



Review Article

Biochemical and clinical features of hereditary hyperprolinemia

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Abstract There are two classifications of hereditary hyperprolinemia: type I (HPI) and type II (HPII). Each type is caused by an autosomal recessive inborn error of the proline metabolic pathway. HPI is caused by an abnormality in the proline-oxidizing enzyme (POX). HPII is caused by a deficiency of Δ -1-pyrroline-5-carboxylate (P5C) dehydrogenase (P5CDh). The clinical features of HPI are unclear. Nephropathy, uncontrolled seizures, mental retardation or schizophrenia have been reported in HPI, but a benign phenotype without neurological problems has also been reported. The clinical features of HPII are also unclear. In addition, the precise incidences of HPI and HPII are unknown. Only two cases of HPI and one case of HPII have been identified in Japan through a questionnaire survey and by a study of previous reports. This suggests that hyperprolinemia is a very rare disease in Japan, consistent with earlier reports in Western countries. The one case of HPII found in Japan was diagnosed in an individual with influenza-associated encephalopathy. This suggests that HPII might reduce the threshold for convulsions, thereby increasing the sensitivity of individuals with influenza-associated encephalopathy. The current study presents diagnostic criteria for HPI and HPII, based on plasma proline level, with or without measurements of urinary P5C. In the future, screening for HPI and HPII in healthy individuals, or patients with relatively common diseases such as developmental disabilities, epilepsy, schizophrenia or behavioral problems will be important.

Key words hyperprolinemia type I, hyperprolinemia type II, inborn error of metabolism, P5C, proline.

Hyperprolinemia type I (HPI) results from a deficiency of proline oxidase (POX; enzyme commission no. [EC] 1.5.99.8), also called proline dehydrogenase (PRODH). The POX gene (*PRODH*) is located on chromosome 22 (22q11.21). This region is deleted in a congenital malformation syndrome known as velocardio-facial syndrome. This gene locus is also related to susceptibility to schizophrenia. Another form of hyperprolinemia is called hyperprolinemia type II (HPII). HPII is caused by a deficiency of Δ -1-pyrroline-5-carboxylate (P5C) dehydrogenase (P5CDh; EC 1.5.1.12) activity.^{1,2} The P5CDh gene (*ALDH4A1*) is located on chromosome 1 (1p36.13).

Schafer *et al.* were the first to document direct involvement of an error in proline metabolism in human disease in 1962.³ The affected family members had hyperprolinemia, cerebral dysfunction, renal anomalies, hereditary nephropathy, and deafness. Subsequently, many families with hyperprolinemia were reported in the literature.^{4–14} Several phenotypes of hyperprolinemia were identified, including those in some asymptomatic family members. Several authors have discussed the potential coincidental association between these clinical features and hyperprolinemia. Recently, our understanding of the associations between proline metabolism, biological function, and disease has

increased dramatically. There are now at least six enzymes, three transporters, and seven structural genes known to be directly involved in the interconversions of proline and its immediate metabolites. The functional characteristics, tissue distribution, regulation, and subcellular location of these proteins have been determined, and, for some, structural information is available.^{1,2} The exact clinical features and the frequencies of HPI or HPII, however, remain unknown. The aim of this study was therefore to assess the incidence of HPI and HPII in Japan, and to evaluate the cases reported previously. In addition, we propose diagnostic criteria for hyperprolinemia and identify a need for future studies on this disorder.

L-Proline metabolism

Tissue and fluid concentrations of L-proline are primarily related to the balance between the enzymatic activities of POX and P5C reductase. In this context, P5C plays a pivotal role in maintaining the concentration of proline in body fluids. Inborn errors of proline metabolism can, therefore, result from a disturbance of P5C-associated enzyme activity. HPI and HPII are differentiated by distinct biochemical and genetic deficiencies in the proline catabolic pathway (Fig. 1). HPI is caused by a deficiency in POX, a mitochondrial inner-membrane enzyme that converts proline to P5C, the first step in the conversion of proline to glutamate. P5CDh, a mitochondrial matrix NAD(+)-dependent dehydrogenase, catalyzes the second step of the proline degradation pathway. Deficiency of this enzyme is associated with HPII, an

