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# Fibroblast Growth Factor 23 Associates with Death in Critically Ill Patients

David E. Leaf, Edward D. Siew, Michele F. Eisenga, Karandeep Singh, Finian R. Mc Causland, Anand Srivastava, T. Alp Ikizler, Lorraine B. Ware, Adit A. Ginde, John A. Kellum, Paul M. Palevsky , Myles Wolf, and Sushrut S. Waikar

## Abstract

**Background and objectives** Dysregulated mineral metabolism is a common and potentially maladaptive feature of critical illness, especially in patients with AKI, but its association with death has not been comprehensively investigated. We sought to determine whether elevated plasma levels of the osteocyte-derived, vitamin D–regulating hormone, fibroblast growth factor 23 (FGF23), are prospectively associated with death in critically ill patients with AKI requiring RRT, and in a general cohort of critically ill patients with and without AKI.

**Design, setting, participants, & measurements** We measured plasma FGF23 and other mineral metabolite levels in two cohorts of critically ill patients ( $n=1527$ ). We included 817 patients with AKI requiring RRT who enrolled in the ARF Trial Network (ATN) study, and 710 patients with and without AKI who enrolled in the Validating Acute Lung Injury biomarkers for Diagnosis (VALID) study. We hypothesized that higher FGF23 levels at enrollment are independently associated with higher 60-day mortality.

**Results** In the ATN study, patients in the highest compared with lowest quartiles of C-terminal (cFGF23) and intact FGF23 (iFGF23) had 3.84 (95% confidence interval, 2.31 to 6.41) and 2.08 (95% confidence interval, 1.03 to 4.21) fold higher odds of death, respectively, after adjustment for demographics, comorbidities, and severity of illness. In contrast, plasma/serum levels of parathyroid hormone, vitamin D metabolites, calcium, and phosphate were not associated with 60-day mortality. In the VALID study, patients in the highest compared with lowest quartiles of cFGF23 and iFGF23 had 3.52 (95% confidence interval, 1.96 to 6.33) and 1.93 (95% confidence interval, 1.12 to 3.33) fold higher adjusted odds of death.

**Conclusions** Higher FGF23 levels are independently associated with greater mortality in critically ill patients.

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## Introduction

Dysregulated mineral metabolism, including hypocalcemia, hyperparathyroidism, and low circulating levels of 25-hydroxyvitamin D (25D) and 1,25-dihydroxyvitamin D (1,25D), is an important feature of critical illness (1,2), and is especially pronounced among patients with AKI (3). These mineral metabolite abnormalities may be causally associated with adverse outcomes in critical illness through a variety of pathways, including nonclassic effects of vitamin D metabolites on immunity and inflammation (4). However, the mechanisms responsible for dysregulated mineral metabolism in critical illness are poorly understood.

Fibroblast growth factor 23 (FGF23), an osteocyte-derived hormone initially discovered for its pathologic role in rare syndromes of urinary phosphate wasting (5), plays a critical role in regulating vitamin D and phosphate homeostasis. FGF23 inhibits the conversion of 25D to its biologically active form, 1,25D, while also stimulating the catabolism of both 25D and 1,25D (6).

Elevated FGF23 levels are now recognized as a key feature of dysregulated mineral metabolism in patients

with CKD (7,8), and are one of the most robust predictors of cardiovascular disease (9) and death (10,11) in this patient population.

In contrast to CKD, few studies investigated FGF23 in critical illness. In a pilot study, we recently reported that higher plasma ( $n=113$ ) and urinary ( $n=243$ ) C-terminal FGF23 (cFGF23) levels are associated with an increased risk of the composite end point of AKI or death in critically ill patients (12). However, the study was focused on incident, rather than established, AKI, and thus included few patients with severe AKI at the time of FGF23 measurement; assessed only cFGF23, but not intact FGF23 (iFGF23) levels; and, most importantly, was underpowered to assess mortality.

We therefore analyzed plasma FGF23 levels in patients who enrolled in two large studies: the ARF Trial Network (ATN) study, which included patients with AKI requiring RRT, and the Validating Acute Lung Injury biomarkers for Diagnosis (VALID) study, which included patients with and without AKI. We hypothesized that in both cohorts, higher plasma cFGF23 and iFGF23 levels are independently and strongly associated with greater 60-day mortality.

Due to the number of contributing authors, the affiliations are provided in the Supplemental Material.

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## Materials and Methods

### Study Population

**ATN Study.** The ATN study, described elsewhere in detail (13), enrolled 1124 critically ill patients with AKI requiring RRT from 27 major United States medical centers into a randomized trial of intensive versus less intensive RRT. Subject enrollment began in November of 2003 and was completed July of 2007. At the time of randomization, EDTA-plasma samples were archived in a biorepository and stored at  $-80^{\circ}\text{C}$ . All patients who provided samples for the biorepository ( $n=817$ ) were included in this study. These patients had similar baseline characteristics and outcomes compared with the overall cohort ( $n=1124$ ) (Supplemental Table 1).

**VALID Study.** The VALID study, described elsewhere in detail (14), is a single-center, prospective cohort study of patients admitted to intensive care units (ICUs) at Vanderbilt University Medical Center (VUMC; Nashville, TN). The VALID study began enrollment in 2006 and is ongoing. Eligible adult patients ( $\geq 18$  years of age) were enrolled within 24 hours of ICU admission, and EDTA-plasma samples were obtained and stored at  $-80^{\circ}\text{C}$ . Patients were excluded from the parent study if they had chronic lung disease requiring oxygen supplementation, pulmonary fibrosis, a history of cardiac arrest, transfer orders written or anticipated within 4 hours, or were admitted for an uncomplicated overdose. Additional exclusion criteria for this study were a history of kidney transplant ( $n=31$ ), baseline eGFR  $< 15$  ml/min per  $1.73\text{ m}^2$  ( $n=15$ ), or receiving dialysis ( $n=150$ ) at the time of enrollment.

**Informed Consent.** In the ATN study, all patients or their surrogates provided written informed consent, and the study was approved by the Human Rights Committee at the West Haven Veterans Affairs Cooperative Studies Program Coordinating Center and by the institutional review boards at each of the participating sites. In the VALID study, the VUMC institutional review board granted a waiver of consent if patients were unable to participate in the consent process and no surrogate was available, due to the minimal risk of the study.

### Study Design

We assessed the association between plasma FGF23 levels and death in the ATN ( $n=817$ ) and VALID ( $n=710$ ) studies using a prospective cohort and case-cohort study design, respectively. Cases were defined as patients who died within 60 days after enrollment. The case-cohort in VALID was selected from a source population of 1732 patients with samples available (Supplemental Figure 1). An advantage of the case-cohort design is that oversampling of cases is an efficient approach that provides nearly the same magnitude of statistical power compared with the full cohort (15).

### Exposures, Measurements, and Outcomes

The primary exposure was plasma FGF23 level, measured on enrollment in both studies. The primary outcome was 60-day mortality. In the VALID study, we also assessed 1-year mortality as a secondary outcome.

