Research Article

U-Shape Relationship between Plasma Leucine Level and Mortality in the Intensive Care Unit

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Patients in the intensive care unit (ICU) are at high risk of mortality which is not well predicted. Previous studies noted that leucine has prognostic value in a variety of diseases. This study investigated whether leucine concentration was a useful biomarker of metabolic and nutritional status and 6-month mortality in ICU. We recruited 454 subjects admitted to ICU (348 and 106 in the initiation and validation cohorts, respectively) with an acute physiology and chronic health evaluation (APACHE II) score ≥ 15. We measured plasma leucine concentrations, traditional biomarkers, and calculated APACHE II and sequential organ failure assessment (SOFA) scores. Leucine levels were weakly correlated with albumin, prealbumin, and transferrin levels ($r = 0.30$, $0.12$, and $0.15$, $p = 0.001$, $0.029$, and $0.007$, respectively). During follow-up, 116 (33.3%) patients died. Compared to patients with leucine levels between 109 and 174 μM, patients with leucine > 174 μM or < 109 μM had a lower cumulative survival rate. Death was also associated with age, higher APACHE II and SOFA scores, C-reactive protein, and longer stays in the ICU, but with lower albumin, prealbumin, and transferrin. Patients with leucine > 174 μM had higher alanine aminotransferase levels, but no significant differences in other variables; patients with leucine levels < 109 μM had higher APACHE II and SOFA scores, higher incidence of using inotropic agents, longer ICU and hospital stays, but lower albumin and transferrin levels. Multivariable analysis demonstrated that leucine > 174 μM was an independent predictor of mortality, especially early mortality. However, among patients who stayed in ICU longer than two weeks, leucine < 109 μM was an independent predictor of mortality. In addition, leucine < 109 μM was associated with worse ventilator weaning profiles. These findings were similar in the validation cohort. Our study demonstrated a U-shape relationship between leucine levels and mortality rate in ICU.

1. Introduction

Patients admitted to the intensive care unit (ICU) are at high risk of morbidity and mortality [1–3]. Catabolic status and/ or malnutrition are associated with poor outcomes. In 1950, Cuthbertson et al. defined distinct catabolic pathophysiology and nutritional deficiency at different phases throughout the trajectory of critical illness [4]. To mitigate these devastating
metabolic disturbances, assessment of patients’ catabolic and nutritional status is required for interventions such as protein supplementation or anabolism-stimulating medications. A simple and timely surrogate biomarker covering both catabolic and nutritional status and prognostic value is thus a critical unmet need, especially for patients in a contemporary ICU with overlapping and cyclical phases of metabolism that were not adequately addressed by Cuthbertson et al. long ago, whose work was based on bed-ridden patients with femoral fractures [5].

Nutritional parameters, including albumin, prealbumin, and transferrin, provide prognostic value in critical patients [6–9]. However, all these nutritional biomarkers are proteins with a complex process of synthesis that depends on the amount of amino acids present; moreover, a variety of factors may interfere with the level of these biomarkers, such as inflammation, caloric intake, and iron storage. Leucine, an essential amino acid, is an element of albumin and prealbumin [10, 11]. Previously, we demonstrated that blood leucine concentrations had a clear correlation with the total amount of essential amino acids, indicating that leucine can potentially represent the quantity of fundamental elements for protein synthesis [12]. Moreover, leucine has been proven to have prognostic value in patients with advanced cardiovascular diseases [12]. Thus, further investigation is merited to understand the value of measuring blood leucine levels in patients with critical illness.

As shown in our previous study of patients with heart failure, the I-shape relationship between leucine level and disease outcomes suggests that leucine might indicate catabolic and malnutrition status associated with outcomes [12–15]. However, how to use leucine levels in a clinical setting remains to be elucidated, especially in critical illness. In this study, we hypothesized that plasma leucine concentration could be used as a biomarker of metabolic and nutritional status and mortality in patients cared in the ICU. The aims of this study were (1) to explore the relationship between blood concentrations of leucine and traditional nutrition parameters, (2) to estimate whether leucine provides prognostic value compared to the other traditional parameters, and (3) to understand how leucine level predicts mortality in patients admitted to the ICU for future clinical application.

