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RESEARCH ARTICLE

U-shaped association between serum triglyceride levels and mortality among septic patients: An analysis based on the MIMIC-IV database

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Abstract

Background

Sepsis is characterized by upregulated lipolysis in adipose tissue and a high blood triglyceride (TG) level. It is still debated whether serum TG level is related to mortality in septic patients. The aim of this study is to investigate the association between serum TG level and mortality in septic patients admitted to the intensive care unit (ICU).

Methods

Data from adult septic patients (\geq 18 years) admitted to the ICU for the first time were obtained from the Multiparameter Intelligent Monitoring in Intensive Care IV (MIMIC-IV) database. The patients' serum TG levels that were measured within the first week after ICU admission were extracted for statistical analysis. The endpoints were 28-day, ICU and inhospital mortality.

Results

A total of 2,782 septic patients were included. Univariate analysis indicated that the relationship between serum TG levels and the risk of mortality was significantly nonlinear. Both the Lowess smoothing technique and restricted cubic spline analyses revealed a U-shaped association between serum TG levels and mortality among septic patients. The lowest mortality rate was associated with a serum TG level of 300–500 mg/dL. Using 300 ~ 500 mg/dL as the reference range, we found that both hypo-TG (<300 mg/dL) and hyper-TG (\geq 500 mg/ dL) were associated with increased mortality. The result was further adjusted by Cox regression with and without the inclusion of some differential covariates. **Funding:** This study was supported by the National Natural Science Foundation of China (No. 82202389, No. 81870441, and No. 82070669). The sponsor had no role in the design of the study, the collection and analysis of the data, or the preparation of the manuscript.

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Conclusions

There was a U-shaped association between serum TG and mortality in septic ICU patients. The optimal concentration of serum TG levels in septic ICU patients is 300–500 mg/dL.

Introduction

Sepsis is an inflammatory disease characterized by life-threatening organ dysfunction that is caused by a dysregulated immune response to infection [1]. Despite significant progress in understanding the underlying mechanisms of the pathogenesis of sepsis and devising clinical therapeutic interventions, sepsis remains a major disease, affecting over 30 million people worldwide every year [2–4]. In the US, the estimated national weighted incidence of sepsis is 6%, and its in-hospital mortality rate is 15.6% [5]. A recent national cross-sectional survey revealed that sepsis affects one-fifth of all patients admitted to ICUs in mainland China, and the 90-day mortality rate for sepsis is 35.5% [6]. One of the reasons for the high mortality rate of sepsis is that current therapeutic approaches, which focus on anti-inflammation, hemodynamic resuscitation and organ support, disregard metabolic disorders in sepsis [7–9].

Alterations in lipid metabolism are typical metabolic disorders that occur during sepsis [10]. Sepsis upregulates lipolysis in adipose tissue [11]. As a result, septic patients have higher levels of serum triglycerides (TGs) in the blood [12]. Many studies have attempted to investigate the relationship between TG levels and mortality in patients with sepsis. Some studies have found no correlation between TG levels and mortality in sepsis patients [13–16]. Some studies have found a negative correlation [16–18], and others have found a positive one [19]. Thus, it is still unclear whether serum TG level is related to mortality in septic patients.

Here, a total of 2,782 septic patients included in an online public database called "Multiparameter Intelligent Monitoring in Intensive Care IV" (MIMIC-IV) were studied. We found a U-shaped association between serum TG level and the risk of mortality. The lowest risk of mortality was associated with a serum TG level of 300–500 mg/dL.

Methods

Database profile

This retrospective observational study involved septic patients included in the MIMIC IV (V.1.0) database. The MIMIC IV database was developed and is maintained by the Massachusetts Institute of Technology (MIT) Laboratory for Computational Physiology. It contains the medical data of more than 257,366 patients admitted to ICUs and emergency departments between 2008 and 2019 at the Beth Israel Deaconess Medical Center [20]. It is publicly accessible to researchers who have completed the 'protecting human subjects' training course of the National Institutes of Health (NIH).

Ethical statement

The establishment of the MIMIC-IV database was approved by the Massachusetts Institute of Technology (Cambridge, MA) and Beth Israel Deaconess Medical Center (Boston, MA), and consent was obtained for the original data collection. The authors are accountable for all aspects of the work and ensure that questions related to the accuracy or integrity of any part of the work were appropriately investigated and resolved. The study adhered to the Declaration of Helsinki. The requirement for patient informed consent was waived due to the study design.

Patient inclusion

Adult septic patients from the MIMIC IV (V.1.0) database were included in this study. The inclusion criteria were as follows: (1) 18 years old or older; (2) diagnosis of sepsis in line with the sepsis 3.0 criteria [21]; (3) admission to the ICU for the first time; and (4) TG level testing within the first week after ICU admission. According to the sepsis 3.0 criteria, the diagnosis of sepsis requires the presence of infection (suspected infection) and organ dysfunction. Infection means that the patient has clear etiological evidence, and suspected infection is defined as undergoing any culture and at least two doses of antimicrobials. Organ dysfunction is defined as an acute change in total Sequential Organ Failure Assessment (SOFA) score of 2 points or higher consequent to the infection. The TG levels were recorded every day if the patients underwent TG testing within the first week after ICU admission. The exclusion criteria were as follows: (1) less than 18 years old; (2) pregnant or lactating or in a permanent vegetative state; or (3) presence of a communicable disease, such as AIDS.

Data extraction

Data were extracted by Deng, who completed the online training course of the NIH (certification number: 39921130). The extracted parameters included age, sex, weight, height, SOFA score, comorbidities, heart rate, mean arterial pressure (MAP), blood glucose level, lactate level, white blood cell (WBC), neutrophil, and platelet counts, hematocrit, hemoglobin, blood urea nitrogen (BUN), creatinine, alanine transaminase (ALT), aspartate aminotransferase (AST), albumin, total bilirubin levels, and renal replacement therapy (RRT) within 7 days. Comorbidities included hypertension, diabetes, hyperlipidemia, chronic pulmonary disease (CPD), myocardial infarct (MI), congestive heart failure (CHF), atherosclerosis, vascular disease, liver disease, renal disease, hypothyroidism, pancreatitis, and tumor. We also collected the patients' serum TG levels measured within the first week after ICU admission. Then, both the maximum value of TG (TG_{max}) and minimum value of TG (TG_{min}) were selected for statistical analysis. The endpoints were 28-day, ICU and in-hospital mortality. Data extraction was performed using PostgreSQL tools V.1.12.3.

