





UREMIC TOXINS: WHAT THEY DO AND HOW TO CLEAR THEM

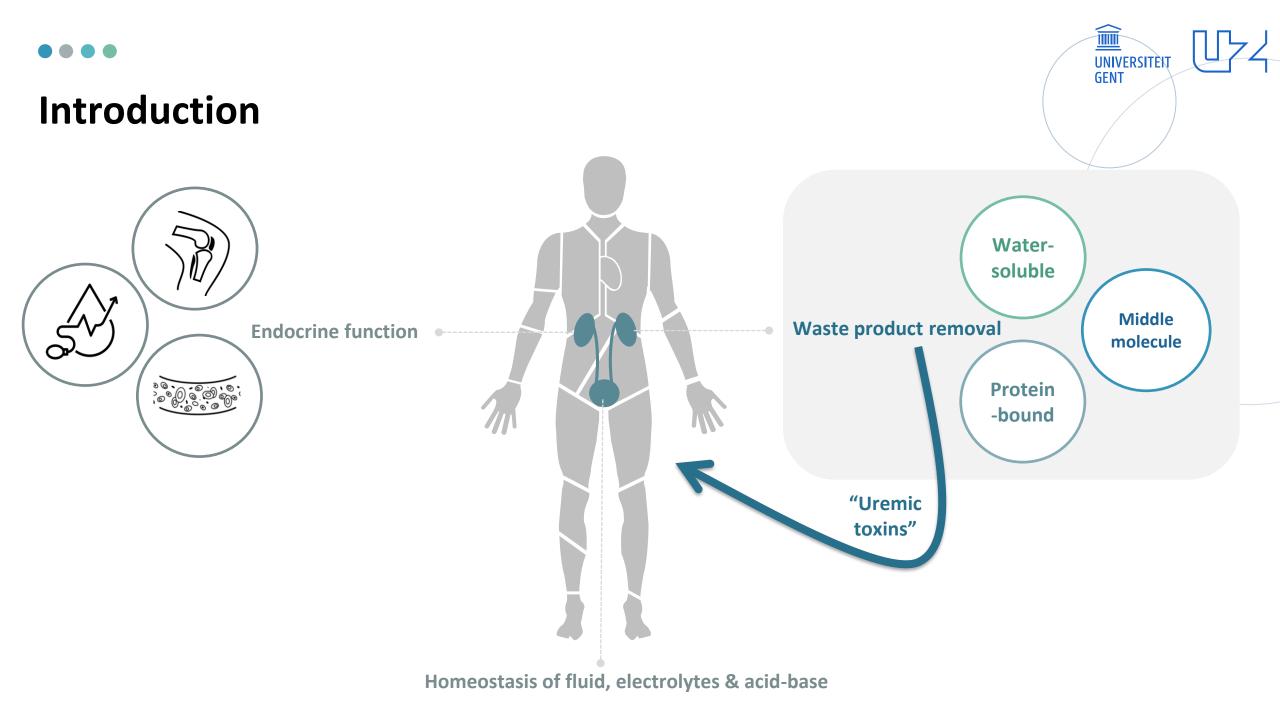
EVELIEN SNAUWAERT, MD, PhD

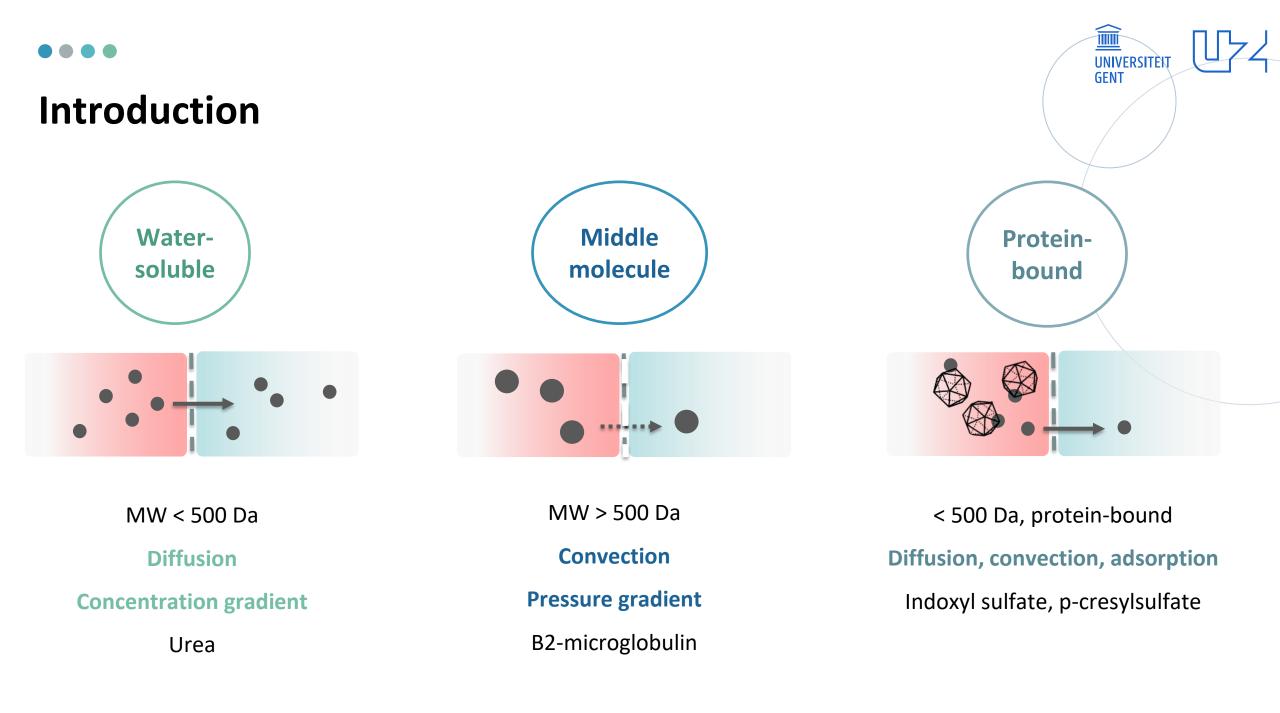
DEPARTMENT OF PEDIATRIC NEPHROLOGY Ghent University Hospital, Belgium







