ORIGINAL ARTICLE

Taurine transporter gene expression in peripheral mononuclear blood cells of type 2 diabetic patients

Loria Bianchi · Riccardo Lari · Roberto Anichini · Alessandra De Bellis · Angela Berti · Zaleida Napoli · Giuseppe Seghieri · Flavia Franconi

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Abstract Taurine acts as antioxidant, cell osmolyte, modulator of glucose metabolism, and plays a role in the retinal function. It is 10³-fold more concentrated in the intracellular than in the extracellular milieu due to a specific taurine-Na-dependent transporter (TauT), which is upregulated by hypertonicity, low extracellular taurine, or oxidative stress and acutely downregulated 'in vitro' by high glucose concentrations. Aim of this study was to investigate whether TauT expression was modified in mononuclear peripheral blood cells (MPC) of type 2 diabetic patients with or without micro/macrovascular complications. Plasma taurine, as well as other sulphurcontaining aminoacids (assayed by HPLC) and TauT gene expression (assayed by real-time PCR analysis) were measured in MPC of 45 controls and of 81 age-and-sex matched type 2 diabetic patients with or without micro/ macrovascular complications. Median value (interquartile range) of plasma taurine was significantly lower in diabetic patients than in controls [28.7 (13.7) µmol/l vs. 46.5 (20.3) μ mol/l; P < 0.05], while median TauT expression, in arbitrary units, was significantly higher in diabetics than in controls [3.8 (3.9) vs. 1 (1.3); P < 0.05) and was related to HbA1c only in controls (r = 0.34; P < 0.05). Patients with

 $\begin{tabular}{ll} \textbf{Keywords} & Taurine \cdot Taurine \ transporter \cdot Type \ 2 \\ diabetes \cdot Retinopathy \cdot Mononuclear \ peripheral \ blood \ cells \end{tabular}$

retinopathy (n = 25) had lower TauT expression than those

who were unaffected [3.1 (2.8) vs. 4.1 (3.4); P < 0.05],

while persistent micro/macroalbuminuria was associated with unchanged TauT expression. A trend toward reduction

in TauT expression was observed in patients with macro-

angiopathy [n = 27; 3.3 (2.5) vs. 4 [3.7]; P = NS]. In

conclusion, TauT gene is overexpressed in MPC of type 2

diabetic patients, while presence of retinopathy is specifi-

cally associated with a drop in TauT overexpression, sug-

gesting its possible involvement in this microangiopathic

L. Bianchi · R. Lari · Z. Napoli Department of Clinical Chemistry, Spedali Riuniti, Pistoia, Italy

R. Anichini · A. De Bellis · A. Berti · G. Seghieri (⊠) Department of Internal Medicine, Spedali Riuniti, Viale Matteotti 9/D, 51100 Pistoia, Italy e-mail: gseghier@tin.it

F. Franconi Department of Pharmacology, University of Sassari, Sassari, Italy

Introduction

lesion.

The semi-essential amino acid taurine has numerous functions (Huxtable 1992; Schaffer et al. 2000) playing also a role in glucose metabolism (Franconi et al. 1996; Schaffer et al. 2009). Taurine deficiency has been observed in experimental and in clinical models of both type 1 and type 2 diabetes mellitus, or in clinical situations of insulin resistance (De Luca et al. 2001; Franconi et al. 1995, 2006; Merheb et al. 2007) and has been associated with the pathogenesis of diabetic complications (Ha et al. 1999; Vilchis and Salceda 1996). Next, taurine administration seems to prevent several alterations induced by hyperglycaemia 'in vitro' (Franconi et al. 2006; Schaffer et al. 2009; Verzola et al. 2002; Wu et al. 1999) and by diabetes in animal models (Franconi et al. 2003, 2006) or in humans (Franconi et al. 1995; Xiao et al. 2008; Schaffer et al. 2009; Ito et al. 2011).



The mechanisms that underlie taurine deficiency in diabetes mellitus are, nevertheless, still unknown, even if it has been suggested that diabetes-specific taurine deficiency could be due, at least in part, to the impairment in net intestinal and/or renal taurine absorption (Merheb et al. 2007), or to disruption of trans-membrane Na-dependent amino acid taurine transporter (TauT) (Stevens et al. 1999).

