

# INDOXYL SULFATE RELATED TO PROTEIN INTAKE AND INTRAHEMODIALYSIS ARTERIAL PRESSURE VARIATION, AND $\beta_2$ -MICROGLOBULIN TO ALBUMIN LEVELS AND INFLAMMATION

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## INTRODUCTION AND AIM

- Hypoalbuminemia is a strong factor for morbidity and mortality in the general and renal population. Inflammation and dietary protein intake exert competing effects on serum albumin over time.
- The nPCR is considered a valid surrogate for dietary protein intake under steady-state conditions, in the face of inflammation may overestimate protein intake because of endogenous nitrogen breakdown.
- High protein intake is associated with increased levels of albumin-bound uremic solutes in hemodialysis patients and this products show concentrations not determined by hemodialysis clearances.
- The objective of the study is to analyze the patterns of relationship of the prototypic uremic toxins with protein metabolism, inflammation, and other candidate clinical variables

## METHODS 1: SAMPLES AND STATISTICS

- Cross-sectional study of 60 chronic hemodialysis (HD) patients.
- Pre/postHD total serum Indoxyl Sulfate (IS) by high-performance liquid chromatography (HPLC).
- Pre/post (post in hemodilution patients-HDF-) serum  $\beta_2$ -microglobulin (B2M) by nephelometry.
- Urea kinetic model calculated by Solute Solver in midweek sessions: KD urea (urea dialyzer clearance), Kd urea (urea difusive dialyzer clearance), Eq Kt/V (equilibrated Kt/V), KoA urea (urea mass transfer-area coefficient), G2P urea (urea double pool generation rate), TAC urea, SAN\_DstdKt/V (weekly Dialysis Standard Kt/V normalized to body Surface Area), dp nPCR (normalized Protein Catabolic Rate double pool), V2P urea double pool.
- Testing of variables by Kolmogorov-Smirnov for normality ( $p>0,20$ ).
- T-test or ANOVA for IS and B2M as dependent variables; and as categorical factors: albumin  $\geq$  or  $<$  38 g/L, dp nPCR categorized by levels, and albumin and dp nPCR profiles. T-test for albumin, dp nPCR and CRP as dependent variables; and as categorical factors: IS higher or lower than mean, and B2M higher or lower than mean.
- Bivariate correlations (Pearson) and multiple regression analyses of IS and B2M as dependent variables; as independent variables: biologic (hemoglobin, lymphocytes, BUN, urate, HCO3-, P, Na, Log CRP, albumin by Brom cresol Green...), kinetic (SAN\_DstdKt/V, dp nPCR, URR, IS reduction rate -RR-, B2M RR-), and other clinical variables (residual diuresis, mean arterial pressure/MAP change, convective volume/CV in HDF, BMI, BSA...).
- MAP change: calculated as the difference between measures at the start and the end of each session. MAP=DAP+1/3(SAP-DAP).

## METHODS 2: PATIENTS AND SESSIONS

- 60 patients of an university regional hospital chronic haemodialysis unit.
- 60±20 (X±SD) years (range 20-84). 32 male (53,3%) / 28 (46,6%) female.
- Height 161,1±1,03 cm. Dry weight 67,5±19,3. BMI 25,5±0,9 kg/m<sup>2</sup>. BSA 1,7±0,03 m<sup>2</sup>.
- Modified Charlson's Index 7,2±2,85.
- Residual diuresis: 34 p null, 21 p <500 ml/24h, 5 p >500 ml/24 h.
- 21,66% (13 p) of the patients with positivity for hepatitis C/B or HIV: C hepatitis 8, B hepatitis 2, HIV 1, HIV and C hepatitis 1, B and C hepatitis 1.
- Haemodialysis session time 244±19'. Q<sub>0</sub> 356,8±6,13 ml/min. Q<sub>0</sub> standard HD 500-800 ml/min. Q<sub>0</sub> HDF 800 ml/min.
- Dialyzers: Membrane: 56 polynephron, 4 cellulose. 34 High flux, 26 medium flux. Surface: 11.17 m<sup>2</sup>; 33 1.9 m<sup>2</sup>; 16 2.1 m<sup>2</sup>.
- 29 p (48,3%) on line hemodilution / 31 p (51,6%) standard hemodialysis.
- Ultrafiltration 2,3±0,93 L. Convective volume in HDF 23,8±2,8 L. Infusion volume in HDF 21,4±2,7 L.
- Mean dialysis prevalence months 46,8 (3,9 years) (range 3-299 months): mean 58,3 months HDF patients and 36 months standard HD patients.

