


## ORIGINAL ARTICLE

# Efficacy of DPP-4 inhibitors, GLP-1 analogues, and SGLT2 inhibitors as add-ons to metformin monotherapy in T2DM patients: a model-based meta-analysis

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**Received** 15 December 2017; **Revised** 19 September 2018; **Accepted** 27 October 2018

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**Keywords** metformin, model-based meta-analysis, population pharmacodynamics, type 2 diabetes mellitus

## AIMS

The aim of the present study was to quantitate the hypoglycaemic effects of dipeptidyl peptidase-4 inhibitors (DPP-4i), glucagon-like peptide-1 receptor agonists (GLP-1r) and sodium glucose cotransporter 2 inhibitors (SGLT2i) as add-on treatments to metformin monotherapy in patients with type 2 diabetes mellitus (T2DM) using a model-based meta-analysis (MBMA).

## METHODS

A systematic literature search of public databases was conducted to develop models that describe the time courses of the fasting plasma glucose (FPG)- and haemoglobin A1c (HbA1c)-lowering effects of three antidiabetic classes using NONMEM 7.3.0.

## RESULTS

Seventy-six publications were eligible for this study, and 873 FPG and 1086 HbA1c values were collected. We developed a physiological indirect response model that described the time courses of FPG and HbA1c and simulated reductions in these values 90 days after the initiation of add-on treatments. FPG and HbA1c reductions with once weekly exenatide, liraglutide and dulaglutide were greater than those with other drugs. Mean changes from baseline FPG and HbA1c with these drugs were as follows: exenatide (−22.5 and −16.6%), liraglutide (−22.1 and −16.3%), and dulaglutide (−19.3 and −14.3%). The hypoglycaemic effects of DPP-4i and SGLT2i were similar.

## CONCLUSIONS

Once weekly exenatide, liraglutide and dulaglutide provided better hypoglycaemic effects among the antidiabetic drugs analysed. Long-acting GLP-1r appears to be more useful for T2DM patients inadequately controlled with metformin monotherapy.

## WHAT IS ALREADY KNOWN ABOUT THIS SUBJECT

- Clinical practice guidelines recommend metformin as first-line therapy; however, we sometimes encounter patients who do not respond well to this treatment. Although some classes of antidiabetic drugs have been identified as candidates for adjunctive treatments to metformin monotherapy, there is no consistent consensus regarding add-on second-line therapy.

## WHAT THIS STUDY ADDS

- The FPG- and HbA1c-lowering effects of three classes of antidiabetics, DPP-4i, GLP-1r and SGLT2i, as add-ons to metformin may be evaluated using an MBMA approach.
- Long-acting GLP-1r appears to be more useful than other drugs for T2DM patients inadequately controlled with metformin monotherapy.

## Introduction

Type 2 diabetes mellitus (T2DM) is characterized by a chronic hyperglycaemic state due to decreases in insulin secretion and sensitivity [1, 2]. The estimated prevalence of diabetes worldwide is more than 400 million, and the total number of patients with diabetes is predicted to increase to 629 million by 2045 [3]. Appropriate glycaemic control based on haemoglobin A1c (HbA1c) and fasting plasma glucose (FPG) is required in order to prevent various complications, such as retinopathy, nephropathy, and neuropathy [4, 5]. The American Diabetes Association (ADA) and the European Association for the Study of Diabetes (EASD) recommend metformin as first-line monotherapy for most T2DM patients [6]. Metformin is a biguanide that decreases blood glucose concentrations by inhibiting gluconeogenesis in the liver [7]. Secondary failure may occur after long-term metformin therapy [8–11]. Therefore, the ADA and EASD recommend dual therapy with metformin and other antidiabetic drugs if glycaemic control is not achieved. There is an extensive list of pharmacological therapies available for the second-line adjunctive treatment of T2DM, including sulfonylureas (SU), thiazolidines (TZD), **dipeptidyl peptidase-4** inhibitors (DPP-4i), **glucagon-like peptide-1 receptor** agonists (GLP-1r), **sodium glucose cotransporter 2** inhibitors (SGLT2i), and basal insulins. DPP-4i, GLP-1r and SGLT2i, which have novel mechanisms of action, are less likely to cause weight gain and hypoglycaemia [12–14]; therefore, the use of these drugs is increasing. In Japan, the share of DPP-4i in the total oral antidiabetic drugs market reached 69% in 2015 [15]. In terms of dual therapy with metformin monotherapy, since few randomized controlled trials (RCTs) have directly compared the efficacy of these drugs, a consistent consensus regarding the most appropriate drugs as add-ons is lacking.

A model-based meta-analysis (MBMA) is an extension of a traditional meta-analysis. A traditional meta-analysis has the following limitations: (1) it may only be applicable when direct head-to-head RCTs exist, and (2) observation periods and doses are limited to specific ranges. In contrast, MBMA, which involves a meta-analysis using mathematical models, has the capacity to perform indirect comparisons even though head-to-head RCTs are lacking. In addition, MBMA may incorporate longitudinal and dose–response data, thereby allowing for the quantification of dose–response relationships and time courses of effects. Therefore, MBMA is more flexible than a traditional meta-analysis and is expected to provide more information [16, 17].

The aim of the present study was to develop a population pharmacodynamic (PPD) model that quantitates the FPG- and HbA1c-lowering effects of DPP-4i, GLP-1r and SGLT2i as add-ons to metformin monotherapy in T2DM patients using an MBMA approach.

