ABSTRACT  Insulin is released from the pancreas in pulses with a period of ~ 5 min. These oscillatory insulin levels are essential for proper liver utilization and perturbed pulsatility is observed in type 2 diabetes. What coordinates the many islets of Langerhans throughout the pancreas to produce unified oscillations of insulin secretion? One hypothesis is that coordination is achieved through an insulin-dependent negative feedback action of the liver onto the glucose level. This hypothesis was tested in an in vitro setting using a microfluidic system where the population response from a group of islets was input to a model of hepatic glucose uptake, which provided a negative feedback to the glucose level. This modified glucose level was then delivered back to the islet chamber where the population response was again monitored and used to update the glucose concentration delivered to the islets. We found that, with appropriate parameters for the model, oscillations in islet activity were synchronized. This approach demonstrates that rhythmic activity of a population of physically uncoupled islets can be coordinated by a downstream system that senses islet activity and supplies negative feedback. In the intact animal, the liver can play this role of the coordinator of islet activity.

INTRODUCTION

Islets of Langerhans secrete insulin in response to elevations in the blood glucose level. As with many endocrine systems, insulin secretion from the pancreas is pulsatile, yielding ~ 5 min oscillations in the plasma insulin concentration (1−3). These oscillations are important, because pulsatile insulin levels in the portal vein are more effective than constant levels at evoking insulin action in the liver (3), and disorganized insulin pulsatility is observed in people with diabetes and their near relatives (4,5).

The activity of individual islets is also pulsatile, exhibiting a ~ 5 min period in electrical activity, intracellular Ca2+ concentration ([Ca2+]i), and insulin secretion (6). The similarity of the periods from individual islet activity and in vivo pulsatility suggests that the hundreds of thousands of islets scattered throughout the pancreas are synchronized. However, they are not physically coupled, so what coordinates their activity enabling the population of oscillators to produce a coherent rhythm in insulin secretion? Islet synchronization must occur, otherwise phase and period differences between the individual islet oscillators would result in an overall blood insulin level that is relatively flat. Yet pulsatile insulin levels have been observed in humans (1), mice (2), rats (7), dogs (8), and monkeys (9).

One mechanism for synchronization is that islets may be coordinated through intrapancreatic ganglia (10,11). The pancreas is innervated by preganglionic vega neurons (12–15), and the ganglia have been shown to exhibit electrical excitability when autonomic nerve trunks were stimulated in the cat (16). It has been shown that a bolus of the cholinergic agonist carbachol can transiently synchronize islet oscillations (17), and application of a pulse of ATP induced a [Ca2+]i response in islets, suggesting that release of ATP from ganglia neurons may help to coordinate islet activity (18). However, there is at present no data suggesting that ganglia neurons exhibit pacemaking activity, as would be required to synchronize the population of islets in the pancreas.

A study using mathematical modeling and computer simulations demonstrated that interactions between pancreatic islets and the liver would be sufficient to coordinate islet rhythmicity into 5 min pulses (19). According to this hypothesis (Fig. 1A), insulin secreted from the pancreas would signal the liver to reduce hepatic glucose output and initiate glucose uptake, causing a reduction in blood glucose. The reduced glucose level would then be sensed by each islet in the population, lowering the collective insulin output from the pancreas. In this way, the liver acts as a negative feedback on the islets by reducing blood glucose levels. The lower insulin levels would then promote a reduced glucose uptake by the liver allowing the levels of glucose to rise again. These variable glucose levels act as the coordinating signal for islet activity promoting oscillations in both insulin and glucose. Although glucose uptake by other tissues, such as muscle, is quantitatively important, these tissues would be exposed to smaller amplitude changes in insulin levels than the liver (1).