## METHODS 3: NORMALITY

**Normality:**  
 Indoxyl sulfate D=0,09 p=0,20, IS RR D=0,08 p<0,20,  $\beta_2$ -microglobulin (HDF) D=0,14 p>0,20,  $\beta_2$ -microglobulin (standard HD) D=0,14 p>0,20,  $\beta_2$ -microglobulin RR D=0,18 p>0,20, urea D=0,07 p>0,20, hemoglobin D=0,05 p>0,20, hematocrit D=0,06 p>0,20, leucocytes D=0,10 p>0,20, lymphocytes D=0,10 p>0,20, residual diuresis D=0,14 p>0,20, convective volume D=0,13 p>0,20, infusion volume D=0,16 p>0,20, HCO3- D=0,09 p>0,20, pH D=0,09 p>0,20, P D=0,11 p>0,20, Log CRP D=0,05 p>0,20, MAP change D=0,07 p>0,20, urea D=0,12 p>0,20, height D=0,05 p>0,20, Charlson index D=0,11 p>0,20, albumin D=0,10 p>0,20, dp nPCR D=0,08 p>0,20, Eq Kt/V urea D=0,06 p>0,20, G2P urea D=0,09 p>0,20, TAC urea D=0,07 p>0,20, BSA D=0,09 p>0,20

### No normality:

URR D=0,14 p<0,15, prevalence months D=0,21 p<0,1, time in minutes D=0,35 p<0,01, Q<sub>0</sub> D=0,28 p<0,1, PTHi D=0,16 p<0,1, Ionic Ca D=0,16 p<0,1, Na D=0,26 p<0,01, CRP D=0,32 p<0,01, BMI D=0,18 p<0,05, KoA urea D=0,25 p<0,01, KD urea D=0,17 p<0,10, Kd urea D=0,15 p<0,15, Kc D=0,19 p<0,05, V2P urea D=0,19 p<0,05, SAN\_DstdKt/V D=0,19 p<0,05

## METHODS 4: VARIABLES

### Regression analysis:

- Indoxyl sulfate
- $\beta_2$ -microglobulin
- Indoxyl sulfate reduction rate
- $\beta_2$ -microglobulin reduction rate
- BUN preHD 78,6±11,8 mg/dl
- Hemoglobin 11,2±0,2 g/dl
- Hematocrit 34,4±0,5 %
- Lymphocytes 1,490±86,9 cell/ml
- Albumin 37,8±0,5 g/L
- HCO3- 23,9±0,3 mmol/L
- pH 7,4±0,01
- G,350,18 mg/dl
- Log CRP
- Urate 5,±0,18 mg/dl
- Eq Kt/V 1,6±0,05
- G2P urea 5,3±0,26 mg/min
- TAC urea 34,3±1,26 mg/dl
- SAN\_DstdKt/V 2,7±0,07
- dp nPCR: 1±0,04 g/kg/day
- Mean arterial pressure:
- Initial: 87,3±2,13 mmHg
- Final: 86,5±2,01 mmHg
- Mean arterial pressure change: -0,6 mmHg
- Modified Charlson index 7,2±2,85 points
- Height 161,1±1,03 cm
- Body surface area (BSA) 1,7±0,03 m<sup>2</sup>