Statistical analysis

All data were statistically analyzed using SPSS 20.0 and R 4.1.2 software. Continuous variables are presented as medians (interquartile ranges, IQRs). Differences were analyzed by the Mann-Whitney U test (for 2 groups) or Kruskal–Wallis H test (for 3 groups). Categorical variables are presented as frequencies (percentages), and their differences were analyzed by chi-square tests. In total, less than 10% of the data were missing for the continuous variables of height and weight. The missing data were imputed using multiple imputation with chain equations. Then, the body mass index (BMI) was calculated by the formula BMI = weight/(height * height). The number of participants with missing data for each variable of interest is shown in S1 Fig.

The relationship among the septic patients was further adjusted by restricted cubic splines with or without some differential covariates. The reference was set at 150 mg/dL (the upper limit of the reference value of serum TGs). Cox regression was used to explore the relationship between TG level and mortality in septic patients. Variables were selected for inclusion in the models based on statistical significance in the univariate analyses, biological plausibility and known associations with in-hospital mortality. Kaplan–Meier (K-M) survival curves were constructed. P < 0.05 was considered statistically significant.

Results

Low serum TG values are associated with higher mortality

In total, 11,263 adult septic patients were included in the MIMIC-IV database. After a rigorous screening process, 2,782 septic patients were included in this study (Fig 1). Their median age (IQR) was 63.9 (51.8, 74.5) years, and female patients accounted for 40.2% of patients. Among the included patients, 605 (21.7%) died within 28 days, 487 (17.5%) died in the ICU, and 651 (23.4%) died in the hospital. We found that the serum TG levels of the patients who died within 28 days were lower than those of the survivors (Table 1). Similar results were found in the patients who died in the ICU (S1 Table) and in the hospital (S2 Table). These data indicated that a low serum TG value was associated with higher mortality.

In total, 1,444 (51.9%) patients had serum TG_{max} levels of less than 150 mg/dL (clinically normal TG level), 1,118 (40.2%) patients had serum TG_{max} levels between 150 mg/dL and 500 mg/dL, and the remaining 220 (7.9%) patients had serum TG_{max} levels above 500 mg/dL. We calculated mortality in the septic patients in the 3 groups defined by the different TG_{max} levels. The results showed that patients with serum TG_{max} levels of 150 ~ 500 mg/dL had the lowest mortality (Table 2). Similar results were found in serum TG_{min} levels (Table 2).

All of these results indicated a nonlinear association between serum TG levels and mortality among septic patients.

U-shaped association between serum TG and mortality

By using the Lowess smoothing technique, we preliminarily explored the relationship between serum TG levels and mortality in patients with sepsis. Three models yielded nonlinear relationships, with the lowest mortality rate at a serum TG level of $300 \sim 500 \text{ mmol/L}$ (Figs 2A-2C and S2A-S2C).



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Variables	Total (n = 2782)	Survivors (n = 2177)	Non-survivors $(n = 605)$	<i>p</i> value
Age (years)	63.9 (51.8, 74.5)	63.2 (50.5, 73.8)	61.2 (55.9, 77.3)	<0.001
Female (n (%))	1119 (40.2)	877 (40.3)	232 (38.4)	0.389
BMI (kg/m ²)	28.4 (24.4, 33.8)	28.4 (24.5, 33.7)	28.2 (24.1, 33.9)	<0.001
SOFA score	8.0 (5.0, 12.0)	7.0 (5.0, 11.0)	11.0 (7.0, 14.0)	<0.001
Comorbidities	·			
Hypertension (n (%))	1075 (38.6)	851 (39.1)	224 (37.0)	0.356
Diabetes (n (%))	834 (30.0)	644 (29.6)	190 (31.4)	0.387
Hyperlipidemia (n (%))	890 (32.0)	689 (31.7)	201 (33.2)	0.463
CPD (n (%))	754 (27.1)	574 (26.4)	180 (29.8)	0.097
MI (n (%))	516 (18.6)	393 (18.1)	123 (20.3)	0.202
CHF (n (%))	902 (32.4)	670 (30.8)	232 (38.4)	<0.001
Atherosclerosis (n (%))	336 (32.4)	247 (11.4)	89 (14.7)	0.025
Vascular disease (n (%))	978 (35.2)	751 (34.5)	227 (37.5)	0.168
Liver disease (n (%))	600 (21.6)	420 (19.3)	180 (29.8)	<0.001
Renal disease (n (%))	593 (21.3)	434 (19.9)	159 (26.3)	0.001
Hypothyroidism (n (%))	348(12.5)	268(12.3)	80(13.2)	0.548
Pancreatitis (n (%))	202(7.3)	170(7.8)	32(5.3)	0.035
Tumor (n (%))	332 (11.9)	228 (10.47)	104 (17.2)	<0.001
During the first 24 hours after ICU a	dmission			
Heart rate (beat/min)	109.0 (95.0, 124.0)	108.0 (95.0, 124.0)	111.0 (96.0, 126.0)	0.480
MAP (mmHg)	59.0 (53.00, 66.0)	59.0 (53.0, 66.0)	58.0 (52.0, 63.0)	<0.001
Blood glucose (mg/dL)	167.0 (131.0, 225.0)	165.0 (130.0, 222.0)	177.0 (137.00, 241.00)	0.003
Lactate (mmol/L)	2.2 (1.6, 3.1)	2.2 (1.5, 2.9)	2.2 (1.9, 4.5)	<0.001
WBC (×10 ⁹ /L)	13.9 (10.1, 19.0)	13.7 (9.9, 18.6)	15.5 (11.2, 21.4)	<0.001
Platelet (×10 ⁹ /L)	209.0 (148.0, 281.0)	212.0 (151.0, 285.0)	197.0 (132.0, 267.0)	0.001
Hematocrit (%)	35.5 (30.5, 40.9)	35.6 (30.7, 41.0)	34.5 (29.3, 40.5)	<0.001
Hemoglobin (g/L)	11.60 (9.90, 13.5)	11.7 (10.0, 13.5)	11.1 (9.5, 13.2)	<0.001
BUN (mg/dL)	24.0 (16.0, 39.0)	22.0 (15.0, 36.0)	30.0 (20.0, 51.0)	<0.001
Creatinine (mg/dL)	1.2 (0.9, 2.0)	1.2 (0.8, 1.9)	1.5 (1.0, 2.4)	<0.001
ALT (IU/L)	31.0 (21.0, 50.0)	31.0 (21.0, 49.0)	31.0 (21.0, 59.0)	0.324
AST (IU/L)	47.0 (32.0, 79.0)	47.0 (31.0, 73.0)	47.0 (37.0, 104.0)	<0.001
Albumin (g/dL)	3.3 (3.0, 3.5)	3.3 (3.1, 3.5)	3.3 (2.8, 3.4)	0.001
Total bilirubin (mg/dL)	0.8 (0.5, 1.4)	0.8 (0.5, 1.2)	0.8 (0.6, 2.2)	<0.001
RRT within 7 days (n (%))	443 (15.9)	288 (13.2)	155 (25.6)	<0.001
TG _{max} level (mg/dL)	145.0 (93.0, 249.0)	148.0 (95.0, 254.0)	136.0 (88.0, 227.0)	0.013
TG _{min} level (mg/dL)	132.5 (88.0, 207.0)	135.0 (90.0, 211.0)	123.0 (84.0, 190.0)	0.001

