

J. Radziuk · S. Pye

## Diurnal rhythm in endogenous glucose production is a major contributor to fasting hyperglycaemia in type 2 diabetes. Suprachiasmatic deficit or limit cycle behaviour?

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**Abstract** *Aims/hypothesis:* An increase in endogenous glucose production (EGP) is a major contributor to fasting morning hyperglycaemia in type 2 diabetes. This increase is dissipated with fasting, later in the day. To understand its origin, EGP, gluconeogenesis and hormones that regulate metabolism were measured over 24 h. We hypothesised that EGP, and therefore glycaemia, would demonstrate a centrally mediated circadian rhythm in type 2 diabetes. *Subjects and methods:* Seven subjects with type 2 diabetes and six age- and BMI-matched control subjects, fasting after breakfast (08.00 h), underwent a further 24-h fast, with the infusion of [U-<sup>13</sup>C] glucose and [3-<sup>14</sup>C]lactate, starting at 14.00 h. The MCR and production of total and gluconeogenic glucose were determined from the tracer concentrations using compartmental analysis. *Results:* MCR was near constant:  $1.73 \pm 0.10$  in control and  $1.40 \pm 0.14$  ml kg<sup>-1</sup> min<sup>-1</sup> in diabetic subjects ( $p=0.04$ ). EGP in diabetes rose gradually overnight from  $8.2 \pm 0.7$  to  $11.3 \pm 0.5$  μmol kg<sup>-1</sup> min<sup>-1</sup> at 06.00 h ( $p<0.05$ ). Glucose utilisation lagged EGP, rising from  $8.5 \pm 0.6$  to  $10.5 \pm 0.4$  μmol kg<sup>-1</sup> min<sup>-1</sup> ( $p<0.05$ ), inducing a fall in glycaemia from a peak of  $8.0 \pm 0.5$  mmol/l to  $6.3 \pm 0.4$  mmol/l ( $p<0.05$ ). Cortisol and melatonin showed diurnal variations, whereas insulin, glucagon and leptin did not. Melatonin was most closely related to EGP, but its secretion was attenuated in diabetes ( $p<0.05$ ). *Conclusions/interpretation:* In type 2 diabetes, EGP and gluconeogenesis display diurnal rhythms that drive the fasting hyperglycaemia and are absent in healthy control subjects. The rise in EGP may be related to a deficit in

suprachiasmatic nucleus activity in diabetes, or result from non-linear behaviour plus a transition from a normal steady state to a limit cycle pattern in diabetes, or both.

**Keywords** Diurnal rhythms · Dynamical diseases · Endogenous glucose production · Fasting plasma glucose · Limit cycles · Metabolic clearance · Non-linear systems · Tracer methods · Type 2 diabetes

**Abbreviations** EGP: endogenous glucose production · MCR: metabolic clearance rate (of glucose) ·  $R_d$ : utilisation rate (of glucose) · SCN: suprachiasmatic nucleus

### Introduction

Fasting hyperglycaemia is one of the principle criteria for the diagnosis of type 2 diabetes [1]. Its aetiology has been attributed to the beta cell, the liver and the peripheral tissues (muscle) [2], with the relative contributions a matter of continuing investigation.

Since it is the principal source of glucose under postabsorptive conditions, the liver has been implicated as an important player [2, 3], with correlations demonstrated between endogenous glucose production (EGP) and fasting glucose concentrations in a large number of laboratories (for review, see [4]). Some uncertainty arose from a number of observations of similar rates of EGP in type 2 diabetes and healthy control subjects [5, 6]. The resolution of this issue was based on the following observations: (1) glycaemia in type 2 diabetes is never constant, and so the assumption of a steady state could lead to quantitative inaccuracies [7, 9]; and (2) EGP is elevated in the morning by 30–50% and decreases to near-normal rates by afternoon [4, 7, 8]. This results in a high degree of correlation between elevated glycaemia and EGP early in the morning and its dissipation by mid-afternoon [7, 8]. The time of its determination is therefore critical to the conclusions drawn, as is the methodology used. The metabolic clearance rate (MCR) of glucose, however, remains constant throughout the day and is decreased in

J. Radziuk (✉) · S. Pye  
Diabetes and Metabolism Research Unit,  
Ottawa Hospital (Civic Campus),  
1053 Carling Avenue,  
Ottawa, ON, K1Y 4E9, Canada  
e-mail: jradziuk@ohri.ca  
Tel.: +1-613-7614112  
Fax: +1-613-7615329

J. Radziuk · S. Pye  
Ottawa Health Research Institute and the University of Ottawa,  
Ottawa, ON, Canada

type 2 diabetes [7]. It therefore accounts for the baseline elevation in glycaemia that persists throughout the day [7, 9].

Gluconeogenesis has been implicated in the fasting increase in EGP [10, 11]. When deuterated water and the relative incorporation of the deuterium into the second and fifth position of glucose is used to estimate fractional gluconeogenesis, both gluconeogenesis and glycogenolysis appear to contribute to the elevated fasting EGP [12–15].

The observations discussed lead to the following questions:

1. How does the EGP increase to its morning level from rates that are near normal the previous evening? Does this occur during fasting?
2. Is gluconeogenesis a primary contributor to this increase?

The aim of this study was therefore to investigate the hypotheses that in type 2 diabetes (1) there is a diurnal rhythm in EGP with a nocturnal increase and daytime fall; (2) the changes in EGP drive those in glycaemia; (3) gluconeogenesis is a major contributor to the changes in EGP; and (4) the cyclicity of these processes occurs independently of exogenous influences such as meals, and, therefore, autonomously.