**FGF23 Measurements.** The immunometric assay for cFGF23 detects both the intact hormone and C-terminal cleavage products, whereas the assay for iFGF23 recognizes the intact, biologically active hormone only (16). Few studies

of patients with critical illness have compared the results from both assays. Therefore, we measured both cFGF23 and iFGF23 levels using second generation ELISA kits (Immutopics, San Clemente, CA) in batched assays by investigators who were unaware of the outcomes. Specifically, in the ATN study we measured cFGF23 levels in all 817 participants, and we measured iFGF23 levels in a random subcohort ( $n=400$ ). Characteristics among participants from the random subcohort ( $n=400$ ) were similar to the full cohort ( $n=817$ ) (data not shown). In the VALID study we measured both cFGF23 and iFGF23 in all 710 participants. Interassay coefficients of variation for cFGF23 and iFGF23, determined from blinded replicate samples, were  $< 10\%$ . We report additional performance characteristics for the cFGF23 and iFGF23 assays elsewhere (17).

**Additional Measurements.** In the ATN study we also measured plasma levels of vitamin D metabolites, including 25D, 1,25D, and 24,25-dihydroxyvitamin  $\text{D}_3$ , (24,25D<sub>3</sub>), and intact parathyroid hormone (PTH) in the same random subcohort ( $n=400$ ) in which we measured iFGF23. Vitamin D metabolites were measured using immunoaffinity enrichment and liquid chromatography–tandem mass spectrometry (18). Serum levels of creatinine, calcium, phosphate, and albumin were measured for clinical purposes by the hospital laboratory in both the ATN and VALID studies. Plasma IL-6 levels were measured in the ATN study only (details reported elsewhere) (19). A summary of the measurements performed in both cohorts, along with assay performance characteristics, is provided in Supplemental Table 2.

### Statistical Analyses

We performed the statistical analyses with SAS Version 9.4 (Cary, NC) and R Version 3.2.3 (Vienna, Austria). Baseline characteristics were compared between patients alive or dead at day 60 using Wilcoxon Rank Sum and chi-squared tests for continuous and categorical variables, respectively. In the ATN study, 1.7% of comorbidity and severity of illness data were missing and were imputed using regression switching with predictive mean matching (20). Five datasets were multiply-imputed, and results were pooled using Rubin's rules (21). In VALID,  $< 0.1\%$  of data were missing and were not imputed. In both cohorts, the primary outcome variable (60-day mortality) was available in all patients.

We assessed the associations between FGF23 and other analyte levels using Spearman correlation coefficients and restricted cubic splines. We used univariable (model 1) and multivariable (models 2 and 3) logistic regression to assess the association between FGF23 and other mineral metabolite levels and 60-day mortality. We assessed FGF23 levels in quartiles to test for nonlinear associations and by natural log–transformation and normalization to 1 SD to allow comparison across biomarkers.

We adjusted for covariates in multivariable models (models 2 and 3) on the basis of univariable associations with 60-day mortality and biologic plausibility. In both cohorts, model 2 was adjusted for age, sex, race, baseline eGFR, diabetes, cancer, congestive heart failure, and chronic liver disease. In the ATN study, model 3 was further adjusted for the following severity of illness covariates assessed on enrollment: ICU type, mechanical ventilation, Acute Physiology and Chronic Health Evaluation (APACHE) II score (22),

RRT before randomization, treatment group, type of RRT, oliguria, sepsis, hypotension, white blood cell count, hemoglobin, serum albumin, serum creatinine, and plasma IL-6 levels. In the VALID study, the severity of illness covariates included in model 3 were ICU type, mechanical ventilation, APACHE II score, sepsis, hypotension, white blood cell count, hemoglobin, serum creatinine, and AKI presence and severity, defined according to serum creatinine–based consensus criteria (23). The covariates included in model 3 differed slightly in the ATN study compared with the VALID study due to differences in the availability of covariates in the two cohorts as well as differences in univariable associations with mortality.

We depicted the association between FGF23 levels and 1-year survival in the VALID study using Kaplan–Meier curves, with differences assessed by the log-rank test. Multivariable adjusted associations were assessed using Cox proportional hazards models. Using Martingale residual plots and Kolmogorov-type supremum tests, we confirmed that the proportional hazards assumption was not violated. We used inverse probability weighting, using the Barlow weighting method (24), to account for the case-cohort design in the VALID study (15). All comparisons are two-tailed, with  $P < 0.05$  considered significant.

## Results

### Patient Characteristics

Demographic and clinical characteristics from participants in the ATN and VALID studies, dichotomized according to survival status at day 60, are shown in Table 1. Demographic and clinical characteristics according to quartiles of cFGF23 and iFGF23 are shown in Supplemental Table 3.

### Laboratory Values

Median cFGF23 levels were 3608 (interquartile range [IQR], 1079–12,983 RU/ml) and 603 (IQR, 483–1183 RU/ml), and median iFGF23 levels were 88 (IQR, 30–239 pg/ml) and 44 (24–87 pg/ml), among participants in the ATN and VALID studies, respectively. Compared with informal reference ranges (cFGF23, 21–82 RU/ml; iFGF23, 3.8–18.7 pg/ml) reported in elderly adults with normal kidney function (25), we observed markedly elevated cFGF23 levels and modestly elevated iFGF23 levels. Additional laboratory values are shown in Table 2.

### Association between FGF23 and Other Mineral Metabolite Levels

The associations between FGF23 and vitamin D metabolite levels are shown in Figure 1, A–F. The strongest association was between higher levels of iFGF23 and lower levels of 1,25D ( $r_s = -0.26$ ,  $P < 0.001$ ; Figure 1D), consistent with FGF23-mediated inhibition of 1- $\alpha$  hydroxylase. Additional correlations are shown in Figure 1G.

### FGF23 Levels and Mortality in the ATN Study

Higher quartiles of cFGF23 were associated with a monotonic increase in the risk of death in both unadjusted and adjusted analyses (adjusted odds ratio [OR], 3.83, for patients in the highest compared with lowest quartiles; 95% confidence interval (CI), 2.27 to 6.47; Figure 2A). Higher

quartiles of iFGF23, by contrast, were associated with a nonlinear increase in the risk of death in unadjusted models, and were associated with a semilinear increase in the risk of death after adjustment for comorbidities and severity of illness (Figure 2A).

Next, we assessed the magnitude of association between each mineral metabolite, expressed as a natural log-transformed continuous variable and normalized to 1 SD, with risk of death. In fully adjusted models, higher levels of cFGF23 and iFGF23 were each associated with an increased risk of death, whereas calcium, phosphate, parathyroid hormone, 25D, 1,25D, and 24,25D<sub>3</sub> were not (Figure 2B).

**Exploratory Analyses.** In exploratory analyses, the associations between cFGF23, iFGF23, and death remained significant after further adjustment for 25D, 1,25D, and serum phosphate levels (Supplemental Table 4). cFGF23 remained associated with death after adjustment for iFGF23 (Supplemental Table 5). The relationship between cFGF23 and death was consistent across a number of subgroups, including age (above versus below 65 years), sex, ICU type, sepsis, shock, APACHE II score (above versus below 25), RRT before randomization, and 25D and 1,25D levels above versus below the median (Supplemental Figure 2).

