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C-Terminal Fragment of Agrin (CAF): A Novel Marker for Progression of Kidney Disease in Type 2 Diabetics

Devetzis, Vasilios; Daryadel, Arezoo; Roumeliotis, Stefanos; Theodoridis, Marios; Wagner, Carsten A; Hettwer, Stefan; Huynh-Do, Uyen; Ploumis, Passadakis; Arampatzis, Spyridon

Abstract: BACKGROUND Diabetes is the leading cause of CKD in the developed world. C-terminal fragment of agrin (CAF) is a novel kidney function and injury biomarker. We investigated whether serum CAF predicts progression of kidney disease in type 2 diabetics. METHODS Serum CAF levels were measured in 71 elderly patients with diabetic nephropathy using a newly developed commercial ELISA kit (Neurotune®). Estimated glomerular filtration rate (eGFR) and proteinuria in spot urine were assessed at baseline and after 12 months follow up. The presence of end stage renal disease (ESRD) was evaluated after 24 months follow-up. Correlation and logistic regression analyses were carried out to explore the associations of serum CAF levels with GFR, proteinuria, GFR loss and incident ESRD. Renal handling of CAF was tested in neurotrypsin-deficient mice injected with recombinant CAF. RESULTS We found a strong association of serum CAF levels with eGFR and a direct association with proteinuria both at baseline ($r = 0.698$, $p < 0.001$ and $r = 0.287$, $p = 0.02$) as well as after 12 months follow-up ($r = 0.677$, $p < 0.001$ and $r = 0.449$, $p < 0.001$), respectively. Furthermore, in multivariate analysis, serum CAF levels predicted eGFR decline at 12 months follow-up after adjusting for known risk factors (eGFR, baseline proteinuria) [OR (95%CI) = 4.2 (1.2-14.5), $p = 0.024$]. In mice, injected CAF was detected in endocytic vesicles of the proximal tubule. CONCLUSION Serum CAF levels reflect renal function and are highly associated with eGFR and proteinuria at several time points. Serum CAF was able to predict subsequent loss of renal function irrespective of baseline proteinuria in diabetic nephropathy. CAF is likely removed from circulation by glomerular filtration and subsequent endocytosis in the proximal tubule. These findings may open new possibilities for clinical trial design, since serum CAF levels may be used as a selection tool to monitor kidney function in high-risk patients with diabetic nephropathy.

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1 **C-terminal fragment of agrin (CAF): a novel marker for progression of kidney disease in type 2**

2 **Diabetics**

3

4 Vasilios Devetzis¹, Arezoo Daryadel², Stefanos Roumeliotis³, Marios Theodoridis³, Carsten A.
5 Wagner², Stefan Hettwer⁴, Uyen Huynh-Do¹, Passadakis Ploumis², Spyridon Arampatzis¹

6

7 ¹ Department of Nephrology, Hypertension and Clinical Pharmacology, Inselspital, Bern University
8 Hospital, Bern, Switzerland

9 ² Institute of Physiology, University of Zurich, Switzerland

10 ³ Department of Nephrology, University Hospital of Alexandroupoli, Alexandroupoli, Greece

11 ⁴ Neurotune AG, Schlieren, Switzerland

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18

19 * Corresponding author: Spyridon Arampatzis, MD

20 spyridon.arampatzis@insel.ch

21 Department of Nephrology, Hypertension and Clinical Pharmacology

22 Inselspital, University Hospital Bern

23 3010 Switzerland

24 Tel: + 41 31 632 31 44

25 **Abstract**

26 **BACKGROUND:** Diabetes is the leading cause of CKD in the industrialized world. C-terminal
27 fragment of agrin (CAF) is a novel kidney function and injury biomarker. We investigated whether
28 serum CAF predicts progression of kidney disease in type 2 diabetics.

29 **METHODS:** Serum CAF levels were measured in 71 elderly patients with diabetic nephropathy
30 using a newly developed commercial ELISA kit (Neurotune®). Estimated glomerular filtration rate
31 (eGFR) and proteinuria in spot urine were assessed at baseline and after 12 months follow up. The
32 presence of end stage renal disease (ESRD) was evaluated after 24 months. Correlation and logistic
33 regression analyses were carried out to explore the associations of serum CAF levels with GFR,
34 proteinuria, GFR loss and incident ESRD. Renal handling of CAF was tested in neurotrypsin-
35 deficient mice injected with recombinant CAF.