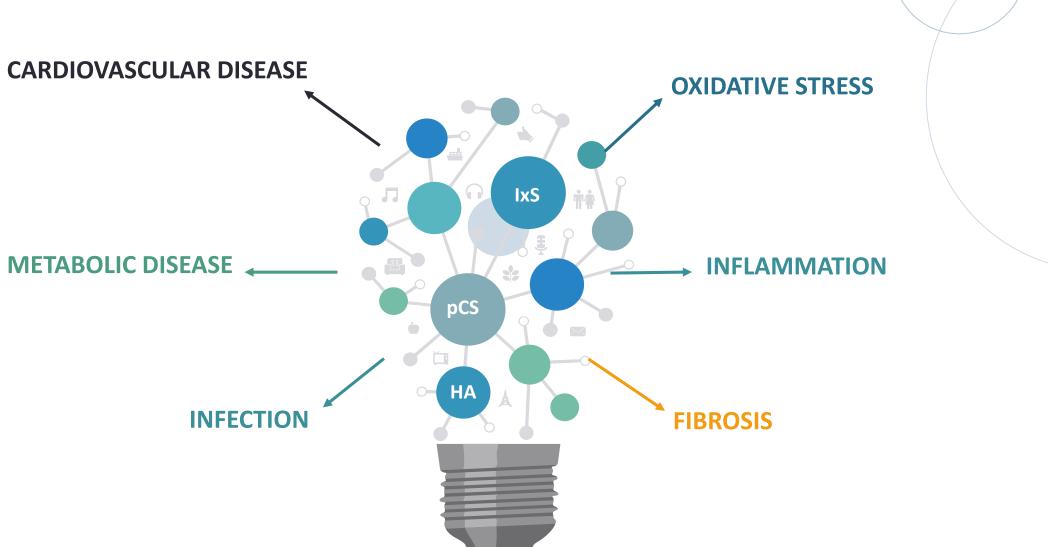






Uremic toxins, what they do?

Pathophysiology



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Which uremic toxins are making our patient sick?









Confounding factors (pre-existing CV disease)





Inability to decrease a single compound



Complex and multifactorial interplay between different key elements, present for longer time



Which uremic toxins are making our patient sick?



No confounding factors or co-morbidities

Isolated kidney disease

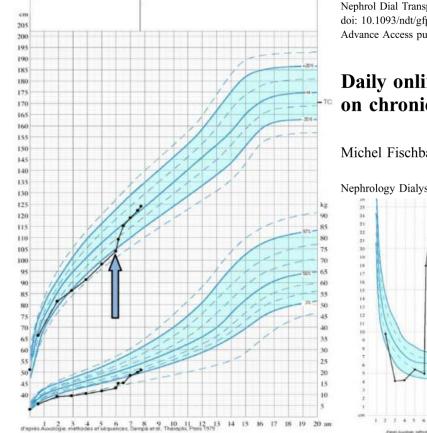
Growth



Opportunities in pediatric population



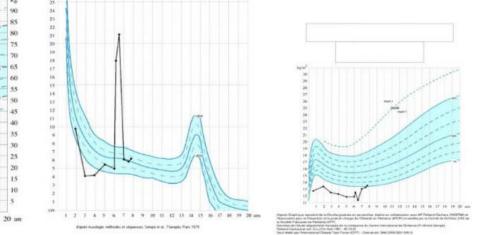
Enhancing uremic solute removal improves growth



Nephrol Dial Transplant (2010) 25: 867–873 doi: 10.1093/ndt/gfp565 Advance Access publication 4 November 2009

Daily online haemodiafiltration promotes catch-up growth in children on chronic dialysis

Michel Fischbach, Joelle Terzic, Soraya Menouer, Céline Dheu, Laure Seuge and Ariane Zalosczic



Nephrology Dialysis Transplantation Children's Unit, University Hospital Hautepierre, Avenue Molière, 67098 Strasbourg, France

Figure 2 Daily intensive online hemodiafiltration (1) promotes catch-up growth (from reference 12).