### FGF23 Levels and Mortality in VALID

In the VALID study, higher quartiles of cFGF23 and iFGF23 were each associated with an increase in the risk of death in both unadjusted and adjusted analyses (adjusted OR, 4.08, for patients in the highest compared with lowest quartiles of cFGF23; 95% CI, 2.19 to 7.58; adjusted OR, 1.95, for patients in the highest compared with lowest quartiles of iFGF23; 95% CI, 1.13 to 3.39; Figure 3A). Higher cFGF23 and iFGF23 levels, assessed as natural log-transformed continuous variables, were also associated with increased 60-day mortality in both unadjusted and fully adjusted models (Figure 3B).

Higher quartiles of cFGF23 and iFGF23 were also associated with increased 1-year mortality in both univariable (Figure 4, A and B) and multivariable analyses (Figure 4C). In exploratory analyses, the associations between FGF23 levels and 1-year mortality were qualitatively similar when stratified by the presence or absence of AKI on enrollment (Supplemental Figure 3).

## Discussion

In this study, which includes two prospective cohorts, we found that plasma FGF23 levels are elevated in critically ill patients, and that higher FGF23 levels are associated with higher mortality. This association was present for both cFGF23 and iFGF23, and was independent of other known risk factors. Additionally, we found that higher iFGF23 associates with lower 1,25D levels, consistent with FGF23-mediated inhibition of 25D-to-1,25D conversion. However, iFGF23 remained significantly associated with death independently of 25D and 1,25D. Cumulatively, these data identify elevated FGF23 as an important feature of dysregulated mineral metabolism in critical illness.

Our data are consistent with and expand on prior studies demonstrating an association between higher FGF23 levels and adverse outcomes in patients with CKD (9–11) and,

**Table 1. Enrollment characteristics**

| Characteristic                                 | ATN Study     |               |                         | VALID Study   |               |                            |
|--|---------------|---------------|-------------------------|---------------|---------------|----------------------------|
|  | All (n=817)   | Alive (n=402) | Dead (n=415)            | All (n=710)   | Alive (n=449) | Dead (n=261)               |
| <b>Demographics</b>                            |               |               |                         |               |               |                            |
| Age, yr, median (IQR)                          | 62 (51–72)    | 59 (48–68)    | 64 (54–76) <sup>a</sup> | 57 (45–68)    | 54 (41–65)    | 61 (51–73) <sup>a</sup>    |
| Male sex, no. (%)                              | 568 (70)      | 275 (68)      | 293 (71)                | 439 (62)      | 271 (60)      | 168 (64)                   |
| White, no. (%)                                 | 628 (77)      | 301 (75)      | 327 (79)                | 625 (88)      | 381 (85)      | 244 (93) <sup>a</sup>      |
| <b>Comorbidities, no. (%)</b>                  |               |               |                         |               |               |                            |
| Diabetes mellitus                              | 199 (25)      | 108 (28)      | 91 (23)                 | 166 (23)      | 103 (23)      | 63 (24)                    |
| Congestive heart failure                       | 198 (25)      | 93 (24)       | 105 (26)                | 90 (13)       | 47 (10)       | 43 (16)                    |
| Chronic liver disease                          | 95 (12)       | 38 (10)       | 57 (14)                 | 77 (11)       | 46 (10)       | 31 (12)                    |
| Chronic lung disease/COPD <sup>b</sup>         | 103 (13)      | 43 (11)       | 60 (15)                 | 107 (15)      | 68 (15)       | 39 (15)                    |
| CKD <sup>c</sup>                               | 293 (38)      | 137 (37)      | 156 (39)                | 183 (26)      | 83 (18)       | 100 (38) <sup>a</sup>      |
| Malignancy                                     | 158 (19)      | 69 (17)       | 89 (21)                 | 109 (15)      | 58 (13)       | 51 (20) <sup>d</sup>       |
| <b>Baseline kidney function</b>                |               |               |                         |               |               |                            |
| Creatinine, mg/dl, median (IQR) <sup>e</sup>   | 1.1 (0.9–1.4) | 1.1 (0.8–1.4) | 1.1 (0.9–1.4)           | 0.9 (0.7–1.2) | 0.8 (0.6–1.0) | 1.0 (0.8–1.4) <sup>a</sup> |
| eGFR, median (IQR) <sup>f</sup>                | 71 (50–94)    | 73 (52–96)    | 70 (49–89)              | 90 (60–112)   | 98 (70–117)   | 72 (50–98) <sup>a</sup>    |
| Medical ICU, no. (%)                           | 361 (44)      | 176 (44)      | 185 (45)                | 329 (46)      | 181 (40)      | 148 (57) <sup>a</sup>      |
| <b>Severity of illness</b>                     |               |               |                         |               |               |                            |
| Sepsis, no. (%)                                | 527 (65)      | 253 (63)      | 274 (66)                | 297 (42)      | 162 (36)      | 135 (52) <sup>a</sup>      |
| APACHE II score, median (IQR) <sup>g</sup>     | 26 (21–31)    | 23 (19–29)    | 28 (24–33) <sup>a</sup> | 26 (20–31)    | 23 (19–29)    | 29 (23–34) <sup>a</sup>    |
| Mechanical ventilation, no. (%)                | 661 (81)      | 293 (73)      | 368 (89) <sup>a</sup>   | 512 (72)      | 330 (74)      | 182 (70)                   |
| Hypotension, no. (%) <sup>h</sup>              | 515 (63)      | 217 (54)      | 298 (72) <sup>a</sup>   | 450 (63)      | 276 (61)      | 174 (67)                   |
| <b>AKI and RRT data on enrollment, no. (%)</b> |               |               |                         |               |               |                            |
| AKI <sup>i</sup>                               | 817 (100)     | 402 (100)     | 415 (100)               | 348 (49)      | 228 (51)      | 120 (46)                   |
| AKI severity                                   |               |               |                         |               |               |                            |
| Stage 1  | 0 (0)         | 0 (0)         | 0 (0)                   | 238 (34)      | 155 (35)      | 83 (32)                    |
| Stage 2  | 0 (0)         | 0 (0)         | 0 (0)                   | 81 (11)       | 54 (12)       | 27 (10)                    |
| Stage 3  | 817 (100)     | 402 (100)     | 415 (100)               | 29 (4)        | 19 (4)        | 10 (4)                     |
| Oliguria <sup>j</sup>                          | 654 (80)      | 300 (75)      | 354 (85) <sup>a</sup>   |               |               |                            |
| RRT before enrollment                          | 561 (69)      | 275 (68)      | 286 (69)                |               |               |                            |
| <b>RRT data in the ATN study, no. (%)</b>      |               |               |                         |               |               |                            |
| Treatment group                                |               |               |                         |               |               |                            |
| Intensive RRT                                  | 403 (49)      | 191 (48)      | 212 (51)                |               |               |                            |
| Less intensive RRT                             | 414 (51)      | 211 (52)      | 203 (49)                |               |               |                            |
| Type of RRT                                    |               |               |                         |               |               |                            |
| CRRT   | 557 (68)      | 237 (58)      | 320 (77) <sup>a</sup>   |               |               |                            |
| Intermittent RRT                               | 266 (32)      | 170 (42)      | 96 (23) <sup>a</sup>    |               |               |                            |

Percentages are on the basis of the number of patients without missing data. ATN, ARF Trial Network; VALID, Validating Acute Lung Injury biomarkers for Diagnosis; IQR, interquartile range; COPD, chronic obstructive pulmonary disease; ICU, intensive care unit; APACHE II, Acute Physiology and Chronic Health Evaluation II; CRRT, continuous renal replacement therapy.

