Enhanced Anti-diabetic Effect of Berberine Combined With Timosaponin B2 in Goto-Kakizaki Rats, Associated With Increased Variety and Exposure of Effective Substances Through Intestinal Absorption

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Objective: Inspired by the traditionally clinical application of herb pair Zhimu-Huangbo to treat diabetes, a combination of plant ingredients, timosaponin B2 (TB-2) and berberine (BBR), was evaluated for their anti-diabetic efficacy and cooperative mechanisms.

Methods: The efficacy and pharmacokinetics of orally administered TB-2 (33.3 mg/kg/day), BBR (66.7 mg/kg/day), and TB-2+BBR (100 mg/kg/day) were evaluated in spontaneously non-obese diabetic Goto-Kakizaki (GK) rats, and metformin (200 mg/kg/day) was used as a positive control. The comparative exposure of the parent drugs, timosaponin A3 (TB-2 metabolite), and M1–M5 (BBR metabolites) was quantified in the portal vein plasma (before hepatic disposition), liver, and systemic plasma (after hepatic disposition) of normal rats on single and combination treatments. Cooperative mechanism of TB-2 and BBR on intestinal absorption and hepatic metabolism was investigated in Caco-2 cells and primary hepatocytes, respectively.

Results: After a 6-week experiment, non-fasting and fasting blood glucose levels and oral glucose tolerance test results showed that TB-2+BBR treatments (100 mg/kg/day) displayed significantly anti-diabetic efficacy in GK rats, comparable to that on metformin treatments. However, no significant improvement was observed on TB-2 or BBR treatments alone. Compared to single treatments, combination treatments led to the increased circulating levels of BBR by 107% in GK rats, and metformin (200 mg/kg/day) was used as a positive control. The comparative exposure of the parent drugs, timosaponin A3 (TB-2 metabolite), and M1–M5 (BBR metabolites) was quantified in the portal vein plasma (before hepatic disposition), liver, and systemic plasma (after hepatic disposition) of normal rats on single and combination treatments. Cooperative mechanism of TB-2 and BBR on intestinal absorption and hepatic metabolism was investigated in Caco-2 cells and primary hepatocytes, respectively.
interaction between TB-2 and BBR was mediated by intestinal absorption, rather than hepatic metabolism.

**Conclusion:** Combining TB-2 and BBR enhanced the anti-diabetic efficacy by increasing the in vivo variety of effective substances, including the parent compounds and active metabolites, and improving the levels of those substances through intestinal absorption. This study is a new attempt to assess the effects of combined plant ingredients on diabetes by scientifically utilizing clinical experience of an herb pair.

**Keywords:** berberine, timosaponin B2, diabetes, combination treatments, drug–drug interaction, metabolites, hepatic metabolism, intestinal absorption

**INTRODUCTION**

Type 2 diabetes (T2D) mellitus, a chronic, complex metabolic disorder with serious and lethal complications, is becoming increasingly common throughout the world. The prevalence of diabetes has been predicted to give rise to 592 million cases by 2035 (Guariguata et al., 2014). Although several oral synthetic anti-diabetic drugs, such as metformin and sulfonylureas, are available, patients taking these medications continue to suffer from poor long-term efficacy and high complication rates (Nathan, 2007; Nathan et al., 2009). Due to the advantages of fewer side effects and positive perceptions of patients, herbal medicines or their active natural ingredients are increasingly popular around the world, especially for chronic metabolic diseases. In ancient China, T2D was highly consistent with “Xiao ke,” a symptom characterized as “three excesses and one loss,” including excessive fluid intake, food consumption, and urination, along with weight loss (Wang, 2003). “Xiao ke” was first described in “Effective Prescriptions of the Past and Present” written during the Tang Dynasty (AD 600), nearly one century earlier than the description of diabetes by Western people in 1674 (Wang, 2003). Through their long history of clinical usage, herbal medicines, predominantly in the form of combinations of herbal medicines, have shown excellent therapeutic efficacy. **Zhimu-Huangbo,** composed of the dried rhizome of *Anemarrhena asphodeloides* Bunge (*Zhimu* in Chinese) and dried bark of *Cortex Phellodendri Chinensis* (*Huangbo* in Chinese), is one of the famous herb pairs for treating “Xiao Ke,” which was originally recorded in “Lan shi mi cang” written during the Jin Dynasty (AD 300). Currently, some relevant commercial products, such as *Zishen Wan* and *Zhibai Dihuang Pill,* are available in the market for treating diabetes or relevant symptoms. Correspondingly, our team firstly confirmed the anti-diabetic effect of *Zishen Wan* in spontaneously T2D mice (Tang et al., 2012). Zhao et al. (2012) demonstrated that *Zhibai Dihuang Pill* ameliorated diabetic symptoms in streptozotocin-induced diabetic rats. Other teams also confirmed the anti-diabetic efficacy of *Zhimu-Huangbo* in diabetic animals (Zhang et al., 2014; Song et al., 2015).

Compared with herbal medicines with complicated compositions and unclear scientific standards, evidence-based active natural ingredients are more likely to be understood and accepted. Due to the unsatisfactory therapeutic effects of monotherapy, combination therapy has become a preferable choice for diabetic patients (Prabhakar et al., 2014). Moreover, the low bioavailability of most natural ingredients calls for improved levels in target organs, and drug–drug interactions (DDIs) may be employed as an applicable strategy for solving this problem (Liu et al., 2016). The combination of several herbs can serve as a good template for developing combination treatments of natural ingredients for diabetes. Thus, our team has made many efforts over years to elucidate the medical material basis of **Zhimu-Huangbo** on diabetic treatments, accompanied by some published papers regarding this herb pair, single herbs, and active ingredients (Ma et al., 2008, 2009; Tian et al., 2014, 2016a,b; Fu et al., 2015; Jia et al., 2015; Sheng et al., 2015). Finally, after many pharmacodynamic comparisons, berberine (BBR) in *Huangbo* and timosaponin B2 (TB-2) in *Zhimu* were selected for combination therapy for diabetes.

Berberine, a well-known isoquinoline alkaloid and an over-the-counter drug for gastrointestinal infection, has received considerable renewed attention due to its anti-diabetic potential via multiple proposed pathways (Yao et al., 2015). Recently, various researchers have confirmed the blood glucose-lowering activities of BBR in patients and diabetic animals, along with its great efficacy against diabetes-induced complications (Zhang and Chen, 2012). However, the major drawback of BBR is the low bioavailability (usually <1% in various species) (Liu et al., 2016), especially considering that high oral doses of BBR (>900 mg/day) carry a risk of clinical gastrointestinal side effects (Yin et al., 2008). Moreover, the major BBR metabolites, M1–M5 (Figure 1A), were also verified to have hypoglycemic activity in cells or animals (Li et al., 2011; Wang K. et al., 2017; Yang et al., 2017). M1 even showed more hypoglycemic capacity than BBR in diabetic animals (Yang et al., 2017).