This hypothesis of a dynamic insulin-glucose feedback system coordinating islet activity to produce coherent 5 min pulses of insulin was postulated some years ago (20–23), yet no experimental test of this theory has been described. Evidence for this hypothesis comes from
oscillations of insulin (1,2,7–9) and glucose (9,24) observed in vivo. In this report, a microfluidic system was used to perfuse a population of islets with glucose and the average response of the population was used to continuously update the glucose level. The updating was achieved with a simple mathematical model for negative feedback. Without the negative feedback, the islets exhibited uncoordinated oscillations in \([\text{Ca}^{2+}]_i\) and insulin secretion. When the feedback was turned on these oscillations quickly synchronized, producing glucose and insulin oscillations with periods of 5 to 8 min. These results indicate that a negative feedback on glucose levels can coordinate pulsatile insulin release.

**MATERIALS AND METHODS**

**Chemicals and reagents**

Potassium chloride (KCl), sodium chloride (NaCl), calcium chloride (CaCl₂), magnesium chloride (MgCl₂), dimethyl sulfoxide (DMSO), tricine, and penicillin-streptomycin were purchased from Sigma-Aldrich (Saint Louis, MO). Pluronic F-127 and fura 2 acetoxymethyl ester (fura 2-AM) were from Life Technologies (Grand Island, NY). Glucose (dextrose) was purchased from Thermo Fisher Scientific (Waltham, MA). RPMI 1640 was purchased from Mediatech (Manassas, VA). Gentamicin sulfate was purchased from Lonza (Basel, Switzerland). Collagenase P (from Clostridium histolyticum) was purchased from Roche Diagnostics (Indianapolis, IN). Poly(dimethylsiloxane) (PDMS) prepolymer (Sylgard 184) was purchased from Dow Corning (Midland, MI). All solutions were made with Milli-Q (Millipore, Bedford, MA) 18 Ω cm deionized water.

**Isolation and culture of islets of Langerhans**

All experiments were performed under guidelines approved by the Florida State University Animal Care and Use Committee (ACUC) (protocol No. 1235). Islets of Langerhans were isolated from male CD-1 mice (30 – 50 g). Islets were isolated using collagenase P digestion as described previously (25–27). The isolated islets from multiple mice were mixed during picking and randomly chosen for the experiment. After isolation, islets were incubated at 37°C, 5% CO₂ in RPMI 1640 media containing 11 mM glucose, 10% calf serum, 100 units mL⁻¹ penicillin, 100 μg mL⁻¹ streptomycin, and 10 μg mL⁻¹ gentamicin. Islets were used within five days after isolation and fresh media was provided on day 3 after isolation.

For \([\text{Ca}^{2+}]_i\) monitoring, 1.0 μL of 5.0 mM fura 2-AM in DMSO and 1.0 μL Pluronic F-127 in DMSO were mixed and transferred into 2 mL of RPMI to form a final fura 2-AM concentration of 2.5 μM. Each batch of islets was incubated in this solution at 37°C and 5% CO₂ for 40 min. After this time, the islets were removed and placed in the microfluidic device and rinsed with 3 mM glucose in balanced salt solution (BSS). The BSS was composed of 2.4 mM CaCl₂, 125 mM NaCl, 1.2 mM MgCl₂, 5.9 mM KCl, varying concentrations of glucose as described below, and 25 mM tricine at pH 7.4 and was used in all experiments.
Insulin quantification

Ten islets were used in the experiment described in the text and [Ca\(^{2+}\)]\(_i\), was monitored as described below. During the experiment, perfusate was collected at 1 min intervals from the top of the chamber. An insulin ELISA (Ultra Sensitive Mouse Insulin ELISA kit, Crystal Chem, Inc., Downers Grove, IL) was used to quantify insulin levels according to the manufacturer’s instructions.

