Automated Adaptation of Insulin Treatment in Type 1 Diabetes

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Abstract— Individuals with type 1 diabetes (T1D) need lifelong insulin therapy to compensate for the lack of endogenous insulin due to the autoimmune damage to pancreatic beta-cells. Treatment is based on basal and bolus insulin, to cover fasting and postprandial periods, respectively, according to three insulin dosing parameters: basal rate (BR), carbohydrate-toinsulin ratio (CR), and correction factor (CF). Suboptimal BR, CR, and CF profiles leading to incorrect insulin dosing may be the cause of undesired glycemic events, which carry dangerous short-term and long-term effects. Therefore, correct tuning of these parameters is of the utmost importance. In this work, we propose a new algorithm to optimize insulin dosing parameters in individuals with T1D who use a continuous glucose monitor and an insulin pump. The algorithm was tested using the University of Virginia/Padova T1D Simulator and led to an improvement in the quality of glycemic control. Future efforts will be devoted to test the algorithm in human clinical trials.

I. INTRODUCTION

In type 1 diabetes (T1D), life-long insulin replacement is needed to compensate for the practically absent internal insulin secretion deriving from the autoimmune destruction of pancreatic beta-cells [1]. Consequently, the quality of glycemic control in T1D is heavily dependent on multiple daily treatment decisions by the patient, to account for a wide variety of factors influencing insulin demand, e.g., circadian rhythms, physical activity, food, and stress. In this context, despite the improving accuracy of glucose monitoring devices [2],[3], the availability of modern insulin analogs [4], and the growing proliferation of new technologies [3],[5]-[7], glycemic control remains a challenge in T1D [8], with complication rates and excess mortality still significantly higher when compared to the general population [9],[10].

Traditionally, intensive insulin treatment implemented through continuous subcutaneous insulin infusion (CSII, also referred to as insulin pump therapy), includes basal insulin administered to cover the overnight and fasting periods, and bolus insulin given with meals to cover carbohydrate consumption and correct postprandial hyperglycemia, in an attempt to mimic insulin secretion in health [11]. However, insulin therapies are not as efficient as the natural endogenous insulin secretion, and suboptimal insulin dosing is still common in the management of T1D, causing potentially life-threatening hypoglycemic episodes [12],[13] and/or sustained hyperglycemia which in turn leads to the development and progression of long-term diabetes comorbidities [14],[15]. Therefore, individuals with T1D face a life-long optimization challenge [16]: to reduce average glucose (BG) while simultaneously avoiding blood

hypoglycemia, which has been indicated as the limiting factor in the management of diabetes [13].

To help individuals with T1D solve this problem, technological options are available, in the form of open-loop decision support systems [17],[18] or closed-loop automated insulin delivery (i.e., the artificial pancreas) [19],[20]. Especially in the former case but also in the latter, the optimal trade-off between glycemic control and iatrogenic hypoglycemia is typically achieved through periodical reviews of BG traces, to adjust daily profiles of basal rate (BR) – i.e., the amount of basal insulin delivered throughout the day – and carbohydrate-to-insulin ratio (CR) and correction factor (CF) – i.e., parameters used to dose prandial insulin. If a glycemic pattern is identified, optimized insulin dosing parameters (i.e., optimized BR, CR, and CF profiles) are calculated and implemented. This can be a timeconsuming and onerous task, requiring data to be downloaded from multiple devices and subjectively evaluated. In this context, innovative digital technologies represent a promising alternative to lessen the burden on health care providers and provide an objective way to analyze data and extract information otherwise not readily available.

In this work, we introduce a novel method to optimize insulin dosing parameters in individuals with T1D who use a continuous glucose monitoring (CGM) system and follow CSII therapy. The method relies on CGM data (i.e., BG measurements collected and stored every five minutes), basal and bolus insulin records, and meal information to provide weekly recommendations on optimal BR, CR, and CF profiles to be deployed by the user. Here, we present results obtained by testing the optimization algorithm in an FDAaccepted simulation environment known as the University of Virginia/Padova T1D Simulator [21]-[23]. The simulation platform includes a large, fully-identified model of glucose metabolism in T1D and a population of 100 virtual adult subjects displaying key metabolic behaviors observed in the general population of individuals with T1D. In simulation, the algorithm was successful at identifying underlying glycemic patterns and correcting insulin therapy, thereby improving the overall quality of glycemic control.