Timosaponin B2 (TB-2), the major bioactive steroid saponin with the highest content in **Zhimu,** has been a focus of our team for many years. Previously, we found that BBR and TB-2 were both the substrates of uptake transporter organic anion-transporting polypeptide (OATP) (Chen et al., 2015; Sheng et al., 2015). Moreover, we clarified the metabolism, hepatic disposition, and DDIs of TB-2 (Fu et al., 2015; Jia et al., 2015; Sheng et al., 2015; Tian et al., 2016b). As reported, TB-2 significantly decreased blood glucose levels and ameliorated diabetic nephropathy in vivo, and it improved insulin sensitivity in vitro (Yuan et al., 2015, 2016). Timosaponin A3 (TA-3, Figure 1B), a major metabolite of TB-2 previously confirmed by our team (Fu et al., 2015; Jia et al., 2015), was found to lower blood glucose levels in diabetic rats and mice, probably
Chemical structures and metabolic pathways of (A) BBR and (B) TB-2.
Biological Products Co., Ltd. (Beijing, China), respectively. Thalifendine (M2) (>98% pure) was synthesized in our laboratory. Berberrubine (M1), demethyleneberberine (M3), jatrorrhizine (M4), columbamine (M5), and tolbutamide (internal standard, IS) were purchased from Chengdu Must Bio-Technology Co., Ltd. (Chengdu, China, >98% pure). Acetonitrile and formic acid were purchased from Merck (Darmstadt, Germany). Triple-deionized water was prepared with a Milli-Q system (Millipore, United States). Dimethyl sulfoxide (DMSO), Dulbecco’s modified Eagle’s medium (DMEM), 2-(N-morpholino)ethanesulfonic acid (MES), fetal bovine serum, and Hanks’ balanced salt solution (HBSS) were purchased from Sigma–Aldrich Co., Ltd. (St. Louis, MO, United States). All other analytical-grade agents were obtained from the Sino Pharm Chemical Reagent Co., Ltd. (Shanghai, China).

Animals
Male GK rats, normal age-matched Wistar rats, and high-fat diet were provided by Shanghai SLAC Laboratory Animal Co., Ltd. (Shanghai, China). All the animals were maintained in air-conditioned animal quarters under controlled temperature (23 ± 1°C), noise (<50 db), and relative humidity (60 ± 10%), with a 12 h light/dark cycle (6:00-18:00). The rats were given ad libitum access to standard diet and water.

Instrumentation
Analyses were performed on a Shimadzu LCMS-8030 triple quadrupole system (Shimadzu Corp., Japan) equipped with an electrospray ionization source. Data processing was performed using Shimadzu LC–MS LabSolutions (version 5.42 SP4, Shimadzu, Japan). Chromatographic separation was carried out on a Waters CORTECS C18+ column (100 mm × 2.1 mm, 2.7 μm, Ireland) with a security guard C18+ column (5 mm × 2.1 mm, 2.7 μm, Ireland). The analysis of BBR and M1–M5 was optimized as follows: the mobile phase consisted of acetonitrile containing 0.2% formic acid (A) and water containing 0.2% formic acid and 2.5 mM of ammonium acetate (B), and the flow rate was set to 0.2 ml/min with a 10-μl injection volume. The gradient elution was 85% B from 0 to 2.5 min, 85% B–83% B from 2.5 to 3.0 min, 83% B from 3.0 to 6.0 min, 83% B–40% B from 6.0 to 6.5 min, 40% B from 6.5 to 7.0 min, 40% B–85% B from 7.0 to 7.2 min, and 85% B from 7.2 to 12 min. The optimized precursor-to-product ion pairs (collision energy) were in the positive mode as follows: m/z 579.20–255.15 (28 V) for TB-2 (at 4.1 min) and timosaponin A3 (at 6.1 min) and 271.50–154.70 (25 V) for tolbutamide (IS, at 5.8 min). The calibration standards were prepared by spiking the standard solutions in blank plasma to achieve the final concentrations of 1–500 ng/ml for TB-2 and TA-3, as well as 0.5–250 ng/ml for BBR and its metabolites (M1–M5). The calibration standards were prepared by spiking the standard solutions into the liver homogenates to yield final concentrations of 3–3000 ng/ml for TA-3, and 1–1000 ng/ml for TB-2, BBR, and M1–M5. If the concentrations of the compounds were over the linear range, the relevant samples were diluted with blank matrix by a factor of 10. All the test compounds showed good linearity with R² > 0.99, and more than 66.7% of the control samples had a target accuracy within ±20% standard deviation (SD).

Anti-diabetic Efficacy and PK in GK Rats
Four-week-old male GK rats were fed a high-fat diet, and age-matched non-diabetic Wistar rats were fed standard chow as the control group. Under these dietary conditions, blood glucose was estimated each week for 7 weeks in 12-h fasted rats using a portable glucometer (One Touch Sure Step Meter, Johnson & Johnson, United States) by drawing the blood from the tail vein. GK rats with fasting blood glucose levels (FBG) above 16.7 mM were considered as diabetic rats, and only uniformly diabetic rats were used in this study. At 11 weeks of age, the GK rats were randomly divided into five groups (n = 8): the vehicle group, TB-2 group (33.3 mg/kg/day), BBR group (66.7 mg/kg/day), TB-2+BBR (33.3+66.7 mg/kg/day) group, and metformin group (200 mg/kg/day) as a positive control. All the drugs were suspended in 0.5% carboxymethylcellulose sodium (CMC-Na) solution for oral gavage (10 ml/kg) once per day (at 9:00) for 6 weeks. Simultaneously, the Wistar and vehicle groups received vehicle over the treatment period.

