Dysfunctional phagocytosis capacity, granulocyte recruitment and inflammatory factor secretion of Kupffer cells in diabetes mellitus reversed by Lidocaine

Rubin Wang,1 Minjia Sheng,2 Feng Shi,1 Yanjie Zhao,4 Lin Zhao,5 Jiangping Wu,6 Guangjiang Wu,6 Qingkun Song2,4

1Department of Emergency, Beijing Shijitan Hospital, Capital Medical University, Beijing 100038, China; 2Department of Gynaecology and Obstetrics, China-Japan Union Hospital of Jilin University, Changchun, Jilin 130033, China; 3Department of Pathology, Beijing Shijitan Hospital, Capital Medical University, Beijing 100038, China; 4Department of Medical Oncology, Beijing Shijitan Hospital, Capital Medical University, Beijing 100038, China; 5Department of Medical Records and Statistics, Xuanwu Hospital, Capital Medical University, Beijing 100045, China; 6Department of Infection Control, Beijing Shijitan Hospital, Capital Medical University, Beijing 100038, China; 7Department of Gynaecology and Obstetrics, Beijing Key Laboratory of Cancer Therapeutic Vaccine, Beijing 100038, China; 8Department of Evidence-based Medicine, Oncology School of Capital Medical University, Beijing 100038, China; 9Department of Science and Technology, Beijing Shijitan Hospital, Capital Medical University, Beijing 100038, China.

Purpose: Kupffer cells (KCs) present dysfunctional immunity capacity among the diabetes mellitus patients. This study aims to investigate whether Lidocaine could reverse dysfunctions of KCs, in terms of phagocytosis, granulocyte recruitment and inflammatory mediator secretion.

Methods: db/db and C57BL/6 mice were employed to establish diabetic and nondiabetic models. Upon intravenous injection of Lidocaine, KCs were isolated and cultured ex vivo. The functions of phagocytosis, recruiting granulocytes and inflammatory mediator secretion in KCs were compared between Lidocaine-treated and untreated (control) groups.

Results: Comparing with nondiabetic mice, KCs in diabetic mice presented reduced phagocytosis, activated granulocyte recruitment, increased expression of intercellular cell adhesion molecule-1 (ICAM-1) and activated levels of inflammatory mediators. With Lidocaine injection, phagocytic functions of KCs in diabetic mice were improved significantly; in contrast, recruitment of granulocytes, expression of ICAM-1 and secretion of inflammatory mediators were reduced markedly. However, Lidocaine intervention did not alter KC functions in phagocytosis, granulocyte recruitment, ICAM-1 expression or inflammatory mediator secretion among nondiabetic mice.

Conclusion: Lidocaine reversed diabetes-related dysfunctions of KCs in terms of phagocytosis, granulocyte recruitment, ICAM-1 expression or inflammatory mediator secretion.

Keywords: macrophages, diabetes, phagocytosis, granulocyte recruitment, inflammatory mediator, Lidocaine

Introduction

Diabetes mellitus (DM) is a chronic metabolic disease with lifetime hyperglycemia. According to 2015 International Diabetes Federation, 415 million people had been diagnosed with DM and 5 million people died of DM worldwide. In China, the number of DM patients accounts for 25% of global cases. In 2010, the prevalence of DM increased to 11.6% of the population in China. The age-standardized incidence of type 2 DM was 9.6 and 9.2 per 1,000 person-years in men and women, reported from the China Multicenter Collaborative Study of Cardiovascular Epidemiology and the China Cardiovascular Health Study. DM is a serious burden to public health, as well as economic and social development in China.

Kupffer cells (KCs) are a group of mononuclear macrophages, engaging in removal of bacteria from the circulation and subsequent elimination of phagocytosed bacteria. KCs played an important role against bacterial infection from the gastrointestinal tract;
however, in DM, the KCs were dysfunctional\(^5\) while the inflammatory factors were significantly activated.\(^5,6\) KCs activation was related with the altered level of tumor necrosis factor $\alpha$ (TNF-$\alpha$), interferon (IFN), ILs and nitric oxide (NO).\(^6,9\) In our previous study, the number of KCs was comparable with non-DM patients; however, ICAM-1 level was significantly higher in DM patients. Lidocaine was reported to affect the inflammatory factors,\(^10\) but whether lidocaine can improve the KCs function of diabetic patients has not been reported.

This study aimed to investigate whether Lidocaine could recover functions of KCs in terms of phagocytosis, granulocyte recruitment and inflammatory factor secretion using diabetes murine models.