II. MATERIALS AND METHODS

A. Optimization of Basal Insulin Dosing

BR is optimized weekly relying on an algorithm which assesses the impact of BR alone (i.e., in the absence of meals and boluses) on the glycemic levels. In summary, the method uses an individualized model of glucose metabolism capable of describing the glycemic response to meals and insulin, thus enabling to parse out the effect of BR and prandial disturbances on BG fluctuations. The model receives as inputs injected insulin (J), consumed meals (M),

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and a residual metabolic signal needed to explain the experimental data (ω). Optimal BR is computed as the insulin input to the model, that allows to maintain the BG response flat at the desired target BG (here, 110 mg/dL), in the presence of ω while setting the meal input to zero.

In detail, the method relies on the following three steps. *1. Individualization of the metabolic model.* The metabolic model used within the algorithm can be written as the following discrete-time, linear, time-invariant model:

$$x(k+1) = Ax(k) + B_{ins}J(k) + B_{meal}M(k) + B_{\omega}\omega(k)$$

$$y(k) = Cx(k)$$
(1)

where k is the discrete timestamp; x is the metabolic state vector including plasma glucose concentration, insulin concentration in the subcutaneous space and in plasma, insulin action, and amount of carbohydrates in the stomach and gut; y is the model output coinciding with plasma glucose concentration; and A, B_{ins} , B_{meal} , B_{ω} , and C are statespace matrices describing the interaction between glucose and insulin in the presence of a meal. To reflect individual metabolic behaviors, the model is identified on the available CGM data, inputting known J and M to the model, while setting ω to zero. Model identification is performed using maximum-a-posteriori Bayesian estimation, relying on prior knowledge on the metabolic parameters available from the literature. The interested reader is referred to [24] for a more detailed description of the model, the meaning of the various parameters, and the formulation of the estimation problem.

2. Estimation of ω . The residual metabolic signal ω is estimated by regularized deconvolution via inversion of the individualized model outlined above, to capture unmodeled phenomena and glucose dynamics, thereby allowing to well-describe the experimental CGM data. The theoretical bases for the computation of ω are detailed in [24].

3. Estimation of the optimal BR. Upon estimation of ω , a second model inversion is performed, again by regularized deconvolution, to determine the optimal insulin input that allows to obtain a BG profile flat at around 110 mg/dL, in the absence of any meal input but considering ω .

The described procedure is performed for each day, providing daily optimal BR traces. Then, at the end of each week, the median BR across days is computed to determine the final BR profile to be deployed over the following week.

B. Optimization of Prandial Insulin Dosing

In addition to BR, the algorithm provides optimal values of prandial dosing parameters (i.e., CR and CF), based on the placement of the average minimum postprandial BG (mPBG) via minimization of its variance while controlling for the amount of tolerated exposure to hypoglycemia.

In the algorithm, *mPBG* is described by a linear model with the classical prandial predictors:

$$mPBG = BG_{TGT} + \beta_{BG}(BG_{meal} - BG_{TGT}) + (2) + \beta_{CHO}CHO_{meal} + \beta_{ins}INS_{meal}$$

where BG_{TGT} is the target postprandial BG (here set at 110 mg/dL); BG_{meal} , CHO_{meal} , and INS_{meal} are mealtime CGM value, meal carbohydrate amount, and meal-related insulin

dose; and the β s are model parameters identified using historical data and a weighted least-square estimator with regularization terms on the parameters based on the subject's current CR and CF values. The model is identified for each 30-min timespan of the day (48 identifications in total), with weights assigned to meal events that decrease exponentially with the datapoint absolute age and its distance from the current identification timespan in terms of time of day.

