

# Evaluation of Renal Blood Flow using PCASL in Chronic Kidney Disease patients and healthy controls: preliminary results from

**Primary:** Contrast Mechanisms - Arterial Spin Labelling   **Secondary:** Body - Kidney   **PowerPitch Oral** · 2 min + **Digital Poster** · 60 min | [MRI of Renal Function](#) · 2:10 PM–2:12 PM · Session: 1:40–3:16 PM · Power Pitch Theatre 1   **Keywords:** RENAL BLOOD FLOW PCASL CHRONIC KIDNEY DISEASE RENAL PERFUSION ARTERIAL SPIN LABELING (ASL)

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## Impact

Harmonized PCASL sequences show promise for renal blood flow measurements in CKD patients and controls in a multi-center and multi-vendor study. RBF is significantly reduced in patients with CKD stage 3 compared to CKD Stage 2 patients and controls

## Synopsis

**Motivation:** Chronic kidney disease (CKD) is associated with reduced renal blood flow (RBF). ASL can estimate RBF, however ASL data acquired with harmonized sequences are limited.

**Goals:** Evaluate RBF differences in CKD patients and healthy volunteers (HV) using harmonized PCASL at 3T systems of different vendors.

**Approach:** HV and CKD (Stage 2 and 3) patients were scanned at two centers using harmonized PCASL sequences across vendors. RBF maps were computed and used in a general linear model to assess differences among groups.

**Results:** RBF values: HV (n=40): 229±50ml/min/100g, CKD2(n=20): 235±41ml/min/100g, CKD3(n=20): 185±53 ml/min/100g. RBF was significantly reduced in CKD3 compared to CKD2 and HV.

## Introduction

Chronic kidney disease (CKD) affects over 800 million people globally, with higher prevalence in older adults, women, and individuals with diabetes or hypertension<sup>1</sup>. It is a leading cause of death and one of the few silent diseases with rising mortality<sup>2</sup>.

As CKD progresses toward end-stage renal disease, altered renal hemodynamics result in diminished perfusion, leading to reduced renal blood flow (RBF)<sup>3</sup>.

Arterial Spin Labeling (ASL) can provide contrast-free assessment of kidney perfusion<sup>4</sup>. Despite promising single-center and single-vendor studies, methodological variability limits its clinical adoption.

This multicenter and multivendor study, conducted as part of the European RESPECT project, aims to characterize and stage CKD using harmonized pseudo-continuous arterial spin labeling (PCASL) sequences in healthy volunteers (HV) and CKD patients.

## Methods

The study was approved by local Ethics Research Committees. Written informed consent was obtained from all subjects.

## Study Design:

Two centers participated in this study, providing MRI data from 3T scanners of two different vendors. Twenty HV and 20 CKD patients (stage 2 and 3) were recruited at each center by participating nephrologists.

HVs were scanned twice on the same day (exams 1, 2), followed by a third scan after 7–14 days (exam 3), while CKD patients were scanned twice within 7–14 days. Blood and urine samples were collected on the first visit. Estimated glomerular filtration rate (eGFR) was calculated using the 2021 CKD-EPI equation based on creatinine<sup>5</sup>.

## MRI Protocol:

ASL sequences employed optimized PCASL configurations. For vendor 1, a research PCASL sequence with FSE readout was used<sup>6</sup>, while for vendor 2, a research PCASL sequence with SE-EPI readout.<sup>7</sup> Sequences were harmonized in terms of labeling parameters, post-labeling delay, labeling duration, the use

of background-suppression pulses and 2D readouts, following PARENCHIMA recommendations<sup>4</sup>. Sequence details are in [Figure 1](#).

### ASL Postprocessing:

CKD and HV data from exam 1 were analyzed. ASL images were motion-corrected using non-rigid transformation algorithms. RBF was quantified using the single-compartment model.<sup>4</sup> The renal cortex was manually segmented in the central slice of the M0 image or T1 maps, avoiding cysts and artifacts.

### Statistical Analysis:

Statistical analysis was performed using R. Normally distributed data are expressed as mean±standard deviation, and non-normally distributed data as median (interquartile range). Age, serum creatinine, eGFR, and urine albumin-to-creatinine ratio (UACR) were compared across groups using one-way analysis of variance (ANOVA) with a between-subject factor: group (3 levels = HV, CKD2, CKD3), while sex differences using Fisher's exact test.

