Transport Kinetics of Amino Acids across the Resting Human Leg

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Abstract

Flux rates of amino acids were measured across the leg after an overnight fast in resting human volunteers. A balanced amino acid solution was, after a primed infusion, continuously infused for 2 h at each of three step-wise and increasing rates corresponding to 8.3, 16.7, 33.2 mg N/kg per h that were equivalent to 0.2, 0.4, 0.8 g N/kg per d. Flux of amino acids across the leg was compared with the flux of glucose, glycerol, lactate, free fatty acids, and oxygen. The size of the muscular tissue pool of amino acids was measured. Whole body amino acid oxidation was estimated by means of the continuous infusion of a ¹⁴C-labeled mixture of amino acids.

Arterial steady state levels were obtained for most amino acids within 30 to 45 min after the primed constant infusion. Leg flux of amino acids switched from a net efflux after an overnight fast to a balanced flux between infusion rates corresponding to 0.2–0.4 g N/kg per d. At 0.8 g N/kg per d essentially all amino acids showed uptake. The infusion of amino acids stimulated leg uptake of glucose and lactate production and decreased FFA release. Oxygen uptake and leg blood flow increased significantly with increased infusion of amino acids. There was significant variability in transport rate among individual amino acids. Branched chain amino acids showed rapid transport and methionine slow transport rate. Only small changes in the muscle tissue concentration of certain amino acids were registered after 6 h of amino acid infusion despite uptake for several hours. When amino acids were infused at a rate corresponding to 0.8 g N/kg per d, the leg uptake of amino acids was 6% and the simultaneous whole body oxidation of infused amino acids was ~ 10%. Net uptake of leucine across the leg per hour was 62% of the muscle pool of free leucine when amino acids were infused at a rate corresponding to 0.4 g N/kg per d. Multiple regression analysis showed that the arterial concentration of an amino acid was the most important factor for uptake, more so than insulin concentration and blood flow.

It is concluded that leg exchange of amino acids is large enough to rapidly change the pool size of the amino acids in skeletal muscle, if not counter-regulated by changes in rates of protein synthesis and degradation. Estimates of the capacity for protein synthesis and transfer RNA acceptor sites in muscles agree in order of magnitude with the net uptake of amino

acids at high infusion rates of amino acids. Therefore, measurements of the balance of tyrosine, phenylalanine, and particularly methionine at steady state may reflect net balance of proteins across skeletal muscles even in short-time experiments.

Introduction

We have recently reported that malnourished patients with and without cancer, who received nasogastric tube feeding, were in positive energy and nitrogen balance (1). However, even after a 2-wk period of enteral nutrition we found that most patients remained in negative amino acid balance across the leg even in the fed state, in spite of the fact that the balance of glucose and FFA across the leg indicated a positive energy balance (2). Such results emphasize a discrepancy between energy and protein balance in peripheral tissues during clinical nutrition as well as between whole body nitrogen balance and amino nitrogen balance in some peripheral compartments.

Although these studies were carried out on hospitalized patients (but not bedridden), such observations inevitably raise questions whether conventional nutritional therapy is insufficient to maintain and replenish lean body mass. Conversely, the results may indicate a more general deficiency in resynthesizing skeletal muscle proteins in malnourished patients which is perhaps related to the underlying disease. Irrespective, prior results show that such patients cannot improve lean body mass on a feeding therapy that is supposed by medical practice to be sufficient for resynthesis of lean tissues. Our observation may partly explain why nutritional support to hospitalized patients often fails to increase lean body mass, but not the replenishment of subcutaneous fat.

Modern artificial nutrition must be able to support the resynthesis of peripheral tissue proteins, particularly skeletal muscles. The most simple prerequisite of this is a positive amino acid balance. Therefore, the purpose of this study was to examine kinetics of amino acids and regulating factors for positive amino acid balance across resting peripheral tissues. For this purpose, the human leg has been used, which contains a considerable portion of the lean body mass in the body.

Methods

Patients. 11 healthy males without medication or a history of diabetes were investigated before surgery for uncomplicated inguinal hernia. They had all been weight stable for at least 1 yr before the examinations. Their body weight was 79±3 kg and height 179±3 cm. Body weight index was 1.02±0.04. Body potassium was 3,650±200 mmol and body potassium index 0.96±0.04. Plasma albumin was 40±3 g/liter, triceps skinfold 8.7±0.7 mm, and mid-arm circumference was 30.1±0.9 cm. All patients had normal body temperature. We regard these patients as healthy volunteers. They all had normal hemoglobin concentration (153±4 g/liter), sedimentation rate of erythrocytes (7±1 mm/h), normal liver function enzyme test, and normal serum electrolytes and creatinine. Informed consent was obtained from all individ-

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uals and the study protocol was approved by the Ethical Committee, University of Gothenburg.