**Methods**

**Ethical approval**

All procedures performed in studies in were in accordance with the ethical standards of the ethical committee of Beijing Shijitan Hospital, Capital Medical University, and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. The ethical committee of Beijing Shijitan Hospital, Capital Medical University approved this study.

**Animal administration**

Ten-week-old male C57BLKS/J db/m and C57BLKS/J db/db mice (Cavens Experimental Animal Co. Ltd.) were divided into nondiabetic (n=5, C57BLKS/J db/m mice), nondiabetic + lidocaine (n=5, C57BLKS/J db/db mice injected with lidocaine), diabetes (n=5, C57BLKS/J db/db mice) and diabetes + lidocaine (n=5, C57BLKS/J db/db mice injected with lidocaine) groups. Lidocaine (1 mg/kg) was injected intravenously once every 5 minutes for four times. After 1 hour, all mice were anesthetized and their livers, femur and tibia were aseptically harvested.

**KCIs isolation and culture and detection**

The liver tissue was smashed to the size of 1 mm\(^3\). 0.1% type IV collagenase was added for digestion at 37°C for 40 minutes. The sample was blown gently with a pipette for about 100 times. The cell suspension was collected by a 100 $\mu$m cell strainer, and the filtrate was centrifuged and re-suspended in the medium of DMEM supplement with 10% FBS and 1% Penicillin-Streptomycin (PS).

**Number of KCs detection**

The number of KCs was detected by immunohistochemistry. With formalin fixation, paraffin-embedded samples were collected from all mice. A serial of 4 $\mu$m thick slides were used to determine the number of KCs of each specimen. CD68 monoclonal antibody (sc-20060; Santa Cruz Biotechnology Inc., Dallas, TX, USA) was purchased from Beijing Zhong Shan Golden Bridge Biotechnology Co. Ltd. Sections were baked at 60°C in a dehydration oven for 60 minutes, dewaxed and rehydrated with xylene and graded alcohol washes. Antigen retrieval and deparaffinization were processed with the EnVision FLEX Target Retrieval Solutions, cooled to room temperature in Tris Buffered Saline Tween wash buffer for 5 minutes and then incubated with the primary antibodies. CD68 detection was visualized with diaminobenzidine. Slides were counterstained with hematoxylin. CD68 was expressed in the cytoplasm of KCs, and the dark brown granules in cytoplasm were taken as CD68-positive reaction cells. The average number of KCs was counted in five randomly selected high-power fields (HPF) under a microscope ($\times$400) for each specimen.

**Electron microscopy to detect cellular ultrastructure**

The liver tissue was transferred to an Eppendorf (Hamburg, Germany) tube containing 2.5% glutaraldehyde solution, fixed for ~1 hour and processed a 5-minute wash for three times by 1 M phosphate buffer. With fixation in 1% osmic acid for 30 minutes, the sample was processed a 5-minute wash twice by 0.1 M phosphate buffer. 3%–4% agar was used to pre-embed the sample and after solidification, the agar was cut into small pieces of about 1 mm\(^3\). 50%, 70% and 90% ethanol solutions, as well as 90% acetone and 10% aceton were applied for dehydration. Propylene oxide transition, epoxy resin impregnation, embedding, polymerization, ultrathin microtome section, uranyl acetate and lead citrate double staining, and drying were performed orderly. The sample was observed under FE I Tecnai G 212 transmission electron microscope at magnification of $\times$5,000.

**Phagocytosis assay**

KCIs (1×10\(^5\)) and polystyrene beads were seeded on coverslips. After incubation for 60 minutes, the coverslips were rinsed three times with PBS, fixed in 4% paraformaldehyde for 30 minutes and stained with Giemsa. KCis phagocytosing polystyrene beads under a microscope were observed. On each coverslip, five fields were selected randomly and ten KCs randomly selected in each field of view. The mean number of polystyrene beads phagocytosed by KCs represented phagocytosis function.

**Granulocyte recruitment**

The recruitment capacity was measured by Transwell assay. The femur and tibia were flushed with culture medium in
a 1 mL sterile syringe to obtain bone marrow, and single
neutrophil cell suspension was prepared at the concentration
of $2 \times 10^6$–$1 \times 10^9$/mL. A suitable amount of separation liquid
was added to the centrifuge tube, and the cell suspension was
spread above the level of separation liquid, and centrifuged
for 20–30 minutes. After centrifugation, a clear stratification
was obtained. The granulocytes were in the middle layer of
separation liquid. Cells were cultured in medium (DMEM +
10% FBS + 1% PS), in humidified atmosphere of 5% CO$_2$
at 37°C.

