Appreciation

Vernon Young and the development of current knowledge in protein and amino acid nutrition
Vernon Young 1937–2004

Personal note

Vernon Young, undoubtedly the best-known and most highly honoured nutritionist globally of his generation and a unique personality in the history of nutrition, died 30 March 2004 of complications from renal cancer. He was 66. At the time of his death he held posts at Massachusetts Institute of Technology (MIT), Massachusetts General Hospital, Harvard Medical School, Tufts University, and was Director of the Mass Spectrometry Facility at Shriners Burns Institute, Boston. His many honours and prizes started in 1973 with the Mead-Johnson Award from the American Institute of Nutrition, included the Bristol-Myers Squibb/Mead Johnson Award for Distinguished Achievement in Nutrition Research in 1995, the Danone International Prize for Nutrition in 1997 and most recently the Danone International Prize for Nutrition in 1997 and most recently the Conrad Elvehjem Award from the American Society for Nutritional Sciences in 2003. He was elected to the National Academy of Sciences USA in 1990 and to the Institute of Medicine in 1993 in recognition of his scientific accomplishments in the field of human nutrition. In 2001, he was named to the Board of Directors of Nestlé Corporation, which sought his nutritional expertise.

Vernon was born in Rhyl, North Wales on 15 November 1937, but lived in Cardiff from an early age. His twin sister says that his interests in agriculture developed during his visits to an uncle’s farm in Nottinghamshire. He studied agriculture at Reading (BSc 1959) and Cambridge (postgraduate diploma 1960). He moved to the University of California at Davis in 1960 to gain his PhD in 1965 for work on Ca and P homeostasis in sheep, moving to MIT to work with Nevin Scrimshaw and meeting Janice his wife-to-be that same year.

His work was characterised by three main interests; human protein and amino acid metabolism and requirements, nutrition and ageing and nutrient bioavailability. His output was quite remarkable (especially given two triple bypass operations), just short of 600 full-length peer-reviewed papers, book chapters and reviews, more than one paper per month for 42 years and at his peak in 1980 nearly one paper per week (Fig. 1). He was a man who lived for his work and thought about little else apart from his wife Janice and his five children.

Throughout this work I have had very similar interests, closely following Vernon’s work having first met him in the late 1960s when a life-long friendship was established with him and his wife. This was notwithstanding the lively debate which characterised our scientific relationship. In what follows I have made no attempt to present a comprehensive review of his work, instead presenting a personal and selective view of his major interests in the context of the development of our understanding of protein and amino acid nutrition.

Defining protein requirements

As detailed in the excellent series on the history of nutritional science (Carpenter, 2003a–d), following the identification of N by Rutherford in the late 18th century, our understanding of the nutritional importance of protein began in Paris.
in the early 19th century. This is when the essentiality of dietary N was established by the feeding trials on dogs by Magendie and the balance studies on herbivores of Boussingault. These studies, together with the chemical identification of protein by Mulder, established protein as the determinant of food ‘quality’ and the food constituent which dominated nutritional science for the rest of the 19th century. Furthermore, on the basis of measuring the protein intakes of working men and soldiers, all meat eaters of course, Voit and Peltenkofer, Atwater and Rubner argued that relatively large amounts of protein, 120–130 g, were needed in the diet. There were some dissenting voices, notably John Harvey Kellogg and most famously Chitten- don, who both advocated less meat and lower protein intakes as part of the healthy diet. Nevertheless, from the early 20th century the majority view was that health and especially fitness depended on generous amounts of protein in the diet, especially meat, and that populations that existed on low levels of protein were usually extremely unfit. This view was reinforced by Cecily Williams’ suggestion that kwashiorkor, which she had described in West Africa in the 1930s, was due to protein deficiency (Williams, 1935). The ultimate statement of the primacy of protein in the diet came in a report published in 1968 under the auspices of the UN identifying widespread protein deficiency and calling for ‘a global strategy to deal with the impending protein crisis’ (United Nations, 1968). Thus, provision of protein came to dominate international aid initiating the green revolution to improve yields of cereals in the developing world and with soyabean production becoming of extraordinary importance within US agricultural production.

