



Arterial and renal consequences of partial genetic deficiency in tissue kallikrein activity in humans

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Tissue kallikrein (TK), the major kinin-forming enzyme, is synthesized in several organs, including the kidney and arteries. A loss-of-function polymorphism of the human TK gene (*R53H*) induces a substantial decrease in enzyme activity. As inactivation of the TK gene in the mouse induces endothelial dysfunction, we investigated the vascular, hormonal, and renal phenotypes of carriers of the *53H* allele. In a crossover study, 30 *R53R*-homozygous and 10 *R53H*-heterozygous young normotensive white males were randomly assigned to receive both a low sodium–high potassium diet to stimulate TK synthesis and a high sodium–low potassium diet to suppress TK synthesis, each for 1 week. Urinary kallikrein activity was 50–60% lower in *R53H* subjects than in *R53R* subjects. Acute flow-dependent vasodilatation and endothelium-independent vasodilatation of the brachial artery were both unaffected in *R53H* subjects. In contrast, *R53H* subjects consistently exhibited an increase in wall shear stress and a paradoxical reduction in artery diameter and lumen compared with *R53R* subjects. Renal and hormonal adaptation to diets was unaffected in *R53H* subjects. The partial genetic deficiency in TK activity is associated with an inward remodeling of the brachial artery, which is not adapted to a chronic increase in wall shear stress, indicating a new form of arterial dysfunction affecting 5–7% of white people.

Introduction

Tissue kallikrein (TK), a serine protease synthesized in many organs, cleaves low- and high-molecular-weight kininogens, thus releasing the vasodilator peptides known as kinins (1–4). The kallikrein-kinin system is present in the endothelium and in the smooth muscle of vascular walls (5–7), where locally generated kinins have potent endothelium-mediated vasodilatory and antithrombotic properties through activation of bradykinin B₂ receptors, triggering NO release and other endothelial mediators (1, 8, 9). TK is also synthesized in large amounts in the kidney connecting tubule and cortical collecting tubule and is released in the urine and the peritubular interstitium (10). The renal kallikrein-kinin system is believed to operate in concert with the renin-angiotensin system to regulate physiologically the distribution of renal blood flow (1, 11, 12) and the metabolism of water and electrolytes (1). Urinary kallikrein activity (UKLKa) is influenced by hereditary factors (13, 14) and by dietary Na⁺ and K⁺ intake (1). Family studies have demonstrated familial aggregation of UKLKa and have suggested that a large part of the observed population variance is attributable to a major gene effect (14). We recently identified a loss-of-function polymorphism in exon 3 of the TK gene. This polymorphism changes an active-site arginine at position 53 to a histidine (*R53H*), resulting in a substantial loss of kallikrein activity in vitro (15). The *53H* allele is found at a frequency of 0.03 in

white people (15), and approximately 5–7% of the population are heterozygous for *53H*. On average, UKLKa is 50% lower in *R53H* heterozygotes than in *R53R*-homozygous subjects, reflecting the strong functional effect of the mutation (15).

The identification of a subset of subjects with genetically reduced kallikrein activity provides an opportunity to study the physiological role of the tissue kallikrein-kinin system in humans and to assess the consequences of constitutively low kallikrein activity. Interestingly, inactivation of the TK gene in the mouse does not alter BP but does induce endothelial dysfunction and a loss of flow-mediated vasodilatation, an important adaptive process controlling blood delivery to organs, in both conductance and resistance arteries (3, 7, 16).

We thus investigated the arterial, renal, and hormonal phenotypes of *53H* allele carriers. We studied these subjects on an ad libitum diet and then on controlled diets with high and low Na⁺ and K⁺ content to modulate the level of kallikrein synthesis and excretion (17, 18) and to monitor physiological adaptation to these diets. We found that a decrease in TK activity resulting from a genetic mutation in humans is associated with a newly discovered form of arterial dysfunction characterized by an inward remodeling of the brachial artery, which is not adapted to a chronic increase in wall shear stress.

Results

Consequences of the *R53H* polymorphism on UKLKa. UKLKa followed a circadian cycle with a large decrease during the night (Table 1), as reported previously (19). The changes in Na⁺/K⁺ intake were accompanied by substantial changes in the 24-hour UKLKa (Table 1): compared with the 24-hour UKLKa of subjects measured at baseline on the ad libitum diet, the 24-hour UKLKa increased by

Nonstandard abbreviations used: ANP, atrial natriuretic peptide; Di, internal diameter; GTN, glyceryl trinitrate; IMT, intima media thickness; MSR, mean shear rate; MSS, mean shear stress; TK, tissue kallikrein; UKLKa, urinary kallikrein activity; WSR, wall shear rate; WBV, whole blood viscosity.