Food consumption and body weight were measured every week throughout the experimental periods. Non-fasting blood glucose (NFBG) levels were determined by collecting blood samples from the tail vein (at 9:00), and FBG levels were determined after a 6-h fast (at 15:00) on days 0, 7, 14, 21, and 42. On day 28, an oral glucose tolerance test (OGTT) was performed after an overnight fast. After the rats received oral administration of glucose solution (2.0 g/kg), the blood glucose levels were measured at 0, 30, 60, 90, and 120 min after glucose administration. On day 35, after different treatments, approximately 250 μl of blood was collected from the tail vein of the GK rats in each group at 0.5, 1, 2, 4, 6, 12, and 24 h. The blood samples were centrifuged at 10,000 rpm for 5 min, and all the samples were stored at −20°C until analysis. On day 42, the rats were anesthetized with urethane (1.4 g/kg, dissolved in saline) after an overnight fast and were sacrificed to collect the liver, kidney, heart, and spleen. The organs were weighed after washing with saline and drying to calculate the organ coefficient, namely, the ratio between the weight of organs and the body in each rat.
**PK and Liver Distribution in Wistar Rats**  
Male Wistar rats (135; 250 ± 20 g) were acclimated for 7 days before use. The rats were given *ad libitum* access to standard diet and water. They were randomly divided into three groups (43 animals per group) and fasted overnight with access to water before treatment. An oral dose of TB-2 (33.3 mg/kg) was administrated to the rats in group A. BBR (66.7 mg/kg) was administrated to the rats in group B through oral gavage. A combination of TB-2 (33.3 mg/kg) and BBR (66.7 mg/kg) was orally administrated to the rats in group C. Forty-five rats per group (n = 5 for each time point) were anesthetized with urethane (1.4 g/kg) and sacrificed to collect hepatic portal vein blood (2 ml), abdominal aorta blood (6–8 ml), and liver (2 g) at 0.167, 0.5, 1, 2, 4, 6, 9, 12, and 24 h after oral administration. The blood samples were centrifuged at 10,000 rpm for 5 min for getting the plasma, and the liver samples were washed with saline three times. All the samples were stored at −20°C until analysis.

**Hepatic Metabolism in Primary Hepatocytes**  
The influence of hepatic metabolism on the DDIs of TB-2 and BBR was investigated using freshly isolated primary rat hepatocytes. Hepatocytes were isolated from male Wistar rats via two-step perfusion, as previously described (Shen et al., 2012; Sheng et al., 2015). The hepatocytes were incubated in six-well plates at a density of 0.5 × 10⁶/well, followed by the addition of TB-2 (1 µM) in the absence or presence of BBR (5, 50 µM), and the addition of BBR (1 µM) in the absence or presence of the CYP enzyme inhibitor 1-aminobenzotriazole (ABT) (1 mM) or TB-2 (0.1, 0.5, 5, and 30 µM). The cells were incubated on an orbital shaker at 37°C for 3 h. Cell suspensions (300 µl) were collected at 0 and 3 h and were immediately mixed with ice-cold methanol (300 µl) to terminate any potential reactions. The cell samples were then lysed by sonication, and each sample (100 µl) was extracted via acetonitrile precipitation, and prepared for analysis by HPLC–MS/MS.

**Transport in Caco-2 Cells**  
Caco-2 cells at passage 39 were seeded onto Millipore Transwell® 24-well culture plate inserts (Corning 0.4-µm pore polycarbonate membrane, Sterile Co., Bedford, MA, United States) at a density of 1.5 × 10⁶ cells/cm². The culture medium was changed every other day for the first 7 days and then daily for the next 14 days. On day 21 of culture, the integrity of the cell monolayers was evaluated via transepithelial electrical resistance (TEER) measurements using a Millicell-ERS epithelial voltohmeter (Millipore Co., Ltd.), with required TEER values of 300–1000 Ω/cm for the subsequent transport experiments. Prior to the transport experiments, the cell monolayers were washed three times with HBSS at 37°C. HBSS–MES transport medium (pH 6.0, containing 10 mM of MES) and HBSS–HEPES transport medium (pH 7.4, containing 10 mM of HEPES) with or without TB-2 (5 µM), BBR (10 µM), or TB-2 (5 µM) + BBR (10 µM) were added to the apical and basolateral sides. At the same time, the corresponding opposite side was filled with medium as the receiving chamber. After incubation for 30, 60, 90, and 120 min, 250-µl aliquots of the medium was taken from the receiving chamber at each time point, and prepared for analysis by HPLC–MS/MS.

**Sample Preparation**  
The liver samples were diluted with 3 volumes (0.5 g/1.5 ml) of saline and homogenized. The liver homogenates and plasma samples (100 µl) were precipitated with three volumes of acetonitrile (containing 10 µg/ml tolbutamide). The samples were then vortexed for 5 min and centrifuged at 13,000 rpm for 10 min, after which the supernatant (250 µl) was removed, and the samples were evaporated to dryness under a vacuum at 40°C. Then, 100 µl of the original mobile phase was added, and the samples were vortexed for 5 min and centrifuged at 13,000 rpm at 4°C for 10 min. Finally, the supernatant was used for HPLC–QQQ–MS/MS analysis.

**Data Analysis**  
A non-compartmental analysis was performed using WinNonlin software (Pharsight 6.2, Raleigh, NC, United States) to calculate the PK parameters. The significance of PK parameters in normal rats, generated by the sparse method in WinNonlin, was assessed with Welch’s unequal variances *t*-test. The significance of the other results was assessed using two-tailed Student’s *t*-tests. Statistical significance was defined as *P < 0.05.*

The liver extraction ratio (ERLiver) indicates the fraction of hepatic clearance from the liver portal vein plasma to the systemic plasma, and it was calculated as follows:

\[
ER_{\text{liver}} = \frac{\text{AUC}_{\text{por}} - \text{AUC}_{\text{sys}}}{\text{AUC}_{\text{por}}} \tag{1}
\]

where AUCpor and AUCsys are the areas under the concentration–time curve calculated in the portal vein plasma and systemic plasma, respectively.

The apparent permeability (*P*app) of drugs across the Caco-2 cell monolayers was calculated from the linear portion (i.e., 30, 60, 90, and 120 min) of a plot of the total amount of drug transported versus time, as follows:

\[
P_{\text{app}} = \frac{\Delta Q/\Delta t}{(A^*C_0)} \tag{2}
\]

where \(\Delta Q/\Delta t\) is the linear slope of the drug concentration in the receptor chamber over time, \(A\) represents the surface area of the membrane (1.12 cm²), and \(C_0\) is the initial concentration of drug in the donor chamber.

The efflux ratio (ERpermeability) for BL to AP and AP to BL transport was defined by the following equation:

\[
ER_{\text{permeability}} = \frac{P_{\text{app (BL-AP)}}}{P_{\text{app (AP-BL)}}} \tag{3}
\]

where \(P_{\text{app (BL-AP)}}\) and \(P_{\text{app (AP-BL)}}\) are the apparent permeability of drugs from the basolateral (BL) to apical (AP) and AP to BL sides of the monolayer, respectively.
FIGURE 2 | Hyperglycemic parameters (A) weight, (B) food intake, (C) organ coefficient, (D) FBG levels, and (E) NFBG levels on the vehicle, TB-2 (33.3 mg/kg/day), BBR (66.7 mg/kg/day), TB-2+BBR (100 mg/kg/day), and metformin (200 mg/kg/day) treatments for 6 weeks in GK rats; Wistar rats were used as the control group. (F) The levels and (G) AUC of the blood glucose during OGTT test in GK rats (n = 8). *P < 0.05, **P < 0.01, and ***P < 0.001 compared with the vehicle group.
RESULTS

Body Weight, Food Intake, and Organ Coefficients in GK Rats

As shown in Figures 2A,B, compared with the age-matched Wistar rats, the GK rats had lower body weights (P < 0.05 or P < 0.01) but similar food intake (P > 0.05) during the experimental period, consistent with published data (Jeong et al., 2009). Compared with the vehicle group, there were no significant differences in the body weight or food intake (P > 0.05) in any drug-treated groups, suggesting that orally administered TB-2 (33.3 mg/kg) and BBR (66.7 mg/kg) had no effect on the body weight or food consumption of GK rats receiving single or combined treatments.