Whilst this dominant role of protein was not universal and seen as ‘a protein myth’ by some (see McLaren, 1974; Waterlow & Payne, 1975), nevertheless this was the climate of opinion within which the modern era of international expert committees to define protein requirements had begun with a report from the FAO in 1957 (Food & Agriculture Organization, 1957). This first report was followed quickly by a second from the FAO and WHO in 1965 (Food & Agriculture Organization/World Health Organization, 1965). It was at this point that Vernon Young’s work, which was to dominate the next 40 years of the development of our understanding of protein and amino acid requirements, began.

The 1965 FAO/WHO report (Food & Agriculture Organization/World Health Organization, 1965) had derived a safe protein requirement value for adults of 0.89 g animal protein/kg per d (1.12 g plant proteins/kg per d). The requirement had been derived by a factorial method based on an obligatory N loss (ONL) value (losses on a protein-free diet), with additions for interindividual variability. The database from which these values were derived was very limited, so Young and Scrimshaw immediately started studies to better define the ONL, publishing initial data in 1968 (Young & Scrimshaw, 1968; Young’s first Nature paper, Young et al. 1968b) and the definitive study in 1972 (Scrimshaw et al. 1972). These data formed the basis of the next Food & Agriculture Organization/World Health Organization (1973) report which was a much more thorough, scientifically based, document than any previous one. Although N-balance data were examined by the Committee, these were quite limited at the time, so the report writers again adopted a factorial method but used the new extensive data from Young and Scrimshaw for the ONL. Because these were much lower than the previous values used in the 1965 report, this resulted in an adult protein requirement, after correction for efficiency of utilisation and interindividual variability, of 0.57 g/kg per d. This lowering of the protein requirement had a dramatic effect within the nutritional community and began a major controversy.

To understand the impact of this report (Food & Agriculture Organization/World Health Organization, 1973), it is important to recognise the major problem within protein nutrition which is still unresolved. This is how to identify protein deficiency in the absence of any definitive markers of nutritional status in relation to protein adequacy, an especially difficult problem in adults. Without such markers, protein deficiency can only be defined in terms of intakes below an assumed requirement level. Thus the prevalence of protein deficiency changes as protein requirements change and the 1973 report, by lowering the protein requirement, reduced the prevalence of protein deficiency; this was contrary to the spirit of the 1968 UN ‘impending protein crisis’ report (United Nations, 1968).

Young and Scrimshaw’s extensive studies at MIT became the dominant influence within this controversy. They reported a major series of N-balance studies on young adult men, both short- and long-term studies. The short-term balances (Young et al. 1973) essentially confirmed the 1973 value (Food & Agriculture Organization/World Health Organization, 1973) consistent with contemporary work from Calloway & Margen (1971) at the University of California at Berkeley, the other major centre working on protein requirements at the time. However, Young and Scrimshaw’s long-term balances (Garza et al. 1977a,b, 1978) carried out specifically to test the safety of the 1973 safe allowance (0.57 g protein) showed that this level of intake resulted in negative N balance, loss of lean body mass, deteriorating values for serum protein and increased serum transaminases unless additional energy or N was fed. These alarming findings led to calls by Scrimshaw (for example, Scrimshaw, 1977) for an upwards re-evaluation of the 1973 report recommendations, and to the initiation of an international programme of short-term balance studies with...
a common methodology organised by Young and Scrimshaw through the UN University World Hunger Programme. The outcome of these studies, reported in early 1981, was a mean requirement of 0·6 g/kg per d (Torun et al. 1981) and a safe allowance 20–30% higher than this. As a result, the agenda of the next FAO/WHO/UN University meeting in 1981 was set and the report (Food & Agriculture Organization/World Health Organization/United Nations University, 1985), on the basis of both short- and long-term balance data, duly raised the safe allowance to 0·75 g/kg per d. This value was adopted by national bodies around the world. Apart from N-balance studies on the elderly (Gersovitz et al. 1982) following up earlier work (Uauy et al. 1983), the early 1980s were to mark the end of Young’s N-balance studies aimed at the adult protein requirement. The magnitude of this part of his work is apparent from his recent meta-analysis of all published data on adults (Rand et al. 2003), used as the basis of the recent US dietary reference intakes report on protein requirements (Institute of Medicine, 2002). Of the 118 assembled references, thirty-two studies were his work, with many more reported by workers he trained at MIT.

