

Figure 6 Detection of 3-NT proteins in mitochondria complex I-IV subunits. The cells were treated with 250 μ M SIN-1 for 1 hr, and P2 fraction was isolated. The samples were subjected to SDS-PAGE and detected by the immunoblot assay using the antibody against 3-NT, or complex I (I), complex II (II), complex III (III), and complex IV (IV), respectively. Molecular markers are shown in the right column. Arrows show the protein bands positively stained with anti-3-NT and anticomplex I antibody.

the subunits of complex I, as assigned based on the staining with anticomplex I antibody (Fig. 6). These results clearly show that the preferential nitration of complex I subunits may contribute to mitochondrial dysfunction observed in the nigro-striatum of parkinsonian brain (30,31).

Treatment of ONOO⁻-generating *N*-morpholino sydnonimine (SIN-1) induced apoptosis in the SH-SY5Y cells (32-35).

A step-wise activation of apoptotic cascade was observed: decline in $\Delta\Psi_m$, activation of caspase 3, and phosphorylation of p38 mitogen-activated phosphokinase (MAP) (33). Nitric oxide and ONOO^- were reported to induce apoptosis by nitration of tyrosine residues to release cytochrome *c* (36), or of cytochrome *c* itself (37). In addition, mitochondrial ATP synthesis was inhibited markedly by ONOO^- , as shown in Fig. 7, which may be due to its reversible binding to cytochrome oxidase or inactivation of complex I and II, and ATPase (38–41).

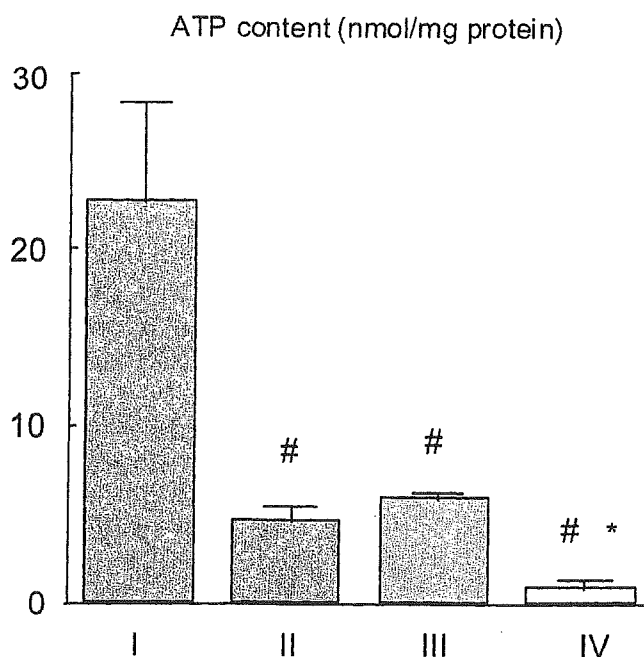


Figure 7 Effect of SIN-1 and PSI on the ATP levels in SH-SY5Y cells. The cells were incubated without or with 250 μM SIN-1 in the absence and presence of 10 μM PSI for 2 hr. The ATP content was measured by the luminofluorometric method. The column and bar represent the mean and SD of triplicate measurements of three independent experiments. I, control cells; II, cells treated with PSI alone; III, SIN-1 alone; and IV, SIN-1 and PSI. # Difference from control (I) is statistically significant ($p < 0.05$). * Difference from SIN-1 alone (III) is statistically significant ($p < 0.05$).

As shown in Fig. 5A, SIN-1 treatment increased 3-NT protein only in distinct protein bands, suggesting the preference of some proteins to ONOO⁻ modification. In addition, increase in the amount 3-NT protein was not so significant.

III. PROTEASOME: ITS ROLE IN ACCUMULATION OF OXIDIZED PROTEINS

Insoluble intracellular protein aggregates, such as Lewy body in PD and senile plaques composed of β -amyloid in AD, are hallmarks of neurodegeneration. Although it remains unclear whether protein aggregates cause directly neuronal cell death or are the results of deteriorated cellular homeostasis in dying neurons, the analysis of the constituents of inclusion bodies may suggest the molecular mechanism leading their formation. Protein aggregation is a manifestation of disturbed cellular protein-folding homeostasis maintained by the ubiquitin-proteasome system. In LB, ubiquitin, and proteasome, subunits are major components (42,43), in addition to α -synuclein, Parkin, and ubiquitin C-terminal hydrolase-L1 (UCH-1) (44-47). They were modified with ROS-RNS as nitrated synuclein (48) and dityrosine (49).