36 **RESULTS:** We found a strong association of serum CAF levels with eGFR and a direct association
37 with proteinuria both at baseline ($r=0.698$, $p<0.001$ and $r=0.287$, $p=0.02$) as well as after 12
38 months follow-up ($r=0.677$, $p<0.001$ and $r=0.449$, $p<0.001$), respectively. Furthermore, in
39 multivariate analysis, serum CAF levels predicted eGFR decline at 12 months follow-up after
40 adjusting for known risk factors (eGFR, baseline proteinuria) [OR(95%CI) 4.2(1.2-14.5), $p=0.024$]. In
41 mice, injected CAF was detected in endocytic vesicles of the proximal tubule.

42 **CONCLUSION:** Serum CAF levels reflect renal function and are highly associated with eGFR and
43 proteinuria at several time points. Serum CAF was able to predict subsequent loss of renal
44 function irrespective of baseline proteinuria in diabetic nephropathy. CAF is likely removed from
45 circulation by glomerular filtration and subsequent endocytosis in the proximal tubule. These
46 findings may open new possibilities for clinical trial design, since serum CAF levels may be used as
47 a selection tool to monitor kidney function in high risk patients with diabetic nephropathy.

48

49 **Introduction**

50 Chronic kidney disease (CKD) represents a global public health problem affecting more than 1 in
51 10 adults worldwide [1]. Diabetes is the leading cause of CKD in the industrialized world and
52 people with both diabetes and chronic kidney disease have a greatly increased risk of all-cause
53 mortality, cardiovascular mortality, and end-stage renal disease (ESRD) [2]. Although several
54 factors have been identified to predict risk of developing progressive nephropathy in diabetic
55 patient populations, none sufficiently predict the risk for individual patients [3].

56 Currently the simultaneous evaluation of albuminuria and glomerular filtration rate (GFR) is
57 recommended by the Kidney Disease Improving Global Outcomes (KDIGO) guidelines for the
58 prediction of progression in diabetic nephropathy [4]. A growing body of evidence challenges the
59 traditional conceptual model of the course of diabetic nephropathy [5,6], since it can present with
60 a rapid decline of renal function and no overt albuminuria or progressive proteinuria [7,8,9,10].
61 Based upon these clinical observations, more reliable biomarkers are urgently needed in the clinic
62 to predict renal outcome in patients with early stages of CKD in diabetic nephropathy [11].

63 Agrin is the major heparin sulfate proteoglycan in the glomerular basement membrane and a
64 ubiquitous component of the extracellular matrix [12,13]. Neurotrypsin, a serine protease, cleaves
65 agrin at two distinct molecular sites generating a 110 kDa fragment (CAF110) at the alpha site,
66 whereas cleavage at the beta site produces the 22 kDa C-terminal fragment (CAF22) [14]. In
67 human urine, CAF22 can be detected, suggesting renal clearance for this small fragment [15,16].
68 Furthermore, serum CAF22 (sCAF) as kidney function biomarker has only recently been explored in
69 septic patients and in renal transplant recipients [17,18]. Both studies indicate that the sCAF
70 concentration is associated and comparable to established parameters of renal function such as
71 creatinine and cystatin C.

72 However, to date, there are no clinical studies which have investigated whether sCAF could serve
73 as biomarker in clinical practice for diabetic nephropathy and no animal studies addressing the
74 renal handling of sCAF. We hypothesize that rising sCAF levels may reflect progression of kidney
75 damage and dysfunction. In this prospective study in a cohort of patients with diabetic
76 nephropathy, we aimed to: 1) explore and validate the cross-sectional associations between sCAF
77 and the currently used clinical markers of kidney damage and dysfunction; estimated glomerular
78 filtration rate (eGFR) and proteinuria (protein to creatinine ratio [PCR]) and 2) examine the
79 independent predictive performance of sCAF for renal function decline and ESRD. In addition we
80 studied the renal handling of CAF in neurotrypsin deficient mice lacking endogenous CAF22
81 production.

82

83 **Methods**

84 *Study design and patient cohort*

85 The present study was designed as a prospective observational cohort study. Study subjects were
86 recruited from the outpatient clinic of Department of Nephrology, University Hospital of
87 Alexandroupoli, Greece. Patients were recruited if they fulfilled the following inclusion criteria; (i)
88 age > 18 years, (ii) ability to provide written, informed consent (iii) type 2 diabetes, defined as the
89 use of oral glucose-lowering treatment , a fasting plasma glucose >7.0 mmol/l (126 mg/dl) or non-
90 fasting plasma glucose >11.1 mmol/l (>200 mg/dl). The presence of diabetic nephropathy was
91 defined by microalbuminuria [albumin to creatinine ratio 3-30 mg/mmol (30-300mg/g)] or
92 persistent albuminuria [(albumin: creatinine ratio >30 mg/mmol (>300mg/g)] in three consecutive
93 measurements in sterile spot urine sample during a 6-month period, presence of diabetic
94 retinopathy, and no clinical or laboratory evidence of kidney or urinary tract disease [19]. Subjects
95 were considered to have diabetic retinopathy if they showed nonproliferative or proliferative

96 stages in funduscopy through dilated pupils or had a history of retinal laser surgery
97 (photocoagulation) for diabetic retinopathy.