Protein metabolism in skeletal muscle

A second main interest which he developed soon after arriving in MIT from Davis was the regulation of protein metabolism in skeletal muscle, in part inspired by his second mentor at MIT, Hamish Munro. Against the background of interest in malnutrition, the mechanisms of growth failure and tissue wasting had emerged as a major focus of interest in the 1960s. From the mid 1960s, the preparation and examination of ribosome fractions from skeletal muscle became a fruitful approach to the study of protein synthesis (Chen & Young, 1968), especially important in muscle where changes in protein synthesis in response to insulin and other factors had been shown to occur very rapidly (for example, Young et al. 1968a). Young published a series of studies of the effects of dietary protein, infection and hormones on muscle ribosomes focusing on the translational phase of protein synthesis at the level of both initiation and elongation. However, at that time there were several competing groups working on muscle and to his great disappointment his National Institutes of Health grant-support for his muscle work, which he saw as his first fully independent research programme, was not renewed. He told me at the time that he believed this reflected a view that nutritionists should not be working within the emerging field of cellular and molecular biology. This aspect of his work ended in 1974 with a publication relating to elongation factor 2 in rat skeletal muscle during protein depletion and re-feeding (Alexis et al. 1974).

However, in parallel with this in vitro work he was developing a new approach to the in vivo study of muscle protein degradation. During the preparation of a major review on protein metabolism in skeletal and cardiac muscle for Munro’s classic series on mammalian protein metabolism (Young, 1970) he became interested in the presence in specific muscle proteins of methylated amino acids of which 3-methylhistidine only occurred in actin and some types of myosin heavy chain. This led to a series of classic studies showing that this was a post-translational modification and that 3-methylhistidine, once released during actin proteolysis, was not re-utilised for protein synthesis. Since it was not further metabolised in vivo its urinary excretion could serve as a measure of skeletal muscle proteolysis (Young & Munro, 1978). This paper was the basis of his Rank Prize in Nutrition in 1989, and became a citation classic in 1992, demonstrating how widely adopted the method became. Although subsequent work has showed that actin turnover in non-muscle tissue is a major fraction of urinary 3-methylhistidine excretion, reducing the specificity of the method (Millward et al. 1980; Millward & Bates, 1983), the idea was a good example of Young’s single-minded and inventive approach to science.

Stable isotopes, protein turnover and amino acid requirements

Young’s best-known work today relates to his development of the use of stable isotopes in his search for methods to build on and replace N balance in assessing protein and amino acid requirements. The use of stable isotopes to probe the dynamic aspects of protein metabolism was initiated by the brilliant studies of Rudolf Schoenheimer in the late 1930s soon after 15N and 2H became available. These studies were developed by Rittenberg and others, after Schoenheimer’s untimely death in 1941, into increasingly mathematically complex compartmental analyses with 14N tracers. Few could understand these methods and development was slow until in the early 1960s John Waterlow working for the UK Medical Research Council in Jamaica rationalised and simplified the methodology by introducing stochastic steady-state tracer methodologies with 15N tracers (glycine, urea and ammonium salts) in human subjects and [13C]lysine studies in animals (Waterlow et al. 1978). The motivation behind this work was that body protein is in a dynamic state, with overall rates of protein synthesis several-fold greater than protein intakes. Thus, the tissue protein mass can be controlled not only by the relative rates of dietary protein supply and body N losses, as measured by N balance, but also by adaptive changes in the rates of re-utilisation and redistribution of the amino acids released by protein breakdown. This would allow adaptation to varying intakes of protein over a wide range, and conditions such as kwashiorkor represented the consequence of a failure to sufficiently adapt to a reduced intake. Early work on albumin turnover in Jamaica strengthened this view (James & Hay, 1968). Young was impressed by this work, and adopted Waterlow as one of his mentors. This led to the establishment of appropriate isotope ratio MS facilities at MIT, a rare example of the Third World exporting leading-edge technology to the USA, enabling a series of studies of protein turnover on neonates (Pencharz et al. 1977), children (Kien et al. 1978a,b,c), adults (Steﬀee et al. 1976), and the
elderly (Winterer et al. 1976; Uauy et al. 1978b), all aimed at better understanding the protein requirement (Young et al. 1975b). However the pace of this work was slow by Young’s standards since isotope ratio MS analysis of 15N is laborious. The introduction of gas chromatography–MS in selective ion monitoring mode, allowing the automated analysis of isotopic enrichments of amino acids, resulted in a dramatic change of pace. This development of the technology occurred through the work of Bier and Matthews at Washington University at St Louis in the late 1970s with whom Young established a long-standing collaboration, as did the present author and colleagues in the UK (for example, Matthews et al. 1981). Thus, by 1980, gas chromatography–MS facilities had also been established by Halliday in the UK Medical Research Council Unit at Northwick Park.