2. Material and Methods

2.1. Patient Enrollment. From April 2017 to February 2020, for the initiation cohort, patients were consecutively enrolled at the contemporary medical and cardiac ICU based on these inclusion criteria: (1) they were at critical status with an APACHE II score ≥ 15, (2) they needed to stay in the ICU > 48 hours, and (3) they were older than 20 years old. The exclusion criteria included patients with comorbid disorders other than the main cause for admission that might compromise their survival within three months, such as terminal stage cancer. All patients provided informed consent. The study was designed and carried out in accordance with the principles of the Declaration of Helsinki and with approval from the Ethics Review Board of Chang Gung Memorial Hospital.

The validation phase set, another independent population including 106 patients, was recruited from March 2020 to September 2020 (Figure 1).

2.2. Scoring Systems. Disease severity was assessed by SOFA scores [16] on the first day of admission to the intensive care unit. APACHE II scores [2] were calculated both on the first day and 2 weeks later after admission to the ICU.

2.3. Blood Sampling and Examination. Fasting blood samples were collected in EDTA-containing tubes in the early morning, the day after obtaining informed consent. Plasma leucine was measured by ultraperformance liquid chromatography (UPLC) workflow. Measurement of other parameters, including prealbumin, transferrin, albumin, estimated glomerular filtration rate (eGFR), C-reactive protein, and hemoglobin, was performed in the central laboratory.

2.4. Leucine Measurement. Plasma leucine concentration was quantified by UPLC. Plasma samples (100 μL) were precipitated with sulfosalicylic acid (10%). After protein precipitation and centrifugation, derivatization was conducted by AQC in acetonitrile. Amino acids were analyzed using the ACQUITY UPLC System, consisting of a Sample Manager, a Binary Solvent Manager, and a Tunable UV detector. EmpowerTM 2 Software was used to control the system and collect data. Separations were conducted on a 2.1 × 100 mm ACQUITY BEH C18 column at a flow rate of 0.70 mL/min. The average intra-assay coefficient of variation was 4.5%. The total coefficient of variation was 4.1%, and the detection limit was 0.9 μM. The linear range was 25–500 μM.

2.5. Measurement of Respiratory Profile. Respiratory profile was estimated within 2 days after admitted to the ICU if the patients were intubated. The content of the profile included respiratory rate, tidal volume, minute ventilation, and maximal inspiratory pressure which were measured with a Mechanics monitor (Respira-dyne II, Sherwood/Covidien, Boulder, Colorado). The rapid shallow breathing index was calculated as respiratory rate divided by tidal volume and recorded as an index for the pulmonary mechanics and weaning procedure.

2.6. Follow-Up Program. Follow-up data were prospectively obtained from hospital records, personal communication with the patients’ physicians, telephone interviews with patients, and patients’ regular visits to staff physician outpatient clinics. Patients were followed until death or a maximum of 6 months. The primary endpoint was death from all causes. In addition, we analyzed the length of stay in ICU and hospital.

2.7. Statistical Analyses. Results are expressed as the mean ± SD for variables with normal distribution, as the median (interquartile range (IQR)) for variables with skewed distribution, and as the number (percentage) for categorical variables. We compared data using t-test, one-way ANOVA (with post hoc analysis by Bonferroni), Chi-square (multiple
3. Results

3.1. Baseline Characteristics and Laboratory Data. In the initiation cohort, we enrolled 348 patients. These patients were admitted to the ICU for the following conditions: (1) 157 patients (45.1%) had coronary artery disease, myocardial infarction, or cardiovascular diseases; (2) 92 patients (26.4%) had infection; (3) 44 patients (12.6%) had chronic obstructive pulmonary disease; (4) 15 patients (4.3%) had gastrointestinal bleeding; (5) 7 patients (2.0%) had kidney disease; (6) 5 patients (1.4%) were admitted for sudden collapse; (7) 2 patients (0.6%) for stroke, and 26 patients (7.4%) for other conditions. The average age was 71.4 years, and the average APACHE II and SOFA scores were 20.2 and 7.07, respectively (Table 1). Generally, the enrolled patients had multiple comorbidities, high C-reactive protein levels, but low hemoglobin, cholesterol, triglyceride, albumin, eGFR, prealbumin, and transferrin levels. The validation cohort included 106 patients. The demographic data are shown in the supplemental Table S1.