## Methods

### Literature search

The ‘targeted drugs’ in the present study included DPP-4i (sitagliptin, vildagliptin, alogliptin, linagliptin, teneligliptin, anagliptin, saxagliptin, trelagliptin and omarigliptin), GLP-1r (liraglutide, exenatide, lixisenatide and dulaglutide), and SGLT2i (ipragliflozin, dapagliflozin, luseogliflozin, tofogliflozin, canagliflozin and empagliflozin), which are all approved in Japan. A systematic literature search of PubMed, the Cochrane library (CENTRAL/CCTR: Cochrane Central Register of Controlled Trials), and ClinicalTrials.gov (<https://clinicaltrials.gov/>) was conducted on 3 March 2016. The words used in the search were (‘metformin’ OR ‘targeted drugs’) AND (‘diabetes’ OR ‘diabetic’). Details of the search terms are provided in Supplementary Table S1. Only clinical trials satisfying the following inclusion criteria were included in the analysis: (1) randomized double-blind clinical trials, (2) patients diagnosed with T2DM, (3) targeted drugs added to metformin monotherapy because of inadequate glycaemic control, (4) HbA1c or FPG values used as clinical indicators, and (5) published in English. We excluded trials focusing on specific populations, such as renal failure and paediatric subjects. MBMA was conducted according to PRISMA (Preferred Reporting Items for Systematic Reviews and Meta-Analyses) [18].

We extracted the following information from eligible studies: mean and median values of HbA1c and FPG at each time point, sample size, dosage, duration of T2DM, duration of metformin therapy, age, sex, race (e.g., Caucasians, Asians, and others), body mass index (BMI) and body weight. Graphical data were converted to numerical data using GetData Graph Digitizer<sup>®</sup> version 2.26 (<http://getdata-graph-digitizer.com>).

### Model development

A PPD analysis was performed using NONMEM 7.3.0 (Icon Development Solutions, Ellicott City, Maryland) with first-order conditional estimation with interaction method

(FOCE-INTER). Graphical processing of the NONMEM output was performed with R (version 3.3.2).

In the present study, a physiological indirect response model was established to describe the time courses of FPG and HbA1c [19]. Overall model structures for FPG and HbA1c are shown as follows:

$$K_{in,FPG} = (\text{Baseline}_{FPG} \times K_{out,FPG}) \times (1 + DP_{FPG} \times \text{Time})$$

FPG (DPP-4i and GLP-1r):

$$\frac{dFPG}{dt} = K_{in,FPG} \times (1 - E_{\text{placebo}}) \times (1 - E_{\text{drug}}) - K_{out,FPG} \times FPG$$

FPG (SGLT2i):

$$\frac{dFPG}{dt} = K_{in,FPG} \times (1 - E_{\text{placebo}}) - K_{out,FPG} \times (1 + E_{\text{drug}}) \times FPG$$

HbA1c:

$$\frac{dHbA1c}{dt} = K_{in,HbA1c} \times (FPG/\text{Baseline}_{FPG})^{\lambda} - K_{out,HbA1c} \times HbA1c$$

where  $K_{in,FPG}$  and  $K_{out,FPG}$  are the FPG production rate constant and FPG elimination rate constant, respectively. Changes in HbA1c were modelled as secondary changes dependent on the baseline ratio of FPG ( $FPG/\text{Baseline}_{FPG}$ ), with the HbA1c production rate constant ( $K_{in,HbA1c}$ ) and HbA1c elimination rate constant ( $K_{out,HbA1c}$ ). The description of HbA1c production also included the use of the power function  $\lambda$  [20–22].  $K_{in,HbA1c}$  is defined by  $K_{in,HbA1c} = \text{Baseline}_{HbA1c} \times K_{out,HbA1c}$ .  $\text{Baseline}_{FPG}$  and  $\text{Baseline}_{HbA1c}$  represent FPG and HbA1c levels before the initiation of dual therapies, respectively.  $\text{Baseline}_{FPG}$  and  $\text{Baseline}_{HbA1c}$  in the  $x^{\text{th}}$  biomarker of the  $j^{\text{th}}$  arm of the  $i^{\text{th}}$  study are given by:

$$\text{Baseline}_{x,i,j} = \text{TVB}_x \times \exp\left(\eta_{B,x,i} + \frac{\kappa_{x,i,j}}{\sqrt{N_{x,i,j}/100}}\right)$$

where  $\text{TVB}_x$  is the estimated typical baseline FPG and HbA1c.  $\eta_B$  and  $\kappa$  are the random effects of inter-study variabilities (ISV) and inter-arm variabilities (IAV) [23], respectively. ISV and IAV were assumed to be symmetrically distributed as random variables with mean zero and variance  $\omega_{ISV}^2$  and  $\omega_{IAV}^2$ . IAV was weighted by the inverse of the square root of the number of patients in the study arm ( $N_{ij}$ ) normalized to 100 patients. The reason why IAV was included in this analysis was that IAV is purely the product of a small sample size, because in a randomized trial with an infinite sample size, there are no random differences across arms. Disease progression for FPG was assumed to be a proportional increase with a slope parameter ( $DP_{FPG}$ ) relative to the baseline. An exponential error model was used to describe ISV on  $DP_{FPG}$ . The placebo effect to FPG levels ( $E_{\text{placebo}}$ ) was described by a constant model. An exponential error model was used to describe ISV on  $E_{\text{placebo}}$ . The drug effect ( $E_{\text{drug}}$ ) to FPG levels was as follows:

$$E_{\text{drug}} = \frac{E_{\text{max}} \times \text{Dose}}{ED_{50} + \text{Dose}}$$

where  $E_{\text{max}}$  is the maximum treatment effect ranging between 0 and 1;  $ED_{50}$  is the dose resulting in 50% of  $E_{\text{max}}$ . The model included the individual potency ( $ED_{50}$ ) of each

drug, but assumed the same  $E_{\text{max}}$  for drugs with the same mechanism of action. The drug effect was assumed to be constant across studies, i.e., ISV on  $E_{\text{drug}}$  was not estimated. These models indicate that DPP-4i and GLP-1r inhibit FPG production, and SGLT2i stimulates FPG elimination. An additive error model was used to describe residual error variability (RUV). RUV was weighted by the inverse of the square root of the number of patients in the study arm normalized to 100 patients. Ideally for mean data, residual variability needs to be weighted by the precision of the mean (the inverse of squared standard errors). However, since we did not obtain standard errors in many studies, residuals were weighted by the sample size.

After establishing the basic models, covariate modelling was conducted. Age, sex, race, BMI and body weight were selected as candidates for the covariate. Covariate selection was conducted based on clinical plausibility and differences in the objective function value (OFV) estimated by NONMEM between hierarchical models. Forward inclusion and backward exclusion were used to develop the covariate model. Significance levels for forward inclusion and backward exclusion were set at 0.01 and 0.001, respectively.

### Model validation

During model building, changes in OFV, Akaike information criterion (AIC), relative standard errors and goodness-of-fit (GOF) plots were used for model evaluation. GOF was investigated using plots of the observation vs. population prediction (PRED) and individual predictions (IPRED), conditional-weighted residuals (CWRES) vs. the treatment duration [24], CWRES vs. PRED, and absolute individual weighted residuals (IWRES) vs. IPRED. In order to assess the robustness of the final PD model, a prediction-corrected visual predictive check (pcVPC) was conducted. An 80% prediction interval (PI) was defined for pcVPC from the 10th and 90th percentiles of simulated dependent data at each time point and was then compared with original data. One thousand simulations were performed for pcVPC. pcVPC was performed with the software package Perl-speaks-NONMEM version 4.8.1.

### Simulation

Using the final models, we simulated reductions in FPG and HbA1c 90 days after the initiation of add-on therapy. The dosage was set to the recommended dose of each drug in Japan. Parameter uncertainty, obtained from the variance-covariance matrix of the final model, was implemented in the simulations. The typical time courses of FPG and HbA1c for the three drugs (vildagliptin, exenatide and canagliflozin), which were selected from each drug class, were simulated.

### Nomenclature of targets and ligands

Key protein targets and ligands in this article are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY [25], and are permanently archived in the Concise Guide to PHARMACOLOGY 2017/18 [26–28].

**Table 1**

Study characteristics for each drug

Drug	Number of trials	Dose (mg)	Number of points		Age <sup>a</sup> (year)	Male <sup>a</sup> (%)	Caucasian <sup>a</sup> (%)	BMI <sup>a</sup> (kg m <sup>-2</sup> )	HbA1c <sup>a</sup> (%)	FPG <sup>a</sup> (mg dl <sup>-1</sup> )	Weight <sup>a</sup> (kg)
			FPG	HbA1c							
<b>Placebo</b>	50	0	250	274	55.9	53.4	72.7	31.1	8.0	165	87.4
<b>DPP-4 inhibitors</b>											
<b>Sitagliptin</b>	24	50/100	132	134	55.1	54.1	57.6	30.9	8.0	160	86.7
<b>Vildagliptin</b>	10	50/100	52	74	55.8	52.8	76.5	30.3	8.1	173	81.8
<b>Alogliptin</b>	4	12.5/25	36	64	53.9	52.7	63.0	31.3	7.9	177	83.9
<b>Linagliptin</b>	6	1/5/10	26	48	58.7	54.0	75.1	30.2	8.0	166	85.0
<b>Teneligliptin</b>	1	20	5	5	55.7	55.1	0	NA	7.8	151	NA
<b>Anagliptin</b>	1	200	3	9	56.8	51.7	0	24.9	7.7	146	66.1
<b>Saxagliptin</b>	10	2.5/5/10	47	73	54.7	53.0	81.0	31.4	8.0	164	87.8
<b>Omarigliptin</b>	2	25	4	4	57.5	50.5	NA	NA	7.5	158	87.5
<b>GLP-1 receptor agonists</b>											
<b>Liraglutide</b>	2	0.5/0.6/1/1.2/ 1.5/1.8/2	11	15	53.5	54.0	0	25.9	8.6	177	68.4
<b>Exenatide</b>	3	0.01/0.02/2	18	30	52.7	53.7	78.5	33.0	8.2	167	94.6
<b>Lixisenatide</b>	4	0.005/0.01/ 0.02/0.03/ 0.04/0.06	38	60	55.4	44.7	81.8	32.3	7.6	162	89.0
<b>Dulaglutide</b>	1	0.75/1.5	24	22	54.0	46.2	52.6	31.3	8.2	175	86.4
<b>SGLT2 inhibitors</b>											
<b>Ipragliflozin</b>	2	12.5/50/150/300	27	27	56.6	50.0	91.3	31.8	7.8	157	89.3
<b>Dapagliflozin</b>	6	2.5/5/9.2/10	93	105	55.0	50.4	81.8	31.6	7.9	161	86.3
<b>Canagliflozin</b>	5	50/100/150/200/300	60	62	55.3	50.8	68.6	31.6	7.8	164	87.7
<b>Empagliflozin</b>	5	1/5/10/25/50	47	80	57.3	55.3	84.1	31.5	7.9	162	88.1

<sup>a</sup>Values are given as medians.

