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Dynamic insulin sensitivity index: importance in diabetes

Gianluigi Pillonetto, Andrea Caumo, and Claudio Cobelli

¹Dipartimento di Ingegneria dell'Informazione, Università degli Studi di Padova, Padova; and ²San Raffaele Scientific Institute, Milan, Italy

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Pillonetto G, Caumo A, Cobelli C. Dynamic insulin sensitivity index: importance in diabetes. Am J Physiol Endocrinol Metab 298: E440-E448, 2010. First published November 17, 2009; doi:10.1152/ajpendo.90340.2008.—The classical minimal model (MM) index of insulin sensitivity, S_I, does not account for how fast or slow insulin action takes place. In a recent work, we proposed a new dynamic insulin sensitivity index, S₁^D, which is able to take into account the dynamics of insulin action as well. The new index is a function of two MM parameters, namely S_I and p_2 , the latter parameter governing the speed of rise and decay of insulin action. We have previously shown that in normal glucose tolerant subjects S^D provides a more comprehensive picture of insulin action on glucose metabolism than S_I . The aim of this study is to show that resorting to S_I^D rather S_I is even more appropriate when studying diabetic patients who have a low and slow insulin action. We analyzed insulinmodified intravenous glucose tolerance test studies performed in 10 diabetic subjects and mixed meal glucose tolerance test studies exploiting the triple tracer technique in 14 diabetic subjects. We derived both S_I and S_I^D resorting to Bayesian and Fisherian identification strategies. The results show that $S_1^{\rm D}$ is estimated more precisely than S_I when using the Bayesian approach. In addition, the less laborintensive Fisherian approach can still be used to obtain reliable point estimates of S₁^D but not of S₁. These results suggest that S₁^D yields a comprehensive, precise, and cost-effective assessment of insulin sensitivity in subjects with impaired insulin action like impaired glucose tolerant subjects or diabetic patients.

model; insulin resistance; parameter estimation; Bayesian estimation; Markov chain Monte Carlo strategy; diabetes

THE MINIMAL MODEL (MM) of glucose kinetics has been used in nearly 600 papers since its inception in the late seventies (4). The reason for such popularity is related to the ability of MM to provide an index of insulin sensitivity, denoted by $S_{\rm I}$, from the analysis of an intravenous glucose tolerance test (IVGTT). An extension of this model is being currently used also to determine $S_{\rm I}$ from an oral/meal glucose tolerance test (OGTT/MGTT; Ref. 8).

Albeit of paramount importance, it must be recognized that $S_{\rm I}$ does not yield an exhaustive picture of insulin action. The reason can be more easily grasped by making reference to the assessment of insulin sensitivity with the gold standard method, which is the euglycemic hyperinsulinemic clamp. The clamp-based insulin sensitivity, of which the MM-based $S_{\rm I}$ is a theoretical counterpart, is measured at the end of the clamp when insulin action is at steady state. However, in those clamp studies that monitored in detail the time course of insulin action from the initial to the final steady state, it became apparent that the speed of rise of insulin action varied considerably among individuals and that such speed was related to the degree of

Address for reprint requests and other correspondence: C. Cobelli, Dipartimento di Ingegneria dell'Informazione, Università degli Studi di Padova, Via Gradenigo, 6/B-35131 Padova, Italy (e-mail: cobelli@dei.unipd.it).

insulin sensitivity measured at steady state (15). This observation indicates that when judging on the individual's ability to dispose glucose, not only is the maximum excursion of insulin action relevant, but also the dynamics of insulin action plays a role. If this is true during a clamp, a fortiori is true during an IVGTT in which insulin action never attains a plateau and its speed of rise and decay certainly influence the return of glucose to the baseline. For this reason, we proposed a new insulin sensitivity index that, at variance with the classic one, also accounts for the dynamics of insulin action. The new index, denoted by S₁^D, is able to account for both the speed and the capacity of response. Thanks to this peculiarity, in Ref. 17 we have shown that in normal glucose tolerant (NGT) subjects $S^{\mathrm{D}}_{\scriptscriptstyle{\tau}}$ provides, compared with S_I, a more comprehensive picture of insulin action on glucose metabolism. In addition, S₁^D is intrinsically more robust than S_I, which makes its identification easier and more reliable with respect to S_I.

The aim of the present study is to compare the estimation of S_I and S_I^D in diabetic patients and verify whether in these subjects the assessment of S₁^D produces even greater benefits than in NGT subjects. There are two reasons for this to happen. The first one is related to the notion that not only are diabetic patients insulin resistant, but, according to the above-mentioned euglycemic hyperinsulinemic studies, their insulin action develops at a slower pace than in NGT subjects (1, 15). As a result, in diabetic patients $S_{\scriptscriptstyle I}^D$ should be more capable than $S_{\scriptscriptstyle I}$ to provide a realistic picture of how a diabetic patient is able to handle the glucose challenge. Moreover, S_{I}^{D} should be more accurate than S_I in ranking diabetic patients. The second reason why S_I^D is expected to perform better than S_I in diabetic patients is related to the numerical identifiability of the MM in such patients. It is well known that in diabetics the MM identification is difficult because it is often associated with large parameters uncertainties (see Refs. 1, 18, 19, 21). The greater robustness of $S_{\scriptscriptstyle I}^{\scriptscriptstyle D}$ with respect to $S_{\scriptscriptstyle I}$ should ensure that S₁^D is precisely estimated even in those patients in which the estimation of S_I is difficult or impossible.