3.2. Relationship between Leucine Levels and Mortality. During the 6-month follow-up period, 116 (33.3%) patients died. In 61 (52.6%) patients, death occurred due to infection, 23 (19.8%) died of critical cardiovascular conditions, 19 (16.4%) due to both conditions combined, 7 (6.0%) due to pulmonary diseases, and 5 (4.3%) due to other reasons. Leucine levels were weakly correlated with albumin, prealbumin, and transferrin levels ($r = 0.30, 0.12, and 0.15$, with $p = 0.001, 0.029$, and $0.007$, respectively). Association between leucine level and mortality risk was shown in the additive Cox regression models (Figure 2(a)). Although lower leucine levels were significantly associated with higher mortality rates, higher leucine levels also correlated with higher mortality rates, represented by a U-shaped curve.

Patients were separated into three groups based on the ROC curve analyses and Youden’s index consisting of patients with baseline leucine levels $< 109 \mu M$, between 109 and $174 \mu M$, and $> 174 \mu M$ (Supplemental Figure S1). In Figure 2(b), the Kaplan-Meier curves demonstrate that patients with leucine levels $> 174 \mu M$ or $< 109 \mu M$ had a lower accumulative survival rate, compared to those with leucine levels of 109 to 174 $\mu M$ (log rank = 9.11 and 6.22, $p = 0.003$ and 0.013, respectively). The mortality rate was 45.5%, 40.0%, and 25.9% in patients with leucine levels $> 174 \mu M$ and $< 109 \mu M$ and between 109 and 174 $\mu M$, respectively. The validation cohort confirmed these findings (Kaplan-Meier curves are shown in supplemental Figure S2).

Patients with leucine levels $< 109 \mu M$ had higher APACHE II and SOFA scores compared to patients with leucine levels between 109 and $174 \mu M$, higher incidence of using inotropic agents, and longer stays in the ICU and hospital, but lower hemoglobin, cholesterol, triglyceride, albumin, and transferrin levels (Table 2). Patients with leucine levels $> 174 \mu M$ had higher levels of alanine aminotransferase, but no significant differences in other parameters.
In addition, compared to patients with leucine levels < 109 μM, patients with leucine levels > 174 μM had similar prealbumin levels, but higher cholesterol, triglyceride, alanine aminotransferase, albumin, and transferrin levels. The demographic and laboratory data in patients with different leucine concentrations in the validation cohort are shown in supplemental Table S2.

3.3. Factors Associated with Mortality. Mortality was associated with age, higher APACHE II and SOFA scores, higher incidence of using inotropic agents, higher C-reactive protein levels, and longer length of stay in the ICU; but also with lower incidence of chronic obstructive pulmonary disease, lower cholesterol, albumin, prealbumin, and transferrin levels (Table 1). In addition, leucine levels between 109 and 174 μM were related to lower mortality rate in both the initiation and validation cohorts (Table 1 and Supplemental Table S1).

3.4. High Leucine Predicted Early Mortality. Of the 25 patients who had leucine levels > 174 μM and died in 6 months, 20 (80%) died within the 2 weeks following enrollment. COX univariate analysis showed that mortality within 6 months was associated with leucine > 174 μM or <109 μM, age, APACHE II and SOFA scores, chronic obstructive pulmonary disease, and levels of C-reactive protein, cholesterol, albumin, prealbumin, and transferrin (Table 3). Next, we conducted multivariable analysis by adjusting for significant demographic variables and by adjusting for significant demographic and laboratory variables according to univariate analysis. Either after adjusting for age and chronic obstructive pulmonary disease or after adjusting for age,
chronic obstructive pulmonary disease, C-reactive protein, cholesterol, albumin, pre-albumin, and transferrin, leucine > 174 μM remained an independent predictor of mortality within 6 months. Furthermore, even after adjusting for all significant variables in the univariate analysis additionally including APACHE II and SOFA scores, leucine > 174 μM was still demonstrated to be an independent predictor (hazard ratio = 1.97, 95% CI = 1.16 – 3.33, \( p = 0.012 \)).