<sup>a</sup> $P < 0.001$ , for comparison between patients alive versus dead at day 60.

<sup>b</sup>Defined in the ATN study as chronic hypoxemia, hypercapnea, pulmonary hypertension, or ventilator dependence; defined in VALID as COPD.

<sup>c</sup>Defined as baseline eGFR  $< 60$  ml/min per  $1.73$  m<sup>2</sup>.

<sup>d</sup> $P < 0.05$ , for comparison between patients alive versus dead at day 60.

<sup>e</sup>Defined in the ATN study as the premorbid serum creatinine (SCr) at the time of screening or, if unavailable, the lowest SCr within 4 d before screening; defined in VALID as the lowest SCr within 365 d before hospitalization. If no SCr values were available before hospitalization, the lowest SCr during hospitalization (excluding values obtained during RRT) was used as the baseline (44).

<sup>f</sup>Reported in ml/min per  $1.73$  m<sup>2</sup> and determined using the Chronic Kidney Disease Epidemiology Collaboration Equation (45).

<sup>g</sup>An ICU severity of illness scoring system ranging from 0 to 71, with higher scores indicating more severe disease.

<sup>h</sup>Defined in the ATN study as requirement for vasopressor support for  $> 1$  h; defined in VALID as systolic BP  $\leq 90$  mm Hg or need for vasopressor support.

<sup>i</sup>All patients in the ATN study had AKI requiring RRT on enrollment; in VALID, AKI on enrollment was defined as an increase in SCr  $\geq 0.3$  mg/dl within 48 h or  $\geq 50\%$  within 7 d before enrollment (23).

<sup>j</sup>Defined in the ATN study as an average urine output  $< 20$  ml/h for  $> 24$  h before enrollment.

to a lesser extent, critical illness and AKI (3,12,26). Prior studies on FGF23 in critical illness were limited by small sample sizes, resulting in lack of power to assess hard outcomes such as death. Additionally, prior studies included few patients with severe AKI at the time of FGF23 measurement, and assessed only cFGF23, but not iFGF23

levels. In contrast, this study is the first to report an independent association between higher plasma FGF23 levels and increased all-cause mortality in a large cohort of critically ill patients. Additionally, we confirmed these findings in critically ill patients both with and without AKI, and with measurement of both cFGF23 and iFGF23 levels.

**Table 2. Enrollment laboratory values**

| Variable                                       | ATN Study         |                   |                   | P Value <sup>a</sup> | VALID Study       |                    |                   | P Value <sup>a</sup> |
|--|-------------------|-------------------|-------------------|----------------------|-------------------|--------------------|-------------------|----------------------|
|  | All (n=817)       | Alive (n=402)     | Dead (n=415)      |                      | All (n=710)       | Alive (n=449)      | Dead (n=261)      |                      |
| <b>Routine labs, median (IQR)</b>              |                   |                   |                   |                      |                   |                    |                   |                      |
| White cell count, per mm <sup>3</sup>          | 13<br>(9–19)      | 13<br>(9–18)      | 14<br>(8–20)      | 0.46                 | 14<br>(9–19)      | 13<br>(9–19)       | 14<br>(9–19)      | 0.51                 |
| Hemoglobin, g/dl                               | 9.8<br>(8.9–10.8) | 9.7<br>(8.8–10.6) | 9.9<br>(9.0–10.9) | 0.04                 | 9.8<br>(8.8–11.0) | 10.0<br>(8.9–11.1) | 9.7<br>(8.8–11.0) | 0.50                 |
| Creatinine, mg/dl                              | 3.9<br>(2.9–5.1)  | 4.3<br>(3.2–5.6)  | 3.6<br>(2.7–4.6)  | <0.001               | 1.1<br>(0.9–1.7)  | 1.0<br>(0.8–1.4)   | 1.4<br>(1.0–2.1)  | <0.001               |
| Albumin, g/dl                                  | 2.3<br>(1.9–2.8)  | 2.4<br>(1.9–3.0)  | 2.3<br>(1.8–2.8)  | 0.03                 |                   |                    |                   |                      |
| <b>cFGF23, RU/ml</b>                           |                   |                   |                   |                      |                   |                    |                   |                      |
| Median   | 3608              | 1918              | 5819              | <0.001               | 603               | 483                | 1183              | <0.001               |
| IQR  | 1079–12,983       | 740–7736          | 1656–18,731       |                      | 212–1882          | 181–1218           | 366–4344          |                      |
| <b>iFGF23, pg/ml</b>                           |                   |                   |                   |                      |                   |                    |                   |                      |
| Median   | 88                | 81                | 97                | 0.36                 | 44                | 38                 | 51                | <0.001               |
| IQR  | 30–239            | 24–266            | 40–224            |                      | 24–87             | 23–76              | 27–107            |                      |
| <b>Other markers, median (IQR)<sup>b</sup></b> |                   |                   |                   |                      |                   |                    |                   |                      |
| Calcium, mg/dl                                 | 7.7<br>(7.1–8.3)  | 7.8<br>(7.2–8.4)  | 7.7<br>(7.0–8.3)  | 0.33                 | 7.8<br>(7.4–8.3)  | 7.8<br>(7.5–8.3)   | 7.8<br>(7.4–8.3)  | 0.52                 |
| Phosphate, mg/dl                               | 5.2<br>(4.0–6.7)  | 5.1<br>(4.0–6.7)  | 5.2<br>(4.0–6.7)  | 0.79                 | 3.3<br>(2.5–4.3)  | 3.2<br>(2.5–4.2)   | 3.4<br>(2.4–4.7)  | 0.44                 |
| Parathyroid hormone, pg/ml                     | 304<br>(118–588)  | 306<br>(118–567)  | 301<br>(120–592)  | 0.77                 |                   |                    |                   |                      |
| 25-hydroxyvitamin D, ng/ml                     | 10<br>(6–14)      | 11<br>(7–14)      | 9<br>(6–13)       | 0.01                 |                   |                    |                   |                      |
| 1,25-dihydroxyvitamin D, pg/ml                 | 12<br>(8–18)      | 11<br>(7–20)      | 12<br>(8–18)      | 0.73                 |                   |                    |                   |                      |
| 24,25-dihydroxyvitamin D <sub>3</sub> , ng/ml  | 0.4<br>(0.3–0.7)  | 0.5<br>(0.3–0.7)  | 0.4<br>(0.3–0.7)  | 0.82                 |                   |                    |                   |                      |
| IL-6, pg/ml                                    | 166<br>(74–539)   | 125<br>(57–281)   | 230<br>(104–911)  | <0.001               |                   |                    |                   |                      |

ATN, ARF Trial Network; VALID, Validating Acute Lung Injury biomarkers for Diagnosis; IQR, interquartile range; cFGF23, c-terminal fibroblast growth factor 23; RU/ml, reference units per milliliter; iFGF23, intact fibroblast growth factor 23.