Investigative procedures. All metabolic studies started in the morning after an overnight fast (12–14 h) with the patients at rest in the bed. Flux of amino acids, glucose, glycerol, lactate, and FFA were measured in relationship to oxygen uptake across the leg in the fasted state in eight of the volunteers. Catheters were inserted in the radial artery and in the femoral vein 1 h before the start of the investigation. The patency of the catheter was achieved by injections of saline after blood sampling. A balanced solution of amino acids (Vamin N; KabiVitrum, Stockholm, Sweden) was then infused as a primed constant infusion at rates corresponding to 0.2, 0.4, and 0.8 g N/kg per d for 2 h at each level. The infusion started with a monoexponentially primed infusion as described by DeFronzo (3) for insulin infusion to rapidly reach stable arterial levels of amino acids (Fig. 1). The arterio-venous (A-V)¹ difference of amino acids was measured during the last 15-min period of 2 h at each steady state level (4). Leg blood flow was measured at rest and at each infusion level by a strain-gauge plethysmograph (2). Flux is determined as the A-V difference multiplied by blood flow. Blood flow values (milliliters per minute per 100 ml) were converted to milliliters per minute per 100 g leg tissue assuming that leg tissue has a specific activity close to one (1, 2). This conversion is useful when comparing leg metabolism with whole body metabolism.

At the highest level of infusion (0.8 g N/kg per d) blood flow was also measured in five patients by the cardio-green dilution technique as described by Jorfeldt and Wahren (5). An additional three volunteers were subjected to a step-wise infusion with the amino acid solution labeled with a mixture of ¹⁴C-amino acids of known specific radioactivity. These individuals were infused according to the same protocol as used for the other eight volunteers but flux measurements were done only at the infusion rate of 0.8 g N/kg per d. Simultaneously, whole body energy expenditure was measured and expired ¹⁴CO₂ was trapped in potassium hydroxide (0.72 mol/liter) over 30 min as described elsewhere (6). The percentage oxidation of the infused labeled amino acid solution was measured from the appearance of radioactivity in the expired carbon dioxide and measured as described for the oxidation rate of tyrosine (6), assuming that 18% of the labeled carbon dioxide was retained in the bicarbonate pool. Muscle needle biopsies (quadriceps) were taken before and during infusions in five of the eight volunteers during step-wise increased infusion. The tissue content of free amino acids was measured after extraction of the free amino acids by methanol.

Biochemical measurements. Whole blood was immediately precipitated by perchloric acid at a final concentration of 1.4 mol/liter. Proteins were discarded after centrifugation. The supernatant was neutralized to pH 7.4 by the addition of 3.6 mol/liter potassium hydroxide. The amino acids were separated by means of high pressure liquid chromatography using a C₁₈ μm Bonda-pak column and precolumn derivatization of amino acids with *O*-phthalaldehyde in an automatic sample injector (7). Eight samples were run over night with a standard before and after each fourth sample. Parafluorophenylalanine (P-PHE) was used as the internal standard in each sample. P-PHE was added in a known concentration to each sample of whole blood before precipitation and correction with regard to all the amino acids in blood was done. The area under the curve of the amino acids was integrated using an automatic computer procedure (Waters Assoc., Div. of Millipore Corp., Milford, MA). According to a chromatogram of all standard amino acids of known concentration, the coefficient of variation for measurements each day of the standard amino acid (P-PHE) over six following weeks was 4.1±0.2% based on 47 chromatograms. When 10 standards were analyzed consecutively the variation coefficient was between 2 and 4% in repeated experiments.

Derivatization with *O*-phthalaldehyde does not give any derivate of secondary amines. Therefore, values of proline are not obtained. In

addition, cysteines are not derivatized under these conditions and are therefore not reported. Threonine and glycine do not separate with a linear gradient of acetonitrile at 25°C as used in this system and concentration of the two amino acids are given together. Values of tryptophan are not reported, since difficulties to integrate the peaks in some chromatograms were observed.

Blood gases were measured according to the manufacturer's recommendation in a Radiometer 2, Acid Base Laboratories, Copenhagen, Denmark. Insulin was measured in plasma with a radioimmunoassay (Phadebas) from Pharmacia Fine Chemicals, Uppsala, Sweden. Glucose, glycerol, and lactate were measured by kits from Boehringer Mannheim Diagnostics, Houston, TX. FFA were quantified according to Ho (8). All samples were kept at -70°C until analyses were performed.

Clinical tests and nutritional assessment. Hemoglobin concentration, sedimentation rate of blood cells, liver function enzyme tests, serum electrolytes, serum creatinine, and albumin were measured according to standard procedures in our hospital. Nutritional assessment included physical examination and measurement of body weight, height, serum albumin, triceps skinfold, and upper-arm circumference. Body potassium was measured in a total body counter. Body weight index is defined as actual body weight divided by reference weight obtained from healthy individuals living in the same area of Sweden as the study individuals (9). Total potassium index is actual body potassium divided by body potassium in reference patients.