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Table 1
UKLKa and Na⁺ and K⁺ excretion

	Baseline	Low Na ⁺ -high K ⁺ diet	High Na ⁺ -low K ⁺ diet	P value between the 2 diets
24-hour UKLKa (ΔFU/s/24 h)				
R53H (n = 10)	5228 [2084;6630]	20017 [14536;24872]	1759 [403;4950]	<0.0001
R53R (n = 30)	14534 [6700;26118]	51515 [32557;84699]	2560 [222;10773]	<0.0001
P value R53H vs. R53R	0.006	0.0045	0.26	
12-hour daytime UKLKa (ΔFU/s/12 h)				
R53H (n = 10)	4490 [1063;6286]	13460 [12052;20276]	1383 [81;3294]	<0.0001
R53R (n = 30)	9336 [3951;20051]	38119 [24806;62239]	2550 [70;10594]	<0.0001
P value R53H vs. R53R	0.053	0.006	0.16	
12-hour night-time UKLKa (ΔFU/s/12 h)				
R53H (n = 10)	15 [11;203]	3854 [735;7805]	18 [13;2437]	<0.0001
R53R (n = 30)	2171 [9;7988]	9027 [1203;27493]	14 [9;28]	<0.0001
P value R53H vs. R53R	0.07	0.13	0.28	
24-hour urine volume (ml)				
R53H (n = 10)	1995 [602;2820]	1931 [810;4225]	1914 [1476;2964]	0.60
R53R (n = 30)	1497 [720;2627]	2362 [849;4326]	2108 [1028;3473]	0.19
P value R53H vs. R53R	0.16	0.40	0.60	
24-hour urinary Na⁺ excretion (mmol/24 h)				
R53H (n = 10)	142 [110;179]	19 [14;22]	242 [210;263]	<0.0001
R53R (n = 30)	160 [133;188]	17 [12;21]	265 [226;300]	<0.0001
P value R53H vs. R53R	0.40	0.60	0.20	
24-hour urinary K⁺ excretion (mmol/24 h)				
R53H (n = 10)	67 [59;85]	104 [96;134]	38 [31;40]	<0.0001
R53R (n = 30)	69 [60;81]	128 [107;139]	40 [31;44]	<0.0001
P value R53H vs. R53R	0.70	0.30	0.50	

Data are expressed as median and interquartile range (in square brackets, separated by a semicolon).

4.5-fold (interquartile range, 2.8–7) when the subjects received the low Na⁺-high K⁺ diet and decreased by 3.2-fold (interquartile range, 1.1–22) when they received the high Na⁺-low K⁺ diet. The low Na⁺-high K⁺ diet had no influence on the circadian cycle of UKLKa, whereas the high Na⁺-low K⁺ diet amplified it by almost abolishing kallikrein excretion at night (Table 1).

The R53H subjects had significantly lower UKLKa values than did the R53R subjects. At baseline (i.e., on the ad libitum diet), the 24-hour UKLKa level was 50–60% lower on average in R53H subjects than in R53R subjects (P = 0.006; Table 1 and Figure 1), as observed previously in other populations (15). With the low Na⁺-high K⁺ diet, the absolute increment in UKLKa was smaller in the R53H subjects than in the R53R subjects, with the consequence that the difference between the genotypes was amplified (Table 1). Similarly, during the high Na⁺-low K⁺ intake, the absolute decrease in UKLKa was less pronounced in the R53H subjects than in the R53R subjects, with the consequence that the genotype-related difference was blunted and was no longer statistically significant (Table 1). The between-subject variability in 24-hour UKLKa was large for the R53R subjects in all experimental conditions and was lower for the R53H subjects. Consequently, the 24-hour UKLKa values for the 2 groups overlapped (Figure 1). UKLKa was not correlated to urine volume (not shown).

Consequences of the R53H polymorphism on brachial artery function and on BP. Figure 2 shows a representative tracing of the brachial artery diameter and shear stress measured continuously. Acute postischemic flow-mediated changes in brachial artery

diameter during hand hyperemia did not significantly differ between the 2 genotypes on either Na⁺/K⁺ diet (percent changes in artery diameter; Tables 2 and 3 and Figure 3). After 20 minutes of recovery, the endothelium-independent vasodilatory response to sublingual glyceryl trinitrate (GTN) was similar for both genotypes (percent changes in artery diameter; Tables 2

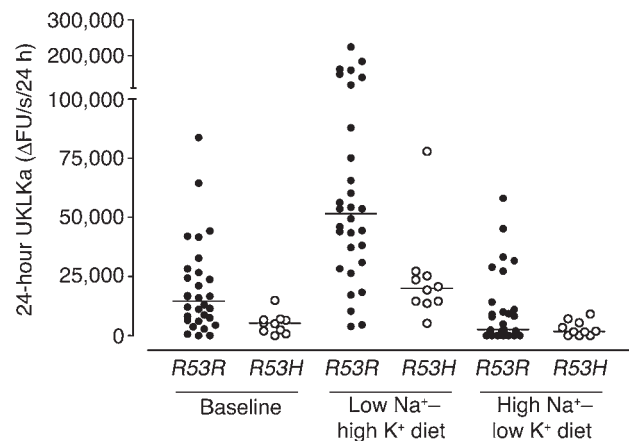


Figure 1
Effect of the R53H polymorphism of the TK gene and of the Na⁺ and K⁺ content of the diet on 24-hour UKLKa. R53R, filled circles; R53H, open circles. The horizontal lines indicate the median values.

