Evidence that glutamine modulates respiratory burst in stressed rat polymorphonuclear cells through its metabolism into arginine

Christophe Moinard1,2*, Florence Caldeﬁ-Chezet1, Stephane Walrand1,2, Marie-Paule Vasson1 and Luc Cynober2,3

1Laboratoire de Biochimie, Biologie Moléculaire et Nutrition EA 2416 and Centre de Recherche en Nutrition Humaine, Faculté de Pharmacie, Clermont-Ferrand, France
2Laboratoire de Biologie de la Nutrition, Faculté de Pharmacie, 4 avenue de l’Observatoire, 75270 Paris Cedex 06, France
3Laboratoire de Biochimie, Hotel-Dieu, AP-HP and Inserm U341, France

(Received 29 October 2001 – Revised 10 July 2002 – Accepted 11 August 2002)

Glutamine (GLN) and arginine (ARG) are recognized for their ability to modulate immune cell function. However, the metabolic pathways involved in their action remain unclear. It was recently shown that GLN- or ARG-enriched diets increased radical oxygen species (ROS) production by neutrophils from stressed rats. Since these two amino acids have a tied metabolism, we hypothesized that conversion between GLN and ARG (and its active metabolites NO† and polyamines) might be involved. To test this hypothesis male Sprague–Dawley rats (n = 117) were randomized into thirteen groups: rats in eleven groups were rendered catabolic by dexamethasone injection (1·5 mg/kg per d for 5 d) and 6·8 mmol either GLN, ARG or non-essential amino acids (NEAA; glycine, alanine and histidine)/kg per d were given by the enteral route; one group was pair-fed to the treated groups. The regimens of all the groups were rendered isonitrogenous by the addition of NEAA. The last group was not treated and was fed ad libitum. For each supplementation three subgroups were formed, each of which received a speciﬁc inhibitor: methionine sulfoximine (inhibitor of GLN synthase; 100 mg/kg per d), S-methylthiourea (inhibitor of inducible NO† synthase (iNOS); 50 mg/kg per d) and difluoromethylornithine (inhibitor of ornithine decarboxylase (ODC); 50 mg/kg per d). Oxidative metabolism, intracellular H2O2, and extracellular O2† production were measured in unstimulated and phorbol myristate acetate-stimulated polymorphonuclear neutrophils. GLN- and ARG-enriched diets increased respiratory burst by neutrophils (oxidative metabolism of 152 (SEM 24) and 138 (SEM 45) v. 57 (SEM 18) mV for GLN-, ARG- and NEAA-enriched diets respectively, P < 0.05). In vivo inhibition of iNOS or ODC decreased ROS production induced by GLN and ARG. In vivo inhibition of GLN synthase did not modify the effect of ARG on ROS production. In conclusion, GLN and ARG modulate ROS production in neutrophils from stressed rats by the same pathway involving polyamine and NO† synthesis.

Glutamine: Arginine: Polymorphonuclear cells: Polyamines: Nitric oxide: Respiratory burst

Response to injury is characterized by a hypermetabolic state and an alteration of protein metabolism (Jeevanandam, 1995). In particular, injury is also associated with marked changes in the concentrations of free amino acids, especially glutamine (GLN) and arginine (ARG), both in plasma and muscles (Cynober, 1989; Ardawi & Jamal, 1990; Ardawi & Majzoub, 1991). One current view is that the decrease in GLN and ARG availability is partly responsible for the dysimmunity often observed in stress (Newsholme et al. 1988; Brittenden et al. 1994; Pastores et al. 1994). Under these conditions, GLN- and ARG-enriched diets may have pharmacological effects on immunity. Many studies (for a review, see De Bandt & Cynober, 1998) have shown the ability of these two

Abbreviations: AL, ad libitum group; ARG, arginine; DEX, dexamethasone; DFMO, α-difluoromethylornithine; GLN, glutamine; MSO, methionine sulfoximine; NEAA, non-essential amino acids; PF, pair-fed group; PMN, polymorphonuclear neutrophils; ROS, reactive oxygen species; SMT, S-methylthiourea.