Microfluidic device

A microfluidic device design was adapted from (28). The PDMS-glass hybrid device was fabricated using conventional photolithography. All channel dimensions were 250 × 40 μm (width × height). The device consisted of two inputs and three outputs. The inputs were connected to 60 mL syringes suspended by a pulley system. The syringes contained BSS described above with either 3 or 13 mM glucose. The height of one of the syringes was actuated by a stepper motor controlled by a LabVIEW (National Instruments, Austin, TX) program while the height of the other syringe was displaced in the opposite direction by an equal amount using a fixed-length belt and pulley system (28). The difference in the heights of the two syringes caused different flow rates of the two glucose solutions to enter the 70 mm mixing channel with excess buffer diverted via the side channels to a waste reservoir. This design maintains a constant total flow rate. By changing the heights of the two syringes, the concentration to homogeneity and delivered to a 0.8 mm diameter chamber that housed 5 to 10 islets. By changing the heights of the two syringes, the concentration of glucose delivered to the islets could be changed. The system was calibrated in a manner similar to our previous reports (25–27). It took ~30 s of glucose delivered to the islets could be changed. The system was calibrated in a manner similar to our previous reports (25–27). It took ~30 s of glucose delivery to the islet chamber. The buffers passing to the mixing channel were mixed to homogeneity and delivered to a 0.8 mm diameter chamber that housed 5 to 10 islets. By changing the heights of the two syringes, the concentration of glucose delivered to the islets could be changed. The system was calibrated in a manner similar to our previous reports (25–27). It took ~30 s of glucose delivery to the islet chamber.

The microfluidic device was fixed on the stage of a Nikon Eclipse Ti inverted microscope. A lamp integrated with filter wheel and shutter (Lambda XL, Sutter Instruments, Novato, CA) containing appropriate filters was used for excitation of fura-2 at 340 and 380 nm. The images were acquired with a 150 ms exposure every 20 s using a CCD (Cascade, Photometrics, Tucson, AZ) controlled by Nikon NIS Elements software (Nikon, Melville, NY). The ratio of fluorescence intensity excited at 340 nm to that at 380 nm (F\(_{340}/F_{380}\)) for all islets were obtained by NIS software and exported to a file. The F\(_{340}/F_{380}\) ratio values were converted to [Ca\(^{2+}\)]\(_i\) using predetermined calibration values that were found by standard methods (29). The LabVIEW program that controlled the syringes then read the [Ca\(^{2+}\)]\(_i\) and calculated C\(_{avg}\) to be used in the feedback.

Mathematical model

The differential equation for the extracellular glucose concentration, Eq. 6, was discretized using the forward Euler method. Let F(G\(_{avg}\), G\(_i\)) = G\(_{avg}\) (I\(_{avg}\)/G\(_{avg}\)) – G\(_i\)/τ\(_G\) be the right-hand side of the differential equation, where the superscript “n” represents “time step n.” Then at each subsequent sampling point, G\(_n\) was updated from its previous value G\(_{n-1}\) according to the following:

\[
G_{e,n+1} = G_{e,n} + F\left(\frac{I_{avg}}{G_{avg}}, G_{e,n}\right) \Delta t
\]  

where \(\Delta t = 1 s\) was the time step. The C\(_{avg}\) and I\(_{avg}\) values were updated every 20 time steps.

Synchronization index (\(\lambda\))

The synchronization index (\(\lambda\)) was calculated as described previously (25,30,31). Briefly, for a group of five islets, the [Ca\(^{2+}\)]\(_i\) oscillations of the first islet are compared with the [Ca\(^{2+}\)]\(_i\) oscillations of each of the remaining four islets, and in each case the degree of phase locking is captured as \(\lambda_{ij}\) for \(j = 2,...,5\). A \(\lambda\) value near 1 means that two [Ca\(^{2+}\)]\(_i\) oscillation traces are nearly phase locked, whereas a \(\lambda\) near 0 means that they are not. The calculation is then repeated by comparing the [Ca\(^{2+}\)]\(_i\) oscillations of islet 2 with each of the others, and computing \(\lambda_{ij}\) for \(j = 1,3,4,5\). This is repeated three more times giving \(\lambda_{ij}, \lambda_{4,j}, \lambda_{5,j}\) and in the end an overall \(\lambda\) is calculated as the average of the 20 individual indices. For the open-loop experiments (Fig. 1 D), \(\lambda\) was calculated in a similar manner by computing the degree of phase locking between an individual islet and the glucose trace.