Vamin N. The L-amino acid solution that was used for infusion contained in grams per liter: Gly, 2.1; Asp, 4.1; Glu, 9.0; Ala, 3.0; Arg, 3.3; cystein-cystin, 1.4; His, 2.4; Ile, 3.9; Leu, 5.3; Lys, 3.9; Met, 1.9; Phe, 5.5; Pro, 8.1; Ser, 7.5; Thr, 3.0; Trp, 1.0; Tyr, 0.5; and Val, 4.3.

Statistics. For independent variables the non-parametric Mann-Whitney U-test was used. Dependent variables were tested by means of Wilcoxon's test for paired differences. A parametric group comparison test was used to evaluate whether metabolic balance differs statistically from zero (10). *P* < 0.05 is regarded as a statistically significant difference. Regression lines were evaluated by means of analyses of variance and multiple regressions were calculated by means of a computer program (IDA 800; Luxor AB, Motala, Sweden).

Results

Arterial concentrations of the sum of all amino acids reached stable values between 30 and 45 min after the start of the primed constant infusion, as shown in Fig. 1. Most of the individual amino acids showed similar kinetics as the sum of all amino acids (results not shown). It was therefore assumed that all samples were taken at steady state levels between 115 and 120 min after the start of constant infusion.

Table I summarizes the mean arterial concentration of individual amino acids at fasting and at steady state levels during infusion. The concentration of most amino acids increased significantly when compared with the concentration at the preceding steady state level. Glutamine, arginine, asparagine, and ornithine did not show such a step-wise increase. Of these four amino acids, only arginine was infused.

Table II summarizes the A-V difference and flux rates across the leg of individual amino acids at fasting and during infusion at steady state. Alanine and glutamine accounted for 70% of the amino nitrogen that was released at fast. Glutamic acid was taken up and ornithine was at balance during the fast. All the other amino acids were released to a significant extent at fast. The infusion of 0.2 g N/kg per d and 0.4 g N/kg per d resulted in balances that were not statistically different from zero, while the infusion of 0.8 g N/kg per d gave a significant uptake of most amino acids. The balances of alanine and glutamine were negative at all infusion rates although glutamine

1. Abbreviations used in this paper: A-V, arterio-venous; P-PHE, parafluorophenylalanine.

Figure 1. Arterial concentrations (conc) of the sum of all amino acids in whole blood during step-wise increased infusion of a balanced amino acid solution corresponding to 0.2, 0.4, and 0.8 g N/kg per d. Results are from two individuals.

release was not statistically different from zero at the highest infusion. Asparagine showed a significant release up to 0.4 g N/kg per d and the balance of threonine–glycine was negative at 0.2 g N/kg per d and continued to show numerical negative values, although they were not statistically different from zero balance at 0.4 and 0.8 g N/kg per d.

Table III summarizes insulin levels, oxygen uptake, and blood flow in relationship to arterial concentrations and bal-

ances of glucose, lactate, glycerol, and FFA. Insulin concentrations and oxygen uptake increased significantly only at the highest infusion rate, while blood flow increased at 0.4 g N/kg per d compared with fasting levels. Arterial concentrations of glucose, glycerol, and FFA were unchanged, while lactate decreased significantly from 0.2 g N/kg per d of infusion and on. Glucose uptake increased continuously and reached a significantly increased uptake at 0.4 g N/kg per d. Glycerol flux was not statistically significantly altered at any amino acid infusion rate, although the A-V difference was at 0.4 and 0.8 g N/kg per d. FFA release showed a significant decrease between infusion of 0.4 and 0.8 g N/kg per d. Amino acid infusion resulted in increased lactate release by the leg, yet arterial lactate levels decreased. This suggests that the uptake of lactate by other tissues, most likely the liver, was increased in response to amino acid infusion.

The muscle content of free amino acids at the various infusion rates is shown in Table IV. The muscle free pool of all amino acids was unchanged in spite of 6 h infusion of amino acids. However, the pool of valine, leucine, isoleucine, phenylalanine, and methionine increased significantly, while that of taurine decreased in five of six patients. The pool of methionine increased, although the change in flux rate across the leg was not statistically significant. Considering the net efflux of amino acids in the fasting state, the time required to decrease the pool size of individual amino acids to zero levels was calculated, provided that the pool was not replenished from net degradation of proteins. Such calculations are shown in Table V. The phenylalanine and methionine pool have the shortest turnover time, in the range of 3 h, while glutamine had 47 h and serine 123 h turnover.