As presented in Figure 2C, the liver, kidney, and spleen coefficients were significantly different (P < 0.001 or P < 0.01) between the vehicle and Wistar groups, indicating that diabetes significantly increased the organ coefficients of the GK rats. The positive control, metformin, had no effect on the decreased organ coefficients compared with the vehicle group. However, a significant decrease in the liver and kidney coefficients (P < 0.05) was observed on TB-2+BRR treatments, while this was limited on TB-2 or BBR treatments alone. These results indicated that the combination of TB-2 and BBR improved the physiological state of liver and kidney induced by diabetes in rats.

NFBG and FBG in GK Rats

As presented in Figures 2D,E, the measured FBG levels were much higher in the GK rats than those in the age-matched Wistar rats on days 0, 7, 14, 21, and 42, as were the NFBG levels, thereby validating the stability of diabetes for pharmaceutical testing in GK rats. Despite slightly reduced trends in the FBG and NFBG levels, there were no significant differences (P > 0.05) during the 6-week experiment on either TB-2 or BBR treatments alone compared with the vehicle group. In contrast, the combination of TB-2 and BBR not only caused a greater decrease in FBG and NFBG levels than any single compound, but also exhibited a significant difference (P < 0.05) from day 14 to the end of the experiment, compared with the vehicle group. The metformin treatments also significantly decreased FBG levels from the first to the sixth week (P < 0.05 or P < 0.01). Nevertheless, metformin did not significantly decrease NFBG levels (P > 0.05) throughout the whole experiment. At the end of the experiment, compared with the vehicle group, oral administration of TB-2 or BBR alone didn’t significantly decrease FBG or NFBG levels (P > 0.05), while co-administration of TB-2 and BBR decreased FBG and NFBG levels by 44.4% and 43.4% with statistical significance (P < 0.05), respectively, comparable to the effect on metformin treatments.

OGTT in GK Rats

After 4 weeks of treatment, an OGTT was performed with oral administration of glucose at 2 g/kg in GK rats, as shown in Figure 2F. Despite the similar levels at the zero time point, the blood glucose levels in GK rats were much higher than those in Wistar rats (P < 0.001) after glucose injection. Unlike the blood glucose levels in the vehicle and Wistar rat groups, which peaked at 30 min, the peak values in the drug-treated groups occurred at 60 min. Compared with the vehicle group, the co-administration of TB-2 and BBR significantly decreased the blood glucose levels by 22% (P < 0.05) and 25% (P < 0.05) at 30 and 120 min, respectively, while the significant decreases of 30% (P < 0.05) and 24% (P < 0.05) occurred only at 30 min on the metformin and BBR treatments, respectively. To further compare the anti-diabetic efficacy, the area under the time–blood glucose curve (AUC) was calculated. As exhibited in Figure 2G, compared with the vehicle group, the AUCs of blood glucose didn’t significantly decrease (P > 0.05) in the GK rats treated with TB-2 or BBR alone. In contrast, the combination of TB-2 and BBR led to a significant decrease of 12.0% in the AUC of blood glucose (P < 0.05), which was slightly better than the decrease (11.3%, P < 0.05) on metformin treatments.

PK in GK Rats

After 5 weeks of repeated treatments, a comparative PK evaluation of TB-2, BBR, and the relevant metabolites was conducted in GK rats treated with orally administered TB-2 (33.3 mg/kg), BBR (66.7 mg/kg), and TB-2+BBR (33.3+66.7 mg/kg), as shown in Figure 3 and Table 1. BBR and M1, a major metabolite, were observed in the systemic plasma after oral administration of BBR and TB-2+BBR in GK rats. Compared with BBR treatments alone, the combination treatments led to obvious increases in the Cmax and AUC of BBR by 695% (P < 0.05) and 107% (P < 0.05), respectively, compared with the vehicle group. The co-administration of TB-2 and BBR led to obvious increases in the Cmax and AUC of BBR by 695% (P < 0.05) and 107% (P < 0.05), respectively.
TABLE 1 | The PK parameters of BBR, TB-2, and M1 in the systemic plasma after oral administration of BBR (66.7 mg/kg), TB-2 (33.3 mg/kg), and their combination in GK rats on day 35 (mean ± SE, n = 8).

| PK parameters | BBR | BBR+TB-2 | M1 | BBR+TB-2 | TB-2 | BBR+TB-2 |
|----------------|-----|----------|----|----------|------|----------|
| T1/2 (h)       | 28.1 ± 11.7 | 9.3 ± 2.5 | 100.1 ± 92.2 | 23.9 ± 7.4 | 14.3 ± 5.3 | 8.6 ± 1.8 |
| Tmax (h)       | 4.6 ± 1.5 | 0.9 ± 0.1* | 10.4 ± 3.1 | 9.4 ± 1.2 | 2.1 ± 0.6 | 1.1 ± 0.3 |
| Cmax (ng/ml)   | 5.3 ± 0.4 | 42.2 ± 17.4* | 4.3 ± 1.7 | 6.6 ± 2.0 | 18.8 ± 5.4 | 47.4 ± 14.5 |
| AUC (h*ng/ml)  | 74.8 ± 3.5 | 154.8 ± 59.7* | 66.3 ± 18.5 | 94.9 ± 26.4 | 77.5 ± 13.6 | 100.4 ± 26.2 |

The significance of PK parameter was assessed with Welch’s unequal variances t-test. The significance of PK parameters was assessed using two-tailed Student’s t-tests. *P < 0.05, **P < 0.01, ***P < 0.001.