Patients with type 2 diabetes were compared with non-diabetic control subjects in terms of the rate of glucose production and an index of gluconeogenesis, measured using tracers during a 24-h fast. The relationship of the diurnal cycle in EGP with circadian rhythms was examined.

## Subjects and methods

**Subjects** Eight patients with type 2 diabetes and eight healthy individuals participated in these studies. Control subjects were matched for age, sex and BMI. All diabetic subjects were treated with diet and exercise ( $n=4$ ), or with sulfonylureas and/or metformin ( $n=4$ ). Any oral hypoglycaemic agents were discontinued 3 days prior to the study. Other medications used were not considered to affect glucose metabolism. The subjects were otherwise healthy on physical and laboratory examination. The studies were approved by the Human Ethics Review Board of the Ottawa Hospital and participants gave their informed consent.

**Protocol** Subjects were prescreened using a polysomnogram, with overnight continuous measurement of four-

channel electroencephalogram, one-lead electrocardiogram, finger oximetry, gastrocnemius electromyogram, and rib cage and abdominal motion by electrical impedance bands. Subjects with sleep apnoea or other sleep disorders were not included in the study because of the known effects of these conditions on metabolism. On day 1, subjects fasted after breakfast (08.00 h). They were admitted to the clinical investigation area at 12:00 h (day 1). Unprimed infusions of [ $U-^{13}C$ ]glucose and [ $3-^{14}C$ ]lactate were initiated at 14.00 h. Sleep was again monitored during the night. The study continued until 14.00 h on day 2. The initiation and termination times were adjusted relative to the normal bedtimes of the subjects so that approximately 9 h of sampling was carried out prior to start of sleep (Table 1). Samples were collected for metabolite, tracer and hormone measurements on a variable schedule related to the anticipated rapidity of changes in the tracer concentrations. At night, samples were collected less frequently, but at least every 40 min.

**Materials** [ $3-^{14}C$ ]lactate was obtained from Amersham (Arlington Heights, IL, USA). [ $U-^{13}C$ ]Glucose was obtained from Cortec (Paris, France). Labelled glucose was purified using ion-exchange HPLC (HPX-87P column; Biorad, Hercules, CA, USA). The [ $^{14}C$ ]lactate was purified on an HPX-87H column (Biorad) All products were then tested for sterility and pyrogenicity.

**Analytical methods** Plasma glucose and lactate were measured using the YSI 2300 (Yellow Springs Instrument Company, Yellow Springs, OH, USA). Plasma NEFA concentrations were determined enzymatically (Wako Chemicals, Richmond, VA, USA). Plasma insulin, glucagon (CV 8%) and leptin were determined using RIAs (with inter-assay CVs of 4%, 8% and 10%, respectively) from Linco Research (St Charles, MI, USA). Melatonin was also measured by RIA (IBL, Hamburg, Germany). Cortisol was measured using an ELISA (Alpco, Salem, NH, USA).

All samples for tracer measurements were prepared by deproteinisation [16]. The supernatant was passed through ion-exchange resin (Dowex 1-X8 and Dowex 50W; Biorad). An aliquot of the neutral fraction was evaporated to dryness and the residue dissolved in water and scintillation fluid added (Scintisafe; Fisher Scientific, Hampton, NH, USA). The lactate was removed from the resin using formic acid (0.5 N), which was then evaporated to dryness, dissolved in water and scintillation fluid added. The radioactivity was determined in a liquid scintillation

**Table 1** Experimental schedule

	Start tracer infusion -9 h		Bedtime 0 h			Termination of study +15 h	
	↓		↓			↓	
Time of tracer infusion (min)	0	240	480	720	960	1,200	1,440
24-h clock time	14.00	18.00	22.00	02.00	06.00	10.00	14.00

counter (Tricarb 2200CA; Perkin Elmer, Wellesley, MA, USA) [16].

The enrichment of plasma [ $^{13}\text{C}$ ]glucose was determined by derivatising the glucose in deproteinised plasma with butyl boronic acid (Alfa Aesar, Ward Hill, MA, USA) and acetic anhydride (BDH, Poole, UK) in the presence of pyridine (BDH) [17]. The butaneboronyl derivative was analysed on a GC/MS (Agilent Technologies, Palo Alto, CA, USA) using a Zebron 1701 column (Phenomenex, Torrance, CA, USA) and selective ion monitoring at 297 and 303 m/z, and a standard curve of known enrichments.

**Calculations** Glucose kinetics were analysed using non-steady-state methods and compartmental analysis, as detailed previously [4, 7]. Concentrations of the infused glucose tracer were fitted to integrated exponential functions [18]. The optimal number of exponentials determined the number of compartments used subsequently ( $\chi^2$  test). MCR, volume of distribution and, if necessary, exchange parameters were evaluated from the infused tracer and used in the calculation of EGP and the rate of glucose utilisation from the glucose concentrations. MCRs were also allowed to vary if indicated by a runs test [7].