Data analysis

For calculation of synchronization index and oscillation periods described above, the initial 5 min of each F\(_{340}/F_{380}\) trace after a change in glucose level were not used for data analysis. For example, during an open-loop experiment, the timing of the glucose delivery was the following: \(t = 0 – 3\) min, 3 mM glucose; \(t = 3 – 23\) min, 11 mM glucose; \(t = 23 – 60\) min, oscillatory glucose. To determine \(T_n\) and \(T_f\), the F\(_{340}/F_{380}\) trace from 8 – 23 min and 28 – 60 min were used, respectively. This was performed to reduce transients in the [Ca\(^{2+}\)]\(_i\) levels associated with the change in glucose level (25). \(T_n, T_p\), and the period of [Ca\(^{2+}\)]\(_i\) oscillations with sinusoidal glucose were calculated by fast Fourier transforms of the [Ca\(^{2+}\)]\(_i\) trace with baseline subtracted.

Phase angle

In Fig. 1 C, the islet response to the forcing glucose signal was shown by a phase angle (\(\Phi\)) between [Ca\(^{2+}\)]\(_i\) oscillations and the glucose oscillograms. \(\Phi_i\) (for \(i = 1, 2, 3,..., n\) oscillations) was found by measuring the difference in the times of the midpoints between each glucose peak (\(t_{g,i}\)) and the closest [Ca\(^{2+}\)]\(_i\) peak (\(t_c,i\)):

\[
\Phi_i = \frac{360^\circ (t_{c,i} – t_{g,i})}{T}
\]  

where \(T\) is the period of the forcing glucose wave. The radius of the point on the polar plot is the amplitude of the [Ca\(^{2+}\)]\(_i\) oscillation.

RESULTS

Effects of oscillatory glucose levels on islets

A prediction of the hypothesis that a dynamic interaction between the pancreas and liver could synchronize an islet population is that the glucose levels sensed by islets will be oscillatory. Initial experiments were, therefore, first performed in an open-loop manner in which the glucose concentration was varied sinusoidally and the range of glucose amplitudes and periods that could effectively entrain islet activity was determined.

Glucose oscillations with periods from 5 to 10 min, a median value of 11 mM, and amplitudes ranging from 0.25 to 3 mM were delivered to islets while monitoring [Ca\(^{2+}\)]\(_i\) changes within individual islets (Fig. 1 B). [Ca\(^{2+}\)]\(_i\) is a good marker for insulin secretion because they are proportional (32,33). We observed slow oscillations of [Ca\(^{2+}\)]\(_i\) with a period of ~5 min, but not faster oscillations because the imaging rate was 0.05 Hz. To illustrate entrainment, the phase angles (\(\Phi\)) between each glucose and [Ca\(^{2+}\)]\(_i\) oscillation from four experiments are shown on the polar plot in Fig. 1 C. Entrainment is evident by convergence of \(\Phi\) at a
particular value, as illustrated by the blue, black, and green points in Fig. 1 C. As shown by the red points, $\Phi$ did not converge for each condition tested, which is discussed below.

To better quantify entrainment, a synchronization index ($\lambda$) was employed (25,30,31). $\lambda$ indicates the degree of synchrony between the imposed glucose oscillations and the resulting oscillations of [Ca$^{2+}$], with values of $\lambda$ near 1 indicating complete synchronization. $\lambda$ was calculated for all islets ($n = 89$) at each combination of glucose amplitude and period tested (188 conditions tested). The results of the experiments are shown in Fig. 1 D with the forcing glucose amplitude on the ordinate and the ratio of the forcing period to the islet’s natural period ($T_f/T_n$) on the abscissa. Islets were deemed entrained if $\lambda \geq 0.7$ (blue circles) whereas $\lambda < 0.7$ indicates no entrainment (red crosses). Entrainment was achieved provided that the forcing amplitude was sufficiently large, as expected in entrainment experiments (34,35).