To clarify the interactions among oxidative stress, dysfunction of the proteasome system, and formation of the inclusion body, the effects of a proteasome inhibitor, carbobenzoxy-L-isoleucyl- γ -*t*-butyl-L-alanyl-L-leucinal (PSI) were examined on the deposition of modified proteins and the cell vulnerability (34,50). PSI increased the amount of 3-NT proteins in the SH-SY5Y cells, but the number of 3-NT protein bands was almost the same as in the control (Fig. 5B). The functional deterioration of mitochondria was also enhanced by PSI, as shown by the severe reduction of ATP synthesis (Fig. 7). In addition, the number of apoptotic cells increased significantly by PSI, whereas that of necrotic cells remained almost the same (Fig. 8A). At the same time, the acrolein-modified protein increased significantly in subcellular fractions of the SH-SY5Y cells after being treated with PSI (Fig. 8B). These results clearly demonstrate that the inhibition of proteasome activity

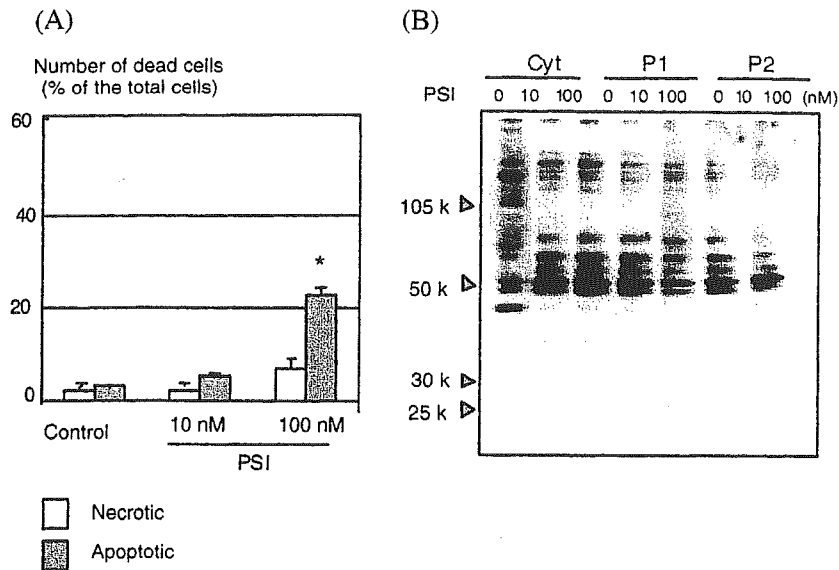


Figure 8 Effect of PSI on the cell death and the levels of acrolein-modified protein. (A) Apoptotic and necrotic cell death after treatment with 10 and 100 nM PSI was quantified by staining the cells with 10 nM Hoechst 33342. The column and the bar represent the mean and SD of the number of dead cells, expressed as the percentage of the total. The open and the filled columns represent necrotic and apoptotic dead cells, respectively. The difference from the control cells is significant ($p < 0.05$) by ANOVA. (B) The effects of PSI on the level of acrolein-modified proteins were examined by immunoblot with antiacrolein antibody. The cytoplasmic (Cyt) P1 and P2 fractions of the cells treated with 10 and 100 nM of PSI for 24 hr were prepared by centrifugation at 800 g and 13,000 g , respectively.

may play a key role in the accumulation of oxidation-modified protein and neuronal death.

IV. MITOCHONDRIAL DYSFUNCTION INHIBITS THE PROTEASOME SYSTEM BY OXIDATIVE MODIFICATION

Systemic reduction in complex I of the mitochondrial electron transfer chains was reported in the nigro-striatum of patients

with PD (30,31,51). Recently, an inhibitor of complex I, rotenone, was reported to induce parkinsonism in rodents, and fibrillar cytoplasmic inclusions containing ubiquitin and α -synuclein were detected after the continuous administration (52). The effects of mitochondrial dysfunction on the proteasome system were studied by the use of rotenone at a concentration to induce apoptosis in the SH-SY5Y cells (50). ATP levels reduced markedly, and apoptosis was induced in the cells after 4–5 days treatment with rotenone. The oxidative modification of proteins was followed by the use of antibody against the acrolein-modified protein (19). The levels of the acrolein-modified protein increased markedly by rotenone treatment (Fig. 9).

The ubiquitin–proteasome system is a major site to remove damaged or mutated proteins and also regulatory proteins controlling cell cycle and signal transduction. In the nigro-striatum of the parkinsonian brain, the decreased activity of the proteasome was reported, suggesting its involvement in the pathological features (53). The oxidized protein is preferentially degraded *in vitro* by 20S proteasome in an ATP-independent way. 20S proteasome is composed of four rings to make a cylindrical structure, and each ring is made of seven α and β 20S subunits (54). Noncatalytic α -heptameric rings form each of the two outer rings and catalytic β -heptamers form the two inner rings. Binding of a regulatory subunit named 19S complex (ATPase, PA700) to both ends of the 20S cylinder produces 26S proteasome with a higher catalytic activity than the 20S proteasome. The 26S proteasome can degrade polyubiquitinated proteins and ornithine decarboxylase in an ATP-dependent process.