An index of gluconeogenesis was obtained from the incorporation of  $^{14}\text{C}$  label from lactate into glucose [19]. The rate of appearance of [ $^{14}\text{C}$ ]glucose was calculated in exactly the same way as that of glucose, i.e. using the MCR determined from the infused glucose label. This rate of [ $^{14}\text{C}$ ]glucose appearance represents the transfer of labelled carbon from circulating lactate to glucose. To convert it to a mass rate, it was divided by the interpolated lactate specific activity at each time point, yielding a gluconeogenetic rate in  $\mu\text{mol kg}^{-1} \text{min}^{-1}$ . Since this is neither corrected for dilution of label by carbon from the tricarboxylic acid cycle [20], nor accounts quantitatively for all substrates, it is considered an index of gluconeogenesis.

Comparisons of (hormone, metabolite, flux) time course data between diabetic patients and control subjects were made using ANOVA with time as a repeated measure. The Greenhouse–Geisser correction was used to adjust the  $p$  values. Contrast analysis was used to compare potential differences in the time profiles of the concentration or flux data between the two groups. Anthropometric data and extracted parameters were compared using  $t$ -tests, after testing for the equality of variances. One-tailed tests were used where appropriate. An autoregressive moving average model was used to compute cross-correlations between variables (melatonin, cortisol and EGP) that displayed cyclic behaviour, after prewhitening with a filter based on modelling the predictor variable. A cross-correlation function was determined for different lag times in each experiment. SAS (SAS Institute, Cary, NC, USA) software was used. Results are presented as means $\pm$ SEM, and flux rates are calculated per kg of body weight.

## Results

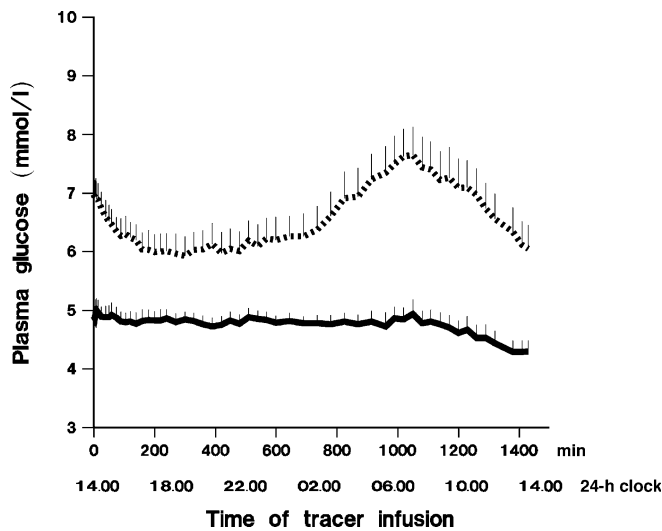
**Subjects** The clinical characteristics of the subjects are summarised in Table 2. All polysomnogram results, both before and during the studies, were within normal limits in the patients. In particular, there was no evidence of abnormal periods of sleep apnoea.

**Glycaemia** In diabetic subjects, glucose concentrations (Fig. 1) initially decreased from  $7.2\pm 0.3$  mmol/l and plateaued near 6.3 mmol/l for several hours. By midnight (600 min), concentrations had started to increase, and rose continuously until 07.00 h (1,020 min), when they peaked at  $8.0\pm 0.5$  mmol/l, gradually decreasing thereafter to  $6.3\pm 0.4$  mmol/l by the end of the study. In contrast, control subjects showed a stable glucose concentration ( $4.9\pm 0.2$  mmol/l) until near the end of the study, when a drift to about 4.4 mmol/l occurred. The two glycaemic curves are significantly different ( $p=0.0003$ ). Contrast analysis of datapoints against a baseline level shows that the profiles of the two curves diverge significantly ( $p<0.05$ ) at  $t=825$  and remain different until  $t=1,410$  min.

**Fluxes** The MCR of glucose (Table 3) was near constant and averaged  $1.73\pm 0.10$  ml  $\text{kg}^{-1} \text{min}^{-1}$  in control subjects and  $1.40\pm 0.14$  ml  $\text{kg}^{-1} \text{min}^{-1}$  in diabetic patients ( $p=0.04$ ). There was a gradual rise in EGP, from  $8.2\pm 0.7$  to  $11.3\pm 0.5$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$  over the course of the night (Fig. 2, Table 3), peaking near 940 min ( $\sim 06.30$  h) in diabetic patients. This is in contrast to control subjects, where EGP remained near constant at  $8.2\pm 0.4$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$ , before drifting down to  $7.2\pm 0.2$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$ , but only after  $\sim 24$  h of fasting. The time course of EGP is different in the two groups ( $p=0.0017$ ), with contrast analysis showing a significant ( $p<0.05$ ) divergence between  $560 < t < 1,100$  min (23.30 and 08.30 h on day 2). The pattern of glucose removal in diabetic subjects ( $R_d$ ; Fig. 2) was similar to that for glucose, peaking ( $10.5\pm 0.4$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$ ) at 07:30 h from a baseline value of  $8.5\pm 0.6$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$ . The time course was different in control subjects ( $p=0.0017$ ), who maintained a removal rate near  $8.2\pm 0.3$   $\mu\text{mol kg}^{-1} \text{min}^{-1}$  for most of the study. The two

**Table 2** Subject characteristics

	Control subjects	Type 2 diabetic patients	$p$ value
$n$	8	8	
Age (years)	49.0 $\pm$ 3.5	53.4 $\pm$ 3.8	0.4
Sex (male/female)	4/4	5/3	
Weight (kg)	79.7 $\pm$ 4.0	88.6 $\pm$ 5.4	0.2
BMI	28.8 $\pm$ 1.2	29.7 $\pm$ 1.5	0.6
HbA <sub>1c</sub>	0.052 $\pm$ 0.001	0.071 $\pm$ 0.006	<0.002
Fasting plasma glucose (mmol/l)	5.4 $\pm$ 0.2	8.0 $\pm$ 0.6	<0.002
Time since diagnosis (years)	NA	5.3 $\pm$ 1.7 (0.5–15)	



