Differential Treatment Response to Insulin Intensification Therapy: A Post Hoc Analysis of a Randomized Trial Comparing Premixed and Basal-Bolus Insulin Regimens

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ABSTRACT

Introduction: Identification of subgroups of patients that may benefit most from certain treatment is important because individual treatment response varies due to multiple contributing factors. The present study used the subgroup identification based on the differential effect search (SIDES) algorithm to identify subgroups with different treatment responses to insulin intensification therapies.

Methods: This was a post hoc analysis of a 24-week, multicenter, open-label, randomized, parallel study comparing prandial premixed therapy (PPT) to basal-bolus therapy (BBT). Patients with type 2 diabetes mellitus were randomized to PPT (insulin lispro mix 50/50 thrice daily with meals) or BBT (glargine at bedtime plus mealtime insulin lispro) insulin intensification therapies. The SIDES algorithm was used to identify the subgroups from at-goal patients [glycated hemoglobin (HbA1c) <7.0% (53.0 mmol/mol) at the end of 24 weeks; n = 182] who could have benefitted from insulin intensification therapies.

Results: Baseline characteristics of overall at-goal patients were comparable between PPT and BBT groups. The SIDES algorithm identified patients with race other than Caucasian (i.e., African–American, Asian, and Hispanic) and baseline fasting blood glucose (FBG) <8.89 mmol/L as a subgroup that could respond better to PPT relative to BBT than the overall at-goal patient population. In this identified subgroup population, the HbA1c mean (standard deviation) changes from baseline to endpoint in PPT and BBT groups were −2.27 (0.88)% versus −2.05 (0.75)%; p = 0.40, respectively; while in the overall at-goal patients, the HbA1c changes were −2.17 (0.79)% versus −2.34 (1.00)%; p = 0.19, respectively.

Conclusions: The preliminary results showed that the subgroup of patients with race other than Caucasian and FBG <8.89 mmol/L may respond better to premixed intensification therapy. This result provides some preliminary information for further investigation in prospective studies.

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INTRODUCTION

By 2040, the prevalence of type 2 diabetes mellitus (T2DM) is estimated to be 642 million cases (10.4%) across the globe [1]. T2DM is a serious health epidemic with multiple contributing factors, including but not limited to diet, lifestyle, and genetic predisposition [2].

Insulin therapy is recommended to compensate for inadequate insulin secretion due to progressive pancreatic beta cell dysfunction after failure on oral antihyperglycemic medications (OAMs) and to combat a worsening glycemic profile. Intensive insulin therapy is recommended in patients who have failed previous insulin therapy, and two main options of intensification are basal-bolus and premixed insulin regimens [3, 4]. According to 2017 American Diabetes Association (ADA) guidelines [5], premixed insulin, thrice daily, is recommended as intensification therapy when glycated hemoglobin (HbA1c) could not be controlled with premixed insulin twice-daily regimen. The guidelines also suggested switching patients from premixed analogue insulin thrice daily to basal-bolus regimen or vice versa if HbA1c targets are not being met and depending on other patient considerations. Intensive glycemic control reduces microvascular and macrovascular complications. However, multiple contributing factors, such as genetic variations and clinical, sociodemographic, comorbidity, and behavioral factors, may affect the outcome of the antidiabetic treatment [6, 7]. Hence, choosing an optimal insulin intensification therapy is important and needs exploration.

In clinical trials, the overall or average treatment effect in the study population may not reflect the subgroup and/or individual treatment outcomes. The overall or average treatment effect obscures the individual meaningful benefits or risks, or both, because some patients may have more beneficial outcomes than average from treatment, whereas others do not benefit from treatment or may be harmed [8]. In this regard, identification of patient subgroups that can benefit the most from certain treatments is of clinical importance for patients and physicians. Thus, personalized medicine or tailored therapeutics has been developed, leading to a substantial improvement in the ratio of benefits to risks for this set of patients.