98 The main exclusion criteria were clinical or laboratory evidence of non-diabetic nephropathy such
99 as active malignancy, infection, or auto-immune disease. The study was approved by the Ethics
100 Committee of the Scientific Council of the University Hospital of Alexandroupoli, was in
101 accordance with Helsinki Declaration of Human Rights and all subjects gave written informed
102 consent.

103 The diagnosis and staging of CKD was assessed based on the Clinical Practice Guidelines from the
104 National Kidney Foundation–Kidney Disease Outcomes Quality Initiative [20]. GFR was estimated
105 (eGFR) using the Chronic Kidney Disease Epidemiology Collaboration equation (CKD-EPI), which is
106 more accurate and less biased than the MDRD Study equation, especially in patients with higher
107 GFR, resulting in reduced misclassification of CKD [21]. The onset of ESRD was defined as the date
108 of first dialysis or transplantation.

109

110 *Biochemical analysis*

111 After an overnight fasting of 8 h blood samples were drawn from a peripheral vein of the patients
112 into vacutainer tubes without anticoagulant in order to obtain serum. Samples for fasting blood
113 glucose levels, potassium, sodium, calcium, phosphate, protein, albumin, urea, creatinine and
114 HbA1c were measured according to routine laboratory methods. Proteinuria was determined in
115 spot urine by an automated biochemistry analyzer (Clinical Chemistry System ADVIA 2400,
116 Siemens). Blood samples were immediately centrifuged at 4,000 rpm for 10 min at ambient
117 temperature, and the extracted serum was aliquoted and stored at 253 K (-20 °C) until use.

118

119 *C-terminal agrin fragments (CAF)*

120 CAF levels were measured using the NTtotalCAF ELISA kit from Neurotune AG
121 (www.neurotune.com). This assay detects both the large C-terminal agrin fragment CAF110
122 generated by alpha cleavage and the small C-terminal agrin fragment generated by beta cleavage
123 (CAF22). The molecular ratio of both fragments is 1:5 (CAF110 vs CAF22). In brief, serum samples
124 were diluted two-fold with sample incubation buffer and heated to 329 K (56°C) for 30 minutes.
125 Samples were diluted 20 fold with dilution buffer. 100 µL of diluted sample was added to an ELISA
126 plate precoated with a monoclonal catcher antibody raised against CAF. The plate was then
127 incubated for 16 hours at room temperature. After rinsing 3 times with washing buffer, the
128 monoclonal detector antibody was added for 30 minutes at room temperature. The plate was
129 then washed as previously described [6]. Streptavidin- conjugated HRP was then added for 30
130 minutes at room temperature. After washing, TMB substrate was added and developed for 30
131 minutes. The reaction was then stopped with an acidic solution and the resulting absorbance was
132 measured on an ELISA plate reader at 450 nm. The results were quantified using the calibration
133 curve prepared independently on each individual plate. The lower detection limit was 40 pmol/L.
134 Independent analysis revealed a mixed intra- and inter-assay %CV < 13 % for serum samples.

135

136 *Statistical analysis*

137 Variables were tested for normality using the Kolmogorov- Smirnov test. Comparisons between
138 categorical variables were performed by the chi- square test or the Fisher's exact test when
139 appropriate. Differences in continuous variables between two groups were assessed using the
140 Student's t-test, the Mann-Whitney's U-test or one-way ANOVA when appropriate. Pearson and
141 Spearman correlation coefficients were calculated as appropriate between CAF or logarithmic
142 transformed CAF (to achieve normal distribution), creatinine, GFR, proteinuria, age, BMI, HbA1c
143 and duration of T2DM at baseline and at 12 months follow-up. Correlation analysis of decline of

144 renal function (Δ GFR) and progression to ESRD with various risk factors was also conducted to
145 determine possible outcome predictors.

146 Multiple logistic regression analysis was carried out to further explore the predictive role of serum
147 CAF levels with GFR decline. The variables entered in the models were known risk factors
148 associated with progression of kidney disease. For both outcomes a basic and an advanced model
149 were used with the following variables:1) Basic model A: CAF, baseline GFR and baseline
150 proteinuria and 2) Advanced model B: CAF, baseline GFR, baseline proteinuria, age and BMI.
151 Variables retained in all final models were chosen according to clinical significance. A two-sided p
152 value < 0.05 was considered significant. The IBM SPSS Statistics 20.0 statistical software package
153 (SPSS Inc, Chicago, Illinois, USA) was used for all calculations.