Table 1. Characteristics of survivors and non-survivors within 28 days.

Note: Continuous variables (age, BMI, MAP, *et al.*) were presented as median (IQR). Categorical variables (hypertension, CPD, MI, *et al.*) were presented as frequencies (percentages). The differences between survivors and non-survivors were analyzed by Mann-Whitney U test, and Chi-square. BMI: body mass index; SOFA: sequential organ failure assessment; CPD: chronic pulmonary disease; MI: myocardial infarct; CHF: congestive heart failure; MAP: mean arterial pressure; WBC: white blood cell; BUN: blood urea nitrogen; ALT: alanine transaminase; AST: aspartate aminotransferase; RRT: renal replacement therapy; TG_{max}: maximum value of triglycerides; TG_{min}: minimum value of triglycerides.

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The relationship between serum TG levels and risk of mortality was further adjusted by restricted cubic splines with or without some differential covariates. When the serum TG_{max} level was examined as a continuous variable in relation to mortality, a significant nonlinear pattern (p < 0.05) was demonstrated, which resulted in a U-shaped curve (Fig 2D–2F). The

Total	< 150 mg/dL	150–500 mg/dL	> 500 mg/dL	<i>p</i> value
2782	1444 (51.9)	1118 (40.2)	220 (7.9)	
605 (21.7)	341 (23.61)	211 (18.87) *	53 (24.09) #	0.011
487 (17.51)	270 (18.70)	169 (15.12) *	48 (21.82) #	0.013
651 (23.40)	363 (25.14)	229 (20.48) *	59 (26.82) #	0.010
2782	1627 (58.5)	1047 (37.6)	108 (3.9)	
605 (21.7)	386 (23.7)	198 (18.9) *	21 (19.4)	0.011
487 (17.51)	313 (19.2)	153 (14.6) *	21 (19.4)	0.008
651 (23.40)	415 (25.5)	211 (20.2) *	25 (23.1)	0.006
	Total 2782 605 (21.7) 487 (17.51) 651 (23.40) 2782 605 (21.7) 487 (17.51) 651 (23.40)	Total < 150 mg/dL 2782 1444 (51.9) 605 (21.7) 341 (23.61) 487 (17.51) 270 (18.70) 651 (23.40) 363 (25.14) 2782 1627 (58.5) 605 (21.7) 386 (23.7) 487 (17.51) 313 (19.2) 651 (23.40) 415 (25.5)	Total< 150 mg/dL150-500 mg/dL27821444 (51.9)1118 (40.2)605 (21.7)341 (23.61)211 (18.87)*487 (17.51)270 (18.70)169 (15.12)*651 (23.40)363 (25.14)229 (20.48)*27821627 (58.5)1047 (37.6)605 (21.7)386 (23.7)198 (18.9)*487 (17.51)313 (19.2)153 (14.6)*651 (23.40)415 (25.5)211 (20.2)*	Total< 150 mg/dL150-500 mg/dL> 500 mg/dL27821444 (51.9)1118 (40.2)220 (7.9) $605 (21.7)$ $341 (23.61)$ $211 (18.87)^*$ $53 (24.09)^*$ $487 (17.51)$ 270 (18.70) $169 (15.12)^*$ $48 (21.82)^*$ $651 (23.40)$ $363 (25.14)$ 229 (20.48) * $59 (26.82)^*$ 2782 $1627 (58.5)$ $1047 (37.6)$ $108 (3.9)$ $605 (21.7)$ $386 (23.7)$ $198 (18.9)^*$ $21 (19.4)$ $487 (17.51)$ $313 (19.2)$ $153 (14.6)^*$ $21 (19.4)$ $651 (23.40)$ $415 (25.5)$ $211 (20.2)^*$ $25 (23.1)$

Table 2. Mortality of different serum TG groups (clinical boundary values).

Note: Data were presented as frequencies (percentages). The differences between survivors and non-survivors were analyzed by Chi-square. *Compared with < 150 mg/dL, P < 0.05

[#]Compared with 150–500 mg/dL, *P* < 0.05. TG_{max}: maximum value of triglycerides; TG_{min}: minimum value of triglycerides; ICU: Intensive Care Unit.

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lowest 28-day mortality rate was observed with a TG_{max} level of 396.5 mg/dL. The lowest ICU mortality rate was observed with a TG_{max} level of 459.6 mg/dL, and the lowest in-hospital mortality rate was observed with a TG_{max} level of 410.5 mg/dL. Similar results were found in serum TG_{min} levels (S2D-S2F Fig). All of these results indicate a U-shaped association between serum TG levels and mortality among septic patients.



