

Determinants of serum levels of surfactant proteins A and B and Clara cell protein CC16

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Increased leakage of surfactant proteins A and B (SP-A and SP-B) and Clara cell secretory protein (CC16) from the air spaces into the circulation occurs in a range of respiratory conditions. However, circulating levels depend not only on the rate of entry into the circulation, but also on the rate of clearance. In order to clarify the role of the kidney in the clearance of these proteins, serum levels were related to markers of glomerular filtration in 54 non-smoking patients with varying degrees of renal dysfunction, none of whom had respiratory disease or were receiving dialysis at the time of sampling. Serum SP-A was related to SP-B ($r = 0.53$, $p < 0.001$) and to CC16 ($r = 0.33$, $p < 0.02$). Similarly, SP-B was related to CC16 ($r = 0.39$, $p < 0.004$). Stepwise multiple linear regression analysis suggested that serum SP-A and SP-B are influenced by age (~ 20 and $\sim 25\%$ of variance, respectively), whereas CC16 is determined by renal function and, to a lesser extent, by body weight ($\sim 63\%$ of variance in total). We conclude that CC16 is cleared from blood by the renal route, whereas SP-A and SP-B are not. Serum SP-A and SP-B are influenced by age, which we speculate reflects increased damage to the alveolocapillary barrier.

Keywords: pneumoproteinaemia, alveolocapillary permeability, peripheral markers, plasma protein clearance.

Introduction

The lung epithelium secretes several specific proteins into the air spaces of the respiratory tract. Among these are the major secretory product of Clara cells, the 16 kDa bronchiolar Clara cell protein (CC16), and the alveolar surfactant-associated proteins A to D. Although it is not known whether surfactant protein (SP) C occurs in blood, small amounts of CC16, SP-A, SP-B and SP-D do. Whereas SP-D is expressed by a number of tissues (Akiyama *et al.* 2002), expression of CC16 and SP-A are largely, and of SP-B is exclusively, confined to the lungs (Hermans and Bernard 1999, Akiyama *et al.* 2002). Consequently, the occurrence of CC16, SP-A and SP-B in the circulation can only be explained by leakage into the vascular compartment. Although the exact mechanisms by which

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these proteins enter the blood remains poorly understood, their concentrations in serum, referred to as 'pneumoproteinaemia', could potentially serve as early and highly sensitive peripheral markers for cellular toxicity and epithelial leakage at different levels of the respiratory tract. Intravascular leakage of the proteins increases in conditions characterized by pulmonary inflammation and/or pulmonary epithelial injury (Kuroki *et al.* 1993, Abe *et al.* 1995, Nomori *et al.* 1995, Lesur *et al.* 1996, Okamoto *et al.* 1997, Doyle *et al.* 1997, Carbonnelle *et al.* 2002, Bernard *et al.* 2003). Moreover, we have recently shown that, whereas tobacco smoking increases serum SP-B (Robin *et al.* 2002), it decreases serum CC16, consistent with increased alveolocapillary leakage and the known toxic effects of tobacco smoke on Clara cells (Bernard *et al.* 1992).

However, circulating levels of pneumoproteins will depend not only on the rate of their entry into the circulation, but also on the rate of their plasma clearance. We have previously shown that in patients with acute lung injury (ALI), levels of CC16, SP-A and SP-B are lower in mixed venous than in arterial plasma, indicating that the proteins are rapidly cleared from the circulation (Doyle *et al.* 1998). The kidney normally accounts for 30–80% of the plasma clearance of low molecular weight proteins (Maack *et al.* 1979). In ALI patients we found that the serum clearance of CC16 was related to that of creatinine, whereas that of SP-A and SP-B was not. Moreover, the arterial–venous gradient for SP-A and SP-B was directly related to their arterial levels, suggesting that temporal increases in their serum levels are due to enhanced alveolocapillary permeability that surpass any increase in plasma clearance. Our findings suggest that, at least in ALI patients, the concentration of CC16 in the vascular compartment is influenced by the glomerular filtration rate (GFR), whereas the levels of SP-A and SP-B reflect flux from the lung that is largely independent of this variable.