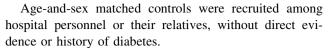
TauT is present in many cells, working against a concentration gradient since taurine is more concentrated (about 10³-fold) in the intracellular compartment than in the extracellular milieu, and has, so far, been both morphologically and functionally well characterized (Tappaz 2004; Han et al. 2006). TauT is regulated by hypertonicity through the TonE (tonicity-responsive element)/TonB (TonE-binding protein) pathway (Uchida et al. 1992; Han et al. 2006; Ito et al. 2004). Acute hyperglycaemia, moreover, reduces the expression of Taut mRNA and protein 'in vitro' in different cellular models (Stevens et al. 1999; Askwith et al. 2009) and, interestingly, oxidation, DNA damage, or dietary manipulation, also affect TauT expression (Matsell et al. 1997; Han et al. 2009; Han and Chesney 2010).

Since no study has ever investigated whether TauT expression is varied 'in vivo' in patients with type 2 diabetes mellitus, we conceived this study to test whether TauT gene expression is modified in blood mononuclear peripheral cells (MPC) of patients with type 2 diabetes with or without micro/macrovascular complications.

Materials and methods

Patients and controls

Patients were chosen among diabetic subjects who consecutively attended the Diabetes Outpatient Clinic of our hospital and were classified as affected by type 2 diabetes since the diagnosis was done after the age of 40 years and did not require insulin therapy at the onset of their illness. Thirty-nine patients were on oral therapy, 16 on insulin, 17 on oral therapy + insulin, and 9 on diet alone. In diabetic patients, presence of retinopathy was ascertained by retinal fluorescein angiography as previously described (Di Simplicio et al. 1995) and nephropathy was diagnosed if 24-h-urinary albumin excretion rate (AER), evaluated as the mean value of three samples during the last 6 months, was $\geq 20 \mu g/min$. Plasma creatinine was $< 124 \mu mol/l$ in all cases. Presence of macroangiopathy was ascertained by anamnestic and clinical-instrumental criteria (clinical evaluation as well as history of previous cardiovascular disease, presence of altered ST-T in ECG and/or presence of atherosclerotic lesions in carotids or lower limbs' vessels evaluated by Echo-Color-Doppler technique).



This study has been approved by the Ethical Committee of our Hospital and all subjects gave their written informed consent prior to their inclusion in the study.

Biochemical assays

All biochemical assays were obtained in the morning in fasting condition. In both patients and controls we measured fasting plasma glucose and glycated hemoglobin (HbA1c) by routine methods. In both patients and controls we also assayed plasma levels of taurine and methionine by separation followed by fluorimetric detection of their *p*-orthophtalaldehyde (OPA) derivatives (Bianchi et al. 1999) and homocysteine by using fluorescence polarization immunoassay (FPIA, Abbott, Italy).

Taurine transport gene expression

In each subject blood mononuclear peripheral cells (MPC, lymphocytes and monocytes) were separated from 4 ml of whole blood by Accuspin System-Histopaque-1077 (Sigma-Aldrich, Italy) and stored in RNA later solution (Qiagen, Italy) at -20° C.

Each sample contained a median count of 2.53×10^6 (interquartile range: 1.1×10^6) cells/ml, of which 2.05×10^6 10^6 (interquartile range: 0.92×10^6) lymphocytes/ml and 0.44×10^6 (interquartile range: 0.19×10^6) monocytes/ ml. Total RNA was purified through Rneasy-Plus-Mini-kit (Qiagen, Italy); its concentration was determined by measuring absorbance at 260 nm and its purity was provided by readings' ratios of 260/280 nm. First-strand cDNA was synthesized from 1 µl of RNA by random examers and SuperScript III reverse transcriptase (Invitrogen, Italy). The amount of c-DNA was quantified by a Qubit spectrophotometer (Invitrogen, Italy). TauT and GAPDH (housekeeping gene) mRNA were quantified by real time RT-PCR (Rotorgene 6000, Corbett, UK), using intercalation of SYBR Green. Primers' sequences for TAUT were: 5'-GAAATCTTCATCGCCTTCG-3' (forward) and 5'-TA GCCAATCATGTCCTCA-3' (reverse), while for GAPDH were: 5'-GAAGGTGAAGGTCGGAGT-3' (forward) and 5'-GAAGATGGTGATGGGATTTA-3' (reverse) (Schöndorf et al. 2002; Mochizuki et al. 2005). To quantify mRNA expression a standard curve was constructed from cDNA generated from human reference RNA (1 µg/1 µl; Stratagene) and standard curves were used for estimation PCR efficiency and as a calibration curve to estimate the starting concentrations of unknown samples (Ramarkers et al. 2003). Amplification was performed in triplicate by the following thermal cycling conditions: 2 min at 52°C,