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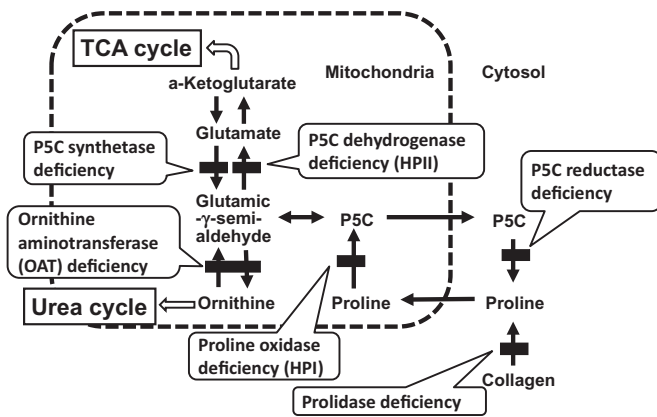


Fig. 1 L-Proline metabolism and enzyme deficiencies. The metabolism of proline in mammals involves two other amino acids, glutamate and ornithine, and six enzymes. Conversion of P5C to glutamic- γ -semi-aldehyde is a non-enzymatic step. HPI, hyperprolinemia type I; HPII, hyperprolinemia type II; P5C, Δ -1-pyrroline-5-carboxylate; TCA, tricarboxylic acid.

autosomal recessive disorder characterized by accumulation of P5C and proline (Fig. 1).^{1,2}

Human disease in proline metabolism

In human inborn errors of metabolism of L-proline, HPI, HPII, P5C synthetase deficiency, P5C reductase deficiency, ornithine aminotransferase deficiency and prolidase deficiency are known as single-gene disorders. The characteristics and molecular biology of these enzymes in humans are summarized in Table 1. Of these diseases, hyperprolinemia is seen only in HPI and HPII.²

Hyperprolinemia type I is diagnosed biochemically on high plasma proline without urinary excretion of P5C. In contrast, the presence of P5C in the urine is indicative of HPII. The clinical phenotype of HPI is not well characterized. Prior to the identification of *PRODH*, more than 15 families were reported in the literature with proline levels ranging from twofold to 10-fold above normal.^{3–16} Several phenotypes were found in these families. Whereas some patients were asymptomatic, others had neurological, renal and/or auditory defects. The potential for a coincidental association between these clinical features and HPI has been discussed.^{10–13}

Hyperprolinemia has been reported in patients with a microdeletion in the 22q11 region.^{15–17} Since the identification of *PRODH*, various mutations of this enzyme (heterozygous and homozygous) have been identified in hyperprolinemic patients with different phenotypes (i.e. patients with schizophrenia, a 22q11 microdeletion and/or early neurological impairment).^{17–20} Jacquet *et al.* identified a complete homozygous deletion of *PRODH* located on chromosome 22q11 in a child with a severe form of HPI.^{17,18} That child had severe psychomotor delay and status epilepticus associated with a very high level of plasma proline (2246 $\mu\text{mol/L}$). These studies show, unambiguously, that the severe form of HPI, characterized by neurological manifestations, results from homozygous inactivating alterations of *PRODH*.

Table 1 Human enzymes involved in proline metabolism

EC	Subcellular location	Subunit size (kDa)	Structure	Cofactor	Activity in tissues	Disease association	Map location	Gene/locus MIM	Proline oxidase	P5C dehydrogenase	P5C synthetase	P5C reductase	OAT	Prolidase
EC 1.5.99.8	Mitochondrial inner membrane	63	Unknown	Unknown	Liver, kidney, brain	HPI	22q11.21	<i>PRODH</i> 606810						
EC 1.5.1.12	Mitochondrial matrix	62	Homodimer	NAD ⁺	Ubiquitous	HPII	1p36.13	ALDH4A1 606811						
EC 1.5.1.2	Cytoplasm	32	Homopolymer	NAD(P)H	Ubiquitous	Cutis laxa	17q25.3	<i>PYCR1</i> 179035						
EC 2.6.1.13	Mitochondrial matrix	49	Homohexamer	Vitamin B6	Ubiquitous	Gyrate atrophy	10q26.13	OAT 613349						
EC 3.4.13.9	Cytoplasm	54.3	Homodimer	Mn ²⁺ , Ca ²⁺ , Mg ²⁺ , Fe ²⁺	Ubiquitous	Prolidase deficiency	19q13.11	<i>PEPD</i> 613230						