In the *mPBG* model of Eq. (2), *INS_{meal}* can be controlled through the typical prandial insulin dosing formula:

$$INS_{meal} = \frac{CHO_{meal}}{CR} + \frac{BG_{meal} - BG_{TGT}}{CF} .$$
 (3)

If Eq. (3) is substituted in Eq. (2), an expression depending on CR and CF is obtained. Thus, the following optimization problem can be formulated to determine CR and CF:

$$\min_{CR,CF} STD(mPBG)$$
t. $AVG(mPBG) - \gamma STD(mPBG) = thHypo$
(4)

where $STD(\cdot)$ and $AVG(\cdot)$ are the standard deviation and average operators, and γ (set to 1.5) and *thHypo* (set to 70 mg/dL) are aggressiveness parameters determining the amount of tolerated hypoglycemia. If BG_{meal} and CHO_{meal} are assumed to be independent random variables with finite variance, STD(mPBG) and AVG(mPBG) can be written as:

$$STD(mPBG) = \sqrt{\theta_{CHO}^2 VAR(CHO_{meal}) + \theta_{BG}^2 VAR(BG_{meal})}$$
$$AVG(mPBG) = BG_{TGT} + \theta_{CHO} AVG(CHO_{meal}) + (5)$$
$$+ \theta_{BG} AVG(BG_{meal})$$

where $VAR(\cdot)$ is the variance operator, AVG and VAR of CHO_{meal} and BG_{meal} are computed empirically using the same weighting schema used above, and θ_{CHO} and θ_{BG} are:

$$\theta_{CHO} = \beta_{CHO} + \frac{\beta_{INS}}{CR}$$
 and $\theta_{BG} = \beta_{BG} + \frac{\beta_{INS}}{CF}$. (6)

Given Eq. (5), the minimization problem outlined in Eq. (4) is a quadratic problem in θ_{CHO} and θ_{BG} with constraints. The constraint gives a conic section that is sampled and from which the minimum-*VAR*(*mPBG*) point is extracted. From the optimal θ_{CHO} and θ_{BG} (one for each 30-min timespan), relying on Eq. (6) and the values of the β s obtained from the identification of the model in Eq. (2) in the same timespans, optimal CR and CF values are obtained.

The interested reader is referred to [25] for a comprehensive description of the outlined method.

C. Simulation Testing

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The optimization algorithm was tested in a 6-week simulation study including 100 virtual subjects with T1D. BG and insulin/meal records were available throughout the study.

Three treatment arms were simulated: a) control arm treated with original insulin dosing parameters built in the simulating platform and factory-designed for each virtual subject (ORIG); b) control arm treated with altered insulin dosing parameters obtained by imposing random deviations with respect to the original parameter configuration (MOD); c) experimental arm treated with dosing parameters optimized at the end of each study week and deployed over the week following the optimization, starting from the altered parameter configuration defined in the previous arm (OPT).

During each study day, each virtual subject received three meals: breakfast (B), lunch (L), and dinner (D). Meals time and amount were sampled from uniform distributions with the following supports: [0600-0800] and [30-50] grams of carbohydrates (grCHO) for B, [1200-1400] and [60-80] grCHO for L, and [1800-2000] and [40-60] grCHO for D.

Metabolic and behavioral variability were included in the simulations to render the virtual study more realistic. Metabolic variability was generated by imposing daily patterns in insulin sensitivity, based on circadian variability observed in real-world data [23]. Behavioral variability was generated by including snacks given between meals ([15-25] grCHO at [0930-1030] and [1530-1630]), carbohydrate counting errors (up to 30% of the real amount), and bolus delays (up to 60 mins after the meal was consumed).

Treatment arms for each virtual subject were identical, with the same meals and metabolic/behavioral variability.

Algorithm performance was assessed using established metrics of glycemic variability and quality of glycemic control, including: average, standard deviation, and coefficient of variation of BG values; time spent in the target glycemic ranges of 70-140 and 70-180 mg/dL; time spent in the hypoglycemic range <54 and <70 mg/dL; and time spent in the hyperglycemic range >180 and >250 mg/dL. For each arm, these metrics were computed weekly from the BG traces. The comparison of week 6 to week 1 in OPT, focusing on time spent in 70-180 mg/dL (TIR), <70 mg/dL (TBR), and >180 mg/dL (TAR), was the primary analysis. All results are reported as mean \pm standard deviation across subjects.