**Closed-loop experiments using a model of insulin-dependent glucose uptake**

After determining the region of entrainment for islets, the feedback loop was closed (Fig. 2) by stimulating groups of 5 to 10 islets with glucose while measuring [Ca$^{2+}$]$_i$ changes within each islet. The [Ca$^{2+}$]$_i$ from all islets was averaged ($Ca_{avg}$) and was converted into a measure of insulin secretion in arbitrary units ($I_{avg}$). Based on previous findings (32,33), this was modeled as an increasing linear function above a threshold value ($Ca_{thr}$) of $Ca_{avg}$ (19):

$$I_{avg}(Ca_{avg}) = \begin{cases} I_{slope}(Ca_{avg} - Ca_{thr}) & \text{for } Ca_{avg} \geq Ca_{thr} \\ 0 & \text{for } Ca_{avg} < Ca_{thr} \end{cases}$$  

$$Ca_{thr} = (Ca_{max} - Ca_{min})\kappa + Ca_{min}$$  

The parameters $I_{slope}$ and $\kappa$ were set at 1 A.U.*nM$^{-1}$ and 0.1, respectively, and $Ca_{max}$ and $Ca_{min}$ were dependent on the individual experiment. Although a sigmoidal fit could have been used to model $I_{avg}$ vs. $Ca_{avg}$, the [Ca$^{2+}$]$_i$ and insulin correlation is nearly linear within the [Ca$^{2+}$]$_i$ ranges we observed (32). That is, the [Ca$^{2+}$]$_i$, is in the linear portion of the sigmoidal curve over the ranges we observed. The asymptotic glucose response function $G_\infty$ was modeled as a decreasing sigmoidal function of $I_{avg}$ (Fig. 2), given by the following:

$$G_\infty(I_{avg}) = G_{min} + \frac{G_{min} - G_{max}}{1 + \exp[(I_{avg} - I_{1/2})/S_G]}$$  

$I_{1/2}$ is the point of inflection of the curve and is the point at which $G_\infty$ is midway between $G_{min}$ and $G_{max}$. $S_G$ is the steepness parameter of the curve, and parameters $G_{min}$ and $G_{max}$ were the minimum and maximum glucose levels of the curve and were set to 7 and 13 mM, respectively.

The extracellular glucose concentration ($G_e$) was updated using the following differential equation:

$$\frac{dG_e}{dt} = \frac{G_e(I_{avg} - G_e)}{\tau_G}$$  

This differential equation was discretized as described in the Materials and Methods section, Eq. 1. All parameters and variables are shown in Tables 1 and 2, respectively. In this model, two key parameters shaped the timing and magnitude of the feedback response. Parameter $\tau_G$ adjusted $G_\infty$, with large values producing a low sensitivity, i.e., a shallow glucose response to $I_{avg}$ (Fig. 2). Parameter $\tau_G$ was the time constant for the feedback response, with large values of $\tau_G$.

| Table 1 Parameters and values used in the experiments |
|---------------------------------|-----------------|
| Parameter | Description | Value |
| $I_{slope}$ | Slope of $Ca_{avg}$ to $I_{avg}$ conversion | 1 A.U.*nM$^{-1}$ |
| $Ca_{thr}$ | Threshold [Ca$^{2+}$], for insulin release | 83-105 nM |
| $Ca_{min}$ | Minimum [Ca$^{2+}$]$_i$ | 70-90 nM |
| $Ca_{max}$ | Maximum [Ca$^{2+}$]$_i$ | 200-240 nM |
| $\kappa$ | Fraction above $Ca_{max}$ to set $Ca_{thr}$ | 0.1 |
| $I_{1/2}$ | Minimum glucose value of $G_\infty$ | 7 mM |
| $G_{min}$ | Maximum glucose value of $G_\infty$ | 13 mM |
| $I_{1/2}$ | Inflection point of $G_\infty$ | 30-40 A.U. |
| $S_G$ | Steepness of $G_\infty$ (smaller is steeper) | 0.1, 1, 10, 50, 100 A.U. |
| $\tau_G$ | Time constant for feedback response | 50, 300, 450, 600, 900 s |

Values that are shown as a range varied between experiments depending on the islets examined.