<sup>a</sup>Comparison between patients alive versus dead at day 60.

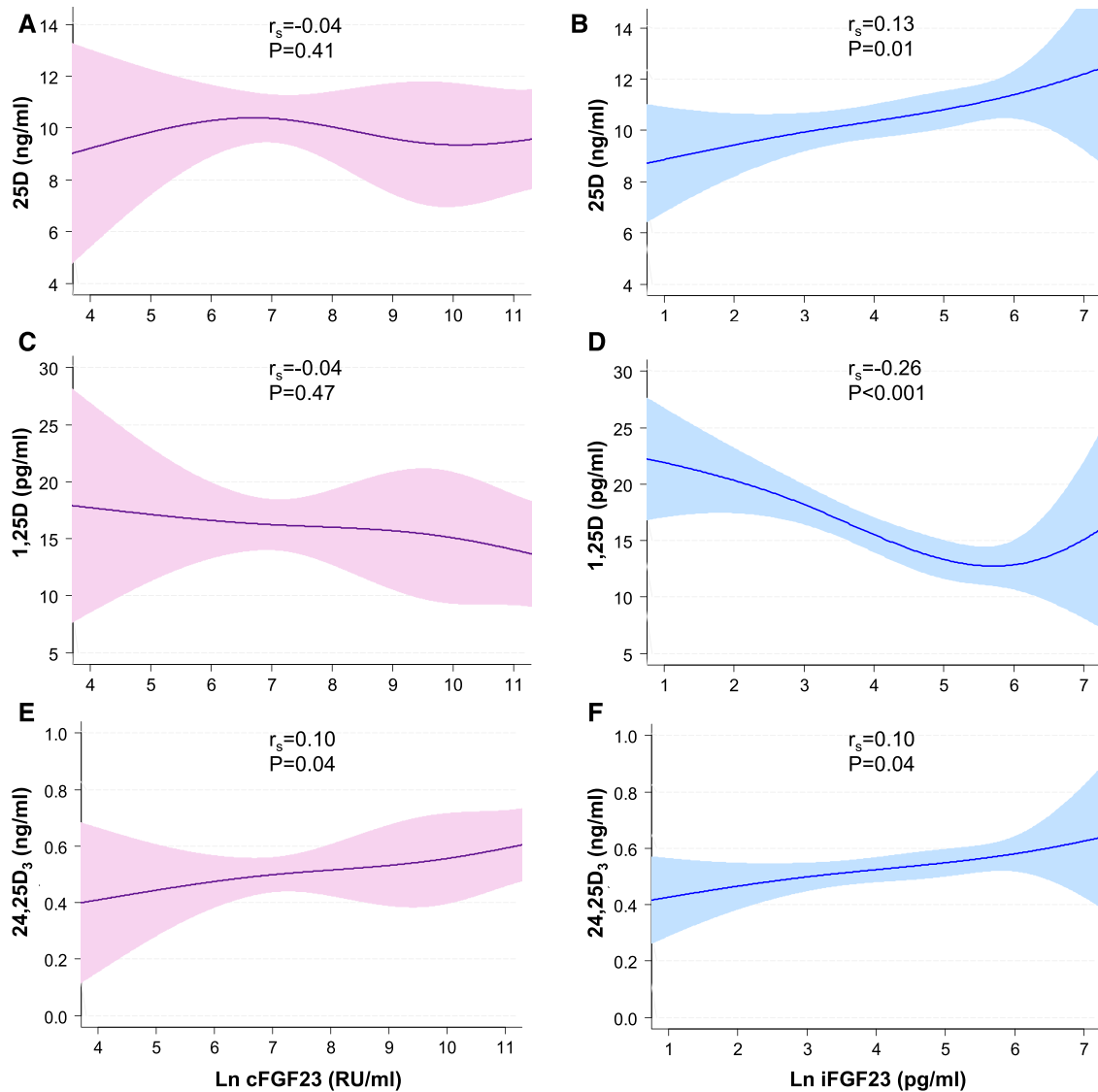
<sup>b</sup>Parathyroid hormone and vitamin D metabolite levels were available in a subcohort (n=400) of patients in the ATN study and were not available in VALID. IL-6 levels were available in all participants in the ATN study and were not available in VALID.

The downstream consequences of FGF23 over-activity have not been fully elucidated. Specifically, whether cFGF23 fragments exert biologic activity is unknown, because conflicting findings have been reported (27–29). In contrast, iFGF23 unequivocally exerts biologic activity through binding to FGF receptors expressed throughout the body. Recent studies found that in addition to classic effects on kidney phosphate handling and vitamin D metabolism, iFGF23 also has myriad “off-target” effects. These include toxic effects on the cardiovascular system through promotion of left ventricular hypertrophy and endothelial dysfunction (30,31), profibrotic effects on the kidneys (32), proinflammatory effects demonstrated by upregulation of hepatic IL-6 production (33), and impairment of immune function (34,35). Specifically, iFGF23’s effects on the immune system include reduced monocyte mRNA expression of the antimicrobial peptide, LL37 (34), and impairment of neutrophil activation and recruitment into inflamed tissues (35). Impaired immunity could increase susceptibility to infection, a leading cause of death in critical illness, and could provide a biologic mechanism for our observations. An alternative mechanism relates to FGF23-mediated upregulation of superoxide, inhibition of nitric oxide bioavailability, and impairment of endothelium-dependent vasorelaxation (31). These effects could increase

the risk of cardiovascular events, an important contributor to death in critical illness (36). Despite these negative consequences of FGF23 over-activity, neutralization of FGF23 with an mAb resulted in hyperphosphatemia, extensive arterial calcification, and increased mortality in a rat model of CKD (37).