activation at 95°C for 10 min, followed by 45 cycles of 15 s at 95°C and 60 s at 60°C. Dissociation analysis of the PCR products was performed by running a gradient from 95 to 55°C (gradient 0.1°C) to confirm the presence of single PCR products.

Changes in mRNA gene expression levels were reported, after normalization to GAPDH, as fold changes relative to controls and values of fold changes in the control sample versus target samples represent averages from triplicate measurements. Relative quantification was performed using the comparative cycle threshold (Ct) method after determining the Ct values for reference (GAPDH) and target gene (TauT) in each sample according to the $2^{-\Delta\Delta Ct}$ method. In short, Δ Ct was obtained by subtracting the Ct of target gene (TauT) from the Ct of the reference gene (GAPDH), and then it was calculated $\Delta\Delta$ Ct = Δ Ct – Δ Ct_Mean. Finally, gene expression level was expressed as $2^{-\Delta\Delta Ct}$ (Pfaffl 2006).

Statistical methods

Data were analyzed by ANOVA (after log transformation for abnormally distributed variables) or via Wilcoxon rank-sum-test to check for statistically significant differences among groups. Relations among variables were tested by univariate analysis, expressing Spearman's correlation coefficients. Values of abnormally distributed variables were expressed as median (interquartile ranges). Significance of P value was set at <0.05.

All statistical analyses were carried out by means of SAS software for Windows, version 8.2 (SAS Institute Inc., Cary, NC, USA).

Table 1 Main characteristics of diabetic patients and of control subjects

	Controls	Type 2 diabetes	P*
No.	45	81	
Age (years)	58.8 ± 9.3	63 ± 10	NS
Sex (% of males)	46	49	NS
BMI (kg/m^2)	23.9 ± 3	29.4 ± 5	< 0.05
Duration of diabetes (years)	_	10.8 ± 8	_
Plasma glucose (mmol/l)	5.1 ± 0.7	8.8 ± 2.9	< 0.05
HbA1c (%)	5.2 ± 0.6	7.7 ± 1.3	< 0.05
Plasma taurine (µmol/l)	46.5 (20.3) ^a	28.7 (13.7) ^a	< 0.05
Plasma methionine (µmol/l)	14.9 (4.4) ^a	$17 (8.4)^{a}$	< 0.05
Plasma homocysteine (µmol/l)	$10.4 (4.4)^a$	13.6 (6.5) ^a	< 0.05
Lymphocites (cells $\times 10^6$ /ml)	2.10 (0.77)	2.19 (1.22)	NS
Monocytes (cells $\times 10^6$ /ml)	0.43 (0.16)	0.45 (0.21)	NS
Taurine Transporter gene expression ^b	1 (1.3) ^a	$3.8 (3.9)^a$	< 0.05
Retinopathy (no.)	_	25	_
Micro-macroalbuminuria (no.)	_	12	-
Macroangiopathy (no.)	_	27	_

^{*} Calculated by Wilcoxon ranksums-test

Results

The median values (interquartile range) of both plasma methionine and homocysteine were higher while, on the contrary, median plasma taurine was significantly reduced in type 2 diabetic patients than in controls (Table 1) and because there was no significant gender difference as to plasma aminoacid levels and to TauT gene expression, data of men and women were pooled in all analyses.

In presence of similar numbers of both lymphocytes and monocytes collected in blood samples (Table 1) diabetes mellitus greatly affected TauT gene expression inducing an increase in its expression about fourfold higher in diabetic patients in comparison with healthy volunteers (Table 1; Fig. 1).