BMI, body mass index; DPP-4, dipeptidyl peptidase-4; FPG, fasting plasma glucose; GLP-1, glucagon-like peptide-1; HbA1c, haemoglobin A1c; NA, not applicable; SGLT2, sodium glucose cotransporter 2

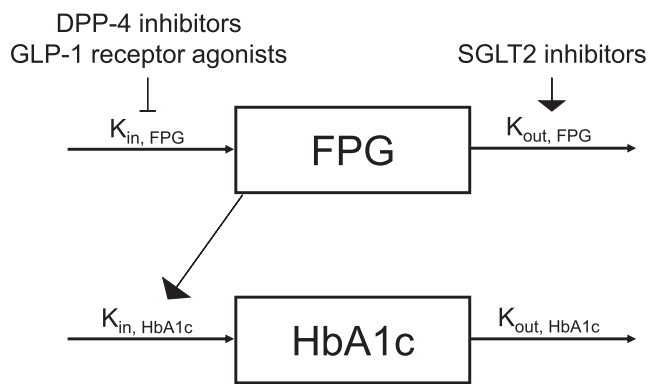
## Results

### Data analysis

A total of 2397 publications were found in the initial literature search. After screening and eligibility evaluations, 76 studies (31 585 patients) were eligible for the analysis, including 55 studies on DPP-4i (eight drugs), 10 on GLP-1r (four drugs), and 18 on SGLT2i (four drugs). However, some targeted drugs (e.g., trelagliptin, luseogliflozin and tofogliflozin) were not included in the analysis because trial data were not available. Detailed literature search results are shown in the PRISMA flow diagram (Figure S1), and a study design summary is provided in Supplementary Table S2. The total numbers of FPG and HbA1c were 873 and 1086, respectively. The medians (ranges) for the baseline values of FPG and HbA1c were 165 mg dl<sup>-1</sup> (138–244 mg dl<sup>-1</sup>) and 8.0% (7.0–9.3%), respectively. The characteristics of each targeted drug are summarized in Table 1.

### PPD models

Figure 1 shows a constructed indirect response model that describes changes in FPG and HbA1c levels over time for all treatments. Table 2 shows the population PPD parameter estimates of FPG and HbA1c. Estimated typical Baseline<sub>FPG</sub>, K<sub>out, FPG</sub>, E<sub>placebo</sub>, and DP<sub>FPG</sub> were 165 mg dl<sup>-1</sup>, 0.0936/day, 0.0168, and 0.0204/year, respectively. The dose–response relationship for each drug was characterized by the E<sub>max</sub> model with a different E<sub>max</sub> for each drug class and a different ED<sub>50</sub> for every drug within each class. ED<sub>50</sub> for teneligliptin was fixed to 0 in the final model because the estimate was close to 0, leading to convergence difficulties. The relationship between FPG and HbA1c was nonlinear and described by a power function with a different λ for the placebo and each drug class. Estimated typical Baseline<sub>HbA1c</sub> and K<sub>out, HbAc</sub> were 7.96% and



**Figure 1**

The final population pharmacodynamic model describing the time course of fasting plasma glucose (FPG) and haemoglobin A1c (HbA1c). DPP-4, dipeptidyl peptidase-4; GLP-1, glucagon-like peptide-1; K<sub>in, FPG</sub>, FPG production rate constant; K<sub>in, HbA1c</sub>, HbA1c production rate constant; K<sub>out, FPG</sub>, FPG elimination rate constant; K<sub>out, HbA1c</sub>, HbA1c elimination rate constant; SGLT2, sodium glucose cotransporter 2