There are multiple approaches for this required subgroup identification. We can define the subgroups with commonly used cutoff values like age of patients (e.g., 60 or 65 years), but the actual cutoff will be data-driven. The other most commonly used subgroup cutoff is median (or quartiles) of the interested factors. Though these two approaches can explore the subgroups of a predefined feature, they fail to identify the subgroup of interest that derives greater benefit compared to the whole population. There has always been a need to identify such subgroups quickly and efficiently. In this regard, we applied the subgroup identification based on the differential effect search (SIDES) algorithm to identify patient subgroups that may benefit more from the prandial premixed therapy (PPT) than from basal-bolus therapy (BBT) in the at-goal patient population that had responded to these treatments [achieved <7.0% (53.0 mmol/mol) HbA1c at the end of treatment period]. SIDES is a novel recursive partitioning method used to identify subgroups of patients with enhanced treatment effects based on a differential effect search algorithm [9].

Identification of subgroups of patients that may benefit most from specific treatment is important for individualized therapy. The present study used the SIDES algorithm to identify subgroups with different treatment responses to insulin intensification therapies. This was the first analysis that used the SIDES algorithm to identify patient subgroups that may respond better to PPT than to BBT. Knowing the subgroup that responds better to specific insulin intensification treatment may provide guidance in generating a preliminary hypothesis to explain heterogeneity that could be tested in confirmatory, prospective, randomized clinical trials.
METHODS

Study Design

These results were a post hoc analysis of a 24-week, multicenter, open-label, randomized, parallel, active-controlled study comparing PPT with BBT [10].

This study included patients (a) with T2DM (b) who were 30–75 years of age, inclusive, at the time of screening, (c) who had been treated for at least 90 days with insulin glargine once daily (≥30 U/day) in combination with OAMs (sulfonylurea or meglitinide, metformin, or thiazolidinedione and, in the opinion of investigator, required further intensification of therapy, and (d) who had inadequate glycemic control [HbA1c ≥7.5% (58.5 mmol/mol) and ≤12.0% (107.7 mmol/mol)] at screening [10].

Major exclusion criteria were (a) patients who had severe hypoglycemia (more than one episode 6 months before study entry), (b) patients with excessive insulin resistance (receiving dose >2.0 U/kg at the time of randomization), (c) patients who were obese [body mass index (BMI) >45 kg/m²], (d) patients who had cardiac diseases with functional status Class III or IV as per the New York Heart Association [11], (e) patients who had renal disorders [serum creatinine ≥135 μmol/L (1.5 mg/dL) for men and ≥100 μmol/L (1.2 mg/dL) for women], or (f) patients who had liver disease (alanine aminotransferase at least three times the upper normal limit of reference range) [10].

Eligible patients, based on inclusion and exclusion criteria, were randomized to receive either PPT or BBT, and either to an aggressive dosing algorithm (based on plasma glucose levels and total daily insulin) or a conservative dosing algorithm (plasma glucose levels). Patients randomized to the PPT group received insulin lispro mix 50/50 (50% insulin lispro and 50% insulin lispro protamine suspension) three times daily before meals. If fasting blood glucose (FBG) targets were not achieved with insulin lispro mix 50/50, the evening preprandial dose could be changed to an injection of insulin lispro mix 75/25 (25% insulin lispro and 75% insulin lispro protamine suspension). The starting dose was based on the insulin glargine total dose at study entry divided into three equal doses of lispro mix 50/50 before meals. Patients randomized to the BBT group received insulin glargine once daily along with preprandial insulin lispro. The starting dose of this regimen was also based on the total glargine dose at study entry, with 50% administered as glargine and the other 50% given in three equal doses of lispro. Patients in both arms continued to receive OAMs; however, insulin secretagogues (sulfonylureas and meglitinides) were discontinued at randomization. The dose of insulin is decided with a goal of achieving preprandial blood glucose of less than 110 mg/dL (6.11 mmol/L) and a 2-h postprandial blood glucose of less than 140 mg/dL (7.78 mmol/L) with no hypoglycemia [blood glucose <72 mg/dL (4.0 mmol/L)]. Detailed study design, dosing schedule, dosing algorithm, efficacy, and safety results of this study were previously published [10].