3.5. Low Leucine Predicted Mortality in Patients Who Survived Longer than Two Weeks. Multivariable analysis showed that leucine < 109 μM was an independent predictor of mortality within 6 months after adjusting for significant demographic variables, including age and chronic obstructive pulmonary disease, according to univariate analysis (Table 3). However, when adjusting for significant demographic and laboratory variables, including age, chronic obstructive pulmonary disease, and nutritional biomarkers, the prognostic value of leucine < 109 μM became insignificant.

In Figure 2(b), comparing patients with leucine levels between 109 and 174 μM and those with leucine levels < 109 μM, the Kaplan-Meier curves show that the mortality rate did not differ in 2 weeks (log rank = 0.46, \( p = 0.49 \)) but began to diverge after 2 weeks. Thus, we performed the secondary analysis. Of patients who survived longer than 2 weeks, those with leucine levels < 109 μM at baseline had a significantly lower survival rate compared to those with leucine levels between 109 and 174 μM (log rank = 7.22, \( p = 0.007 \)) (Figure 3(b)). Survival rates were similar between patients with leucine between 109 and 174 μM and those with leucine > 174 μM. Compared to patients with baseline leucine levels between 109 and 174 μM, patients with a leucine level < 109 μM at baseline had higher respiratory rate (23.2 ± 7.24 breaths/min vs. 19.9 ± 6.99 breaths/min, \( p = 0.003 \)), rapid shallow breathing index (69.0 (45, 92) vs. 53.5 (33, 75.3), \( p = 0.003 \)), but lower tidal volume (358 ± 158 ml vs. 424 ± 210 ml, \( p = 0.027 \)) in the respiratory weaning profile at baseline and longer length of ICU and hospital stay (Table 2).

Based on these results, we further investigated the prognostic value of leucine concentrations in patients who survived longer than two weeks and were still in the ICU (\( n = 103 \)). Leucine levels were measured again at two weeks after enrollment. In Figure 3(b), Kaplan-Meier curves show that leucine < 109 μM at 2 weeks was significantly associated with a lower survival rate, compared to leucine ≥ 109 μM (log rank = 6.28, \( p = 0.012 \)). COX univariate analysis showed that leucine < 109 μM significantly predicted mortality (hazard ratio = 2.41, 95% CI = 1.18 – 4.89, \( p = 0.016 \)). The prognostic value of leucine < 109 μM was independent of age and APACHE II at two weeks (hazard ratio = 2.34, 95% CI = 1.15 – 4.76, \( p = 0.020 \)).

4. Discussion

Our study reveals that the amino acid, leucine, can be used as a biomarker of disease outcomes. Interestingly, we noted a U-shaped relationship between leucine levels and mortality rates; both lower and higher leucine levels predicted higher mortality rates. In addition, low leucine levels became the main prognostic factor when patients stayed in the ICU longer than two weeks. The validation cohort confirmed these findings.

Albumin and prealbumin are two widely used biomarkers of nutrition. Low levels of albumin and prealbumin have been shown to indicate poor outcomes in critical patients [6–8]. However, these biomarkers represent the capacity for protein synthesis associated with nutritional status, and thus, their clinical interpretation is limited by the speed of synthesis and their relatively long half-life (12 and 3 days for albumin and prealbumin, respectively). Leucine is an essential component for albumin and prealbumin synthesis [10, 11], and its blood concentration is assumed to represent the amount of elementary resources for protein synthesis. Intriguingly, our study demonstrated that
The concept of ebb and flow phases is based on Cuthbertson and colleagues’ observations on the catabolic state in response to serious injury and other pathophysiological conditions [4, 5]. Most situations we encounter in the contemporary medical ICU involve the acute and adaptive responses during the flow phase. However, these phases and responses often happen repeatedly and may overlap, making them difficult to distinguish by clinical judgement. It is also impractical to identify different metabolic states by performing an instant immune panel or estimating protein synthesis and breakdown using the indicator amino acid oxidation method.

Our data provide a solution for these complex and interdependent metabolic states. Based on existing literature, both protein breakdown and synthesis increase remarkably in response to the tremendous stress at the acute catabolic phase [3, 5]. However, protein synthesis often cannot catch up with the speed of protein breakdown, since a substantial portion of the amino acids released from skeletal muscle, such as branched chain amino acids, is used to synthesize acute phase protein and produce local energy by oxidation, or goes unused due to anabolic resistance [3, 17]. Plasma
leucine concentrations may represent the net balance of these interacting mechanisms.