**Fig. 1** Glucose concentrations in fasted subjects with type 2 diabetes mellitus (*dashed line*) compared with control subjects (*solid line*). The *abscissa* is given both as the tracer infusion time and the corresponding nominal time on a 24-h clock

curves diverged significantly ( $p < 0.05$ ) after  $t = 600$  min (midnight) until the end of the study. The mismatch between EGP and  $R_d$  explains the diurnal rise and fall of glucose concentrations in type 2 diabetes.

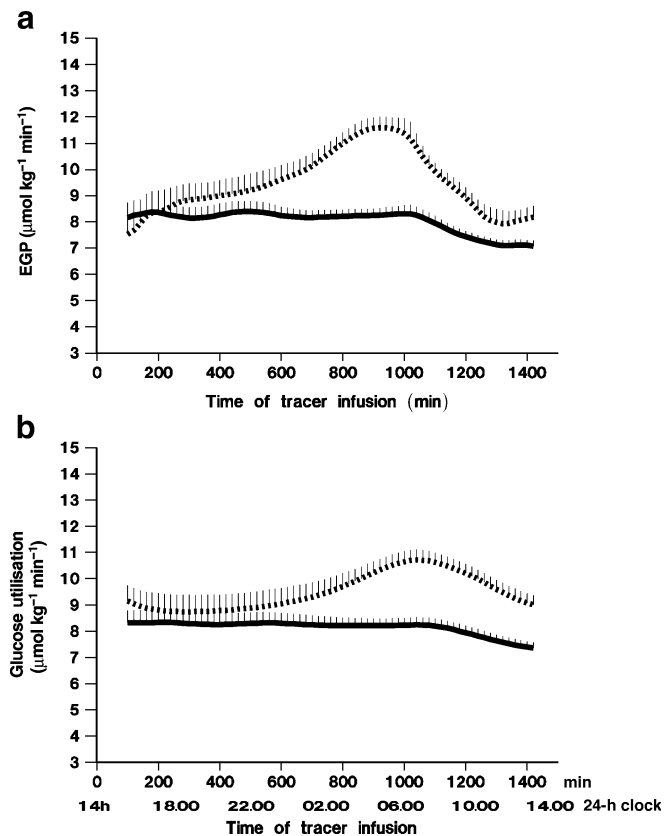
An index of gluconeogenesis (Fig. 3) was obtained from the rate of incorporation of labelled carbon from [ $^{14}\text{C}$ ] lactate into glucose, which was infused simultaneously in these studies. It increased significantly with time in both

groups ( $p < 0.0001$ ), representing the well-known rise with fasting. This increase was about 45% from the beginning to the end of the study (24 h of fasting). Contrast analysis further demonstrated differences in profile between the two groups that were significant at the  $p < 0.05$  level from 840–1,020 min. This difference translates into a peak in gluconeogenesis when the control gluconeogenic time course was subtracted from that in diabetic subjects (Fig. 3). This peak, superimposed on a rising baseline, parallels the EGP during its nocturnal rise and morning fall, and strongly suggests that gluconeogenesis is a major contributor to the nocturnal increase in EGP. Concentrations of lactate (Fig. 3), the major glucogenic substrate, were elevated in diabetes ( $p = 0.0023$ ), but neither of the two groups showed an interaction with the time course ( $p = 0.2$ ) that might reflect the additional nocturnal rise in EGP. The concentrations of NEFA (Fig. 3), a promoter of gluconeogenesis, were not different between the two groups ( $p = 0.4$ ).

**Hormones** Figure 4a–c shows glucagon and insulin concentrations and the ratio of the two. Glucagon levels remained near constant throughout the study in both groups ( $p = 0.06$ ), but were elevated in diabetes ( $p = 0.04$ ). The differences in insulin concentrations between diabetic and control subjects did not reach significance, but the levels fell over time in both groups ( $p = 0.0007$ ). This fall was translated into a rising glucagon:insulin ratio ( $p = 0.006$ ), which was otherwise not distinguishable between the two groups ( $p = 0.8$ ). Leptin concentrations showed variation among individual subjects, but a similar