In the SH-SY5Y cells, rotenone treatment increased ROS–RNS levels only transitionally and slightly, whereas the oxidized protein levels increased progressively, suggesting that degradation of modified proteins may be mainly involved in the deposition. The enzymatic activity of proteasome measured with an artificial fluorescent substrate, carbobenzoxy-L-leucyl-L-leucyl-L-glutamic acid α -(4-methyl-coumaryl-7-amide) (Z-Leu-Leu-Glu-MCA), reduced in a time- and dose-dependent way, and virtually was not detected after 4 days treatment with

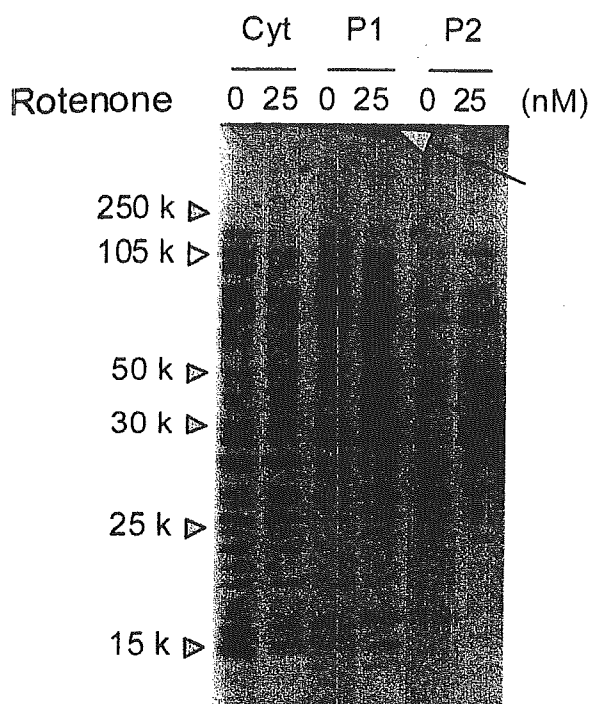


Figure 9 Effect of rotenone on oxidized protein level in SH-SY5Y cells. The cells were treated without or with 25 nM of rotenone for 5 days. The cells were fractionated into the cytoplasmic (Cyt), P1 and P2 fractions, and each fraction was subjected to immunoblotting, using antibody against acrolein-modified proteins. The aggregated proteins were observed in the stacking gel of the P1 fraction (arrow).

rotenone, as shown in Table 2. The reduction in the enzymatic activity was neither due to ATP depletion nor to the reduction of the protein level, indicating that the homospecific activity of proteasome decreased by the treatment. Indeed, the 20S β subunit protein was modified by acrolein, as shown by immunoprecipitation with anti-20S β subunit and by western blot analysis with the antiacrolein antibody. In this immunoprecipitant coprecipitated with 20S β subunit, oxidized proteins immunoreactive to antiacrolein antibody were detected in the cells treated with rotenone, suggesting that the acrolein-conjugated protein

Table 2 Effects of Rotenone Treatment on the Enzymatic Activity of Proteasome in SH-SY5Y Cells

Rotenone (25 nM)	ATP (2 mM)	Proteasome activity in SH-SY5Y cells treated with rotenone (pmol/min/ μ g protein)	
		After 6 hr	After 96 hr
Control	-	0.89 \pm 0.11*	5.19 \pm 0.47
	+	1.76 \pm 0.02	8.21 \pm 0.11
Sample	-	0.97 \pm 0.02	0.96 \pm 0.04
	+	2.25 \pm 0.02	1.80 \pm 0.13

Proteasome activity in the cytoplasmic fraction of the cells treated with 25 nM rotenone was measured fluorometrically in the absence or presence of exogenously added ATP (2 mM).

*Mean \pm SD of four experiments.

may be a substrate of 20S proteasome. However, further studies are required to clarify whether the acrolein-modified protein can directly inhibit proteasome activity, as in the case of the 4-HNE proteins (55). The increase in acrolein adducts after rotenone treatment may be comparable with that observed in the presence of PSI, as described earlier.

V. CONCLUSION

The results of this study suggest that mitochondrial dysfunction may induce degeneration of dopamine neurons through modification and inactivation of the proteasome system and subsequent aggregation of the oxidized proteins. The precise mechanism behind the cell death induced by oxidative stress requires further investigation, but it may be quite relevant to consider that the inactivation of proteasome may play a critical role in the activation of apoptotic signal. In neurodegenerative disorders, the environmental and genetic factors may induce a malignant cycle between mitochondrial dysfunction, increased oxidative stress, and reduced activity of proteasome in neurons, leading to the typical pathological

characteristics, cell death and formation of inclusion body. Our recent studies on the ubiquitin–proteasome system in the substantia nigra of human brains present further support to our hypothesis. In the parkinsonian brain, the 20S β subunit precipitated with the specified antibody was modified with acrolein, whereas that from control was not modified. Future studies may give a new insight into the mechanism of neuronal cell death in age and age-related disorders, and may give us clues for a new therapeutic strategy to protect neurons from apoptosis.