In pediatrics?

Pediatric Nephrology (2018) 33:921–924 https://doi.org/10.1007/s00467-018-3920-8

EDITORIAL COMMENTARY

CrossMark

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A plea for more uremic toxin research in children with chronic kidney disease

Evelien Snauwaert¹ · Wim Van Biesen² · Ann Raes³ · Griet Glorieux² · Raymond Vanholder² · Johan Vande Walle³ · Sunny Eloot²

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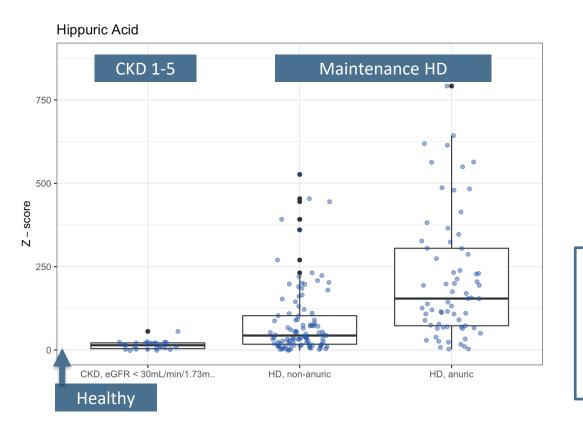
UToPaed study: uremic toxins in pediatric CKD

more advanced and appropriate tools to improve management of children with CKD



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Pediatric reference frame



50 healthy children 65 children CKD stage 1-5 (not on dialysis) 170 children hemodialysis



†††

6 protein-bound uremic toxins 4 small water-soluble uremic toxins 2 middle molecules

Ensure use of biologically relevant uremic toxin levels in experimental studies

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ht st

KOF,

Allow proper design of studies



Results from observational study to be announced soon



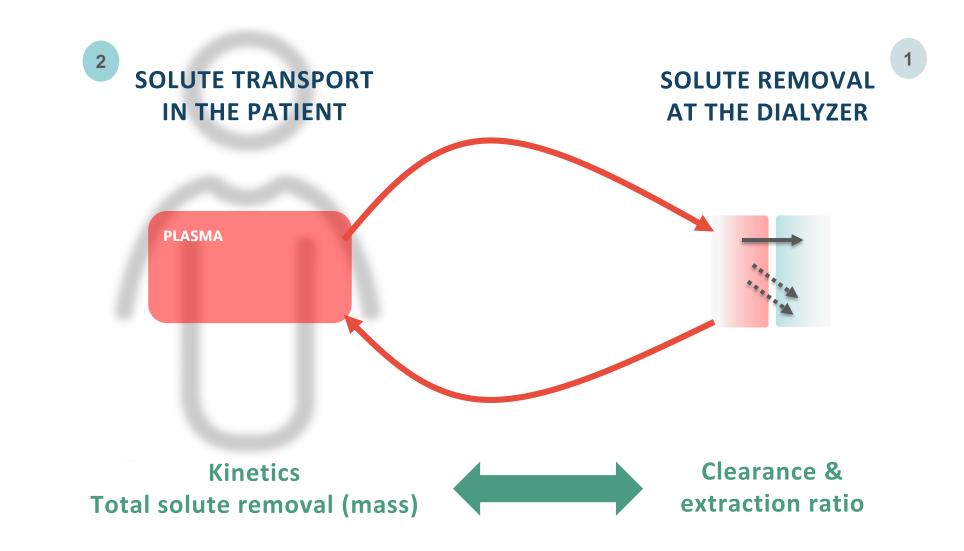


Uremic toxins, how to clear them?

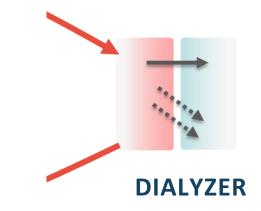




How to clear uremic toxins?