In Fig. 2 the relationship between the arterial concentration and the balance of branched chain amino acids are shown. The branched chain amino acids together represented the major component of amino nitrogen uptake. The flux of individual amino acids was, therefore, compared with that of the branched chain amino acids. The slopes for uptake of individual amino acids are shown in Table VI. The slopes were calculated by fitting the data to linear regressions which were tested by analyses of variance. The coefficient of the slopes were then compared statistically as a measure of uptake rate versus the arterial concentration of the amino acid. Lysine and serine had a significantly more rapid uptake than the branched chain amino acids together. Tyrosine, phenylalanine, methionine, histidine, and threonine–glycine had slower uptake. The arterial concentration for zero balance of the individual amino acids was calculated from the regressions in Table VI. These results were for (in micromoles per liter) branched chain, 543; Tyr, 130; Phe, 34; Met, 57; Ser, 133; His, 134; Thr-Gly, 690; Leu, 229; and for sum of all amino acids, 4,123 μmol/liter. When these values are compared with fasting values of amino acids, one can obtain an estimate of the magnitude of arterial concentrations of individual amino acids that must be increased in order to switch from negative balance to net uptake. These indexes (times fasting levels) were for: branched chain, 1.3; Tyr, 1.5; Phe, 2.2; Met, 3.6; Ser, 1.0; His, 1.6; and Thr-Gly, 1.5; the sum of all amino acids was 1.3.

The percentage uptake of administered amino acids in peripheral tissues was calculated by dividing the infusion rate of individual amino acids (micromoles per kilogram per minute) by the flux across the leg (micromoles per kilogram per minute) at the highest infusion rate (0.8 g N/kg per d) for all the

Table I. Arterial Concentrations of Amino Acids

Amino acids	0	0.2	0.4	0.8
Ala	273±13	333±14*	376±17†	474±20*
Gln	530±32	562±37	479±68	563±27
Glu	197±16	249±12*	332±30†	565±39*
Val	227±12	325±14*	505±26*	841±32*
Leu	125±9	205±10*	343±25*	575±25*
Ile	53±2	112±4*	208±15*	370±20*
Tyr	89±5	123±9†	163±12†	198±13*
Phe	43±2	97±3*	172±15*	298±12*
Met	16±1	33±1*	53±4*	84±4*
Asp	118±10	136±8†	162±18	217±14*
Asn	125±18	123±18	130±19	115±25
Ser	136±5	214±11*	314±14*	466±28*
His	85±7	107±8*	128±10†	159±15†
Thr-Gly	452±13	535±18*	608±27†	716±34†
Arg	55±1	60±2	70±6	84±7
Tau	267±15	310±25†	352±34†	408±28
Orn	113±13	119±11	132±5	167±12*
Lys	180±8	231±13*	274±18†	332±17†
Total	3,088±99	3,874±134*	4,799±259*	6,639±265*

In whole blood (micromoles per liter) after an overnight fast (0), and at steady state during infusion of 0.2, 0.4, and 0.8 g N/kg per d of an amino acid solution. There were eight subjects. Mean±SE.

* $P < 0.01$ vs. the preceding rate of infusion.

† $P < 0.05$ vs. the preceding rate of infusion.

Table II. A-V Differences* and Flux (FL)[‡] across the Leg after an Overnight Fast (0) and at Steady State[§]

	0	0.2	0.4	0.8
Ala A-V	-63±10	-68±6	-92±11	-70±16
FL	-163±21	-216±33	-423±82	-300±96 [†]
Gln A-V	-53±22 [†]	-40±22	-163±49	-51±33
FL	-96±63	-99±85	-547±112	-91±156
Glu A-V	29±5	70±6	113±28	131±23
FL	85±21	238±49	477±137	526±123
Val A-V	-13±3	12±3	32±17	76±13
FL	-30±5	40±14 [†]	130±77	305±67
Leu A-V	-9±5	3±2	29±15	53±10
FL	-28±14 [†]	16±8	117±65	216±56
Iso A-V	-7±1	9±1	24±10 [†]	42±7
FL	-19±5	31±6	107±44 [†]	173±41
Tyr A-V	-4±6	-3±3	-3±6	10±4 [†]
FL	-8±14	-6±7	10±26	37±21
Phe A-V	-6±1	3±1 [†]	10±8	20±5
FL	-21±7 [†]	11±4 [†]	36±35	79±28 [†]
Met A-V	-4±0.3	-1±1	0±3	2±1
FL	-10±1	-3±3	-4±12	6±5
Asp A-V	-6±3	6±6	23±8 [†]	33±9
FL	-14±10	31±18	89±42 [†]	121±45
Asn A-V	-11±3	-9±3 [†]	-30±8	-14±9
FL	-29±8	-28±9 [†]	-147±43	-58±48
Ser A-V	-5±6	22±4	43±14 [†]	44±12
FL	-4±13	64±12	194±78 [†]	170±57 [†]
His A-V	-8±2	-3±1 [†]	0±4	2±5
FL	-19±4	-6±5	-2±15	2±19
Thr-Gly A-V	-37±10	-15±4	-10±19	-2±13
FL	-97±20	-44±18 [†]	-63±69	-24±63
Arg A-V	-6±1 [†]	-4±2	-3±3	2±2
FL	-16±5 [†]	-8±5	-11±9	3±11
Tau A-V	-13±4	-3±4	-16±7 [†]	1±3
FL	-32±7	-3±14	-84±32 [†]	-3±15
Orn A-V	-3±4	1±4	8±7	6±6
FL	0±10	8±13	44±31	29±27
Lys A-V	-17±7 [†]	-13±9	18±10	16±8 [†]
FL	-29±20	-35±30	78±60	63±37
Branched A-V	-28±7	24±4	85±42 [†]	171±29
FL	-33±27	62±20 [†]	516±178 [†]	769±152
Total A-V	-232±67	-28±43	14±140	301±96 [†]
FL	-522±124	9±191	3±593	1,136±448 [†]