The most important feature of this phase of Young’s work was his early identification of the importance of stable-isotope studies of amino acid oxidation and consequent balance, with 13C-labelled amino acids such as leucine, as a means of improving on N balance. Thus the balance technique, which had hitherto been used to assess the protein or amino acid intake which allowed body N equilibrium, was extended by utilising the balance of the individual amino acid from measurement of its oxidation as 13CO2 production. This allows balance measurements over a period of hours as opposed to days with N balance. By 1982 Young was starting to report his amino acid oxidation balance studies with 13C tracers as a new approach for the estimation of human amino acid requirements (Meredith et al. 1982). The complete series of 13C balance studies with leucine, valine, lysine and threonine were reported in 1986 over thirty-three pages of a single issue of the American Journal of Clinical Nutrition (Meguid et al. 1986ab; Meredith et al. 1986; Zhao et al. 1986), an extraordinary amount of work. Protein turnover, as opposed to balance, remained an important part of Young’s work for several years, enabling him to identify lower limits of leucine intake in terms of adaptation and accommodation (Young et al. 1987). However, by the end of the 1980s protein and amino acid turnover per se had become of much less importance within his work. Indeed, notwithstanding the explosion of protein-turnover studies in the 1980s and 1990s as gas chromatography–MS selected ion monitoring technology became widespread, Young believed that the initial promise that protein turnover would transform our understanding of protein requirements had not been realised.

The final phase of this aspect of his work involved recognising the limitations of his first 13C amino acid oxidation balance studies and refining the technique to its ultimate limits in terms of 24 h 13C amino acid oxidation balance studies with his student Antoine El Khoury in MIT which began in 1994 (El-Khoury et al. 1994), and then with Anura Kurpad in India from 1998. With this approach he was able to refine his estimates of the requirements for leucine (Kurpad et al. 2001b, 2003b), lysine (Kurpad et al. 2001a, 2002b, 2003a,c), threonine (Kurpad et al. 2002a) and cystine and methionine (Kurpad et al. 2003d, 2004).