MATERIALS AND METHODS

Estimates of S_I and S_I^D were obtained in diabetic patients under two different experimental conditions: an insulin-modified IVGTT and a MGTT. The IVGTT studies were performed in 10 type 2 diabetic patients whose characteristics have been already described in Refs. 1 and 19. Briefly, the IVGTT consisted of a glucose dose of 300 mg/kg injected at *time 0* followed by a short insulin infusion (0.05 U/kg) administered between 20 and 25 min. The MGTT studies were performed in 14 type 2 diabetic subjects who underwent a mixed meal test in which three tracers were simultaneously infused to allow an accurate determination of the rate of appearance of the orally administered glucose into the circulation (see Ref. 3 for details about the experimental protocol and the subjects' characteristics). The protocol was approved by The Mayo Clinic Institutional Review Board.

To compare the ability of $S_{\rm I}$ and $S_{\rm I}^{\rm D}$ to discriminate between diabetic and NGT subjects, we performed a simulation study. Estimates of $S_{\rm I}$ and $S_{\rm I}^{\rm D}$ were obtained from synthetic insulin modified IVGTT experiments performed in 1,000 diabetic and 1,000 NGT subjects. Synthetic data were generated using two log-normal priors for MM parameters. The first prior was obtained from the estimates in diabetic subjects reported in Table 1, while the second prior was derived from the estimates obtained in 10 NGT individuals in a previous study (details are reported in Ref. 17). Then, the typical insulin profile that can be observed during an insulin-modified IVGTT in a diabetic subject was used as the insulin input to the model. We sampled all the generated glucose profiles at 5, 8, 10, 12, 15, 20, 25, 30, 40, 60, 80, 100, 120, 140, 160, 180, and 240 min and finally corrupted them by white normal noise with a coefficient of percentage variation equal to 2.

To validate the new insulin sensitivity index, we used correlation analysis to compare the estimates of $S_i^{\vec{D}}$ obtained in 21 subjects [11 impaired glucose tolerant (IGT) and 10 NGT] undergoing both an OGTT and a euglycemic hyperinsulinemic clamp. Details about the experimental procedures are reported in the Methods in Ref. 9. Here, we just recall that, during the clamp, regular human insulin was infused at a constant rate (25 mU·m⁻²·min⁻¹) for 180 min. A hyperinsulinemic plateau level was achieved within 30-45 min from the initiation of the insulin infusion. Due to the natural variability of the insulin clearance among subjects, the hyperinsulinemic plateau level differed slightly among individuals (means \pm SD close to 50 and 10 μU/ml, respectively). However, the clamp-based index of insulin sensitivity $[S_I(clamp)]$ and the new $S_I^D(clamp)$, as well as the MMbased indices S_I and S_I^D, have the intrinsic ability to properly account for different hyperinsulinemic levels among individuals (e.g., see Eqs. 2 and 5).

MM and derivation of dynamic insulin sensitivity. In the following we refer, without loss of generality, to the MM equations describing glucose disappearance during an IVGTT. The equations describing glucose disappearance during an MGTT are the same with the only difference that the exogenous glucose input is the rate of appearance of orally administered glucose that has been accurately estimated thanks to the triple-tracer technique (see Ref. 3 for details). MM equations are as follows:

$$\dot{G}(t) = -(p_1 + X(t))G(t) + p_1G_b \quad G(0) = G_0$$
 (1)

$$\dot{X}(t) = -p_2 X(t) + p_3 (I(t) - I_b) \quad X(0) = 0$$
 (2)

where G(t) (mg/dl) and I(t) (μ U/ml) are glucose and insulin concentrations in plasma, respectively, while G_b and I_b are their baseline values. *Equation 1* represents glucose kinetics, whereas *Eq. 2* describes insulin action exerted from a compartment remote from plasma. Uniquely identifiable model parameters are p_1 , p_2 , p_3 , and G_0 . Parameters of interest provided by the model are glucose effectiveness ($S_G = p_1$, min⁻¹), which measures the effect of glucose per se on both

glucose disappearance and endogenous glucose production, and insulin sensitivity ($S_I = p_{3/p_2}$, min⁻¹ $\mu U^{-1}ml$), which measures insulin's ability to enhance glucose effectiveness by both increasing glucose disappearance and inhibiting endogenous glucose production. Parameter p_2 is the time constant of the remote insulin compartment and thus governs the speed of rise and decay of insulin action. Parameters S_1 and p_2 can be used to derive a dynamic index of insulin sensitivity, as already demonstrated in Ref. 17. For the sake of clarity and completeness, we will briefly state without proof the steps leading to the derivation of S₁^D. The first step consists of obtaining a closed form of the MM glucose prediction, as a function of model parameters and insulin profile. This aim can be accomplished by noticing that Eq. 2 can be integrated independently from $Eq. \ 1$. We use Z(t) to denote the integral function of remote insulin. We see from Eq. 2 that the impulse response relating the deviation of insulin from basal and X(t) is $p_3 e^{-p_2 \sigma} =$ $S_I p_2 e^{-p_2 \sigma}$ for $\sigma \ge 0$ and zero otherwise. Thus Z(t) is the output of a system with impulse response

$$\int_{0}^{t} S_{I} p_{2} e^{-\sigma p_{2}} d\sigma = -S_{I} e^{-\sigma p_{2}} \Big|_{0}^{t} = S_{I} (1 - e^{-p_{2}t}), i.e.$$

$$Z(t) = \int_{0}^{t} X(\sigma) d\sigma = S_{I} \int_{0}^{t} (1 - e^{-p_{2}(t-\sigma)}) (I(\sigma) - I_{b}) d\sigma$$
(3)

