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Lymphocyte Cell Ratios and Mortality among Incident Hemodialysis Patients

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Abstract

Background—Neutrophil-to-lymphocyte ratio (NLR) and platelet-to-lymphocyte ratio (PLR) have previously been suggested as oncologic prognostication markers. These are associated with malnutrition and inflammation, and hence, may provide benefit in predicting mortality among hemodialysis patients.

Methods—Among 108,548 incident hemodialysis patients in a large U.S. dialysis organization (2007–2011), we compared the mortality predictability of NLR and PLR with baseline and timevarying covariate Cox models using the receiver operating characteristic curve (AUROC), net reclassification index (NRI), and adjusted \mathbb{R}^2 .

Results—During the median follow-up period of 1.4 years, 28,618 patients died. Median (interquartile range) NLR and PLR at baseline were 3.64 (2.68–5.00) and 179 (136–248), respectively. NLR was associated with higher mortality, which appeared stronger in the timevarying vs. baseline model. PLR exhibited a J-shaped association with mortality in both models. NLR provided better mortality prediction in addition to demographics, comorbidities, and serum albumin; AUROC and NRI for 1-year mortality $(95\%$ CI) were 0.010 $(0.009-0.012)$ and 6.4% $(5.5\% - 7.3\%)$, respectively. Additionally, adjusted R² (95%CI) for the Cox model increased from 0.269 (0.262–0.276) to 0.283 (0.276–0.290) in the non-time-varying model and from 0.467

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 $(0.461 - 0.472)$ to 0.505 $(0.500 - 0.512)$ in the time-varying model. There was little to no benefit of adding PLR to predict mortality.

Conclusions—High NLR in incident hemodialysis patients predicted mortality, especially in the short-term period. NLR, but not PLR, provided additional benefit in predicting mortality along with demographics, comorbidities, and serum albumin, and should be included in prognostication approaches.

Keywords

hemodialysis; chronic hemodialysis; end stage kidney disease; neutrophil; platelet; lymphocyte; mortality; nutrition; inflammation

Introduction

Low-grade systemic and persistent inflammation may play an important role in the development and progression of various chronic conditions and diseases such as obesity, diabetes, atherosclerotic cardiovascular disease, cancer, chronic obstructive lung disease, and chronic kidney disease (CKD) .^{1–3} Among patients with end-stage renal disease (ESRD), chronic inflammation is highly prevalent and associated with adverse clinical outcomes including cardiovascular disease and all-cause death. $4-8$ Inflammation in this context is attributed to various underlying factors including the uremic milieu, infections, decreased clearance of proinflammatory cytokines, volume overload, endotoxinemia, oxidative stress, carbonyl stress, and dialysis-related factors. $9-12$ A recent study found increased arterial wall inflammation even in non-dialysis patients with $CKD₁¹³$ which may explain the cardiovascular risk associated with CKD.14 Inflammation can also interact with malnutrition and lead to wasting and derangements in protein–energy nutritional status, i.e., proteinenergy wasting (PEW), resulting in the excessively high mortality in the dialysis population. 10,12 C-reactive protein, interleukin-6, and tumor necrosis factor-α are among representative inflammatory markers, but their measurements are costly and/or inaccessible in current clinical practice.

The neutrophil-to-lymphocyte ratio (NLR) and platelet-to-lymphocyte ratio (PLR), readily available biomarkers calculated from complete blood count, are closely associated with inflammation and were originally suggested as oncologic prognostic factors.^{15,16} Subsequently higher values in NLR and PLR have been linked to greater inflammatory status and worse prognosis among patients with various disease conditions including cardiovascular disease, 17 chronic obstructive pulmonary disease, 18 liver cirrhosis, 19 and CKD.20 Furthermore, recent studies have indicated that NLR and PLR were associated with inflammation and may predict mortality among hemodialysis patients as well, 2^{1-26} but their small study sample sizes make it difficult to draw a definite conclusion regarding their detailed associations with mortality and comparative usefulness in mortality-predictability when compared to other clinically accessible markers related to CKD. To address this knowledge gap, we studied a 5-year large national cohort of incident hemodialysis patients in the United States and investigated the relationship of NLR and PLR with all-cause mortality.

METHODS

This study was approved by the Institutional Review Committees of the Los Angeles Biomedical Research Institute at Harbor-UCLA, University of California Irvine Medical Center, and the University of Washington, with the exemption of obtaining written consent given the large sample size, anonymity of the patients studied, and nonintrusive nature of the research.