All procedures followed were in accordance with the ethical standards of the responsible committee on human experimentation (institutional and national) and with the Helsinki Declaration of 1964, as revised in 2013. Informed consent was obtained from all patients for being included in the study.

Statistical Analysis

Of 547 patients who entered the study, 374 were randomized by an interactive telephone system at a 1:1 ratio to the PPT and BBT groups, respectively [10]. Of these 374 patients, data for 182 patients (81 on PPT and 101 on BBT) who provided a baseline HbA1c and achieved a target 24-week HbA1c [<7.0% (53.0 mmol/mol); responders, also referred to as at-goal patients] were examined and explored. The data were analyzed by the SIDES algorithm [9], instead of specifying subgroup by anecdotal evidence, to explore a subgroup of patients (out of those 182 at-goal patients) who could show favorable responses to PPT.

The SIDES method, a novel recursive partitioning procedure, allows direct evaluation of the treatment effect in subgroups and is
effective in evaluating modest-sized and large databases. This approach helps in searching for subgroups with a large differential effect relative to the study population. Also, this method can be applied as a part of prospectively defined tailored-therapy strategy and retrospectively throughout drug development across early- and late-phase trials combined to detect important predictive subgroups as early as possible [9]. The algorithm directly searches for large treatment effect simultaneously among many covariates. The traditional way of performing subgroup analyses involves prespecifying the subgroup of interests by specifying a set of variables and the corresponding cutoff points. The differences of treatment effects between comparator groups can then be examined by a parametric regression. Such an approach necessitates predefining subgroups of interests, which is often difficult when there are many covariates. Also, for any particular dataset, the subgroup suggested by our experience may not be the subgroup that had the maximal differential treatment effect. The SIDES algorithm instead allows the data to guide the process of identifying the subgroups, thus eliminating the need to predefine potential cutoff value. Insights suggested by data in this manner can provide insights that might not be evident on an a priori basis. This would provide a unique advantage for exploratory analyses, for which the primary interest is often to discover new evidence instead of confirming a priori hypotheses. The SIDES algorithm was implemented by a subgroup identification tool (SIT). This tool was developed by Eli Lilly and Company and was used to support the refinement of clinical development and post hoc analyses of study results [9]. Few studies using the SIDES method to identify subgroups of interest have already been published [12–15].

For the purposes of this analysis, the SIDES algorithm search criteria incorporated stratification by baseline HbA1c to control for the differences between patients in baseline HbA1c because it was well accepted that baseline HbA1c would have a direct effect on the change in HbA1c from baseline. Without this stratification, the search may likely end up with subgroups that have the largest difference in baseline HbA1c. The stratification levels used were quintiles estimated by the SIT for the distribution of baseline HbA1c for the 182 patients included in the analysis dataset. The variable importance score was calculated as the sum of the negative natural log of the splitting criterion [9] over all explored subgroups where the covariate was one of the splitters. It may be viewed as a measure of the relative importance; the covariate contributed to the subgroup identification process. It also provided a benchmark value computed using permutation-based maximum variable importance. To perform the permutation, first, the treatment labels for the 182 at-goal patients were randomly permuted. Second, new analyses using the SIDES algorithm were performed using the same set of covariates, and the maximum variable importance was recorded (this maximum variable importance may correspond to different covariate at different runs). These two steps were repeated 500 times which yielded 500 maximum variable importance scores. We then calculated the mean and standard deviation (SD) of those 500 scores and used them as a benchmark value. Covariates with variable importance greater than the mean of permutation-based maximum variable importance may be considered a strong indication for an important subgroup effect, and covariates with variable importance greater than the mean plus SD of permutation-based maximum variable importance may be considered a very strong indication of an important subgroup effect. The detailed statistical method was described in detail in Lipkovich et al. [9] and Hardin et al. [12].

All p values were provided as a descriptive measure for the difference between treatments and were computed using a two-sample t test for continuous variables and Fisher’s exact test for categorical variables. Such summaries and comparisons were made between PPT and BBT treatment groups in the overall at-goal patients and in the identified subgroup.

