

# Pentoxifylline ameliorates subclinical atherosclerosis progression in patients with type 2 diabetes and chronic kidney disease: a randomized pilot trial



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# Abstract

**Background** Diabetic kidney disease (DKD) is associated with a higher risk of cardiovascular disease (CVD). Pentoxifylline (PTF), a nonselective phosphodiesterase inhibitor with anti-inflammatory, antiproliferative, and antifibrotic actions, has demonstrated renal benefits in both clinical trials and meta-analyses. The present work aimed to study the effects of PTF on the progression of subclinical atherosclerosis (SA) in a population of patients with diabetes and moderate to severe chronic kidney disease (CKD).

**Methods** In this open-label, randomized controlled, prospective single-center pilot study the evolution of carotid intima-media thickness (CIMT) and ankle-brachial index (ABI) were determined in 102 patients with type 2 diabetes mellitus and CKD assigned to PTF, aspirin or control groups during 18 months. We also determined the variations in the levels of inflammatory markers and Klotho (KL), a protein involved in maintaining cardiovascular health, and their relationship with the progression of SA.

**Results** Patients treated with PTF presented a better evolution of CIMT, increased *KL* mRNA levels in peripheral blood cells (PBCs) and reduced the inflammatory state. The progression of CIMT values was inversely related to variations in KL both in serum and mRNA expression levels in PBCs. Multiple regression analysis demonstrated that PTF treatment and variations in mRNA *KL* expression in PBCs, together with changes in HDL, were significant determinants for the progression of CIMT (adjusted  $R^2 = 0.24$ , *P* < 0.001) independently of traditional risk factors. Moreover, both variables

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constituted protective factors against a worst progression of CIMT [OR: 0.103 (P=0.001) and 0.001 (P=0.005), respectively].

**Conclusions** PTF reduced SA progression assessed by CIMT variation, a beneficial effect related to KL gene expression in PBCs.

**Trial registration** The study protocol code is PTF-AA-TR-2009 and the trial was registered on the European Union Drug Regulating Authorities Clinical Trials (EudraCT #2009–016595–77). The validation date was 2010-03-09.

**Keywords** Pentoxifylline, Diabetic kidney disease, Subclinical atherosclerosis, Klotho, Inflammation, Carotid intimamedia thickness

# Background

Diabetic mellitus (DM) is one of the leading causes of chronic kidney disease (CKD) and end-stage renal disease (ESRD) worldwide [1]. DM and CKD are recognized risk factors for cardiovascular disease (CVD), and the presence of both synergistically increases cardiovascular risk in patients with diabetic kidney disease (DKD) [2]. In these patients, traditional and non-traditional risk factors for atherosclerosis coexist, contributing to the excess of mortality. Non-traditional risk factors related to CKD include alterations in mineral metabolism and chronic inflammation [3, 4]. This scenario favors subclinical atherosclerosis (SA) to be more prevalent in people with CKD and DM, with a higher plaque load and faster progression. [5]. Furthermore, SA burden has been described as the most powerful predictor of cardiovascular events in this population [6]. Therefore, early detection of SA burden but also new, effective and safe therapeutic approaches starting at the earliest stages in patients with DKD are needed to prevent atheromatous disease and reduce cardiovascular events.

Pentoxifylline (PTF) is a methylxanthine derivative non-selective phosphodiesterase inhibitor currently indicated for peripheral arterial disease that exhibits antiinflammatory, antiproliferative, and antifibrotic actions [7]. Moreover, clinical trials and meta-analyses have demonstrated renal protection secondary to anti-inflammatory and antiproteinuric effects in patients with DM and CKD treated with PTF when added to renin-angiotensin system (RAS) blockade [7].

The anti-aging protein  $\alpha$ Klotho (hereinafter Klotho, KL) is an important regulator of mineral metabolism expressed predominantly in the kidneys but also, to a lesser extent, in the parathyroid glands, choroid plexus, vascular tissue, and peripheral blood cells (PBCs) [8–10]. Two forms of KL can be found: a single-pass transmembrane protein and a soluble form generated from the proteolytic cleavage of the extracellular domain of the former. Soluble KL levels in blood and urine decrease from the early stages of CKD, being a characteristic finding of renal involvement in DM since renal KL expression is markedly decreased in DKD compared to other forms of kidney disease [11]. Furthermore, clinical

studies suggest that this reduction is associated with the prevalence and severity of CVD [10, 12–14] and all-cause mortality [15], being related to markers of vascular dysfunction and the incidence of atherosclerosis [10, 16–19]. We have recently conducted a cross-sectional case-control study showing independent associations between reduced serum and PBCs expression levels of KL and the presence of SA in patients with CKD [19]. A post hoc analysis of the Pentoxifylline for Renoprotection in Diabetic Nephropathy (PREDIAN) trial revealed that administration of PTF to patients with type 2 DM (T2DM) and CKD stages 3–4 significantly increased KL concentrations in both serum and urine [20, 21].

In this pilot randomized controlled trial, we have explored the potential benefits of PTF therapy on the progression of SA in patients with DKD. We also determined variations in the expression levels of KL and inflammatory cytokines in serum and PBCs.