* Micromoles per liter. [‡] Nanomoles per minute per 100 g. [§] During infusion of 0.2, 0.4, and 0.8 g N/kg per d of an amino acid solution. There were eight subjects. Mean±SE. ^{||} *P* < 0.01 vs. fasted or different from zero balance at fasting. [†] *P* < 0.05 vs. fasted or different from zero balance at fasting.

subjects. The figures were for individual amino acids in percent of infusion rate: Glu, 146; Val, 139; Leu, 91; Ile, 97; Tyr, 207; Phe, 41; Met, 7; Asp, 66; Ser, 41; His, < 2; Thr-Gly, < 1;

and Lys, 39. The results show that ~ 35% of the infused amino acid nitrogen were taken up in peripheral tissues and that one leg disposal was ~ 6% of the infused amino acids.

Table III. Arterial Concentration (conc),* A-V Differences,[‡] and Flux Across the Leg[§]

	0	0.2	0.4	0.8
Glucose conc	6,039±105	5,966±69	5,900±79	6,466±208
Flux	371±138	751±275	1,200±245	1,814±698 [†]
A-V	144±54	209±67	274±36	366±92 [†]
Lactate conc	779±84	597±61	500±30	597±67
Flux	-160±129	-685±103	-697±123	260±231
A-V	-84±45	-206±28	-181±30 [†]	-102±48
Glycerol conc	65±9	74±6	80±5	70±6
Flux	-136±32	-139±33	-159±67	-66±37
A-V	-45±8	-37±5	-30±9	-21±9
FFA conc	513±34	474±29	533±37	452±55
Flux	-298±77	-119±70	158±163	8±150
A-V	-122±38	-39±17 [†]	+30±32	-6±29
Insulin conc	16±3	18±3	18±2	34±5
Oxygen flux	8.5±1.6	11.0±1.9	14.1±3.9	15.5±3.1
Blood flow	3.06±0.58	3.34±0.57	4.62±0.57	4.62±0.78

* Micromoles per liter. [‡] Micromoles per liter. [§] In relationship to oxygen uptake and leg blood flow after an overnight fast (0) and at steady state during infusion of 0.2, 0.4, and 0.8 g N/kg per d of an amino acid solution. There were eight subjects. Mean ± SE. ^{||} *P* < 0.01 vs. fasted state. [†] *P* < 0.05 vs. fasted state.

Table IV. Muscle Content of Amino Acids*

	0	0.4	0.8
Ala	4.64±0.41	4.40±0.39	5.01±0.20
Gln	22.98±3.07	20.33±2.40	22.29±2.77
Glu	5.29±0.89	4.06±0.74	4.01±0.30
Val	0.74±0.07	1.04±0.07 [‡]	1.74±0.12 [‡]
Leu	0.57±0.05	0.71±0.15 [‡]	1.30±0.07 [‡]
Ile	0.29±0.04	0.49±0.04 [‡]	0.81±0.03 [‡]
Phe	0.18±0.03	0.41±0.04 [‡]	0.74±0.04 [‡]
Met	0.10±0.02	0.15±0.02 [‡]	0.24±0.02 [‡]
Asp	1.08±0.27	1.44±0.53	1.15±0.50
Aspn	0.82±0.13	0.68±0.14	0.54±0.16
Ser	1.47±0.32	1.67±0.22	2.02±0.17
His	0.61±0.05	0.62±0.11	0.60±0.09
Thr-Gly	4.25±0.36	3.96±0.32	4.11±0.38
Arg-Tyr	19.12±1.53	15.95±2.17	16.70±4.64
Tau	40.98±2.85	33.32±2.14 [‡]	36.16±2.49
Orn	0.20±0.05	0.18±0.08	0.18±0.10
Lys	1.03±0.25	0.86±0.22	1.01±0.44
Total	104.4±7.4	89.71±8.01	98.49±10.27
Muscle dry weight (mg)	9.26±1.85	6.79±1.97	8.37±3.30
<i>n</i>	5	5	4

n, Number of patients.