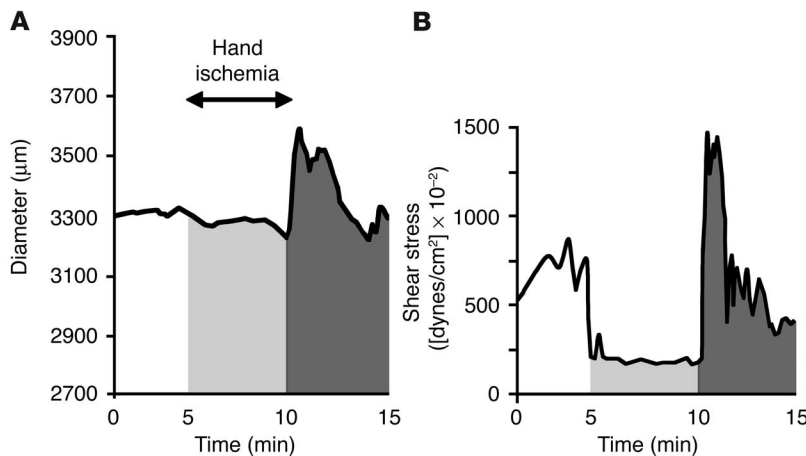


Figure 2

Evaluation of flow-mediated and endothelium-independent vasodilation. (A and B) Typical recording of the mean values of brachial artery diameter (A) and shear stress (B) obtained at baseline (white areas) and during 5 minutes of hand ischemia (light gray areas) and hyperhemia (dark gray areas) in a healthy volunteer.

and 3 and Figure 3). On average, GTN-induced vasodilation was greater than the flow-mediated vasodilation.

Despite showing similar acute vasodilatory responses to flow and GTN, the *R53H* subjects differed from the *R53R* subjects in terms of shear stress and artery morphology in all experimental conditions (baseline, hand ischemia, hand hyperhemia, and after GTN). Indeed, on the high Na⁺-low K⁺ diet, the *R53H* subjects displayed higher shear stress values and had smaller brachial artery diameter associated with similar intima media thickness, wall-to-lumen ratio, and wall cross-sectional area, thus indicating a smaller lumen than that of the *R53R* subjects (Table 2 and Figure 3). When subjects received the low Na⁺-high K⁺ diet, similar results were obtained, with the *R53H* subjects having significantly higher shear stress values in all experimental conditions than the *R53R* subjects. However, with this diet, the difference in brachial artery diameter was no longer significant between the 2 genotypes (Table 3 and Figure 3). Body surface area, known to be correlated with arterial diameter, was similar for the 2 genotypes (not shown). All results were obtained in the absence of genotype-induced differences in blood pressure (Tables 2 and 3).

Consequences of the R53H polymorphism on the renal and hormonal adaptation to the Na⁺/K⁺ diets.

On the ad libitum Na⁺/K⁺ diet, 156 mmol of Na⁺ (interquartile range, 126–187 mmol) and 68 mmol of K⁺ (interquartile range, 59–81 mmol) were excreted in the urine in 24 hours. By controlling the Na⁺/K⁺ intakes, it was possible to achieve the desired Na⁺ and K⁺ balances, as reflected by 24-hour urinary NaCl and KCl excretion rates (Table 1). The 24-hour urinary Na⁺ and K⁺ excretion levels were identical for subjects of both genotypes for all of the diets (Table 1).

Plasma active renin, prorenin, aldosterone, and atrial natriuretic peptide (ANP), and urinary aldosterone concentrations on the ad libitum

Na⁺/K⁺ diet were within the physiological range and did not differ according to genotype (Table 4). As expected, plasma active renin and prorenin, and plasma and urinary aldosterone concentrations increased with the low Na⁺-high K⁺ diet and decreased with the high Na⁺-low K⁺ diet. The changes in plasma ANP concentrations were in the opposite directions (Table 4).

Table 2

Flow-mediated and endothelium-independent vasodilatation in *R53H* and *R53R* subjects receiving the high Na⁺-low K⁺ diet