From a standard result on linear differential equations (see Ref. 6) glucose prediction is given by

$$G(t) = G_0 e^{-S_G t - Z(t)} + S_G G_b \int_0^t e^{-S_G (t - \tau) - Z(t) + Z(\tau)} d\tau$$
 (4)

Equation 4 allows one to appreciate that, according to the MM, glucose concentration is a function of two signals, i.e., $S_G t$ and Z(t). The term $G_0e^{-S_Gt-Z(t)}$ is a single decaying exponential with a coefficient proportional to the sum of these two signals. In particular, the first of these two signals, i.e., S_Gt, is linearly dependent on glucose effectiveness, S_G . The second signal, i.e., Z(t), describes the ability of insulin to enhance glucose effectiveness and depends in a more complex way on both S_I and p_2 . In fact, from Eq. 3 one can see that Z(t) is the output of a time-invariant linear system having $I(t) - I_b$ as input and $h(t) = S_I(1 - e^{-p_2 t})$ as unit impulse response. This impulse response, denoted by integrated insulin action impulse response (IAIR; Refs. 17, 18), describes the modalities by which incremental (above basal) insulin concentration is able to influence the glucose profile. Dynamic insulin sensitivity, S₁^D, is defined as the mean value of the IAIR in a time interval of duration T (how T is chosen will be clarified in a moment). In mathematical terms, we have:

$$S_{I}^{D} = \frac{\int_{0}^{T} S_{I}(1 - e^{-p_{2}t})dt}{T} = S_{I} \left[1 - \frac{1 - e^{-p_{2}T}}{p_{2}T} \right]$$
 (5)

It can be verified that S_1^D is always less or equal to S_1 . Thus S_1^D can be conveniently expressed as a fraction, comprised between 0 and 1, of S_1 . Such fraction is denoted as efficiency:

Table 1. IVGTT in diabetic subjects: Bayesian estimates of minimal model parameters

Subject Nos.	$S_{\rm I}, 10^{-4} \; min^{-1} \mu U^{-1} ml$	p_2 , 10^{-2} min^{-1}	$S_{I}^{D},\ 10^{-4}\ min^{-1}\mu U^{-1}ml$	S_G , $10^{-2} min^{-1}$	Go, mg/dl
1	1.2 (0.61–2.7)	1.3 (0.13–3.3)	0.3 (0.09–0.65)	1.59 (1.07–1.98)	299.7 (284.1–313.3)
2	1.5 (1.4–1.6)	7 (5.7–84)	1.17 (1.1–1.23)	2.1 (1.8–2.38)	382 (369.9–394.7)
3	1.7 (0.4–4.1)	0.18 (0.04-0.68)	0.06 (0.028-0.1)	1.06 (0.78–1.35)	434.6 (421.9–448.4)
4	0.77 (0.58-1.1)	0.8 (0.33–1.3)	0.15 (0.09-0.23)	1.02 (0.76–1.25)	318 (307.4–329.2)
5	2.1 (1.8–2.2)	9.8 (7.8–12)	1.7 (1.52–1.91)	0.89 (0.48-1.32)	411 (395.6–428.4)
6	0.39 (0.032-2.3)	0.7 (0.014–7)	0.02 (0.006-0.058)	1.35 (1–1.62)	563 (543.3–581.3)
7	1.51 (0.36-4.1)	0.19 (0.033-0.8)	0.05 (0.03-0.087)	1.48 (1.32–1.62)	314 (304.8–323.5)
8	0.98 (0.84-1.11)	35 (20–68)	0.93 (0.8-1.06)	0.92 (0.66-1.19)	325.4 (315.3–336.2)
9	0.8 (0.18-2.7)	0.6 (0.034-2.4)	0.06 (0.018-0.2)	1.61 (1–1.85)	350.9 (335.4–363.2)
10	0.6 (0.46-0.72)	2.5 (2.1–3)	0.29 (0.22–0.36)	0.84 (0.52–1.18)	263 (252.5–275.2)

Nos. in parentheses show 95% confidence intervals. IVGTT, intravenous glucose tolerance test; S_1 , insulin sensitivity; p_2 , rate parameter, S_1^D , dynamic insulin sensitivity index; S_G , glucose effectiveness, G_0 , glucose at *time 0*.

$$\eta(p_2, T) = \frac{S_I^D}{S_I} = \left[1 - \frac{1 - e^{-p_2 T}}{p_2 T}\right] \tag{6}$$

Efficiency η measures which fraction of the potentially available insulin sensitivity, denoted by S_1 , is translated into effective glycemic control by means of a rapid increase of insulin action. The rapidity of insulin action is governed by parameter p_2 and η increases monotonically as a function of p_2 (once T has been fixed). Notice that the closer η to 1, the larger the efficiency of the metabolic system in converting S_1 into an effective insulin control on glucose. Conversely, a very low η renders S_1^D close to zero, i.e., in this case insulin action is virtually unavailable during the experiment. This observation explains why a high S_1 value does not necessarily imply an effective control of insulin on glucose, since it can be associated with a low p_2 and thus with a low efficiency.

As far as the choice of T is concerned, such a parameter does not represent the length of the IVGTT but is the duration of a "thought experiment" where insulin action has time to develop and IAIR can be directly observed. The "optimal" value of T should allow η to vary in the largest possible range of values so as to magnify differences between η values of the subjects under study. The reader is referred to Ref. 17 for all the details. Here, we just recall that after considering a wide population of NGT, IGT, and diabetic individuals, a robust choice consists of setting T to 60 min.