After stratifying diabetic patients according to micro and macroangiopathic complications, patients with retinopathy had significantly lower median TauT gene expression when compared to other non retinopathic diabetic patients, even if it was still higher when compared with the expression of healthy volunteers [3.1 (2.7) AU vs. 1 (1.3) AU; P < 0.05 (Table 2; Fig. 2a)].

TauT gene expression was unaffected by presence of persistent micro/macroalbuminuria (Table 2; Fig. 2b), whereas a trend toward a decrease was observed in patients with macroangioapthy.

Lymphocyte and monocyte counts were unchanged in diabetic patients with or without micro/macroangiopathic complications (data not shown).

Plasma levels of taurine, homocysteine, and methionine were not varied in diabetic groups with or without evidence of micro and macroangiopathic complications (Table 2).



^a Median (interquartile range)

^b Expressed in arbitrary units (AU)

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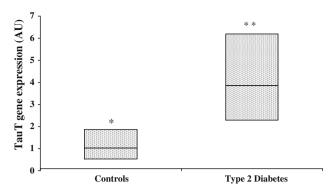


Fig. 1 Median with 1st and 3rd quartile values of TauT gene expression in arbitrary units (AU) in MPC of controls and of type 2 diabetic patients. * versus **P < 0.05**

Finally, we also investigated the correlations among TauT gene expression and age, BMI, duration of diabetes fasting plasma glucose, HbA1c, plasma taurine, methionine, and homocysteine (Table 3). Significant correlations have been found only in controls: in these, latter TauT gene expression was positively related with HbA1c and inversely associated with fasting plasma glucose whereas plasma taurine was negatively related with HbA1c (r = -0.44; P < 0.05). Additionally, in controls, taurine/methionine ratio was inversely related with HbA1c (r = -0.38; P < 0.05) and, reciprocally, methionine/taurine ratio and HbA1c were each other positively related (r = 0.40; P < 0.05).

Discussion

This study evidences that TauT gene is expressed in adult human MPC. These results are in line with previous data that described TauT gene expression in foetal peripheral lymphocytes and platelets (Iruloh et al. 2007). Indeed, here we show that TauT expression: (1) is not sexual dimorphic, and (2) in normal subjects is inversely related with fasting plasma glucose (r = -0.36; P < 0.05) and positively associated with Hba1c (r = 0.34; P < 0.05). This speculatively suggests that TauT gene is acutely downregulated by incremental variations of plasma glucose in its physiological range (3.7-6.8 mmol/l), and, on the contrary, seems chronically upregulated at plateau by higher daily glycemic milieu in the non diabetic HbA1c range: 3.6-6.5%. These findings could tentatively indicate a dimorphic glucose behaviour suggesting a fine acute adjustment of MPC TauT, associated with its upregulation at chronic glycemic plateau.

The most important novelty of this paper, however, resides in the fact that TauT gene is overexpressed in the MPC derived from type 2 diabetic patients indicating that it is associated with an alteration of TauT expression

gene expression in MPC of control subjects, compared to diabetic patients with or Median plasma values (interquartile range) of taurine, homocysteine, methionine as well as of TauT without micro-macroangiopathic lesions

	Taurine (µmol/l)	(1/1)		Homocysteine (µmol/l)	(mool/l)		Methionine (µmol/l)	(Momuli)		TauT (AU) ^a		
	Presence Absence	Absence	Ь	Presence	Absence	Ь	Presence	Absence	Ь	Presence	Absence	Ь
Retinopathy	27.3 (15.4)	28.9 (13.6)	SN	14 (9.5)	13.5 (6.4)	SN	16.8 (5.9)	16.9 (9.2)	SN	3.1 (2.7)	4.1 (3.4)	<0.05
Micro-macroalbuminuria	25.8 (8.6)	29.1 (13.8)	SN	14.5 (7.3)	8.9 (6.4)	SN	16.8 (9.9)	16.9 (8.7)	SN	4.2 (5.8)	3.8 (3.4)	NS
Macroangioapthy	30.6 (13.4)	28.4 (14.4)	SN	13.4 (10.1)	13.6 (6.3)	SN	16.1 (4.4)	17.1 (9.9)	SN	3.3 (2.5)	4 (3.7)	NS

Expressed as arbitrary units (AU)