To assess the ability of cortical RBF to distinguish between HV, CKD2, and CKD3 patients, a general linear model (GLM) was employed with group (3 levels) as the independent variable, and age, sex, and center as potential confounding factors<sup>8</sup>. GLM assumptions were checked.

In all analyses, a p-value < 0.05 was considered statistically significant, with correction for multiple comparisons where applicable.

### Results and discussion

Data from 40 HV, 19 CKD2, and 21 CKD3 patients were analyzed. Demographic and clinical characteristics are summarized in [Figure 2](#). All parameters were normally distributed except for UACR. No significant differences were found in age or sex among groups, however age was slightly elevated in CKD3 group. Significant differences among groups were observed in creatinine and eGFR. UACR was significantly lower in healthy controls compared to CKD2 and CKD3 patients.

[Figure 3](#) shows RBF maps from representative subjects. Boxplots of eGFR and RBF measurements are shown in [Figure 4](#). RBF values ([Figure 2](#)) for all groups were similar across both centers and consistent with the literature<sup>7</sup>. Group mean RBF values: HV=229±50 ml/min/100g, CKD2=235±41 ml/min/100g, CKD3=185±53 ml/min/100g.

GLM analysis revealed a significant main effect of group (p-value=0.0092), after adjusting for confounding variables. Age was a significant co-variate (p-value=0.0002), while sex (p-value=0.1583) and center (p-value=0.0664) were not. Post-hoc comparisons showed that RBF in healthy volunteers was significantly higher than in CKD3 patients (p=0.0278) and RBF in CKD2 was significantly higher than in CKD3 (p=0.0412). No significant difference in RBF between controls and CKD2 patients was found (p-value=0.79).

Limitations should be noted. Motion correction, which is critical for accurate RBF estimation, was not consistently successful during all the free-breathing acquisitions. Cortical segmentation were drawn manually, which may have introduced a bias in the measurements. Sample size was relatively small; however, recruitment and data collection are still in progress. Future work will include automatic/semi-automatic approaches in anatomical images co-registered with ASL data. Additionally, the data of all the centers and exams will be analyzed.