MM identification: IVGTT data. The four unknown MM parameters that are estimated from IVGTT data are hereby defined by the parameter vector $\theta = [S_1, S_G, p_2, G_0]$. These parameters are a priori uniquely identifiable given G(t) and I(t). As usual, we assume that I(t) in $Eq.\ 2$ is known at any t by linearly interpolating its measured plasma concentration samples. G(t) is instead known in sampled and noisy form on a grid $(t_1, t_2, \dots t_N)$. The measurement error is a zero-mean Gaussian with uncorrelated components and a 2% coefficient of variation.

The MM parameters were estimated using either a Bayesian or a Fisherian approach. The Bayesian approach (10) was implemented by using the Markov chain Monte Carlo (MCMC) strategy (11, 13), already used in Refs. 17, 18, 19. This powerful approach allows one to reconstruct (in sampled form) the entire a posteriori probability density function of S_I^D and S_I, thus permitting a more effective comparison of the performance of the new index with respect to the classic one. The Bayesian approach requires a probabilistic description of the a priori knowledge concerning the parameters to be estimated. The MM parameters S_I, S_G, p₂, and G₀ were assumed independent of each other and since all of them are nonnegative, priors whose support extends only in the positive axis were used. The a priori probability density functions of S_G and G₀ were assumed to be uniform in [0,a] with $a \to +\infty$. The choice of the a priori probability density function of S_I was based on the information on diabetic subjects reported in literature, e.g., Refs. 16 and 19. Specifically, we assumed for S_I an a priori probability density function where S_I values $<2 \times 10^{-4} \text{ (min}^{-1}/\mu\text{Uml)}$ are equally probable, whereas S_I values $>2 \times 10^{-4}$ are less and less probable according to a decreasing exponential law (with exponent equal to 10⁴ μUml⁻¹/min). The choice of the a priori probability density function of p_2 was based on the rationale described in Ref. 18; that is, we assumed a uniform distribution in the interval [0,5] min⁻¹. It is worth anticipating that the prior information on the MM parameters did not influence the general conclusions obtained in the study.

The MM was also identified using a Fisher parameter estimation scheme, which, albeit less sophisticated than the Bayesian one, makes it possible the adoption of mode-finding algorithms that are much less demanding than MCMC. This can have practical relevance in large-scale or epidemiological studies in which S₁^D has to be estimated in many subjects. To carry out the Fisherian estimation of the MM parameters, a maximum likelihood (ML) estimator (2, 7) was adopted.

MM identification: MGTT data. As already observed in Ref. 3, when the MM is identified from MGTT data, some difficulties arise

because the slow changes that glucose and insulin concentrations exhibit under postprandial conditions make it difficult to distinguish the individual contribution of insulin sensitivity and glucose effectiveness to glucose disappearance. To overcome this drawback, we followed the same strategy that was adopted in Ref. 3 to improve the numerical identifiability of the model, i.e., we exploited the fact that S_G , measuring glucose effectiveness at basal insulin, can be factored out in the sum of the glucose effectiveness at zero insulin (GEZI) and the product of insulin sensitivity times basal insulin concentration:

$$S_{G} = GEZI + S_{I}I_{b} \tag{7}$$

The advantage of expressing S_G in these terms lies in the possibility of introducing some reasonable assumptions about GEZI that guarantee an easier numerical identification of the MM. GEZI was assumed Gaussian with a mean of 0.025 and SD of 0.0025. The MM parameters were estimated by maximizing a penalized likelihood accounting for Eq. 7 and the prior on GEZI. To investigate the sensitivity of the MM estimates to the prior on GEZI, the MM identification was also carried out using a much less informative prior on GEZI (the SD of GEZI was increased from 0.0025 to 0.01).

Estimation of S_1^P from an OGTT and a euglycemic hyperinsulinemic clamp. We assessed the validity of S_1^D by comparing its estimates obtained in 21 subjects who underwent both an OGTT and a euglycemic hyperinsulinemic clamp.

OGTT-based estimates of \hat{S}_1^D were computed from Eq. 5 using the estimates of S_1 and p_2 obtained in Ref. 9. Of note is that the OGTT-based S_1^D estimate also accounts for the glucose distribution volume, V (see Eq. 7 in Ref. 9).

For what concerns computation of $S_1^D(\text{clamp})$, we simply refer to the theoretical definition of S_1^D . During an ideal clamp, the glucose infusion rate, R(t), is proportional to IAIR (see Appendix of Ref. 17) and equals insulin action multiplied by the distribution volume. Thus from $Eq.\ 2$ we have:

$$\dot{R}(t) = -p_2 R(t) + V \times p_3 (I(t) - I_b)$$
 $R(0) = 0$ (8)

Parameters p_2 and $V \times p_3$ governing the dynamics of R(t) were estimated by fitting the samples of R(t) via nonlinear least squares (the forcing input was defined by linearly interpolating the samples of insulin collected during the clamp). Finally, the dynamic insulin sensitivity was computed as follows:

$$S_1^D(\text{clamp}) = \frac{Vp_3}{p_2} \left[1 - \frac{1 - e^{-p_2 T}}{p_2 T} \right] \text{ with } T = 60 \text{ min}$$
 (9)

RESULTS

The MM identification results of the 10 diabetic patients who underwent the IVGTT are reported in Table 1 (Bayesian approach) and Table 2 (Fisherian approach). In particular, Table 1 reports the minimum variance estimates of S_I , p_2 , S_I^D , S_G , and G_0 , and the 95% confidence intervals (obtained as the