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Oxidative Stress in Mitochondria

Decision to Survival and Death of Neurons in Neurodegenerative Disorders

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Abstract

In mitochondria, oxidative phosphorylation and enzymatic oxidation of biogenic amines by monoamine oxidase produce reactive oxygen and nitrogen species, which are proposed to cause neuronal cell death in neurodegenerative disorders, including Parkinson's and Alzheimer's disease. In these disorders, mitochondrial dysfunction, increased oxidative stress, and accumulation of oxidation-modified proteins are involved in cell death in definite neurons. The interactions among these factors were studied by use of a peroxynitrite-generating agent, *N*-morpholino sydnonimine (SIN-1) and an inhibitor of complex I, rotenone, in human dopaminergic SH-SY5Y cells. In control cells, peroxynitrite nitrated proteins, especially the subunits of mitochondrial complex I, as 3-nitrotyrosine, suggesting that neurons are exposed to constant oxidative stress even under physiological conditions. SIN-1 and an inhibitor of proteasome, carbobenzoxy-L-isoleucyl- γ -*t*-butyl-L-alanyl-L-leucinal (PSI), increased markedly the levels of nitrated proteins with concomitant induction of apoptosis in the cells. Rotenone induced mitochondrial dysfunction and accumulation and aggregation of proteins modified with acrolein, an aldehyde product of lipid peroxidation in the cells. At the same time, the activity of the 20S β -subunit of proteasome was reduced significantly, which degrades oxidative-modified protein. The mechanism was proved to be the result of the modification of the 20S β -subunit with acrolein and to the binding of other acrolein-modified proteins to the 20S β -subunit.

Increased oxidative stress caused by SIN-1 treatment induced a decline in the mitochondrial membrane potential, $\Delta\Psi_m$, and activated mitochondrial apoptotic signaling and induced cell death in SH-SY5Y cells. As another pathway, p38 mitogen-activated protein (MAP) kinase and extracellular signal-regulated kinase (ERK) mediated apoptosis induced by SIN-1. On the other hand, a series of neuroprotective propargylamine derivatives, including rasagiline [*N*-propargyl-1(*R*)aminoindan] and (-)-deprenyl, intervened in the activation of apoptotic cascade by reactive oxygen species-reactive nitrogen species in mitochondria through stabilization of the membrane

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potential, $\Delta\Psi_m$. In addition, rasagiline induced antiapoptotic Bcl-2 and glial cell line-derived neurotrophic factor (GDNF) in SH-SY5Y cells, which was mediated by the ERK–nuclear factor (NF)- κ B pathway. These results are discussed in relation to the interaction of oxidative stress and mitochondria in the regulation of neuronal death and survival in neurodegenerative diseases.

Index Entries: Oxidative stress; mitochondria complex I; proteasome; 3-nitrotyrosine; acrolein; Parkinson's disease; apoptosis; rasagiline; transcription factors.

Oxidative-Modified Protein as the Marker of Oxidative Stress

Oxidative stress has been proposed as one of the major causes inducing neuronal death in aging and age-associated disorders (1), and mitochondria produce most of reactive oxygen and nitrogen species (ROS and RNS, respectively) in the cells. The superoxide anion radical is generated by oxidative phosphorylation in mitochondria and reacts with nitric oxide (NO) to produce peroxynitrite (ONOO⁻), one of the most potent radicals. On the other hand, oxidation of biogenic amines by monoamine oxidase in the mitochondrial outer membrane generates hydrogen peroxide. Mitochondria are now considered to play a pivotal role in apoptosis (2), which is the common death type of neurons in Parkinson's (PD) and Alzheimer's diseases (AD) (3). The role of mitochondria in the process of apoptotic commitment is recognized, and impairment of energy charge and redox, permeability transition (PT), disruption of membrane potential ($\Delta\Psi_m$), and release of cytochrome-*c* are observed prior to fragmentation of nuclear DNA, a hallmark of apoptotic morphological features.

Neurodegenerative disorders are characterized by a decline of specified neurons in selected brain regions, associated with protein deposits specific for each disease. In PD, dopamine neurons in the substantia nigra degenerate progressively with formation of Lewy bodies (LBs). The pathogenesis of PD remains as an enigma, but PD is considered a consequence of various genetic and environmental interactions. The vulnerability of the neurons might be the result of increased generation of ROS and RNS, reduced antioxidant

capacity, high contents of iron and dopamine, and mitochondrial dysfunction in nigral dopamine neurons. ROS and RNS generated in mitochondria modify bioactive molecules, such as lipids, proteins, DNA, and carbohydrates, either directly or indirectly with peroxidation products of lipid or carbohydrates. 4-Hydroxynonenal (4-HNE) and acrolein, aldehyde products of lipid peroxidation, are cytotoxic, and aldehyde-modified proteins were increased in dopamine neurons of the nigro-striatum in PD (4), neurofibrillary tangles in AD (5), and the spinal cord in amyotrophic lateral sclerosis (ALS) (6). Figure 1 shows the accumulation of 4-HNE-modified protein in the substantia nigra of parkinsonian brain. Dopamine neurons containing neuromelanin were selectively modified with 4-HNE more markedly in the PD brain than those in the control or in cells other than dopamine neurons. This suggests the essential role of dopamine in increased oxidative stress.