# Methods

# Patients and study design

Single-center, open-label, randomized and prospective pilot study conducted at the University Hospital Nuestra Señora de Candelaria (Santa Cruz de Tenerife, Spain). Between May and November 2010, a total of 222 patients with T2DM and CKD stage 3 without clinical CVD were considered for initial enrollment. During this 7-month period, no significant variations were observed in the background therapies used in any of the final study groups. Patients who finally entered the study were randomly assigned to a control group, a PTF group (1200 mg/day), or to a group treated with aspirin (100 mg/day). The dose and frequency of oral PTF were 600 mg extended-release tablets twice daily with meals. Patients were asked to begin treatment by taking a single tablet after dinner. After four weeks, one tablet was administered every 12 h, thus reaching the recommended daily dose. The duration of the study was 18 months. Inclusion criteria: CKD patients in stage 3 [estimated glomerular filtration rate (eGFR) 30-59 ml/min/1.73 m<sup>2</sup> according to the Modification of Diet in Renal Disease Study-4 (MDRD-4) equation]; older than 18 years; therapy with RAS blockers and statins; and no history of clinical atherosclerotic CVD. Exclusion criteria: anklebrachial index (ABI) values  $\geq$  1.3; history of heart failure; chronic inflammatory, immunologic, or tumoral disease; positive serology to hepatitis B, hepatitis C, or HIV; acute inflammatory or infectious intercurrent episodes in the previous month; institutionalization; receipt of immunotherapy or immunosuppressive treatment; previous therapy with PTF; and inability or unwillingness to provide informed consent. The study protocol was approved by the Institutional Ethics Committee of the University Hospital Nuestra Señora de Candelaria and complied with ethical standards of the Declaration of Helsinki. Written informed consent was obtained from all participants. Participants medication-adherence and evolution were followed up with clinical visits every 3 or 6 months.

#### Vascular and biochemical assessments

The primary outcome was SA progression, which was assessed by measuring ABI and carotid intima-media thickness (CIMT) at inclusion and at the end of the study. The ABI was calculated using a portable pulse detector (Ultrasonic Mini Doppler ES-100VX; Hayashi Denki Co., Ltd., Kawasaki, Japan) with an 8 mHz probe. CIMT measurement was performed by a single reader in a blinded fashion by ultrasonography of the carotid arteries employing a high-resolution ultrasound (Philips ATL 5000 HDI; Royal Philips Electronics, Amsterdam, The Netherlands) equipped with a 6-13 MHz linear array transducer. According to the Guidelines for the management of arterial hypertension released by the Task Force for the Management of Arterial Hypertension of the European Society of Hypertension (ESH) and the European Society of Cardiology (ESC) [22], we defined patients with SA as those having an ABI<0.9 and/or CIMT≥0.9 mm.

Secondary outcomes included the progression of CKD and the variations in the levels of inflammatory parameters and KL. Blood samples were drawn in the morning after an 8- to 12- overnight fasting at baseline and at the end of the study. Serum samples were retrieved, aliquoted, and immediately frozen at -80°C. Routine biochemical parameters were measured using standard methods. The serum levels of the cytokines tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) and interleukin (IL) 10 were measured by ELISA methods (Quantikine<sup>®</sup>, R&D Systems, Abingdon, UK). Minimum detectable concentrations were 0.10 pg/mL and 0.09 pg/mL, respectively. Intraand inter-assay coefficients of variability were <10.8%. High-sensitivity serum C-reactive protein (hsCRP) was measured by a high-sensitivity particle enhanced immunoturbidimetric fully automated assay (Roche Diagnostics GmbH, Mannheim, Germany) in a Cobas 6000 analyzer with a sensitivity of 0.3 mg/L and intra- and inter-assay coefficients of variation of 1.6% and 8.4%,

respectively. Concentrations of serum KL protein were measured by a solid phase sandwich ELISA using the human soluble  $\alpha$ KL assay kit (Immuno-Biological Laboratories, Takasaki, Japan) according to manufacturer's instructions. The assay sensitivity was 6.15 pg/mL and the intra- and inter-assay coefficients of variation were 2.7–3.5% and 2.9–11.4%, respectively.

#### Gene expression analysis in PBCs

Whole blood samples (2.5 mL) were collected in PAXgene blood RNA tubes (BD, Franklin Lakes, NJ) at the same time as serum samples. Total RNA was isolated using a PAXgene blood RNA kit (Qiagen, Valencia, CA) according to manufacturer's specifications, quantified using a Thermo Scientific NanoDrop Lite Spectrophotometer (Thermo Fisher Scientific, MA, USA) and retrotranscribed by using a High-Capacity RNA-to-cDNA kit (Thermo Fisher Scientific). Transcripts of KL gene, TNF, IL10, and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) as constitutive gene, were measured by real-time TaqMan quantitative PCR (qRT-PCR) with TaqMan Fast Universal PCR master mix (Thermo Fisher Scientific) in a 7500 Fast Real-Time PCR System (Thermo Fisher Scientific). TaqMan gene expression assays for each transcript were: Hs00183100\_m1 [KL], Hs00174128\_ml [TNF], Hs0961622\_m1 [IL10], and Hs99999905\_m1 [GAPDH]. The level of target mRNA was estimated by relative quantification using the comparative method  $(2^{-\Delta\Delta Ct})$  by normalizing to *GAPDH* expression. Quantification of each sample was tested in triplicate and expressed as arbitrary units (a.u.).

# Statistical analysis

Sample size calculation was based on previous studies reporting a mean CIMT of  $0.68\pm0.15$  mm in patients with CKD. The number of patients needed to detect a 15% CIMT difference between groups over 18 months with longitudinal measurements, with a power of 90% and a 5% level of significance, and allowing a dropout percentage of 10%, was 96 (32 subjects in each group).