Table 2. IVGTT in diabetic subjects: maximum likelihood S_I and S_I^D estimates

Subject Nos.	$S_{I},10^{-4}\;min^{-1}\mu U^{-1}ml$	$S_{\rm I}^{\rm D},~10^{-4}~\text{min}^{-1}\mu\text{U}^{-1}\text{ml}$		
1	0.92	0.31		
2	1.51	1.16		
3	0.22	0.029		
4	0.86	0.15		
5	2.1	1.74		
6	0.14	0.027		
7	0.98	0.046		
8	0.99	0.935		
9	0.52	0.017		
10	0.58	0.29		

interval between quantiles 2.5 and 97.5%) in parentheses. Table 2 shows the ML estimates of S_I and $S_{_{\rm I}}^{\rm D}$. Figure 1 shows the marginal posteriors (obtained in sampled form by MCMC) of the parameters $S_I,~p_2,~{\rm and}~S_{_{\rm I}}^{\rm D}$ for subjects~2~(top) and 3 (bottom). Figure 2 displays S_I and $S_{_{\rm I}}^{\rm D}$ point estimates obtained in all the 10 subjects with 95% confidence intervals. Figure 3 compares $S_I~(left)$ and $S_{_{\rm I}}^{\rm D}~(right)$ estimates obtained using the Fisherian and the Bayesian approach. The correlation coefficient between ML and Bayes estimates of S_I and of $S_{_{\rm I}}^{\rm D}$ turns out to be 0.67 and 0.99, respectively.

Table 3 reports the identification results of the 14 diabetic patients who underwent the MGTT. In this table, the estimates of S_I , p_2 , and S_I^D obtained using both the full as well as the less informative prior on GEZI are shown. Figure 4 compares the S_I (*left*) and S_I^D (*right*) estimates obtained with the two priors.

Figure 5 reports the results of the simulation study designed to compare the performance of S_I and S_I^D in NGT and diabetic subjects. The true values of S_I and S_I^D are plotted against the values estimated by ML in the two groups. The correlation coefficients between true and estimated S_I are ~ 0.1 and 0.4 in diabetic and NGT subjects, respectively, while those concerning S_I^D are 0.98 and 0.99, respectively.

Figures 6 and 7 concern the results obtained in the 21 subjects who underwent both an OGTT and a clamp experiment. Figure 6 shows the fit of clamp data in one representative IGT subject: the glucose infusion rate values during the clamp are denoted by the open circles, while the solid line describes the curve, proportional to IAIR, obtained after fitting the samples using Eq. 8 and nonlinear least squares. The area under the curve in the first 60 min (shaded area) is proportional to S_1^D (clamp). Figure 7 compares the S_1^D (clamp) and S_1^D estimates obtained in the two experimental situations via correla-

tion analysis. In particular, the correlation coefficient between OGTT-based estimate of S_{τ}^{D} and S_{τ}^{D} (clamp) is 0.82.

DISCUSSION

IVGTT in diabetic subjects: S_I vs. S_I^D using Bayes estimation. To comment on the results of the Bayesian estimation, it is helpful to examine the MM identification results obtained in two paradigmatic subjects. Figure 1 plots the marginal posteriors (obtained in sampled form by MCMC) of the parameters S_I , p_2 , and S_I^D for subject 2 (top) and 3 (bottom). As far as S_I is concerned (first column of Fig. 1), one can see that this parameter is well estimated in subject 2. In fact, its posterior appears well concentrated around its mean. From Table 1, one can see that the point estimate of S_I is 1.5 (1.4–1.6) in this subject. The situation is completely different for subject 3. In this subject, S_I suffers of poor numerical identifiability, as indicated by its long-tailed marginal posterior (this phenomenon is mathematically explained by Proposition 2 in Ref. 18). In addition, the point estimate of S_I is 1.7 (0.4–4.1) and thus very close to the mean of the prior (whose value is ~ 1.67). This suggests that the IVGTT data provide little information to estimate S_I in this subject. All in all, it is difficult to tell whether the S_I of *subject 3* is really higher than that of *subject* 2. The examination of the S_i^D results obtained in the same subjects offers a completely different scenario. In fact, the marginal posteriors of S. (third column of Fig. 1), are well concentrated around their means in both subjects. The S^D point estimates are 1.17 (1.1-1.23) and 0.06 (0.028-0.1), respectively, and this suggests that insulin is in all likelihood more effective to enhance glucose disappearance in subject 2 than in subject 3. To understand why the point estimate of S_i^D is much

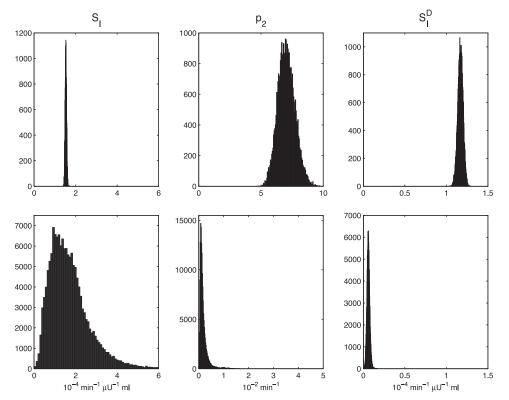


Fig. 1. Intravenous glucose tolerance test (IVGTT): diabetic *subjects* 2 (*top*) and 3 (*bottom*). Marginal posterior [obtained in sampled form by Markov Chain Monte Carlo (MCMC)] of the index of insulin sensitivity (S_1 ; *left*), parameter p_2 (*middle*), and new dynamic insulin sensitivity index (S_1^D ; *right*).