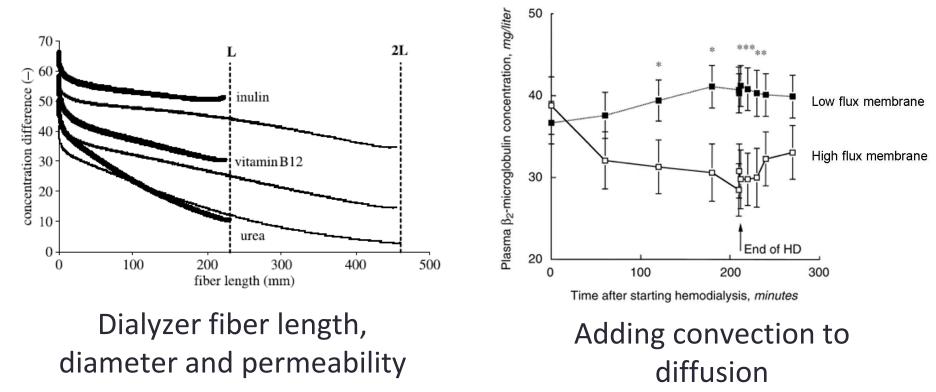


Solute transport in the dialyzer



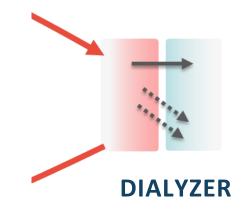


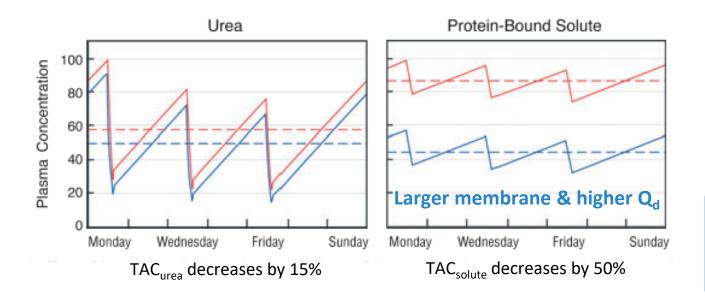
Clearance and extraction ratio ≈ **blood flow** + membrane + dialysate flow



Eloot et al., Comp Meth Biomech Biomed Eng 2006; Leypoldt et al., Kidney Int, 1999

Solute transport in the dialyzer





Diffusion = well established removal strategy of free fraction protein-bound uremic toxins

Table 2. Instantaneous clearance (mL/min) at 60 min

	Post-HDF	Pre-HDF	Pre-HF
Urea	243.0 ± 18.7	$230.1 \pm 10.5^{*}$	$150.7 \pm 15.0^{\circ \circ, \S\S}$
Creatinine	179.4 ± 48.3	$148.9 \pm 22.3^{*}$	$103.7 \pm 19.9^{\circ \circ , \S\S}$
Uric acid	166.4 ± 14.1	$153.4 \pm 9.8^{*}$	$104.8 \pm 8.9^{\circ \circ , \S\S}$
β2Μ	82.8 ± 16.1	$67.2 \pm 18.5^{*}$	$87.5 \pm 9.6^{\S}$
Hippuric acid	131.2 ± 15.6	$121.4 \pm 13.1^{*}$	$68.7 \pm 23.9^{\circ, \S}$
Indole acetic acid	66.6 ± 8.6	67.5 ± 9.3	$38.8 \pm 5.4^{\circ \circ , \S\S}$
Indoxylsulfate	33.4 ± 7.4	34.7 ± 9.9	$18.7 \pm 6.6^{\circ \circ , \S\S}$
<i>p</i> -Cresylsulfate	23.5 ± 4.6	24.6 ± 6.4	$12.9 \pm 2.5^{\circ \circ , \S\S}$
CMPF			

Pre-HDF versus post-HDF: *P < 0.017, **P < 0.001; pre-HF versus post-HDF: °P < 0.017, °°P < 0.001; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P < 0.001$; or P < 0.001; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P < 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P < 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P < 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P < 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P < 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P = 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P = 0.017$, ${}^{\$}P = 0.001$; pre-HF versus pre-HDF: ${}^{\$}P = 0.001$; pre-HF versus pre-HF versus pre-HDF versus pre-HDF versus pre-HF versus pre-HF versus pre-HF versus pre-HF versus pre-HF versus pre-HF versu

Solute transport in the dialyzer



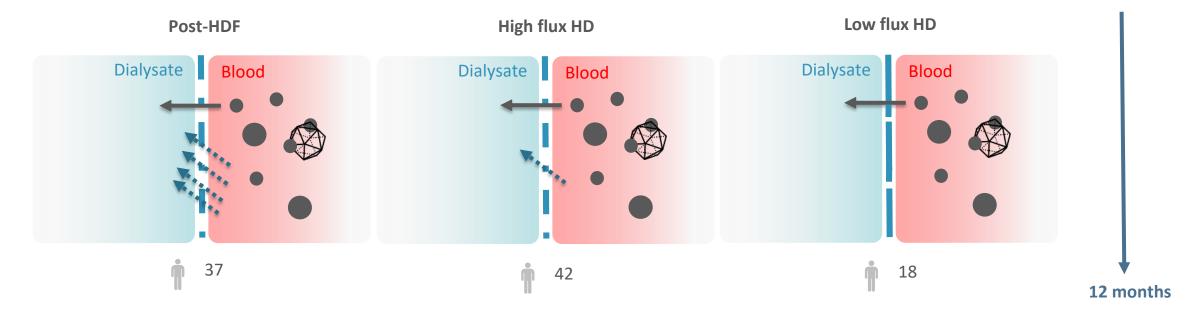




Research question: does post-HDF decrease levels of protein-bound uremic toxins ?