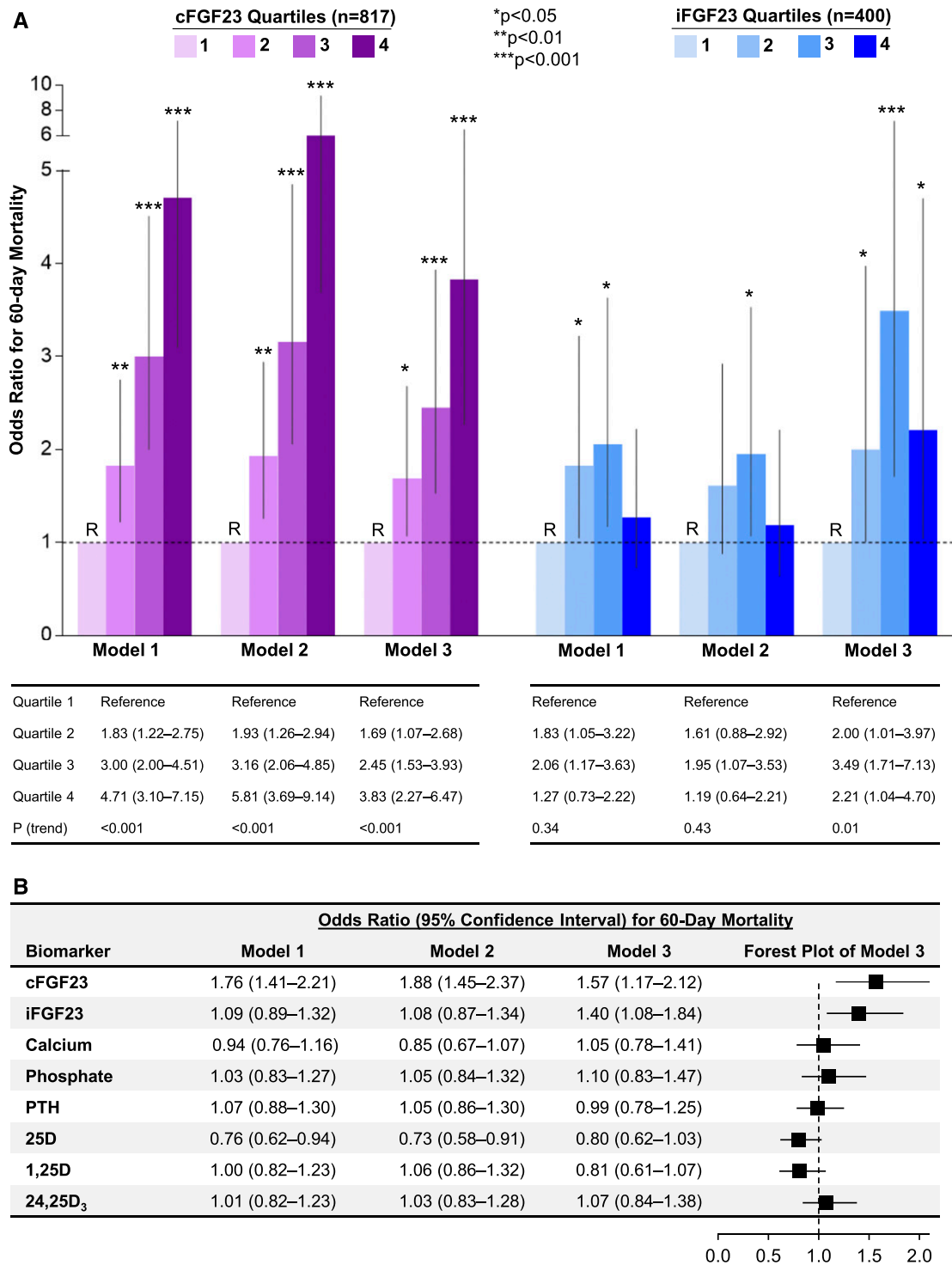
We investigated FGF23’s effects on vitamin D metabolite activation and catabolism as a potential unifying mechanism for our findings, because lower levels of 25D and 1,25D have been associated with higher mortality in critically ill patients (38,39). We found that higher iFGF23 levels are associated with lower 1,25D levels, consistent with FGF23-mediated inhibition of 25D-to-1,25D conversion in the kidneys (6) and perhaps elsewhere (34). To investigate whether impairment of 25D-to-1,25D conversion could partially account for our findings with respect to FGF23 and mortality, we adjusted for both 25D and 1,25D levels and found that iFGF23 remained significantly associated with death in the ATN study. Accordingly, the association between FGF23 and mortality could be mediated by pathways independent of vitamin D metabolism. Alternatively, tissue-level paracrine effects of FGF23 on vitamin D metabolism can occur in the absence of major changes in circulating vitamin D levels (40).

An alternative explanation for our findings could be confounding due to inflammation, because inflammation

**G**

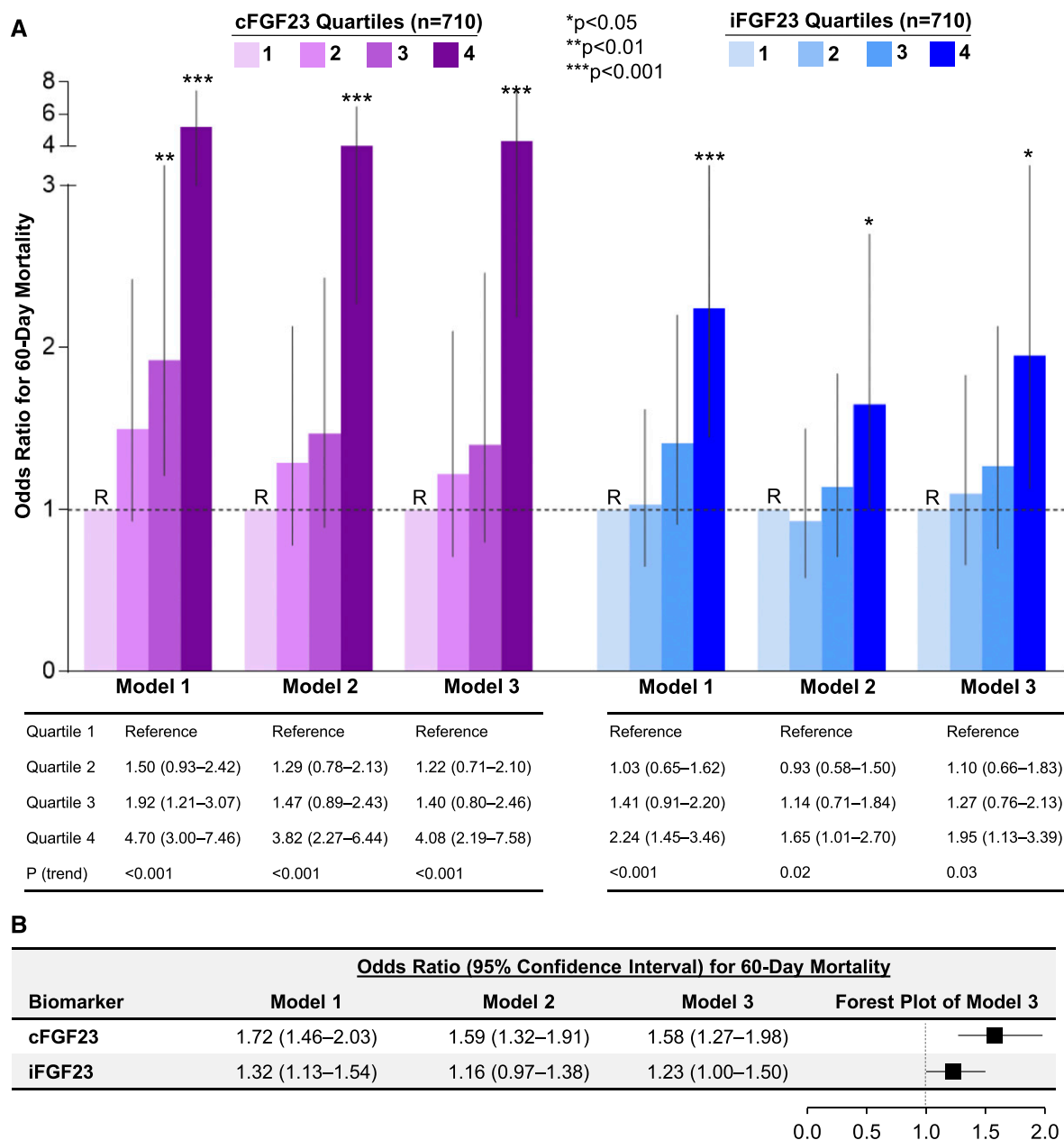
|                     | cFGF23  | Calcium | PO <sub>4</sub> | PTH    | 25D    | 1,25D    | 24,25D <sub>3</sub> | IL-6     |
|---------------------|---------|---------|-----------------|--------|--------|----------|---------------------|----------|
| iFGF23              | 0.30*** | 0.29*** | 0.33***         | -0.09  | 0.13** | -0.26*** | 0.10*               | -0.14**  |
| cFGF23              | ---     | 0.06    | 0.13***         | 0.16** | -0.04  | -0.04    | 0.10*               | 0.24***  |
| Calcium             | ---     | ---     | -0.09**         | -0.04  | 0.13** | -0.25*** | 0.13**              | -0.13*** |
| PO <sub>4</sub>     | ---     | ---     | ---             | 0.03   | -0.03  | 0.04     | 0.009               | -0.04    |
| PTH                 | ---     | ---     | ---             | ---    | -0.02  | 0.04     | 0.00                | 0.14**   |
| 25D                 | ---     | ---     | ---             | ---    | ---    | 0.40***  | 0.77***             | -0.11*   |
| 1,25D               | ---     | ---     | ---             | ---    | ---    | ---      | 0.31***             | 0.05     |
| 24,25D <sub>3</sub> | ---     | ---     | ---             | ---    | ---    | ---      | ---                 | 0.05     |