Cell migration assays were operated with a modified
Boyden chamber (Costar-Corning, New York, NY, USA).
Five hundred microliter KC suspensions with a density of
$2 \times 10^5$ cells/mL were added in the lower chamber. When cells
growing to 80% confluence, the supernatant was discarded,
and 500 µL complete medium (DMEM + 10% FBS + 1% PS)
was added. Two hundred microliter neutrophil suspension
with cells at a $2 \times 10^5$/mL density were added to the upper
chamber and incubated at 37°C, 5% CO$_2$ and saturated
humidity for 2 hours. Liquid in the upper chamber was
discarded, and the cells were washed twice with PBS. The
chamber was inverted and air dried. Five hundred microliter
0.1% crystal violet dye solution was added in the lower
chamber to submerge the chamber in the dye solution. After
incubated for 30 minutes at 37°C, the crystal violet staining
solution was discarded. The cells were washed twice with
PBS and captured using photographed under a microscope
(Olympus IX51) equipped with an Olympus color 3 digital
camera (Olympus). Migration was assessed by counting
the number of stained cells from five random HPF at ×200
magnification.

ICAM-1 detection
ICAM-1 expression in KCs was detected by Western blot.
Cells were collected and lysed on ice with RIPA lysis buffer
containing protease inhibitors. The extracted proteins were
quantified with bicinchoninic acid quantification assay. Total
cellular proteins were subjected to SDS-PAGE gel and trans-
ferred to nitrocellulose membranes. The membranes were
blocked with 5% non-fat milk for 2 hours and then incubated
with respective primary antibody overnight at 4°C. Follow-
ing washing for three times with TBS-T for 10 minutes, the
membranes were incubated with appropriate horseradish
peroxidase-conjugated secondary antibody for 1.5 hour at
room temperature. The bands were captured with SuperSignal
West Pico substrate (Thermo Fisher Scientific, Rockford, IL,
USA). β-Actin expression was used as the internal reference,
and the expression of ICAM-1 was estimated by the absorb-
bance (OD) ratio to β-actin, semiquantitatively.

Detection of TNF-α, IL-6 and IFN-γ
Quantitation of TNF-α, IL-6 and IFN-γ was conducted with
ELISA kits. Supernatant was obtained by centrifugation
of cell culture medium at 1,000 g for 20 minutes. Equal
amounts of cell culture medium were added in ELISA kits
and reacted for 20–25 minutes at room temperature. The OD
of TNF-α, IL-6 and IFN-γ was measured with a microplate
reader at a wavelength of 450 nm to calculate the sample
concentration.

Detection of NO
Equal amounts of cell culture medium and Griess reagent
were mixed and reacted for 20–25 minutes at room tempera-
ture. The OD of NO was read at 546–550 nm wavelength to
calculate the concentration.

Statistical analysis
GraphPad Prism (GraphPad Software, Inc., La Jolla, CA,
USA) was used for data analyses. All data were expressed as
the median and IQR. Kruskal–Wallis test was used to estimate
the difference between groups. Pair-wise comparisons were
conducted by Wilcoxon tests with Bonferroni adjustments.
All tests were two-sided, and a $P<0.05$ was considered sta-
tistically significant.

Results
Number of KCs in DM
The median number of KCs was 6/HPF (IQR = 1), comparable
between diabetic and nondiabetic mice (Figure 1A). However,
KCs in diabetic mice had less mitochondria and endoplasmic
reticulum (ER) than nondiabetic mice (Figure 1B, C).

Phagocytosis capacity
The phagocytic capacity was reduced in KCs from diabetic
mice, in contrast to nondiabetic mice ($P<0.05$, Figure 2A,
B). The number of beads phagocytosed by KCs was 10
(IQR = 4) and 59 (IQR = 5) in diabetic and nondiabetic mice
($P<0.05$, Figure 2C). KCs from nondiabetic mice had a simi-
lar phagocytic ability, the number of beads phagocytosed by
KCs from nondiabetic and nondiabetic plus Lidocaine mice
being comparable (Figure 2C, D). Phagocytic ability was
improved significantly in diabetic mice receiving Lidocaine
administration (Figure 2E), with a median phagocytosis of
49 (IQR = 5) (Figure 2C). The phagocytic capacity of KCs
from diabetic mice treated with Lidocaine was still lower than nondiabetic mice (Figure 2C).