**Table 3** Details of flux calculations

Experiment number	Volume of distribution (l)	Mean MCR (ml kg <sup>-1</sup> min <sup>-1</sup> )	EGP (μmol kg <sup>-1</sup> min <sup>-1</sup> )					
			240 min	480 min	720 min	960 min	1,200 min	1,440 min
Control subjects								
1	17.7	1.86	8.35	8.26	8.54	8.67	7.63	7.14
2	21.3	1.46	8.12	7.90	7.88	8.50	7.65	6.98
3	27.9	2.05	9.19	9.39	8.58	8.86	8.03	7.61
4	31.8	1.71	8.79	9.32	8.66	9.22	7.67	7.60
5	26.4	1.59	7.52	7.57	7.47	7.48	6.77	6.69
6	19.9	1.88	8.68	8.94	8.66	8.86	8.11	7.79
7	14.7	2.06	9.47	9.57	8.91	7.96	7.28	7.06
8	30.7	1.25	5.94	6.18	6.71	6.77	6.37	5.97
Mean	23.8	1.73	8.26	8.39	8.18	8.29	7.44	7.10
SEM	2.2	0.10	0.40	0.41	0.27	0.29	0.21	0.21
Type 2 diabetic subjects								
1	37.5	1.49	7.28	7.92	9.24	9.08	5.45	6.84
2	30.9	1.33	8.94	10.18	11.08	11.95	8.31	8.31
3	27.0	1.10	9.10	10.00	10.01	10.96	8.72	8.22
4	25.1	2.23	8.71	8.50	11.48	12.43	9.76	7.41
5	38.3	1.46	9.43	8.87	10.23	10.65	9.36	8.65
6	24.6	0.86	5.86	6.42	8.32	9.65	6.91	6.46
7	20.6	1.61	11.57	11.70	11.79	12.25	10.11	9.90
8	39.3	1.17	6.52	8.44	9.24	12.92	9.26	8.01
Mean	30.4	1.41	8.43	9.00	10.17	11.24	8.49	7.97
SEM	2.5	0.15	0.64	0.57	0.43	0.49	0.56	0.38



**Fig. 2** EGP (a) and glucose utilisation (b) rates in subjects with type 2 diabetes (dashed line) and control subjects (solid line), calculated over the time course of the study using non-steady-state compartmental analysis

overall decrease was seen in the two groups ( $p < 0.0001$ ) during the 24-h study period. The rate of decrease tended to be faster in control subjects (first-order polynomial contrast  $p = 0.08$ ). Cortisol and melatonin both displayed a cyclicity (changes with time  $p = 0.006$  for cortisol and  $p = 0.02$  for melatonin) similar to that in EGP. The time course for cortisol levels was indistinguishable between the diabetic and control groups ( $p = 0.1$  and contrast analysis). However, the rise of EGP in diabetic subjects precedes the increase in cortisol by  $200 \pm 60$  min, and the fall in EGP is not accompanied by a corresponding decrease in cortisol. This is consistent with an average cross-correlation function for which the absolute value is less than 0.15 across the spectrum of lag times. Although there is individual variation, the increase in melatonin appears attenuated and delayed in diabetes (vs control subjects). Differences between the two groups are seen for an eighth order polynomial contrast when all the data are used ( $p = 0.001$ ), and for a cubic when only data up to  $t = 800$  are used ( $p = 0.01$ ). This is consistent with the near-constant and similar time course of the two curves at the beginning and end of the study, and the night-time excursion where differences between them occur. Cross-correlation analysis between melatonin levels and EGP demonstrated an average of 0.4 at zero lag time and less

than 0.1 for all other lag times, suggesting a degree of synchrony between the two curves.