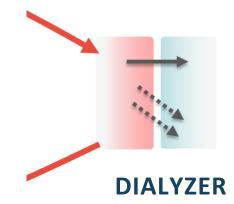
baseline



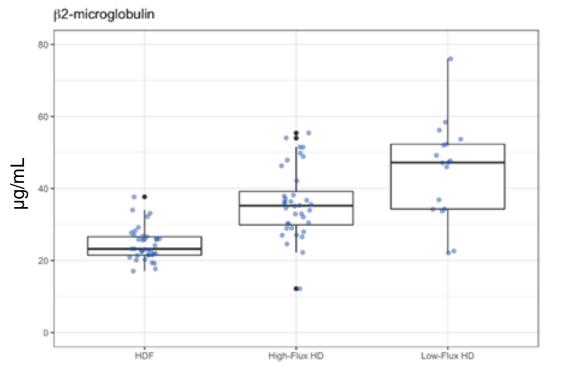
Snauwaert et al., NDT, 2020 Apr 1;35(4):648-656.

Solute transport at the dialyzer

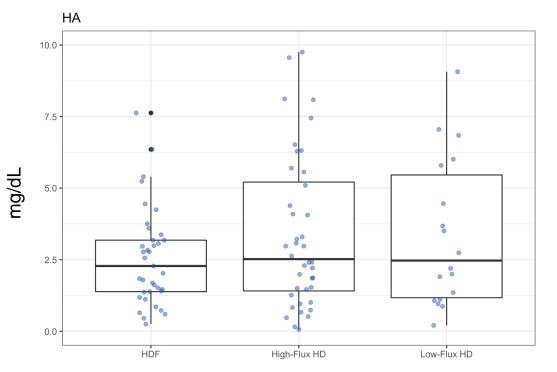




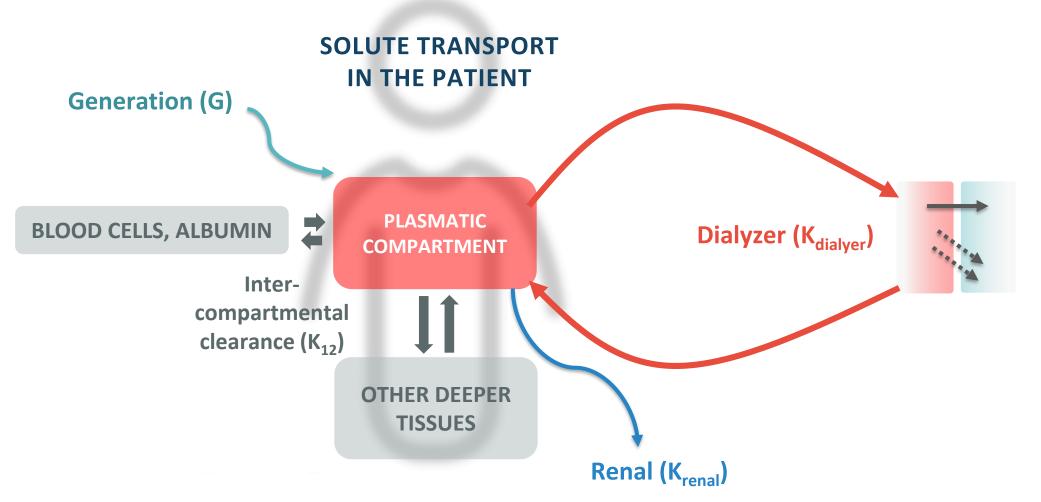
MIDDLE MOLECULES



PROTEIN-BOUND UREMIC TOXINS

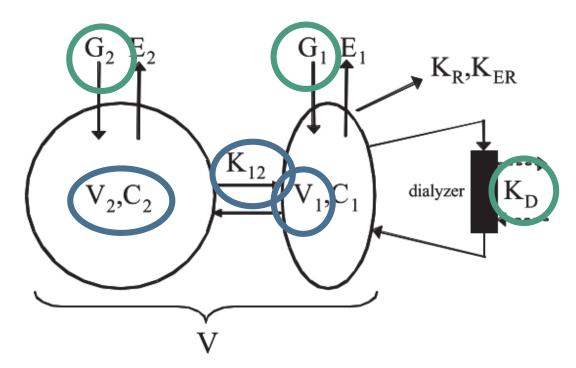


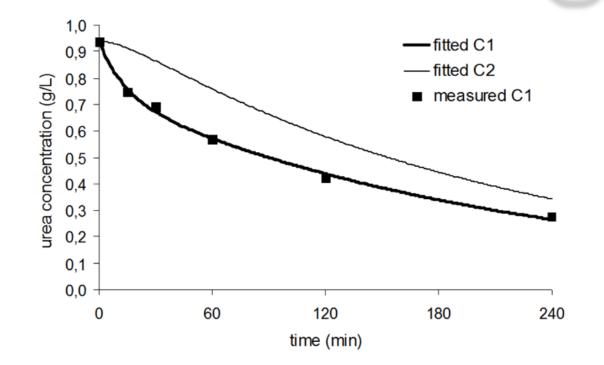
Snauwaert et al., NDT, 2020 Apr 1;35(4):648-656.