ALDH, aldehyde dehydrogenase; EC, enzyme commission number; HPI, hyperprolinemia type I; HPII, hyperprolinemia type II; MIM, mendelian inheritance in man; OAT, ornithine aminotransferase; P5C, Δ -1-pyrroline-5-carboxylate; PEPD, peptidase; *PRODH*, proline dehydrogenase; *PYCR1*, pyrroline-5-carboxylate reductase 1.

The *PRODH* mutations were divided into three groups: those leading to mild (<30%); moderate (30–70%); and severe (>70%) reductions of POX activity. Serum proline level seems to correlate with the severity of POX deficiency.^{19,20} Higher serum proline was associated with lower IQ in patients with the 22q11 deletion (including the region of *PRODH*) syndrome.^{19,21} Recently, Guilmatre *et al.* also reported genotype/phenotype correlations in HPI.²²

Hyperprolinemia type II is due to the absence of P5CDh activity. Valle *et al.* measured P5CDh activity in normal fibroblasts and fibroblasts from patients with HPII.²³ The HPII fibroblast cells had no detectable P5CDh activity over a range of reaction conditions, whereas normal cells had easily measurable activity. This enzymatic defect accounts for the biochemical abnormalities in HPII, which is usually characterized by mental retardation and convulsions.

Geraghty *et al.* surveyed *ALDH4A1* from four affected members of a family with HPII.²⁴ Four mutant alleles were found. A 1 bp deletion (G) at nucleotide 21 (A7fs(-1)) and a 1 bp insertion of a T following nucleotide 1563 (G521fs(+1)) cause a frameshift. The other two are missense mutations (S352L and P16L). To test the functional consequences of three of these mutations, they were expressed in a P5CDh-deficient strain of *Saccharomyces cerevisiae*. In contrast to yeast expressing wild-type human P5CDh, yeast expressing S352L and G521fs(+1) failed to grow on proline and had no detectable P5CDh activity. Interestingly, the G521fs(+1) allele, segregated in the large Irish traveler pedigree reported by Flynn *et al.*,²⁵ was used to define the HPII phenotype. To our knowledge, this is the first description of the molecular basis for this inborn error.

Hyperprolinemia type II was identified in a pedigree of Irish travelers (nomads), among whom consanguineous marriage and high fertility are common. Flynn *et al.* studied 312 subjects in 71 families closely related to a proband with HPII. Thirteen additional cases of HPII were discovered. A further 50 subjects were found to have mild hyperprolinemia. Flynn *et al.* found a strong association between HPII and seizures during childhood but no significant association with cognitive function. Most adults in this group with HPII enjoyed normal health.²⁵

Some neuropsychiatric disorders appear to be related to hyperprolinemia. The prevalence of schizophrenia among patients with the 22q11 interstitial deletion associated with DiGeorge syndrome was high. This suggests the existence of a susceptibility gene for schizophrenia within the DiGeorge syndrome chromosomal region. This deletion was associated with hyperprolinemia in schizophrenia patients. In addition, two heterozygous mutations (L441P and L289M) in *PRODH*, identified in three of 63 patients with schizophrenia, but not in the 68 controls, were also associated with increased plasma proline.^{17,18} Bender *et al.* identified at least 16 missense mutations in *PRODH* in studies of HPI and schizophrenia, 10 of which were present at polymorphic frequencies.²⁶ Although there is limited information on plasma proline level in the individuals with known *PRODH* genotype, extant data suggest that severe hyperprolinemia (>800 mol/L) and the most severe effect on function occurs in individuals with large deletions and/or missense mutations

(L441P and R453C) in *PRODH*; modest hyperprolinemia (300–500 mol/L) is associated with *PRODH* alleles exhibiting a moderate reduction in POX activity. Interestingly, three of the four alleles associated with, or found in, schizophrenia (V427M, L441P, and R453C) resulted in a severe reduction of POX activity and hyperprolinemia.