### Conclusion

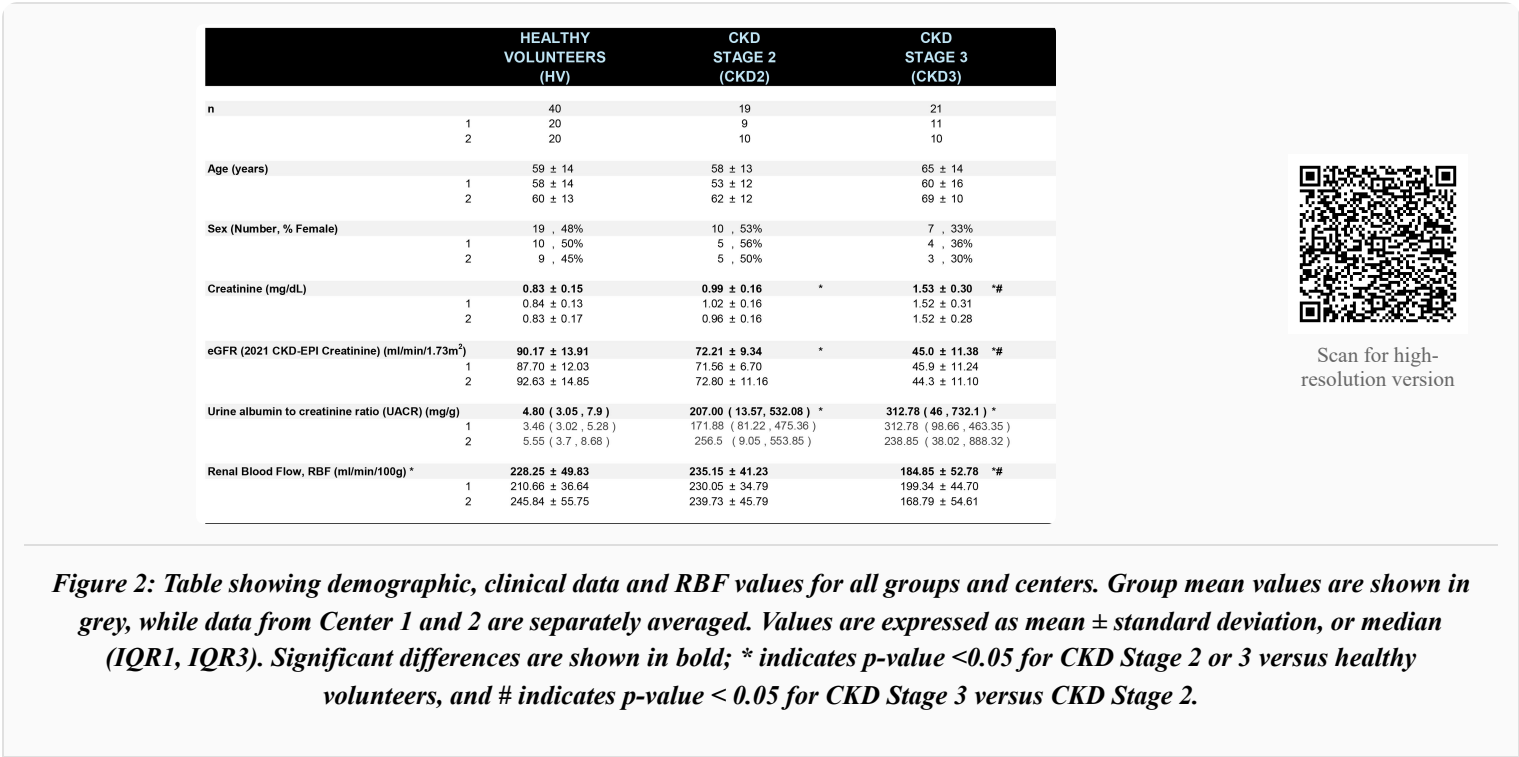
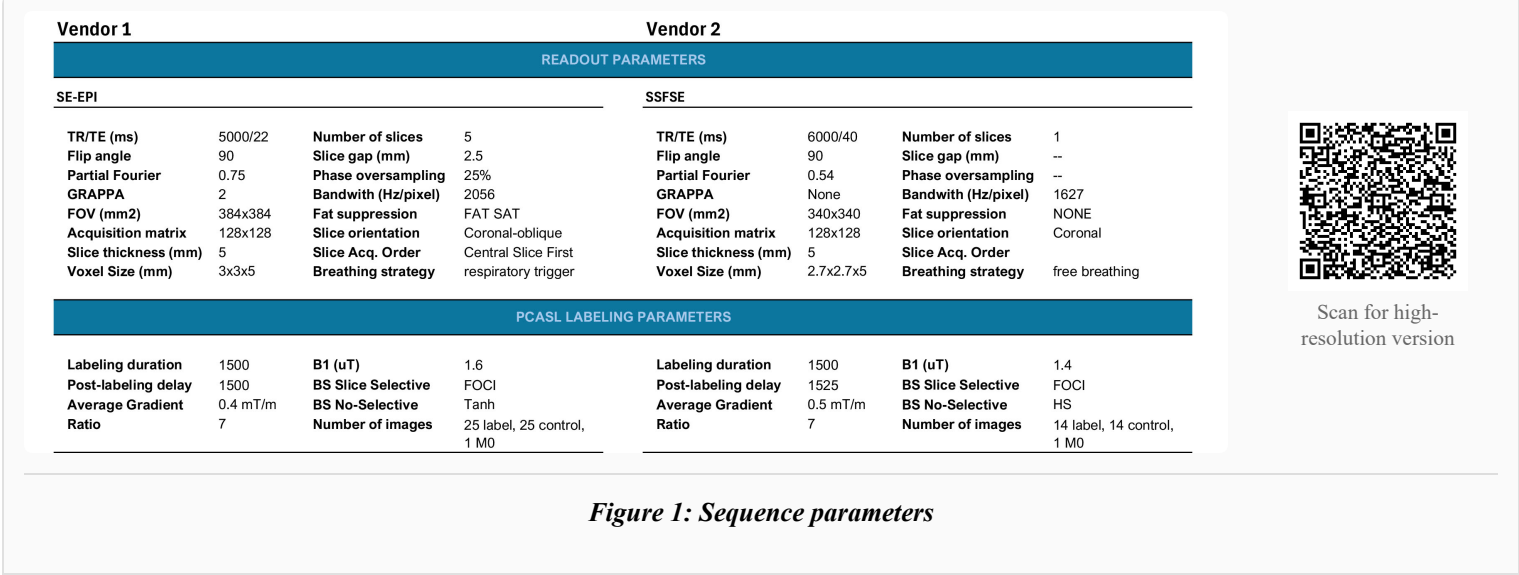
In conclusion, RBF measurements with harmonized sequences among centers showed similar RBF values across vendors. RBF at Stage 3 of CKD is significantly reduced compared to healthy controls and Stage 2 of CKD.