* Corresponding author: Dr C. Moinard, fax + 33 1 53 73 97 56, email christophe.moinard@nutrition-paris5.org
amino acids to modulate immune functions under physiological or pathological conditions. They are known to improve T-cell immunity (Alexander, 1995; Robinson et al. 1999), to stimulate macrophage functions (Calder & Yaqoob, 1999; Wu et al. 2000) and polymorphonuclear neutrophil (PMN) bactericidal power (Moffat et al. 1996; Furukawa et al. 1997, 2000; Holm et al. 1999; Moinard et al. 1999). However, the mechanisms involved in these actions are still unclear. In a recent study (Moinard et al. 1999) we showed the ability of an enterally administered diet supplemented with GLN and ARG to stimulate H₂O₂ production by PMN. However, the mechanisms involved were not elucidated. We (Moinard et al. 2000) and others (Murphy & Newsholme, 1998) have shown that GLN may be an ARG precursor for NO* synthesis in immune cells. These data suggest that these two amino acids have a tied metabolism and so their immunomodulatory effects may be related to a common pathway. We hypothesized that GLN may act via ARG or that ARG may act through GLN neosynthesis.

The aim of the present study was therefore to determine the metabolic pathways involved in GLN and ARG action on radical oxygen species (ROS) production by PMN from stressed rats. The catabolic state was obtained by daily intraperitoneal injection of glucocorticoids (i.e. dexamethasone; DEX). This model has been demonstrated to lead to a highly reproducible catabolic state in rats, including anorexia, protein breakdown and negative N balance (Odedra et al. 1983; Ardawi & Jamal, 1990; Parry-Billings et al. 1990; Minet-Quinard et al. 2000). In addition, glucocorticoids have immunosuppressive effects; they are responsible for changes in the distribution of leucocytes observed during stress (Shijo et al. 1998). These changes may have a significant impact on the functioning and effectiveness of the immune response (Shijo et al. 1998).

We evaluated the enhancing effect of GLN and ARG on respiratory burst using three different methods (chemiluminescence, flow cytometry and ferricytochrome reduction) in glucocorticoid-treated rats. We also used metabolic inhibitors in GLN- and ARG-supplemented stressed rats in order to block different metabolic pathways at the whole body level (Fig. 1): methionine sulfoximine (MSO), a specific inhibitor of GLN synthase (Heeneman & Deutz, 1993), to determine whether ARG acts through neosynthesis of GLN; S-methylthiourea (SMT), a specific inhibitor of inducible NO* synthase (Ruetten & Thiemermann, 1996), to determine whether GLN and ARG act by a process dependent on NO* synthesis (the active metabolite of ARG); α-difluoromethylornithine (DFMO), a specific inhibitor of ornithine decarboxylase (Nsi-Envo et al. 1996) to determine whether GLN and ARG act via polyamine synthesis (deriving from ARG).

Materials and methods

Chemicals

All chemicals were purchased from Sigma (St-Quentin-Fallavier, France) except for 2′,7′-dichlorofluorescin diacetate (DCFH-DA; Acros, Noisy-le-Grand, France), Plasma-gel (Belon, Neuilly-sur-Seine, France), and DEX (Soludecedron®; Merck Sharp & Dohm, Riom, France).

Animals and study design

A total of 117 male Sprague–Dawley rats (3 months old) were used (Iffa Credo, L’Arbresle, France). After their arrival at our animal facility the rats were maintained on a standard chow diet (A04: 170 g proteins, 30 g fat, 590 g carbohydrates, and 210 g water/kg, fibre, vitamins and minerals; Usine d’Alimentation Rationnelle, Villemoisson-sur-Orge, France) and received water ad libitum. They were
kept in individual cages in a controlled environment (constant temperature of 24°C) and light cycle (lights off 8.00–20.00 hours, lights on 20.00–8.00 hours). After this acclimatization period, the rats were randomized into thirteen groups (nine per group) as shown in Table 1.