	<i>R53H</i> heterozygotes (n = 10)	<i>R53R</i> homozygotes (n = 30)	P value
Blood pressure			
Systolic blood pressure (mmHg)	115 [112;117]	113 [108;118]	0.73
Diastolic blood pressure (mmHg)	60 [57;64]	59 [56;62]	0.85
Heart rate (bpm)	57[56;66]	56 [53;62]	0.10
Flow-mediated vasodilatation			
Baseline			
Brachial artery diameter (µm)	3647 [3453;4005]	4095 [3848;4222]	0.053
Intima media thickness (µm)	411 [377;431]	405 [380;412]	0.74
Wall-to-lumen ratio (%)	22.1 [18.4;25.6]	20.2 [18.1;21.6]	0.16
Wall cross-sectional area (mm ²)	4.35 [3.93;4.82]	4.61 [4.42;5.25]	0.15
Shear stress ((dynes/cm ²) × 10 ⁻²)	399 [321;614]	296 [234;386]	0.017
Hand ischemia			
Brachial artery diameter (µm)	3640 [3355;3966]	4063 [3820;4202]	0.036
Shear stress ((dynes/cm ²) × 10 ⁻²)	271 [194;410]	158 [137;187]	0.0009
Hand hyperhemia			
Brachial artery diameter (µm)	3691 [3450;3980]	4114 [3889;4238]	0.036
Shear stress ((dynes/cm ²) × 10 ⁻²)	488 [352;756]	371 [268;461]	0.03
Change in artery diameter (%)	3.7 [1.7;4.6]	2.9 [1.6;4]	0.70
Endothelium-independent vasodilatation			
Baseline			
Brachial artery diameter (µm)	3667 [3365;3960]	4097 [3836;4261]	0.046
Shear stress ((dynes/cm ²) × 10 ⁻²)	408 [346;644]	288 [247;355]	0.005
After GTN			
Brachial artery diameter (µm)	4393 [4254;4668]	4742 [4463;5021]	0.07
Shear stress ((dynes/cm ²) × 10 ⁻²)	373 [292;526]	261 [238;338]	0.049
Change in artery diameter (%)	20 [17;25]	17 [13;21]	0.18

Data are expressed as median and interquartile range.

**Table 3**Flow-mediated and endothelium-independent vasodilatation in *R53H* and *R53R* subjects receiving the low Na⁺-high K⁺ diet

	<i>R53H</i> heterozygotes (n = 10)	<i>R53R</i> homozygotes (n = 30)	P value
Blood pressure			
Systolic blood pressure (mmHg)	115 [113;119]	114 [110;120]	0.89
Diastolic blood pressure (mmHg)	58 [56;63]	61 [59;64]	0.18
Heart rate (bpm)	64 [58;68]	59 [54;65]	0.12
Flow-mediated vasodilatation			
Baseline			
Brachial artery diameter (μm)	3724 [3458;4065]	3967 [3669;4384]	0.25
Intima media thickness (μm)	411 [377;432]	405 [380;413]	0.75
Wall-to-lumen ratio (%)	0.21 [0.19;0.25]	0.21 [0.18;0.22]	0.91
Wall cross-sectional area (mm ²)	4.38 [3.90;4.87]	4.56 [4.16;5.00]	0.51
Shear stress ([dynes/cm ²] × 10 ⁻²)	468 [339;636]	277 [226;383]	0.025
Hand ischemia			
Brachial artery diameter (μm)	3652 [3431;4031]	3942 [3600;4354]	0.20
Shear stress ([dynes/cm ²] × 10 ⁻²)	246 [179;286]	171 [128;225]	0.027
Hand hyperemia			
Brachial artery diameter (μm)	3751 [3496;4106]	4002 [3691;4398]	0.30
Shear stress ([dynes/cm ²] × 10 ⁻²)	535 [414;759]	346 [274;490]	0.02
Change in artery diameter (%)	4.4 [1.8;6.8]	2.8 [2;4.6]	0.50
Endothelium-independent vasodilatation			
Baseline			
Brachial artery diameter (μm)	3753 [3491;4130]	3979 [3725;4367]	0.30
Shear stress ([dynes/cm ²] × 10 ⁻²)	486 [452;633]	239 [217;348]	0.003
After GTN			
Brachial artery diameter (μm)	4538 [4339;4683]	4683 [4466;4914]	0.16
Shear stress ([dynes/cm ²] × 10 ⁻²)	405 [262;572]	250 [190;336]	0.027
Change in artery diameter (%)	17 [13;21]	18 [14;24]	0.88

Data are expressed as median and interquartile range.

The diet-induced changes in these hormonal parameters did not significantly differ between the *R53H* and the *R53R* subjects. The changes in 24-hour UKLKa were positively related to the changes in plasma active renin and urinary aldosterone excretion and were inversely related to changes in plasma ANP (not shown). The physiological relationship between these parameters was not influenced by the *53H* allele.

These data collectively suggest that renal and hormonal adaptation to variations in the Na⁺/K⁺ diet was normal in *R53H* subjects.

Discussion

Our results confirm that the *R53H* kallikrein gene polymorphism (15) and the Na⁺ and K⁺ content of the diet (17, 20–22) are major determinants of the level of UKLKa in physiological conditions. They also show that a 50% deficiency in TK activity that is genetically determined does not alter the capacity of the kidneys to adapt to large variations in Na⁺ and K⁺ intake, but triggers a peculiar form of arterial dysfunction.