Plasma protein clearance is greatly altered in patients with ALI as systemic disease is frequently present (Doyle *et al.* 1998). In order to further elucidate the factors influencing the levels of pneumoproteins in the circulation and to clarify the role of the kidney in the plasma clearance of these proteins, we have related the levels of the pneumoproteins in the vascular compartment to glomerular filtration and to the levels of two other low molecular weight proteins both known to be cleared by a renal route (Donadio *et al.* 2001) – β_2 -microglobulin (β_2 M), an 11.8 kDa ubiquitous protein (Kin *et al.* 1977, Riska *et al.* 1982), and retinol-binding protein (RBP), a 21 kDa liver protein (Rask *et al.* 1980) – in 55 non-smoking patients with varying degrees of renal dysfunction, none of whom had respiratory disease or were receiving dialysis at the time of sampling.

Material and methods

Study population

The protocol of the study was approved by the Ethical Committee of the Faculty of Medicine, Université Catholique de Louvain, Brussels, Belgium. A single blood sample was drawn from the antecubital vein of 17 non-smoking females with a median age of 53 years (range 28–83 years) and 37 non-smoking males with a median age of 51 years (range 16–87 years) with varying degrees of chronic renal failure, none of whom were yet receiving dialysis. An additional blood sample was taken on a separate occasion approximately 4 weeks later from one of each group of patients with diabetic nephropathy, nephrosclerosis and chronic glomerulonephritis.

Analytical methods

Samples were assayed in a blind, randomized manner.

Glomerular filtration. Serum creatinine was measured using Jaffé's technique (Kabanda *et al.* 1994), and its clearance corrected for age and weight was calculated by the Cockcroft formula and used as an index of renal function (Cockcroft and Gault 1976). Creatinine clearance in women was corrected by multiplying by 0.72.

CC16, β_2 M and RBP. These proteins were all measured using latex immunoassay (Bernard *et al.* 1991, Kabanda *et al.* 1994). Briefly, to avoid possible interference by complement, rheumatoid factor or chylomicrons, the serum samples were pretreated by heating at 56°C for 30 min and the addition of polyethylene glycol (16%, v/v, 1:1) and trichloroacetic acid (10%, v/v, 1:40). After overnight precipitation at 4°C, the samples were centrifuged (2000 *g* for 10 min). CC16, RBP and β_2 M were determined in the supernatants using an automatic agglutination technique with purified standards and antibodies from Dakopatts (Glostrup, Denmark). All samples were analysed in duplicate at two different dilutions. The assays have an average analytical recovery of 95%, with intra- and inter-assay coefficients of variation ranging from 5–10% (Bernard and Lauwerys 1983, Bernard *et al.* 1991).

SP-A and SP-B. SP-A and SP-B were measured using inhibition enzyme-linked immunosorbent assays (ELISAs) (Doyle *et al.* 1997). Briefly, SP-A and SP-B were first freed from any associated components using ethylene diamine tetra-acetic acid (EDTA), sodium dodecyl sulphate (SDS) and Triton X-100. The proteins were then measured with inhibition ELISAs using polyclonal antibodies raised against alveolar proteinosis-derived SP-A and mature SP-B. Absorbance was measured at 405 nm using a Dynatech MR5000 reader (Dynatech Laboratories, Chantilly, Virginia, USA). All samples were assayed in duplicate at four serial dilutions. To compensate for differences in incubation times inherent when manually pipetting across the ELISA plate, each plate included two standard curves, one positioned at the beginning of the plate and one at the end, comprising eight serial dilutions in duplicate (SP-A, 1.95–250.00 ng ml⁻¹; SP-B, 7.8–1000.0 ng ml⁻¹; $r > 0.99$). AssayZap software (Biosoft, Ferguson, Missouri, USA) was used to generate standard curves and to weight the analysis of each sample according to its position within the ELISA plate.

The SP-A assay has intra- and inter-assay coefficients of variation (CVs) of <7% and 10%, respectively (Doyle *et al.* 1994). The SP-B assay has intra- and inter-assay CVs of <9 and 10%, respectively, as determined by repeated measures ($n = 16$) of four plasma samples ranging in concentration from 11.7 to 81.0 µg ml⁻¹. The detection limit of the assay is 4.1 ng ml⁻¹ (Ekins 1991) ($n = 30$), with an analytical recovery (\pm SD) of $105 \pm 27\%$ as determined by repeated 'spiking' of the same four plasma samples with 0.75, 1.5, 3.0 and 6.0 µg ml⁻¹ of SP-B.