FIGURE 4 | Time–concentration curves of the parent compounds and metabolites after single oral administration of (A) BBR (66.7 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg), (B) TB-2 (33.3 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg) in the portal vein plasma, (C) BBR (66.7 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg), and (D) TB-2 (33.3 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg) in the systemic plasma of Wistar rats (n = 5).

along with shortened T_{max} and T_{1/2}, while it didn’t significantly increase the AUC or C_{max} of M1 (P > 0.05). The levels of M1 accounted for 75% and 61% of the parent drug in BBR and TB-2+BBR treatments, respectively, indicating a non-negligible amount of BBR metabolite in the plasma after repeated administration. Only TB-2 was detected in systemic plasma after oral administration of TB-2 and TB-2+BBR in GK rats. Compared with TB-2 treatments alone, there were no significant improvement on the C_{max} or AUC of TB-2 (P > 0.05) on TB-2+BBR treatments.

PK Before and After Hepatic Clearance in Wistar Rats
As shown in Figure 4 and Table 2, BBR and M1 were both observed in the portal vein plasma (before hepatic clearance) and systemic plasma (after hepatic clearance), and the presence of TB-2 caused increased levels of BBR and M1 following the oral administration of BBR (66.7 mg/kg) and TB-2+BBR (33.3+66.7 mg/kg) in Wistar rats. Compared with BBR treatments alone, co-administration of TB-2 increased the AUC of BBR by 114% (P < 0.01) in the portal vein plasma. In the systemic plasma, the presence of TB-2 only prolonged the T_{max} of BBR from 0.167 to 2 h, but it didn’t increase the AUC of BBR with statistical significance (P > 0.05). The AUCs of BBR were 6.9- and 9.1-fold higher in the portal vein plasma than those in the systemic plasma on BBR and TB-2+BBR treatments, corresponding to the ER_{liver} of BBR at 86.8 and 90.1%, respectively, indicating that BBR underwent very serious liver first-pass effect.

Concomitantly, in the portal vein plasma, the AUC and C_{max} of M1 were increased by 89% (P < 0.001) and 105% (P < 0.05), respectively. In the systemic plasma, the AUC and C_{max} of M1 were similarly enhanced by 95% (P < 0.05) and 54% (P < 0.05), respectively. Compared with that in the systemic plasma, the AUC of M1 was approximately fivefold higher in the portal vein plasma, corresponding to the ER_{liver} of M1 at 81.9 and 81.3% on BBR and TB-2+BBR treatments, respectively. The AUC ratios between M1 and BBR were 28.1 and 24.9% in the portal vein plasma, and 40.2 and 46.8% in the systemic plasma on BBR and TB-2+BBR treatments, respectively.

Timosaponin B2 and its metabolite TA-3 were observed in the portal vein plasma, whereas only TB-2 was detected in the systemic plasma following oral administration of TB-2
Results indicated that co-administration of TB-2 had the greatest impact on the hepatic collection of M1. On TB-2 treatments, the total exposure of BBR metabolites was 246% of that of BBR. This value was mainly attributed to the contribution of M2, which accounted for 182% of the exposure of BBR, along with minor contributions from M1 (29%), M3 (7%), M4 (16%), and M5 (11%). On TB-2+BBR treatments, the total exposure of BBR metabolites was 260% of that of the parent drug, and M2 made the largest contribution (144%), followed by M1 (77%), M3 (12%), M4 (15%), and M5 (11%). In addition, the presence of TB-2 had little effect on the AUC ratio between BBR and its metabolites in the liver, except for M1. The concentration of most metabolites peaked at 2 h, consistent with that of BBR, except those of M1 and M2.

Similar to the phenomenon in the portal vein plasma, TB-2 and TA-3 were found in the liver in the TB-2- and TB-2+BBR-treated groups. In the presence of BBR, the AUC of TB-2 were significantly increased by 230% (P < 0.05%) with prolonged $T_{\text{max}}$ and shortened $T_{1/2}$. Similar to those in the portal vein plasma, the PK parameters of TA-3 were minimally affected by co-administration with BBR. Moreover, compared with the values in the portal vein plasma, TA-3 showed considerably higher accumulation than TB-2 in the liver, with 259- and 57-fold higher levels than TB-2 in the absence and presence of BBR, respectively.

|TABLE 2| The PK parameters of BBR, TB-2, and their corresponding metabolites in the portal vein plasma and systemic plasma after oral administration of BBR (66.7 mg/kg), TB-2 (33.3 mg/kg), and their combination (100 mg/kg) in Wistar rats (n = 5). |

|PK parameters | BBR | BBR+TB-2 | M1 | M1+TB-2 | TB-2 | TB-2+TA-3 | TA-3 |
|---------------|-----|----------|----|---------|------|-----------|------|
| Portal vein plasma | | | | | | | |
| $T_{1/2}$ (h) | 9.8 | 10.5 | 9.7 | 7 | 8.5 | 22.8 | 0.8 | 0.5 |
| $T_{\text{max}}$ (h) | 2 | 2 | 6 | 9 | 1.0 | 2.0 | 2.0 | 2.0 |
| $C_{\text{max}}$ ± SE (ng/ml) | 63.1 ± 18.5 | 145.3 ± 50.3 | 3.5 ± 1.0 | 7.2 ± 1.5* | 37.1 ± 6.5 | 132.6 ± 16.4*** | 31.1 ± 4.8 | 47.7 ± 7.2 |
| AUC ± SE (h * ng/ml) | 198.0 ± 22.2 | 424.4 ± 60.5** | 55.7 ± 3.5 | 105.5 ± 6.2*** | 105.7 ± 7.6 | 254.8 ± 21.4*** | 73.9 ± 5.2 | 88.3 ± 9.6 |
| Systemic plasma | | | | | | | |
| $T_{1/2}$ (h) | 1.9 | 2.3 | 5.8 | 1.9 | 12.8 | 23.2 |
| $T_{\text{max}}$ (h) | 0.2 | 2 | 6.0 | 9.0 | 0.2 | 2.0 |
| $C_{\text{max}}$ ± SE (ng/ml) | 11.1 ± 3.4 | 10.8 ± 3.9 | 1.0 ± 0.2 | 2.0 ± 0.2** | 11.0 ± 2.9 | 34.2 ± 3.0*** |
| AUC ± SE (h * ng/g) | 25.1 ± 4.1 | 42.1 ± 11.7 | 10.1 ± 0.6 | 19.7 ± 2.9* | 20.3 ± 2.9 | 56.5 ± 4.6*** |
| ER_{liver} | 87.3% | 90.0% | 81.9% | 81.3% | 80.6% | 77.8% |

The calculation of PK parameters used “sparse” method in WinNonlin software (Pharsight 6.2, Raleigh, NC, United States). The significance of PK parameters was assessed with Welch’s unequal variances t-test. *P < 0.05, **P < 0.01, ***P < 0.001.