Patients

We retrospectively extracted, refined, and examined data from all ESRD patients aged 18 years who initiated dialysis from January 1, 2007 to December 31, 2011and who received dialysis treatment in one of the facilities operated by a large dialysis organization in the United States.27 Patients were followed up from the of first date dialysis until death, kidney transplantation, loss to follow-up, or the date of final follow-up assessment for all patients (i.e., December 31, 2011). The follow-up time was divided into patient-quarters (i.e., 91-day periods from the first date of dialysis).

Out of the 156,229 patients who were initially treated with conventional hemodialysis, we excluded 30,493 patients who did not receive dialysis treatment for
 60 consecutive days, 13,719 patients who were ever treated with other modalities (e.g., less frequent hemodialysis, frequent hemodialysis, nocturnal hemodialysis, home hemodialysis, or peritoneal dialysis), 2,593 and 579 patients with missing data or NLR values >14 (i.e., >99.5 percentile), respectively on complete blood count markers. We also excluded 297 patients without data on serum albumin, phosphorus, and alkaline phosphatase (ALP) at baseline (Figure 1).

Demographic, clinical, and laboratory measures

Information on death, race/ethnicity, primary insurance, access type, the presence of comorbidities, and laboratory variables were obtained from the administrative electronic database of the dialysis provider. To minimize measurement variability, all repeated measures for each patient were averaged within each patient-quarter and then used in all analyses. The averaged values during the first patient-quarter (the first 91-days of dialysis) served as baseline data.

Blood samples were drawn using uniform techniques in all dialysis clinics and were transported to the central laboratory in Deland, Florida, typically within 24 hours. All laboratory values were measured by automated and standardized methods. Most laboratory values were measured monthly with the exception of serum ferritin and intact parathyroid hormone (PTH) that were measured at least quarterly. Hemoglobin was measured at least monthly in essentially all patients and weekly to bi-weekly in most patients. NLR was calculated using percent neutrophil divided by percent lymphocyte whereas PLR was calculated using platelet count divided by lymphocyte count. Single-pool Kt/V (spKt/V) was calculated using urea kinetic modeling equations.²⁸

Statistics

Differences in baseline characteristics between two groups were compared by standardized differences, of which 0.8, 0.5, and 0.2 in absolute value were considered large, medium, and small differences, respectively, due to the relatively large sample size of this study.^{29,30} NLR and PLR, the primary exposures of this study, were categorized into 6 groups (<2.0, 2.0– $\langle 3.0, 3.0 - \langle 4.0, 4.0 - \langle 5.0, 5.0 - \langle 6.0, \text{ and } 6.0; \text{ and } \langle 100, 100 - \langle 150, 150 - \langle 200, 200 - \langle 250, \rangle \rangle$ 250–<300, and ≥300, respectively). Their associations with each patient characteristic were evaluated using a non-parametric trend test. Logistic regression was used to identify the predictors of high NLR or PLR using arbitrary cut offs (i.e., 4.0 and 200, respectively). Cox regression models with baseline (non-time-varying) or time-varying covariates were used to estimate associations with all-cause mortality. Models were examined with three levels of adjustment as follows:

- **I.** Unadjusted model that included NLR or PLR;
- **II.** Case-mix adjusted models that included the above plus age, sex, race and ethnicity, primary insurance, 12 comorbidities as in Table 1, vascular access type, body mass index (BMI), and spKt/V; and
- **III.** MICS adjusted models which included all of the covariates in the case-mix model plus nine laboratory variables related to the malnutrition-inflammationcachexia complex syndrome (MICS) [i.e., white blood cell (WBC) count, hemoglobin, serum albumin, creatinine, albumin-corrected calcium, phosphorus, iron saturation, total bicarbonate, ALP, total iron binding capacity (TIBC), and natural log-transformed intact PTH and ferritin].

Time-varying covariates included vascular access type, BMI, spKt/V, and laboratory variables. In order to compare mortality-predictabilities of NLR and PLR with those of albumin and ALP, the mortality risk was estimated according to the deciles of each variable. The reason for selecting serum albumin and ALP for the comparison is that both indices have been consistently and linearly associated with mortality among hemodialysis patients. 31,32 Serum albumin is a representative MICS marker while ALP is not in the realm of MICS but is an abnormality observed in ESRD as a part of CKD-MBD. The area under the receiver operating characteristic curve (AUROC) and net reclassification index (NRI) were examined for 1-year mortality using logistic regression with different sets of variables at baseline. Additionally, explained variation (i.e., adjusted \mathbb{R}^2 , a transformation of Royston and Sauerbrei's D measure of discrimination of a survival model) 33 was evaluated in both time-varying and non-time-varying Cox models to compare the overall performance measure across models. The frequency of missing data was low [<1% for BMI and most laboratory tests, except for ferritin (1.2%), nPCR (1.5%), and creatinine (4.6%)], and mean imputation was used in all regression analyses. All analyses were conducted using STATA MP version 13.1 (StataCorp, College Station, TX).