Continuous variables were checked for the normal distribution assumption using the Kolmogorov–Smirnov statistic. Normally and non-normally distributed variables are expressed as mean±standard deviation (SD) or as median and interquartile range (IQR), respectively. Categorical data are presented as absolute values and percentages. Differences between groups or after treatment were analyzed using Fisher's exact test, Student's t-test, the Mann-Whitney U-test, or Wilcoxon signed rank test as appropriate. Spearman rank correlation was used to determine correlations between changes in SA markers and variations in inflammatory parameters or KL. A stepwise multiple regression analysis was performed to determine the independent association between changes in potential predictive variables and variations in ABI and CIMT. Tolerance and variance inflation factor (VIF) were analyzed in order to exclude collinearity. Finally, multiple logistic regression was performed to evaluate whether PTF therapy and changes in KL and inflammatory cytokines were independent predictors of changes in CIMT values above the median. Statistical analyses were conducted with SPSS software version 25 (IBM Corp.

# Results

#### **Study population**

be statistically significant.

Figure 1 shows the flow of participants through the trial. A total of 222 patients were initially screened for eligibility. Of them, 8 refused to participate, 74 did not meet the inclusion criteria and 32 met the exclusion criteria. Therefore, 108 patients were randomized and 102 completed the study protocol.

Armonk, NY, USA). A value of P < 0.05 was considered to

# **Clinical data**

Baseline characteristics of participants are shown in Table 1 and in supplementary material (Table S1, respectively. Demographic, clinical, and biochemical characteristics and concomitant treatments were balanced between the 3 study groups, with similar proportions of patients treated with antihypertensive, antidiabetic, and antiplatelet medications. A total of 102 patients were randomly assigned to the control group or one of the two treatment groups. All participants were white with a mean age of  $67.3\pm7.9$  years, a similar distribution by sex (male, 49%), hypertension (92.2%), dyslipidemia (74.5%), and smoking habits (34.5%). MedianeGFR was 38.6 (IQR, 35.5-43.9)ml/min/1.73m<sup>2</sup>, with CKD stages 3a (76.5%) and 3b (23.5%) similarly distributed between groups. The median values for ABI and CIMT were 0.95 (IOR, 0.88-1.06) and 0.75 (IOR, 0.7–0.82) mm, respectively. Fifty-six patients (54.9%) had macroalbuminuria (urinary albumin-to-creatinine ratio [UACR] > 300 mg/g). SA was present in 42 patients (41.2%), with no differences in distribution among groups. Likewise, the three groups presented similar levels of HbA1c, fasting glucose, hs-CRP, and serum and PBCs-mRNA levels of TNFα, IL10 and KL.

BP, blood pressure; BMI, body mass index; ABI, anklebrachial index; CIMT, carotid intima-media thickness;



# Table 1 Clinical characteristics, biochemical assessments and gene expression analyses according to the groups at baseline

	Control	ontrol Pentoxyfilline		Р
Characteristics				
Ν	34	34	34	
Age (years)	65.4±8.2	67.8±7.6	68.7±7.7	0.191
Male (%)	17 (50)	16 (47)	17 (50)	0.962
Systolic BP (mm Hg)	143.1±9.5	142.2±8.5	141.2±8.4	0.739
Diastolic BP (mm Hg)	86±8.5	85.8±9.2	86±7.4	0.993
BMI (kg/m <sup>2</sup> )	28 (25.8–30.3)	30 (26-31.3)	29.5 (27.8–32)	0.123
ABI	0.95 (0.88–1.06)	0.92 (0.86-1.04)	0.97 (0.89–1.07)	0.483
CIMT (mm)	0.735 (0.7-0.823)	0.76 (0.71–0.813)	0.75 (0.7–0.84)	0.928
Comorbidities (%)				
Hypertension	31 (91.2)	31 (91.2)	32 (94.1)	0.873
Current smokers	13 (38.2)	10 (29.4)	12 (35.3)	0.738
Macroalbuminuria	19 (55.9)	20 (58.8)	17 (50)	0.758
CKD Stage 3a/3b	23/11	29/5	26/8	0.23
ABI < 0.9	12 (35.3)	15 (44.1)	11 (31.4)	0.536
CIMT≥0.9	4 (11.8)	1 (2.9)	5 (14.3)	0.249
Dyslipidemia	26 (76.5)	24 (70.6)	26 (76.5)	0.813
SA	13 (38.2)	16 (47)	13 (38.2)	0.658
Laboratory data				
Hemoglobin (g/dl)	11.9 (11.6–12.4)	11.8 (11.1–12)	11.8 (11.3–12.4)	0.191
Creatinine (mg/dl)	1.79 (1.52–2.41)	2.03 (1.46-2.73)	1.99 (1.4–2.8)	0.849
eGFR (ml/min/1.73 m <sup>2</sup> )	39 (35.8–49.7)	38.9 (34.9–42.1)	38.5 (34.6–42.7)	0.546
UACR (mg/g)	349.5 (153–1001)	544.5 (176.3–1200)	340 (190–1000)	0.612
Albumin (g/dL)	3.8 (3.5–4)	3.7 (3.5–3.9)	3.9 (3.63-4)	0.155
T-cholesterol (mg/dL)	168.5±31.3	171.6±42.2	176.7±35.3	0.665
HDL-C (mg/dL)	41.5 (35.8–45)	42 (38–46.3)	42 (35.8–46)	0.644
LDL-C (mg/dL)	95.9±23.5	91.7±27.5	100.6±26	0.392
Triglycerides (mg/dL)	153.7±44.5	147.1±35.3	151.2±43.4	0.784
Uric acid (mg/dL)	6.8±1.62	$6.5 \pm 1.56$	$6.63 \pm 1.2$	0.648
Glucose (mg/dL)	89.5 (79.8–121.5)	96.5 (80.3–145.5)	85.5 (77–141)	0.71
HbA1c (%)	7.3 (6.9–7.6)	7.3 (6.9–7.5)	7.2 (6.8–7.8)	0.957
Calcium (mg/dL)	9.29 (8.98–9.9)	9.2 (8.88–9.6)	9.2 (8.9–9.8)	0.628
Phosphorus (mg/dL)	4 (3.7–4.8)	4.3 (3.8–5)	4.4 (3.9–5.1)	0.121
KL (pg/mL)	628±211.3	$602 \pm 169.6$	669.6±208.8	0.375
hs-CRP (mg/L)	$5.23 \pm 2.14$	$5.25 \pm 2.45$	$5.4 \pm 2.6$	0.981
TNFa (pg/mL)	15.5 (12.6–17.5)	15.3 (12.5–17.5)	15.6 (10.9–18.8)	0.937
IL10 (pg/mL)	37.6 (28.8–50.7)	31.8 (24.6–40)	34.7 (24.5–54.2)	0.343
KL mRNA (a.u.)	1.75 (1.4–2.62)	1.8 (1.6–2.7)	2.2 (1.7–2.9)	0.082
TNF mRNA (a.u.)	2 (1.4–2.8)	1.8 (1.6–2.4)	1.9 (1.6–2.5)	0.578
<i>IL10</i> mRNA (a.u.)	$1.23 \pm 0.2$	$1.26 \pm 0.162$	$1.27 \pm 0.25$	0.779
Medication (%)				
Insulin	15 (44.1)	18 (52.9)	16 (47.1)	0.811
Oral antidiabetics	18 (52.9)	18 (52.9)	17 (50)	0.872
ß-Blockers	16 (47.1)	17 (50)	17 (50)	0.801
CCB	21 (61.8)	18 (52.9)	18 (52.9)	0.731
Antiplatelet drugs	31 (91.2)	24 (70.6)	26 (76.5)	0.573