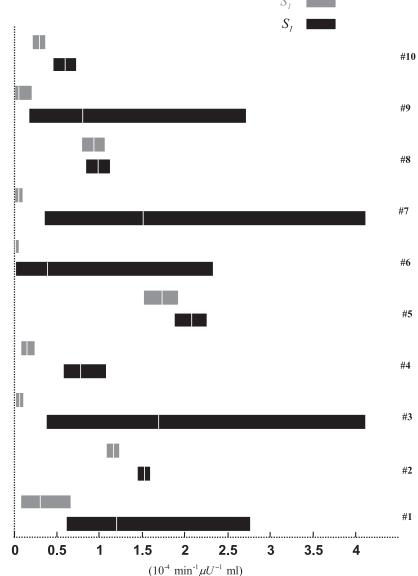


Fig. 2. IVGTT: Bayes S_1 and S_1^D in diabetic subjects. Point estimates are shown with the 95% confidence intervals (horizontal bars)

lower in *subject 3* than in *subject 2*, one must consider that S_1^D embodies the information brought about by both S_I and p_2 (the latter parameter is reported in the second column of Fig. 1). It can be seen that in *subject 3* the S_I 95% confidence interval includes many S_I values that are coupled with very low values of p_2 . This means that in this subject S_I is frequently associated with very low values of the efficiency η , which translates into a low point estimate of S_I^D .

The usefulness of the new index is also apparent when the whole group of diabetic subjects is considered. The results reported in Fig. 2 displaying the S_I and S_I^D estimates together with their 95% confidence intervals indicate that S_I^D never exhibited numerical identifiability problems. In contrast, the S_I results found in *subjects 1*, 3, 6, 7, and 9 are rather difficult to interpret.

IVGTT in diabetic subjects: Bayes vs. Fisher estimation of S_i^D . In Ref. 19, the Bayes approach was shown to be much more robust than the Fisher one in estimating S_1 and p_2 in NGT subjects. Bayes superiority over Fisher can be even more

appreciated in the present study because diabetic patients exhibit low values of S_I that are often coupled with low values of p_2 . In this case, the profile of insulin action becomes so low and slow so as to degrade the precision of the MM parameter estimates. In fact, ML often yields an estimate of S_I that is affected by a large uncertainty and cannot be discriminated from zero (this is known in the literature as the " $S_I=0$ " problem; Refs. 1, 18, 19). As a consequence, the histogram describing the S_I frequency distribution is bimodal showing an artifactual peak at $S_I=0$ and another peak at a positive S_I value, which leads to interpretative difficulties. Another problem is also that S_I estimates can turn out to be unrealistically high. Bayes estimation, implemented by using MCMC, is more robust than Fisher estimation since it provides a nonzero minimum variance estimate of S_I that hinges on the entire marginal posterior of S_I. In addition, rigorous confidence intervals for S_I can be calculated. The question arises as to whether the same results also hold for S₁^D. Since S₂^D depends on both p_2 and S_I (see Eq. 5), one would not be surprised to find

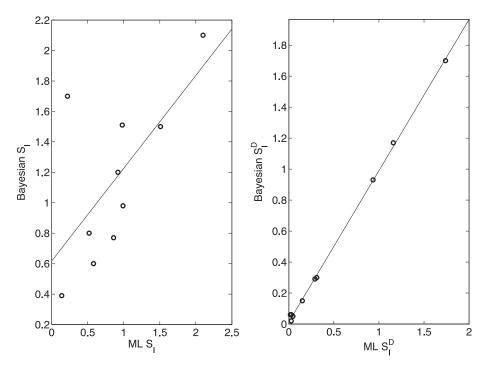


Fig. 3. IVGTT. *Left*: maximum likelihood vs. Bayesian S_I estimates in diabetic subjects. *Right*: ML vs. Bayesian S_I^D in diabetic subjects.

out that also the Fisherian point estimate of S₁^D is plagued by the same problem described above for S_I. Fortunately, however, the Fisherian estimate of S₁^D was often very close to the one obtained with the Bayesian approach. This can be appreciated by looking at Fig. 3 where the Bayesian and Fisherian estimates of S_I a S_I^D are reported. As said, the correlation coefficient between ML and Bayes estimates of S_I and of S_{i}^{D} was 0.67 and 0.99, respectively. This favorable outcome for S_{i}^{D} is related with the functional relationships among S_I , p_2 , IAIR, and S_1^D . While it is difficult to estimate precisely S_1 and p_2 , due to the presence of numerical nonidentifiability regions that make many different combinations of S_I and p_2 capable of generating virtually the same IAIR on a finite time interval, IAIR itself and its integral (and thus S₁^D) can be precisely determined. Thus, although Bayes is more robust in estimating S_1 and p_2 separately, Bayes and Fisher behave similarly as regards S₁^D.

In summary, the Fisher estimator often provides $S_{\scriptscriptstyle I}^D$ estimates comparable to the Bayesian ones with a reduced computational effort (few seconds in place of minutes, since Markov chain generation and Monte Carlo integration are more demanding than mode-finding algorithms). This advantage of $S_{\scriptscriptstyle I}^D$ over $S_{\scriptscriptstyle I}$ may be particularly relevant in large-scale studies in which many model identifications must be carried out.