RESULTS

Baseline demographic, clinical and laboratory characteristics

Baseline characteristics were compared between included and excluded patients and we found that the 108,548 included patients were less likely to be Caucasian; and were more likely to have a CV catheter as their initial vascular access. Compared to the 47,681 excluded patients who lacked NLR or PLR data, included patients had higher hemoglobin, lower white blood cell count, lower neutrophil, higher lymphocyte, higher albumin, higher creatinine, higher phosphorus, higher intact PTH, higher TIBC, lower ferritin, lower bicarbonate, and higher prevalence of hypertension, diabetes, and congestive heart failure at baseline (Supplemental Table 1). The mean \pm SD age of the final study cohort was 63 \pm 15 years of whom 56% were male, 47% were Caucasian, 15% were Hispanic, and 59% were diabetic (Table 1). The median (interquartile range [IQR]) of NLR and PLR were 3.64 (2.68, 5.00) and 179 (136, 248), respectively. As NLR values increased, the absolute lymphocyte count decreased while the absolute counts of white blood cell and neutrophil increased. The absolute platelet count values were not different between the high and low NLR groups. Patients with higher NLR values were more likely to be older and Caucasian, but less likely to be African American. Likewise, As PLR values increased, the absolute lymphocyte count decreased while the absolute neutrophil and platelets counts increased. There were no meaningful differences in demographics, comorbid conditions, and the absolute white blood cell count between the high and low NLR groups.

Biochemical Associations of NLR and PLR

After adjustment for case-mix variables, baseline laboratory variables associated with higher NLR (vs. below the median) included lower levels of hemoglobin, albumin, creatinine, albumin-corrected calcium, phosphorus, intact PTH, TIBC, and iron saturation and higher levels of WBC, ALP, ferritin, and bicarbonate (Supplemental Table 4). PLR exhibited similar association patterns for those laboratory variables except for WBC, where lower levels of WBC were associated with higher PLR. Calcium and phosphorus levels were not associated with PLR.

Associations of NLR and PLR with all-cause mortality

Out of 108,548 patients, a total of 28,618 patients died during the median follow-up period of 1.4 years (IQR, 0.6 to 2.5 years) with an incidence rate of 15.6 (95%CI, 15.4 to 15.7) per 100 patient-years. Baseline characteristics of the study cohort stratified by 6 NLR groups and 6 PLR groups are presented in Supplemental Table 2 and Supplemental Table 3, respectively. Unadjusted non-time-varying (baseline) Cox models demonstrated that higher NLR levels were linearly associated with all-cause mortality risk. Those who had greater NLR experienced incrementally higher mortality, which was robust against adjustments for case-mix and MICS characteristics (P_{trend} <0.001 for all; Figure 1A and Supplemental Table 5). Similarly, the time-varying covariate model demonstrated incremental mortality risk across greater NLR irrespective of adjustment model, which appeared stronger than those observed in the baseline model (P_{trend} <0.001 for all; Figure 1B). When compared to NLR of 2.0 to <3.0, a hazard ratio (HR) of $\,$ 6.0 NLR was 1.94 (95%CI, 1.87 to 2.02) and 4.54

(95%CI, 4.34 to 4.75) in the baseline vs. the time-varying covariate Cox model with adjustment for case-mix variables, respectively.

Baseline PLR exhibited a J-shaped association with all-cause mortality in all adjustment models; the lowest risk was observed in the second lowest group (i.e., PLR 100 to <150), and higher levels were associated with all-cause mortality risk (Figure 1C and Supplemental Table 5). As with NLR, the time-varying covariate model exhibited a consistent yet stronger association between PLR and all-cause mortality, when compared to the baseline model (Figure 1D). Using PLR of 100 to 150 as reference, a case-mix adjusted hazard ratio (HR) of ≥300 PLR was 1.43 (95%CI, 1.37 to 1.48) and 2.95 (95%CI, 2.82 to 3.09) in the baseline vs. time-varying covariate Cox model, respectively.