One of the most active RNS, ONOO⁻, nitrates sulfhydryl and hydroxyl residues in cysteine, methionine, phenylalanine, and tyrosine, and the modification inactivates the membrane function and key enzymes (7). 3-Nitrotyrosine (3-NT) is synthesized by nitration of tyrosine residues in protein and a marker for the oxidative stress induced by ONOO⁻ in vivo (8). An increase in 3-NT-containing protein (3-NT protein) was observed in atherosclerosis (9), ALS (6), AD (10), and PD (11). Another oxidation product of tyrosine is dityrosine, which is produced from free and protein-bound tyrosine in the presence of hydrogen peroxide and myeloperoxidase and was detected in atherosclerotic plaques and lipofuscin pigments in the aged human brain (12).

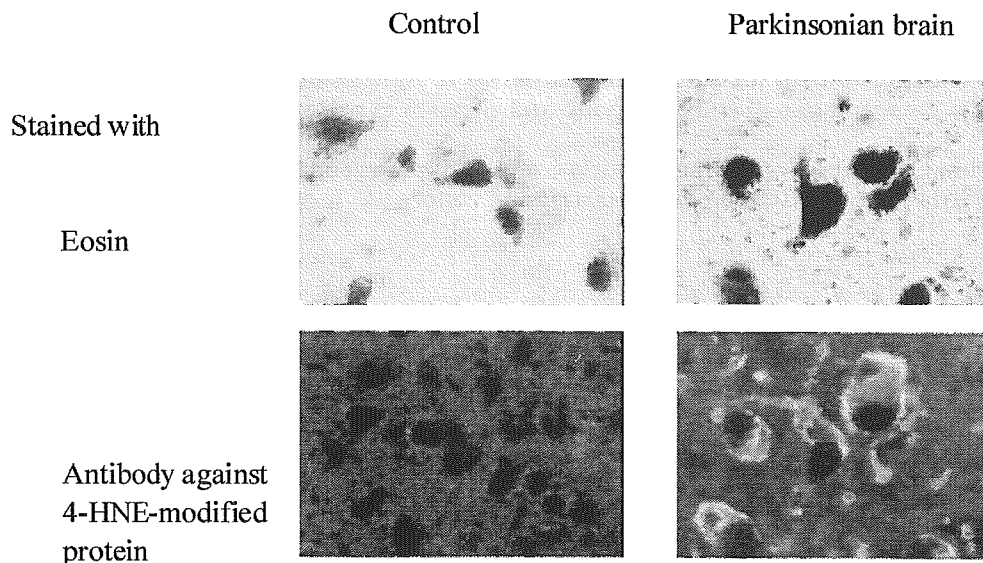


Fig. 1. 4-HNE-modified protein in the substantia nigra of the parkinsonian brain. In dopamine neurons containing neuromelanin, proteins stained with the anti-4-HNE antibody increased in the parkinsonian brain, but were not detected in the age-matched control.

Oxidative modification produces aggregated and crosslinked proteins, which are resistant to proteolytic degeneration and difficult to removed from cells. Accumulation of modified proteins might impact on a variety of cellular functions by changing the enzymatic, regulatory, and transporting potencies of specific proteins, in addition to occupying space in the limited cellular volume. The level of oxidized proteins might reflect the balance between the generation of ROS-RNS and degradation of modified protein, in which the ubiquitin-proteasome system plays a key role (13), as discussed in the following sections.

Mitochondrial Complex I Subunits Are Major Targets of ONOO⁻

In the brain, NO has been thought to be produced in microglia and astrocytes and transported to neurons, where it reacts with superoxide, yielding ONOO⁻. However, NO is synthesized also *in situ* in the neurons and functions as a neuromodulator (14). Using an

antibody against 3-NT protein (15), nitrated proteins were detected in control brains. In brains from AD and PD patients, the same proteins stained with anti-3NT antibody as in control increased markedly, indicating that ONOO⁻ preferentially modified selective proteins in neuronal cells under physiological and pathological conditions. Figure 2 shows the presence of 3-NT protein in control SH-SY5Y cells almost at the same levels as in the cells treated with ONOO⁻-generating *N*-morpholino sydnomine (SIN-1), indicating that the cells are exposed to constant oxidative stress. However, the control cells are intact in growth and proliferation, and under physiological conditions, an active mechanism functions to eliminate modified protein from the cells and to protect cellular function in neuronal cells.

As shown in Fig. 3, the presence of nitrated protein in control cells was confirmed further by the Western blot analysis of the subcellular fractions of SH-SY5Y cells. In mitochondrial fraction, 3-NT protein was detected in the subunits of complex I stained with that against mitochondria complexes I, II, III, and IV,