IVGTT and MGTT in diabetic subjects: sensitivity of S_I and S_I^D to prior. To assess the sensitivity of the results to the prior information on S_I used in the MM Bayesian identification from IVGTT in diabetics, the estimation process was repeated using larger thresholds (3 and 4 in place of 2). As already found in Ref. 19, S_I and p_2 were particularly sensitive to prior information. In particular, the prior information exerted a profound influence on the tails of the S_I and S_I marginal posteriors. Quite to the contrary, we found (results not shown) that the new index S_I^D is scarcely sensitive to the prior information. MM

Table 3. MGTT in diabetic subjects: S_I , S_I^D , and p_2 estimates obtained using prior information on GEZI

Subject Nos.	$S_{I}(1)$	$S_{I}(2)$	$S_{\rm I}^{ m D}(1)$	$S_{\rm I}^{ m D}(2)$	$p_2(1)$	$p_2(2)$
1	5.9	12.2	0.31	0.29	0.18	0.081
2	2.0	2.0	0.41	0.39	0.78	0.758
3	6.8	6.6	0.73	0.74	0.38	0.404
4	2.5	5	0.12	0.11	0.167	0.078
5	2.1	4.3	0.16	0.14	0.258	0.116
6	4.4	6.5	0.349	0.33	0.28	0.177
7	1.5	1.38	0.55	0.23	1.7	0.63
8	6.3	5.74	1.46	1.87	0.92	1.42
9	6.9	6.9	0.72	0.72	0.37	0.377
10	2.0	2.7	0.0149	0.015	0.025	0.019
11	4.6	0.8	0.17	0.17	0.13	0.053
12	5.8	2.5	0.42	0.39	0.26	0.107
13	9.9	20	0.31	0.3	0.107	0.05
14	24.1	24.3	0.0064	0.0057	0.00088	0.00078

Estimates are denoted by $S_I(1)$, $S_I^D(1)$, and $p_2(1)$ when using the full prior and by $S_I(2)$, $S_I^D(2)$, and $p_2(2)$ when using the less informative prior; GEZI, glucose effectiveness at zero insulin; MGTT, meal glucose tolerance test; measurement units of MM purameters are the same as in Tables 1 and 2.

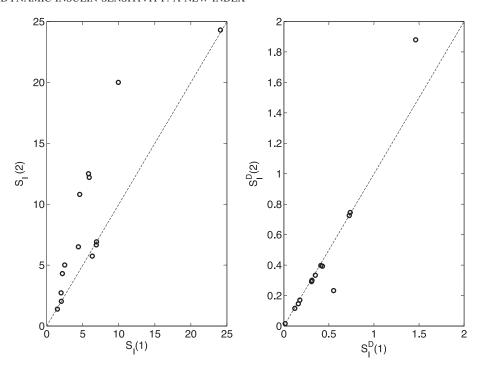


Fig. 4. Meal glucose tolerance test (MGTT). Left: insulin sensitivity using full prior information on glucose effectiveness at zero insulin [GEZI; $S_I(1)$] vs. less informative [$S_I(2)$]. Right: dynamic insulin sensitivity using full prior information on GEZI [$S_I^D(1)$] vs. less informative [$S_I^D(2)$].

identification from MGTT data in the 14 diabetic patients provides further support to these findings. In fact, inspection of Table 3 and Fig. 4 indicate that, at variance with S_I and p_2 , the estimates of S_I^D are virtually insensitive to the prior information on GEZI.

The reasons for the higher robustness of S_1^D with respect to S_1 and p_2 are the same already put forth to discuss the Bayesian vs. Fisherian identification results. The parameters S_1 and p_2 are more sensitive to the prior information because of the existence of numerical nonidentifiability regions in the param-

eter space where changes in these two parameters only produce negligible changes in the likelihood (18). Within such numerical nonidentifiability regions, many different combinations of S_I and p_2 may lead to similar IAIR. Since S_I^D is the mean of IAIR in the interval (0,T) with T=60 min, many different combinations of S_I and p_2 are mapped into the same S_I^D . Therefore, whereas S_I and p_2 may be very uncertain, S_I^D is not. This peculiarity of S_I^D can be easily grasped by resorting to a clamp thought experiment, where S_I depends on the asymptotic value of the exogenous glucose infusion. Large and unrealistic

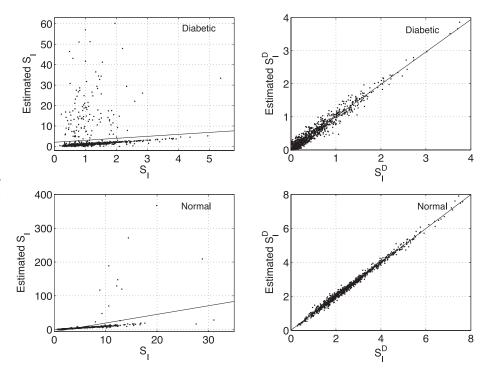


Fig. 5. Left: true S_I vs. estimated S_I in diabetic (top) and NGT subjects (bottom). Right: true S_I^D vs. estimated S_I^D in diabetic (top) and NGT subjects (bottom).

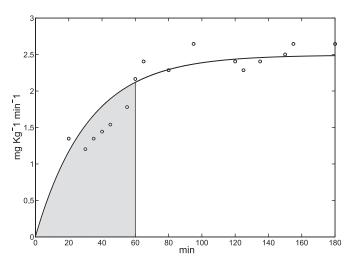


Fig. 6. Clamp model. Samples of glucose infusion measured in the first IGT subject (circles) and curve (proportional to IAIR) obtained after fitting the samples using Eq. 8 (solid line). Area under the curve in the first 60 min is proportional to S₁^D(clamp).