**Table 2**

Population PPD parameter estimates

Parameter	Mean	RSE (%)
<b>Baseline<sub>FPG</sub> (mg dl<sup>-1</sup>)</b>	165	1.0
<b>K<sub>out, FPG</sub> (/day)</b>	0.0936	33.9
<b>Baseline<sub>HbA1c</sub> (%)</b>	7.96	0.6
<b>K<sub>out, HbA1c</sub> (/day)</b>	0.0393	14.3
<b>E<sub>placebo</sub></b>	0.0168	23.2
<b>λ<sub>placebo</sub></b>	0.893	15.0
<b>DP<sub>FPG</sub> (/year)</b>	0.0204	17.9
<b>DPP-4 inhibitors</b>		
<b>E<sub>max, DPP-4i</sub></b>	0.116	5.2
<b>ED<sub>50, sitagliptin</sub> (mg day<sup>-1</sup>)</b>	11.0	41.9
<b>ED<sub>50, vildagliptin</sub> (mg day<sup>-1</sup>)</b>	9.51	55.6
<b>ED<sub>50, alogliptin</sub> (mg day<sup>-1</sup>)</b>	2.36	52.1
<b>ED<sub>50, linagliptin</sub> (mg day<sup>-1</sup>)</b>	1.80	26.7
<b>ED<sub>50, teneligliptin</sub> (mg day<sup>-1</sup>)</b>	0 fixed	–
<b>ED<sub>50, anagliptin</sub> (mg day<sup>-1</sup>)</b>	31.4	33.8
<b>ED<sub>50, saxagliptin</sub> (mg day<sup>-1</sup>)</b>	1.47	48.2
<b>ED<sub>50, omarigliptin</sub> (mg day<sup>-1</sup>)</b>	3.15	66.7
<b>λ<sub>DPP-4i</sub></b>	0.831	3.7
<b>GLP-1 receptor agonists</b>		
<b>E<sub>max, GLP-1r</sub></b>	0.266	6.4
<b>ED<sub>50, liraglutide</sub> (mg day<sup>-1</sup>)</b>	0.377	34.7
<b>ED<sub>50, exenatide BID</sub> (mg day<sup>-1</sup>)</b>	0.0120	12.7
<b>ED<sub>50, exenatide QW</sub> (mg week<sup>-1</sup>)</b>	0.498	9.2
<b>ED<sub>50, lixisenatide</sub> (mg day<sup>-1</sup>)</b>	0.0392	21.8
<b>ED<sub>50, dulaglutide</sub> (mg week<sup>-1</sup>)</b>	0.328	5.4
<b>λ<sub>GLP-1r</sub></b>	0.777	8.7
<b>SGLT2 inhibitors</b>		
<b>E<sub>max, SGLT2i</sub></b>	0.199	13.6
<b>ED<sub>50, ipragliflozin</sub> (mg day<sup>-1</sup>)</b>	14.3	52.5
<b>ED<sub>50, dapagliflozin</sub> (mg day<sup>-1</sup>)</b>	4.06	35.5
<b>ED<sub>50, canagliflozin</sub> (mg day<sup>-1</sup>)</b>	30.0	28.1
<b>ED<sub>50, empagliflozin</sub> (mg day<sup>-1</sup>)</b>	4.54	62.3
<b>λ<sub>SGLT2i</sub></b>	0.654	5.2
<b>Inter-study variability and inter-arm variability</b>		
<b>ISV on Baseline<sub>FPG</sub> (%)</b>	8.4	13.2
<b>ISV on K<sub>out, FPG</sub> (%)</b>	105.4	19.0
<b>ISV on Baseline<sub>HbA1c</sub> (%)</b>	4.4	10.1
<b>ISV on K<sub>out, HbA1c</sub> (%)</b>	48.3	67.6
<b>ISV on E<sub>placebo</sub> (%)</b>	113.1	16.6
<b>ISV on DP<sub>FPG</sub> (%)</b>	92.1	22.4

(continues)

**Table 2**

(Continued)

Parameter	Mean	RSE (%)
<b>IAV on Baseline<sub>FPG</sub> (%)</b>	2.6	10.7
<b>IAV on Baseline<sub>HbA1c</sub> (%)</b>	1.2	15.8
<b>Residual error variability</b>		
<b>Additive error FPG (mg dl<sup>-1</sup>)</b>	3.35	4.4
<b>Additive error HbA1c (%)</b>	0.0721	9.3

DP<sub>FPG</sub>, the coefficient of disease progression; DPP-4i, dipeptidyl peptidase-4 inhibitors; ED50, dose resulting in 50% of E<sub>max</sub>; E<sub>max</sub>, maximum drug effect; E<sub>placebo</sub>, placebo effect; FPG, fasting plasma glucose; GLP-1r, glucagon-like peptide-1 receptor agonists; IAV, inter-arm variability; ISV, inter-study variability; K<sub>out, FPG</sub>, FPG elimination rate constant; K<sub>out, HbA1c</sub>, HbA1c elimination rate constant; RSE, relative standard error; SGLT2i, sodium glucose cotransporter 2 inhibitors

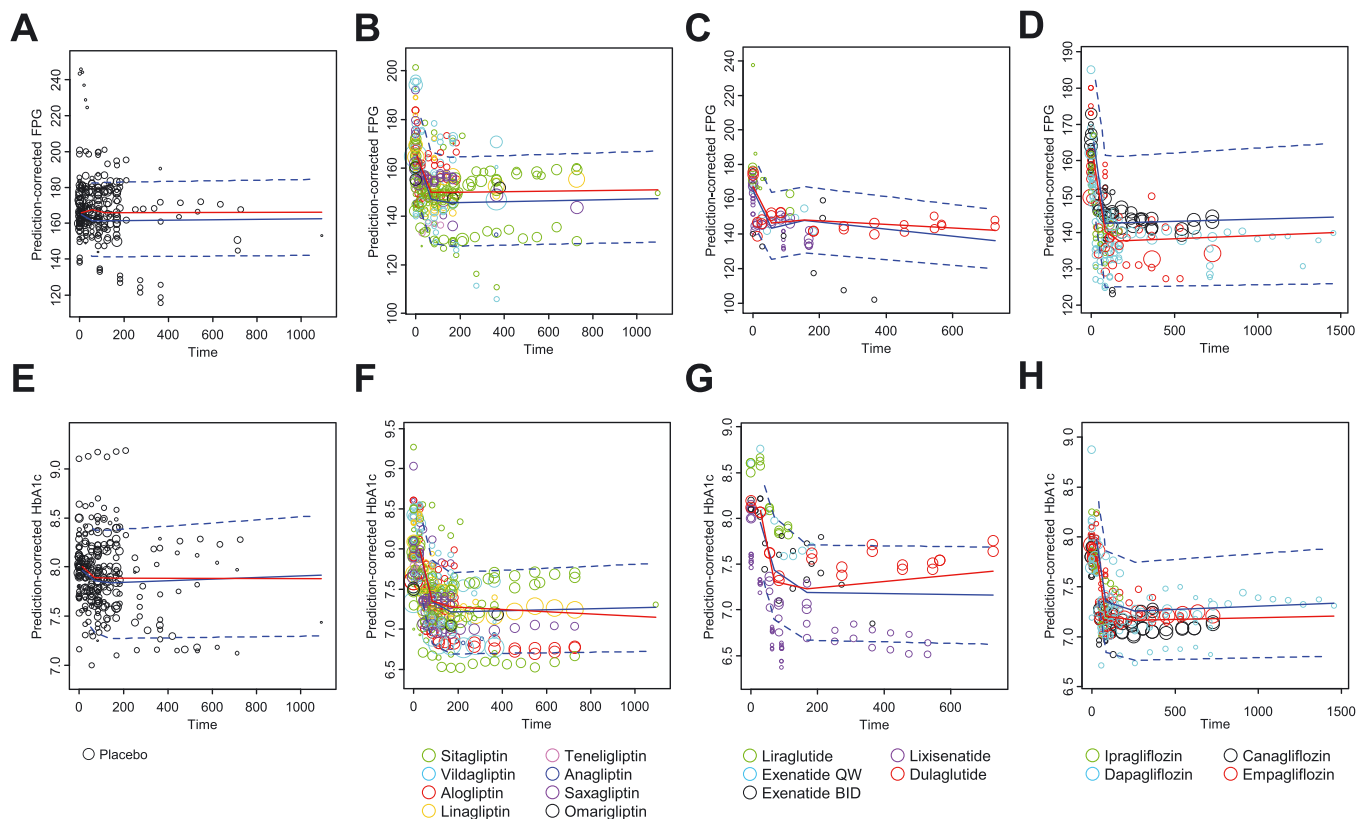
0.0393/day, respectively. None of the covariates were found to significantly improve the PPD model.