Fig 2. U-shaped association between serum TG_{max} **levels and mortality among septic patients.** A. Lowess smoothing technique revealed that the lowest 28-day mortality (12.22%) was associated with TG_{max} levels of 400–500 mg/dL. B. Lowess smoothing technique revealed that the lowest ICU mortality (10.00%) was associated with TG_{max} levels of 400–500 mg/dL. C. Lowess smoothing technique revealed that the lowest hospital mortality (13.33%) was associated with a TG_{max} level of 400–500 mg/dL. D. Restricted cubic splines revealed that the relationship between serum TG_{max} level and risk of 28-day mortality was significantly nonlinear (p = 0.039), resulting in a U-shaped association. The lowest hospital mortality (HR: 0.78, 95% CI: 0.61, 0.99; p < 0.05) was associated with a TG_{max} level of 396.5 mg/dL. E. Restricted cubic splines revealed that the relationship between serum TG_{max} level and risk of ICU mortality was significantly nonlinear (p = 0.002), resulting in a U-shaped association. The lowest ICU mortality (HR: 0.70, 95% CI: 0.53, 0.93; p < 0.05) was associated with a TG_{max} level of 459.6 mg/dL. F. Restricted cubic splines revealed that the relationship between serum TG_{max} level and risk of hospital mortality was significantly nonlinear (p = 0.002), resulting in a U-shaped association. The lowest ICU mortality (HR: 0.70, 95% CI: 0.53, 0.93; p < 0.05) was associated with a TG_{max} level of 459.6 mg/dL. F. Restricted cubic splines revealed that the relationship between serum TG_{max} levels and the risk of hospital mortality was significantly nonlinear (p = 0.013), resulting in a U-shaped association. The lowest hospital mortality (HR: 0.78, 95% CI: 0.66, 0.99; p < 0.05) was associated with a TG_{max} level of 410.5 mg/dL. TG_{max}: the maximum value of serum triglyceride; ICU: intensive care unit.

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In this study, both Lowess smoothing technique and restricted cubic splines indicated that the boundaries of TG levels were less than 300 mg/dL, 300–500 mg/dL and greater than 500 mg/dL. However, clinically normal TG levels are less than 150 mg/dL. To investigate the effect of mildly elevated TG levels on the mortality of septic patients, a total of 2,272 septic patients who had TG levels of less than 300 mg/dL were divided into a normal group (TG_{max} less than 300 mg/dL) and an elevated group (TG_{max}: 300–500 mg/dL). We found that there was no significant difference in mortality (including 28-day mortality, ICU mortality and in-hospital mortality) between the 2 groups (S3 Table). Therefore, we artificially divided TG values into 3 boundaries: less than 300 mg/dL, 300 to 500 mg/dL, and greater than 500 mg/dL.

Both hypo-TG and hyper-TG were associated with increased mortality

According to the U-shaped association, TG_{max} values were selected and categorized into 3 groups: hypo- TG_{max} (< 300 mg/dL), moderate- TG_{max} (300 ~ 500 mg/dL), and hyper- TG_{max} (\geq 500 mg/dL). The primary clinical outcomes unadjusted for covariates for the 3 TG_{max} groups are presented in Table 3. The results showed that hypo- TG_{max} or hyper- TG_{max} was associated with increased mortality. To explore the effect of TG levels on mortality, moderate- TG_{max} was selected as the reference group for the Cox regression models. The results showed that both hypo- TG_{max} and hyper- TG_{max} were associated with increased mortality among septic patients (Table 4). Survival analysis (Kaplan–Meier) further indicated that patients with hypo- TG_{max} and hyper- TG_{max} had higher mortality rates (Fig 3).

Discussion

The amount of TGs is directly proportional to the severity of sepsis [22], but evidence of how TG levels affect the prognosis of sepsis is limited. In this study, 2,782 septic patients with serum TG levels measured within the first week after ICU admission were selected from the MIMIC-IV database. Our results first revealed a U-shaped association between serum TG levels and mortality among ICU-admitted septic patients. The lowest mortality rate was associated with a serum TG level of 300–500 mg/dl. Both hypo-TG (<300 mg/dL) and hyper-TG (\geq 500 mg/dL) were associated with increased mortality among septic patients. To the best of our knowledge, this study is the first retrospective observational study to suggest a link between serum TG (within the first week after ICU admission) imbalance and mortality among septic patients.

TGs are biomolecules that were discovered many years ago. They are derived from glycerol and three fatty acids and have key roles in the storage and transportation of fatty acids in cells and plasma [23-25]. Under normal physiological conditions, a small amount of TGs (<150 mg/dL) can be detected in peripheral blood. Sepsis is accompanied by severe metabolic alterations. In 1992, Kenneth et al. demonstrated that lipopolysaccharide (LPS) can rapidly induce hypertriglyceridemia [26]. They found that low doses of LPS can stimulate hepatic TG production, while high doses of LPS may reduce TG hydrolysis in cells and plasma. Further study demonstrated a shift from glucose metabolism to lipid metabolism in LPS-challenged mice [27]. They found that in sepsis, lipids are transported to the liver and serve as the dominant energy source for septic mice. In addition, a marked elevation in serum TG levels has been observed in septic models and patients [11, 17, 18, 28]. Furthermore, a recent published study reported that baseline TG levels were significantly higher in patients with septic shock than in patients with sepsis [29]. All of these results indicate that sepsis is usually accompanied by elevated serum TG levels, and TGs are the main energy supply substrate for sepsis patients. In this study, we found that nearly half of the septic patients had serum TG levels at the limit of the reference value (150 mg/dL), and these patients had higher mortality rates. Moreover,