Mortality Predictability Comparisons of Selected Laboratory Variables

In order to evaluate the clinical usefulness in predicting mortality, we compared the hazard ratios (Figure 2) of deciles among serum albumin, serum ALP, NLR, and PLR, and then compared AUROC (Table 2), NRI (Table 2 and Supplemental Table 6–12), and adjusted \mathbb{R}^2 (Table 3) among lymphocyte, neutrophil, and platelet counts in addition to those variables above.

Serum albumin appeared to be the strongest predictor of all-cause mortality in both the nontime-varying and time-varying Cox models with adjustment for the case-mix variables (Figure 2), with the greatest AUROC and the highest NRI for 1-year mortality (Table 2). Adjusted \mathbb{R}^2 of serum albumin was also highest in both the non-time-varying and timevarying Cox model (Table 3). Baseline serum ALP and neutrophil, when compared to lymphocyte and platelet, showed higher AUROC, NRI, and adjusted R^2 , and neutrophil showed a higher adjusted R^2 than serum ALP in the time-varying model. However, NLR and the combination of neutrophil and lymphocyte demonstrated equivalent mortality predictability, which was better than serum ALP especially in the time-varying model. Neither PLR nor the combination of platelet and lymphocyte provided clinically relevant gain in these indices. When baseline NLR was added into the model including the case-mix variables and serum albumin, AUROC and NRI was 0.010 (95%CI, 0.009 to 0.012) and 6.4% (95%CI, 5.5% to 7.3%), respectively, which was greater than those observed with serum ALP and PLR. It also increased adjusted \mathbb{R}^2 from 0.269 (0.263, 0.277) to 0.286 (0.279, 0.293) in the non-time-varying model and from 0.484 (0.479, 0.491) to 0.510 (0.504, 0.516) in the time-varying model.

Sensitivity analysis including patients with baseline NLR values >14 (i.e., >99.5 percentile) showed consistent results with a slight attenuation in the mortality predictability of NLR (data not shown)

DISCUSSION

We demonstrated that high NLR was a strong predictor of all-cause death among hemodialysis patients. Patients with malnutrition and inflammation were likely to have high NLR. Baseline NLR was incrementally associated with higher mortality and provided modest benefit to mortality prediction in addition to the case-mix variables and serum

albumin. The NLR-mortality association was more pronounced in the time-varying model, suggesting a stronger association of NLR with short-term mortality than with long-term mortality. The mortality predictability of NLR appeared to be similar to or better than that of serum ALP or the combination of neutrophil and lymphocyte. Although PLR was associated with MICS variables, it exhibited a J-shaped association with mortality, and there was little, if any, improvement in mortality-predictability by adding either PLR or the combination of platelet and lymphocyte to the model including the case-mix variables and serum albumin.

Previous studies of dialysis patients demonstrated that both NLR and PLR had moderate correlations with direct inflammatory markers such as C-reactive protein, interleukin-6, and tumor necrosis factor- $\alpha^{21,26,34}$ and that higher NLR and PLR values were associated with greater mortality.23,25 Some studies also suggested that PLR may have a stronger correlation with inflammation and better mortality-predictability than NLR .^{25,26} However, these studies had small sample sizes (<200 patients), and hence, were susceptible to over-adjustment in multivariable analyses. Additionally, it remained difficult to judge the benefit of adding NLR or PLR as a new predictor to an existing model because these studies did not evaluate performance measures such as AUROC, NRI, and adjusted \mathbb{R}^2 . The large sample size of our cohort allowed us to rigorously adjust for clinically relevant factors and to evaluate several model performances, all of which supported the better mortality-predictability of NLR than PLR.

Serum albumin has good correlations with NLR^{34} because inflammatory status leads to high NLR by increasing neutrophils and decreasing lymphocytes, and because it also results in low serum albumin levels by downregulating albumin gene transcription.^{35–38} Additionally, both serum albumin and lymphocyte count have been used as indices of nutritional status. Thus, serum albumin and NLR share common pathophysiological rationale for survival prediction. However, a previous study highlighted the lack of correlation between total lymphocyte count and nutritional status in the elderly³⁹ while recent randomized clinical trials demonstrated that protein supplementation increased serum albumin levels, which may explain the stronger association of serum albumin with survival than NLR in our study. Nevertheless, NLR modestly improved mortality prediction in addition to the case-mix variables and serum albumin. This additional benefit of adding NLR might be related to its shorter time to response to inflammation than serum albumin; neutrophils are rapidly released from the bone marrow into the bloodstream with a short half-life (\sim 6–8 hours)⁴⁰ whereas serum albumin decreases through reduced synthesis rate and has a longer half-life (~19–21 days).