**Model validation**

GOF plots show the high predictive performance of the constructed models, and systematic deviations were not observed (Figure S2). pcVPCs for each drug class are shown in Figure 2. These models captured most of the observed data, indicating the good predictive performance of the models. These results suggest that the final models adequately describe the time courses for the FPG- and HbA1c-lowering effects of the targeted drugs.

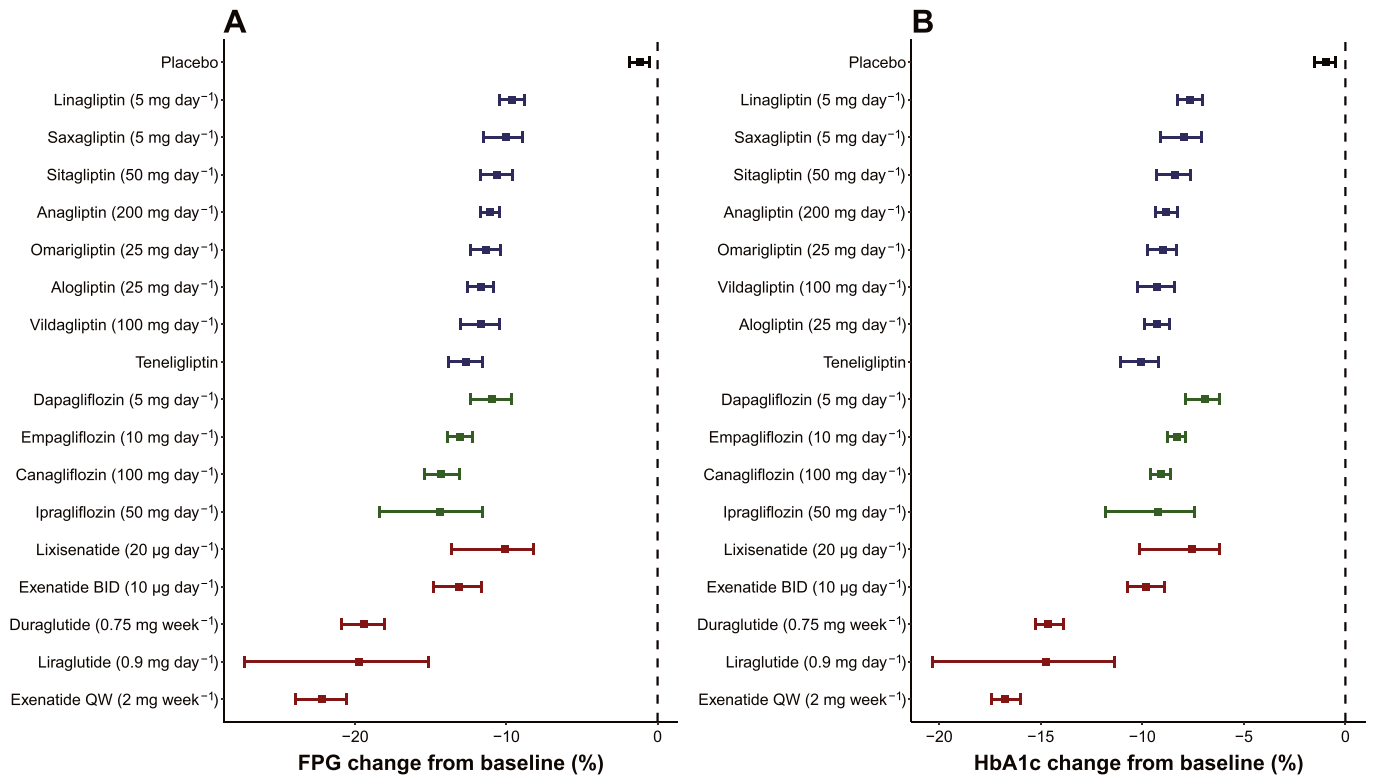
**Simulation**

Based on the final models, we simulated reductions in FPG and HbA1c 90 days after the initiation of add-on therapy (Figure 3). The dosage was set to the recommended dose of each drug in Japan. Parameter uncertainty was implemented in the simulations because some parameters (e.g., ED<sub>50</sub> for