* Nanomoles per milligram lipid free dry weight. After an overnight fast and after 2 h of infusion of amino acids at 0.2 g N/kg per d plus 2 h at 0.4 g N/kg per d and after further 2 h infusion at 0.8 g N/kg per d, altogether equalling 6 h of infusion.

[‡] *P* < 0.01 vs. fasted state.

[§] *P* < 0.05 vs. fasted state.

Values of amino acid uptake across the leg were compared with the estimated whole body oxidation of infused amino acids during a constant infusion of radioactive amino acids in three of the subjects (Table VII). The results show that the immediate oxidation rate of infused amino acid carbons corresponded to ~ 7–10% of the infused amino acids.

Table VIII shows multiple regression analysis for phenylalanine balance (A-V balance) versus arterial concentration of phenylalanine, arterial insulin levels, and leg blood flow. The

Table V. Calculated Turnover Time for the Skeletal Muscle Pool of Individual and Soluble Amino Acids*

	Hours
Ala	9.5
Gln	47
Val	8
Leu	7
Ile	5
Phe	3
Met	3
Asp	26
Asn	9
Ser	123
His	11
Thr-Gly	15
Tau	427
Lys	12

* The tissue pool (nanomoles per gram) after an overnight fast is divided by the efflux rate (nanomoles per minute per gram) of the amino acid after an overnight fast.

Figure 2. The relationship between arterial concentration and leg exchange of branched chain amino acids. Pooled observations from eight individuals investigated in fasting state and during steady state infusion at 0.2, 0.4, and 0.8 g N/kg per d of the amino acid solution. The regression is given in Table VI.

results demonstrate that only the arterial level of phenylalanine had a positive and significant influence on the variance of phenylalanine balance, whereas insulin and blood flow in themselves did not significantly explain the variability in phenylalanine balance. The two together did, however, contribute significantly to the multifactorial analysis. Similar results were obtained for tyrosine and methionine. In a single factor analysis, insulin correlated to the balance of phenylalanine and many other amino acids at the level of $r = 0.40-0.45$, $P < 0.05$, but blood flow did not. The transport capacity of amino acids was not limiting for uptake, which was evaluated by calculations of blood clearance of the amino acids such as branched chain, methionine, tyrosine, and phenylalanine across the leg (results not shown). That amino acid uptake was proportional to the arterial concentration can also be seen in Fig. 2.

Discussion

This study has evaluated the net balance of blood amino acids across the leg in human adults. Flux of amino acids across the

resting leg after an overnight fast was compared to flux at steady state during the constant infusion of increasing quantities of amino acids. Most of the blood amino acids reached steady state levels within 30 to 45 min after a primed constant infusion of an amino acid solution. Therefore, we believe that a time period of 100 min was enough to reach steady state in the plasma and blood compartment (4). We have assumed that interstitial amino acids are in rapid equilibrium with the intravascular pool of amino acids.

The ultimate goal of our experiments is to obtain a clinical model to evaluate regional net protein balance. It is obvious that flux measurements of amino acids across the leg do not necessarily give information on protein balance in acute experiments (11). Rapid changes in flux may in theory only reflect changes in the intracellular pool of free amino acids (12). However, seen over time, net balance of amino acids must give predictive information with regard to net protein balance. In a condition of continuous efflux of amino acids, the tissue pool must soon approach a state where amino acid exchanges reflect protein balance. This is illustrated by calculations of the fractional replacement time for the tissue pool of individual amino acids (Table V). Accounting for the efflux rate of methionine from the leg after an overnight fast, it can be calculated that the intracellular pool of methionine should be near zero value within 3 h if not replenished. In this sense, it can be claimed that the intracellular availability of certain amino acids may under some conditions be limiting for protein synthesis, although it is generally recognized that transport rates or the transport capacity of amino acids in themselves is never limiting for protein synthesis. In this study it was confirmed that the transport capacity was not limiting for amino acid uptake. It has been reported that the tRNA pool of amino acids is fully charged, even in starvation (13). The complete charging of tRNA in substrate-deficient conditions and Michaelis constant (K_m) values in the range of 2–6 μM for amino acid acylation of tRNA have been taken to indicate that amino acids are not limiting for protein synthesis (14).