We studied only white males because gender and ethnic origin are known to influence urinary kallikrein excretion (23, 24). We used a new kallikrein-specific enzymatic assay (25) to monitor the level of UKLKa that reflects both the activity and excretion

of the renal enzyme. We altered Na⁺ and K⁺ intake (a) to modify renal kallikrein synthesis and to study the interaction between the *R53H* polymorphism and this regulation and (b) to monitor the renal and vascular adaptations of kallikrein-deficient subjects after large variations in ionic balance. Salt restriction is known to stimulate the excretion of renal kallikrein by an unknown mechanism, possibly mediated by aldosterone (17, 18). K⁺ intake also upregulates kallikrein synthesis and excretion either directly or by triggering aldosterone secretion (10, 18). As expected, the low Na⁺-high K⁺ diet strongly stimulated renal kallikrein synthesis and excretion, whereas the high Na⁺-low K⁺ diet largely suppressed this excretion. The UKLKa varied by a factor of 20 between the 2 controlled diets. This allowed us to study the interaction between the *R53H* polymorphism and the physiological regulation of renal kallikrein.

The *53H* allele suppressed UKLKa in all conditions of Na⁺/K⁺ intake and at all kallikrein synthesis levels, consistent with the observation that the *R53H* mutation inactivates the catalytic activity of the enzyme in vitro (15). Our data also show that the diet-induced variations in UKLKa are blunted in *R53H* subjects compared with that in *R53R* subjects, as this activity varied by a factor of only 11.3 between the 2 controlled regimens in the *R53H* subjects and by more than 20-fold in the *R53R* subjects. This might be because only the non-mutated allele contributes to the catalytic activity of kallikrein in *R53H* subjects. These observations indicate that the *R53H* mutation has a considerable effect on kallikrein activity levels in vivo.

We observed that the renal adaptation to Na⁺ and K⁺ intake was not impaired in *R53H* subjects, as their urinary electrolyte and blood pressure levels after 7 days of controlled Na⁺/K⁺ regimens were similar to those of the *R53R* subjects. Furthermore, the regulation of the renin-angiotensin-aldosterone system and ANP was not altered in *R53H* subjects. These results suggest that despite its abundance and high turnover rate in the kidney tubule (26) and its strong regulation by Na⁺ and K⁺ intake, kallikrein does not play a major role in the renal adaptation to these dietary ions, at least in the short term. An alternative explanation could be the persistence of enough residual kinin generating activity due to kallikrein in *R53H* heterozygous subjects and/or a physiological readjustment of the kininogen-kallikrein-kinin system. Changes in Na⁺ diet in both rats and humans have indeed been shown to induce opposite changes in low-molecular-weight kininogen and in UKLKa, resulting in unchanged renal kinin concentrations (27–29). Investigation of homozygous *53H* subjects should theoretically resolve this question, but no such subjects have been identified so far. The low frequency of the *R53H* polymorphism indicates that only 1 in 2,000 white people are homozygous for the mutation. Interestingly, the renal adaptation to large variations in Na⁺ intake was

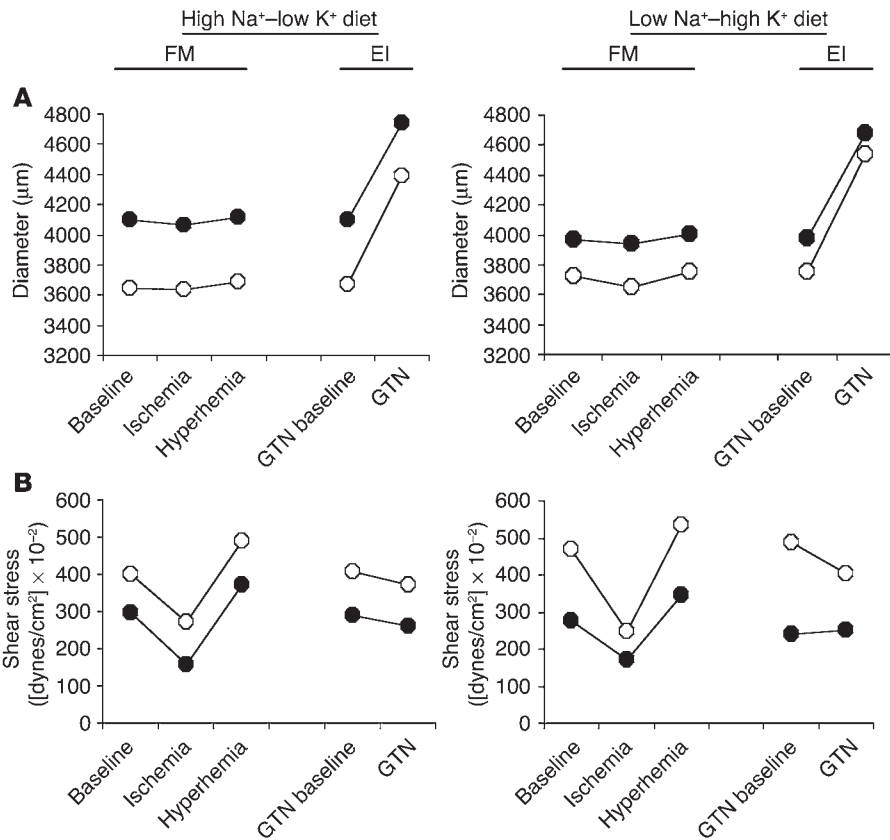


Figure 3 Diameter and shear stress responses of the brachial artery to flow and GTN administration. (A and B) Flow-mediated (FM) and endothelium-independent (EI) changes in brachial artery diameter (A) and shear stress (B) according to genotype and to the Na⁺ and K⁺ content of the diet. *R53R*, filled circles; *R53H*, open circles.