For the next 5 d (day 0 to day 4) all the rats (except those that were pair-fed) had free access to the chow diet as described above. DEX-treated groups (i.e. groups 3 to 13, see Table 1) received a daily intraperitoneal injection of DEX (1.5 mg/kg per d) for 5 d (day 0 to day 4) as previously described (Minet-Quinard et al. 2000). The pair-fed group (PF) was pair-fed to DEX groups and received a daily isovolumic intraperitoneal injection of NaCl (9 g/l). The rationale for the study of a PF was the fact that DEX induces anorexia (Minet-Quinard et al. 2000); studying a PF thus isolates the effect of the treatment itself from the effects induced by marked anorexia. In addition to their oral chow diet the supplemented groups received enteral supplementation of GLN or ARG (6.8 mmol/kg per d) (day 0 to day 4) by oral administration as previously (Lasnier et al. 1996). The dose of 6.8 mmol/kg per d corresponds to 1.00 g GLN/kg per d and 2.19 g ARG/kg per d. These doses are in the range of those used in the literature (Barbul, 1995; Fürst & Stehle, 1995; Farges et al. 1999; Moinard et al. 1999) and are an extrapolation of dosages used in human subjects allowing for the high metabolic rate of rats (Cynober, 1989). The enteral route was chosen to provide supplements, rather than mixing them with chow (Le Bricon et al. 1995), to be certain that rats suffering from anorexia received the targeted amount of pharmaceutrients. The enteral supplementation of all the groups was rendered isonitrogenous (up to 0.5 g N/kg per d) by addition of a mixture of non-essential amino acids (NEAA; glycine, alanine, histidine in equimolar amounts). DEX-NEAA and PF groups received only the NEAA mixture (0.5 g N/kg per d) as a supplement to their regimen. In our experimental conditions, the NEAA mixture exerted neither pharmacological nor adverse effects on metabolic pathways (Minet-Quinard et al. 1999). MSO-, SMT- and DFMO-treated groups received MSO (100 mg/kg per d), SMT (50 mg/kg per d) and DFMO (50 mg/kg per d) respectively by the enteral route for 6 d (day –1 to day 4). The use of NEAA groups allowed us to determine whether the metabolic inhibitors are able to modify per se the parameters measured. We did not determine the effect of MSO in the GLN-supplemented group since GLN synthase is not involved in GLN catabolism. Dosages of inhibitors were selected on the basis of literature values (Nsi-Emvo et al. 1996; Ruetten & Thiemermann, 1996; Minet et al. 1997). Concerning the effect of inhibitors per se, GLN synthase activity in muscle (extensor digitorum longus) was determined using a colorimetric assay (Minet et al. 1997) and we previously demonstrated that GLN synthase activity was not detectable in muscles from MSO-treated rats (Moinard et al. 2002). In our experimental conditions, SMT abolished NO secretion in stimulated macrophages (Moinard et al. 2000), which assess that SMT block NO synthase in vivo. In the case of DFMO, ornithine decarboxylase was not measured since this enzyme has a very short half-life and 24 h had elapsed since the last injection. However, it is well known that DFMO blocks ornithine decarboxylase activity in our experimental conditions (Messina et al. 1992; Morgan, 1994; Raul et al. 1995; Walters et al. 1998).

The ad libitum group (AL) received no treatment, and was fed ad libitum.

Animal care and experimentation complied with the rules of our institution, and two of us (L.C. and M.-P.V.) are authorized by the French Ministry of Agriculture and Forestry to use this type of experimental model. After 24 h following the last injection, rats in the

Table 1. Treatment and nutritional supplementation of the thirteen groups studied

<table>
<thead>
<tr>
<th>Groups</th>
<th>Injection</th>
<th>Amino acid Supplementation (6.8 mmol/kg per d and up to 0.5 g N/kg per d by addition of NEAA)</th>
<th>Metabolic inhibitors</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>NEAA</td>
<td>GLN</td>
</tr>
<tr>
<td>1. Ad libitum (AL)</td>
<td>None</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>2. Pair-fed (PF)</td>
<td>NaCl</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>3. DEX-NEAA</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>4. DEX-GLN</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>5. DEX-ARG</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>6. DEX-NEAA-MSO</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>7. DEX-NEAA-SMT</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>8. DEX-NEAA-DFMO</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>9. DEX-GLN-SMT</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>10. DEX-GLN-DFMO</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>11. DEX-ARG-SMT</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
<tr>
<td>12. DEX-ARG-DFMO</td>
<td>DEX</td>
<td>*</td>
<td>*</td>
</tr>
</tbody>
</table>

NEAA, non-essential amino acids; GLN, glutamine; ARG, arginine; MSO, methionine sulfoximine; SMT, S-methylthiourea; DFMO, N-α-difluoromethylornithine; NaCl, intraperitoneal injection of NaCl (9 g/l) during 5 d; DEX, intraperitoneal injection of dexamethasone (1.5 mg/kg per d) during 5 d.

* Amino acids were administered by the enteral route (from day 0 to day 4). All diets were made isonitrogenous (up to 0.5 g N/kg per d) by addition of a mixture of non-essential amino acids (glycine, histidine and alanine in equimolar amounts).

† Metabolic inhibitor was administered by the enteral route (day –1 to day 4).
post-absorptive state were anaesthetized and killed by decapitation.