**The Massachusetts Institute of Technology amino acid requirement pattern and protein quality**

The most fascinating feature of Young’s work on adult amino acid requirements over the final 25 years of his life was the way he organised and reported his stable-isotope work in relation to his proposal for what came to be called the MIT amino acid requirement pattern. Recognition of the concept of amino acid essentiality as the determinant of protein quality arose early in the 19th century through feeding trials with collagen extracted from bones. This showed that collagen protein was demonstrably nutritionally inferior to proteins such as albumin extracted from serum. The advances in chemical analysis of protein and protein hydrolysates in the latter part of the 19th century resulted in the identification of many individual amino acids by the early years of the 20th century. Hopkins isolated tryptophan, which in turn enabled the unusual composition of collagen and especially its lack of any tryptophan to be demonstrated by Kaufman, who showed that gelatin + tryptophan + cystine + tyrosine maintained N balance in dogs and man. The other protein which aroused interest through its inability to allow the growth of animals was zein, the principal protein extract from maize. Hopkins had shown that mice fed on zein (which contains very low levels of tryptophan) lived longer if they also received a supplement of tryptophan. Thus, the concept of protein quality determined by its amino acid content was established in animals. Also, N-balance studies with adults, such as those by Thomas in 1909 with different foods, rapidly established the importance of protein quality in human nutrition. The final refinements of the basic principle that amino acid content was an important determinant of the nutritional quality of protein were in place by 1915 with the experiments of Osborne & Mendel (1916), who demonstrated the different amino acid requirement patterns for maintenance and growth. Thus, they not only confirmed the Hopkins findings that rodents fed diets based on zein lost weight and died without tryptophan, but also showed that by supplementing first with tryptophan and then with additional lysine the rats not only maintained weight but also grew (i.e. zein was limited by tryptophan for maintenance and by lysine for growth). After this, the pioneering work of Rose identified all eight essential amino acids needed to maintain N balance in adult man (Rose, 1957). This work was followed up by studies throughout the USA which quantified the magnitude of amino acid requirements in adult men and women and these data formed the basis of both the Food & Agriculture Organization/World Health Organization (1973) and the Food & Agriculture Organization/World Health Organization/United Nations University (1985) reports on human amino acid requirements. Unfortunately a major flaw in all of these studies resulted in the controversy which raged for 20 years and which dominated Young’s work up to the time of his death.

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Nations University, 1985), had been critically reviewed by Hegsted (Hegsted, 1963; Irwin & Hegsted, 1971). Hegsted pointed out that because the plots of amino acid intake on N balance were curvilinear with very shallow slopes at N equilibrium, the intake for balance critically depended on the very small changes in balance which related to the assumed value for miscellaneous N losses (i.e., skin, sweat, hair, etc.). Because these losses are so small (4–8 mg N/kg per d) they had been ignored. In fact if small additions to N losses were made to account for these losses this had significant and in some cases quite marked effects on the intake for N equilibrium, i.e., the amino acid requirement. However, neither the 1973 nor the 1985 report took this into account and accepted uncorrected values which were, as a consequence, too low. The 1985 report went on to calculate age-related amino acid requirements scoring patterns on the basis of the safe level of proteins, which meant that the scoring pattern for adults involved such low levels of essential amino acids per g protein that all protein sources, including cereal proteins, previously assumed to be deficient in lysine or tryptophan, were nutritionally adequate. Thus protein quality was identified as being unimportant for adults.

In retrospect, given that the major focus of the 1985 report (Food & Agriculture Organization/World Health Organization/United Nations University, 1985) was on increasing the adult protein requirement, the abolition of the importance of protein quality in adults might be thought of as a very surprising decision of the committee. In fact the long-standing controversy of the lysine requirement and the nutritional importance of its low level in cereal proteins, especially wheat, has never been entirely straightforward. As recently highlighted by Carpenter (2003a), wartime studies in the UK by Widdowson and McCance had shown that 9–10-year-old orphanage children studied over 6 months grew in weight and height some 25 % faster than the general population on a bread-based diet with only 14 % from animal (milk) foods, with no advantage observed when nearly one half the protein was from milk. In fact, in children, with energy needs much higher than adults, their protein needs can be satisfied by much less protein-dense foods (Millward & Jackson, 2004), as evident by the low level of protein in breast milk (6 % energy) which mediates rapid postnatal growth. Because wheat is a relatively high-protein cereal (12–16 % energy) and the energy intakes of children of this age are high, the protein content of the diet of the orphanage children (12 % energy) provided them with two to three times their protein needs so that protein quality ceased to be important. More importantly, studies examining the nutritional quality of wheat protein in adults also concluded that wheat-based diets maintained N balance and fitness (Bolourchi et al. 1968), but these studies were criticised by Young because of their high energy intakes (Rand & Young, 1999). Furthermore Young’s own work comparing isonitrogenous intakes of wheat and beef showed clear differences in their utilisation (Young et al. 1975a).