**Figure 1. | Associations between FGF23 and other mineral metabolites in the ARF Trial Network study.** Restricted cubic spline graphs depict the relationships between cFGF23, iFGF23, and (A and B) 25D, (C and D) 1,25D, and (E and F) 24,25D<sub>3</sub> ( $n=394$  for each graph). Four knots were specified at the fifth, 35th, 65th, and 95th FGF23 percentiles. The 95% confidence intervals are indicated by the shaded areas. (G) The Spearman correlation coefficient for each comparison. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ . 25D, 25-hydroxyvitamin D; 1,25D, 1,25-dihydroxyvitamin D; 24,25D<sub>3</sub>, 24,25-dihydroxyvitamin D<sub>3</sub>; cFGF23, C-terminal fibroblast growth factor 23; iFGF23, intact fibroblast growth factor 23; Ln, natural log-transformed; PO<sub>4</sub>, phosphate; PTH, parathyroid hormone;  $r_s$ , Spearman coefficient; RU/ml, reference units per milliliter.



**Figure 2. | Higher cFGF23 and iFGF23 levels associate with increased 60-day mortality in the ARF Trial Network study.** (A) Odds ratios for 60-day mortality according to quartiles of cFGF23 and iFGF23 levels. Quartile 1 was the reference (R) group in all models. cFGF23 levels by quartile: quartile 1, <1072 RU/ml; quartile 2, 1078–3587 RU/ml; quartile 3, 3608–12,951 RU/ml; quartile 4, >12,983 RU/ml. iFGF23 levels by quartile: quartile 1, <30 pg/ml; quartile 2, 30–88 pg/ml; quartile 3, 88–240 pg/ml; quartile 4, >241 pg/ml. (B) Odds ratios for 60-day mortality according to natural log-transformed mineral metabolite levels standardized to 1 SD. Model 1 is unadjusted. Model 2 is adjusted for demographics and comorbidities (age, sex, race, baseline eGFR, diabetes, cancer, congestive heart failure, and chronic liver disease). Model 3 is further adjusted for severity of illness (intensive care unit type, mechanical ventilation, Acute Physiology and Chronic Health Evaluation II score, RRT before randomization, treatment group, type of RRT, oliguria, sepsis, hypotension, white blood cell count, hemoglobin, and serum/plasma levels of albumin, creatinine, and IL-6). 25D, 25-hydroxyvitamin D; 1,25D, 1,25-dihydroxyvitamin D; 24,25D<sub>3</sub>, 24,25-dihydroxyvitamin D<sub>3</sub>; cFGF23, C-terminal fibroblast growth factor 23; iFGF23, intact fibroblast growth factor 23; PTH, parathyroid hormone; RU/ml, reference units per milliliter.