BP, blood pressure; BMI, body mass index; ABI, ankle-brachial index; CIMT, carotid intima-media thickness; SA, subclinical atherosclerosis; eGFR, estimated glomerular filtrate rate; UACR, urinary albumin-to-creatinine ratio; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; hs-CRP, high sensitivity C-reactive protein; TNFα, tumor necrosis factor; IL, interleukin; KL, Klotho; ACEI, angiotensin converting enzyme inhibitor; ARB, angiotensin II receptor antagonist; CCB, calcium channel blockers

SA, subclinical atherosclerosis; eGFR, estimated glomerular filtrate rate; UACR, urinary albumin-to-creatinine ratio; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol; hs-CRP, high sensitivity C-reactive protein; TNF $\alpha$ , tumor necrosis factor; IL, interleukin; KL, Klotho; ACEI, angiotensin converting enzyme inhibitor; ARB, angiotensin II receptor antagonist; CCB, calcium channel blockers

# Evolution of SA, biochemical data, and gene expression in PBCs

After 18 months, the three groups experienced significant increments in CIMT values respect to baseline (Table 2). Control group increased CIMT by 0.015 mm (95% CI, 0.008 to 0.024; P<0.01), the PTF group by 0.006 mm (95% CI, -0.001 to 0.007; *P*<0.01), and aspirin group by 0.014 mm (95% CI, 0.004 to 0.019; P<0.01) resulting in percent variations of 1.63%, 0.58% and 1.48%, respectively (Fig. 2A). Thus, the increase in CIMT was significantly lower in the PTF group than in the control and aspirin groups (P < 0.001). Only patients in the PTF group showed significant variation in ABI values at the end of the study respect to baseline, with an increment of 0.08 (95% CI, -0.03 to 0.11; P<0.01). These variations represented a 7.11% increase of ABI values and a 17.6% reduction in the incidence of patients with ABI values < 0.9 in the PTF group. The differences in the variation of ABI values among groups did not reach statistical significance (P=0.093).

At the end of the study, hs-PCR levels increased in the control group from 5.23±2.14 mg/L to  $6.05\pm2.62$  mg/L (P<0.01); conversely, it was reduced from 5.25±2.45 mg/L to 4.61±2.31 mg/L in the PTF group (P < 0.001), with no differences in the aspirin group (Table 2 and Supplementary Table S2). This resulted in a mean percent difference between control and PTF groups at 18th month for hs-PCR of 28.5% P<0.001) in favor to the first (Fig. 2B). TNFα levels were also reduced from 15.3 pg/mL (IQR, 12.5-17.5) to 13.3 pg/mL (IQR, 12.4–16.3) (P<0.01) in those patients treated with PTF, with no significant changes in the control or in the aspirin groups. This resulted in a mean percentage difference for TNF $\alpha$  changes with respect to the PTF group of 6.5% (95% CI, 0.83-12.1%; P<0.01) and 5.63% % (95% CI, 0.49–10.8%; P<0.01) in favor of the control group and the aspirin groups, respectively (Fig. 2B). Similarly, IL10 only increased significantly in the PTF group: from 31.8 pg/mL (IQR, 24.6-40 pg/mL) to 38.2 pg/mL (IQR, 29.2–48 pg/mL) (P<0.001). Serum KL levels were significantly reduced in control and in aspirin groups (-3.29% [95% CI, -7.09 to 0.51] and -2.4% [95% CI, -4.8% to -0.001%], respectively; P < 0.05 for both) (Fig. 2B). Conversely, patients in the PTF group presented a significant increment in soluble KL levels by 2.1% (95% CI,