Eloot et al., Seminars in Dialysis, 2012

Two-compartiment kinetic model

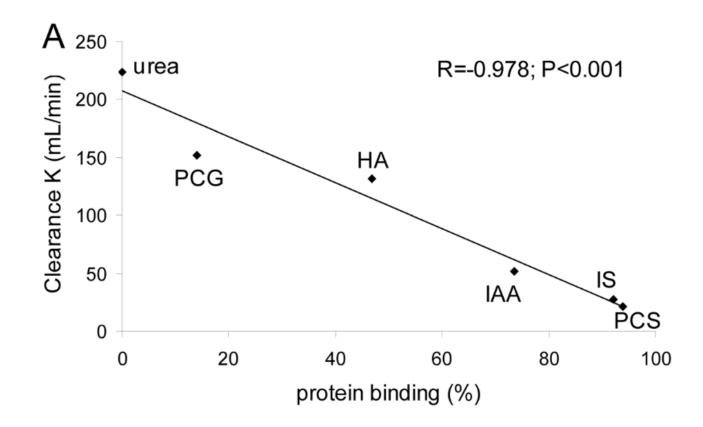




PLASMA

PATIENT

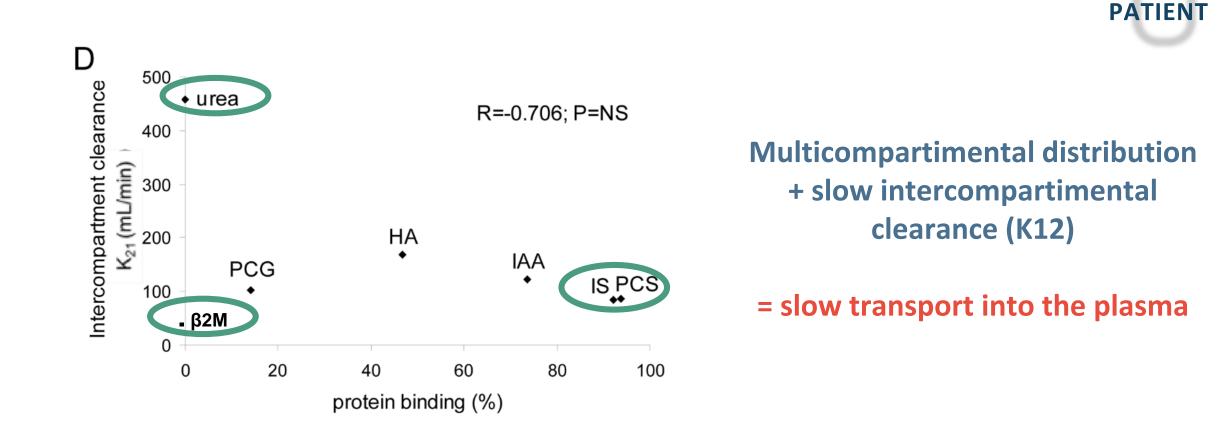
Eloot et al., Seminars in Dialysis, 2012; Eloot et al., PLOS one, 2016





Dialyzer clearance shows an inverse relation with % protein-binding

Only free fraction can be removed!



PLASMA

Solute transport at the dialyzer



Haemodiafiltration does not lower protein-bound uraemic toxin levels compared with haemodialysis in a paediatric population

Evelien Snauwaert D¹, Wim Van Biesen¹, Ann Raes¹, Griet Glorieux¹, Johan Vande Walle¹, Sanne Roels², Raymond Vanholder¹, Varvara Askiti³, Karolis Azukaitis⁴, Aysun Bayazit⁵, Nur Canpolat⁶, Michel Fischbach⁷, Krid Saoussen⁸, Mieczyslaw Litwin⁹, Lukasz Obrycki⁹, Fabio Paglialonga¹⁰, Bruno Ranchin¹¹, Charlotte Samaille¹², Franz Schaefer¹³, Claus Peter Schmitt¹³, Brankica Spasojevic^{14,15}, Constantinos J. Stefanidis³, Rukshana Shroff^{16,*} and Sunny Eloot^{1,*}

Snauwaert et al., NDT, 2020 Apr 1;35(4):648-656.







A Sad but Forgotten Truth: The Story of Slow-Moving Solutes in Fast Hemodialysis

Sunny Eloot, Wim Van Biesen, and Raymond Vanholder Nephrology Section, Department of Internal Medicine, Ghent University Hospital, Gent, Belgium

Does the Adequacy Parameter Kt/V_{urea} Reflect Uremic Toxin Concentrations in Hemodialysis Patients?