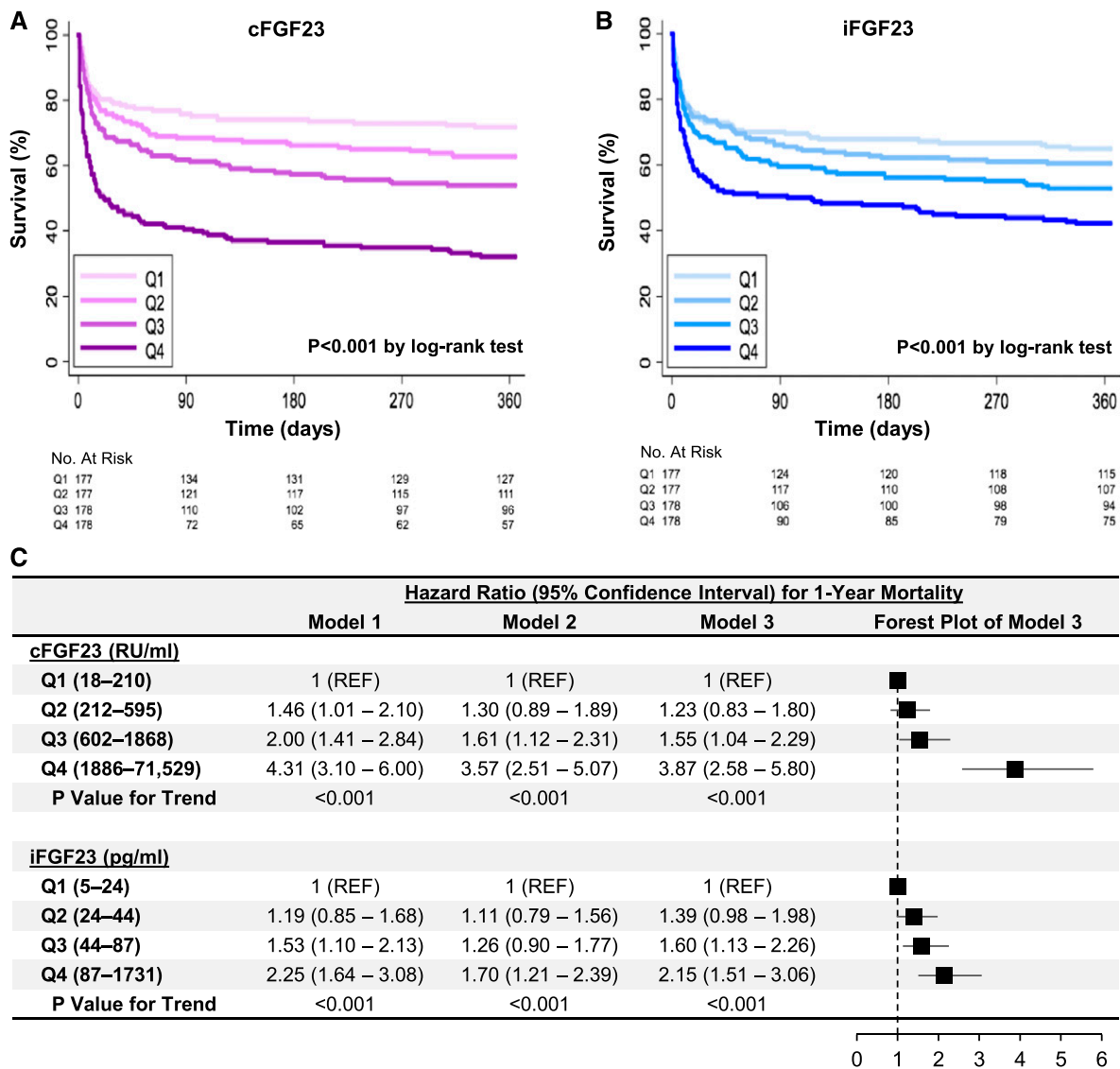




**Figure 3. | Higher cFGF23 and iFGF23 levels associate with higher 60-day mortality in the Validating Acute Lung Injury biomarkers for Diagnosis (VALID) study.** (A) Odds ratios for 60-day mortality according to quartiles of cFGF23 and iFGF23 levels. Quartile 1 was the reference (R) group in all models. cFGF23 levels by quartile: quartile 1, <211 RU/ml; quartile 2, 212–595 RU/ml; quartile 3, 602–1868 RU/ml; quartile 4, >1886 RU/ml. iFGF23 levels by quartile: quartile 1, <24 pg/ml; quartile 2, 24–44 pg/ml; quartile 3, 44–87 pg/ml; quartile 4, >87 pg/ml. (B) Odds ratios for 60-day mortality according to natural log-transformed FGF23 levels standardized to 1 SD. Model 1 is unadjusted. Model 2 is adjusted for demographics and comorbidities (age, sex, race, baseline eGFR, diabetes, cancer, congestive heart failure, and chronic liver disease). Model 3 is further adjusted for severity of illness (intensive care unit type, mechanical ventilation, Acute Physiology and Chronic Health Evaluation II score, sepsis, hypotension, white blood cell count, hemoglobin, serum creatinine, and AKI presence and severity on enrollment). cFGF23, C-terminal fibroblast growth factor 23; iFGF23, intact fibroblast growth factor 23; RU/ml, reference units per milliliter.

upregulates FGF23 production (41). Interestingly, we found that IL-6 associates positively with cFGF23 and negatively with iFGF23 (Figure 1G), consistent with inflammation as a stimulus for both FGF23 production and cleavage. This pattern of increased FGF23 production matched by a near commensurate increase in FGF23 cleavage has also been observed in patients with iron deficiency anemia (41). In multivariable models in the ATN study,

adjustment for IL-6 and other severity of illness covariates attenuated the association between cFGF23 and death, but strengthened the association between iFGF23 and death, consistent with inflammation acting as a positive and negative confounder of cFGF23 and iFGF23, respectively. Accordingly, confounding due to inflammation could be responsible for the differences we observed in the associations between cFGF23, iFGF23, and death in the ATN study.



**Figure 4. | Higher cFGF23 and iFGF23 quartiles associate with higher 1-year mortality in the Validating Acute Lung Injury biomarkers for Diagnosis (VALID) study.** (A and B) Kaplan–Meier curves showing survival during the first year after enrollment, stratified according to quartiles of cFGF23 and iFGF23. (C) Hazard ratios for 1-year mortality according to cFGF23 and iFGF23 quartiles. Model 1 is unadjusted. Model 2 is adjusted for demographics and comorbidities (age, sex, race, baseline eGFR, diabetes, cancer, congestive heart failure, and chronic liver disease). Model 3 is further adjusted for severity of illness (intensive care unit type, mechanical ventilation, Acute Physiology and Chronic Health Evaluation II score, sepsis, hypotension, white blood cell count, hemoglobin, serum creatinine, and AKI presence and severity on enrollment). cFGF23, C-terminal fibroblast growth factor 23; iFGF23, intact fibroblast growth factor 23; Q, quartile; REF, reference group; RU/ml, reference units per milliliter.

Levels of cFGF23 showed a stronger association with death than levels of iFGF23 in both cohorts. Thus, future studies of FGF23 as a prognostic biomarker for adverse outcomes in critical illness could reasonably focus on cFGF23 alone. However, future studies aimed at understanding the biology of FGF23 in critical illness and other acute inflammatory conditions should continue to measure both cFGF23 and iFGF23 to fully characterize the nature of FGF23 production and cleavage in these settings.

Limitations of this study include its observational design, measurement of FGF23 levels at a single time point, and lack of cause-specific mortality. The contribution of changes in unmeasured variables, such as klotho, the

coreceptor for FGF23, is unknown and requires additional study. Finally, although we adjusted for multiple covariates, we cannot exclude potential residual confounding by variables that may not have been ascertained. Although previous studies found that FGF23 levels increase in patients with AKI (12,42), we aimed to minimize confounding due to varying degrees of AKI severity by evaluating FGF23 levels in patients enrolled in the ATN study, all of whom had severe AKI requiring RRT. In VALID, we conducted sensitivity analyses stratified by the presence or absence of AKI on enrollment, and found similar results in both groups.

In conclusion, we found that higher plasma FGF23 levels are strongly and independently associated with an increased

risk of death in critically ill patients. These associations could be mediated through local effects of FGF23-mediated inhibition of 25D activation, which may not be reflected by plasma vitamin D metabolite levels, off-target effects independent of vitamin D signaling, or confounding by inflammation, which may not have been fully accounted for in multivariable models. Given substantial interest in vitamin D administration as a therapeutic strategy for improving outcomes in critical illness (43), future studies are needed to determine whether elevated FGF23 levels could identify patients less likely to respond to vitamin D precursors due to impaired 25D-to-1,25D conversion. Future studies are also needed to determine the physiologic regulators of FGF23 homeostasis in critical illness, whether FGF23 could serve as a useful prognostic tool, and whether FGF23 could be targeted for therapeutic benefit in interventional studies.

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#### Disclosures

None.

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