Statistics

Since serum CC16, SP-A and SP-B levels are not normally distributed (Doyle *et al.* 1997, Robin *et al.* 2002), all analyses were made using log-transformed data. The mean and 95% confidence intervals were determined using log-transformed data and then back-transformed. The independent samples *t*-test was used for all comparisons. The associations between the measured variables were tested using the Pearson correlation test.

Multiple linear regression analysis of the serum variables. The levels of serum CC16, SP-A and SP-B must be determined by both their rate of entry into, and clearance from, the circulation. The kidney normally accounts for the majority of plasma clearance of low molecular weight proteins (Maack *et al.* 1979). The GFR is influenced by body weight and is reflected in the serum creatinine clearance. In addition, there is an age- and gender-related influence on the level of many circulating antigens, including CC16, SP-A and SP-B (Robin *et al.* 2002). Therefore, we hypothesized that in subjects without lung injury, the independent variables influencing serum β_2 M, RBP, CC16, SP-A and SP-B would include age, body weight, serum creatinine clearance, gender and total protein level. All data was log-transformed for inclusion in the stepwise backwards regression analysis model, where independent variables with *p* values <0.05 were sequentially excluded.

Results

Renal function

Patients with diabetic nephropathy and nephrosclerosis were older than patients with chronic glomerulonephritis ($p = 0.005$ and 0.000 , respectively), chronic

interstitial nephritis ($p = 0.010$ and 0.004) or polycystic kidney disease ($p = 0.060$ and 0.015), although there were no other statistical differences between the groups with regards to indices of renal function (table 1).

Gender

Even though serum creatinine was similar in the male and female subjects, serum creatinine clearance was $\sim 60\%$ higher in males (table 2). Creatinine clearance was ~ 4 -fold lower than normal (120 ml min^{-1}) (Cockcroft and Gault 1976) in the male subjects and ~ 7 -fold lower in the females. CC16 was $\sim 45\%$ less in the male subjects. The males subjects were $\sim 24\%$ heavier than the females. There were no other gender-based differences in any of the other determinants.

Serum variables and indices of renal and lung function

Whereas serum $\beta_2\text{M}$, RBP and CC16 were all strongly related to serum creatinine and inversely related to its serum clearance ($p = 0.000$), SP-A and SP-B were not (table 3). Serum creatinine clearance was also related to body weight and inversely related to age.

Pneumoprotein determinates

The serum levels of the pneumoproteins were strongly co-related (table 3). However, whereas serum CC16 was also strongly related to $\beta_2\text{M}$ and RBP ($p = 0.000$), SP-A and SP-B were not. $\beta_2\text{M}$, CC16 and RBP was positively influenced by serum creatinine clearance and, to a lesser extent, by body weight (table 4). RBP was also weakly influenced by age. Notably, SP-A and SP-B were only influenced by age.

Discussion

Although the alveolocapillary barrier is normally extraordinarily effective at partitioning the plasma from the alveolar epithelial lining fluid, increased bidirectional protein flux occurs across the barrier in conditions characterized by pulmonary inflammation and/or pulmonary epithelial injury. The determination of the levels of pneumoproteins in the vascular compartment represents a new and original approach for the non-invasive assessment of the integrity of the lung epithelium. However, circulating levels depend not only on the rate of entry into the circulation, but also the rate of clearance. Of particular significance is the kidney, which is the major organ responsible for the plasma clearance of low molecular weight proteins. Our previous work has suggested that, at least in so far as patients with ALI are concerned, whereas plasma clearance is a determinant of CC16 levels, SP-A and SP-B levels reflect lung function independent of this variable. The present study in patients with varying degrees of renal dysfunction, none of whom had respiratory disease or were receiving dialysis at the time of sampling, confirms that the kidney is an important determinant of intravascular levels of CC16, but is not a determinant of circulating levels of SP-A or SP-B.

Table 1. Indices of renal function in the study population.