Note that SIDES is just a data-mining algorithm that can efficiently explore the features of the observed data and provide a data-driven subgroup of interest. It does not alter the observed data, and the summaries and
comparisons are generated using the common and conventional statistical tools.

**Outcome and Predictor Variables**

The outcome variable input into the SIT was change from baseline in HbA1c for patients who achieved control at 24 weeks \(<7.0\% (53.0 \text{ mmol/mol})\). This measure was used as the response variable for the subgroup identification analysis with the SIT, and the following variables were used as candidate predictors of treatment effect: baseline FBG (from self-monitored blood glucose), race/ethnicity, baseline morning 2-h postprandial glucose (from self-monitored blood glucose), duration of diabetes, age at study entry, gender, baseline BMI, baseline use of thiazolidinedione (Yes or No), baseline use of metformin (Yes or No), and randomized treatment. The SIT compared this change in HbA1c between the two groups to identify the subgroup that maximized the treatment difference (PPT versus BBT).

**RESULTS**

The demographics and characteristics of the at-goal patient population are presented in Table 1. All of the baseline characteristics were comparable between the treatment groups. A greater number of patients were male (55.56% and 56.44%) and Caucasian (59.26% and 59.41%) in the PPT and BBT groups, respectively. The mean (SD) age was 56.19 (9.84) and 54.88 (9.45) years with a mean (SD) baseline HbA1c of 8.46 (0.73)% and 8.67 (1.03)% [69.0 (7.98) and 71.3 (11.26) mmol/mol] in the PPT and BBT groups, respectively. At the end of the study, the mean (SD) total insulin dose was significantly higher for the BBT group compared to the PPT group [1.37 (0.88) versus 1.16 (0.48) U/kg; \(p = 0.05\)]. The mean (SD) increase in weight from baseline to endpoint was similar between treatment groups. The 30-day hypoglycemic rate over the 24-week treatment period was similar in the PPT group compared to the BBT group in the subgroup of at-goal patients.

No significant difference in change in HbA1c from baseline to endpoint was reported between the treatment groups in the at-goal patient population. The mean (SD) decrease in HbA1c was numerically greater in the BBT group (Table 1; Fig. 1) in at-goal patients [PPT −2.17 (0.79)% versus BBT −2.34 (1.00)%; −23.7 (8.63) mmol/mol versus −25.6 (10.93) mmol/mol], whereas the decrease in HbA1c was numerically greater in the PPT group (Table 3; Fig. 1) in the subgroup of at-goal patients with race other
Table 1 Demographics and characteristics of the at-goal (24 weeks) population (who provided a baseline glycated hemoglobin) in the trial