Variables	Hypo-TG _{max} (n = 1,444)	Moderate-TG _{max} (n = 1,118)	Hyper-TG _{max} ($n = 220$)	<i>p</i> value
Age (years)	65.4 (53.9, 76.3)	56.5 (42.7 66.5)	54.75 (42.5, 66.2)	< 0.001
Female (n (%))	935 (41.2)	99 (34.1)	85 (38.6)	0.063
BMI (kg/m ²)	27.8 (23.9, 32.8)	30.9 (26.5, 37.1)	32.4 (27.7, 37.1)	< 0.001
SOFA score	8.0 (5.0, 11.0)	9.0 (6.0, 13.0)	11.0 (7.0, 13.0)	< 0.001
Comorbidities				·
Hypertension (n (%))	878 (38.6)	121 (41.7)	76 (34.6)	0.257
Diabetes (n (%))	660 (29.15)	92 (31.7)	82 (37.3)	0.031
Hyperlipidemia (n (%))	743 (32.7)	80 (27.6)	67 (30.5)	0.187
CPD (n (%))	609 (26.8)	90 (31.0)	55 (25.0)	0.239
MI (n (%))	455 (20.0)	34 (11.7)	27 (12.3)	< 0.001
CHF (n (%))	790 (34.8)	72 (24.8)	40 (18.2)	< 0.001
Atherosclerosis (n (%))	266 (11.7)	38 (13.1)	32 (14.6)	0.398
Vascular disease (n (%))	852 (37.5)	65 (22.4)	61 (27.7)	< 0.001
Liver disease (n (%))	493 (21.7)	55 (19.0)	52 (23.6)	0.419
Renal disease (n (%))	491 (21.6)	53 (18.3)	49 (22.3)	0.399
Tumor (n (%))	266 (11.7)	39 (13.5)	27 (12.3)	0.681
During the first 24 hours after ICU	J admission			·
Heart rate (beat/min)	108.0 (94.0, 123.0)	114.5 (100.0, 130.3)	117.0 (101.0, 128.5)	< 0.001
MAP (mmHg)	59.0 (53.0, 66.0)	59.0 (54.0, 65.0)	58.0 (52.0, 65.0)	0.501
Blood glucose (mg/dL)	164.0 (130.0, 219.0)	175.0 (133.8, 257.3)	182.5 (141.3, 266.5)	< 0.001
Lactate (mmol/L)	2.2 (1.6, 3.0)	2.2 (1.5, 3.8) 2.2 (1.4, 4.4)		0.567
WBC (×10 ⁹ /L)	13.9 (9.9, 18.7)	14.8 (10.6, 20.1)	13.9 (10.7, 20.9)	0.026
Platelet (×10 ⁹ /L)	210.0 (149.3, 280.0)	0) 216.0 (151.8, 296.0) 191.5 (130.0, 265.8)		0.027
Hematocrit (%)	35.5 (30.6, 41.0)	35.5 (29.6, 41.2)	34.7 (29.8, 40.1)	0.223
Hemoglobin (g/L)	11.6 (9.9, 13.4)	11.6 (9.7, 13.7)	11.4 (9.7, 13.2)	0.347
BUN (mg/dL)	23.0 (15.0, 38.0)	24.0 (17.0, 46.3)	26.0 (17.3, 51.8)	0.004
Creatinine (mg/dL)	1.2 (0.8, 1.8)	1.3 (0.9, 2.4)	1.6 (1.0, 2.6)	< 0.001
ALT (IU/L)	31.0 (20.0, 47.0)	31.0 (24.0, 65.0)	36.5 (26.0, 98.0)	< 0.001
AST (IU/L)	47.0 (30.0, 73.0)	47.0 (38.8, 88.0)	47.0 (42.3, 1.5)	< 0.001
Albumin (g/dL)	3.3 (3.1, 3.6)	3.3 (2.9, 3.5)	3.3 (2.7, 3.3)	< 0.001
Total bilirubin (mg/dL)	0.8 (0.5, 1.3)	0.8 (0.5, 1.5)	0.8 (0.5, 2.0)	0.284
RRT within 7days (n (%))	306 (13.5)	58 (20.0) *	79 (35.9) *#	< 0.001
28-day mortality (n (%))	507(22.32)	45(15.52)*	53(24.09) #	0.021
ICU mortality (n (%))	401 (17.7)	38 (13.1)	48 (21.8) #	< 0.001
Hospital mortality (n (%))	543 (23.9)	49 (16.9) *	59 (26.8) #	0.014

Table 3. Both hypo-TG_{max} and hyper-TG_{max} are associated with higher mortality.

Note: Hypo-TG_{max}: <300 mg/dL, Moderate-TG_{max}: $300 \sim 500 \text{ mg/dL}$, Hyper-TG_{max} $\geq 500 \text{ mg/dL}$. Continuous variables (age, BMI, MAP, *et al.*) were presented as median (IQR). Categorical variables (hypertension, CPD, MI, *et al.*) were presented as frequencies (percentages). The difference among the 3 groups was analyzed by Kruskal-Wallis H test.

^{*}Compared with Hypo-TG_{max}, P < 0.05

[#]Compared with Moderate- TG_{max} , P < 0.05. BMI: body mass index; SOFA: sequential organ failure assessment; CPD: chronic pulmonary disease; MI: myocardial infarct; CHF: congestive heart failure; MAP: mean arterial pressure; WBC: white blood cell; BUN: blood urea nitrogen; ALT: alanine transaminase; AST: aspartate aminotransferase; RRT: renal replacement therapy; ICU: Intensive Care Unit; TG: Triglycerides.

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similar results were observed in patients with excessively high serum TG levels (>500 mg/dL). Furthermore, we revealed that the nonsurvivors had lower serum TG values than the survivors. These findings indicate that the serum TG level is directly proportional to the severity of sepsis but inversely proportional to the mortality rate in septic patients.

Models	28-day mortality			ICU mortality			In-hospital mortality		
	HR	95% CI	<i>p</i> value	HR	95% CI	<i>p</i> value	HR	95% CI	<i>p</i> value
Model 1									
Hypo-TG	1.51	1.11, 2.04	0.008	1.96	1.40, 2.74	< 0.001	1.78	1.33, 2.39	< 0.001
Hyper-TG	1.63	1.09, 2.42	0.016	1.81	1.18, 2.77	0.007	1.71	1.17, 2.50	0.005
Model 2									
Hypo-TG	1.52	1.11, 2.07	0.009	1.90	1.35, 2.68	< 0.001	1.71	1.27, 2.30	< 0.001
Hyper-TG	1.52	1.02, 2.26	0.039	1.77	1.16, 2.71	0.009	1.68	1.15, 2.46	0.007
Model 3									
Hypo-TG	1.43	1.05, 1.96	0.025	1.81	1.29, 2.56	0.001	1.59	1.18, 2.15	0.002
Hyper-TG	1.52	1.02, 2.27	0.040	1.84	1.20, 2.82	0.006	1.69	1.16, 2.48	0.007
Model 4									
Hypo-TG	1.46	1.06, 2.00	0.019	1.86	1.31, 2.63	< 0.001	1.60	1.18, 2.17	0.002
Hyper-TG	1.57	1.05, 2.35	0.029	1.84	1.19, 2.84	0.006	1.71	1.16, 2.52	0.006
Model 5									
Hypo-TG	1.48	1.08, 2.04	0.014	1.90	1.34, 2.69	< 0.001	1.62	1.20, 2.20	0.002
Hyper-TG	1.50	1.00, 2.25	0.049	1.79	1.16, 2.78	0.009	1.67	1.13, 2.46	0.009

Table 4. Cox regression based on serum TG_{max} groups.