Liver Distribution in Wistar Rats
As shown in Figure 5 and Table 3, BBR and its five metabolites, M1–M5, all showed considerable levels in the liver in the BBR- and TB-2+BBR-treated groups. Co-administration of TB-2 improved the $C_{\text{max}}$ of BBR by 136% (P < 0.05), while the increased AUC of BBR didn’t elicit statistical significance. Compared to BBR treatments alone, the $C_{\text{max}}$ values of BBR metabolites were significantly increased by one- to twofold in the presence of TB-2: 109% for M1 (P < 0.05), 117% for M2 (P < 0.01), 174% for M3 (P < 0.01), 156% for M4 (P < 0.01). The corresponding AUCs of BBR metabolites were also significantly improved: 282% for M1 (P < 0.001), 144% for M3 (P < 0.001), 31% for M4 (P < 0.05), and 35% for M5 (P < 0.05). These results indicated that co-administration of TB-2 had the greatest impact on the hepatic collection of M1. On TB-2 treatments, the total exposure of BBR metabolites was 246% of that of BBR. This value was mainly attributed to the contribution of M2, which accounted for 182% of the exposure of BBR, along with minor contributions from M1 (29%), M3 (7%), M4 (16%), and M5 (11%). On TB-2+BBR treatments, the total exposure of BBR metabolites was 260% of that of the parent drug, and M2 made the largest contribution (144%), followed by M1 (77%), M3 (12%), M4 (15%), and M5 (11%). In addition, the presence of TB-2 had little effect on the AUC ratio between BBR and its metabolites in the liver, except for M1. The concentration of most metabolites peaked at 2 h, consistent with that of BBR, except those of M1 and M2.

Similar to the phenomenon in the portal vein plasma, TB-2 and TA-3 were found in the liver in the TB-2- and TB-2+BBR-treated groups. In the presence of BBR, the AUC of TB-2 were significantly increased by 230% (P < 0.05%) with prolonged $T_{\text{max}}$ and shortened $T_{1/2}$. Similar to those in the portal vein plasma, the PK parameters of TA-3 were minimally affected by co-administration with BBR. Moreover, compared with the values in the portal vein plasma, TA-3 showed considerably higher accumulation than TB-2 in the liver, with 259- and 57-fold higher levels than TB-2 in the absence and presence of BBR, respectively.

Hepatic Metabolism in Primary Hepatocytes
As shown in Figure 6A, the positive control midazolam (20 μM) was significantly metabolized after incubation for 3 h in the primary rat hepatocytes, and the presence of ABT, a broad-spectrum CYP enzyme inhibitor, completely abrogated midazolam metabolism. Those indicated the favorable metabolic...
FIGURE 5 | Comparison of time–concentration curves of the parent drugs and metabolites in the liver of Wistar rats after the oral administration of (A) BBR (66.7 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg) and (B) TB-2 (33.3 mg/kg) vs. TB-2+BBR (33.3+66.7 mg/kg) (n = 5).

TABLE 3 | The PK parameters of BBR, TB-2, and their corresponding metabolites in the liver after oral administration of BBR (66.7 mg/kg), TB-2 (33.3 mg/kg), and their combination in Wistar rats (n = 5).

| PK parameters | BBR | M1 | M2 | M3 |
|---------------|-----|----|----|----|
|               | BBR | BBR+TB-2 | BBR | BBR+TB-2 | BBR | BBR+TB-2 | BBR | BBR+TB-2 |
| $T_{1/2}$ (h) | 14.8 | 6.9  | 4.6 | 4.2  | 6.8 | 5.0  | 18.5 | 6.2  |
| $T_{\text{max}}$ (h) | 2   | 2   | 4.0 | 6.0  | 4.0 | 2   | 2.0  | 2.0  |
| $C_{\text{max}}$ ± SE (ng/g) | 794.7 ± 156.8 | 1876.0 ± 417.6* | 114.0 ± 10.6 | 238.0 ± 39.5* | 766.0 ± 54.3 | 1659.0 ± 254.9** | 38.8 ± 5.6 | 106.2 ± 18.2** |
| AUC ± SE (h * ng/g) | 3255.1 ± 247.5 | 4600.3 ± 645.1 | 932.5 ± 155.4 | 3566.3 ± 256.7*** | 5935.8 ± 164.9 | 6651.2 ± 469.0 | 234.5 ± 29.3 | 572.9 ± 34.7*** |

| PK parameters | M4 | M5 | TB-2 | TA-3 |
|---------------|----|----|------|------|
|               | BBR | BBR+TB-2 | BBR | BBR+TB-2 | BBR | BBR+TB-2 | BBR | BBR+TB-2 |
| $T_{1/2}$ (h) | 9.7 | 7.0  | 9.5 | 6.0  | 28.3 | 0.4  | 6.4  | 7.6  |
| $T_{\text{max}}$ (h) | 2.0 | 2.0  | 2.0 | 2.0  | 0.2 | 2.0  | 4.0  | 4.0  |
| $C_{\text{max}}$ ± SE (ng/g) | 78.2 ± 10.7 | 200.5 ± 33.7** | 89.4 ± 15.6 | 183.8 ± 38.4 | 73.1 ± 36.4 | 151.2 ± 35.7 | 2255.2 ± 531.4 | 2140.2 ± 380.7 |
| AUC ± SE (h * ng/g) | 527.5 ± 25.0 | 690.8 ± 59.6* | 362.8 ± 18.7 | 489.2 ± 59.2* | 89.4 ± 19.7 | 295.0 ± 59.2* | 23002.9 ± 3364.2 | 17099.4 ± 885.5 |

The calculation of PK parameters used “sparse” method in WinNonlin software (Pharsight 6.2, NC, United States). The significance of PK parameter was assessed with Welch’s unequal variances t-test. *P < 0.05, **P < 0.01, ***P < 0.001.
Digoxin was used as a positive control. The significance of \( P \) values was assessed using two-tailed Student’s t-tests. *\( P < 0.05 \), **\( P < 0.01 \), ***\( P < 0.001 \).

**Intestinal Absorption Transportation in Caco-2 Cells**

Transepithelial electrical resistance values were determined before (712 ± 111 \( \Omega/cm^2 \)) and after (762 ± 109 \( \Omega/cm^2 \)) transport experiments, with a deviation of <15% from the initial values, confirming that epithelial barrier function was maintained. As displayed in Table 4, an approximately 130-fold greater permeability coefficient of digoxin (5 \( \mu M \)) was observed in the secretory (BL→AP) direction (26.515 ± 1.91 \( \times 10^{-6} \) cm/s) than in the absorptive (AP→BL) direction (0.191 ± 0.032 \( \times 10^{-6} \) cm/s), thus confirming the well-behaved efflux function of Caco-2 cells.