Blood sampling
Blood was collected on calcium heparin to isolate PMN on a gradient of Ficoll-Hypaque (for determination of oxidative metabolism and superoxide anion generation) or by sedimentation (for H$_2$O$_2$ determination).

Procedures and analytical methods
Separation of polymorphonuclear cells for determination of superoxide anion generation. Blood was layered carefully on a double gradient of Ficoll-Hypaque (Histopaque 1083 and 1119; St Quentin Fallavier, France) with equal volumes. After centrifuging (700g, 30 min, +20°C), the Ficoll-Hypaque layers were removed and the PMN were transferred to a polystyrene tube (Falcon; Elvetec, Clermont Ferrand, France). The residual erythrocytes were destroyed by a short treatment with an ammonium chloride solution (ammonium chloride 16 mM, sodium hydrogen carbonate 1 mM, EDTA 0.01 mM).

The PMN were then washed twice with PBS, resuspended and adjusted to $10^6$ cells/ml with Hank’s balanced salt solution. Cell viability was determined by trypan blue exclusion and was more than 95%. May-Grunwald-Giemsa staining of leucocytes was carried out to determine the efficiency of our separation, and gave more than 80% PMN in each preparation.

Determination of oxidative metabolism by chemiluminescence assay. Oxidative metabolism was measured by chemiluminescence. Luminol-dependent chemiluminescence was measured using a luminometer (Model 1250; LKB Pharmacia, Trappes, France) with disposable polypropylene tubes at 37°C. PMN ($5 \times 10^9$ cells) were pre-incubated for 5 min at 37°C. Luminol was then added to the medium at a final concentration of 10$^{-6}$ M. Time was recorded from the addition of the stimulating agent (phorbol myristate acetate, 10$^{-8}$ M). All the results are expressed as the maximum peak values (mV) corresponding to the maximal activation state.

Determination of extracellular superoxide generation by ferricytochrome C reduction. Extracellular O$_2^•^-$ generation was measured by reduction of ferricytochrome C (horse-heart type III) as previously described (Vasson et al. 1994). Briefly, PMN ($5 \times 10^8$ cells) pre-incubated for 5 min were added to a ferricytochrome C (0.4 mM) solution and stimulated with phorbol myristate acetate (10$^{-6}$ M) for 10 min at 37°C. The final volume of the reaction mixture was adjusted to 0.5 ml with PBS. Incubation was stopped by placing the tubes in an ice bath for 10 min. After centrifuging (400g, 10 min, 4°C), the absorbance of the supernatant fractions was measured at 545 nm (DU540 spectrophotometer; Beckman-Coulter, Gagny, France). The amounts of extracellular O$_2^•^-$ produced were calculated from the absorbance using an extinction coefficient of 20 nmol/cm. The results are expressed in picomoles of released O$_2^•^-$/min per $10^8$ cells.

Measurement of intracellular hydrogen peroxide production by polymorphonuclear cells. Leucocytes were isolated and intracellular H$_2$O$_2$ was measured as previously described (Moinard et al. 1999). In brief, leucocytes were isolated after gravity sedimentation. During leucocyte oxidative burst non-fluorescent intracellular dichlorofluorescin is oxidized by H$_2$O$_2$ to highly fluorescent 2’-7’-dichlorofluorescin. PMN were discerned among leucocyte populations and 2’-7’-dichlorofluorescin fluorochrome of PMN was measured using a flow cytometer (Epics, Coulter, USA). PMN were selected on the basis of their size and granularity by the combination of low-angle forward scattered and right-angle scattered laser light. For each determination, 2,500 PMN were counted.

Results are expressed as the ratio of fluorescence produced by phorbol myristate acetate-stimulated leucocytes to fluorescence produced by non-stimulated cells.

Statistical analysis
Data are expressed as mean values and SEM. Comparisons between sets of data were made using the Kruskal–Wallis H test followed by the Newman–Keuls test when appropriate as indicated in Table and Fig. legends. Programme Conversationnel de Statistiques pour les sciences et le Marketing (PCSM) software was used (Deltasoft, Grenoble, France). Values of $P<0.05$ were considered significant.