| Variable                                      | Treatment group | p value |
|-----------------------------------------------|-----------------|---------|
|                                               | PPT (N = 81)    | BBT (N = 101) |
| Age (years), mean (SD)                        | 56.19 (9.84)    | 54.88 (9.45)    | 0.36* |
| Gender, n (%)                                 |                 | 1.00*     |
| Male                                          | 45 (55.56)      | 57 (56.44)     |       |
| Female                                        | 36 (44.44)      | 44 (43.56)     |       |
| Race, n (%)                                   |                 | 0.97*     |
| African                                       | 7 (8.64)        | 6 (5.94)      |       |
| Caucasian                                     | 48 (59.26)      | 60 (59.41)     |       |
| East/Southeast Asian                          | 4 (4.94)        | 5 (4.95)      |       |
| Hispanic                                      | 22 (27.16)      | 29 (28.71)     |       |
| Other                                         | 0 (0.0)         | 1 (0.99)      |       |
| Baseline BMI (kg/m²), mean (SD)               | 35.42 (5.29)    | 34.61 (5.62)   | 0.32* |
| Duration of diabetes (years), mean (SD)       | 10.37 (5.90)    | 11.35 (6.13)   | 0.28* |
| Baseline HbA1c (%), mean (SD)                 | 8.46 (0.73)     | 8.67 (1.03)    | 0.10* |
| Baseline HbA1c (mmol/mol), mean (SD)          | 69.0 (7.98)     | 71.3 (11.26)   | 0.10* |
| Baseline FBG (mmol/L), mean (SD)              | 9.41 (3.53)     | 9.71 (3.52)    | 0.57* |
| Baseline weight (kg), mean (SD)               | 103.80 (21.47)  | 99.69 (22.31)  | 0.21* |
| Morning 2-h PPG (mmol/L), mean (SD)           | 14.91 (4.23)    | 14.05 (3.68)   | 0.15* |
| Baseline OAMs, n (%)                          |                 | 0.07*     |
| MET                                           | 50 (61.73)      | 52 (51.49)     |       |
| TZD                                           | 2 (2.47)        | 12 (11.88)     |       |
| MET + TZD                                     | 19 (23.46)      | 20 (19.80)     |       |
| None                                          | 10 (12.35)      | 17 (16.83)     |       |
| Total insulin dose at endpoint (U/kg), mean (SD) | 1.16 (0.48)    | 1.37 (0.88)    | 0.05* |
| Weight at endpoint (kg), mean (SD)            | 108.3 (22.37)   | 104.3 (23.62)  | 0.24* |
| Change in weight from baseline (kg), mean (SD)| 4.53 (4.78)     | 4.59 (4.25)    | 0.93* |
| Change in HbA1c from baseline (%), mean (SD)  | -2.17 (0.79)    | -2.34 (1.01)   | 0.19* |
| Change in HbA1c from baseline (mmol/mol), mean (SD) | -23.7 (8.63)   | -25.6 (10.93)  | 0.19* |
| Change in FBG from baseline (mmol/L), mean (SD)| -1.58 (3.32)    | -2.04 (3.35)   | 0.37* |
than Caucasian and FBG <8.89 mmol/L [PPT −2.27 (0.88)% versus BBT: −2.05 (0.75)%; −24.8 (9.62) mmol/mol versus −22.4 (8.20) mmol/mol]. Although reduction in HbA1c was not statistically significant between PPT and BBT groups, the trend reversed in the identified subgroup.

In at-goal patients, the baseline and end-point mean (SD) FBG were 9.41 (3.53) and 7.71 (1.43) mmol/L in the PPT group; and 9.71 (3.52) and 7.48 (1.86) mmol/L in the BBT group. The mean (SD) change in FBG in the PPT group was −1.58 (3.32) mmol/L and in BBT group was −2.04 (3.35) mmol/L (Table 1; Fig. 2). In the subgroup of at-goal patients, the baseline and endpoint mean (SD) FBG were 6.81 (1.23) and 6.95 (1.52) mmol/L in the PPT group; and 6.59 (1.57) and 7.13 (1.88) mmol/L in the BBT group. The mean (SD) change in FBG was 0.15 (1.94) mmol/L in the PPT group and 0.54 (2.23) mmol/L in BBT group (Table 3; Fig. 2). This method of subgroup identification is robust and identified the subgroup of at-goal patients with race other than Caucasian and FBG <8.89 mmol/L in a sensitivity analysis by adding average blood glucose excursions as an additional candidate predictor variable (data not shown).

Table 1 continued

| Variable | Treatment group | p value |
|----------|----------------|---------|
|          | PPT (N = 81)  | BBT (N = 101) |
| 30-day hypoglycemic rate (number of events), mean (SD) | 5.31 (4.20) | 5.09 (4.11) | 0.72a |

* p ≤ 0.05
a p values are from two-sample t test with Satterthwaite approximation. p values are intended as descriptive measure only
b p values are from Fisher’s exact test. p values are intended as descriptive measure only
c n = 78
d n = 80
e n = 98
f n = 100
g n = 92