found to be normal in the homozygous TK-deficient mouse (N. Picard et al., unpublished results), further suggesting that kallikrein has little, if any, role in this function.

Finally, although renal kallikrein has been suggested to be a prorenin-activating enzyme (30), the fact that plasma prorenin and active renin concentrations were not influenced by the *R53H* polymorphism, despite a markedly reduced UKLKa, does not suggest a physiological role for kallikrein in prorenin activation. However, because of residual kallikrein activity in *R53H* subjects, the possibility of this role cannot be completely ruled out.

By showing that the activity and regulation of the renin-angiotensin-aldosterone system were not altered in kallikrein-deficient subjects, our study reveals a physiological imbalance between the renin-angiotensin-aldosterone and kallikrein-kinin systems in these subjects. Although this physiological imbalance appears to have no consequence on the short-term renal adaptation to dietary ion intake in young male adults, it may have long-term consequences. This hormonal imbalance has indeed been observed in hypertensive patients and their normotensive offspring (31). Low UKLKa has also been associated with “non-modulating” salt-sensitive hypertension, a phenotypic trait characterized by failure to modulate renal blood flow in response to angiotensin II after high Na⁺ intake (32). The *R53H* polymorphism may therefore predispose, in the long term, to hypertension and

renal disorders associated with abnormal regulation of renal blood flow, such as diabetic nephropathy (33).

Vascular consequences of the R53H polymorphism. The main finding of this study is the occurrence of arterial dysfunction in *R53H* subjects.

We first showed that the acute physiological vasodilatory response to flow and the effect of direct stimulation in the smooth muscle of the cGMP-dependent pathway by GTN did not significantly differ between the 2 genotypes. Contrasting with those findings, we observed that compared to the *R53R* subjects, the *R53H* subjects exhibited a permanent chronic increase in shear stress associated with paradoxical reductions of the lumen and diameter of the brachial artery. The increase in shear stress was consistent across all experimental conditions. The reduction of the artery diameter was especially evident on the high Na⁺-low K⁺ diet. This reduction in diameter or the simple absence of a widening of the artery despite a chronic increase in shear stress in the *R53H* subjects indicates an inappropriate inward remodeling of the brachial artery in these subjects. The similarity of the intima media thickness, the wall-to-lumen ratio, and the wall cross-sectional area between the 2 genotypes combined with the reduction in both the lumen and diameter of the brachial artery indicates that this artery under-

went eutrophic remodeling in *R53H* heterozygous subjects. When subjected to long-term changes in blood flow, arteries remodel themselves through endothelium-dependent mechanisms, thus limiting changes in wall shear stress (34). When subjected to an increase in blood flow, vascular endothelial cells release growth factors and vasoactive substances able to induce cell proliferation, migration, death, and matrix deposition (35, 36). In general, chronic increases in blood flow lead to outward arterial remodeling, with a widening of the lumen diameter, thus normalizing wall shear stress (37, 38), whereas sustained reductions in blood flow lead to inward arterial remodeling with a narrowing of lumen diameter (34). Thus, the higher wall shear stress in *R53H* than in *R53R* subjects, together with a paradoxical reduced internal diameter, indicates a defect in the endothelium-dependent pathways involved in arterial remodeling (35).

Our results thus show a dissociation in *R53H* subjects between the acute endothelium-dependent flow-mediated response of the brachial artery (39), which appeared unchanged, and the chronic shear stress-induced endothelium-dependent arterial remodeling (35, 36), which appeared to be inappropriate. It remains possible that acute endothelial dysfunction occurs in *R53H* subjects but is too mild to be detected in our experimental conditions, while having significant long-term consequences on arterial remodeling. Furthermore, acute endothelial dysfunction linked to genetic

