Supplementary material

Supplementary Figure 1: Midkine stability is ascertained by storage at -80°C and preclusion of repeated freeze-thaw cycles Supplementary Figure 2: Correlation of applied heparin with delta midkine levels Supplementary Figure 3: Serum midkine levels before and after dialysis treatment Supplementary Figure 4: Kaplan-Meier survival curves for tenascin-C and galectin Supplementary Figure 5: Correlative analyses of midkine, NTproANP and uPAR Supplementary Figure 6: Kaplan-Meier survival curves for ADMA Supplementary Figure 7: Midkine release in patients with hypervolemia. Supplementary Table 1: Fluid homeostasis assessment Supplementary Table 2: ADMA, SDMA, DGMV, L-arginine serum levels did not correlate with delta midkine serum levels Supplementary Table 3: Hazard Ratio for ΔMidkine, uPAR, and ANP adjusted for

age, gender, fluid status, and co-morbidity diabetes.

Supplementary Figure 1: Midkine stability is ascertained by storage at -80°C and preclusion of repeated freeze-thaw cycles. (A) Serum samples from 3 different patients were stored at 4°C, -20°C, or -80°C. Quantification of midkine was carried out after 4 days following immediate thawing before performance of the ELISA. (B) Serum samples from the same patients were stored at 4°C, -20°C, or -80°C for four days. Before quantification by midkine ELISA samples were thawed and subjected to another freezing/thawing cycle (1st thawing) and two more freezing/thawing cycles (2nd thawing). (C) Long term stability was tested with serum samples from two patients that were included for ELISA quantification after 2 and 3 years storage at -80°C.

Supplementary Figure 2: Correlation of applied heparin with delta midkine levels. (A) The doses of applied non-fractionated heparin were correlated with changes in serum midkine levels after a short ($r^2=0.06$, p=0.03) and long dialysis-free interval ($r^2=0.17$, p<0.001). (B) For non-fractionated heparin similar correlations of delta midkine levels after a short ($r^2=0.002$, p=0.88) and long dialysis-free interval ($r^2=0.01$, p=0.76) were performed in 12 patients (average 2,958 ± 1,389 IU). The increase of midkine levels was similar after both intervals, 17.6 ± 17 ng/ml and 14.5 ± 11.9 ng/ml, respectively, and did not positively correlate with the dose of fractionated heparin.

Supplementary Figure 3: Serum midkine levels before and after dialysis treatment. (A) Δ midkine values (midkine post – pre-hemodialysis) were calculated for the short and long interval, yielding no significant difference. Furthermore the $\Delta\Delta$ midkine levels (Δ midkine short interval - Δ midkine long interval) were calculated to assess the variability of midkine changes. (B) For the subgroups of diabetic (n=33) versus non-diabetic (n=50) patients analyses were performed. Δ midkine (midkine levels after dialysis - before dialysis) were calculated for the short and long dialysis-free interval. (C) For the subgroups of patients diagnosed with hypervolemia (n=51) versus euvolemia (n=32), Δ midkine (serum midkine levels after dialysis - before dialysis) were calculated for the short and long dialysis-free interval. Here, significant differences for the subgroup comparisons were confirmed. (D) ROC curve analyses showing the prognostic power of absolute midkine values or Δ midkine values to predict diabetes (cohort with n=70 diagnosed with diabetes; n=101 without diabetes),

hypervolemia (cohort with n=83 diagnosed with hypervolemia; n=88 without hypervolemia). In the last panels prediction of both conditions combined is tested (cohort with n=32 diagnosed with diabetes and hypervolemia; n=51 without diabetes and hypervolemia). AUC, area under the curve. **(E)** ROC curve analyses showing the prognostic power of Δ midkine values with binary logistic regression performed following short and long dialysis-free intervals to diagnose hypervolemia. Selected cut-off points for Δ midkine of the short and long intervals are indicated.

Supplementary Figure 4: Kaplan-Meier survival curves for tenascin-C and galectin. Serum concentrations of tenascin-C (A, B) and galectin (C,D) were determined. Kaplan-Meier survival curves indicate that both parameters are not predictive for overall survival.

Supplementary Figure 5: Correlative analyses of midkine, NTproANP and uPAR. NTproANP and uPAR serum concentrations were correlated (Pearson) with absolute midkine serum values in the whole dialysis cohort (A,C) or selected patients diagnosed with diabetes (B,D).

Supplementary Figure 6: Kaplan-Meier survival curves for ADMA. (A) Patients with less-than-average ADMA serum levels before the long dialysis-free interval (group 1) were compared with those having above-average ADMA levels (group 2). Over a 36 months observation period, censoring was performed for overall mortality and **(B)** cardiovascular mortality.

Supplementary Figure 7. Effect of fluid removal on midkine release in patients with hypervolemia. In patients diagnosed with hypervolemia (n=21) appropriate fluid management was planned by increasing net fluid removal during dialysis sessions. Serum midkine levels after long dialysis-free intervals were quantified. 8/21 patients successfully removed additional 0.5-1.0 kg (n=3) or >1.0 kg (n=5). (A) Δ Midkine levels were calculated as maximal difference to baseline in the 3 week intervention period with subgrouping according to weight loss: >0.5 to 1.0 kg and >1.0 kg. (B) Change of midkine values calculated as [%] change to baseline values again with subgrouping according to weight loss: >0.5 to 1.0 kg.







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Supplementary Table 1

Fluid homeostasis assessment

	hypervolemia defined by		
	comet tail sign lungs	elevated weight (> +0.5 kg)	V. cava diameter > 20mm
number of patients (n)	11	38	22

Delta Midkine	3 days interval above mean [24,9 ng/ml] n=42	3 days interval below mean [24,9 ng/ml] n=41	p-value
ADMA [µmol/l]	0.49 ± 0.99	0.49 ± 0.09	0.88
SDMA [µmol/l]	1.28 ± 0.36	1.33 ± 0.1	0.65
DGMV [mmol/l]	334.96 ± 218.34	356.07 ± 385.85	0.77
L-arginine [µmol/l]	22.78 ± 9.4	24.11 ± 20.7	0.72

ADMA, SDMA, DGMV, L-arginine, delta midkine serum levels

Supplementary Table 3. HR for ΔMidkine, uPAR, and ANP adjusted for age, gender, fluid status, and diabetes. (*Abbreviations: CI, confidence intervall;* HR, hazard ration.)

	HR	95% CI			
∆Midkine					
	1.059	1.026 - 1.094			
HR _{Gender}	0.905	0.447 - 1.830			
HRHypervolemia	1.367	0.652 - 2.863			
HR _{Diabetes}	0.898	0.457 - 1.762			
HR∆Midkine	2.204	1.244 - 3.641			
uPAR					
	1.052	1.017 - 1.088			
HR _{Gender}	1.083	0.533 - 2.199			
HR _{Hypervolemia}	1.720	0.824 - 3.592			
HR _{Diabetes}	0.987	0.501 - 1.943			
	2.421	1.153 - 5.086			
ANP					
	1.055	1.022 - 1.090			
HR _{Gender}	0.951	0.468 - 1.934			
HR _{Hypervolemia}	1.729	0.833 - 3.590			
HR _{Diabetes}	1.036	0.529 - 2.029			
	1.352	0.678 - 2.697			