Table 2 Variable importance score based on subgroup significance

| Variable                  | Importance score based on subgroup significance |
|---------------------------|-------------------------------------------------|
| Baseline FBG              | 2.36                                            |
| Race/ethnicity            | 1.40                                            |
| Morning PPG               | 1.07                                            |
| Duration of diabetes      | 0                                               |
| Age                       | 1.23                                            |
| Gender                    | 0.28                                            |
| Baseline BMI              | 0.75                                            |
| Baseline TZD              | 0                                               |
| Baseline MET              | 0                                               |
| Permutation-based maximum | 1.25                                            |
| variable importance (mean)|                                  |
| Permutation-based maximum | 0.54                                            |
| variable importance (SD)  |                                  |

Importance scores greater than mean plus SD of the permutation-based maximum variable importance is highlighted in bold
BMI body mass index, FBG fasting blood glucose, MET metformin, PPG postprandial glucose, SD standard deviation, TZD thiazolidinedione

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Table 3 Demographics and characteristics of at-goal (24 weeks) population (who provided a baseline glycated hemoglobin) subgroup with race other than Caucasian and baseline fasting blood glucose (<8.89 mmol/L) in the trial

| Variable                        | Treatment group       | p value |
|---------------------------------|-----------------------|---------|
|                                 | PPT (N = 21)          | BBT (N = 21) |        |
| Age (years), mean (SD)          | 56.56 (6.67)          | 55.06 (9.25) | 0.55a  |
| Gender, n (%)                   |                       | 1.00b    |
| Male                            | 9 (42.86)             | 10 (47.62) |        |
| Female                          | 12 (57.14)            | 11 (52.38) |        |
| Race, n (%)                     |                       | 1.00b    |
| African                         | 4 (19.05)             | 3 (14.29)  |        |
| Caucasian                       | 0 (0)                 | 0 (0)     |        |
| East/Southeast Asian            | 3 (14.29)             | 4 (19.05)  |        |
| Hispanic                        | 14 (66.67)            | 14 (66.67) |        |
| Other                           | 0 (0)                 | 0 (0)     |        |
| Baseline BMI (kg/m²), mean (SD) | 32.26 (5.21)          | 29.68 (4.62) | 0.10a  |
| Duration of diabetes (years), mean (SD) | 10.90 (7.27)          | 14.29 (6.79) | 0.13a  |
| Baseline HbA1c (%), mean (SD)   | 8.45 (0.69)           | 8.50 (0.73) | 0.81a  |
| Baseline HbA1c (mmol/mol), mean (SD) | 68.9 (7.54)           | 69.4 (7.98) | 0.81a  |
| Baseline FBG (mmol/L), mean (SD) | 6.81 (1.23)           | 6.59 (1.57) | 0.62a  |
| Baseline weight (kg), mean (SD) | 87.18 (18.16)         | 81.38 (15.15) | 0.27a  |
| Morning 2-h PPG (mmol/L), mean (SD) | 14.27 (3.82)c          | 13.78 (2.94) | 0.65a  |
| Baseline OAMs, n (%)            |                       | 0.38b    |
| MET                             | 12 (57.14)            | 9 (42.86)  |        |
| TZD                             | 1 (4.76)              | 1 (4.76)   |        |
| MET + TZD                       | 5 (23.81)             | 3 (14.29)  |        |
| None                            | 3 (14.29)             | 8 (38.10)  |        |
| Total insulin dose at endpoint (U/kg), mean (SD) | 1.10 (0.39)          | 1.25 (0.45) | 0.26a  |
| Weight at endpoint (kg), mean (SD) | 90.69 (18.70)         | 85.12 (15.80) | 0.30a  |
| Change in weight from baseline (kg), mean (SD) | 3.51 (3.89)           | 3.75 (3.29) | 0.83a  |
| Change in HbA1c from baseline (%), mean (SD) | -2.27 (0.88)         | -2.05 (0.75) | 0.40a  |
| Change in HbA1c from baseline (mmol/mol), mean (SD) | -24.8 (9.62)         | -22.4 (8.20) | 0.40a  |
DISCUSSION

The goal of this analysis was to provide preliminary information suitable for insulin intensification therapy. This post hoc analysis used the SIDES algorithm to identify the subgroups that benefitted most from different insulin intensification therapies (PPT versus BBT).