## METHODS 5: VARIABLES

### Student-t / ANOVA:

- Dependent variables:  
 • Indoxyl sulfate  
 •  $\beta_2$ -microglobulin  
 Categorical factors:  
 • CRP  $\geq$  1 mg/dl / < 1 mg/dl  
 • Albumin  $\geq$  38 g/L / < 38 g/L  
 • dp nPCR  $\geq$  1 g/kg/day / < 1 g/kg/day  
 • dp nPCR (g/kg/day): Figure 1a  
 • <0,6; 5 p (8,3%)  
 • 0,6-0,79; 11 p (18,3%)  
 • 0,8-0,99; 14 p (23,3%)  
 • 1-1,19; 17 p (28,3%)  
 • >1,2; 13 p (21,6%)  
 • Albumin and dp nPCR profiles (Figure 1b)  
 • Catabolic (Albumin<38 g/L)  
 • dp nPCR  $\geq$  1 g/kg/day; 12 p (20%)  
 • Low intake (Albumin<38 g/L)  
 • dp nPCR < 1 g/kg/day; 21 p (35%)  
 • High intake (Albumin>38 g/L)  
 • dp nPCR  $\geq$  1 g/kg/day; 19 p (31,6%)  
 • Anabolic (Albumin>38 g/L)  
 • dp nPCR < 1 g/kg/day; 8 p (13,3%)
- Dependent variables:  
 • Albumin  
 • dp nPCR  
 Categorical factors:  
 • IS<sub>2</sub> mean / < mean  
 • B2M mean / < mean

## RESULTS 6: SIGNIFICANT BIVARIATE CORRELATIONS

- Indoxyl sulfate:  
 $P: r=+0,11 / p=0,007$   
 $BUN: r=+r^2=0,11 / p=0,009$   
 $Albumin: r=+r^2=0,10 / p=0,01$   
 $TAC urea: r=+r^2=0,09 / p=0,01$   
 $PCRN_2P: r=+r^2=0,08 / p=0,02$   
 $MAP change: r=+r^2=0,08 / p=0,02$
- Indoxyl sulfate reduction rate:  
 $G2P urea: r=+r^2=0,17 / p=0,0008$   
 $Eq Kt/V: r=+r^2=0,15 / p=0,002$   
 $Height: r=+r^2=0,11 / p=0,007$   
 $MAP change: r=+r^2=0,11 / p=0,008$   
 $TAC urea: r=+r^2=0,11 / p=0,009$   
 $BSA: r=+r^2=0,08 / p=0,02$   
 $HCO3-: r=+r^2=0,19 / p=0,01$   
 $BUN: r=+r^2=0,07 / p=0,04$   
 $TAC urea: r=+r^2=0,17 / p=0,02$   
 $BSA: r=+r^2=0,16 / p=0,02$   
 $TAC urea: r=+r^2=0,16 / p=0,02$
- $\beta_2$ -microglobulin:  
 $Residual diuresis: r=+r^2=0,17 / p=0,0008$   
 $Albumin: r=+r^2=0,11 / p=0,007$   
 $Log CRP: r=+r^2=0,07 / p=0,03$
- $\beta_2$ -microglobulin reduction rate:  
 $G2P urea: r=+r^2=0,32 / p=0,001$   
 $Log CRP: r=+r^2=0,30 / p=0,001$   
 $Eq Kt/V: r=+r^2=0,27 / p=0,003$   
 $Infusion volume: r=+r^2=0,21 / p=0,001$   
 $BSA: r=+r^2=0,17 / p=0,02$   
 $TAC urea: r=+r^2=0,16 / p=0,02$

## RESULTS 5: ANOVA

IS and B2M as dependent variables  
 vs dp nPCR categorized by levels (Figure 1a)  
 and vs albumin/dp nPCR profiles (Figure 1b)

