterin in animals. Zool Sci 3: 745–757
- Katoh S, Nagai M, Nagai Y, Fukushima T, Akino M (1970) Some new biochemical aspects of sepiapterin and sepiapterin reductase. in "Chemistry and Biology of Pteridines" Ed by Iwaki K, Akino M, Goto M, Iwanami Y, International Printing, Tokyo, pp. 225–234
- Katoh S, Sueoka T, (1984) Sepiapterin reductase exhibits a NADPHdependent dicarbonyl reductase activity. Biochem Biophys Res Commun 118: 859–866
- Katoh S, Sueoka T, Yamamoto Y, Takahashi S (1994) Phosphorylation by Ca²⁺/calmodulin-dependent protein kinase II and protein kinase C of sepiapterin reductase, the terminal enzyme in the biosynthetic pathway of tetrahydrobiopterin. FEBS Lett 341: 227– -232
- Kaufman S, Fisher DB (1974) Pterin-requiring aromatic amino acid hydroxylases in "Molecular mechanisms of oxygen activation" Ed by Hayaishi O, Academic Press, New York, pp 285–369
- Kaufman S, Levenberg B (1959) Further studies on the phenylalanine-hydroxylation cofactor. J Biol Chem 234: 2677–2682
- Kiuchi K, Kiuch K, Titani K, Fujita K, Suzuki K, Nagatsu T (1991) Limited proteolysis of tyrosine hydroxylase by Ca²⁺-activated neutral protease (calpain). Biochemistry 30: 10416–10419
- Korri KK, Chippel D, Chauvin MM, Tirpak A, Scrimgeour KG (1977) Quinonoid dihydropterin reductase from beef liver. Can J Biochem 55: 1145–1152
- Laemmli UK (1970) Cleavage of structural proteins during the assembly of the head of bacteriophage T4. Nature 227: 680-

685

- Lockyer J, Cook RG, Milstien S, Kaufman S, Woo SL, Ledley FD (1987) Structure and expression of human dihydropteridine reductase. Proc Natl Acad Sci USA 84: 3329–3333
- Nagatsu T, Mizutani K, Nagatsu I, Matsuura S, Sugimoto T (1972) Pteridines as cofactor or inhibitor of tyrosine hydroxylase. Biochem Pharmacol 21: 1945–1953
- Nishiura I, Tanaka K, Yamato S, Murachi T (1978) The occurrence of an inhibitor of Ca²⁺-dependent neutral protease in rat liver. J Biochem 84: 1657–1659
- Pant HC, Terakawa S, Gainer H (1979) A calcium activated protease in squid axoplasm. J Neurochem 32: 99–102
- Phillips DR, Jakabova M (1997) Ca²⁺-dependent protease in human platelets.Specific cleavage of platelet polypeptides in the presence of added Ca²⁺. J Biol Chem 252: 5602–5605
- Rechsteiner M, Rogers SW (1996) PEST sequences and regulation by proteolysis. TIBS 21: 267–271
- Rogers S, Wells R, Rechsteiner M (1986) Amino acid sequences common to rapidly degraded proteins: The PEST hypothesis. Science 234: 364–368
- Schlaepfer WW (1974) Calcium-induced degeneration of axoplasm in isolated segments of rat peripheral nerve. Brain Res 69: 203– 215
- Shahbaz M, Hoch JA, Trach KA, Hural JA, Webber S, Whiteley JM (1987) Structural studies and isolation of cDNA clones providing the complete sequence of rat liver dihydropteridine reductase. J Biol Chem 262: 16412–16416

- Su Y, Varughese KI, Xuong NH, Bray TL, Roche DJ, Whiteley JM (1993) The crystallographic structure of a human dihydropteridine reductase NADH binary complex expressed in *Escherichia coli* by a cDNA constructed from its rat homologue. J Biol Chem 268: 26836–26841
- Sueoka T, Katoh S (1982) Purification and characterization of sepiapterin reductase from rat erythrocytes. Biochim Biophys Acta 717: 265–271
- Sueoka T, Katoh S (1985) Carbonyl reductase activity of sepiapterin reductase from rat erythrocytes. Biochim Biophys Acta 843: 193– 198
- Suzuki K (1993) In search of physiological function of calpain activation and tissue specific novel calpain. Seikagaku 65: 537–552
- Waxman L, Krebs EG (1978) Identification of two protease inhibitors from bovine cardiac muscle. J Biol Chem 253: 5888–5891
- Webber S, Deits TL, Snyder WR, Whiteley JM (1978) The purification of rat and sheep liver dihydropteridine reductase by affinity chromatography on methotrexate-sepharose. Anal Biochem 84: 491– 503
- Yamauchi T, Fujisawa H (1983) Purification and characterization of the brain calmodulin-dependent protein kinase (kinase II), which is involved in the activation of tryptophan 5-monooxygenase. Eur J Biochem 132: 15–21
- Yang N, Hanssen K, Armarego WRF (1996) The sequence of mouse dihydropteridine reductase cDNA, and comparison with human and rat sequence. Pteridines 7: 14–23

(Received September 20, 1999 / Accepted November 10, 1999)