As a result of the progressive diminution in insulin secretory capacity, patients will eventually require insulin intensification therapy by adding prandial insulin to basal insulin (BBT) or by switching to premixed insulins [16]. Basal-bolus therapy is often considered to be the gold standard in achieving target goals of glycemic control. However, evidence suggests that thrice-daily premixed insulin could be an alternative to BBT [17]. The pharmacodynamics, as measured by glucose infusion rates, is similar between these two regimens [18]. Also, a recent systemic review of 13 randomized clinical trials comparing the effects of basal-bolus and premixed insulin regimens in patients with T2DM who had not responded to previous antidiabetes medications showed that premixed

| Variable | Treatment group | p value |
|----------|-----------------|---------|
|          | PPT (N = 21) | BBT (N = 21) |         |
| Change in FBG from baseline (mmol/L), mean (SD) | 0.15 (1.94) | 0.54 (2.23) | 0.55 a |
| 30-day hypoglycemic rate (number of events), mean (SD) | 7.24 (5.26) | 5.80 (3.82) | 0.31 a |

BBT basal-bolus therapy, BMI body mass index, FBG fasting blood glucose, HbA1c glycated hemoglobin, MET metformin, N number of patients in each group, n number of affected patients, OAM oral antihyperglycemic medications, PPG postprandial glucose, PPT prandial premixed therapy, SD standard deviation, TZD thiazolidinedione

a p values are from two-sample t test with Satterthwaite approximation. p values are intended as descriptive measure only
b p values are from Fisher’s exact test. p values are intended as descriptive measure only
c n = 20

![Graph](graph.png)

**Fig. 1** HbA1c at baseline, endpoint, and change from baseline to endpoint (24 weeks of treatment) in both at-goal patients and subgroup of at-goal patients. BBT basal-bolus therapy, HbA1c glycated hemoglobin, n number of patients without missing data, PPT prandial premixed therapy
insulins are noninferior to basal-bolus regimens with respect to reduction of HbA1c from baseline, with similar rates of hypoglycemia, weight change, and daily insulin dose. Meta-analysis of change in HbA1c level between basal-bolus and premixed insulin regimens result in a small and nonsignificant difference of 0.09% (95% CI -0.03 to 0.21) [19]. Thus, to mix or to separate the insulins for the correct patient is still an important clinical question.

Our study revealed that patients with both race other than Caucasian and FBG \(<8.89 \text{ mmol/L}\) may have a different response to intensification therapies compared to the at-goal patient population. The numerically larger decrease in HbA1c in the PPT group compared to the BBT group in the subgroup of at-goal patients than total at-goal patients indicated that subgroup patients responded better than at-goal patients with the PPT regimen.

Though evidence showed the role of genetic susceptibility, thrifty phenotype, diet, and lifestyle in development of diabetes with racial disparities, evidence evaluating the effect of race on the efficacy of certain insulin regimens is limited [2, 20–23]. From the pathophysiology perspective, impaired beta cell function and insulin resistance are important in the development of diabetes [24]. Some studies showed that Hispanic and African population may be more susceptible to T2DM and more insulin resistant, although they have greater insulin secretion ability [25–27], while development of diabetes in the East Asian population is characterized by early beta cell dysfunction and insulin resistance [20, 28]. In the identified subgroup of patients, most of the patients belong to the Hispanic race; African and East Asian patients were of small proportion. Most of the clinical trials evaluating the efficacy of insulins were conducted in Caucasians and the racial diverse data are limited. A post hoc pooled analysis of 11 multinational clinical trials explored the impact of race/ethnicity on the efficacy and safety of commonly used insulin regimens. The analyses showed that patients of Latino/Hispanic, Asian, and African descent had different responses to insulin therapy: Latino/Hispanic patients had greater glycemic improvement than Caucasians with once-daily basal insulin treatment, while Caucasians had better glycemic control than Latino/Hispanic and Asian patients with twice-daily premix insulin therapy. For premixed insulin thrice-daily therapy, the HbA1c control was comparable across patients groups [29], but there was no information related to BBT. Also, a
recently conducted systematic review supported the use of premixed insulins as an initiation and intensification regimen in East Asian patients, indicating that the efficacy profile of premixed insulins is similar to that of other insulin treatments [30].