	Control	Pentoxyfilline	Aspirin	P value between groups
Systolic BP (mm Hg)	-2 (-3.81 to 0.73)	-1.7 (-3.5 to 0.36)	-1.15 (-2.59 to 0.89)	0.783
Diastolic BP (mm Hg)	-2.4 (-8.2 to 4.1)	-1.5 (-5.1 to 0.77)	-0.85 (-4.2 to 3.5)	0.811
BMI (kg/m²)	0.7 (-0.1 to 1.5)	0.2 (-0.5 to 1.1)	0.9 (-0.3 to 1.7)	0.451
T-cholesterol (mg/dL)	6.6 (-8.3 to 24.3)	17.6 (-15.8 to 35)	5.9 (-24.3 to 38.9)	0.798
HDL-C (mg/dL)	2.5 (-3.23 to 14.2)	3 (-10.5 to 16.52)	1.5 (-12.2 to 12.4)	0.296
LDL-C (mg/dL)	3.1 (-12.2 to 23)	6.3 (-11.9 to 22.6)	5.3 (-17.2 to 34.8)	0.206
Triglycerides (mg/dL)	13.8 (-18.4 to 41.4)	27.1 (7.5 to 61.1)	10.1 (-20.4 to 37.9)	0.29
Uric acid (mg/dL)	-0.35 (-1.54 to 0.79)	-0.57 (-2.28 to -0.67)	0.49 (-0.62 to 1.61)	0.417
Glucose (mg/dL)	7 (-33.2 to 34.9)	-2.4 (-10 to 9.5)	6.5 (-19.9 to 35)	0.305
HbA1c (%)	0.1 (-0.27 to 0.63)	0.17 (-0.39 to 0.73)	0.12 (-0.57 to 0.82)	0.986
ABI	0.035 (-0.013 to 0.12)	0.08 (-0.03 to 0.11) <sup>b</sup>	-0.01 (-0.07 to 0.08)	0.093
CIMT (mm)	0.015 (0.008 to 0.024) <sup>b</sup>	0.006 (–0.001 to 0.007) <sup>b</sup>	0.014 (0.004 to 0.019) <sup>b</sup>	< 0.001
hs-CRP (mg/L)	0.83 (0.31 to 1.35) <sup>b</sup>	–0.63 (–0.95 to –0.31) <sup>c</sup>	-0.2 (-0.57 to 1.47)	< 0.001
TNFa (pg/mL)	0.4 (-0.96 to 0.86)	−2.01 (−3.04 to −1.26) <sup>b</sup>	0.9 (-0.62 to 1.46)	0.001
IL10 (pg/mL)	1.5 (-9.89 to 12.8)	6.4 (0.18 to 9.6) <sup>c</sup>	-0.4 (-4.8 to 3.6)	0.041
KL (pg/mL)	-37.5 (-93.3 to 13.3) <sup>a</sup>	10.32 (-0.05 to 41.8) <sup>a</sup>	-15.3 (-30.2 to 11.9) <sup>a</sup>	0.005
TNF mRNA (a.u.)	0.2 (-0.36 to 0.54)	-0.02 (-0.31 to 0.09)	-0.14 (-0.91 to 0.63)	0.825
<i>IL10</i> mRNA (a.u.)	0.02 (-0.17 to 0.31)	0.02 (-0.055 to 0.13)	0.007 (-0.18 to 0.19)	0.774
KL mRNA (a.u.)	-0.18 (-0.39 to 0.07) <sup>a</sup>	0.47 (0.11 to 1.05) <sup>b</sup>	-0.09 (-0.19 to -0.01) <sup>b</sup>	< 0.001
eGFR (ml/min/1.73 m <sup>2</sup> )	−3.45 (−4.73 to −2.71) <sup>c</sup>	-1.93 (-2.34 to 0.66) <sup>c</sup>	−3.58 (−6 to −3.4) <sup>c</sup>	0.008
UACR (mg/g)	-14.6 (-53 to 81.3)	-116.6 (-300 to 155)	40.8 (-154 to 164)	0.045

Table 2 Variations in diverse atherosclerosis parameters, kidney function, inflammatory cytokines and KL after 18 months

Data represents mean or median values and 95% confidence intervals (95% CI). Abbreviations: BP, blood pressure; BMI, body mass index; LDL-C, low- density lipoprotein cholesterol; HDL-C, high-density lipoprotein cholesterol PTF, pentoxifylline; SA, salicylic acid; ABI, ankle-brachial index; CIMT, carotid intima-media thickness; eGFR, estimated glomerular filtrate rate; UACRE, urinary albumin-to-creatinine ratio; hs-CRP, high sensitivity C-reactive protein; TNF $\alpha$ , tumor necrosis factor; IL, interleukin; KL, Klotho. P values for differences from baseline: a, P < 0.05; b, P < 0.01; c, P < 0.001



**Fig. 2** Percent variations from baseline for CIMT (**A**), serum levels of inflammatory parameters and circulating KL (**B**), mRNA expression levels of cytokines and KL in PBCs (**C**), and variations in eGFR (**D**). Bars and ranges represent mean values and SEM. CIMT, carotid intima-media thickness; PTF, pentoxifylline; hs-CRP, high sensitivity C-reactive protein; TNFa, tumor necrosis factor; IL, interleukin; KL, Klotho; eGFR, estimated glomerular filtrate rate P values: f, < 0.001 vs. control group; #, < 0.001 vs. PTF group;  $\alpha$ , < 0.01 vs. control group; &, < 0.05 vs. control group

0.7–4.14%; P<0.05). A trend was observed for the difference in serum KL variatons between control and PTF groups at the end of the study: 5.4% (95% CI, 1.14–9.64%; P=0.057) in favor to PTF group (Fig. 2B). PBCs *KL* gene expression significantly increased 0.47 a.u. (95% CI, 0.11 to 1.05; P<0.01) in patients receiving PTF and decreased –0.18 (95% CI, –0.39 to 0.07; P<0.05) in the control and –0.09 (95% CI, –0.19 to –0.01; P<0.01) in the aspirin groups. Therefore, PBCs *KL* gene expression increased by 28.2% in the PTF group, whereas decreased by –5.27% and –9.54% in the control and the aspirin groups, respectively (Fig. 2C).