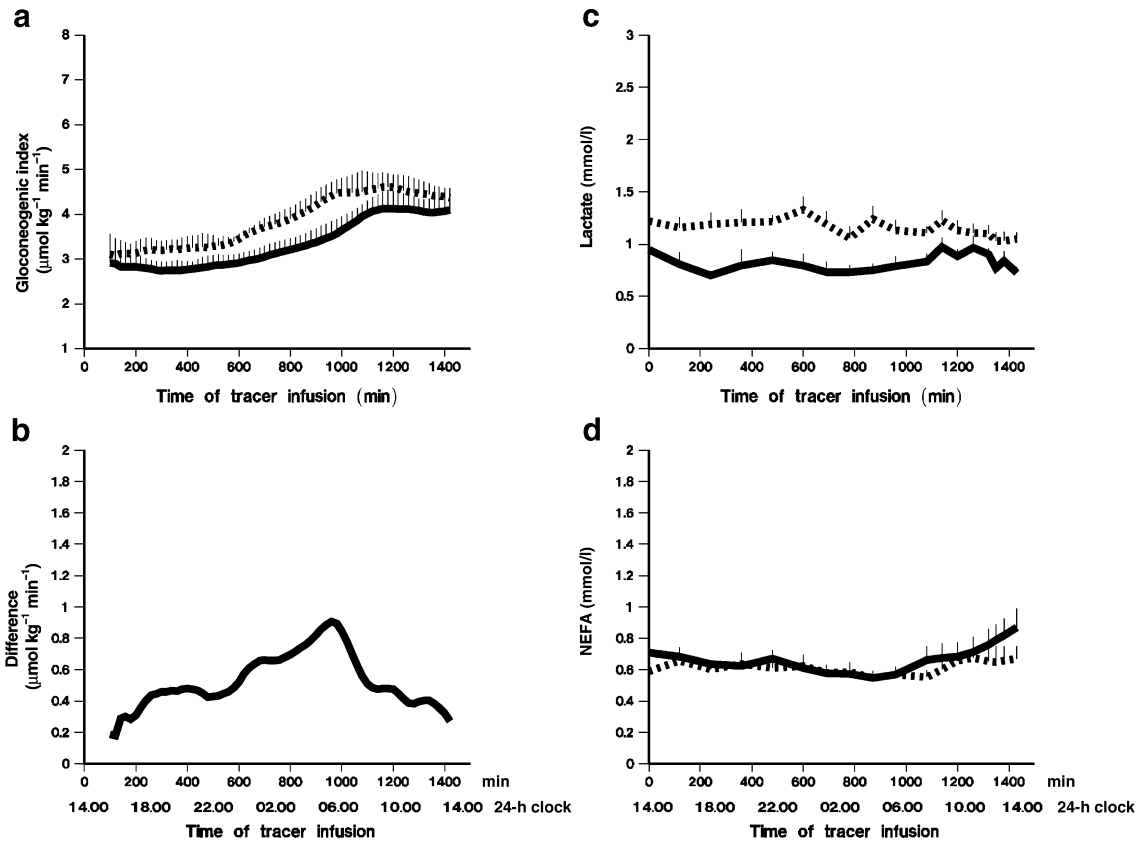
## Discussion

This study demonstrates a spontaneous diurnal variation of glucose concentrations in type 2 diabetes, with a gradual transition from a nadir in the late evening to a peak in the morning. Such phasic behaviour was not detectable in healthy control subjects. It is, moreover, driven by parallel changes in EGP, specifically, gluconeogenesis. The results answer the question of the provenance of the morning increase in EGP, and a significant portion of the hyperglycaemia. Juxtaposition with studies demonstrating a repeated cycle in EGP during a near isoglycaemic clamp in diabetes [21] over a 3-day period, strongly suggest that the 24-h pattern of EGP can be extended to conditions where the EGP provides the hyperglycaemic drive. Since subjects were fasting from the previous morning, it is highly unlikely that the morning fasting hyperglycaemia depends on preceding meals. The lack of a detectable increase in insulin secretion in response to the nocturnal rise in glycaemia further accentuates this rise, and emphasises the glucose insensitivity of insulin secretion (which might also undergo diurnal exacerbations) in the diabetic state.

These results can be explained by:

1. *A circadian rhythm* in EGP, probably driven by the suprachiasmatic–paraventricular axis and mediated by the autonomic nervous system and perhaps by hormones such as cortisol; and
2. *Limit cycle behaviour*: a conversion from normal to diabetes, characterised by a transition from a stable steady state to another stable but cyclic state (of EGP and glycaemia), in a metabolic system that is inherently complex and non-linear.

*Circadian rhythms in glucose metabolism and the suprachiasmatic nucleus* Studies on variations in glucose levels have produced a range of results. One study reported diurnal variations in the concentrations and fluxes of glucose and other metabolites in diabetic, but not control, subjects [22]; another found elevated morning glucose levels in both diabetic and control subjects, although these were significantly attenuated in the latter group [23]; and still others observed no additional increase in glycaemia in overnight-fasted diabetic subjects [24, 25]. These nocturnal increases in glucose may be related to the dawn phenomenon [26, 27], although this usually starts later than the rise in EGP described here and is likely related to postprandial counter-regulatory effects. The precise glycaemic profile seen could therefore depend on the preceding meal schedule and content [28]. The observations in the present study, however, are independent of meals because of the extended fasting period. Daily variations in metabolism have been related to the hypothalamic biological clock [29, 30]. In rodents, a rise



**Fig. 3** **a** Gluconeogenic index calculated from the incorporation of  $^{14}\text{C}$  from [ $^{14}\text{C}$ ]lactate into glucose. Both control (*solid line*) and diabetic (*dashed line*) groups demonstrate a gradual increase in the gluconeogenic index as fasting progresses although this occurs more rapidly in diabetes. **b** Subtraction of the control index from that

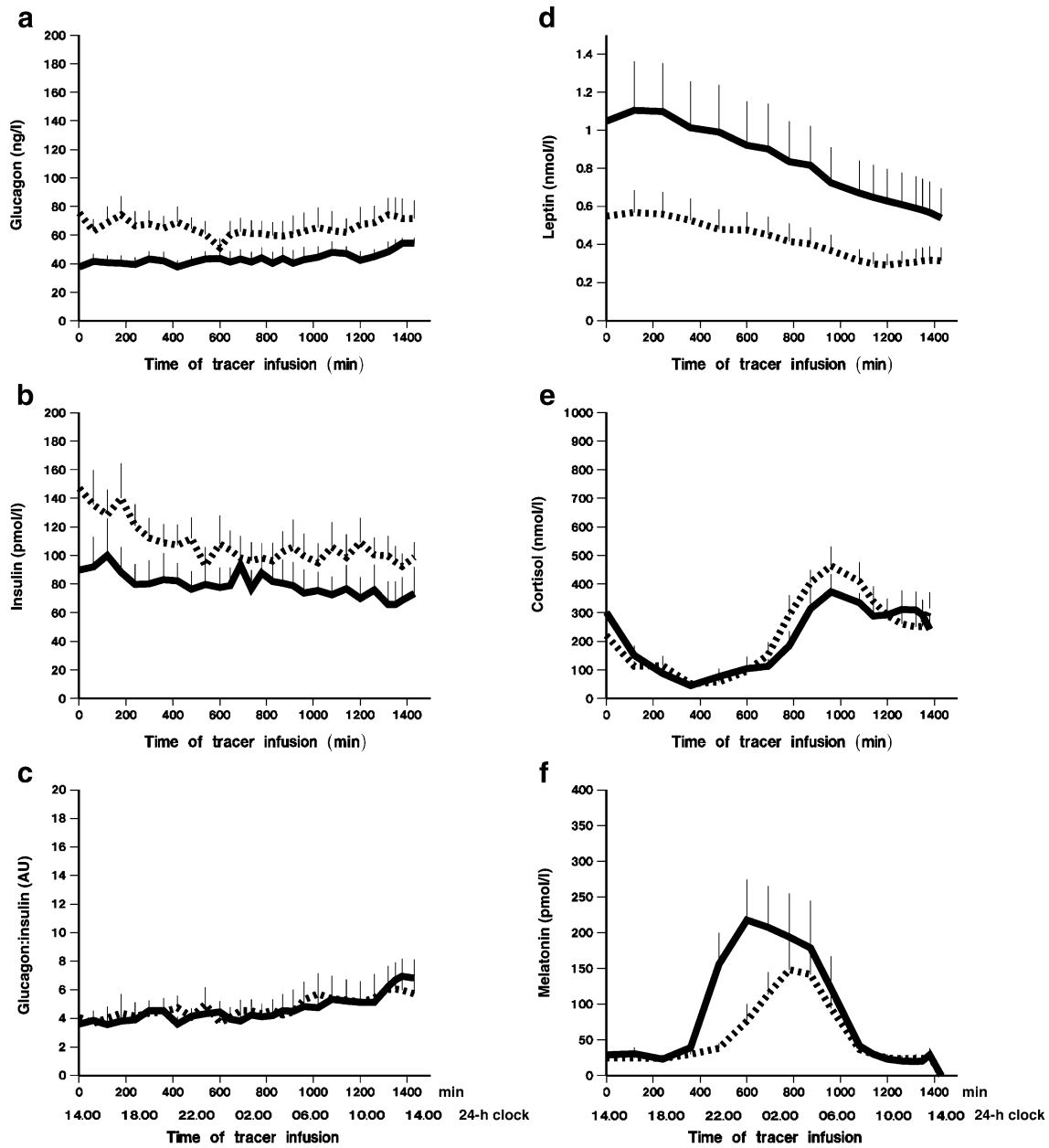
obtained in diabetes reveals a superimposed excursion in the gluconeogenic index in diabetes that matches that in EGP. Lactate concentrations (**c**) and free fatty acid levels (**d**) over the 24-h period in control (*solid line*) and diabetic (*dashed line*) subjects