The FBG plays a very important role in overall glycemic control [31, 32]. On the trial level, more than half of the patients switched from evening insulin lispro mix 50/50 to insulin lispro mix 75/25 [10]. A recently published study compared the treatment effects of insulin lispro mix thrice daily (insulin lispro mix 50 before breakfast and lunch plus insulin lispro mix 25 before dinner) with BBT in East Asian patients with T2DM who were inadequately controlled with twice-daily premixed insulin. The results showed that premixed insulin lispro mix thrice daily was noninferior to BBT in the East Asian population, and the frequency of adverse events and the incidences of overall and nocturnal hypoglycemia were comparable between groups [17]. If FBG could not be controlled by the mid mixture, the evening injection should be switched to low mixture. The identified subgroup (race other than Caucasian and lower baseline FBG level) responded better to PPT treatment; the reason might be that the insulin lispro protamine suspension in premix was enough to control the FBG levels, and the patients could benefit from this simplified insulin intensification therapy with fewer injections.

Insulin intensification is not a one-way process. According to the American Diabetes Association/European Association for the Study of Diabetes (ADA/EASD) statement, it is important to individualize treatment targets and treatment strategies with an emphasis on a patient-centered approach while making treatment decisions [16]. The important aspects to consider while choosing or titrating insulin intensification regimens, according to the ADA/EASD statement, are the number of injections needed and the complexity and flexibility of the regimens [16]. For patients who wanted to limit the number of injections per day, premixed insulins remained the preferable option because of convenience of administration [33, 34]. Nonadherence to insulin therapy is the major reason for worsening of glycemic control [35]. Nonadherence to insulin therapy also depends on perceived burden of insulin therapy [36], which may lead to increased risk of long-term complications [37–39]. Fewer injections might promote patients’ adherence to the treatment. Because premixed insulins require few injections, PPT may potentially improve treatment adherence. Although premixed insulin analogues are more convenient, they are less adaptable because of their inability to titrate the shorter- from the longer-acting components of the formulation. However, these proved to be a viable treatment option owing to their less-complicated injection schedule and need of fewer injections [4].

There are some limitations for consideration in this analysis. First, the present results are a post hoc analysis of a study that compared two intensive insulin regimens in patients who failed on basal insulin plus OAM therapy [10]. Second, in the identified subgroup, though patients treated with the premixed, thrice-daily regimen had a numerically larger decrease in HbA1c than with the basal-bolus regimen, the results were not statistically significant. The nonsignificance may be due to the small sample size of only 42 patients in the identified group. However, our study sought to explain the heterogeneity and provided a preliminary hypothesis for tailored individualized therapy. The analysis was exploratory in nature and may need to be further confirmed with other prevailing evidence or future studies. Third, the SIDES algorithm extensively explores the structure of the observed data and identifies a differentiate group suggested by the data. The advantage of this novel statistical tool is that the result is driven by data, without any human intervention. However, with a different set of study data, a separate group may have been identified. If possible, this tool could be used in a large dataset to test and validate the hypothesis.

CONCLUSIONS

In the present post hoc analysis, a preliminary hypothesis of patient segmentation that will get the maximum benefit from insulin
intensification regimens was identified. Patients with both race other than Caucasian and lower levels of FBG were identified as the subgroup that might benefit from a premixed thrice-daily regimen. Though this subgroup analysis provides some information for tailored patient-centered therapy, it needs further exploration in future prospective, randomized trials.

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Disclosures. Li Xin Shi has nothing to disclose. Peng Fei Li is an employee of Eli Lilly and Company. Jia Ning Hou is an employee of Eli Lilly and Company.

Compliance with Ethics Guidelines. All procedures followed were in accordance with the ethical standards of the responsible committee on human experimentation (institutional and national) and with the Helsinki Declaration of 1964, as revised in 2013. Informed consent was obtained from all patients for being included in the study.

Data Availability. The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

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