FIGURE 2 Overview of closed-loop feedback system. Beginning from the top left, a microfluidic device produced varying concentrations of glucose and delivered to batches of 5 to 10 islets shown in the inset. [Ca$^{2+}$]$_i$ from individual islets is measured using Fura-2 and the average calcium level of the population ($Ca_{avg}$) is converted to an average insulin secretion level ($I_{avg}$) using a linear proportionality (bottom right). $I_{avg}$ is the sole input to the asymptotic glucose response function, $G_\infty$ (bottom left). Using $G_\infty$, the extracellular glucose level ($G_e$) is delivered to the islets at a rate defined by $\tau_G$. This new $G_e$ value would induce a new $Ca_{avg}$ that closes the feedback loop.
TABLE 2 Variables used in the experiments

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Description</th>
<th>Unit</th>
</tr>
</thead>
<tbody>
<tr>
<td>( \bar{C}<em>{a</em>{\text{avg}}} )</td>
<td>Average ([Ca^{2+}]), measured every 20 s</td>
<td>nM</td>
</tr>
<tr>
<td>( l_{\text{avg}} )</td>
<td>Average insulin secretion</td>
<td>A.U.</td>
</tr>
<tr>
<td>( G_e )</td>
<td>Asymptotic glucose response function</td>
<td>mM</td>
</tr>
</tbody>
</table>

producing a slow feedback response. Experiments were performed using \( S_G \) values of 0.1, 1, 10, 50, and 100 and \( \tau_G \) values of 50, 300, 450, 600, and 900 s.

A group of 10 islets was initially stimulated with a constant 10 mM glucose (Fig. 3 A, red line) which induced uncoordinated \([Ca^{2+}]_i\) oscillations among islets. The color density map at the bottom of Fig. 3 A shows that all islets responded to the increased glucose with \([Ca^{2+}]_i\) oscillations, but these oscillations were out of phase. Accordingly, the \( \bar{C}_{a_{\text{avg}}} \) response (black line) did not exhibit coherent rhythmicity. After 46 min, the extracellular glucose level was adjusted dynamically, as described above, closing the feedback loop. With the parameter values used in Fig. 3 A (\( S_G = 1, \tau_G = 50 \) s), the individual islets synchronized as evident by the vertical stripes in the color density map. Oscillations in both \( \bar{C}_{a_{\text{avg}}} \) and the extracellular glucose level were produced with a period of 4.7 and 4.8 min, respectively. The insulin concentration (Fig. 3 A, blue curve) was measured simultaneously with \([Ca^{2+}]_i\), and a similar rhythmic pattern was observed (4.9 min), and was in phase with \( \bar{C}_{a_{\text{avg}}} \). Fura-2 ratiometric images of the islet population from different times are shown in Figs. 3 B and C. The images in Fig. 3 B were taken at two points during the delivery of constant glucose and show heterogeneous \([Ca^{2+}]_i\) values for the different islets, indicating that the islets were unsynchronized. In Fig. 3 C, the images were acquired when the feedback was on, and show a homogeneous \([Ca^{2+}]_i\) as would be expected from a synchronized population of islets. The movie that the images in Fig. 3 B and C were taken is provided in the Supporting Material (Movie S1).

To examine how the dynamic glucose feedback varied with the two feedback parameters, other combinations of \( S_G \) and \( \tau_G \) were used and representative experiments are shown in Fig. 4 A–D. For the same \( S_G \), increasing \( \tau_G \) decreased the overall glucose amplitude. For example, Fig. 4 A shows an experiment that used five islets with parameters \( S_G = 10, \tau_G = 50 \) s and produced \(~ 2 \) mM amplitude glucose oscillations with large amplitude oscillations in \( \bar{C}_{a_{\text{avg}}} \) indicating the islets were synchronized (\( \lambda = 0.97 \)). In Fig. 4 B, five different islets were used with \( S_G = 10, \tau_G = 600 \) s. No pulsatility in \( \bar{C}_{a_{\text{avg}}} \) was apparent and the glucose level was relatively flat, indicating that the islets were not synchronized (\( \lambda = 0.33 \)).