 S_I values may be obtained in diabetic subjects since they are associated with low p_2 values that define virtual glucose infusions similar to straight lines, which "almost diverge" at infinity. On the other hand, by definition, S_I^D is proportional to the area of glucose infusion in the first 60 min and it is thus much less sensitive to poor numerical identifiability of p_2 . As a practical consequence, S_I^D can be reliably estimated from IVGTT/MGTT experiments also under conditions where S_I and p_2 suffer from poor numerical identifiability. The small sensitivity of S_I^D to the prior information contributes to make S_I^D a much more robust and objective measure of insulin ability to control glucose.

IVGTT: S_I vs. S_I^D ability in discriminating NGT and diabetic subjects. The results reported in Fig. 5 provide further support to the notion that S_{I}^{D} can be estimated with much more precision than S_I during an IVGTT experiment, especially when diabetic subjects are under study. As already said, the correlation coefficients between true and estimated S_I are ~ 0.1 and 0.4 in diabetic and NGT subjects, respectively, while those concerning S₁^D are 0.98 and 0.99, respectively. It is of interest that in Fig. 5, top left, almost 10% of estimated S_I values turn out to be 10 times larger than the true ones. This means that, in a nonnegligible number of cases, a diabetic patient is erroneously classified as a subject with a normal insulin sensitivity. In contrast, the discriminatory power of S₁^D is always excellent. This new evidence of the superior performance of S_I^D against S_I adds to the one previously reported in Refs. 17 and 20, where we showed that the new dynamic index is better correlated than S_I with the time by which glucose concentration reattains its basal level after a glucose perturbation.

Validation of S_i^D against the euglycemic hyperinsulinemic clamp. Figure 6 shows the area under the curve in the first 60 min of the clamp experiment, proportional to S_i^D (clamp). This clearly shows how the new index measures the ability of insulin to control glucose by accounting not only for the final value achieved by the glucose infusion but also for the speed of rise of the glucose infusion.

The results displayed in Fig. 7 show that the OGTT-based estimates of S_1^D and S_1^D (clamp) are well correlated (r = 0.82).

Such degree of correlation is almost identical to the one between S_I and S_I (clamp) (r=0.81) obtained from the analysis of the same experimental data in a previous report (9). It is worth noticing that if we focus our attention on the results obtained in the 11 IGT subjects, the correlation index between S_I^D and S_I^D (clamp) is 0.91, whereas that between S_I^D and S_I^D (clamp) is only 0.7. Instead, in the 10 NGT subjects, the two correlation indexes are similar and \sim 0.6. This dissimilarity also points in the direction that the benefits of S_I^D become more and more relevant as the degree of glucose tolerance decreases.

Conclusions

The present study is a follow up of previous paper in NGT subjects (17) where we pointed out the limitations of the classical MM index of insulin sensitivity, S_I, and proposed a novel index, S.D. The new index yields a more comprehensive picture of insulin's ability to control glucose metabolism because it hinges upon a theoretically based combination of p_2 and p_3 that better accounts for the dynamic properties of insulin action. For instance, S₁^D has been shown to be better correlated than S_I with the time by which glucose concentration reattains its basal level after a glucose bolus (17). Further, by incorporating the concept of efficiency, the new index provides an elegant answer to those perplexing situations in which, despite an elevated value of S_I, the rate of glucose disappearance seems to be scarcely affected by insulin. This occurs when the elevated S_I cannot be promptly translated into effective insulin action due to a low p_2 value. This situation is exemplified by the results of patient 14 in Table 3. In fact, in this subject, the large S_I value would suggest that insulin control is in the normal range, whereas the S₁^D value unmasks a low ability to restore glucose homeostasis. The simulation study results reported in Fig. 5 further corroborate the notion that S₁^D can considerably improve the reliability of the assessment of insulin sensitivity in diabetic patients. Furthermore, the issue of the validity of S₁^D against a reference measure has been tackled in the present study by comparing the estimates of S^D obtained from an OGTT with those obtained in the same subjects from a euglycemic hyperinsulinemic clamp, i.e., the gold standard technique for the assessment of insulin sensitivity.

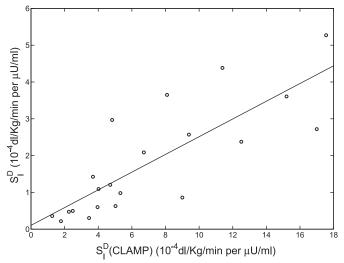


Fig. 7. Clamp and oral minimal model. $S_{\rm I}^{\rm D}$ vs. $S_{\rm I}^{\rm D}({\rm clamp})$ estimates in IGT and NGT subjects.

The results of the present study indicate that the new dynamic index is especially useful in those diabetic subjects having a slow timing of insulin action. It is well known that, when analyzing IVGTT data, the estimation of p_2 in diabetic subjects is often problematic. One of the most relevant findings of the present study is that S_I^D is much less affected than S_I by such a drawback. Thus S_I^D has the additional advantage over S_I in that its estimation is much more robust and can be accomplished by resorting to the Fisher approach. In fact, at variance with what happens with S_I, the Fisherian and Bayesian point estimates of \hat{S}_{r}^{D} are similar, with the latter being virtually insensitive to the prior information incorporated in the estimator. Since ML is much less computationally demanding than MCMC, this makes S_{i}^{D} more appealing than S_{I} in studies where many subjects have to be analyzed. S_i^D thus appears particularly suitable to be employed in population studies where glucose intolerant subjects are likely to be present, as well as in studies seeking for diabetes-relevant genes in the human genome (12, 14, 21, 22).

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DISCLOSURES

No conflicts of interest are declared by the author(s).

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