

Nutritional biochemistry

A new redox-cofactor vitamin for mammals

Nicotinamides and flavins are essential cofactors in enzyme-catalysed reduction–oxidation (redox) reactions and are classified as vitamins because they must be supplied in the diet. Another redox cofactor, pyrroloquinoline quinone (PQQ), was first discovered in bacteria¹ and is also likely to be important in mammals^{2,3}, but the biochemical pathways in which it participates are unknown. Here we identify a PQQ-dependent dehydrogenase enzyme that is crucial for the degradation of the amino acid lysine in mice. PQQ is acting as a mammalian redox cofactor in this reaction, and therefore qualifies as a newcomer to the B group of vitamins.

In animals, lysine is an essential amino acid (that is, one that cannot be made in the body) that is degraded after use to 2-aminoadipic 6-semialdehyde (AAS), which is subsequently oxidized to 2-aminoadipic acid (AAA; Fig. 1a, right arrows)⁴. The molecular features of the AAS-dehydrogenase (AASDH; EC 1.2.1.31) enzyme that catalyses this oxidation of AAS are unknown.

In yeast, lysine is synthesized from AAA by the reverse pathway (Fig. 1a, left arrows)⁵. The reduction of AAA to AAS requires two enzymes that are encoded by the *Lys2* and *Lys5* genes: the *LYS2* protein has NADPH-dependent AAA-reductase activity, whereas *LYS5* catalyses the post-translational 4'-phosphopantetheinylation of *LYS2* (ref. 6). A human homologue of *Lys5* has been cloned⁷, raising the possibility that there is a structural analogue of *LYS2* in animals.

To find the analogue of *LYS2*, we searched the *Drosophila* genome databases for gene product(s) containing functional domains, adenylation and thiolation domains, which serve to capture the substrate in *LYS2* (ref. 6). A BLAST search detected two gene products, EBONY (DDBJ/EMBL/GenBank accession number, CAA11962) and U26 (accession number, AAF52679); EBONY is β -alanyl-dopamine synthetase, but the function of U26 was unknown. In contrast to yeast *LYS2*, which contains a binding domain for the reducing cofactor NADPH, *Drosophila* U26 contains several repeats of the binding motif for PQQ (an oxidizing cofactor); this motif is conserved among bacterial PQQ-dependent dehydrogenases⁸.

To investigate whether U26 could be AAS-dehydrogenase, we cloned mouse U26 complementary DNA (accession number, AB095954), which encodes a protein of 1,100 amino acids that is 23% identical and 43% similar to *Drosophila* U26. Mouse U26 contains one set of adenylation/thiolation