154

155 *Animal experiments*

156 Experiments were performed in 8-10 weeks old male neurotrypsin knock-out mice (NT KO). The
157 generation and genotyping of neurotrypsin KO mice has been previously described [22]. All animal
158 experiments were conducted according to Swiss laws for the welfare of animals and were
159 approved by the local Zurich Veterinary Authority (Kantonales Veterinäramt Zürich). The animals
160 had free access to food and tap water.

161 Mice were pretreated 60 min before experiments with an injection of leupeptin (5 mg/mouse,
162 Sigma Aldrich, Buchs, Switzerland), an inhibitor of lysosomal degradation. Wildtype and
163 neurotrypsin KO mice were anesthetized by i.p. injection of xylazin and ketamin and were injected
164 with 100 μ l of a mixture containing recombinant hCAF22 (100 ng/mouse, produced by
165 Neurotune) dissolved in 0.9 % NaCl and FITC-labeled sinistrin (100 ng/ mouse). FITC-sinistrin is
166 cleared from circulation exclusively by glomerular filtration and served as control for successful
167 injection. Before injection of hCAF, the urinary bladder was completely emptied through a small

168 abdominal incision to allow collection of urine produced during the experimental period. Mice
169 received also a bolus i.p. of 0.5 ml of 25 mM NaHCO₃/150 mM NaCl to prevent dehydration and to
170 promote diuresis. Mice were kept warm at 37 °C by placing animals on a heating tablet for the rest
171 of the experiment. Twenty or 60 minutes after hCFA injections, mice perfused with PBS and PLP
172 (see below) through the heart to obtain fixed kidneys for later immunohistochemistry and
173 localization of recombinant hCAF.

174

175 *Immunohistochemistry*

176 Mice were anesthetized with ketamine/xylazine (i.p.) and systemically perfused through the left
177 ventricle with phosphate-buffered saline (PBS) to remove blood followed by paraformaldehyde-
178 lysine-periodate (PLP) fixative (50 ml/mouse) [23]. Kidneys were removed and fixed overnight at
179 4°C by immersion in PLP. Kidneys were washed 3 times with PBS and 5 µm cryosections were cut
180 after cryoprotection with 2.3 M sucrose in PBS for at least 12 h. Immunostaining was carried out
181 as described previously [24]. Briefly, sections were shortly incubated in microwave with Tris-HC
182 [pH 10], following 1% (wt/vol) SDS for 5 min for retrieval of antigenic sites, washed 3 times with
183 PBS and incubated with PBS containing 1% bovine serum albumin for 15 min prior to the primary
184 antibody. The primary antibodies were mouse monoclonal anti cleaved Agrin Abs (CAF; 14B7B8;
185 1:1000) and (CAF; 12A11D11; 1:1000) (provided by Neurotune diluted in PBS and applied
186 overnight at 4°C. Sections were then washed twice for 5 min with high NaCl PBS (PBS + 18 g
187 NaCl/l), once with PBS, and incubated with dilutions of the secondary antibodies (donkey anti-
188 rabbit 594 (1:500), donkey anti-mouse 488 (1:200), donkey anti-mouse Alexa 594 (1:500), donkey
189 anti-rabbit Alexa 488 (1:500) (Molecular Probes, Oregon, USA) and from DAPI (1 mg/ml) (1:500)
190 for 1 h at room temperature. Sections were again washed twice with high NaCl PBS and once with
191 PBS before mounting with VectaMount (Vector Laboratories, Burlingame, CA). Sections were

192 viewed with a Zeiss LSM 410 confocal microscope or a Leica DFC490 charged-coupled device
193 camera attached to a Leica DM 6000 fluorescence microscope (Leica, Wetzlar, Germany). Images
194 were processed (overlays) using Adobe Photoshop.

195

196 **Results**

197 *Study population*

198 A total of 71 consecutive patients with long standing (>10 years) diabetic nephropathy, were
199 recruited from January 2010 to December 2014. A full medical history and physical examination
200 was performed by a trained physician at baseline (time point 0; T0) and after 12 months (time
201 point 1; T1). All patients were of Caucasian origin. Demographic, clinical and laboratory data of all
202 enrolled patients at T0 and T1 as well as of groups of progressors (loss of GFR > 1ml/min/1.73m²)
203 and non progressors (stable GFR) are given in Table 1.

204 There was no significant difference in age, BMI and duration of T2DM across the categories of
205 patients with and without GFR decline. Urea and phosphate concentrations were significantly
206 different across groups and as expected, increased in the group of patients with decreased GFR
207 (p=0.004, 0.03 respectively). There were significantly more males than females among GFR
208 decliners compared with those with stable or increased GFR (62,5% vs 35,4%, p=0.024). sCAF
209 levels showed a trend towards higher values in those with deterioration of renal function (p=0.09).