After 18 months, the eGFR decreased by a mean±SEM of  $3.45\pm0.47$  and  $3.58\pm0.43$  ml/min/ $1.73m^2$  in the control and aspirin groups, respectively. By contrast, eGFR only decreased  $1.93\pm0.33$  ml/min/ $1.73m^2$  in patients treated with PTF (Fig. 2D). This resulted in significant mean differences respect to the control and the aspirin groups of 1.52 ml/min/ $1.73m^2$  (95% CI, 0.37 to 2.67 ml/min/ $1.73m^2$ ; P<0.01) and 1.65 ml/min/ $1.73m^2$  (95% CI, -2.75 to -0.55 ml/min/ $1.73m^2$ ; P<0.01), respectively, in favor to PTF in both cases. Regarding albuminuria,

UACR experienced a mean percent decrease of 9.6% in the PTF group, whereas it increased by 3.6% and 12.6% in the control and aspirin groups, respectively. Three patients (8.8%) progressed from stage 3b to stage 4 in the control group *versus* 2 patients (5.9%) in the PTF group.

#### Correlations and multivariate analysis

Variations in CIMT were inversely correlated with changes in HDL-C concentrations (r=0.25, P=0.012), in serum and PBCs-mRNA levels of KL (r = -0.302, P=0.002; r = -0.441, P<0.001, respectively), as well as with variations in eGFR (r = -0.217, P=0.028) (Table 3). Variations in ABI did not correlate with changes in none of the parameters included in the analysis. Among inflammatory markers, only variations in hs-CRP presented significant associations with markers of SA, being positively related with CIMT (r=0.195, P=0.049).

Multiple forward stepwise regression analysis was performed with changes in CIMT as the dependent variable. Results showed that variations in PBCs mRNA *KL* levels and PTF therapy, together with variations in HDL levels, were significantly associated with the evolution of CIMT

**Table 3** Bivariate correlations between changes in ABI or CIMTand changes in diverse atherosclerosis parameters after 18months

	ABI		CIMT	
	r	Р	r	Р
Systolic BP (mm Hg)	-0.046	0.644	0.066	0.511
Diastolic BP (mm Hg)	0.037	0.712	0.098	0.326
BMI (kg/m²)	0.045	0.621	0.072	0.372
Uric acid (mg/dL)	-0.007	0.948	0.052	0.605
T-cholesterol (mg/dL)	-0.041	0.685	-0.024	0.813
HDL-C (mg/dL)	-0.045	0.655	0.25	0.012
LDL-C (mg/dL)	0.024	0.812	-0.027	0.786
Triglycerides (mg/dL)	0.052	0.601	-0.05	0.618
Glucose (mg/dL)	-0.037	0.715	0.062	0.535
HbA1c (%)	0.078	0.634	-0.006	0.972
hs-CRP (mg/L)	-0.170	0.088	0.195	0.049
TNFa (pg/mL)	0.043	0.664	0.174	0.08
IL10 (pg/mL)	0.153	0.124	-0.075	0.451
KL (pg/mL)	-0.089	0.372	-0.302	0.002
TNF mRNA (a.u.)	0.022	0.823	0.048	0.634
<i>IL10</i> mRNA (a.u.)	0.08	0.423	0.001	0.993
<i>KL</i> mRNA (a.u.)	0.11	0.269	-0.441	< 0.001
eGFR (ml/min/1.73 m <sup>2</sup> )	-0.085	0.398	-0.217	0.028
UAF (ma/a)	-0.061	0.546	0.191	0.054

ABI, ankle-brachial index; CIMT, carotid intima-media thickness; BP, blood pressure; BMI, body mass index; HbA1c, glycated hemoglobin; SA, subclinical atherosclerosis; hs-CRP, high sensitivity C-reactive protein; TNFa, tumor necrosis factor; IL, interleukin; KL, Klotho; eGFR, estimated glomerular filtrate rate; UAE, urinary albumin excretion; HDL-C, high-density lipoprotein cholesterol; LDL-C, low-density lipoprotein cholesterol

**Table 4** Multiple backward stepwise regression analysis for changes in CIMT as the dependent variable

	Ad- justed R <sup>2</sup>	ß	Standard error	t	Ρ
$\Delta$ CIMT (mm)	0.24				< 0.001
∆ mRNA KL (a.u.)		-0.301	0.02	-3.01	0.004
Pentoxifylline		-0.212	0.02	-2.11	0.0231
$\Delta$ HDL-C (mg/dL)		0.181	$0.85 \times 10^{-3}$	1.99	0.041

CIMT, carotid intima-media thickness; KL, Klotho; HDL-C, high-density lipoprotein cholesterol

(adjusted  $R^2=0.24$ , P<0.001) (Table 4). Tolerance and VIF values were higher than 0.60 and lower than 1.5 for all variables in any of the analysis.

In the multivariate logistic regression modelling (Table 5), delta CIMT $\geq$ 0.01 mm, which corresponds to the median value of the CIMT variations, was used as the dependent variable to define SA progression. Traditional risk factors for atherosclerosis (sex, baseline hypertension, smoking, DM, and dyslipidemia, together with variations in BMI, systolic BP, T-cholesterol and HbA1c) were entered as covariates (model 1), with additional models in which markers of renal function (eGFR and macroalbuminuria) (model 2), and inflammatory cytokines (model 3) were added. Results of this analyses showed that PTF treatment and variations in *KL* mRNA levels in PBCs were covariates associated with the progression of CIMT.