Figs. 4 C and D show representative results of the effect of \( S_G \) on synchronization. In Fig. 4 C, six islets were used with parameters \( S_G = 50, \tau_G = 50 \) s, and in Fig. 4 D, eight islets were used with \( S_G = 100, \tau_G = 50 \) s. Synchronization only occurred with the smaller \( S_G \) value (\( \lambda = 0.93 \) and 0.53 for Fig. 4 C and D, respectively). The effects of \( S_G \) and \( \tau_G \) on the average \( \lambda \) and average induced glucose amplitudes are summarized in Fig. 4 E. The shapes of the points indicate values of \( \tau_G \), whereas the colors indicate values of \( S_G \), with error bars corresponding to +/- 1 standard deviation.

FIGURE 3 Islet synchronization occurs when the feedback loop is closed. (A) In this example, the feedback loop was closed using parameter values of \( S_G = 1, \tau_G = 50 \) s with 10 islets in the microfluidic chamber. At time 46 min, the feedback was turned on and \( G_e \) changed in response to \( \bar{C}_{a_{\text{avg}}} \). The glucose level is shown at the top in red, whereas \( \bar{C}_{a_{\text{avg}}} \) (black line) and insulin (blue line) are shown below the glucose. The \([Ca^{2+}]_i\), from all islets is shown in the color density map at the bottom. Scale bars for all measurements are shown adjacent to the graphs. (B) Two fluorescent ratiometric images of Fura-2 taken of the islets shown in part A before the onset of feedback. The image on the left was taken at time 25 min and the one on the right at 32 min. (C) The two images correspond to times 51 min (left) and 63 min (right), when feedback was on. The color scale bar for B and C are shown below C.
feedback would be both oscillations of insulin and glucose, negative feedback onto islets, all of which sense the glucose could indeed entrain oscillations in the islet Ca$^{2+}$, appropriate period and amplitude, here imposed on the islets, loop manner to verify that glucose oscillations of appropriate period and amplitude, here imposed on the islets, could indeed entrain oscillations in the islet Ca$^{2+}$ concentration. Similar experiments have been performed with single (20,26,36) and groups (25,37,38) of islets, but the ranges of glucose periods and amplitudes were not investigated. We found that islets were entrained by sinusoidal glucose levels with periods near the natural oscillation period and with glucose amplitudes above 0.5 mM. It has been reported that even small amplitude oscillations of glucose (~0.3 mM, or 5%) (39) entrains in vivo pulses of insulin, and our results are similar in this manner. Recent work also demonstrates that islets can be entrained at harmonics of the glucose oscillation, so that a 10 min glucose wave can entrain two pulses of [Ca$^{2+}$]i (38). We too observed a range of harmonics, ranging from 3:1 to 1:2 but only show the 1:1 entrainment in Fig. 1 A.

After confirmation that islets can be entrained to glucose oscillations with the appropriate combinations of amplitudes and periods, the closed-loop experiments were performed where the response of the islet population was used to update the glucose level, via Eqs. 3 to 6, to be delivered to the islets. The two parameters we tested, S, and r, were based on liver glucose absorption and the time the new glucose level was delivered to the islets, respectively. Because physiological values are not known for these parameters, a range of values was tested. As shown in Fig. 3, insulin release and $Ca_{avg}$ became synchronized with a concomitant production of oscillatory glucose levels using the appropriate model parameters. In all closed-loop experiments performed, the glucose concentration was 180° out of phase with both insulin and $Ca_{avg}$. This phase difference is attributable to the asymptotic glucose response function and was also observed when this dynamic feedback interaction was mathematically modeled (see Fig. 10 of (19)). Fig. 3 also confirms that the use of $Ca_{avg}$ is a suitable marker for $I_{avg}$ in later experiments because oscillations of insulin and $Ca_{avg}$ were in phase.

The oscillations seen in Fig. 3 A in the glucose trace, color density map, and the $Ca_{avg}$ trace were the result of mutual interactions between the islets in the microfluidic chamber and the dynamic glucose feedback. That is, with the glucose feedback action, the majority of islets in the chamber synchronized their activity to produce a rhythm whose period was determined by the intrinsic oscillatory periods of the islet population. As expected, the amplitude and period of the glucose oscillation were within the entrainment window measured in the open-loop experiments (Fig. 1 D).