Results
Effect of arginine and glutamine on oxidative metabolism in stimulated polymorphonuclear cells
Oxidative metabolism evaluated by chemiluminescence was increased under GLN and ARG supplementation.
Table 3. Oxidative metabolism, extracellular O$_2^{**}$ and intracellular H$_2$O$_2$ production by stimulated polymorphonuclear cells (PMN) from dexamethasone (DEX)-treated rats (1·5 mg/kg per d during 5 d) and supplemented with either non-essential amino acids (NEAA; DEX-NEAA), NEAA + methionine sulfoximine (MSO) (DEX-NEAA-MSO), NEAA + S-methylisothiourea (SMT) (DEX-NEAA-SMT) or NEAA + α-difluoromethylornithine (DFMO) (DEX-NEAA-DFMO)* (Mean values and standard errors of the mean for nine rats per group)

<table>
<thead>
<tr>
<th></th>
<th>Oxidative metabolism (mV/5·10$^5$ cells)</th>
<th>Extracellular O$_2^{**}$ (pmol/10$^6$ PMN per min)</th>
<th>Intracellular H$_2$O$_2$ (arbitrary units)</th>
</tr>
</thead>
<tbody>
<tr>
<td>DEX-NEAA</td>
<td>Mean 57 SEM 18</td>
<td>0·56 SEM 0·11</td>
<td>7·0 SEM 0·7</td>
</tr>
<tr>
<td>DEX-NEAA-MSO</td>
<td>69 SEM 23</td>
<td>0·25 SEM 0·06</td>
<td>5·0 SEM 0·7</td>
</tr>
<tr>
<td>DEX-NEAA-SMT</td>
<td>42 SEM 6</td>
<td>0·34 SEM 0·09</td>
<td>6·0 SEM 0·8</td>
</tr>
<tr>
<td>DEX-NEAA-DFMO</td>
<td>31 SEM 20</td>
<td>0·54 SEM 0·10</td>
<td>5·0 SEM 0·4</td>
</tr>
</tbody>
</table>

* For details of study groups and procedures, see Table 1 and p. 690.

Table 5. Oxidative metabolism, extracellular O$_2^{**}$ and intracellular H$_2$O$_2$ production by stimulated polymorphonuclear cells (PMN) from dexamethasone (DEX)-treated rats (1·5 mg/kg per d during 5 d) and supplemented with either glutamine (GLN; DEX-GLN), GLN + methionine sulfoximine (SMT) (DEX-GLN-SMT) or GLN + α-difluoromethylornithine (DFMO) (DEX-GLN-DFMO)† (Mean values and standard errors of the mean for nine rats per group)

<table>
<thead>
<tr>
<th></th>
<th>Oxidative metabolism (mV/5·10$^5$ cells)</th>
<th>Extracellular O$_2^{**}$ (pmol/10$^6$ PMN per min)</th>
<th>Intracellular H$_2$O$_2$ (arbitrary units)</th>
</tr>
</thead>
<tbody>
<tr>
<td>DEX-GLN</td>
<td>Mean 152 SEM 24</td>
<td>1·41 SEM 0·20</td>
<td>11·2 SEM 0·6</td>
</tr>
<tr>
<td>DEX-GLN-SMT</td>
<td>40* SEM 11</td>
<td>0·33* SEM 0·11</td>
<td>3·3* SEM 0·4</td>
</tr>
<tr>
<td>DEX-GLN-DFMO</td>
<td>40* SEM 9</td>
<td>0·29* SEM 0·10</td>
<td>4·8* SEM 0·6</td>
</tr>
</tbody>
</table>

† For details of study groups and procedures, see Table 1 and p. 690.

In the control groups (DEX-NEAA) the inhibitors used had no effect on oxidative metabolism, extracellular O$_2^{**}$ or H$_2$O$_2$ production by stimulated PMN (Table 3).

In the ARG-supplemented groups, oxidative metabolism was decreased in rats treated with SMT and DFMO (DEX-ARG-SMT, DEX-ARG-DFMO v. DEX-ARG; P<0·05). However, the three inhibitors did not significantly modify extracellular O$_2^{**}$ generation. Concerning intracellular H$_2$O$_2$ production by PMN, only SMT and DFMO abolished the enhancing effect of ARG (DEX-ARG-SMT, DEX-ARG-DFMO v. DEX-ARG; P<0·05) (Table 4).

In GLN-supplemented groups the use of SMT and DFMO abolished the GLN-induced increase in oxidative metabolism, extracellular O$_2^{**}$ generation and intracellular H$_2$O$_2$ production by PMN (DEX-GLN-SMT, DEX-GLN-DFMO v. DEX-GLN; P<0·05) (Table 5).