In the adaptive phase, substantial amounts of amino acids are needed for tissue repair and protein synthesis. Low leucine concentrations may indicate deficient amino acids and may be associated with impaired wound healing, tissue repair, muscle synthesis, and immune function [18]. Previously, in patients with COPD, Labaki et al. found that lower concentrations of serum leucine were independently associated with one-year incidence of respiratory exacerbation [19]. Labaki’s observations are further supported by our data in the secondary analysis that, in patients who survived longer than two weeks after admitted in the ICU, lower leucine levels predicted a poorer weaning profile, longer length of ICU and hospital stays, and an even higher mortality rate compared to those with relatively normal plasma leucine concentrations. In our study, leucine < 109 μM appeared to be associated with chronic malnutrition, since the values of all nutritional parameters were the lowest of the three subgroups. This notion was further supported by our finding that the prognostic value of leucine < 109 μM disappeared after adjusting for albumin or other nutritional biomarkers. Whether aggressive supplementation of protein can improve the outcomes of these patients is an urgent matter in need of clarification.

| Leucine            | Univariate | Multivariable | Multivariable |
|--------------------|------------|---------------|---------------|
|                    | HR (95% CI)| p value       | HR (95% CI)   | p value       | HR (95% CI)   | p value       |
| Leucine 109-174 μM (reference) |            |               |               |               |               |               |
| Leucine > 174 μM   | 2.16 (1.30-3.59) | 0.003         | 2.03 (1.22-3.37) | 0.006         | 2.06 (1.22-3.47) | 0.007         |
| Leucine < 109 μM   | 1.66 (1.10-2.51) | 0.015         | 1.52 (1.01-2.31) | 0.043         | 0.95 (0.60-1.50) | 0.828         |
| Age (years)        | 1.02 (1.00-1.03) | 0.046         |               |               |               |               |
| APACHE II score    | 1.12 (1.09-1.16) | <0.001        | 1.08 (1.06-1.11) | <0.001        | 1.09 (1.05-1.12) | <0.001        |
| SOFA score         | 1.23 (1.17-1.28) | <0.001        | 1.14 (1.07-1.21) | <0.001        | 1.18 (1.11-1.24) | <0.001        |
| COPD               | 0.27 (0.09-0.86) | 0.027         |               |               |               |               |
| C-reactive protein (log) | 2.21 (1.62-3.01) | <0.001        | 2.26 (1.65-3.10) | <0.001        |               |               |
| Cholesterol (mg/dL) | 0.99 (0.98-0.99) | <0.001        | 0.98 (0.98-0.99) | <0.001        |               |               |
| Albumin (g/dL)     | 0.392 (0.29-0.53) | <0.001        | 0.41 (0.29-0.54) | <0.001        |               |               |
| Prealbumin (mg/dL) | 0.92 (0.89-0.95) | <0.001        | 0.92 (0.89-0.95) | <0.001        |               |               |
| Transferrin (mg/dL)| 0.99 (0.98-0.99) | <0.001        | 0.98 (0.98-0.99) | <0.001        |               |               |

APACHE: acute physiology and chronic health evaluation; CI: confidence interval; COPD: chronic obstructive pulmonary disease; HR: hazard ratio; SOFA: sequential organ failure assessment. *, adjusting for age and COPD. †, adjusting for age, COPD, C-reactive protein, cholesterol, albumin, prealbumin, and transferrin.