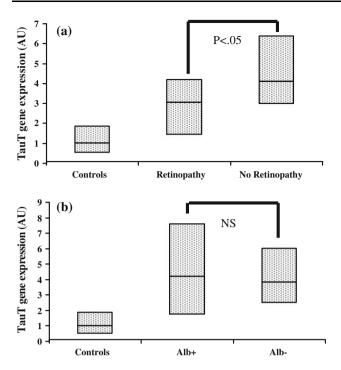


Fig. 2 Median with 1st and 3rd quartile values of TauT gene expression in arbitrary units (AU) in MPC of controls and of type 2 diabetic patients with or without retinopathy (a) or micro-macroal-buminuria (Alb+ and Alb-) (b)

Table 3 Correlation coefficients between TauT expression in MPC and other variables in controls and in type 2 diabetic patients

	Controls $(n = 45)$	P	Type 2 Diabetes $(n = 81)$	Р
	r		r	
Age	0.24	NS	0.005	NS
BMI	-0.04	NS	0.12	NS
Duration of diabetes	-	NS	-0.13	NS
Plasma glucose	-0.36	0.02	0.03	NS
HbA1c	0.34	0.02	0.06	NS
Plasma taurine	0.16	NS	-0.10	NS
Plasma methionine	-0.12	NS	-0.02	NS
Plasma homocysteine	0.19	NS	-0.16	NS

'in vivo', at least in these cells. It is known that TauT expression is adaptively upregulated by low extracellular taurine (Tappaz 2004; Han et al. 2006; Uchida et al. 1992) and taurine levels are indeed decreased in the group of diabetic patients, in agreement with most previous observations (De Luca et al. 2001; Franconi et al. 1995; Merheb et al. 2007). Low plasma taurine in diabetes has been ascribed to reduced taurine reabsorption at renal or intestinal level due to the impairment of cell TauT expression (Merheb et al. 2007). In this context, however, increased MPC TauT expression, as we observed, cannot

automatically mirror an equal TauT upregulation at renal tubular cells, where taurine handling seems to be a complex mechanism controlled by adaptive regulation of TauT in response to taurine availability as well as by its upregulation due to increased osmolality (Satake et al. 2010).

Finally, although mammals placed on taurine-restricted diet markedly reduce taurine urinary excretion enhancing renal TauT (Han et al. 2006), here the lack of correlation between TauT expression and plasma taurine suggests that the decrease in plasma taurine levels has a scarce effect on the adapting mechanisms that produce TauT upregulation.

Acute hyperglycemia in vitro downregulates TauT in cultured rat cardiomyocytes (Shi et al. 2003), in cultured retinal pigment epithelial cells (Stevens et al. 1999; Nakashima et al. 2005) and in human Schwann cells (Askwith et al. 2009). Glucose-induced TauT downregulation in human Schwann cells is reversed by inhibition of aldose reductase as well as by protein kinase C (PKC) inhibitors indicating the importance in the process of both osmolality (Askwith et al. 2009) and of glucose-driven intracellular pathways (Stevens et al. 1999; Shi et al. 2003). All this could be viewed as deeply conflicting with our results obtained in vivo. To explain this apparent contradiction, however, we should consider some points.

The first point is that in almost every study the inhibitory effect exerted by high glucose 'in vitro' on cell TauT expression is obtained with glucose medium concentrations equal to or above 20 mmol, which largely exceed the glucose concentrations observed in our patients (7–16 mmol/l). As to this aspect it is to note, moreover, that no studies have, so far, specifically investigated TauT cell expression and/or function 'in vitro' in such a lower range of glucose concentrations.

The second point is the deeply different time length of high glucose exposure in these two different situations: relatively short (from 24 h to a maximum of 7 days) in previous studies 'in vitro', as compared to a much longer chronical high glucose exposure as happens in diabetic patients.

The third point is characterized by the fact that when we consider diabetic patients we must take into account that diabetes is just not only hyperglycemia: in other words increased TauT expression which we have observed could be related with other constitutive aspects of the diabetic 'milieu' such as, for instance, the presence of various degrees of insulin resistance or the presence of increased concentration of other metabolites such as, for instance, lipids, free radicals, etc. In this context, previous studies concerning retinal pigment epithelial cells obtained from diabetic rats showed that cell taurine uptake was elevated, rather than decreased, suggesting that diabetes is able to stimulate 'in vivo', at least in some cell lines, the activity of the taurine transporter (Vilchis and Salceda 1996).