**Table 4**
Hormonal parameters

	Baseline	Low Na ⁺ - high K ⁺ diet	High Na ⁺ - low K ⁺ diet	P value between the 2 diets
Plasma active renin (pg/ml)				
<i>R53H</i> (n = 10)	12 [9;17]	41 [27;47]	5 [4;9]	<0.0001
<i>R53R</i> (n = 30)	15 [11;22]	36 [30;48]	9 [6;15]	<0.0001
P value <i>R53H</i> vs. <i>R53R</i>	0.14	0.9	0.0502	
Plasma prorenin (pg/ml)				
<i>R53H</i> (n = 10)	130 [115;143]	150 [136;194]	87 [53;106]	<0.0001
<i>R53R</i> (n = 30)	90 [74;114]	109 [73;145]	60 [47;101]	<0.0001
P value <i>R53H</i> vs. <i>R53R</i>	0.046	0.11	0.5	
Plasma ANP (pg/ml)				
<i>R53H</i> (n = 10)	21 [18;23]	16 [15;19]	28 [25;35]	<0.0001
<i>R53R</i> (n = 30)	20 [18;26]	16 [14;20]	21 [18;28]	<0.0001
P value <i>R53H</i> vs. <i>R53R</i>	0.95	0.7	0.07	
Plasma aldosterone (pg/ml)				
<i>R53H</i> (n = 10)	68 [63;83]	327 [254;430]	30 [24;38]	<0.0001
<i>R53R</i> (n = 30)	70 [58;89]	309 [253;416]	40 [26;55]	<0.0001
P value <i>R53H</i> vs. <i>R53R</i>	0.9	0.95	0.15	
24-hour urinary aldosterone excretion (μg/24 h)				
<i>R53H</i> (n = 10)	7.5 [6.5;12.1]	51 [36;65]	2.5 [1.3;3.9]	<0.0001
<i>R53R</i> (n = 30)	9 [7.1;11.6]	43 [36;51]	2.2 [1.5;3.3]	<0.0001
P value <i>R53H</i> vs. <i>R53R</i>	0.9	0.2	0.8	

Data are expressed as median and interquartile range.

variations may be apparent only in some vascular beds; for example, at the femoral but not at the brachial artery site (40).

The observation of an abnormal arterial phenotype in subjects with a 50% deficiency in kallikrein activity is consistent with the hypothesis that the kallikrein activity level is low in arteries compared to the kidney tubule, but nevertheless plays a critical role in arterial function (3, 7). The physiological regulation of vascular kallikrein remains largely unknown.

Like in all human genetic association studies, we cannot completely rule out the possibility that the phenotype observed was due to a genetic variation in another gene located at the same locus. However, the observation of arterial dysfunction in mice carrying a specifically inactivated TK gene (3, 7, 16) strongly suggests that the arterial phenotype observed in *R53H* subjects is causally linked to kallikrein deficiency. Given the recent observation of impaired cardioprotection against ischemia in TK-deficient mice (V. Griol-Charhbili et al., manuscript submitted for publication), it would be interesting to study whether TK-deficient humans are at increased risk of cardiovascular diseases.

Methods

Subjects

We calculated that 10 *R53H* heterozygous subjects and 30 *R53R* homozygous subjects were necessary to detect a 4% difference in acute flow-mediated dilatation between *R53H* and *R53R* subjects, with a SD of 2.9%, an α risk of 5% and a β risk of 20%.

We therefore screened 206 white male subjects 18–35 years of age for the *R53H* genotype and the following inclusion criteria: nonsmoker, normotensive (BP less than 140/90 mmHg in the supine position after 5

minutes of rest), normal clinical examination, normal plasma creatinine and serum cholesterol concentrations (less than 100 μmol/l and less than 5.2 mmol/l, respectively), and no proteinuria. Among the 11 *R53H* heterozygotes (5.3%) detected, all except 1 smoker were included in the study. Among the remaining *R53R*-homozygous subjects, we included the first 30 consecutive subjects who matched all the inclusion criteria. All subjects completed the study after giving written informed consent. The protocol was approved by the “Comité Consultatif de Protection des Personnes se prêtant à des Recherches Biomédicales” (Paris-Cochin, France). All procedures were in accordance with institutional guidelines.

Study design

UKLKa and hormonal parameters were first studied for all participants while they received an ad libitum diet. Subjects were then randomly assigned to receive the first of 2 experimental diets for 7 days: (a) low Na⁺ (less than 20 mmol NaCl/day) and high K⁺ (more than 140 mmol KCl/day; low Na⁺-high K⁺ diet) or (b) high Na⁺ (more than 250 mmol NaCl/day) and low K⁺ (less than 50 mmol KCl/day; high Na⁺-low K⁺ diet). The order of assignment of the diets was randomized according a crossover study design. Controlled Na⁺/K⁺ diet periods were separated by a 7-day wash-out period. Protein and calorie intakes were kept constant during the 2 periods. All meals were provided by the metabolic kitchen of the Hôpital Européen Georges Pompidou Clinical Investigation Center and were taken in the unit. On the ad libitum Na⁺ and K⁺ diet and on day 7 of each controlled Na⁺/K⁺ diet period, blood was sampled at 0900 in the fasting state after 1 hour of rest in the sitting position for plasma active and total renin and plasma aldosterone and ANP determinations. Urine was collected in 2 12-hour periods from 0800 to 2000 and from 2000 to 0800 and was used for hormone and electrolyte determinations. BP (the mean of 10 measurements performed at 2-minute intervals after 40 minutes of rest in the sitting position) was determined using an automatic validated BP recorder (Press Mate BP 8800; Colin Co.).