Sunny Eloot*, Wim Van Biesen, Griet Glorieux, Nathalie Neirynck, Annemieke Dhondt, Raymond Vanholder

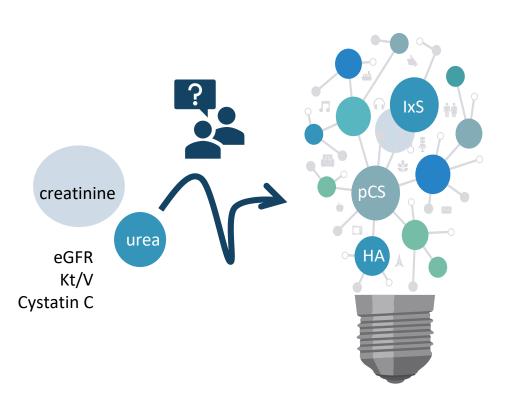
Nephrology Section, Department of Internal Medicine, Ghent University Hospital, Gent, Belgium

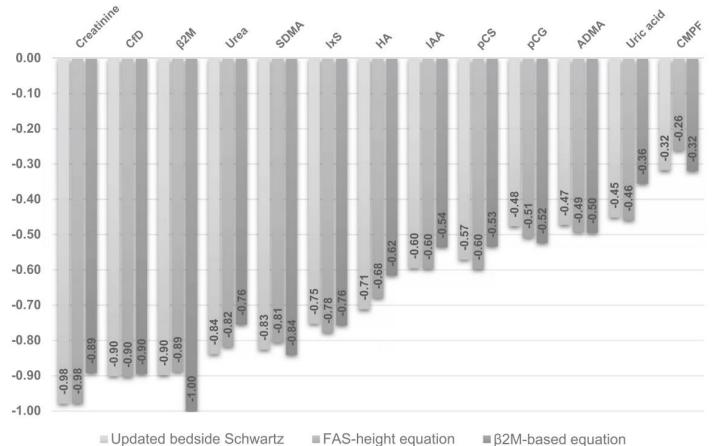


Abstract

Hemodialysis aims at removing uremic toxins thus decreasing their concentrations. The present study investigated whether Kt/Vurea, used as marker of dialysis adequacy, is correlated with these concentrations. Predialysis blood samples were taken before a midweek session in 71 chronic HD patients. Samples were analyzed by colorimetry, HPLC, or ELISA for a broad range of uremic solutes. Solute concentrations were divided into four groups according to guartiles of Kt/Vurea, and also of different other parameters with potential impact, such as age, body weight (BW), Protein equivalent of Nitrogen Appearance (PNA), Residual Renal Function (RRF), and dialysis vintage. Dichotomic concentration comparisons were performed for gender and Diabetes Mellitus (DM). Analysis of Variance in guartiles of Kt/Vurea did not show significant differences for any of the solute concentrations. For PNA, however, concentrations showed significant differences for urea (P<0.001), uric acid (UA), p-cresylsulfate (PCS), and free PCS (all P<0.01), and for creatinine (Crea) and hippuric acid (HA) (both P<0.05). For RRF, concentrations varied for β_2 -microglobulin (P<0.001), HA, free HA, free indoxyl sulfate, and free indole acetic acid (all P<0.01), and for p-cresylglucuronide (PCG), 3-carboxy-4-methyl-5-propyl-2-furanpropionic acid (CMPF), free PCS, and free PCG (all P<0.05). Gender and body weight only showed differences for Crea and UA, while age, vintage, and diabetes mellitus only showed differences for one solute concentration (UA, UA, and free PCS, respectively). Multifactor analyses indicated a predominant association of concentration with protein intake and residual renal function. In conclusion, predialysis concentrations of uremic toxins seem to be dependent on protein equivalent of nitrogen appearance and residual renal function, and not on dialysis adequacy as assessed by Kt/V_{urea}. Efforts to control intestinal load of uremic toxin precursors by dietary or other interventions, and preserving RRF seem important approaches to decrease uremic solute concentration and by extension their toxicity.

Current markers are poor predictors of overall uremic toxin accumulation





Alternative strategies to decrease uremic toxicity?