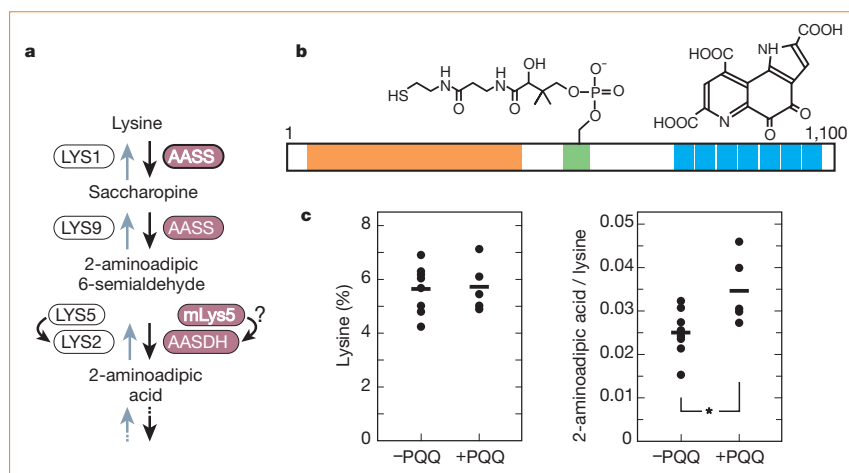


Figure 1 The role of pyrroloquinoline quinone (PQQ) in mammalian lysine degradation. **a**, Metabolic pathways of lysine: right arrows, lysine degradation in mammals; left arrows, lysine biosynthesis in yeast. In yeast, *LYS2* is post-translationally 4'-phosphopantetheinylated in a reaction catalysed by *LYS5*. AAS, AAS-synthetase; AASDH, AAS-dehydrogenase; mLys5, mammalian homologue of *LYS5*. **b**, Arrangement of putative binding domains in murine U26 protein. The adenylation domain (Pfam 00501), thiolation domain (00550), and PQQ-binding motifs (01011) are shown in orange, green and blue, respectively. Serine 592 in the thiolation domain is 4'-phosphopantetheinylated and PQQ (right) is non-covalently bonded by the seven repeats of the PQQ-binding motif near the carboxy terminus. **c**, Peripheral blood concentrations of lysine and 2-aminoadipic acid in PQQ-supplemented and PQQ-depleted mice. Lysine is quantified as a percentage of the total amino acids (left; determined using an amino-acid analyser) and 2-aminoadipic acid is normalized with respect to lysine (right). Horizontal bars, average for each group of mice. Asterisk denotes $P = 0.010$.

domains and seven PQQ-binding motifs (Fig. 1b). The six to eight tandem repeats are also present in bacterial PQQ-dependent dehydrogenases⁸, indicating that mouse U26 could be a PQQ-dependent dehydrogenase.

Northern-blot analysis showed that mouse U26 messenger RNA is expressed in all of the tissues that we tested, with expression being notably high in the heart, liver and kidney (results not shown). In mice fed on a lysine-rich diet (containing 2,000-fold more free lysine than normal), blood concentrations of lysine and AAA were markedly increased (2.4- and 1.5-fold, respectively). The level of U26 mRNA (normalized to that of glyceraldehyde-3-phosphate dehydrogenase) in the liver and heart of lysine-loaded mice was significantly higher (1.6- and 1.4-fold, respectively; $P = 0.008$ for each) than in control mice. Mouse AAS-synthetase (AAS in Fig. 1a) and *Lys5* transcripts were also increased in the lysine-loaded mice (results not shown). These results indicate that U26, as well as AAS-synthetase and *LYS5*, is involved in the lysine-degradation pathway in mice.

To test the importance of PQQ in AAS-dehydrogenase activity, we examined the effects of a PQQ-deficient diet on mice. After being fed on a chemically defined diet^{2,3} that was either devoid of PQQ or supplemented with PQQ (1,000 ng PQQ-2Na per g of food), we measured the blood levels of lysine and AAA in the two groups of mice. AAA concentrations were significantly decreased in PQQ-depleted mice, whereas lysine levels remained the same (Fig. 1c); the intermediates saccharo-

pine (Fig. 1a) and AAS were not detected. This decrease in AAA may be indicative of impaired AAS-dehydrogenase (U26) activity resulting from a PQQ deficiency in deprived mice, pointing to a requirement for PQQ as a redox cofactor for the proper functioning of AAS-dehydrogenase.

PQQ is found in various foods, including vegetables and meat^{9,10}. When mice are fed a PQQ-deficient diet, they grow slowly, have fragile skin and a reduced immune response, and do not reproduce well^{2,3}. On the basis of our demonstration of its molecular function, we propose that PQQ should be classified as a new B vitamin, joining niacin/nicotinic acid (vitamin B3) and riboflavin (vitamin B2), the redox-cofactor derivatives of which are NAD^+ / NADP^+ and FAD/FMN, respectively.

Takaoki Kasahara, Tadaaki Kato

Laboratory for Molecular Dynamics of Mental Disorders, Brain Science Institute, RIKEN, Wako-shi, Saitama 351-0198, Japan
e-mail: kato@brain.riken.go.jp

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