III. RESULTS

Out of the 100 subjects, 50/50 subjects had their original therapy modified to generate insulin over/under dosing.

In OPT, TIR improved from $57.5\% \pm 21.9\%$ in week 1 to $80.4\% \pm 13.8\%$ in week 6; TBR improved from $9.3\% \pm 13.6\%$ to $1.5\% \pm 3.1\%$; and TAR improved from $33.2\% \pm 27.6\%$ to $18.1\% \pm 14.1\%$. These results are also summarized in Figure 1, where the evolution of TIR, TBR, and TAR across the six study weeks is displayed for the three treatment arms. An exhaustive comparison between week 1 and week 6 in the OPT treatment arm is presented in Table I.

TABLE I.	ALGORITHM PERFORMANCE

	Week 1	Week 6
BG average [mg/dL]	156.9 ± 47.4	143.4 ± 19.4
BG standard deviation [mg/dL]	44.9 ± 15.6	38.3 ± 14.2
BG coefficient of variation [%]	29.9 ± 10	26.1 ± 6.7
Time in 70-140 mg/dL [%]	35.7 ± 22	55.7 ± 14.4
Time in 70-180 mg/dL [%]	57.5 ± 21.9	80.4 ± 13.8
Time below 54 mg/dL [%]	2.2 ± 3.5	0.2 ± 0.6
Time below 70 mg/dL [%]	9.3 ± 13.6	1.5 ± 3.1
Time above 180 mg/dL [%]	33.2 ± 27.6	18.1 ± 14.1
Time above 250 mg/dL [%]	9.4 ± 15.2	2.9 ± 6.3



Figure 1. Evolution of TIR (Panel A), TBR (Panel B), and TAR (Panel C) across the six study weeks, comparing the three treatment arms (ORIG: solid line and triangles; MOD: solid line and diamonds; OPT: dashed line and stars).

IV. DISCUSSION

Insulin dosing parameters are needed to make daily treatment decisions in T1D. Wrong parameters are common due to the difficulty of extracting relevant information from the large amount of available data (e.g., CGM, insulin records, meal records). Suboptimal insulin dosing resulting from incorrect dosing parameters may lead to dangerous glycemic events, like hypoglycemia [13] – which if not promptly treated can lead to coma and death – and sustained hyperglycemia and elevated glycemic variability [14] – both identified as independent risk factors for the development and progression of long-term diabetes comorbidities.

New technologies like the artificial pancreas and more general decision support systems have helped the management of T1D and improved the quality of glycemic control [17]-[20]. At the core of these technologies, optimal insulin dosing parameters are needed to maximize system performance and reduce the risk for low/high-BG episodes.

In this work, we proposed an adaptive method to optimize insulin dosing parameters in individuals with T1D who use CGM and follow CSII therapy. The method relies on CGM data, insulin records, and meal information to provide recommendations to patients about optimal dosing parameters to use, based on the analysis of glycemic patterns extracted from previous individual data collected in the field. A proof-of-concept in silico study demonstrated that the method performs well in simulation, increasing time spent in the target euglycemic range, while reducing exposure to hypoglycemia, hyperglycemia, and glycemic variability.

The presented algorithm has large potential if embedded within decision support systems to provide suggestions to patients upon request or on a regular schedule (e.g., weekly), regarding needed therapy modifications. The method can also be personalized to reflect individual treatment preferences (e.g., overnight and postprandial glycemic targets). Further, the method can be extended to work in the presence of different insulin therapies (e.g., multiple daily insulin injections or within an artificial pancreas).

V. CONCLUSIONS

This paper introduces a novel algorithm to optimize insulin dosing parameters in individuals with T1D using CGM and CSII therapy. The method was successful in simulation studies. Clinical testing will be performed next, to further validate the technology.

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