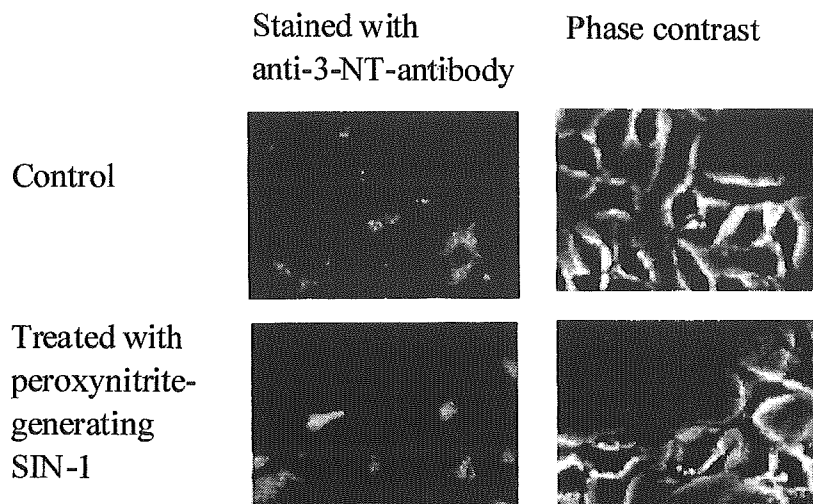


Fig. 2. 3-NT protein detected in control SH-SY5Y cells and cells treated with ONOO⁻-generating SIN-1. In cytoplasm, 3-NT proteins were detected by staining with anti-3-NT antibody (15). Note that the 3-NT protein was detected also in the control.

respectively (16), (Fig. 3). These results suggest that complex I subunits are nitrated preferentially and the modification might contribute to mitochondrial dysfunction observed in the nigro-striatum of the parkinsonian brain (17).

ONOO⁻-generating SIN-1 induced apoptosis in SH-SY5Y cells (14,18–20) and inhibited ATP synthesis in mitochondria. The inhibition might be the result of the binding of NO and ONOO⁻ to cytochrome oxidase or inactivation of complexes II and III and ATPase (21). However, SIN-1 treatment did not increase 3-NT protein as markedly. These seemingly contradicting results indicate again that the levels of oxidative stress and oxidative-modified protein are regulated by the degradation rate rather than the production of ROS-RNS.

Proteasome Plays a Key Role in Accumulation of Oxidized Proteins

Insoluble protein aggregates, such as LBs in PD and senile plaques composed of β -amyloid in AD, are hallmarks of neurodegeneration. However, it remains to be clarified whether

Stained with antibody against

3-NT Mitochondrial complex

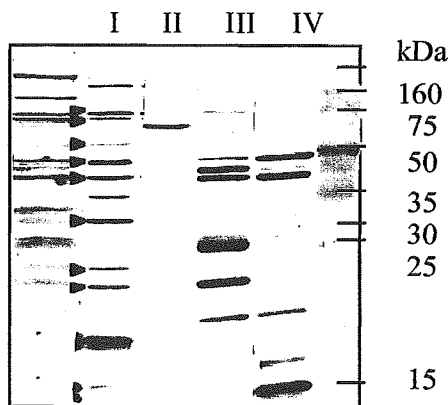


Fig. 3. 3-NT proteins in the subunits of mitochondrial complex I. Mitochondria were prepared from SH-SY5Y cells and subjected to Western blot analysis using antibody against 3-NT protein or mitochondrial complex I, II, III, and IV, respectively.

protein aggregates cause neuronal cell death directly or are the results of deteriorated cellular homeostasis in dying neurons. Protein aggregation is considered a manifestation of

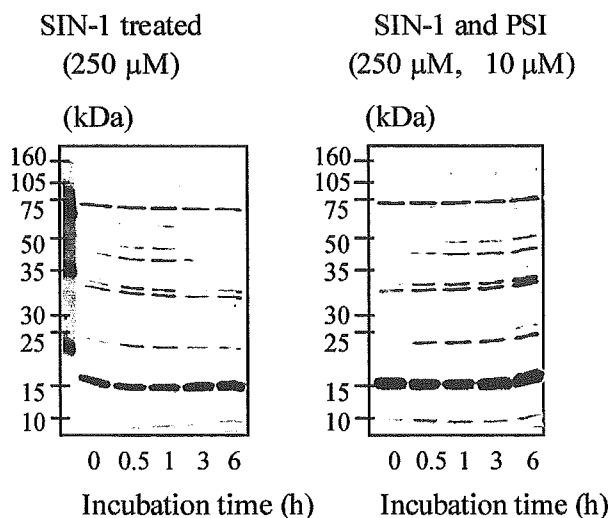


Fig. 4. Effects of ONOO⁻-generating SIN-1 and a proteasome inhibitor, PSI, on 3-NT levels in SH-SY5Y cells. After incubated for 0.5, 1, 3, and 6 h, the cells were subjected to the immunoblotting analysis with anti-3-NT antibody. PSI increased 3-NT in the same proteins as in control and SIN-1-alone-treated cells.

disturbed cellular protein-folding homeostasis maintained by the ubiquitin-proteasome system. Ubiquitinated proteins and proteasome subunits (22,23), in addition to α -synuclein, Parkin and ubiquitin C-terminal hydrolase-L1 (UCH-1) (24–27), are the components of LBs and are sometimes modified with ROS-RNS.