Note: Among the 3 TG groups, moderate-TG ($300 \sim 500 \text{ mg/dL}$) was adopted as the reference group. Hypo-TG: <300 mg/dL, Hyper-TG $\geq 500 \text{ mg/dL}$. Model 1: unadjusted model.

Model 2: Adjusted for age, gender, BMI and SOFA score.

Model 3: Adjusted for variables included in model 2 + hypertension, diabetes, hyperlipidemia, CPD, MI, CHF, atherosclerosis, vascular disease, liver disease, renal disease, and tumor.

Model 4: Adjusted for variables included in model 3 + heart rate, MAP, blood glucose, lactate, WBC, platelet, hematocrit, hemoglobin. BUN, creatinine, ALT, AST, albumin, and total bilirubin.

Model 5: Adjusted for variables included in model 4 + RRT within 7days.

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The main function of TG is to store unused calories and provide the body with energy. Previous studies showed that lower serum TG levels were significantly associated with higher hospital mortality rates among septic patients [18, 24, 30–32]. Furthermore, Maile et al. reported that the pre-sepsis TG level was associated with the severity of illness, and high TG values led to higher in-hospital mortality among septic patients [19]. A recent quantitative proteomic analysis



Fig 3. Survival curve based on serum TG groups. Both hypo-TG and hyper-TG were associated with higher 28-day mortality (A), ICU mortality (B) and hospital mortality (C) among septic patients. TG_{max}: the maximum value of serum triglyceride; ICU: intensive care unit.

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showed that TG/cholesterol homeostasis was associated with sepsis-related acute kidney injury [33]. However, many other studies, which involved simple comparisons between survivors and nonsurvivors, reported similar TG levels in deceased patients and survivors [13–16].

Unlike previous studies, our study revealed a 'U'-shaped relationship between changes in serum TG levels and mortality among septic patients. Further analysis revealed that both hypo-TG (<300 mg/dL) and hyper-TG ($\geq 500 \text{ mg/dL}$) were associated with increased mortality. These results were similar to those of Choi et al., who reported that both hypertriglyceridemia and hypotriglyceridemia may be associated with poor early outcomes of acute ischemic stroke [34]. While the specific mechanism remains unclear, we speculate that it may be as follows. First, the body of a septic patient enters a state of stress, resulting in hypermetabolism and higher energy consumption. Therefore, an extremely low level of TGs, which serve to store energy, may lead to poor prognoses [35-37]. Second, lipid metabolism disorders may seriously damage immune function, leading to uncontrolled infection and inflammatory damage to tissues and organs [38-40]. Recent studies have pointed out that TG-based markers are associated with many inflammatory conditions including type 2 diabetes mellitus, coronary heart disease and liver steatosis [41-43]. The initial phase of sepsis is characterized by overwhelming inflammation [44]. Interestingly, a recent study revealed that a combination of bis (α -furancarboxylato) oxovanadium and metformin ameliorates hepatic steatosis in high-fat diet-induced obese mice by alleviating hepatic inflammation and enhancing the insulin signaling pathway [45]. Thus, we speculate that TG may be associated with sepsis mortality via the inflammation pathway. This should be further investigated soon. In addition, hyper-TG is associated with elevated blood viscosity and reduced blood flow velocity [46]. As a result, hypertriglyceridemia restricts the blood supply to tissue and organs. All of these factors result in a poor prognosis in septic patients.

There are some limitations to this study. First, this is a retrospective trial, and the data in the MIMIC-IV database were not collected for this purpose. Therefore, our findings should be considered purely hypothesis-generating rather than confirmatory and need to be further validated in clinical practice. Second, the risk of sampling bias should be considered. In this study, our data were extracted from the MIMIC-IV database. The database is not a routine laboratory for sepsis management. Therefore, the act of simply having one drawn is likely to add bias to the sample. Third, the nutritional statuses of the patients before ICU admission were not clear. We extracted only height and weight parameters and calculated BMI values. No significant difference was found in BMI between the survivors and nonsurvivors. Fourth, we did not obtain data on inflammatory cytokines (including interleukin-1, interleukin-6 and tumor necrosis factor α) that may have contributed to lipid metabolism [47]. Inflammatory cytokines were not available in the MIMIC-IV (1.0) database. Fifth, our study is a single-center study that may have potential bias. Therefore, a better-designed prospective study involving different ethnic groups is required to reconfirm the role of the TG level in the prognosis of septic patients.

Conclusions

In conclusion, this study is the first retrospective observational study to suggest a U-shaped association between serum TG (within the first week after ICU admission) imbalance and mortality among septic patients. According to the link between serum TG (within the first week after ICU admission) imbalance and mortality among septic patients, we should maintain serum TG levels in septic patients at 300–500 mg/dL.

Supporting information

S1 Table. Characteristics of survivors and non-survivors within ICU. (DOCX)

S2 Table. Characteristics of survivors and non-survivors in hospital. (DOCX)

S3 Table. Mortality of different serum TG_{max} groups (clinical boundary values). (DOCX)

S1 Fig. Number of participants with missing data for each variable of interest. (DOCX)

S2 Fig. U-shaped association between serum TG_{min} levels and mortality among septic patients. (DOCX)