# Discussion

Treatment with PTF for 18 months led to a slowdown in the rate of SA progression among patients with T2DM and stage 3 CKD. This beneficial effect was reflected in a nearly threefold smaller percentage increase in CIMT in patients receiving PTF compared with patients in the control and aspirin groups (0.58% vs. 1.63% and 1.48%, respectively). The better SA outcome in the PTF group was accompanied by increased KL mRNA expression in PBCs and reduced systemic levels of the inflammatory markers hs-CRP and TNFa. Moreover, patients in the PTF group also presented a lower decline in eGFR, with a significant mean difference of 1.52 ml/min/1.73m<sup>2</sup> and 1.65 ml/min/1.73m<sup>2</sup> with respect to control and aspirin groups, respectively, and a tendency to a significant antialbuminuric effect. The beneficial effect of PTF on CIMT progression was related to variations in KL levels independently of other cardiovascular risk factors. In deep, changes in both KL protein and PBCs KL mRNA levels were significantly related with the progression of CIMT. Specifically, both determinations inversely correlated with CIMT. Moreover, PTF therapy and mRNA KL expression in PBCs were found to be independent determinants for the variation in CIMT values, and protective factors against a worst progression of carotid SA.

**Table 5** Multivariate logistic regression analysis for variations of CIMT ≥ 0.01 mm with PTF administration and variations in serum Klotho and KL PBCs mRNA levels as independent variables. OR values are expressed per unit variation

	Unadjusted		Model 1		Model 2		Model 3	
	OR (95% CI)	Р	OR (95% CI)	Р	OR (95% CI)	Р	OR (95% CI)	Р
PTF administration	0.222 (0.75 to 0.659)	0.007	0.128 (0.089 to 0.311)	0.005	0.115 (0.091 to 0.231)	0.003	0.103 (0.022 to 0.41)	0.001
∆ mRNA <i>KL</i> (a.u.)	0.004 (0.94×10 <sup>-3</sup> to	0.005	0.002 (0.27 × 10 <sup>-3</sup> to	0.003	0.002 (0.21 × 10 <sup>-3</sup> to	0.003	0.001 (0.12×10 <sup>-3</sup> to	0.005
	0.192)		0.131)		0,152)		0.133)	

Model 1 was adjusted by sex, baseline hypertension, current smokers, diabetes mellitus, hyperlipidemia, and by changes in BMI, systolic BP, T-cholesterol, and HbA1c values Model 2 was Model 1 adjusted by macroalbuminuria and eGFR. Model 3 was Model 2 adjusted by serum levels and PBCs mRNA levels of TNF/IL10. Abbreviations: PTF, pentoxifylline; KL, Klotho; BMI, body mass index; BP, blood pressure; HbA1c, glycated hemoglobin

Our results point to PTF as a kidney and vascular protective therapy in patients with T2DM and moderate to severe CKD. Whereas intermittent claudication is the main indication for PTF therapy, several studies suggests that this drug may have a broader role to play in the promotion of kidney and cardiovascular health. The PREDIAN trial [20] is the most important randomized clinical trial (RCT) evaluating the kidney protective effects of PTF in patients with T2DM and CKD stages 3-4 and residual albuminuria despite RAS blockade. Similar to the results of the present work, patients randomized to PTF in the PREDIAN trial had a smaller decline in eGFR (3.9%) and a higher reduction in albuminuria (12.5%) as compared to patients assigned to the control group (eGFR decreased by 9% and albuminuria increased by 4.4%; *P*<0.01 and *P*<0.001 between groups, respectively).

From the cardiovascular perspective, studies with PTF are scarce. Experimental works show that oral administration of PTF (40 mg/kg) to rabbits fed with a cholesterol-enriched diet to induce atherosclerotic plaque decreased the area of aortic atherosclerotic plaque by 38% without changes in serum lipids [23]. In the clinical scenario, two 6-months RCTs evaluated the effects of PTF in patients experiencing transient ischemic attacks (TIAs) compared with a control group receiving aspirin+dipyridamole [24, 25]. During the follow-up, proportion of recurrent TIAs was 14% in the PTF group, and 24.1% in the group receiving aspirin+dipyridamole. However, to our knowledge, none of these studies evaluated the progression of SA. In our study, we employed two methods widely used for assessing the presence of SA. ABI is considered as an indirect indicator of general atherosclerosis and an independent predictor of cardiovascular and all-cause mortality in CKD patients [26, 27]. CIMT is a widely accepted direct marker of atherosclerotic disease independently associated with increased cardiovascular risk, impaired kidney function, and longterm mortality in different stages of CKD [22, 27, 28].

Improved outcomes of PTF may be related with its anti-inflammatory effects. Clinical trials evaluating the anti-inflammatory properties of this drug report considerable modulating effects on the production of inflammatory cytokines and adhesion molecules in patients with acute coronary syndrome (ACS) and atherosclerosis [29–31]. In line with the results of our present work, administration of PTF (1200 mg/day) to patients with ACS reduced both hs-CRP and TNF $\alpha$  levels [29, 30]. In CKD patients, where inflammation is prevalent and constitutes a major cardiovascular risk factor, the anti-inflammatory effects of PTF have been also determined [20, 32, 33]. In the PREDIAN trial, inflammatory markers were reduced in the PTF group; specifically, urine TNF $\alpha$  decreased by a 10.6%, with no changes in the control group [20].