There were no significant differences in the \( P_{\text{app}} \) values of TB-2 in the absorptive (0.057 ± 0.032 \( \times 10^{-6} \) cm/s) and secretory (0.061 ± 0.043 \( \times 10^{-6} \) cm/s) directions on TB-2 (5 \( \mu M \)) treatments, corresponding to an \( \text{ER}_{\text{permeability}} \) at 1.1. Co-incubation with BBR (10 \( \mu M \)) had no significant effect on the absorptive and secretory permeability of TB-2. In contrast, the \( P_{\text{app}} \) of BBR was markedly higher in the secretory side (2.714 ± 0.355 \( \times 10^{-6} \) cm/s) than that in the absorptive (0.038 ± 0.011 \( \times 10^{-6} \) cm/s) side on BBR treatments, and the \( \text{ER}_{\text{permeability}} \) of BBR was up to 71.4.

The presence of TB-2 resulted in a significant reduction in the \( P_{\text{app}} \) (BL→AP) from 2.714 ± 0.355 \( \times 10^{-6} \) cm/s to 1.495 ± 0.499 \( \times 10^{-6} \) cm/s, while this reduction was limited for \( P_{\text{app}} \) (AP→BL) (0.038 ± 0.011 \( \times 10^{-6} \) cm/s vs. 0.048 ± 0.014 \( \times 10^{-6} \) cm/s). Correspondingly, the \( \text{ER}_{\text{permeability}} \) of BBR was up to 71.4.

![Figure 6](image-url)
of BBR was decreased from 71.4 to 31.1 in the presence of TB-2 (5 μM). In brief, TB-2 could significantly inhibit the excretory permeability of BBR, whereas BBR barely affected the limited intestinal absorption of TB-2 in Caco-2 cells.

**DISCUSSION**

Based on the long history of clinical usage of Zhimu-Huangbo to treat “Xiaoke” symptom, TB-2 isolated from Zhimu and BBR derived from Huangbo were used as a combination therapy for diabetes in GK rats, a preferable model to mimic the hyperglycemic state of humans (Akash et al., 2013). Previously reported studies verifying the hypoglycemic efficacy of BBR or TB-2 were conducted in chemically induced diabetic rats (Liu et al., 2008); however, none were performed in GK rats. In the current study, no obvious side effects were observed throughout the whole experiment, indicating the short-term safety of these compounds in GK rats. We found there were no significant improvements on treating diabetes on either BBR (66.7 mg/kg) or TB-2 (33.3 mg/kg) treatments (Figure 2). In contrast, BBR (66.7 mg/kg) co-administered with TB-2 (33.3 mg/kg) achieved significantly anti-diabetic efficacy, as evidenced by the parameters of FBG, NFBG, and OGTT in GK rats, and those parameters were much better than those on single compound treatments (Figure 2).

Berberine is often compared with metformin, the first-line anti-T2D drug (Lee et al., 2006); both are typical AMPK activators to suppress hepatic gluconeogenesis. However, to match the similar anti-diabetic efficacy of metformin, the in vitro and in vivo studies verified the requirements for the same or higher dosage of BBR (Xu et al., 2014; Liu et al., 2015), likely due to its low bioavailability (<1%) (Liu et al., 2016). However, at oral dosages of BBR over 0.9 g/day, diabetic patients were prone to suffer gastrointestinal side effects (Yin et al., 2008). Therefore, reducing the oral dosage of BBR while sustaining efficacy is of great clinical significance. In our study, when combined with TB-2, only one-third of the oral dosage (66.7 mg/kg) was required for BBR to achieve comparable hypoglycemic efficacy to metformin (200 mg/kg), as evidenced by OGTT and FBG results. Some parameters (NFBG, liver, and kidney coefficients) were even better (Figure 2). These results indicated the superiority of combination treatments for treating diabetes.

On the one hand, an enhancement in pharmacology is associated with introducing more effective compounds to cause additive or synergistic effects on combination treatments (Prabhakar et al., 2014). BBR elicited anti-diabetic efficacy through multiple target pathways, such as activating AMPK, inhibiting protein-tyrosine phosphatase 1B (PTP1B), increasing InsR, and inhibiting liver gluconeogenesis (Zhang et al., 2008; Dong et al., 2012; Pirillo and Catapano, 2015). The gut microflora may also play a role in the anti-diabetic efficacy of BBR (Wang Y. et al., 2017). The metabolites of BBR, M1–M5, have all shown some potentials for diabetic management through different pathways, involved of InsR, AMPK and succinate dehydrogenase activation, and glucose-6-phosphatase and hexokinase inhibition (Li et al., 2011; Wang K. et al., 2017; Yang et al., 2017). TB-2 was able to improve insulin sensitivity through the insulin receptor substrate-1/phosphatidylinositol 3-kinase/Akt pathway (Yuan et al., 2016). The hypoglycemic efficacy of TA-3 was likely associated with hepatic gluconeogenesis or glycogenolysis inhibition (Nakashima et al., 1993). Notably, many of the above-mentioned anti-diabetic pathways happened in the liver, such as activating AMPK, increasing InsR expression, and inhibiting liver gluconeogenesis (Lee et al., 2006; Turner et al., 2008; Xie et al., 2011; Yao et al., 2015). Following oral administration of TB-2 or BBR on single or combination treatments, we found that the circulating levels of BBR and TB-2 were both low in rats (Figures 3, 4), in line with the reported poor bioavailability of TB-2 (1.2%) and BBR (<1%) (Cai et al., 2008; Liu et al., 2016). Compared to the poor circulating levels of BBR, the hepatic levels of BBR were two orders of magnitude higher. The discrepancy can be explained by the AUCs of BBR that were approximately 10-fold higher in the portal vein plasma, responsible for transporting compounds to liver, than those in the systemic plasma (Figure 4). Besides, five metabolites of BBR also showed substantial collection in the liver, where their total exposure was approximately 200-fold of circulating levels of BBR (Figure 5). Of note, the AUCs of M1, a metabolite that exhibited more hypoglycemic capacity than BBR (Yang et al., 2017), also showed very high hepatic collection (Figure 7B). Interestingly, the hepatic collection of TA-3, an active metabolite of TB-2, was 259- and 57-fold higher than those of TB-2 on TB-2 and TB-2+BBR treatments, respectively (Figure 7B). In sum, compared with the single compound treatments, the combination of TB-2 and BBR can introduce more compounds, involving the parent drugs and active metabolites in vivo, to cause additive or synergistic effects, leading to the enhanced efficacy in GK rats. On the other hand, DDIs can improve the exposure of drugs and active metabolites in vivo, especially for the target organs, which also bring about potentiation effects (Prabhakar et al., 2014). In normal rats, co-administration also improved the levels by 41–114% for BBR, 141–230% for TB-2, and 12–282% for BBR metabolites in the plasma and liver (Figure 7). Similarly, the improved circulating levels of BBR were verified after repeated oral administration in GK rats (Figure 7D). Thus, the improved exposure of parent drugs and active metabolites in the plasma and target organ also contributed to the enhanced anti-diabetic efficacy on TB-2+BBR treatments.