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References

- Rhodes A, Evans LE, Alhazzani W, et al., Surviving Sepsis Campaign: International Guidelines for Management of Sepsis and Septic Shock: 2016. Intensive Care Med 2017; 43(3): 304–377. <u>https://doi.org/10.1007/s00134-017-4683-6 PMID: 28101605</u>
- Evans L, Rhodes A, Alhazzani W, et al., Surviving sepsis campaign: international guidelines for management of sepsis and septic shock 2021. Intensive Care Med 2021. https://doi.org/10.1007/s00134-021-06506-y PMID: 34599691
- Fleischmann-Struzek C, Mellhammar L, Rose N, et al., Incidence and mortality of hospital- and ICUtreated sepsis: results from an updated and expanded systematic review and meta-analysis. Intensive Care Med 2020; 46(8): 1552–1562. https://doi.org/10.1007/s00134-020-06151-x PMID: 32572531
- Rudd KE, Johnson SC, Agesa KM, et al., Global, regional, and national sepsis incidence and mortality, 1990–2017: analysis for the Global Burden of Disease Study. Lancet 2020; 395(10219): 200–211. https://doi.org/10.1016/S0140-6736(19)32989-7 PMID: 31954465
- Rhee C, Dantes R, Epstein L, et al., Incidence and Trends of Sepsis in US Hospitals Using Clinical vs Claims Data, 2009–2014. JAMA 2017; 318(13): 1241–1249. https://doi.org/10.1001/jama.2017.13836 PMID: 28903154

- Xie J, Wang H, Kang Y, et al., The Epidemiology of Sepsis in Chinese ICUs: A National Cross-Sectional Survey. Crit Care Med 2020; 48(3): e209–e218. <u>https://doi.org/10.1097/CCM.00000000004155</u> PMID: 31804299
- Chang P and Liu Z, Metabolic Resuscitation Therapy for Sepsis: Where to Go. Intensive Care Research 2021; 1(1–2). https://doi.org/10.2991/icres.k.210622.002
- Reitsema VA, Star BS, de Jager VD, et al., Metabolic Resuscitation Strategies to Prevent Organ Dysfunction in Sepsis. Antioxid Redox Signal 2019; 31(2): 134–152. https://doi.org/10.1089/ars.2018.7537 PMID: 30403161
- Leite HP and de Lima LF, Metabolic resuscitation in sepsis: a necessary step beyond the hemodynamic? J Thorac Dis 2016; 8(7): E552–7. https://doi.org/10.21037/jtd.2016.05.37 PMID: 27501325
- Amunugama K, Pike DP and Ford DA, The lipid biology of sepsis. J Lipid Res 2021; 62(100090. <u>https://doi.org/10.1016/j.jlr.2021.100090 PMID: 34087197</u>
- Golucci A, Marson FAL, Ribeiro AF, et al., Lipid profile associated with the systemic inflammatory response syndrome and sepsis in critically ill patients. Nutrition 2018; 55–56(7–14. <u>https://doi.org/10.1016/j.nut.2017.10.022</u>
- 12. Wasyluk W and Zwolak A, Metabolic Alterations in Sepsis. J Clin Med 2021; 10(11). https://doi.org/10. 3390/jcm10112412 PMID: 34072402
- 13. Tanaka S, Stern J, Bouzid D, et al., Relationship between lipoprotein concentrations and short-term and 1-year mortality in intensive care unit septic patients: results from the HIGHSEPS study. Ann Intensive Care 2021; 11(1): 11. https://doi.org/10.1186/s13613-021-00800-0 PMID: 33469739
- Guirgis FW, Leeuwenburgh C, Moldawer L, et al., Lipid and lipoprotein predictors of functional outcomes and long-term mortality after surgical sepsis. Annals of Intensive Care 2021; 11(1). https://doi.org/10.1186/s13613-021-00865-x PMID: 34018068
- Nabavi A, Allami A and QasemiBarqi R, Changes in plasma lipid and in-hospital deaths in patients with sepsis. Med J Islam Repub Iran 2020; 34(45. https://doi.org/10.34171/mjiri.34.45 PMID: 32884920
- Delirrad M, Gharebaghi N, Mobarhan S, et al., Relationship Between Lipid Profile and Sepsis Outcome in Intensive Care Unit. Archives of Clinical Infectious Diseases 2020; 15(2). https://doi.org/10.5812/ archcid.93533
- Lee SH, Park MS, Park BH, et al., Prognostic Implications of Serum Lipid Metabolism over Time during Sepsis. Biomed Res Int 2015; 2015(789298. https://doi.org/10.1155/2015/789298 PMID: 26351639
- 18. Cetinkaya A, Erden A, Avci D, et al., Is hypertriglyceridemia a prognostic factor in sepsis? Ther Clin Risk Manag 2014; 10(147–50. https://doi.org/10.2147/TCRM.S57791 PMID: 24600230
- Maile MD, Sigakis MJ, Stringer KA, et al., Impact of the pre-illness lipid profile on sepsis mortality. J Crit Care 2020; 57(197–202. https://doi.org/10.1016/j.jcrc.2020.01.016 PMID: 32182565
- Johnson A, Bulgarelli L, Pollard T, et al., MIMIC-IV (version 1.0). PhysioNet 2021. <u>https://doi.org/10.13026/s6n6-xd98</u>.
- Singer M, Deutschman CS, Seymour CW, et al., The Third International Consensus Definitions for Sepsis and Septic Shock (Sepsis-3). JAMA 2016; 315(8): 801–10. https://doi.org/10.1001/jama.2016.0287 PMID: 26903338
- 22. Fahila R, Kembaren T and Rahimi A. The effect of bacterial sepsis severity on triglyceride value. in 1st International Conference on Tropical Medicine and Infectious Diseases (ICTROMI). 2017. Univ Sumatera Utara, Fac Med, Medan, INDONESIA.
- Spitler KM and Davies BSJ, Aging and plasma triglyceride metabolism. J Lipid Res 2020; 61(8): 1161– 1167. https://doi.org/10.1194/jlr.R120000922 PMID: 32586846
- Barcia AM and Harris HW, Triglyceride-rich lipoproteins as agents of innate immunity. Clin Infect Dis 2005; 41 Suppl 7(S498–503. https://doi.org/10.1086/432005 PMID: 16237653
- AbouRjaili G, Shtaynberg N, Wetz R, et al., Current concepts in triglyceride metabolism, pathophysiology, and treatment. Metabolism 2010; 59(8): 1210–20. https://doi.org/10.1016/j.metabol.2009.11.014 PMID: 20060141
- 26. Feingold KR, Staprans I, Memon RA, et al., Endotoxin rapidly induces changes in lipid metabolism that produce hypertriglyceridemia: low doses stimulate hepatic triglyceride production while high doses inhibit clearance. Journal of Lipid Research 1992; 33(12): 1765–1776. <u>https://doi.org/10.1016/s0022-2275(20)41334-3 PMID: 1479286</u>
- Irahara T, Sato N, Otake K, et al., Alterations in energy substrate metabolism in mice with different degrees of sepsis. J Surg Res 2018; 227(44–51. <u>https://doi.org/10.1016/j.jss.2018.01.021</u> PMID: 29804861