We have reported that the maximum rate of leucine incorporation into proteins is $\sim 50-100$ nmol/min per 100 g in incubated human muscle tissue (15–19). This value can be compared with the maximum acceptor site (pool-size) for

Table VI. Regressions Calculated from the Arterial Levels of Amino Acids versus Net Balance of Amino Acids at each Level of Steady State*

Amino acids	Regression	$P <$	r	Coeff of the slope	SD coeff	Total coeff	n	Vs. branched chain amino acids
Branched chain	$y = 0.6154, x - 334$	0.01	0.76	0.6154	0.0950	6.4752	32	—
Tyr	$y = 0.4934, x - 86$	0.01	0.51	0.4934	0.1507	3.2753	32	$2 P < 0.001$
Phe	$y = 0.4479, x - 42$	0.01	0.63	0.4479	0.1007	4.4461	32	$2 P < 0.001$
Met	$y = 0.2980, x - 17$	0.01	0.45	0.2980	0.1092	2.7283	32	$2 P < 0.001$
Ser	$y = 0.7124, x - 95$	0.01	0.61	0.7124	0.1674	4.2558	32	$2 P < 0.01$
His	$y = 0.4268, x - 57$	0.01	0.48	0.4268	0.1435	2.9733	32	$2 P < 0.001$
Thr + Gly	$y = 0.5119, x - 353$	0.05	0.45	0.5119	0.1837	2.7866	32	$2 P < 0.01$
Lys	$y = 0.7591, x - 174$	0.05	0.44	0.7591	0.2830	2.6820	32	$2 P < 0.02$
Total	$y = 0.5527, x - 2,312$	0.05	0.71	0.5527	0.1009	5.4794	32	$2 P < 0.025$

Significance of regressions were tested by analysis of variance. * At fast, at infusion of 0.2, 0.4, and 0.8 g N/kg per d. Total is the sum of all amino acids.

Table VII. Whole Body Gas Exchange and Amino Acid Metabolism*

Patients	O ₂ uptake	CO ₂ production	RQ	Infusion rate of AA	Uptake across the leg of AA	Oxidation of infused AA
	$\mu\text{mol} \times \text{min}^{-1}$	$\mu\text{mol} \times \text{min}^{-1}$	$\mu\text{mol} \times \text{min}^{-1}$	$\mu\text{mol} \times \text{min}^{-1}$	$\mu\text{mol} \times \text{min}^{-1}$	$\mu\text{mol} \times \text{min}^{-1}$
S.F.	12,300	9,712	0.79	3,287	248	321
J.B.	13,709	11,278	0.84	3,044	103	207
S.S.	12,803	9,698	0.75	3,079	184	309

AA, amino acid. * In three subjects who were infused with a mixture of ¹⁴C-amino acids to estimate the percentage of infused amino acids being oxidized in relation to flux. For details see Methods.

highly purified leucyl-tRNA, which is in the range of 100–120 nmol/100 g tissue weight, whereas the maximum capacity for aminoacyl-tRNA synthetase activity (the enzyme) is ~ 100-fold higher (20–22). The above values shall be compared with leucine uptake across the leg, which was 216±25 nmol/min per 100 g at the highest infusion rate (0.8 g N/kg per d, Table II). Such values, thus, indicate that flux rate of amino acids in vivo is within the order of magnitude for previously estimated capacity of protein synthesis in vitro. Low K_m values for tRNA acylation may be important for creating a necessary downhill gradient to promote the intracellular flux of amino acids towards protein synthesis. The intracellular concentration of phenylalanine and methionine will decline within 2 to 4 h (Table V) unless altered protein synthesis and degradation maintain the concentration above K_m for tRNA charging. If the intracellular pool is not refilled enough at regular intervals, protein synthesis must be supported by amino acids made available by proteolysis, or protein synthesis will be rapidly decreased. A negative protein balance may thus exist at different levels of protein turnover (23–25). This theory does, however, not explain why there should be a concentration threshold for amino acids to be taken up in peripheral tissues. A pull phenomenon due to a down-hill gradient for amino acid uptake across muscle membranes without a threshold phenomenon across the cell membrane could divert amino acids away from vital organs in conditions with low postabsorptive

plasma levels of amino acids. If so, the skeletal muscles could be in serious competition with other tissues and their protein synthesis just by means of the size of the skeletal muscle pool. A threshold gradient across the cell membrane may then prevent this situation from occurring and allow amino acids to be taken up in muscles only in the fed situation.