Whatever the explanation of the Food & Agriculture Organization/World Health Organization/United Nations University (1985) report’s abolition of protein quality contrary to the stated view of one of its authors, by the time that it appeared in 1985 Young was well into the stable-isotope studies described earlier. He was aiming to rectify what he saw as a major flaw in the report and had completed the 13C amino acid oxidation balance studies, described earlier, which were published in 1986. These demonstrated much higher requirement values for all indispensable amino acids examined, most importantly for an increase in the lysine requirement from 12 mg/kg per d in the 1973 and 1985 reports (Food & Agriculture Organization/World Health Organization, 1973; Food & Agriculture Organization/World Health Organization/United Nations University, 1985) to 30 mg/kg per d. The MIT amino acid requirements pattern was published in 1989 (Young et al. 1989) in what Young believed to be his most important paper entitled ‘A theoretical basis for increasing current estimations of the amino acid requirements in adult man, with experimental support’. The key idea was that amino acid requirements for maintenance could be predicted by converting the obligatory N losses into an isonitrogenous equivalent pattern of obligatory oxidative losses of amino acids (on the basis that amino acids derived from tissue protein gave rise to the obligatory losses), scaled up assuming a 70 % efficiency of dietary amino acid utilisation. Thus, after some adjustments on the basis of his stable-isotope studies the requirements pattern was essentially that of 0·48 g tissue protein with lower values for lysine (71 %), threonine (71 %) and valine (83 %). In effect this metabolic model was similar to that of the 1973 FAO/WHO report, i.e., a factorial model of a dietary requirement derived from the ONL adjusted for a 70 % efficiency of utilisation.

The MIT amino acid requirements pattern was immediately very controversial in its theoretical origins (Millward, 1990). It was especially controversial because its basic idea, of a maintenance pattern based on tissue protein, meant that requirements for maintenance and growth were similar, contrary to the considerable body of work, largely animal studies, going back to Osborne & Mendel (1916), pointing to quite different amino acid patterns for maintenance and growth. However Young defended it vigorously both with argument (for example, Young, 1992; Young & El-Khoury, 1995) and with further experimental work (Marchini et al. 1993; Raguso et al. 1999; Young & Borgonha, 2000). He was quite happy to organise a session at the 1992 Federation meetings at Anaheim where the arguments could be aired in public in front of a packed auditorium containing the entire US nutrition academic leadership (Millward, 1994; Young, 1994). He remained convinced to the end that his stable-isotope studies in the USA and in India supported the MIT pattern especially in terms of the lysine requirement. He had the satisfaction that the Nutrition Board expert committee, in writing the Institute of Medicine (2002) US dietary reference intakes report, defined a pattern of amino acid requirements which was very similar to the MIT pattern, consequently restoring the importance of protein quality for human adults. Whatever the ultimate verdict on the MIT amino acid requirements pattern, there can be no doubt that the published experimental work aimed at its verification represents a body of knowledge which is invaluable in terms
of extending both our understanding of amino acid requirements and the methodology and technology necessary for the further work which undoubtedly will be required.

Other unresolved controversies

Adaptation

During the period after the Food & Agriculture Organization/World Health Organization (1973) report the problem of adaptation to low intakes emerged. It was discussed in the Food & Agriculture Organization/World Health Organization/United Nations University (1985) report and has been often revisited especially by Waterlow (1999) and Jackson (1999, 2000).

According to Scrimshaw (1996) the MIT view at the time, and subsequently, was that the individual adult’s requirement is genetically fixed over time with inter- but no intra-individual variability. In contrast to this, at UCLA Berkeley, a different view emerged based on the work of Sukhatme & Margen (1978). On the basis of a limited number of long-term N-balance studies, they argued for a different definition of the protein requirement. This allowed for intra-individual variability, i.e., adaptation, with a range of intakes within which protein homeostasis could be maintained.