Proline oxidase functions are also related to tumor growth, hypoalimentation stress reaction, and fetal growth. Recent studies found that POX can regulate tumor growth both by tumor suppression and increasing tumor survival.^{27,28}

Hyperprolinemia in Japan

The present study investigated hyperprolinemia patients in Japan in 2009. The study was approved by the Ethics Committee of the Faculty of Life Science, Kumamoto University. A questionnaire was sent to the Departments of Pediatrics, Neonatology, Genetics, and Transplant surgery in 928 institutions in Japan. Institutions were asked to confirm that they had diagnosed and/or treated HPII. A secondary detailed questionnaire concerning HPII was then sent to those confirming institutions. We also examined previous patient reports on hyperprolinemia in Japan. Two reports of HPI and one report of HPII were identified and analyzed. According to the secondary questionnaire, the HPII patient was a 4-year-old boy. His parents were not consanguineous, and his brother died at 11 months of age due to sudden infant death syndrome. The HPII patient had no seizures and had normal development until 4 years of age, when he suddenly developed high fever progressing to generalized convulsions and unconsciousness. He was admitted to hospital and diagnosed with influenza encephalopathy. His blood proline was elevated (2767 $\mu\text{mol/L}$), while his blood ornithine was normal. Urinary P5C was detected. After treatment for encephalopathy, the patient recovered, although a high level of blood proline and urinary P5C excretion continued. He was then diagnosed with HPII. He currently is not undergoing any treatment, has no convulsions, his IQ score is 96, and his growth is within the normal range.^{29,30}

Two cases of HPI have been reported in Japan; one in 1987 and one in 1991. The 1987 case was accompanied by chromosomal abnormality. This patient had below-normal activity of POX and hyperprolinemia, although no relation between blood proline level and clinical features was established.³¹ The 1991 case involved a 9-year-old girl who had photosensitive epilepsy. She had developmental delay and developed epilepsy at 7 years of age. Her blood proline level was three–fourfold higher than normal. P5C was not detected in the urine. POX activity in the liver was 23.5% of normal. A proline-restricted diet was not effective in preventing convulsions.³² Overall, details of these two cases are limited because the reports are more than 20 years old.

Amino acids were analyzed in plasma samples from 200 severely handicapped patients living in a special institution in Kumamoto, as well as in plasma samples from 500 healthy individuals. None of the samples indicated hyperprolinemia, although some of the samples from normal individuals had plasma proline 2.5 SD higher than normal. Analysis of plasma proline in samples from elderly patients in a nursing home identified one case of suspected hyperprolinemia in a 90-year-old

patient but, despite additional analysis, the patient was too old to confirm causality (data not shown).

Most newborn screening tests are done by measuring amino acids in whole blood samples collected on specialized filter paper. Almost all amino acids can be analyzed by this method. In Korea, blood proline is analyzed as part of newborn screening, and four neonates were identified as having HPI.³³ Similar newborn screening for proline is needed to determine the incidence of hyperprolinemia in Japan.

Diagnostic criteria for HPI and HPII

The exact clinical features of HPI are unclear. Nephropathy, uncontrolled seizures, mental retardation, or schizophrenia have been associated with HPI, but HPI without neurological manifestations is also known. The clinical features of HPII are similarly variable. Therefore, clinical signs and symptoms cannot be used to diagnose HPI or HPII. Patients who have uncontrolled seizures or developmental delay or encephalopathy of unknown origin should undergo blood amino acid analysis. When hyperprolinemia is identified, the patient may be diagnosed with HPI or HPII. A flow chart for the diagnosis of HPI and HPII is given in Figure 2. In HPI, blood proline is usually fivefold–10-fold higher than the normal range of 51–271 $\mu\text{mol/L}$. In HPII, blood proline is even higher than in HPI, usually exceeding 1500 $\mu\text{mol/L}$. Lactic acidosis can lead to secondary hyperprolinemia. Thus, any diagnosis of HPI or HPII must rule out lactic acidosis. If either HPI or HPII is suspected, analysis of urinary organic acids on gas chromatography–mass spectroscopy is useful. If P5C is detected in urinary samples, a diagnosis of HPII is confirmed. If P5C is not detected, the diagnosis is HPI. The activities of the different enzymes affected in HPI and HPII can be determined using cultured lymphocytes, peripheral leukocytes, or cultured fibroblast. Each affected gene has been cloned