**Table 3: COX univariate and multivariable analysis of factors for predicting mortality in the initiation cohort.**

**Figure 3:** Prognostic value of plasma leucine concentration in patients who survived longer than two weeks. (a) The Kaplan-Meier curves for three different groups (for all-cause death). Leucine levels were measured at baseline. (b) The Kaplan-Meier curves for patients with leucine < 109 μM versus ≥109 μM (for all-cause death). Leucine levels were measured at two weeks after admission to the intensive care unit in patients who remained in the intensive care unit.
In critical illness, a subgroup of patients develops anabolic resistance [20, 21], which is probably associated with insulin resistance, interstitial edema, and dysfunction of amino acid transporters on the cell membrane. In anabolic resistance, increased feeding of calories and protein does not give rise to increased protein synthesis. High plasma leucine levels may represent this metabolic phenotype, consistent with our previous report that high leucine levels in critically ill patients correlated with poor mitochondrial function leading to impaired beta-oxidation and increased unmetabolized acylcarnitines in the circulation [12]. High leucine levels may thus indicate a state of metabolic storm in response to critical stress. These conclusions are further supported by our current findings that leucine concentration > 174 μM was associated with a higher incidence of mortality which mostly occurred during the acute phase within 2 weeks after being hospitalized to the ICU. Patients with leucine >174 μM could not be easily distinguished from other patients by clinical characteristics, such as APACHE II and SOFA scores or laboratory parameters. Interestingly, although the leucine concentration was high in the circulation, the prealbumin level in these patients was low, which could mislead clinicians to attempt protein supplementation. Anabolic resistance might be mechanistically related to impaired prealbumin production. Although inadequate protein supplementation is associated with a poor prognosis [22], overfeeding is also harmful [23–26], and leucine concentration could help clinicians avoid making mistakes. These findings strongly imply that measuring leucine concentration in critically ill patients could differentiate the causes of malnutrition identified by traditional biomarkers.

4.1. Limitations. There are a few limitations in this study. First, although we and others have previously demonstrated significant prognostic value of leucine compared to other essential amino acids in critically ill patients [3, 12, 15, 18], it is possible that using a multimino acid panel can provide additional value. However, doing so would negatively impact cost-benefit considerations. Second, for estimation of the patients survived longer than 2 weeks, since this was a secondary analysis, other nutritional biomarkers were not measured in the current study, such as albumin, prealbumin, and transferrin. To further confirm the independent prognostic value of low leucine level in this population, a prospective design should be designed in the future. Third, whether serial measurement of patients’ leucine levels throughout the critical and subacute phases is helpful has not been proven and warrants further interventional investigation. Finally, we did not explore the prognostic value by combining leucine and other independent predictors of adverse outcomes in ICU. There is a strong need for improving outcome prediction by contemporary techniques such as machine learning models [27].

5. Conclusions

Our study demonstrated a U-shaped relationship between plasma leucine levels and mortality rate in patients admitted to the ICU due to critical status. Our findings provide a way to scientifically delineate the complex phases of catabolism and malnutrition initially defined by Cuthbertson et al. in 1950. Monitoring leucine levels could provide a meaningful reference to guide optimal nutrition support for patients hospitalized in the ICU.

Data Availability

The authors declare that the data underlying this article will be shared on reasonable request to the corresponding author.

Conflicts of Interest

The authors declare that there is no conflict of interest regarding the publication of this paper.

Authors’ Contributions

Chao-Hung Wang, Wei-Siang Chen, Min-Hui Liu, and Huei-Jen Wang contributed to the conceptualization. Chien-Ming Chu, Huang-Ping Wu, and Mei-Ying Wang contributed to the validation. Min-Hui Liu, Wei-Siang Chen, and Chao-Hung Wang contributed to the formal analysis. Yi-Tsen Lin, Chien-Ming Chu, Huang-Ping Wu, Kuo-Chin Kao, Chung-Yu Liang, Wen-Hsin Chen, and Shu-Chiu Lee contributed to the investigation. Chien-Ming Chu, Huang-Ping Wu, Kuo-Chin Kao, Chung-Yu Liang, Wen-Hsin Chen, and Shu-Chiu Lee contributed to the data curation. Chao-Hung Wang and Min-Hui Liu contributed to the writing—review and editing. Chao-Hung Wang contributed to the funding acquisition. All authors contributed to the conduction of the study and reviewed and approved the final version of the manuscript.

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Supplementary Materials

Supplementary Table S1: demographic and laboratory data in the validation cohort. Supplementary Table S2: comparisons of demographic and laboratory data in patients with different leucine concentrations in the validation cohort. Supplementary Figure S1: an inverse-sigmoidal receiver operating characteristic curve for the leucine level to predict mortality. Supplemental Figure S2: the Kaplan-Meier curves for patients with leucine < 109 μM, 109-174 μM, >174 μM (for all-cause death). Leucine levels were measured at baseline. (Supplementary Materials)
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