in plasma glucose concentrations at dusk (the end of the sleep period) has been shown to result from a rise in EGP [31, 32]. Elegant studies in mice demonstrated, that this rise in EGP was generated via the suprachiasmatic–paraventricular pathway inducing a sympathetic stimulation of EGP [33]. This neural pathway resembles that controlling melatonin secretion: photic information is conveyed from the retina to the suprachiasmatic nucleus (SCN), relayed through the paraventricular nucleus and a multisynaptic pathway to the pineal gland, which secretes melatonin [34], providing the rationale for using melatonin as a surrogate measure for SCN activity in our studies. Although causality cannot be established by analysis of correlations, changes in melatonin concentrations in the present studies are correlated with the changes in EGP in diabetic subjects, suggesting the possibility that this flux, too, may be regulated by the biological clock resident in the SCN. Interestingly, however, the nocturnal excursion of melatonin in the diabetic subjects is postponed and attenuated relative to that in control subjects (Fig. 4), raising the possibility that, unlike the case in rodents, the SCN may be involved in suppressing a cyclic nocturnal increase in EGP, which is then unmasked in diabetes. Parenthetically, the decrease of melatonin itself in diabetes may contribute to the metabolic syndrome, since its ad-

ministration in rats decreased body weight and suppressed visceral fat and plasma leptin [35].

Since SCN activity appears diminished in diabetes, a straightforward linkage of the cycle in EGP (and glycaemia) to the hypothalamic biological clock appears less likely. Furthermore, such an association in rodents relates to arousal and occurs at dusk (dawn phenomenon in humans), with a shorter lived cycle than that seen in our study. It occurs in normal animals, rather than being restricted to diabetes as seen here. The clock-related rise in EGP is also caused by glycogenolysis [36], which is controlled by the sympathetic innervation to the liver [33, 37], whereas gluconeogenesis plays an important part in the increased glucose flux in our study. Although a deficit in SCN activity in type 2 diabetes may explain the changes seen, it is also possible that the circadian association of EGP may be, to some extent, fortuitous.

The rodent model described above [33] also demonstrated peaks in glucocorticoid levels, glucagon or adrenaline, which, although parallel to the glucose excursions, were not responsible for the hyperglycaemia [33]. In our study, only cortisol followed a diurnal pattern (Fig. 4). Its nocturnal rise has been implicated in glucoregulation [38]. Our results show a phase delay with cortisol following increases in both melatonin and EGP (Figs. 2,



**Fig. 4** Hormone levels over the course of the 24-h fast in patients with type 2 diabetes (*dashed line*) and control subjects (*solid line*). The panels show the levels of glucagon (a) and insulin (b), the glucagon:insulin ratio (c), and the leptin (d), cortisol (e) and melatonin (f) concentrations

4), obviating a possible role for cortisol in the stimulation of the nocturnal rise in glucose production in diabetes. This is consistent with the lack of correlation of cortisol with EGP also seen here, and with the observation that suppression of corticosterone synthesis did not alter morning hyperglycaemia in diabetes [39].

*Limit cycle behaviour: is type 2 diabetes a dynamical disease?* It is possible that, in diabetes, insulin resistance (for example) might unmask a more fundamental regulation arising from the SCN. However it remains puzzling why the SCN does not appear to influence glucose fluxes in normal subjects. The need for an alternative explanation is therefore

suggested. Variations in EGP have also been seen over a 72-h period in the context of a glycaemic clamp to 11 mmol/l (near isoglycaemic) in type 2 diabetes [21]. The present study demonstrates that the diurnal changes in EGP drive the corresponding changes in glucose concentrations under physiological fasting conditions. The combination of the two results suggests the stable nature of the cyclic changes in glucose and a degree of autonomy in this rhythm, which would be characteristic of a system property, rather than a response to a regulatory drive.