210

211 *Correlation analysis*

212 Correlation matrix analysis showed that values of creatinine and eGFR at both time-point 0 and
213 time-point 1 were significantly associated with sCAF levels (p<0.001) (Table 2). sCAF concentration
214 was also correlated with proteinuria at baseline and at T1 (p=0.02 and p<0.001, respectively)
215 (Table 2). Furthermore, sCAF levels were strongly correlated with progression to ESRD (r=0.314,

216 $p=0.008$) (Table 2). At 24 months follow-up (time point 2; T2) 6 of the 71 patients progressed to
217 ESRD (8.5%). Baseline sCAF levels were significantly elevated in those who reached ESRD
218 compared to the group of patients that did not ($p=0.009$). sCAF also showed a significant
219 association with the rapid increase of proteinuria (>500 mg/day) at T1 ($r=0.340$, $p=0.004$) (Table
220 2). The baseline sCAF level was higher in those who presented a rapid increase of proteinuria than
221 in patients with stable, decreased or mild increase (<500 mg/day) of proteinuria ($p=0.005$). A
222 significant positive association was observed between rapid increase of proteinuria and
223 progression to ESRD (χ^2 , $p=0.005$).

224

225 *Logistic regression analysis*

226 Logistic regression models of the association of various renal outcomes (normal and rapid GFR
227 loss) with baseline sCAF levels and other clinically significant regressors, assessed by 2
228 complementary models: Basic model A (CAF, baseline GFR and baseline proteinuria) and advanced
229 model B (CAF, baseline GFR, baseline proteinuria, age, BMI) are shown in Table 3. CAF was the
230 only variable which was significantly associated with GFR loss (≥ 1 ml/min/1.73m²) in both model
231 A [OR(95%CI) 4.2(1.2-14.5), $p=0.024$] and model B [OR(CI) 4.09(1.2-14.4), $p=0.028$].

232

233 *Animal experiments*

234 In order to test for the role of the kidney in determining serum CAF levels, we injected
235 recombinant human CAF22 into neurotrypsin KO mice. These mice lack endogenous CAF22 due to
236 the absence of the specific protease required for the cleavage of agrin [22]. In kidneys from NT KO
237 mice injected with saline, no recombinant hCAF22 could be observed providing evidence for the
238 specificity of the antibody. However, full length agrin was detected in the glomerulum and in the
239 basement membrane of all tubules (Figure 1A). Twenty and sixty minutes after hCFA22 injections,

240 hCAF22 accumulated in subapical vesicles along the proximal tubule consistent with the
241 reabsorption of hCAF22 from urine via apical endocytosis (Figure 1B-F).

242

243

244

245 **Discussion**

246 The major findings in this prospective study of predominantly elderly patients with long-term
247 diabetic nephropathy are that sCAF levels are strongly associated with eGFR and proteinuria. Of
248 note these associations were preserved and evident in subsequent measurements for both
249 parameters at T1. Higher baseline sCAF levels were significantly correlated with progression to
250 ESRD and notably elevated in subjects with an overt increase of proteinuria at T1. Furthermore,
251 sCAF levels predicted loss of renal function, as mirrored by eGFR decline at T1 even after adjusting
252 for multiple well established risk factors, and in particular, for baseline GFR and proteinuria.

253 Our results are consistent with previous studies that have reported associations between sCAF
254 levels and renal function. Steubl et al. showed that sCAF was highly correlated with kidney
255 function in a renal transplant cohort [18]. Additionally, Drey and colleagues demonstrated that
256 sCAF levels are strongly associated and comparable to established renal function parameters
257 (creatinine and cystatin C) in critical ill patients [17]. In the present study, serum levels of CAF at
258 baseline were significantly higher in diabetics, which progressed to ESRD. Importantly, the
259 association between sCAF levels and the different aspects of kidney dysfunction were also evident
260 irrespective of proteinuria. Although sCAF levels reflect renal function and is highly associated with
261 eGFR and proteinuria at several time points, it predicted loss of renal function also in the absence
262 of proteinuria in diabetic nephropathy. sCAF seems to overcome a significant prognostic

263 restriction of proteinuria in a subset of patients with rapid decline of renal function and no
264 progressive proteinuria, a finding that to our knowledge has not been reported before.