Granulocyte recruitment
Compared to nondiabetic mice, KCs recruited more granulocytes in diabetic mice (Figure 3A, B). The median of granulocytes recruited by KCs in nondiabetic and diabetic mice was 15/HPF (IQR =7) and 96/HPF (IQR =19), respectively (P<0.05, Figure 3C). Lidocaine administration did not change the number of granulocytes in nondiabetic mice (Figure 3C, D). However, diabetic mice received Lidocaine had a significant reduction of granulocytes with a median of 33/HPF (IQR =11) (Figure 3C, E). The median number of granulocytes recruited by KCs in diabetic mice treated with Lidocaine was still higher than nondiabetic mice (Figure 3C, E).

ICAM-1 production
The level of ICAM-1 was 2.41 (IQR =0.03) and 1.33 (IQR =0.02) in diabetic and nondiabetic mice, respectively (P<0.05, Figure 4A, B). The ICAM-1 level was reduced to 1.51 (IQR =0.03) upon Lidocaine intervention in diabetic mice (P<0.05, Figure 4B), which was not affected in nondiabetic mice (P>0.05, Figure 4A, B).

NO production
The level of NO was 1.32 mmol/L (IQR =0.22) in nondiabetic mice, significantly lower than that of diabetic mice (4.97 mmol/L, IQR =0.10) (Figure 5A). In response to Lidocaine administration, NO was reduced significantly to 3.37 mmol/L (IQR =0.54) in diabetic mice, which was still higher than nondiabetic mice (P<0.05, Figure 5A). The level of NO did not change significantly in nondiabetic mice with Lidocaine intervention (Figure 5A).
Lidocaine reversed dysfunctional Kupffer cells of diabetes mellitus

**TNF-α production**

TNF-α level was significantly higher (776.91 pg/mL, IQR =14.52) in diabetic mice compared with nondiabetic mice (461.61 pg/mL, IQR =3.35) \((P < 0.05, \text{Figure 5B})\). In response to Lidocaine intervention, TNF-α level was reduced to 631.44 pg/mL (IQR =7.70) in diabetic mice \((P < 0.05)\), still higher than nondiabetic mice (Figure 5B). Lidocaine intervention did not affect TNF-α level in nondiabetic mice (Figure 5B).

**IL-6 production**

The baseline level of IL-6 was 516.32 pg/mL (IQR =8.58) and 741.84 pg/mL (IQR =15.21) in nondiabetic and diabetic mice, respectively \((P < 0.05, \text{Figure 5C})\). In diabetic mice, in response to Lidocaine administration, IL-6 level was reduced to 687.66 pg/mL (IQR =64.54), still higher than nondiabetic mice (Figure 5C). Lidocaine intervention did not change IL-6 level in nondiabetic mice (Figure 5C).

**INF-γ production**

The level of INF-γ was 541.60 pg/mL (IQR =11.66) in nondiabetic mice, significantly lower than that of diabetic mice (845.33 pg/mL, IQR =25.01, Figure 5D). Upon Lidocaine treatment, INF-γ level was reduced to 704.46 pg/mL (IQR =6.86) in diabetic mice (Figure 5D). Lidocaine intervention did not influence INF-γ level significantly in nondiabetic mice (Figure 5D).

**Discussion**

Worldwide, 284 million people have been affected by DM and have an increasing prevalence.\(^1\) DM patients are vulnerable

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**Figure 3** Granulocyte recruitment by Kupffer cells in nondiabetic and diabetic mice. Notes: (A) Baseline granulocyte recruitment in nondiabetic mice; (B) baseline granulocyte recruitment in diabetic mice; (C) comparison of granulocyte recruitment by Kupffer cells among different groups (median ± IQR, n=5); (D) granulocyte recruitment in nondiabetic mice treated with Lidocaine; (E) granulocyte recruitment in diabetic mice treated with Lidocaine. Magnification ×200.