Discussion

The amino acids GLN and ARG are considered immuno-nutrients (De Bandt & Cynober, 1998; Calder & Yaqoob, 1999; Wu et al. 2000). This term was introduced to describe the effects of specific nutrients on immune functions. The present study, in agreement with the literature (Moffat et al. 1996, Moinard et al. 1999; Furukawa et al. 2000), shows the ability of GLN and ARG to increase respiratory burst of PMN from stressed rats. GLN is utilized by immune cells. In particular, Pithon Curi et al. (1997) have already characterized the glutaminase activity in PMN. This glutaminase activity was higher than that reported for lymphocytes and macrophages, which confirms the importance of GLN for the functioning of these cells. These authors also reported that less than 1 % GLN was totally oxidized, confirming that GLN is not only a fuel for these cells but may generate other metabolites (i.e. ARG or polyamines, which may play a key role in the cells). Concerning ARG, two pathways exist in PMN. First, the arginine pathway in which ARG is converted into urea and ornithine, which generates aliphatic polyamines...
by the action of ornithine decarboxylase. Second, ARG is the substrate for NO’ synthesis; NO’ is synthesized by the action of NO* synthase resulting in the formation of NO* and citrulline (Cynober et al. 1995). In the present study we determined which of these metabolic pathways were involved at the whole body level in GLN- and ARG-mediated actions. For this purpose enzymic inhibitors were used. Their utilization did not modify respiratory burst in control groups (i.e. DEX-NEAA-treated groups). This result indicates that in basal conditions the metabolic inhibitors did not affect the parameters measured and that only the modifications induced by GLN or ARG were affected by the metabolic inhibitors.

The use of MSO showed that the action of ARG is not related to GLN synthesis. However, ARG stimulates respiratory burst through NO’ and polyamine pathways. Several studies have shown the ability of NO’ to control NADPH-dependent oxidase activity (the first enzyme involved in radical oxygen species production) (Seth et al. 1994; Catz et al. 1995; Fujii et al. 1997). However, this effect of NO’ on NADPH oxidase is controversial. These differences may be explained by the results of Pieper et al. (1994), who showed a biphasic effect of NO’ on NADPH-oxidase activity. For these authors this mediator has a stimulatory effect at low dose and an inhibitory effect at high dose. The effect of polyamines on ROS production was previously observed in vitro (Walters et al. 1998); treatment with DFM0 produced dose-dependent inhibition of the respiratory burst in PMN. Our results confirm that in vivo ARG may be a polyamine precursor that in turn stimulates ROS production in stimulated PMN. Concerning GLN, we showed that it stimulates oxidative metabolism via NO’ and polyamines generation. But how does GLN generate NO’ and polyamines?

Recently, several studies have investigated the relationship between GLN and NO’. O’Dowd & Newsholme (1997) have evidenced the involvement of GLN in NO’ production by stimulated PMN. They showed that the PMN possess the ability to produce ARG from GLN. However, these observations were made in vitro. For the first time our study has shown the ability of GLN supplementation in vivo to be an NO’ precursor in neutrophils.

To the best of our knowledge there are few available data on the relationship between GLN and polyamine pathways; Kandil et al. (1995) have established that GLN stimulates ornithine decarboxylase dose- and time-dependently in a porcine jejunal enterocyte cell line. These data suggest that GLN may be able to stimulate ornithine decarboxylase in neutrophils. Also, GLN could be a precursor of polyamine synthesis (see Fig. 1). This last point deserves further study.

In conclusion, we demonstrate that both ARG and GLN stimulate oxidative metabolism in stressed rats via NO’ and polyamine generation. However, the present study did not allow us to determine which one of these two pathways is predominant. Further study, using labelled molecules, is required to quantify the metabolic flux at the whole-body level and in PMN. It would be useful to determine whether these observations are specific to neutrophils or may be extended to other cells and tissues. GLN and ARG have many common actions on immunity and protein metabolism. In practice, they are known to improve patient outcome, to enhance immunity and post-traumatic N metabolism. However, there are few data available on the mechanistic action of these nutrients in catabolic patients. Knowledge of their action would help optimize their use in clinical practice. Further studies are necessary to determine which of these two amino acids is more efficacious in catabolic patients.

Acknowledgements

This work was supported in part by funds from the Ministry of Research and Technology (EA2416). C. Moinard, S. Walrand and F. Caldefie are supported by fellowships from the Ministry of Research and Technology. We are indebted to Professor J. Chassagne (CJIP, Hospital) and Dr A. Tridon (Laboratory of Immunology) for helpful collaboration.

References


Modulation of respiratory burst by glutamine

695