Cause of chronic renal failure	<i>n</i>	Age (years)		Body weight (kg)		Creatinine clearance (ml min ⁻¹)		Total serum protein (g dl ⁻¹) ^a		Serum creatinine (mg dl ⁻¹) ^a		Serum RBP (µg ml ⁻¹) ^a		Serum β ₂ M (µg ml ⁻¹) ^a	
		Mean	95% CI	Mean	95% CI	Mean	95% CI	Mean	95% CI	Mean	95% CI	Mean	95% CI	Mean	95% CI
Diabetic nephropathy	8	63	52–76	86	76–98	27.3	10.1–74.0	6.7	5.7–7.9	3.0	1.6–5.6	101	74–141	9.1	4.4–18.9
Chronic glomerulonephritis	20	43	30–73	73	49–110	25.6	9.5–69.5	6.7	5.9–7.6	3.5	1.1–11.1	135	66–278	8.3	3.6–18.9
Chronic interstitial nephritis	12	37	16–83	64	46–91	30.5	14.4–67.7	6.8	5.8–7.9	2.6	1.1–6.0	82	38–176	5.3	1.2–22.8
Polycystic kidney disease	6	50	36–70	76	58–99	22.4	7.0–71.3	7.1	6.6–7.6	4.4	2.4–7.8	106	88–127	9.0	4.5–18.2
Nephrosclerosis	7	68	57–80	76	65–89	18.7	4.1–85.3	6.7	6.5–7.0	4.0	2.0–8.0	89	42–189	9.8	3.9–24.5
Multiple myeloma	1	77		70		22.9		6.4		2.7		88		11.9	
Hereditary	1	28		47		19.9		6.5		3.1		92		9.7	
Polyarteritis nodosa	1	79		65		11.9		6.1		4.7		100		15.3	
Wegener's granulomatosis	1	63		82		28.6		6.5		3.1		64		6.3	
Unknown	1	87		73		13.1		5.4		4.1		91		8.9	

95% CI, 95% confidence interval.

The patients with polyarteritis nodosa and Wegener's granulomatosis had no lung involvement of their disease.

^a The reference intervals used were as follows: total protein, 6.0–8.0 g dl⁻¹; serum creatinine, 0.5–1.2 mg dl⁻¹; serum RBP, 30–60 µg ml⁻¹; serum β₂M, 1.0–3.0 µg ml⁻¹ (from Dade Behring Diagnostics Pty Ltd).

Table 2. Variables measured in the study population.

	Females (n = 17)		Males (n = 37)		p value
	Mean	95% CI	Mean	95% CI	
Age (years)	52	32–83	49	30–80	0.626
Body weight (kg)	63	41–98	78	62–100	0.001
Creatinine clearance (ml min ⁻¹)	18.1	6.2–53.2	28.9	11.2–74.5	0.008
Total serum protein (g dl ⁻¹)	6.7	5.9–7.7	6.7	5.7–7.9	0.971
Serum creatinine (mg dl ⁻¹)	3.5	1.6–7.6	3.3	1.3–8.3	0.691
Serum RBP (µg ml ⁻¹)	115	64–206	100	50–198	0.226
Serum β ₂ M (µg ml ⁻¹)	10.4	4.8–22.8	7.0	2.4–21.0	0.046
Serum CC16 (ng ml ⁻¹)	73.8	24.7–220.8	43.2	7.5–247.2	0.020
Serum SP-A (ng ml ⁻¹)	381	162–897	360	156–832	0.630
Serum SP-B (ng ml ⁻¹)	3573	1318–9682	3304	1175–9290	0.617

95% CI, 95% confidence interval.

Study design and validation of methods

Pneumoproteins. The assays employed were validated using independently sourced monoclonal and polyclonal (Bernard and Lauwerys 1983, Bernard *et al.* 1991, Doyle *et al.* 1995, 1997) antibodies. Our CC16 assay includes an initial precipitation step to eliminate serum lipids likely to interfere with its determination (Hermans *et al.* 1998). Similarly, our assay procedure for SP-A and SP-B incorporates EDTA to disrupt the Ca²⁺–lipid bridges, and SDS and Triton-X 100 to dissociate the proteins and free them from any associated components. We have recently analysed plasma from an infant independently confirmed by two laboratories to have an autosomal recessive frame shift mutation at 121 bp in the SP-B gene that resulted in a premature stop codon and no SP-B mRNA or protein. Analysis of this definitive negative control by ELISA has confirmed the specificity of our assay.

In addition, given that the pneumoproteins share few structural similarities, it is difficult to conceive that the strong correlations we observed with our serum determinations are fortuitous or influenced by non-specific reactivities.