The database on which Sukhatme and Margen developed their ideas was a small series of long-term N balances at fixed intakes, in which an individual’s daily N output showed serial correlations rather than random fluctuations. Young and Scrimshaw responded to the Sukhatme and Margen model by analysing their own long-term balance data for serial correlations and were unable to identify them (Rand et al. 1979), reporting only random variations in urinary output after correcting for long-term trends. Then, before the 1981 FAO/WHO/UN University meeting, Margen reported further data on the extent of serial correlations in long-term N balances in six subjects on a very low intake similar to the ONL. The intake was 0.36 g egg protein/kg per d, which they interpreted as indicating that this level of N intake was within the ‘adequate’ range (N Durkin, DA Ogar, SG Tilve and S Margen, unpublished results). However, most of the subjects lost weight during the study and since no measurements were made of body composition the extent of lean tissue loss was not indicated. Importantly there were no adverse physiological (work performance) or biochemical changes, unlike the MIT studies at a higher intake. Since these further studies of Margen presented to the 1981 FAO/WHO/UN University meeting had not been (and were never) published, with no other similar evidence for the cyclical patterns of N loss having been published (at least to my knowledge), and since the arguments of Sukhatme were particularly mathematically abstruse, they have had relatively little impact. In fact this particular argument is a complex one. The existence or not of the Sukhatme and Margen regulatory mechanism controlling N balance within a range of intakes is an interesting theoretical question. Yet, in one sense the key difference between MIT and Margen was the magnitude of the minimum intake at which N balance can be achieved at no obvious cost, i.e., 0.35 g protein/kg per d or >0.6 g protein/kg per d.

Young firmly rejected the idea of long-term adaptation to intakes much below 0.6 g/kg per d, arguing that ‘the limits of adaptation are bounded by the safe range of nutrient intakes, . . . are similar among healthy populations worldwide, . . . and are reached over a relatively short time frame’ (Young & Borgonha, 1998). However, Young has never rested on his well-deserved laurels and his approach to those who have challenged his view (for example, Hegsted, 2000), and myself (Millward, 2003; Millward & Jackson, 2004) has been to perform more studies. With Kurpad he has examined whether adaptive reductions in the lysine requirement occurs in chronically undernourished adult Indian men, showing that, on the contrary, the lysine requirement is increased (Kurpad et al. 2003c). Not prepared to leave it at that they then went on to show that this reflected the presence of intestinal parasites, which when treated, resulted in a value for the lysine requirement ‘in the range of 30 mg/kg per d or possibly less’ (Kurpad et al. 2003a). This means that the argument about the importance or relevance of adaptation will no doubt continue, not least by the very large number of researchers worldwide who Young trained and will no doubt continue to carry his flag.

Urea recycling and the de novo synthesis of lysine

Young’s interest in the use of stable isotopes to probe N metabolism and his close relationship with Waterlow’s unit in Jamaica meant that he knew of, and had great interest in, work by David Picou on urea recycling (Picou & Phillips, 1972). The idea that urea hydrolysis in the lower gut, liberating NH₃ to be utilised for bacterial amino acid synthesis, could also be of importance as a source of amino acids for the host organism had been suggested by evidence from several studies (Giordano et al. 1968; Furst, 1972; Sheng et al. 1977). However, the suggestion that the utilisation of urea-N in this way in Papua New Guinea highlanders could improve their N balance on low- and poor-quality-protein diets (Rikimaru et al. 1984) was something Young was very sceptical of. He was also sceptical of Jackson’s idea that urea salvage was an important control point in the regulation of N balance (see Jackson 1999, 1999) since this was contrary to his own work on urea kinetics (El-Khoury et al. 1996). Nevertheless as further evidence emerged from Malcolm Fuller’s work with pigs in Aberdeen (for example, Torraldana et al. 2003) and our own work (Millward et al. 2000), Young set up the appropriate experiments with Fuller and went on to show with [¹⁵N]urea and [¹⁵N]NH₃ that lysine deriving from intestinal microbial de novo synthesis did indeed appear in the circulating pool (Metges et al. 1999a,b).

The intriguing issue which remains is whether the de novo synthesis of lysine and other indispensable amino acids in the lower gut and absorption into the systemic pool is of any nutritional significance. Young believed that it wasn’t. The tragedy of his death deprives us of the person who undoubtedly could have resolved the issue one way or another.
References


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