Furthermore raised TauT gene expression by MPC can, additionally, be explained by an increase in MPC overall activation in type 2 diabetes, since, as recently described, peripheral B cells contribute to insulin resistance by producing pathogenic IgG antibodies (Winer et al. 2011), suggesting that these circulating cells can be activated by diabetic-related mechanisms.

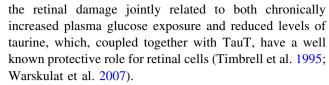
A further factor possibly implicated in stimulating TauT expression could be insulin itself, even if there are no evidences in this sense from previous reports, and, in the present study, no difference in TauT expression was noted between patients on insulin therapy, when compared with those on other treatments (data not shown).

Finally TauT promoter contains an antioxidant response element (ARE), activated in response to prooxidants (Nakashima et al. 2005) and oxidants, whose production is raised in diabetes, increase TauT expression in human Schwann cells 'in vitro' (Askwith et al. 2009). It is also interesting to note that nitric oxide, which is a promoter of TauT expression in cultured human retinal pigment epithelial cells, is increased in diabetes and its action can be amplified by the presence of oxidative stress (Bridges et al. 2001).

A separate consideration should be done about the relation linking high glucose with hypertonicity and TauT expression. It is indeed also known that TauT is generally upregulated by hypertonicity through the TonE/TonEBP signalling pathway (Ito et al. 2004, 2009). Nevertheless conflicting results have been described when different cells have been acutely challenged with high glucose concentrations 'in vitro': under such a circumstance extracellular and intracellular hypertonicity seem to have a sort of 'paradoxical' inhibitory effect on TauT expression and function, both restored, even if in not all cases, by treatment with aldose reductase inhibitors (Stevens et al. 1999; Askwith et al. 2009). This represents a further piece of evidence for possible discrepancies between what observed 'in vitro' and the effect of chronic exposure to higher glucose levels 'in vivo': proving that hypertonicity cannot be assumed to be the sole or most important mechanism involved in the regulation of TauT expression (Han et al. 2006).

In summary, considering that taurine is an osmolyte and an antioxidant and that its intracellular content is mainly regulated by TauT, the hypothesis stemming by our data is that cells chronically exposed to diabetic milieu might attempt to improve their homeostasis increasing TauT gene expression.

Furthermore, it is of interest to note that TauT overexpression is significantly reduced in patients with retinopathy while nephropathy and macroangiopathic complications do not affect TauT, thus suggesting that retinopathic patients have a minor capacity to counteract



Finally, plasma levels of other sulphur-containing aminoacids such as methionine and homocysteine were elevated in diabetic patients. Increased levels of homocysteine in diabetes have previously been described (Hoogeveen et al. 2000). And, notably, the increase in homocysteine observed in diabetic patients could decrease the substrate for taurine biosynthesis decreasing the availability of cysteine (Wu 2009). Previous data on plasma methionine are scarce and one paper (Tessari et al. 2005) shows that it is not changed in diabetic patients, while we found a significant increase in methionine. However, considering that taurine inhibits methionine uptake in some cell models such as intestinal Caco-2 cells (Martin-Venegas et al. 2009) the observed increase in methionine could be biologically plausible.

In conclusion, this study describes, for the first time, a raised TauT gene expression in MPC of type 2 diabetic patients. Whether this specific upregulation is the response to a number of diabetes-associated factors such as increase in circulating oxidants, hypertonicity, chronic hyperglycemia, reduction in plasma taurine or eventually other unidentified factors remains to be further elucidated. The increment of TauT expression is reduced in MPC of patients with retinopathy, suggesting a possible role for TauT in the protective mechanisms against the retinal damage of diabetes. In any case, even if these data have some limitations and warrant further confirmation, they indicate TauT as another candidate target of diabetic illness, further confirming the role of taurine in glucose homeostasis both in normoglycemic subjects and diabetic patients.

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Conflict of interest The authors declare that they have no conflict of interest.

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