To clarify the interactions among oxidative stress, dysfunction of the proteasome system, and formation of the inclusion body, the effects of a proteasome inhibitor, carbobenzoxy-L-isoleucyl- γ -*t*-butyl-L-alanyl-L-leucinal (PSI) were examined on the aggregation of oxidative-modified proteins and the cell vulnerability (20,28). PSI increased the amount of 3-NT proteins in SH-SY5Y cells, but the number of 3-NT proteins was almost the same as in the control (Fig. 4). At the same time, the number of apoptotic cells increased significantly by PSI, but that of necrotic cells was not (Fig. 5). These results indicate that inhibition of proteasome activity might play a key role in the accumulation of oxidative-modified protein and the induction of cell death.

Rotenone, a Complex I Inhibitor, Inactivates Proteasome by Oxidative Modification

The activity and protein of complex I of the mitochondrial electron transfer chains reduced in the nigro-striatum of patients with PD (17). The systemic administration of rotenone, an inhibitor of complex I, induced parkinsonism in rodents, and fibrillar cytoplasmic inclusions containing ubiquitin and α -synuclein were detected in dopamine neurons (29). The effects of mitochondrial dysfunction on the proteasome system were studied by use of rotenone (28). Apoptosis was induced in the cells after 4–5 d treatment with rotenone. The oxidative modification of proteins was followed by the use of an antibody against acrolein-modified protein (30). As shown in Fig. 6, the levels of acrolein-modified protein increased markedly by the rotenone treatment. In the lysate of rotenone-treated cells, aggregation of acrolein-modified protein with high molecular mass was also detected.

The ubiquitin-proteasome system is a major site for removal of damaged or modified proteins and also regulatory proteins controlling cell cycle and signal transduction. In the nigro-striatum of the parkinsonian brain, the decreased activity of proteasome was reported, suggesting its involvement in the pathological features (31). Oxidized protein is preferentially degraded *in vitro* by 20S proteasome in an ATP-independent way. Binding of the regulatory subunit, 19S complex (ATPase, PA700), to both ends of the 20S cylinder produces 26S proteasome with higher catalytic activity than 20S proteasome. 26S proteasome degrades polyubiquitinated proteins and ornithine decarboxylase in an ATP-dependent process.

In SH-SY5Y cells, the rotenone treatment increased ROS-RNS levels detected with 2,7-dichlorofluorescence diacetate only transitionally and slightly, whereas the oxidized protein levels increased progressively, suggesting that protein might function as “the second scavenger” of ROS-RNS and that degradation of modified proteins might be impaired. The

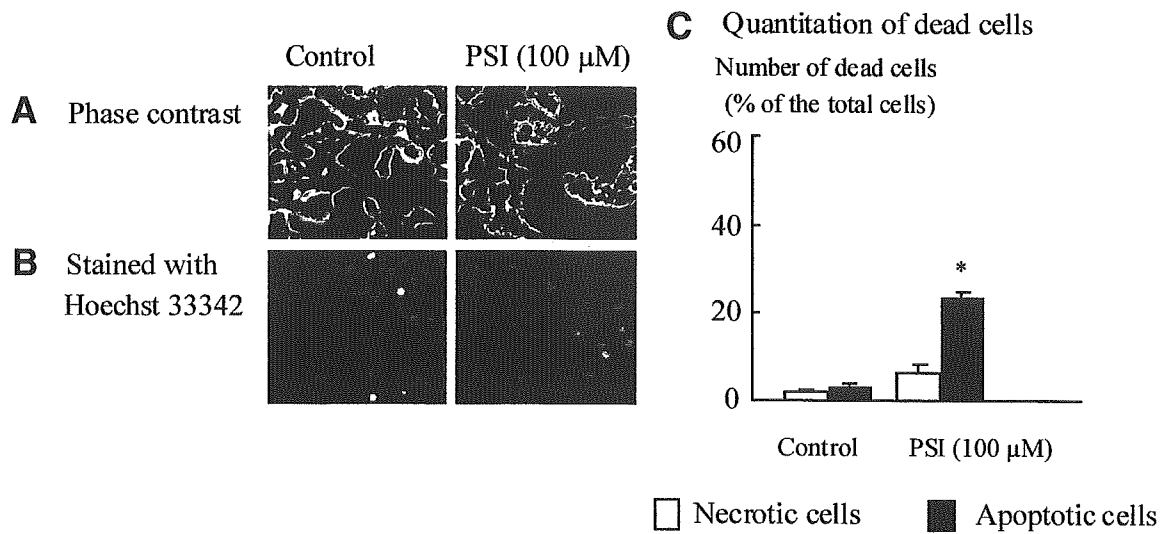


Fig. 5. Cytotoxic effect of PSI treatment on SH-SY5Y cells. The cells were treated with or without PSI for 24 h. (A) Phase-contrast microscopy; (B) nuclear staining with Hoechst 33342, (C) quantitative measurement of apoptotic and necrotic cells. Apoptotic cells were assessed from condensed nuclei stained with Hoechst 33342. The column and the bar represent the mean and SD of the number of dead cells, respectively, expressed as the percentage of the total. The open and the filled columns represent necrotic and apoptotic dead cells, respectively. * $p < 0.05$ from control.