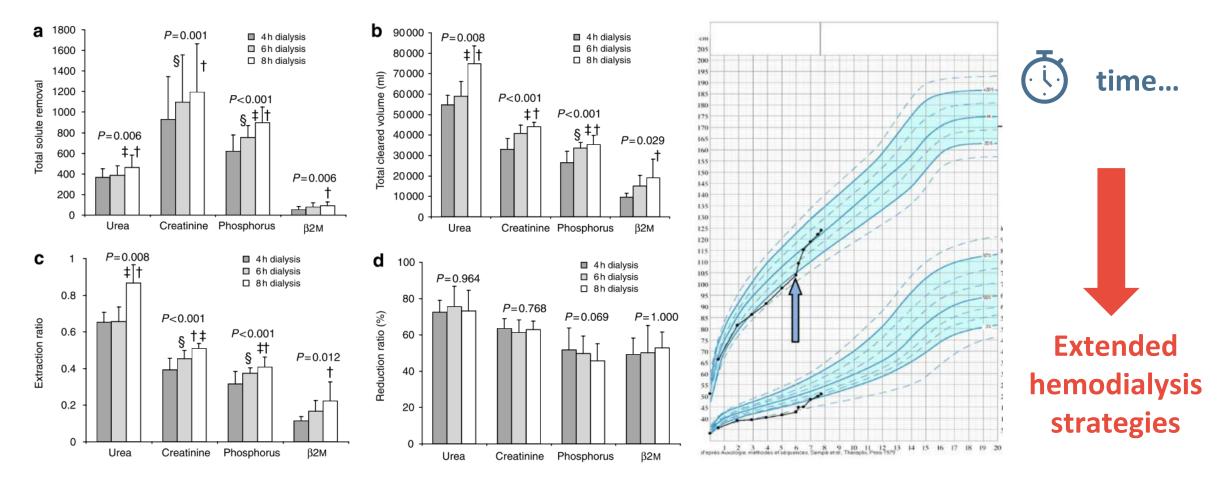


Figure 2 | **Removal parameters.** (a) Total solute removal (mg, except for urea in 0.1 g), (b) total cleared volume (ml), (c) dialyzer extraction ratio, (d) and reduction ratio (%) of urea, creatinine, phosphorus, and β 2-microglobulin for the 4, 6, and 8 h dialysis session.

Alternative strategies to decrease uremic toxicity?

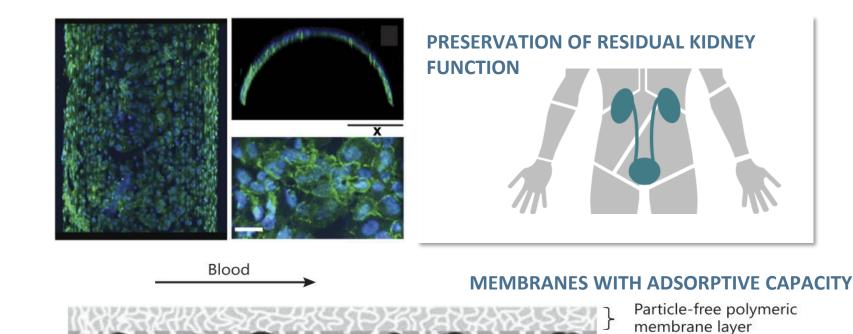
Polymeric

membrane matrix

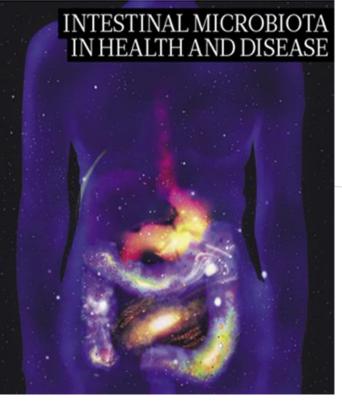
≻ MMM layer

Embedded activated

carbon particle



natureINSIGHT



DIETARY FIBER, PRE-, PRO, and SYNBIOTICS

а

Dialysate

Jansens et al. 2016; Tijink et al. 2014



Question 1: Which statement is true?

- A. Solute transport at the dialyzer is solely dependent on Q_d and Q_b
- B. Solute removal at the dialyzer is the only determinant of uremic solute removal
- c. With even the most recent advances, it seems that small solute removal in the dialyzer is close to its optimum.
- D. The main removal strategy of protein-bound uremic toxins is convection

Answers Question 1: Which statement is true?

- A. Solute transport at the dialyzer is solely dependent on Q_d and Q_b + membrane
- B. Solute removal at the dialyzer is the only determinant of uremic solute removal
 FALSE + Patient kinetics
- c. With even the most recent advances, it seems that small solute removal in the dialyzer is close to its optimum.

D. The main removal strategy of protein-bound uremic toxins is convection

FALSE = Diffusion of free fraction

Question 2: Which statement is true?

- A. eGFR is a good predictor of overall uremic toxin accumulation
- B. Kt/V is a good marker of dialysis adequacy as it reflects overall uremic toxin removal
- c. Solute transport in the patient limits the increase of performance with the traditional dialytic approaches.
- Intercompartiment clearance (K₁₂) of urea and β2-microglobuline are similar, e.g. 800mL/min

Answers Question 2: Which statement is true?

- A. eGFR is a good predictor of overall uremic toxin accumulation Only partially!
- B. Kt/V is a good marker of dialysis adequacy as it reflects overall uremic toxin removal
 FALSE
- c. Solute transport in the patient limits the increase of performance with the traditional dialytic approaches.
- D. Intercompartiment clearance (K_{12}) of urea and β 2-microglobuline are similar, e.g. 800mL/min β 2M = 50mL/min, only 6% of K_{12} of urea



TEAMWORK DIVIDES THE TASK & DOUBLE THE SUCCESS

Mentors & Promotors

Sunny Eloot Johan Vande Walle Wim Van Biesen Ann Raes Griet Glorieux Raymond Vanholder Rukshana Shroff

Lab & data

Sophie Lobbestael Tom Mertens Maria Vanlandschoot Sofie Vermeiren Els Holvoet Sanne Roels An Desloovere Elke De Bruyne

Charlotte Vanherzeele Christel Beerens Nathalie Polfliet Katty Van Cauwenberghe Katya Degroote Karlien Dhondt Members of advisory board

Collaborators

Elena Levtchenko Nathalie Godefroid Koen Van Hoeck Laure Collard Chiodini Benedetta Rukshana Shroff

& their local team

All patients and families participating







THANK YOU!