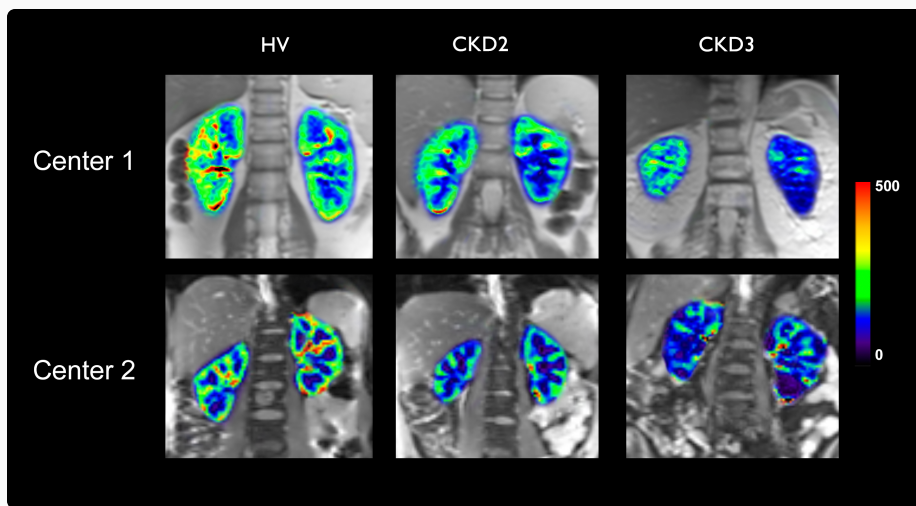
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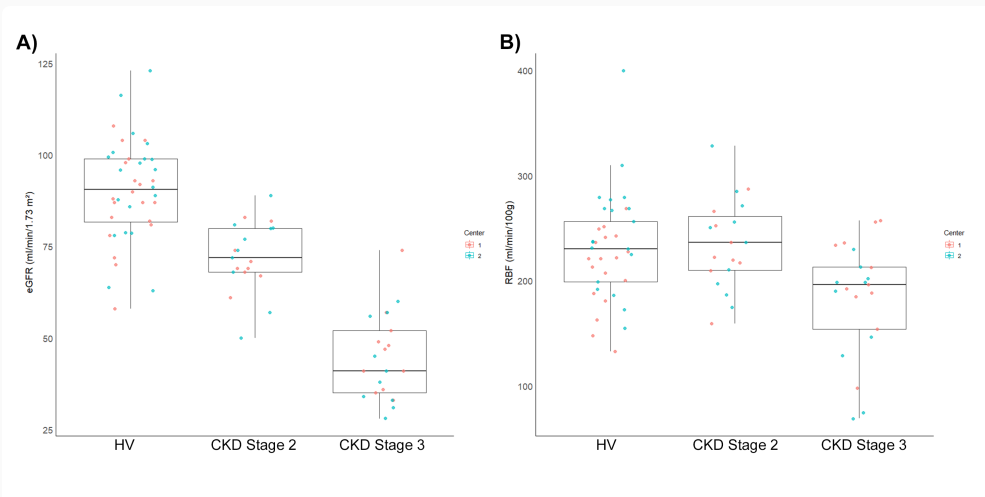
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**Figure 3: RBF maps of a representative subject of each group (HV, CKD2 and CKD3) and center (1,2).**



**Figure 4: Group mean boxplots for eGFR (panel A) and RBF (panel B) for each group and center.**