- Khovidhunkit W, Kim MS, Memon RA, et al., Effects of infection and inflammation on lipid and lipoprotein metabolism: mechanisms and consequences to the host. J Lipid Res 2004; 45(7): 1169–96. https://doi.org/10.1194/jlr.R300019-JLR200 PMID: 15102878
- Karampela I, Chrysanthopoulou E, Skyllas G, et al., Alterations in serum lipid profile in critically ill septic patients: a prospective study. European Respiratory Journal 2020; 56 https://doi.org/10.1183/ 13993003.congress-2020.2753
- Chenaud C, Merlani PG, Roux-Lombard P, et al., Low apolipoprotein A-I level at intensive care unit admission and systemic inflammatory response syndrome exacerbation. Crit Care Med 2004; 32(3): 632–7. https://doi.org/10.1097/01.ccm.0000114820.47460.0a PMID: 15090939
- Harris HW, Grunfeld C, Feingold KR, et al., Human very low density lipoproteins and chylomicrons can protect against endotoxin-induced death in mice. J Clin Invest 1990; 86(3): 696–702. https://doi.org/10. 1172/JCI114765 PMID: 2394827
- Black LP, Puskarich MA, Henson M, et al., Quantitative and Qualitative Assessments of Cholesterol Association With Bacterial Infection Type in Sepsis and Septic Shock. J Intensive Care Med 2021; 36 (7): 808–817. https://doi.org/10.1177/0885066620931473 PMID: 32578468
- Liang X, Wu T, Chen Q, et al., Serum proteomics reveals disorder of lipoprotein metabolism in sepsis. Life science alliance 2021; 4(10). https://doi.org/10.26508/lsa.202101091 PMID: 34429344
- Choi KH, Park MS, Kim JT, et al., Serum triglyceride level is an important predictor of early prognosis in patients with acute ischemic stroke. J Neurol Sci 2012; 319(1–2): 111–6. https://doi.org/10.1016/j.jns. 2012.04.018 PMID: 22578636
- Alverdy JC, Hypermetabolism and Nutritional Support in Sepsis. Surg Infect (Larchmt) 2018; 19(2): 163–167. https://doi.org/10.1089/sur.2017.313 PMID: 29394142
- 36. Vincent JL, Metabolic support in sepsis and multiple organ failure: more questions than answers. Crit Care Med 2007; 35(9 Suppl): S436–40. <u>https://doi.org/10.1097/01.CCM.0000278601.93369.72</u> PMID: 17713390
- Pravda J, Metabolic theory of septic shock. World J Crit Care Med 2014; 3(2): 45–54. <u>https://doi.org/10.5492/wjccm.v3.i2.45 PMID: 24892019</u>
- Bermudes ACG, de Carvalho WB, Zamberlan P, et al., Changes in lipid metabolism in pediatric patients with severe sepsis and septic shock. Nutrition 2018; 47(104–109. <u>https://doi.org/10.1016/j.nut.2017.</u> 09.015 PMID: 29429528
- Sharma NK, Ferreira BL, Tashima AK, et al., Lipid metabolism impairment in patients with sepsis secondary to hospital acquired pneumonia, a proteomic analysis. Clin Proteomics 2019; 16(29. <u>https://doi.org/10.1186/s12014-019-9252-2 PMID: 31341447</u>
- Martinez BA, Hoyle RG, Yeudall S, et al., Innate immune signaling in Drosophila shifts anabolic lipid metabolism from triglyceride storage to phospholipid synthesis to support immune function. PLoS Genet 2020; 16(11): e1009192. https://doi.org/10.1371/journal.pgen.1009192 PMID: 33227003
- Bilgin S, Aktas G, Atak T, et al., Triglyceride to high density lipoprotein cholesterol ratio is elevated in patients with complicated type 2 diabetes mellitus. Acta Facultatis Medicae Naissensis 2022; 39(1): 66–73. https://doi.org/10.5937/afmnai39-33239
- Hadaegh F, Khalili D, Ghasemi A, et al., Triglyceride/HDL-cholesterol ratio is an independent predictor for coronary heart disease in a population of Iranian men. Nutr Metab Cardiovasc Dis 2009; 19(6): 401– 8. https://doi.org/10.1016/j.numecd.2008.09.003 PMID: 19091534
- Kurtkulagi O, Bilgin S, Kahveci GB, et al., Could triglyceride to high density lipoprotein-cholesterol ratio predict hepatosteatosis? Experimental Biomedical Research 2021; 4(3): p224–229. https://doi.org/10. 30714/j-ebr.2021370081
- Nedeva C, Menassa J and Puthalakath H, Sepsis: Inflammation Is a Necessary Evil. Front Cell Dev Biol 2019; 7(108. https://doi.org/10.3389/fcell.2019.00108 PMID: 31281814
- 45. Liu Q, Li L, Gao L, et al., Combination of bis (alpha-furancarboxylato) oxovanadium (IV) and metformin improves hepatic steatosis through down-regulating inflammatory pathways in high-fat diet-induced obese C57BL/6J mice. Basic Clin Pharmacol Toxicol 2021; 128(6): 747–757. <u>https://doi.org/10.1111/</u> bcpt.13573 PMID: 33599105
- 46. Rosenson RS, Shott S and Tangney CC, Hypertriglyceridemia is associated with an elevated blood viscosity Rosenson: triglycerides and blood viscosity. Atherosclerosis 2002; 161(2): 433–9. https://doi.org/10.1016/s0021-9150(01)00656-6 PMID: 11888528
- Chen Y, Yu CY and Deng WM, The role of pro-inflammatory cytokines in lipid metabolism of metabolic diseases. Int Rev Immunol 2019; 38(6): 249–266. <u>https://doi.org/10.1080/08830185.2019.1645138</u> PMID: 31353985