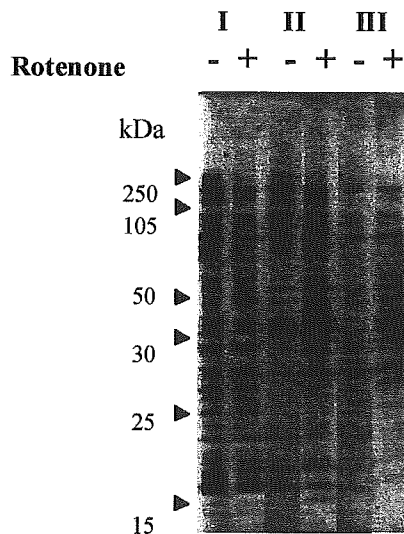


Fig. 6. Acrolein-modified protein in rotenone-treated SH-SY5Y cells. The cells were treated with 5 μ M rotenone (+) or without (-) for 3 d and the sub-cellular fractions were prepared. I: cytoplasm; II: P₁ fraction (nuclei, plasma membrane); III: P₂ fraction (mitochondria). Proteins were immunoblotted with an antibody against acrolein-modified protein. In the P₁ fraction, aggregated acrolein-modified proteins were detected at the top of the gel.

enzymatic activity of proteasome (post-glutamyl peptidase-like activity) measured with a synthetic fluorescent substrate, carbobenzoxy-L-leucyl-L-leucyl-L-glutamic acid α -(4-methylcoumaryl-7-amide) [Z-Leu-Leu-Glu-MCA],

reduced in a time- and dose-dependent way and virtually was not detected after 4 d treatment with rotenone, as shown in Table 1.

The lysate of rotenone-treated cells was immunoprecipitated with antibody against the

Table 1
Effects of Rotenone Treatment on Proteasome Activities in SH-SY5Y Cells

Rotenone (25 nM)	ATP (2 mM)	Proteasome activity in SH-SY5Y cells treated with rotenone (pmole/min/ μ g protein)	
		After 6 h	After 96 h
Control	-	0.89 \pm 0.11 ^a	5.19 \pm 0.47
	+	1.76 \pm 0.02	8.21 \pm 0.11
Sample	-	0.97 \pm 0.02	0.96 \pm 0.04
	+	2.25 \pm 0.02	1.80 \pm 0.13

Cytoplasmic fraction was prepared from the cells treated with 25 nM rotenone. Postglutamyl peptidase-like proteasome activity was measured fluorometrically using Z-Leu-Leu-Glu-MCA as a substrate with or without the addition of ATP (2 mM).

^a Mean \pm SD of four experiments.

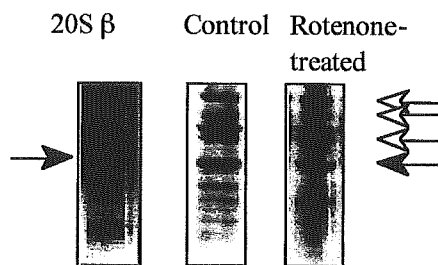


Fig. 7. Modification of the 20S β -subunit of proteasome with acrolein after treatment with rotenone. The cells were treated with 25 μ M rotenone (rotene-treated) or without (control) for 4 d, lysed, and precipitated with antibody against the 20S β -subunit. The immunoprecipitant was subjected to Western blot analysis with antibody against acrolein-modified protein. The 20S β -subunit in the precipitant was visualized with anti-20S β -antibody (20S β). A filled arrow shows that 20S β was modified with acrolein.

20S β -subunit and the subunit protein was modified by acrolein, as shown by Western blot analysis with anti-acrolein antibody (Fig. 7). Acrolein-modified proteins other than the 20S β -subunit were also detected in the immunoprecipitant, suggesting that the oxidized proteins might serve as substrates of 20S proteasome and the aggregated ones might inhibit the activity. However, it requires further

studies to clarify whether the acrolein-modified protein can directly inhibit the proteasome activity, as in the case of 4-HNE proteins (32).

Apoptotic Signal Activated by ROS-RNS

Both SIN-1 and NO induce apoptosis in SH-SY5Y cells by opening a megachannel called mitochondrial permeability transition pore (mPTP) with a decline in mitochondrial membrane potential ($\Delta\Psi_m$), as shown in Fig. 8. Opening of mPTP induces matrix swelling and physical disruption of the outer membrane in mitochondria and the release of Ca^{2+} , cytochrome-*c*, and apoptosis-inducing factors into cytoplasm. The mPTP is a protein complex composed of a voltage-dependent anion channel and other components located at the contact site between the mitochondrial outer membrane and inner membrane, and the opening is regulated by Bcl-2 in the outer membrane and the adenine nucleotide translocator in the inner membrane (2). Bcl-2 family proteins regulate cell death induced by oxidative stress and other stimuli, either in a preventing way (Bcl-2 and Bcl-xL) or a promoting way (Bax, Bad, and Bid). The antiapoptotic function of Bcl-2 is mediated by maintaining