The regulation of KL expression in CKD after exposure to PTF has been explored in clinical and experimental studies. In a post-hoc analysis of the PREDIAN trial, serum (5.9%; P<0.05) and urine (9.3%; P<0.001) KL levels increased after 12 months of PTF treatment; by contrast, KL decreased by 0.2% in serum and by 3.9% in urine in control subjects [21]. The effects of PTF on KL expression in kidney tubular cells were also determined, finding a direct dose-dependent increase of KL protein and mRNA levels after treatment [21].Experimental and clinical studies suggest that both reduced serum an PBCs-mRNA levels of KL might play a role in the pathogenesis of CVD [10, 12-19, 34]. However, there are few works on the relationship between KL and SA markers. Polymorphisms of KL have been associated with increased CIMT in hypertensive patients [35] and with the risk of early-onset occult coronary artery disease [36]. In CKD patients, decreased soluble KL levels are associated with increased brachial-ankle pulse wave velocity (adjusted OR=0.60; 95% CI,0.39 to 0.98, P=0.0075) [16]. We recently observed an association between reduced KL levels in serum and mRNA expression in PBCs with the presence of SA in CKD patients, being directly related with ABI and inversely with CIMT (P<0.0001 for both) [19].

Macrophages, monocytes, lymphocytes, and other PBCs play a central role in the development of the inflammatory response associated with the atherogenic process. Therefore, the modulation of these cells constitutes an interesting approach for developing new therapeutic strategies in atherosclerosis. Marked reductions in KL expression in PBCs has been related with aging and with the development of pathologies with an inflammatory component including atherosclerosis [19, 37–39]. In a cross-sectional case-control study recently published by our group, which included patients with atherosclerosis and a control group of cadaveric organ donors without a history of CVD, we found significantly lower gene expression of KL in PBCs in the first group (56.4% difference, P < 0.001) and a higher methylation of the promoter region of KL ( $34.1 \pm 4.1\%$  vs.  $14.6 \pm 3.4\%$ , P<0.01) [10]. Although the intimate mechanisms of action deserve further investigation, the up-regulation of KL in these cells could antagonize the progression of atherosclerotic lesions by exerting anti-inflammatory effects, suppressing the stress response of the Golgi apparatus and the endoplasmic reticulum, reducing the levels of oxidant radicals, and preserving the immune function in senescent monocytes [40, 41, 42, 43]. All these mechanisms play key roles in the inflammatory response exerted by PBCs in the atherosclerotic process and, therefore, make KL expression in these cells an interesting target in such scenario.

Although our study provides novel information about the potential benefits of PTF on the progression of SA in patients with T2DM and CKD, we acknowledge several limitations. First, this was a single-center study, and therefore, generalizability and reproducibility will require further validation. Second, the study was not designed in a double-blinded fashion, so the open-label design may have inherent bias. In addition, because this study was an independent clinical trial, a placebo was not used in the control group as a result of limited resources. Nevertheless, the main study outcomes were performed blinded to the study group allocation of patients. Third, although the sample size needed to detect differences was calculated, we recognize that the small sample size is a limitation of this study. Finally, serum concentrations of vitamin D, fibroblast growth factor-23, and parathyroid hormone -factors related to KL and calcium/phosphate metabolism, with potential impact on atherosclerosiswere not measured in our study, and therefore a possible influence on the relationship between KL and CVD cannot be completely ruled out.

# Conclusions

In conclusion, PTF administration to patients with T2DM and CKD stage 3 reduced the progression of SA evaluated by the determination of CIMT. This beneficial effect was related to a stimulation of *KL* mRNA expression in PBCs. Furthermore, PTF treatment was associated with a slowing of the decline in eGFR. Whether PTF administration is an appropriate clinical approach to reduce the impact of atherosclerosis in patients with T2DM and CKD and if this is achieved through a KL-preserving effect is an intriguing question that requires further study.

#### Abbreviations

DM	Diabetes mellitus
T2DM	Type 2 diabetes mellitus
CKD	Chronic kidney disease
ESRD	End-stage renal disease
DKD	Diabetic kidney disease
PBCs	Peripheral blood cells
CVD	Cardiovascular disease
PTF	Pentoxifylline
PBCs	Peripheral blood cells
RAS	Renin-angiotensin system
BP	Blood pressure
ABI	Ankle-brachial index
CIMT	Carotid intima-media thickness
SA	Subclinical atherosclerosis
eGFR	Estimated glomerular filtrate rate
UACR	Urinary albumin-to-creatinine ratio
HDL-C	High-density lipoprotein cholesterol
LDL-C	Low-density lipoprotein cholesterol
hs-CRP	High sensitivity C-reactive protein
TNFa	Tumor necrosis factor
IL	Interleukin
KL	Klotho
GAPDH	Glyceraldehyde-3-phosphate dehydrogenase
VIF	Variation inflation factor
TIAs	Transient ischemic attacks
ACS	Acute coronary syndrome

#### **Supplementary Information**

The online version contains supplementary material available at https://doi. org/10.1186/s12933-024-02393-x.

Supplementary Material 1

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Not applicable.

#### Author contributions

J.F.N.-G., and C.M.-F. equally contributed to the study concept and design. J.D.-C., A.G.-L., C.F., and N.P.-D. processed the blood samples and performed ELISA and qPCR measurements. J.F.N.-G., J.D.-C., C.F., and C.M.-F. analyzed and interpreted the data. J.F.N.-G. and J.D.-C. performed the statistical analysis and wrote the manuscript. J.F.N.-G., J.D.-C., C.F., N.P.-D., A.G.-L., and C.M.-F. approved the final version of the manuscript for publication.

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#### Data availability

The datasets used and/or analyzed during the current study are available from the corresponding authors on reasonable request.

#### Declarations

#### Ethics approval and consent to participate

The trial was approved by the Institutional Review Board at the Hospital Universitario Nuestra Señora de Candelaria. The study protocol code is PTF-AA-TR-2009 and the trial was registered on the European Union Drug Regulating Authorities Clinical Trials (EudraCT #2009–016595–77). The validation date was 2010-03-09.

#### **Consent for publication**

All authors approve the manuscript and give their consent for submission and publication.

#### **Competing interests**

The authors declare no competing interests.

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