Indices of renal function. Serum creatinine was used as an index of renal function. It is known that serum creatinine is influenced by other factors; indeed, in the present study serum creatinine was slightly lower in females than males and was inversely related to age. We believe that these small differences probably reflect variations in muscle mass rather than renal function. Although the use of single serum creatinine measurements to determine creatinine clearance is limited when renal function changes acutely, all of our patients had stable chronic renal failure in which such changes are unlikely. Indeed, in the three patients where blood was sampled on a second occasion approximately 4 weeks later, creatinine clearance, as estimated by this method, remained similar to that on the previous occasion, and serum CC16, SP-A and SP-B levels were essentially unchanged. As a further precaution, we also measured RBP and β₂M as additional indices of renal function. As with CC16, our assays for RBP and β₂M also included an initial precipitation step to prevent interference by serum lipids. Although we did not correct for multiple comparisons and partial correlation coefficients were not applied, the

Table 3. CROSS CORRELATION BETWEEN THE MEASURED VARIABLES

	β 2M	CC16	Creat Clear	Creat	RBP	SP-A	SP-B	Protein	Body Weight
Age	0.31 0.017	0.31 0.019	−0.33 0.013	0.15 0.266	0.10 0.438	0.45 0.001	0.50 0.000	−0.26 0.050	0.30 0.024
β 2M		0.77 0.000	−0.86 0.000	0.82 0.000	0.58 0.000	0.26 0.058	0.34 0.010	−0.09 0.496	−0.20 0.144
CC16			−0.78 0.000	0.73 0.000	0.47 0.000	0.33 0.014	0.39 0.003	0.00 0.970	−0.18 0.173
Creat				−0.88 0.000	−0.49 0.000	−0.09 0.518	−0.35 0.010	0.04 0.758	0.42 0.001
Clear					0.58 0.000	0.08 0.575	0.19 0.168	0.06 0.644	−0.09 0.499
Creat						0.21 0.119	0.13 0.331	0.10 0.455	−0.00 0.992
RBP							0.53 0.000	−0.07 0.615	0.31 0.023
SP-A								0.24 0.075	−0.01 0.967
SP-B									−0.13 0.329
Protein									

Cross correlation between the measured variables where the “r” value is given in the top and the “p-value” in the bottom line. Relations where $p < 0.01$ are underlined. Legend as in Table 1.

Table 4. Determinants of serum proteins.

β_2 M	F = 91.0, $p = 0.0000$, $r^2 = 0.77$ Creatinine clearance: $p = 0.0000$, $\beta = -0.94$ Body weight: $p = 0.0070$, $\beta = 0.20$
RBP	F = 8.8, $p = 0.0001$, $r^2 = 0.33$ Creatinine clearance: $p = 0.0000$, $\beta = -0.73$ Age: $p = 0.0757$, $\beta = -0.25$ Body weight: $p = 0.0104$, $\beta = 0.38$
CC16	F = 45.9, $p = 0.0000$, $r^2 = 0.63$ Creatinine clearance: $p = 0.0000$, $\beta = -0.85$ Body weight: $p = 0.0602$, $\beta = 0.18$
SP-A	Age: $r^2 = 0.20$ F = 13.1, $p = 0.0007$, $\beta = 0.45$
SP-B	Age: $r^2 = 0.25$ F = 17.1, $p = 0.0001$, $\beta = 0.50$

We hypothesized that independent variables influencing serum β_2 M, CC16, RBP, SP-A and SP-B would include age, body weight, serum creatinine clearance, gender and total protein.

cross correlations between our indices of renal function (serum creatinine, serum creatinine clearance, RBP and β_2 M) were very strong ($p = 0.000$), further strengthening our findings.

β_2 M has a diagnostic accuracy very similar to that of creatinine, whereas RBP has been purported to be a less adequate marker of glomerular filtration (Donadio *et al.* 2001), possibly because its serum level varies with hormonal fluctuations and nutritional status. Consistent with this, β_2 M and RBP were both predicted by serum creatinine clearance, our index of GFR, but to different extents. Whereas multiple linear regression analysis suggests that GFR, as reflected by serum creatinine clearance, and body weight together accounted for $\sim 77\%$ of the serum β_2 M, RBP was also influenced by age, which together accounted for only $\sim 33\%$ of its serum level. Indeed, whereas CC16 (Robin *et al.* 2002) and β_2 M (Donadio *et al.* 2001) were elevated approximately in proportion to the decrease in GFR (~ 5 -fold), RBP (Donadio *et al.* 2001) was increased by a much lesser amount (~ 2 -fold), consistent with our own (Doyle *et al.* 1998) and other observations (Maack *et al.* 1979, Kabanda *et al.* 1994) that, when the GFR is reduced, some proteins that would otherwise cleared by the kidney are removed by unknown pathways that may partially, or even totally, compensate for the reduced renal protein clearance.