**Figure 4** ICAM-1 expression on the surface of Kupffer cells in nondiabetic and diabetic mice. Notes: (A) Detection of ICAM-1 using Western blot; (B) comparison of ICAM-1 expression levels among different groups (median ± IQR, n=5).
to infections due to dysfunctional immunity, neuropathy and poor circulation.\textsuperscript{12} Hyperglycemia contributes to dysfunctional immunity and bacterial growth.\textsuperscript{5} 35.3\% of DM patients have an increased risk for pyogenic liver abscess, compared with normal people.\textsuperscript{13} KCs are the first line of defense for pathogens entering the liver and involved in elimination of resisting bacteria and viruses invading into liver sinus.\textsuperscript{6,14,15} The number of KCs was similar between diabetic and non-diabetic mice. The amounts of mitochondria, lysosome and ER in KCs in the DM group were significantly lower than the control group. ER connects inflammatory and stress signals with metabolism of cells, engaged in the development of type 2 DM.\textsuperscript{16} Reduction in mitochondria and ER may imply diminished phagocytosis of KCs.\textsuperscript{17} The functions rather than the number of KCs are significantly affected in DM. The KCs of diabetic mice were dysfunctional, with markedly reduced phagocytic ability than nondiabetic mice. This was consistent with previous reports that the function of diabetic

\begin{figure}[h]
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\caption{Secretion of inflammatory mediators by Kupffer cells (median ± IQR, n=5).}
\textbf{Notes:} (A) Nitric oxide secretion; (B) tumor necrosis factor alpha secretion; (C) interleukin-6 secretion; (D) interferon-gamma secretion.
\end{figure}
macrophages was depressed.\textsuperscript{5,18,19} Hyperglycemia may lead to dysfunctional macrophages through anoxic stress.\textsuperscript{18–20} Therefore, Lidocaine may improve phagocytosis of KCs by inhibiting inflammatory response in diabetic mice. Lidocaine may restore phagocytosis of KCs in DM through suppressing the release of inflammatory factors.

KCs involve in direct clearance and recruit neutrophils to eliminate invading bacteria.\textsuperscript{21,22} Pathological exacerbation of endotoxin, TNF-\(\alpha\), TGF, IFN, ILs, oxygen-free radicals, NO and other inflammatory mediators contribute to the activation of KCs.\textsuperscript{6,9}

The inflammatory mediators upregulate ICAM-1 expression in KCs.\textsuperscript{23} Higher expression of ICAM-1 on the surface of KCs was observed only in DM. KCs were significantly activated among DM patients. DM patients always have chronic inflammation\textsuperscript{24,25} under an aberrant immunity microenvironment.\textsuperscript{26,27} Our study identified increased secretion of NO, IL-6, INF-\(\gamma\) and TNF-\(\alpha\), as well as enhanced expression of ICAM-1 in KCs of diabetic mice compared with nondiabetic mice. Therefore, KCs of diabetic mice were activated and recruited more granulocytes. We found that Lidocaine could inhibit the release of inflammatory mediators, downregulate expression of ICAM-1 on the surface of KCs and reduce granulocyte recruitment in diabetic mice.

Lidocaine is widely used as a local anesthetic and antiarrhythmic agent. Recent studies have indicated that Lidocaine exerts a protective effect on vasculature system,\textsuperscript{28} inhibits the expression of inflammatory factors\textsuperscript{30} and presents an anti-tumor effect.\textsuperscript{29} These previous findings were similar as our study. Amide-linked local anesthetics have well-established anti-inflammatory effects, attenuate inflammatory Src signaling and disrupt the PI3K–Akt–NO pathway, thus blocking Src-dependent neutrophil adhesion and endothelial hyperpermeability.\textsuperscript{28} Transcription factor NF-\(\kappa B\) is indicated to be involved in the development of DM and chronic inflammation.\textsuperscript{30,31} Lidocaine could regulate NF-\(\kappa B\) signaling pathway and inhibit cellular inflammatory response.\textsuperscript{10} Lidocaine could protect endothelial cell function by inhibiting inflammatory mediators.\textsuperscript{28}

Major limitations of this study include a relatively small sample size and lack of functional validation for hypothesized molecular mechanisms. ICAM-1 expression was only detected at the translation level but not the transcription.

In DM, KCs actively recruit granulates, markedly release inflammation mediators and dysfunctionally phagocytose. Lidocaine was capable of recovering phagocytosis and inhibiting the release of inflammatory mediators.

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**Author contributions**

Study design: QS, MS and RW. Data acquisition: FS, YZ, LZ and GW. Data analysis: QS, LZ and JW. Manuscript writing and modification: QS, FS, RW, YZ, LZ, JW, MS and GW. Submission approval: QS, FS, RW, YZ, LZ, JW, MS and GW. All authors contributed to data analysis, drafting and revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

**Disclosure**

The author reports no conflicts of interest in this work.

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