**Figure 2**

Prediction-corrected visual predictive check plots for fasting plasma glucose (FPG) of the placebo (A), dipeptidyl peptidase-4 inhibitors (DPP-4i) (B), glucagon-like peptide-1 receptor agonists (GLP-1r) (C), and sodium glucose cotransporter 2 inhibitors (SGLT2i) (D) as well as haemoglobin A1c (HbA1c) of the placebo (E), DPP-4i (F), GLP-1r (G) and SGLT2i (H). Red solid lines represent the observed median. Blue solid and dashed lines represent the predicted median and 80% prediction intervals, respectively. Open circles represent observed data, and the symbol size is proportional to the number of subjects in each studyQW, once weekly; BID, twice daily



**Figure 3**

Reductions in FPG (A) and HbA1c (B) 90 days after the initiation of add-on therapy. Each square and bar represent the median and 90% confidence interval from model simulation ( $n = 1000$ ) for each drug. Red, green, and blue squares and bars represent changes from the baseline in FPG for GLP-1r, SGLT2 and DPP-4, respectively. The dosage was set to the recommended dose of each drug in Japan QW, once weekly; BID, twice daily

omarigliptin and empagliflozin, Table 2) were estimated with poor precision. Among these drugs, GLP-1r (exenatide QW, liraglutide and dulaglutide) showed superior FPG- and HbA1c-lowering effects ( $-22.2$ ,  $-19.7$  and  $-19.4\%$  for FPG, and  $-16.8$ ,  $-14.7$  and  $-14.7\%$  for HbA1c, respectively). FPG- and HbA1c-lowering effects were similar between DPP-4i and SGLT2i. Median reductions in FPG were  $-10.9$  to  $-14.4\%$  for SGLT2i and  $-9.6$  to  $-12.6\%$  for DPP-4i. Median reductions in HbA1c were  $-6.9$  to  $-9.2\%$  for SGLT2i and  $-7.6$  to  $-10.0\%$  for DPP-4i. The typical time courses of FPG and HbA1c of the three drugs, which were selected from each drug class, were simulated and are shown in Figure 4.

## Discussion

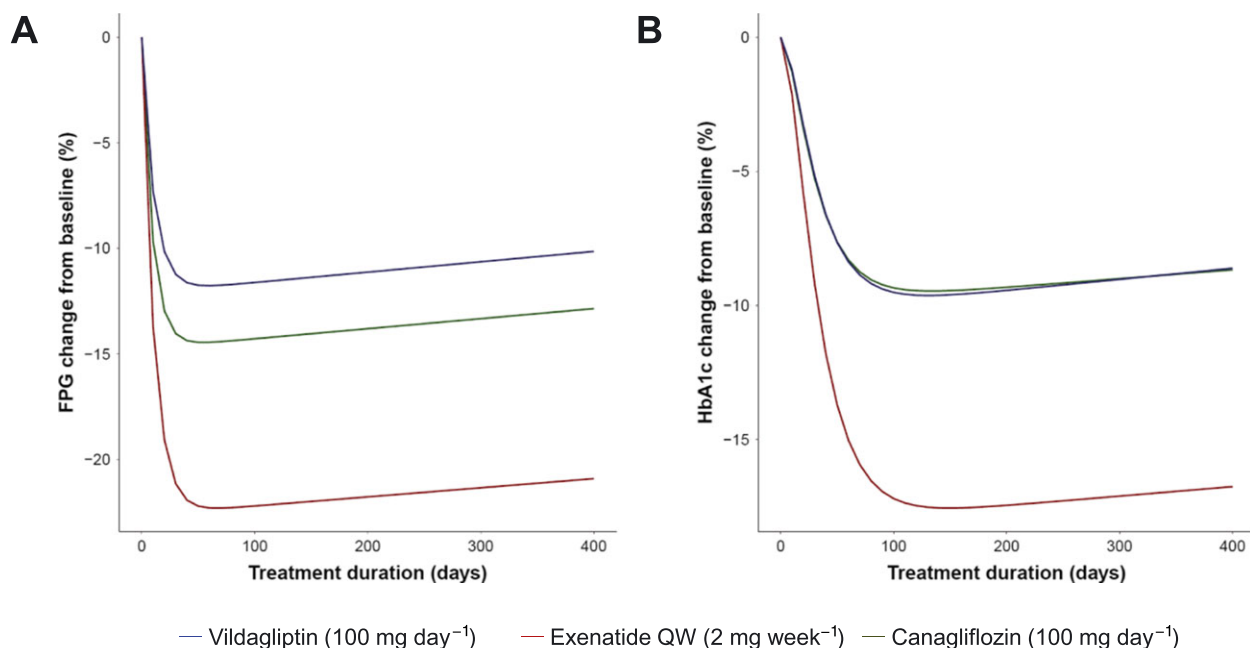
The aim of the present study was to quantitate the hypoglycaemic effects of DPP-4i, GLP-1r and SGLT2i as add-on treatments to metformin monotherapy in T2DM patients using MBMA.