265 Our animal experiments are consistent with the hypothesis that CAF22 is cleared from the
266 circulation by glomerular filtration. Injected human recombinant hCAF22 accumulated in subapical
267 vesicles along the proximal tubule. These vesicles likely represent endocytic vesicles containing
268 low molecular weight proteins that escaped the glomerular filtration barrier. CAF22 has only 22
269 kDa size and is expected to be filtered to a large extent similar to other proteins of this size [25].
270 The subapical localization of vesicles containing hCAF22 strongly suggests that hCAF22 is taken up
271 from urine by endocytosis, possibly receptor-mediated endocytosis. It is unlikely that hCAF22 is
272 secreted into urine by proximal tubules in quantitatively relevant amounts since we never
273 detected hCAF22 positive vesicles close to the basolateral membrane. Reabsorption of hCAF22
274 along with other low molecular weight proteins will determine its urinary excretion. Thus, reduced
275 glomerular filtration in kidney disease would increase serum CAF22 levels whereas a selective
276 reduction in proximal tubular reabsorption may increase urinary CAF22 levels. While the exact
277 mechanism of CAF production and trafficking in the kidney is currently under investigation, our
278 findings further support the notion that CAF may reflect both structural (proteinuria) and
279 functional (glomerular filtration rate) alterations.

280 CKD has a major public impact worldwide, with a global prevalence of 10% and diabetic
281 nephropathy is the primary reason of CKD [1,26,27]. sCAF may represent a promising biomarker of
282 kidney damage and progression to ESRD in diabetic nephropathy. However, more studies with
283 sufficient follow up are needed to evaluate the clinical value of sCAF measurements for: 1) the
284 detection of kidney damage, 2) the prediction of GFR decline and 3) the development of ESRD. The
285 correlation between sCAF levels and the progression of kidney dysfunction in other cohorts [28] as
286 well as in the present study indicates that sCAF measurements may be of clinical importance. This

287 finding may open new possibilities for clinical trial design, since sCAF levels could be used as a
288 selection tool for high-risk patients (individuals with CKD and in particular with diabetic
289 nephropathy and no overt proteinuria at baseline) and for monitoring treatment effects in
290 diabetic nephropathy.

291 Limitations of the present study include the single center design, the short-term follow-up and
292 unknown cross-relevance to other age and ethnic groups. While a multitude of relevant variables
293 were adjusted in our multivariate analyses, as with all observational studies, it is possible that
294 unmeasured confounders may have influenced our estimates and results. No conclusions
295 regarding causality should be drawn from our prospective observational data. However, the strong
296 associations with eGFR and proteinuria in diabetics are of interest because sCAF has been shown
297 to be a prognostic marker of kidney dysfunction in other cohorts including patients with CKD and
298 in critically ill patients with acute kidney injury [17,18].

299 Our findings suggest that sCAF, a novel kidney function and injury biomarker, is associated with
300 subsequent renal function loss irrespective of proteinuria at baseline. The association we observed
301 was strong and was not altered by adjusting for several risk factors. sCAF levels provide important
302 information on the long-term outcome of patients with diabetic nephropathy, which exceed a
303 simple reflection of glomerular filtration rate and proteinuria. sCAF measurements may have a
304 substantial impact on clinical trial design as a selection tool since it holds the potential to identify
305 individuals with the higher progression risk for diabetic nephropathy. Further studies are needed
306 in order to replicate our results and to confirm sCAF as a monitoring tool for rapid disease
307 progression in diabetic kidney disease.

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309

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387 **TABLES**

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389 **Table 1:** Anthropometric, clinical, and biochemical characteristics of patients with TD2M and CKD