The kidney as an organ of serum protein clearance

The filtration of proteins across the renal glomerular capillary membrane occurs with an approximate log-log relationship to the molecular weight (Maack *et al.* 1979, Kabanda *et al.* 1995, Hermans *et al.* 1998). Although steric considerations also play a major role, most proteins smaller than 50 kDa normally pass relatively freely through the glomerulus, and virtually all (95–99%) are taken up and catabolized by fusion with lysosomes in the heterolysosomes of proximal tubular cells (Guder 1988, Kabanda *et al.* 1995).

Consistent with this, CC16 was strongly predicted by serum creatinine clearance, our index of glomerular filtration. In contrast, and consistent with our findings in ALI patients (Doyle *et al.* 1998), SP-A and SP-B levels were not predicted by serum creatinine clearance. In blood, native SP-A appears to be complexed with immunoglobulin G (IgG) and immunoglobulin M (IgM) (Kuroki

et al. 1993, Doyle *et al.* 1995). It is possible that such complexes are too large to be filtered in the glomerulus. It is not apparent why SP-B is not cleared by the renal route, since the immunoreactive forms detected in plasma, at least under denaturing electrophoretic conditions, are sufficiently small ($\sim 18\text{--}45\text{kDa}$) (Doyle *et al.* 1997). One possibility is that the hydrophobicity of SP-B is such that it complexes with serum lipids, and the derived complexes are too large to be filtered through the glomerulus.

Systemic clearance of SP-A and SP-B

The pathway by which SP-A and SP-B are cleared from the circulation remains controversial. SP-A may be phagocytosed through binding directly to specific receptor(s) found on macrophages (Chroneos *et al.* 1996), monocytes (Geertsma *et al.* 1994) and neutrophils (Hartshorn *et al.* 1998), or through Fc-receptor-mediated phagocytosis in association with complexed IgG and IgM. Although it is not known whether SP-B specifically binds to phagocytes, SP-B is phagocytosed by alveolar macrophages (Phelps and Floros 1991). Moreover, infants deficient in SP-B die of alveolar proteinosis (deMello and Lin 2001), probably caused by impaired alveolar macrophage clearance of surfactant lipids, while airway administration of SP-B-containing vesicles can enhance the induction of IgM-mediated systemic immunity (van Iwaarden *et al.* 2001). Systemic phagocytes may also clear SP-B.

The effect of age

The present study confirms our recent findings that CC16, SP-A and SP-B increase with age. The levels of CC16, SP-A and SP-B in our patients are similar to those recently reported by us in healthy, aged-matched subjects (Robin *et al.* 2002) and were $\sim 50\%$ higher than we previously reported in subjects aged on average 25 years younger (Bernard *et al.* 1994, Hermans and Bernard 1999). Age-related structural and functional changes in the lung have been previously described. Gas trapping increases with age (Lee *et al.* 2000), as do air space abnormalities (Soejima *et al.* 2000) and diameter (Zeman and Bennett 1995), whereas lung compliance and shear modulus decrease (Lai-Fook and Hyatt 2000) in association with increased lung elastin content (Foster and Curtiss 1990). Moreover, whereas the internal surface area of the lung increases to a maximum of $\sim 80\text{ m}^2$ at the age of ~ 20 years, it then decreases $\sim 2.7\text{ m}^2$ per decade thereafter (Thurlbeck 1967). Finally, increased albumin levels have been reported in bronchoalveolar lavage fluid from older subjects (Roberts *et al.* 1993). We have no information concerning the effect of age on the alveolar levels of SP-A or SP-B. However, since CC16 mRNA expression is not related to age (Jensen *et al.* 1994), we suggest that the age-related pneumoproteinaemia in non-smokers reflects increased alveolocapillary leakage and the effects of non-specific deterioration of the lung.

Conclusion

Our findings suggest that, whereas serum CC16 is influenced by GFR, serum SP-A and SP-B levels are largely independent of this variable. Serum SP-A and SP-B are influenced by age, which we speculate reflects increased damage to the alveolocapillary barrier with ageing.

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