A lack of uptake or balance of amino acids may be manyfold, such as substrate availability, hormonal milieu, and low physical activity, but the extracellular concentration of amino acids may also be a critical factor (26). We have no reason to believe that lack of energy was a factor of importance in our present experiments. The individuals were well nourished and assumed to have normal glycogen stores. Besides, the infusion of amino acids increased oxygen consumption and gave a net uptake of glucose concomitant with increased efflux of lactate. Previous studies have also demonstrated a consistently negative amino acid balance in traumatized and septic patients on intravenous nutrition (27), and in patients on enteral nutrition (2). It is, however, obvious that it is possible to obtain a net uptake of amino acids across skeletal muscle in normal individuals simply by increasing the infusion rate of amino acids, which leads to increased arterial levels (Table III [28, 29]).

It was recently reported that 80% of intravenously infused amino acids are taken up across the splanchnic bed when amino acids were given at a constant rate of 0.2 g amino acids/kg per h (30). In the present study, the peripheral tissue uptake of amino acids was estimated to be ~ 35% at our highest infusion rate. The leg disposal of amino acids was 6–8% (30), which agrees with 6% in our experiments (Table VII). The fate of such amino acids may either be degradation, protein synthesis, or intracellular accumulation. The results in the present study show that ~ 10% of the infused amino acids were immediately oxidized when amino acids were infused at 0.8 g N/kg per d. This oxidation percentage of infused amino acids is considerably less than the percentage of overall whole body protein oxidation, which is ~ 30% as measured by indirect calorimetry. This discrepancy emphasizes that several amino acid pools may not have equilibrated within 6 h of infusion in spite of steady state in the intravascular compartment. In the skeletal muscle tissue we measured that the intracellular concentration of the branched chain, phenylalanine, and methionine increased significantly already at the lower infusion rate. The precise fate of the amino acids taken up cannot be decided in the present study. Future experimentation may answer the questions of whether such amino acids end up in protein, to what extent individual amino acids are

Table VIII. Multiple Linear Regression Analysis*

	y variable (AV-phe)		
	x variable (A-phe)	Insulin	Blood flow
Intercept	-5.7344		
Coefficients	0.1287	-0.1092	-1.3560
t value of coefficients	4.5648	-0.4715	-1.3030
SD of coefficients	0.0282	-0.2316	1.0406

$R = 0.78$ is significantly different from the simple correlation coefficient $r = 0.61$ ($P < 0.05$). The simple regression between AV-phe and A-phe satisfied the equation $y = 0.0963x - 8.9000$ ($n = 35$, $P < 0.01$). The multiple regression coefficient for AV-phe versus A-phe, insulin, and blood flow. $R = 0.78$. $N = 26$. $P < 0.01$.

* Dependent variable is A-V difference of phenylalanine (AV-phe). Independent variables are arterial concentrations of phenylalanine (A-phe), arterial plasma insulin, and leg blood flow.

oxidized, or whether they are stored in a labile pool intracellularly ready to be mobilized when the amino acid infusion is interrupted.

Multiple regression analysis was performed to evaluate the source of variation in A-V balance of phenylalanine. The arterial concentration of phenylalanine, insulin, and leg blood flow were selected on a theoretical basis as possible factors regulating amino acid balance. The arterial concentration of phenylalanine was the most important factor for phenylalanine uptake within the observed range for insulin levels up to 58 μ U/ml and blood flow values within the normal range, from 2.0 to 2.5 ml/min per 100 g up to 9 ml/min per 100 g. Only insulin, and not blood flow, correlated to the A-V differences of phenylalanine in a simple correlation analysis, although the two together contributed significantly to explain the variation in phenylalanine balance in a multifactorial analysis (Table VIII). Insulin and blood flow were not interrelated, but insulin and arterial levels of amino acids were ($P < 0.05$). It has been reported that amino acid uptake by splanchnic and leg tissues is relatively independent of insulin levels (31), although this conclusion was recently questioned (32). 38% of the variation of phenylalanine balance was thus explained by the arterial concentration alone, while 61% was explained by the three together ($P < 0.05$). Multiple regressions for tyrosine and methionine gave similar results. The addition of the A-V-differences of the branched chain amino acids increased the multiple regression coefficient to 0.91. This means that 83% of the uptake of phenylalanine was related to the uptake of the four factors together. This may be interpreted to indicate that the transport of several amino acids is dependent to some degree on the balance of the branched chain.

To summarize, this study provides us with information on amino acid transport kinetics during step-wise increased infusion of a complete amino acid solution used in intravenous nutrition. By the primed constant infusion technique, it was possible to obtain constant arterial concentrations of most of the amino acids. This fact made it possible to evaluate flux of amino acids and to translate such results into transport kinetics for individual amino acids. Estimates of the maximum capacities for protein synthesis in human muscles according to our previous studies and values of tRNA acceptor sites taken from the literature agree in order of magnitude with net uptake of amino acids in vivo in the fed situation as presented in this study. The transport capacity of amino acids was not a limiting step for amino acid uptake across the leg. Future studies are aimed to relate flux of amino acids directly to protein synthesis in vivo.

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