We demonstrated that GLP-1r was associated with greater reductions in FPG and HbA1c than the other treatments tested within the approved dosages in Japan (Figure 3). The significant superiority of GLP-1r to DPP-4i as add-on therapy to metformin has been suggested in most RCTs. For example, exenatide ( $2 \text{ mg week}^{-1}$ ) resulted in significantly greater

improvements in HbA1c than sitagliptin ( $100 \text{ mg day}^{-1}$ ) in the DURATION-2 study [29]. Furthermore, the network meta-analysis (NMA) conducted by Zintzaras *et al.* indicated a higher proportion achieving the HbA1c goal with GLP-1r than other combination therapies with metformin [30]. The DURATION-8 study, which compared the efficacy and safety of exenatide ( $2 \text{ mg week}^{-1}$ ) vs. dapagliflozin ( $10 \text{ mg day}^{-1}$ ), showed that reductions in HbA1c at week 12 were greater in patients given exenatide [31].

In comparisons between long- and short-acting GLP-1r, treatments with long-acting GLP-1r (i.e., exenatide QW, dulaglutide and liraglutide) have been associated with greater reductions in FPG and HbA1c (Figure 3); long-acting GLP-1r provide relatively stable drug concentration–time profiles in the long term, leading to stable glycaemic control [32–34]. The NMA conducted by Kayaniyil *et al.* showed that the administration of exenatide QW led to a slightly higher proportion of patients achieving the glycaemic target than exenatide BID and lixisenatide [35].

The present study developed a PPD model that combined with the physiological relationship between FPG and HbA1c. During the model building process, we combined the mechanism of action of each drug class into the model: DPP-4i and GLP-1r inhibit FPG production, and SGLT2i stimulates FPG elimination. A large number of physiological models have been developed to describe the relationship between FPG and HbA1c. Our PPD parameters were similar to those



**Figure 4**

Typical time courses of fasting plasma glucose (FPG) (A) and haemoglobin A1c (HbA1c) (B)

reported previously [21, 22]. The FPG progression rate was estimated to be 2.04%/year and was similar to that reported by Stringer *et al.* (1.7%/year) [22]. In the present study, disease progression for FPG was assumed to be a proportional increase with  $DP_{FPG}$ . Several different disease progression models have been investigated (e.g., log-linear and exponential), but were not found to be superior.

The relationship between FPG and HbA1c was found to be nonlinear and was described by a power function with a different  $\lambda$  for the placebo and each drug class. Our estimated  $\lambda$  was similar to values reported in previous studies (0.74 and 0.71) [20, 21]. The nonlinear relationship between FPG and HbA1c may result from the contribution of postprandial glucose because the value for HbA1c is the result of FPG and postprandial glucose [36]. In addition, previous studies demonstrated that mean plasma glucose (the arithmetic mean of FPG and postprandial glucose) correlated better with HbA1c than FPG alone [37, 38].

This MBMA has several limitations. For example, covariate information obtained from the literature was limited. Some information, such as the metformin dose and durations of T2DM and metformin monotherapy, was not consistently reported; therefore, we were unable to include these as candidates for the covariate analysis. Since this information may contribute to patient heterogeneity, the results of the covariate analysis need to be interpreted with caution.

In conclusion, this MBMA quantified the hypoglycaemic effects of DPP-4i, GLP-1r and SGLT2i when they were added to metformin monotherapy. The simulations based on PPD models suggested that long-acting GLP-1r (i.e., exenatide QW, liraglutide and dulaglutide) were more effective than other drugs for T2DM patients inadequately controlled with metformin monotherapy.

## Competing Interests

There are no competing interests to declare.

## Contributors

H.I., Y.K., T.H., and I.I. wrote the manuscript. H.I., Y.T., Y.K., S.M., M.K., T.H., and I.I. designed the research. H.I., Y.T., and Y.K. analysed the data. The authors confirm that the PI for this paper is Ichiro Ieiri and that he had direct clinical responsibility for patients.

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## Supporting Information

Additional supporting information may be found online in the Supporting Information section at the end of the article.

<http://onlinelibrary.wiley.com/doi/10.1111/bcp.13807/supinfo>

**Table S1** Name of drugs used in the systematic literature search

**Table S2** Characteristics of studies included in the model-based meta-analysis dataset

**Figure S1** Results of the literature search

CENTRAL/CCTR, Central Register of Controlled Trials; DPP-4i, dipeptidyl peptidase-4 inhibitors; FPG, fasting plasma glucose; GLP-1r, glucagon-like peptide-1 receptor agonists; HbA1c, haemoglobin A1c; SGLT2i, sodium glucose cotransporter 2 inhibitors

**Figure S2** Goodness-of-fit plots of the final pharmacodynamic model for fasting plasma glucose (FPG) (A–E) and haemoglobin A1c (HbA1c) (F–J)

Population predictions were made using population mean parameters. Individual predictions were obtained using individual empirical Bayesian estimated parameters. Delta ( $\Delta$ ) is defined as the difference between the baseline values and observed or predicted values of FPG (A–D) and HbA1c (F–I). Black lines represent the line of identify (A, B, F and G) and  $y = 0$  (C, D, E, H, I and J). Red lines represent spline curves. DPP-4, dipeptidyl peptidase-4; GLP-1, glucagon-like peptide-1; SGLT2, sodium glucose cotransporter 2