The ranges of r and S that produced a synchronized population were explored, and it was found that increasing either r or S produced a decrease in the glucose amplitude with a concomitant drop in λ. With large values of S (≥100), the glucose response function was too flat to reflect changes in $I_{avg}$ and the islets did not synchronize. With large values of r, the rate of response of the feedback system was reduced and the extreme values of the response function ($G_{max}$ and $G_{min}$) were never reached, reducing the amplitude of the glucose oscillations. As shown by Fig. 4 E, it was essential for the amplitude of the glucose oscillations to be >0.4 mM. Combinations of S and r that produced glucose amplitudes between 0.40 to 3 mM were successful at synchronizing islet populations (λ ≥ 0.7), but below

DISCUSSION

We set out to test the hypothesis that a dynamic interaction between the pancreas and liver could synchronize the oscillations of activity and insulin secretion across an islet population (19). By taking up glucose, the liver provides negative feedback onto islets, all of which sense the glucose level and respond accordingly. The result of this negative feedback would be both oscillations of insulin and glucose, both of which have been observed in vivo (1,2,7–9,24). Initial tests of the hypothesis were performed in an open-loop manner to verify that glucose oscillations of appropriate period and amplitude, here imposed on the islets, resulted from the feedback interactions for all values in the feedback system. (A) $S = 10$ and $r = 50$ s; (B) $S = 10$ and $r = 600$ s; (C) $S = 50$ and $r = 50$ s; and (D) $S = 100$ and $r = 50$ s. In traces A and C, the islet population was synchronized (λ= 0.97 (five islets) and 0.93 (six islets), respectively). With the large $r$ value in B and the large $S$ value in D, the amplitudes of the glucose oscillations were small and islets were not synchronized (λ = 0.33 (five islets) and 0.53 (eight islets), respectively). (E) The average λ are shown versus the average glucose amplitude that resulted from the feedback interactions for all experiments. The colors and shapes of the data points correspond to the values of $S$ and $r$ as shown in the legend and error bars correspond to ±1 standard deviation. At low values of $S$ and $r$, glucose amplitudes > 0.4 mM are produced and the islets synchronize (high λ).
0.40 mM were not. These results are consistent with the open-loop experiments in which glucose oscillations were imposed; amplitudes below 0.5 mM were largely outside of the islet entainment window (Fig. 1 D).

Although our dynamic glucose model is simple, it captures the key element of negative feedback provided by the liver, and results in islet oscillations with a period similar to those observed in vivo (1–3,7–9). For combinations of $S_G$ and $\tau_G$ that produced $\lambda \geq 0.7$, the $Ca_{avg}$ and glucose oscillation periods were within the range of 5 to 8 min. This is important because it is what would be expected if the rhythmicity were attributable to the intrinsic period of the islets, rather than the glucose feedback system. The liver is known to take up 25% to 35% of an oral glucose load (40–42) with blood circulation taking ~1 min. Combined with the inhibitory effect of insulin on hepatic glucose production, the liver has the capability of making large changes in the blood glucose level and the effects of these changes can be rapid.

Negative feedback provided by the liver is not the only plausible mechanism for islet synchronization. Intrapancreatic ganglia could provide pacemaking activity to entrain islet oscillations (10,17,43), but at present there are no data suggesting that ganglia neurons exhibit coordinated pacemaking activity. It is also possible that islets communicate through chemical messengers in a paracrine fashion, and this would account for observations of pulsatile insulin in the perfused pancreas (23,44,45) and from superfused islets (46,47). However, no such paracrine factor has been isolated, and we find no coordination of islet activity with a constant glucose level (Fig. 3). Given the physiological importance of insulin pulsatility, it is likely that there are overlapping mechanisms for synchronization, and indeed, no one mechanism can explain all the reports of synchronization in different preparations. Our data support the hypothesis that a dynamic interaction between the pancreas and the liver could synchronize intrinsic islet oscillations, and can provide one mechanism for islet synchronization in a number of species, including humans.

**SUPPORTING MATERIAL**

One movie is available at [http://www.biophysj.org/biophysj/supplemental/S0006-3495(14)00398-1](http://www.biophysj.org/biophysj/supplemental/S0006-3495(14)00398-1).

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