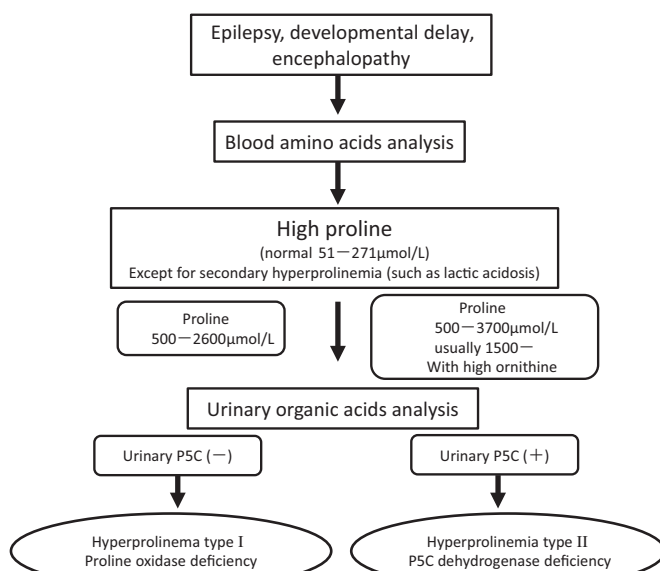


Fig. 2 Diagnostic criteria for hyperprolinemia types I and II. P5C, Δ -1-pyrroline-5-carboxylate.

previously, and, therefore, gene analysis is also possible. Many of the mutations on these genes are described in the previous sections.

Treatment and prognosis

Treatment of the hyperprolinemia seen in HPI has included dietary therapy such as the restriction of proline. Unfortunately, this results in only modest control of plasma proline level and has no impact on clinical phenotype.³⁴ As a result, dietary therapy may not be necessary.¹

The seizures associated with the rare inherited disorder HPII are usually precipitated by an infection during childhood. The control of seizures in HPII is clearly important. It is necessary to protect against febrile convulsions and status epilepticus. As in HPI, dietary therapy is unable to significantly reduce plasma proline level in HPII. P5C is a unique endogenous vitamin antagonist. The inactivation of vitamin B6 by P5C may contribute to seizures in HPII, which are so far unexplained. The evidence for this largely rests on the diagnosis of HPII and vitamin B6 deficiency in a well-nourished child with seizures. Pyridoxine deficiency was implicated in her seizures and may have been the result of inactivation of vitamin B6 by the proline metabolite, P5C. Long-term vitamin B6 supplementation may therefore prevent these seizures.^{35,36}

Recently, oxidative stress was detected in the brain tissue from rats with hyperprolinemia. Antioxidants such as vitamin E, vitamin C, and glutathione may be effective therapeutic agents in this disorder. van de Ven *et al.* suggested that these agents should be used for hyperprolinemia in patients as young as possible.³⁷

If acute manifestations of hyperprolinemia such as convulsions are controlled safely, the prognosis for these disorders is good. Normal development can be expected, although the long-term outcome in terms of neurological disturbances is unknown. In general, individuals with hyperprolinemia should be monitored closely due to their frequent behavioral problems.³⁸

Conclusions

Hereditary hyperprolinemia is a very rare disease in Japan. The patients that have been identified have a variety of neurological abnormalities. Screening and investigation of healthy individuals as well as patients with relatively common disorders such as developmental disabilities, epilepsy, schizophrenia, and behavioral problems will be important in Japan. Analysis of plasma amino acids can be used to identify HPI and HPII. The current study has presented diagnostic criteria for these diseases. In addition, a strong relationship between proline metabolism and tumor growth, as well as hypoalimentation stress reaction and fetal growth has been reported. Further studies will provide additional insights into the problems associated with hyperprolinemia, and may lead to new therapeutic strategies.

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