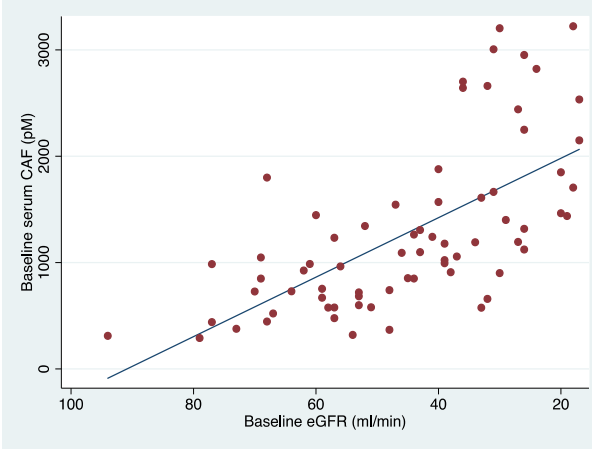
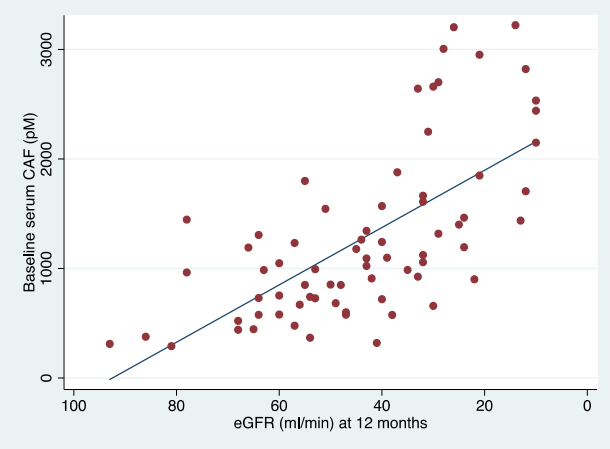
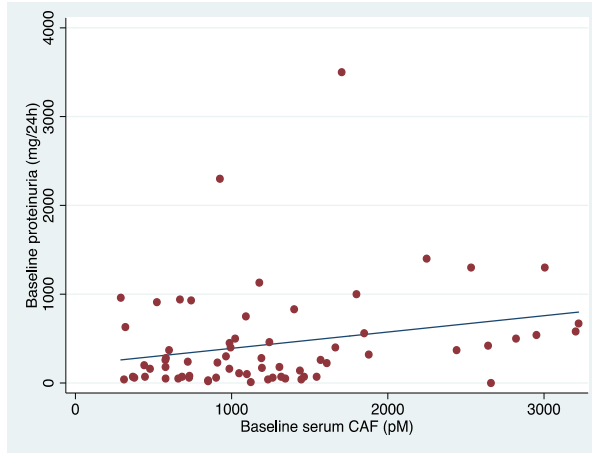
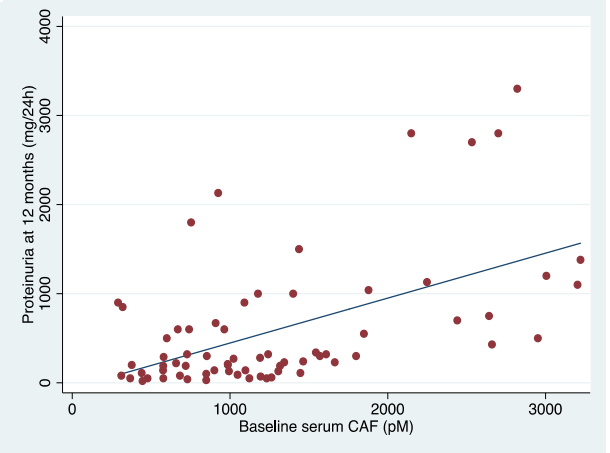
390 with stable (or increased) and with decreased GFR at one year follow-up.

	All patients (N=71)	Stable GFR (N=31)	Decreased GFR (N=40)	p
Age (years)	70.9 (8.8)	71.8 (6.4)	70.2 (10.3)	0.79
Gender (M/F)	36/35	11/20	25/15	0.024*
BMI (kg/m ²)	32.4 (5.9)	33.0 (5.9)	31.9 (6.0)	0.49
Duration of T2DM (years)	17.4 (9.7)	16.5 (7.6)	18.1 (11.2)	0.83
RAAS blockade (%)	49 (69)	25 (80.6)	24 (60)	0.06
Urea (mg/dl)	13.1 (6.76)	10.5 (3.98)	15.1 (7.77)	0.004*
Potassium (mEq/L)	4.8 (0.5)	4.7 (0.3)	4.9 (0.6)	0.34
Sodium (mEq/L)	138.3 (3.0)	138.2 (3.0)	138.4 (3.0)	0.71
Calcium (mmol/L)	2.35 (0.15)	2.38 (0.13)	2.33 (0.18)	0.35
Phosphate (mmol/L)	1.23 (0.26)	1.16 (0.19)	1.29 (0.29)	0.03*
Protein (g/L)	73 (6)	74 (07)	72 (6)	0.34
Albumin (g/L)	43 (4)	44 (3)	42 (4)	0.06
HbA1c (%)	8.6 (9.3)	7.5 (1.3)	9.5 (12.3)	0.88
Crea T0 (mg/dl)	123.8 (53)	114.9 (35.4)	132.6 (61.9)	0.19
Crea T1 (mg/dl)	123.8 (79.6)	106.1 (35.4)	150.3 (79.6)	<0.001*
eGFR T0 (ml/min/1.73 m ²)	43 (27)	44 (23)	40.5 (30.5)	0.62
eGFR T1 (ml/min/1.73 m ²)	43 (27)	54 (24)	34 (23)	<0.001*
ΔGFR T0-T1 (ml/min/1.73 m ²)	-2.0 (11)	5.0 (8)	-6.0 (8)	<0.001*
PU T0 (mg/24h)	270 (600)	245 (430)	370 (705)	0.28
PU T1 (mg/24h)	300 (770)	240 (490)	465 (985)	0.059
ΔPU T0-T1 (mg/24h)	10 (270)	0 (200)	45 (390)	0.19
ESRD (%)	6 (8.5)	0 (0)	6 (15)	0.024*
Total CAF (pmol/L)	1091.4 (890.2)	993.4 (737.8)	1145.9 (1571.1)	0.09