The existence of two stable states for EGP (and glycaemia), one of which represents a disease state (type 2 diabetes), is highly reminiscent of the description of a

'dynamical' disease. The concept arises from the generally complex, non-linear nature of many biological systems—a description applicable to the metabolic system. Under appropriate circumstances, the behaviour of such systems can be characterised by the coexistence of dual (or even multiple) stable states. For example, either a steady state (e.g. near-constant glycaemia) or a limit cycle (e.g. a diurnal variation in glycaemia) may occur [40]. When a key control parameter (e.g. an enzyme activity) achieves a critical value, or an appropriate perturbation takes place, the system may pass through a 'bifurcation' or 'switching' point and alter its stable state. Such transitions between stable states are characteristic of dynamical diseases [41, 42] such as Cheyne–Stokes respiration [43] or aplastic anaemia [44]. Interestingly, the onset of type 1 diabetes has been described as a collective dynamic instability within a complex framework [45], rather than a single aetiological factor.

This also appears to be what happens in type 2 diabetes: patients with a (genetic) predisposition to diabetes (and therefore closer to a switching point) may pass from a steady state (constant and normal fasting EGP and glycaemia) to a limit cycle solution (with cyclic changes in EGP and glycaemia and thus elevated morning fasting glucose concentrations) in response to an environmental or lifestyle challenge.

1. Such a cyclic state, which is then necessarily restricted to diabetes, is demonstrated in our study, while explaining an important part of fasting hyperglycaemia.
2. It occurs at the level of gluconeogenesis, which is integrated into the intermediary metabolic network and thus likely to exhibit complex behaviour.
3. Since the transition to the cyclic state represents a 'switch' between two system behaviours, it would be expected to occur rapidly rather than gradually. Indeed, it has been shown that fasting glycaemia does increase abruptly (within a 3-year timeframe) at the onset of diabetes [46]. These complementary data strengthen the argument for non-linear system behaviour. In turn, the transition to a limit cycle provides an explanation for the rapidity of onset of fasting hyperglycaemia.
4. Although it can be influenced by hormones and the central nervous system, the cyclic state is dependent on the entire system and therefore behaves autonomously, explaining both the refractoriness to meal-related regulation [47] and the inexorable clinical progression of diabetes, which is only delayed by treatment (e.g. [48]).

*Overall contribution of fluxes to glycaemia* In diabetes, glucose concentrations do not fall to normal levels after the nocturnal excursion has dissipated (Fig. 1). In this group of subjects, the increase over normal at the glycaemic nadir was 1.5 mmol/l and the increase over this baseline at peak morning values was 1.7 mmol/l. Since EGP is normal at the nadir (Fig. 2), this indicates that at peak values, the decrease in metabolic clearance and the rise in glucose production

respectively account for 47% and 53% of the increase in glycaemia. This is consistent with previous findings (e.g. [7, 49]). Interestingly, when glucose concentrations were at their lowest in the diabetic subjects, both EGP and  $R_d$  were matched and normal (the same as control subjects; Fig. 2), and consistent with a normal glucagon:insulin ratio. The only defect at this point is the low MCR, which necessitates an increase in glycaemia to maintain  $R_d$ . The increase in EGP, occurring near midnight, then initiates the cycle of increased glycaemia and  $R_d$ , which allows the return to nadir conditions once EGP decreases.

The gluconeogenic index demonstrates a dual time course. The first concerns the expected increase of about 50% with fasting, which occurs in both control and diabetic subjects. The value at 22 h of fasting in control subjects was approximately  $3.8 \mu\text{mol kg}^{-1} \text{min}^{-1}$ . If a correction factor (for label dilution in the tricarboxylic acid cycle) of  $\sim 1.5$  is used [20], gluconeogenesis is 70% of EGP ( $8.1 \mu\text{mol kg}^{-1} \text{min}^{-1}$ ), almost exactly the same number as found using the deuterated water method in normal subjects [14]. This increase is compensated by a decrease in glycogenolysis (autoregulation) [50]. The second time course involves a cyclic increase in diabetes that corresponds to the rise in EGP, discerned from the difference in the gluconeogenic index between control and diabetic subjects. This is not compensated for by a decrease in glycogenolysis, suggesting a different aetiology for the two gluconeogenic fluxes.

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## Conclusion

In conclusion, we have found that EGP in type 2 diabetes is never in steady state, displaying rather a cyclic (diurnal) temporal pattern. This phasic behaviour is restricted to diabetes and is not seen in normal control subjects. It is most closely correlated with melatonin levels and therefore SCN activity, although these are decreased in diabetes, suggesting the possibility of reduced suppression of EGP, a process that might normally be mediated by the SCN. The cyclic behaviour of EGP determines the periodic pattern of glycaemia in diabetes and is, at least partly, determined by the same pattern in the rate of gluconeogenesis, suggesting that this behaviour is fundamentally related to intermediary metabolism. These changes could therefore reflect the intrinsic non-linear behaviour of (primarily hepatic) metabolism that determines glucose levels, but which could also subsume central regulatory pathways. The transition from the normal to the diabetic state would then be characterised by the switch in EGP from steady-state behaviour to the stable cyclic behaviour, demonstrated here.

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