We acknowledge several limitations in this study. First, due to the observational nature of our study, we cannot prove causality between NLR or PLR and death, nor exclude the possibility of residual confounding. Second, the administrative database from a large dialysis organization does not enable us to individually identify specific condition(s) for high NLR or PLR, such as infection, malnutrition, central venous catheter, stress, metabolic syndrome, and atherosclerosis. Third, there are currently no established reference ranges for NLR and PLR in the general population or the dialysis population although a previous study of healthy individuals reported lower NLR values $(2.5th-97.5th$ percentiles, $0.8-3.5)⁴¹$ than those in our study. However, our survival analyses showed a linear association between

higher NLR and higher mortality without clear thresholds indicating its optimal ranges. Finally, we included exclusively incident hemodialysis patients, and it remains unclear whether our findings can be extrapolated to prevalent hemodialysis patients or peritoneal dialysis patients. Nevertheless, our study is the largest study of hemodialysis patients evaluating model performance measures of NLR and PLR, and both non-time-varying and time-varying models yielded consistent results.

In conclusion, a high NLR in incident HD patients predicted mortality, especially in the short-term period, and provided modest benefit to predict mortality in addition to demographics, comorbidities, and serum albumin. PLR demonstrated a weaker association with mortality, and there was small or virtually no gain in mortality-predictability. NLR, together with serum albumin, may be useful as a surrogate of nutritional and inflammatory status in both clinical and epidemiological research settings given that it is readily available without additional cost.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

Statement of Competing Financial Interests:

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This study is based on data provided by DaVita Clinical Research. The manuscript has been reviewed and approved by DaVita. The interpretation and conclusions are those of the authors and do not represent the views of DaVita.

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Figure 1.

Mortality risk associated with neutrophil-to-lymphocyte ratio (NLR) and platelet-tolymphocyte ratio (PLR) after initiating dialysis among 108,548 incident hemodialysis patients (2007–2011) in the (A and C) baseline and (B and D) time-varying Cox models with three-level hierarchical adjustments, respectively.

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Figure 2.

Case-mix adjusted all-cause mortality hazard ratios of deciles of (A) baseline and (B) timevarying serum albumin, serum alkaline phosphatase (ALP), neutrophil-to-lymphocyte ratio (NLR), and platelet-to-lymphocyte ratio (PLR) among 108,548 patients. Only the deciles of serum albumin were ordered in reverse (i.e., the highest on the left and the lowest on the right) for the comparison with the other indices.

Table 1

Baseline Demographic and Clinical Characteristics of 108,548 incident hemodialysis patients. Baseline Demographic and Clinical Characteristics of 108,548 incident hemodialysis patients.

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and 0.2 in absolute value were considered large, medium, and small differences, respectively. Abbreviations: COPD, chronic obstructive pulmonary disease; Human Immunodeficiency Virus; AIDS, acquired immune deficiency syndrome; WBC, white blood cell; PTH, parathyroid hormone; TIBC, total iron binding capacity. Conversion factors for units: albumin and hemoglobin in g/dL to g/L, 10;

creatinine in mg/dL to mmol/L, 88.4; calcium in mg/dL to mmol/L, 0.2495; phosphorus in mg/dL to mmol/L, 0.3229. No conversion is necessary for ferritin in ng/mL and mg/L.

acquired immune deficiency syndrome; WBC, white blood cell; PTH, parathyroid hormone; TIBC, total iron binding capacity. Conversion factors for units: albumin and hemoglobin in g/dL to g/L, 10;
creatinine in mg/dL to mmol/ and 0.2 in absolute value were considered large, medium, and small differences, respectively. Abbreviations: COPD, chronic obstructive pulmonary disease; Human Immunodeficiency Virus; AIDS,

Table 2

Area under receiver operating characteristic curve (AUROC) and net reclassification index (NRI) for 1-year mortality by adding baseline levels of serum albumin, serum alkaline phosphatase (ALP), lymphocyte, neutrophil, platelet, neutrophil-to-lymphocyte ratio (NLR), and platelet-to-lymphocyte ratio (PLR) in addition to the case-mix variables.

Table 3

Adjusted \mathbb{R}^2 in the non-time-varying and time-varying Cox model by adding serum albumin, serum alkaline phosphatase (ALP), lymphocyte, neutrophil, platelet, neutrophil-to-lymphocyte ratio (NLR), and platelet-tolymphocyte ratio (PLR) in addition to the case-mix variables.