Given that pharmacokinetic and metabolic interactions usually occur through intestinal absorption or hepatic metabolism, those influencing factors were evaluated in vitro. Caco-2 cells, derived from human colonic cancer cells, are appropriate for studying intestinal absorption in vitro due to their ability to mimic intestinal epithelial cell patterns (Proctor et al., 2010). The ER_permability of BBR was as high as 71.4, in agreement with that BBR is a typical substrate of P-gp (Zhang et al., 2011). However, there was no significant efflux of TB-2, with an ER_permability of only 1. This seemed to contradict our previous finding that TB-2 was a substrate of both multidrug resistance-associated protein (BCRP) and multidrug resistance-associated protein 2 (MRP2) (Sheng et al., 2015). As reported, despite the detection of the mRNAs of many intestinal transporters, only P-gp expression in Caco-2 cells was stable.
FIGURE 7 | The compared AUCs of TB-2, BBR, and their metabolites in the (A) portal vein plasma, (B) liver, and (C) systemic plasma of normal rats, as well as (D) systemic plasma of GK rats, following oral administration of BBR (66.7 mg/kg), TB-2 (33.3 mg/kg), and TB-2+BBR (100 mg/kg).

and comparable (within twofold) to that in the human intestine (Troutman and Thakker, 2003). In contrast, the expression of MRP-2 and BCRP in Caco-2 cells is unstable and much lower than those in the human intestine, with one study even describing a 100-fold lower expression of BCRP in Caco-2 cells (Taipalensuu et al., 2001; Proctor et al., 2010). Therefore, the low ER permeability of TB-2 was possibly due to the poor BCRP and MRP2 expression in the Caco-2 cells, and TB-2 was very likely a P-gp inhibitor, rather than a P-gp substrate. This can explain why TB-2 significantly decreased the ER permeability of BBR from 71.4 to 31.1, and co-incubation with BBR did not improve TB-2 excretion in Caco-2 cells (Table 4). If the compounds functioned as P-gp inhibitors by interfering with ATP hydrolysis or altering the integrity of cell membrane lipids, an inconsistency between compounds as inhibitors and substrates might appear (Silva et al., 2015). Likewise, BBR was also reported to be a BCRP inhibitor, rather than a substrate (Zhang et al., 2011; Tan et al., 2013), and BBR was neither an inhibitor nor a substrate of MRP2 (Gu et al., 2015; Qian et al., 2016). Overall, the mutual inhibition of TB-2 and BBR, as P-gp and BCRP inhibitors, respectively, hampered the intestinal absorption of each other, in agreement with the improved TB-2 and BBR levels in the portal vein plasma on combination treatments (Figures 4, 5). After incubation with rat primary hepatocytes, BBR underwent extensive CYP 450-mediated metabolism to generate M2–M5, but such metabolism was limited for TB-2. Furthermore, neither TB-2 nor BBR affected the other’s hepatic metabolism. These findings were consistent with the phenomenon that the presence of TB-2 resulted in the increased BBR and M2–M5 levels, but rarely affected the exposure ratio between BBR and M2–M5 in the liver of rats (Table 3). Hence, the improved accumulation of M2–M5 simply resulted from an increased quantity of metabolizable BBR in the liver, rather than DDIs mediated by the CYP 450 enzyme. Unlike M2–M5, the generation of M1 was attributed to microbiota- rather than liver-mediated BBR metabolism (Wang K. et al., 2017), in line with its absence in primary hepatocytes after BBR incubation (Figure 6), considerable exposure in the portal vein plasma (Figure 4A), and longer $T_{\text{max}}$ than the other BBR metabolites in the liver (Figure 5). Co-administration of TB-2 increased the $AUC_{M1}/AUC_{BBR}$ from 28.6% to 77.5% in the liver (Table 3), which was probably correlated with the microbiota-mediated interaction between BBR (or M1) and TB-2 (or TA-3); this awaits further study. Briefly, the DDIs between BBR and TB-2 were probably mediated by intestinal absorption, rather than hepatic metabolism.

This study has some limitations that warrant further examination. For instance, questions still need to be answered regarding whether the co-administration of TB-2 and BBR influences glycosylated hemoglobin levels, insulin, and lipid parameters. Whether gut microbiota plays a role in the enhanced anti-diabetic efficacy on combination treatments, and the long-term safety and diabetic efficacy in other species requires further verification.

The famous Chinese scientist Youyou Tu, a winner of the Nobel Prize for Medicine in 2015 for the discovery of artemisinin inspired by historical records in ancient China, declared that artemisinin was a gift from Traditional Chinese Medicine (TCM) to the world (Shen, 2015). By comparison, the combination of herb medicines, namely, the TCM compatibility, contained more complicated and valuable ancestral wisdom through a long history of clinical trials in China. However, it has not been scientifically explored and utilized, recently. This study represented a primary attempt to develop combined plant ingredients, inspired by historical clinical experience with the compatibility of herb pairs, to treat diabetes. We hope it can motivate more researchers to scientifically explore the TCM compatibility, which may be the next gift from TCM to the world.

CONCLUSION

In this study, co-administration of TB-2 (33.3 mg/kg/day) and BBR (66.7 mg/kg/day) displayed significantly enhanced...
anti-diabetic efficacy in GK rats, which surpassed that of either compound alone, and was comparable to that of metformin (200 mg/kg/day). On the one hand, this enhanced efficacy was associated with the sum of the effective substances with multiple targets in vivo, including the active parent compounds (TB-2, BBR) and the relevant metabolites (TA-3 and M1–M5), especially for the substantial exposure of active metabolites and BBR in the target organ, liver. On the other hand, except for TA-3, co-administration improved the levels of those effective substances in the plasma and liver via intestinal absorption, rather than hepatic metabolism, which also played an important role in the potentiation effects on diabetes. Briefly, this study represented a successful attempt to assess the effect of combined plant ingredients on diabetes, inspired by historical clinical experience with herb pairs.

**ETHICS STATEMENT**

The animal studies were approved by the Institutional Animal Care and Use Committee of the Shanghai Institute of Materia Medica, China Academic Science, which has been described in the manuscript.

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**AUTHOR CONTRIBUTIONS**

XT, FL, YL, ZL, GP, and CH conceived and designed the experiments. HL, PH, MC, ZL, LH, and YZ assisted with the experiments. ZS and YZ analyzed the data. XT wrote the paper. YZ, GP, and CH critically revised the manuscript. All the authors read and reviewed the final manuscript.

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