391 Continuous variables are presented as mean (S.D.) or median (interquartile rangeIQR). *P* values of
392 Mann-Whitney *U* or one-way ANOVA test for differences of variables among patients with stable
393 or decreased eGFR at one year follow-up. * statistical significance at the 0.05 level (two-tailed)
394 BMI, body mass index; HbA1c, glycosylated hemoglobin A1c; RAAS blockade, use of medicines
395 affecting renin–angiotensin system; Crea T0, serum creatinine levels at timepoint 0 (baseline);
396 Crea T1, serum creatinine levels at timepoint 1 (12 months); eGFR T0, estimated GFR assessed by
397 the CKD-EPI formula at timepoint 0 (baseline); eGFR T1, estimated GFR assessed by the CKD-EPI
398 formula at timepoint 1 (12 months); PU T0, Proteinuria assessed by protein to creatinine ratio at
399 timepoint 0 (baseline); PU T1, Proteinuria assessed by protein to creatinine ratio at timepoint 1
400 (12 months); Δ GFR T0-T1, Algebraic difference between eGFR at timepoint 1 and eGFR at timepoint
401 0; Δ PU T0-T1, Algebraic difference between Proteinuria at timepoint 1 (12 months) and Proteinuria
402 at timepoint 0 (baseline); ESRD, progression to end stage renal disease; Total-CAF, serum levels of
403 C-terminal agrin fragment; Decreased GFR, patients with loss of eGFR ≥ 1 ml/min /1.73m² during
404 the 12 month follow-up.

TABLE 2: Correlation analysis of CAF levels with variables of laboratory and clinical significance and scatterplots of CAF with GFR and proteinuria.

CAF levels (pmol/L)	r	p
Age	,032	0.79
Body Mass Index	,118	0.32
HbA1c	-,098	0.42
Duration of T2DM	,035	0.77
Creatinine T0	,705 ^b	<0.001*
Creatinine T1	,644 ^b	<0.001*
Estimated GFR T0	-,698 ^b	<0.001*
Estimated GFR T1	-,677 ^b	<0.001*
Proteinuria T0	,287 ^a	0.02*
Proteinuria T1	,449 ^b	<0.001*
End Stage Renal Disease	,314	0.008*
Δ-Proteinuria T0-T1	,340	0.004*

Values represent Spearman's correlation coefficients.

^a Correlation is significant at the 0.05 level (2-tailed).

HbA1c, glycosylated hemoglobin A1c; T0, Time point 0 (Baseline); T1, Time point 2 (12 months); Δ -

Proteinuria T0-T1, Increase of proteinuria (≥ 500 mg/day) during the first year of follow-up.

TABLE 3 : Association between GFR-decline and rapid GFR-decline with serum CAF and additional clinical regressors

		GFR decline at 1 year ≥ 1 ml/min/1.73m²		
		OR	CI	P
Model A	logCAF	4.18	1.2-14.5	0.024*
	eGFR T0	1.03	0.99-1.08	0.102
	PU T0	1.0002	0.99-1.0006	0.299
Model B	logCAF	4.09	1.16-14.4	0.028*
	eGFR T0	1.03	0.98-1.08	0.161
	PU T0	1.0002	0.99-1-0007	0.342
	Age	0.99	0.93-1.07	0.950
	BMI	0.96	0.88-1.05	0.438

OR, odds ratio; CI, 95% confidence interval, *significance levels at 0.05

BMI, body mass index; eGFR T0, estimated GFR assessed by the CKD-EPI formula at baseline; PU T0, Proteinuria assessed by protein to creatinine ratio at baseline; logCAF, serum CAF levels (log-transformed). GFR, estimated glomerular filtration rate

FIGURE 1

Accumulation of recombinant hCAF22 in mouse proximal tubule.

Mice were injected with saline or hCAF22 and all sections stained in parallel with the antibodies as indicated. **A.** Localization of full length agrin (red) and hCAF22 (green) in NT KO kidney injected with saline. **B.** Localization of full length agrin (red) and hCAF22 (green) in NT KO mouse kidney 20 min after hCAF22 injections. **C.** Localizations of hCAF22 (green) and actin (red) in NT KO mouse kidney 20 min after hCAF22 injections. **D.** Localization of full length agrin (red) and hCAF22 (green) in NT KO mouse kidney 60 min after hCAF22 injections. **E-F.** Localizations of hCAF22 (green) and actin (red) in NT KO mouse kidney 60 min after hCAF22 injections. A- E Original magnifications between 400 - 630 x, for F original magnification 1000 x.