Noninvasive measurement of flow-mediated, endothelium-dependent vasodilatation

Endothelial function was evaluated by comparison of the endothelium-dependent vasodilatation in response to flow induced by release of a wrist cuff after a period of 5 minutes hand ischemia versus the endothelium-independent response to sublingual GTN (39).

On day 7 of each controlled Na⁺/K⁺ diet, after a light, caffeine-free breakfast at 0930, subjects were instructed to lie down in a temperature-controlled room (21 °C to 23 °C) with their right arm secured comfortably on a support. The diameter and wall thickness of the brachial artery and wall shear rate were continuously measured with a high-resolution radio frequency-based vessel wall tracking system (Wall Track System; Esaote Pie-Medical) using a 7.5-Mhz ultrasound probe. Internal diameter (Di) and intima media thickness (IMT) were measured as described previously (41, 42). Briefly, this system uses optimized algorithms to improve the precision of interface localization, from stored high-frequency digitization of the radio frequency signal generated by reflection of ultrasound on vascular structures. The derived parameters are wall-to-lumen ratio ($2 \times \text{IMT}/\text{Di}$) and wall cross-sectional area ($\pi \times [(\text{Di} + 2 \times \text{IMT})^2 - \text{Di}^2] / 4$).



Wall shear rate (WSR) in the brachial artery was determined directly (39, 43) from the radial derivative of the velocity distribution, obtained with a high-resolution multigate pulsed Doppler ultrasound system incorporated into the Wall Track system, allowing high-precision assessment of low blood flow velocities close to the wall. This method does not make the assumptions required by the Hagen-Poiseuille equation (39). Whole blood viscosity (WBV; mPa × s) was estimated using the formula derived from Weaver (44–46), using hematocrit, plasma viscosity, and WSR. Mean shear stress (MSS; [dynes/cm²] × 10⁻²) was then calculated as MSS = mean shear rate (MSR; s⁻¹) × WBV. MSR and MSS were averaged within each cardiac cycle and were acquired during 4-second periods.

Diameter and shear stress were measured at baseline (after 15 minutes of rest in the supine position), during 5 minutes of hand ischemia, during 60 and 120 seconds of reactive hand hyperemia (produced by releasing a pneumatic wrist cuff inflated to 50 mmHg above the systolic BP for 5 minutes), and 3 minutes after the administration of 150 µg GTN (Natispray 0.15 mg/dose; Procter & Gamble Pharmaceuticals). Post-ischemic, flow-mediated, endothelium-dependent vasodilatation was defined as the maximal increase in the brachial artery diameter during reactive hyperemia, determined as the moving average of 3 consecutive diameter measurements. All hemodynamic measurements returned to baseline 20 minutes after hand hyperemia.

The interobserver variations in baseline diameter and percentage of post-occlusion vasodilatation were 0.7% and 19%, respectively. To reduce bias in the reading and interpretation of the hemodynamic results, all measurements were recorded on a hard drive using random codes and were analyzed independently by 2 readers who were blinded to the genotype, the diet, and the temporal order of the measurements.

Laboratory methods

R53H genotyping. The initial population (n = 206) was genotyped for the R53H polymorphism by the mutagenically separated PCR technique (15). Each participant was genotyped a second time using a second blood sample and by direct sequencing of exon 3 of the *hKLLK1* gene.

Urinary kallikrein measurements. UKLKa was quantified by an amidolytic assay with the chimeric, fluorogenic peptide substrate D-PFF-Nmec (a gift from F. Gauthier, INSERM-François Rabelais University U10, Tours, France), as described previously (25). This assay is highly specific for human TK, and the substrate is not cleaved by other kallikrein-related enzymes (15, 25). Briefly, hydrolysis of D-PFF-Nmec was measured with a Spectra Max Gemini microplate reader (Molecular Devices) by incubation of 0.02 ml of urine for 15 minutes at 37°C with 0.01 mM D-PFF-Nmec in 200 µl of 20 mM Tris-HCl buffer (pH 9.0) containing 1 mM ethylenediaminetetraacetate, as described previously (15, 25). Results are expressed in the

change in fluorescent units per second of incubation, for the total urine collected (ΔFU/s/12- or 24-hour urine).

Hormone measurements. The methods used for collecting blood samples and for quantifying plasma active renin, total renin, ANP and aldosterone and urinary aldosterone extractable at a pH of 1 were as described previously (47). Prorenin level is calculated as the difference between total renin and active renin levels. All enzyme and hormone determinations were performed by experimenters blinded to genotype and order of diet.

Statistical analysis. We compared R53H and R53R subjects within each period of diet using the Mann-Whitney U-test. We compared the effects of diet on UKLKa and on plasma and urinary hormones using the non-parametric Wilcoxon paired test. Stata Statistical Software (version 7.0; StataCorp.) was used for statistical analysis. Data are expressed as medians and interquartile ranges. A P value of less than 0.05 was considered to be significant.

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