## BIBLIOGRAPHY

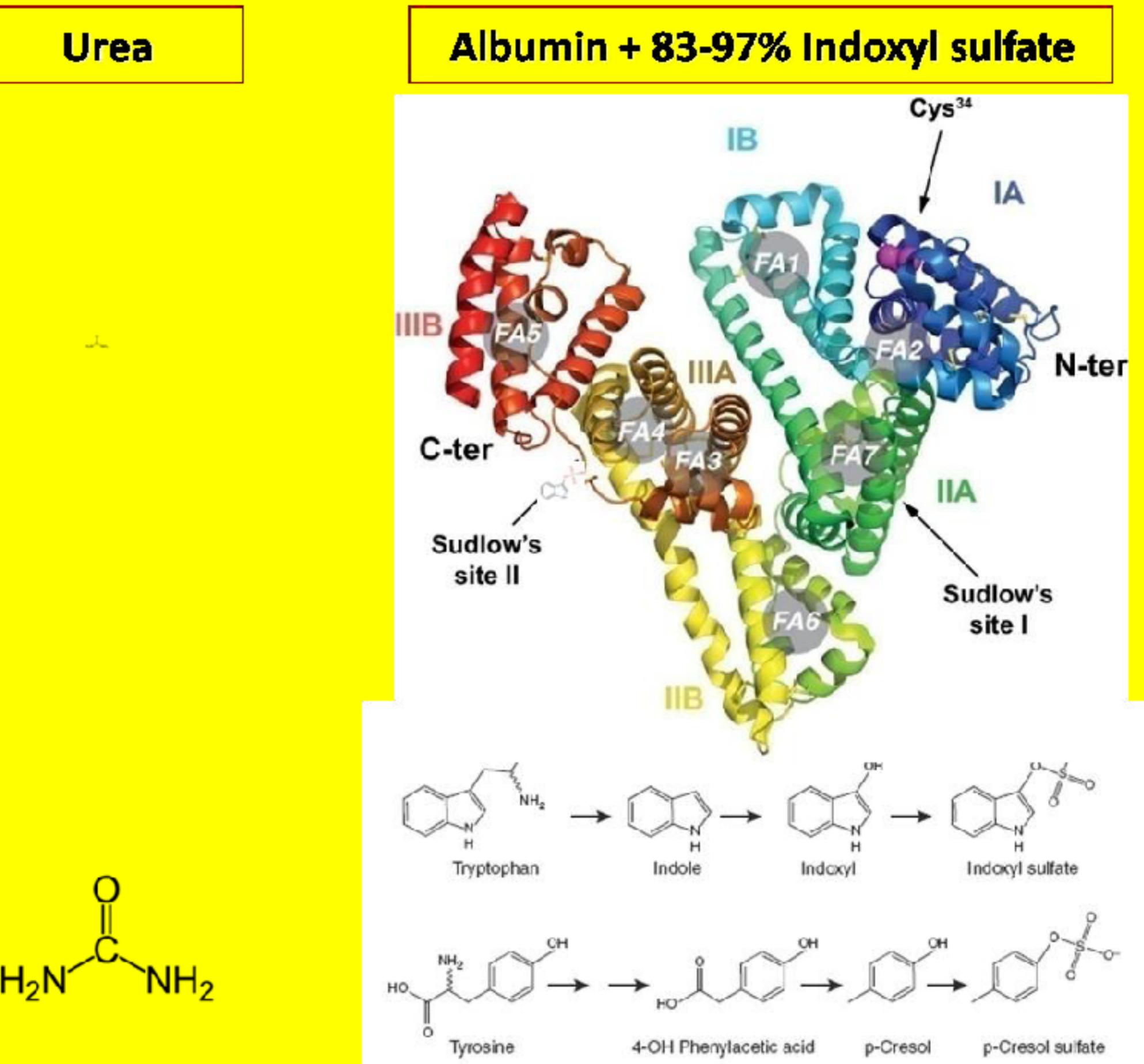
- Meyer T et al. Uremic solutes from colon microbes. *Kidney International* 81: 949-954, 2012 / Kayser G et al: Inflammation and dietary protein intake exert competing effects on serum albumin and creatinine in hemodialysis patients. *Kidney Int* 60: 333-340, 2001 / Viernes I et al: Albumin is the main plasma binding protein for Indoxyl sulfate and p-cresyl sulfate in hemodialysis patients. *Kidney Int* 60: 763-767, 2001 / Meijers B et al: The gut-kidney axis: Indoxyl sulfate and p-cresyl sulfate induce CKD-associated hypertension via a 1-hydroxy-2-oxo-3-oxo-4-phenylpropanoate receptor in the RAAS activation. *Am J Physiol Renal Physiol* 311:435-441, 2011 / Yu M et al: Indoxyl sulfate-induced endothelial dysfunction in patients with chronic kidney disease via an induction of oxidative stress. *Circ J Am Soc Nephrol* 75:39-44, 2011 / Sun Ch et al: Suppression of Klotho expression by protein-bound uremic toxins is associated with increased DNA methyltransferase expression and DNA hypermethylation. *Kidney Int* 81: 640-650, 2012 / Barreto F et al: Serum indoxyl sulfate is associated with vascular disease and mortality in hemodialysis patients. *Kidney Int* 81: 651-658, 2012 / Meyer T et al: p-Cresyl sulfate and indoxyl sulfate induce inflammatory responses in cultured proximal renal tubular cells. *Nephrol Dial Transplant* 1: 1-9, 2012 / Sun Ch et al: Suppression of Klotho expression by protein-bound uremic toxins is associated with increased DNA methyltransferase expression and DNA hypermethylation. *Kidney Int* 81: 640-650, 2012 / Meyer T et al: Indoxyl sulfate and p-cresyl sulfate induce vascular calcification in hemodialysis patients. *Kidney Int* 81: 651-658, 2012 / Schwabe E et al:  $\beta_2$ -microglobulin amyloidosis: a vanishing complication of long-term hemodialysis? *Kidney Int* 30:385-390, 1986 / Kazama J et al: Reduction of circulating  $\beta_2$ -microglobulin level for the treatment of dialysis-related amyloidosis. *Nephrol Dial Transplant* 16 (Suppl 4): 31-35, 2001 / Niwa T et al: Amyloid  $\beta_2$ -microglobulin is modified with N-epicarboxymethyl-lysine in dialysis-related amyloidosis. *Kidney Int* 50:1303-1309, 1996 / Miyata T et al: Involvement of  $\beta_2$ -microglobulin with N-epicarboxymethyl-lysine in the formation of dialysis-related amyloidosis. *Nephrol Dial Transplant* 16: 31-35, 2001 / Miyata T et al: Reactive carbonyl compounds related to uremic toxicity ("carboxyl stress"). *Kidney Int* 59: 525-531, 2001 / Okuno S et al: Serum  $\beta_2$ -microglobulin, a useful marker with a double role in hemodialysis patients. *Nephrol Dial Transplant* 17: 174-175, 2002 / Cheung A et al: Serum  $\beta_2$ -microglobulin levels predict mortality in dialysis patients: results of the HEMO study. *J Am Soc Nephrol* 17: 546-555, 2006 / Cheung A et al: Association between serum  $\beta_2$ -microglobulin level and infection and mortality in hemodialysis patients. *Clin J Am Soc Nephrol* 2008: 3: 69-77.

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## Urea

## Albumin + 83-97% Indoxyl sulfate

## MHC-I $\beta_2$ -microglobulin



X±SE (mg/L)	Range (mg/L)	Hemodiafiltration p	Standard HD p	Normal values	Mortality risk augmented
$\beta_2$ M 35,6±1,9	11,5-72,1	32,8±1,9 mg/L	38,2±3,1 mg/L	<3,5mg/L	>25mg/L
IS 18,91±1,63	0,28-55,12	22,4±2,7 mg/L	16,04±1,6 mg/L	<1,2mg/L	>4mg/L

## RESULTS 1:

$\beta_2$ M and IS means in the whole population and in the hemodiafiltration and standard hemodialysis patients

Total	Hemodiafiltration p	Standard HD p
Urea Reduction Rate	78,6±1,1 %	82,04±1,1 %
$\beta_2$ microglobulin Reduction Rate	79,8±1,1 %	
Indoxyl sulfate Reduction Rate	47,1±2,1 %	49,07±2,8 %

## RESULTS 2:

Urea,  $\beta_2$ M and IS reduction rates

Mean IS (CI 95%)	p	Mean  $\beta_2$ M (CI 95%)	p





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