## Renal Diffusion and Perfusion in Cardiorenal Syndrome.

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**Background** Despite its clinical importance, the pathophysiology of chronic cardiorenal syndrome (CRS) is currently not fully understood [1]. Renal dysfunction (RD) is one of the most important independent risk factors for poor outcomes and all-cause mortality in patients with congestive heart failure (HF). However, it is not understood whether this renal dysfunction is caused by changes in tissue structure (fibrosis) or haemodynamic changes.

<u>Aim</u> To assess MR diffusion and perfusion of the kidney in patients with cardiorenal syndrome compared to healthy volunteers (HV), to assess pathophysiology. <u>Methods</u> 35 subjects were enrolled into four groups: Group 1: 10 HV <40yrs; Group 2: 10 HV >50yrs (BSA corrected ejection fraction, EF=36±2%); Group 3: 8 stable

HF without RD (57-82yrs, eGFR>60mL/min, EF=16±2%); Group 4: 7 stable HF with RD (62-78yrs, eGFR<60mL/min, EF=25±3%). All imaging was performed using a 1.5T Philips Achieva scanner (body transmit coil, 16-channel SENSE torso receive coil) in a single session. Multi-slice True-FISP data was acquired in 3 orthogonal planes to locate organs/vessels. *Diffusion*: Respiratory triggered spin echo (SE) DWI data (288×324 mm FOV, 5 or 10 coronal-oblique slices (3×3x8 mm or 3×3x5 mm voxel) was collected with an EPI readout (TR/TE = 3.2 s/71 ms) at 10 b-values (b = 0, 50, 100, 150, 200, 250, 300, 500, 750, 800 s/mm<sup>2</sup>) and 6 orthogonal diffusion directions. *Perfusion*: Respiratory-triggered ASL data (matched in geometry to DWI, label delay = 1100 ms, in-plane pre-saturation, 30 pairs) were collected with a True-FISP readout (TE/TR 2.1/4.1 ms, SENSE 2, FA 60°, centric half-Fourier acquisition) [2]. Base magnetization images and inversion recovery data were acquired to form M<sub>0</sub> and T<sub>1</sub> maps. *Vessel Flow*: PC data were collected for both renal arteries (RA) using a single slice TFE sequence perpendicular to each vessel (15 phases, TR/TE 6.9/3.7 ms, FA 25°, NEX 2, 1.17x1.17x6 mm<sup>3</sup>, TFE factor 4-6 (dependent on subjects' heart rate), V<sub>ENC</sub> = 100cm/s, single breath hold per vessel).

Data Analysis: Diffusion: DWI images were averaged across directions for each b-value. A renal cortex mask was created for each kidney/slice by thresholding the T1

map. This was used to generate average DW cortex data for each b-value. The ROI data was then fit to a monoexponential diffusion model for ADC, and a biexponential model for D, D\* and  $f_p$ , the product of D\* and  $f_p$  was also calculated as a marker of total flow. Monte Carlo simulations (1000 repeats, SNR 50:1) were performed simulating the diffusion data (ADC: 2.4, D: 1.8, D\*: 15x10<sup>-3</sup> mm<sup>2</sup>/s,  $f_p$ : 30 %) to assess the accuracy and standard deviation of the mean fitted value.

*Perfusion*: ASL images were motion corrected to the base  $M_0$  image using FSL (FMRIB Software Library) and difference images (label-control) calculated and averaged to create a single difference map ( $\Delta M$ ). A perfusion map was formed using a kinetic model using the individuals'  $\Delta M$ , base  $M_0$  image and tissue  $T_1$  [3]. The renal cortex mask was used to calculate the mean perfusion across both kidneys. *Vessel Flow*: Philips Q-flow software (Philips Medical

Systems) was used to draw a region of interest (ROI) over the vessel, and the mean area, flow velocity (cm/s) and flux (ml/s) over the cardiac cycle, across the vessel, calculated.

*Statistics*: (SPSS 18) Independent paired tests were made for







ADC, D, D\*,  $f_p$ ,  $f_pD*$ , flux and perfusion for groups 1v2 (HV <40yrs and >50yrs), groups 3v4 (HF with and without RD) and groups 2v3 (HV v HF w/o RD). Pearson correlation coefficients (R) were assessed.

## Results

Figure 1 shows the cortex ADC, D, D\*,  $f_p$ ,  $f_pD*$ , renal artery flux and renal cortex perfusion, with a reduction in ADC,  $f_pD*$ , flux and perfusion across groups 1-4. Monte carlo simulations show an accuracy of (-0.1,1.5, 5.9,3.0, 2.3)% and stdev of (2.6,9, 31, 18, 12)% for ADC, D, D\*,  $f_p$  and  $f_pD*$  respectively. There is a significant difference in ADC, D, D\*,  $f_pD*$  and flux between the two HV groups, and between the HV >50yrs group and HF w/o RD group for ADC and perfusion. The only significant difference between HF patients with and without renal dysfunction is for T<sub>1</sub>. Figure 2 shows significant correlations of ADC,  $f_pD*$ , flux and perfusion with eGFR. D is also highly correlated with eGFR (R=0.483, p=0.005, not shown).  $f_pD*$  is correlated with both renal artery flux and renal cortex perfusion (R=0.562, p=0.001 and R=0.407, p=0.021 respectively). T<sub>1</sub> correlated negatively with eGFR values (R=-0.436, p=0.01) and was prolonged in HF patients with renal dysfunction compared to those HF without renal dysfunction. There was no significant correlation of ADC or D with T<sub>1</sub>.

## **Discussion and Conclusion**

Results suggest that flow, as demonstrated by  $f_pD^*$ , contributes significantly to the changes in ADC, and that these changes are correlated to PC flux and ASL perfusion measures. However, although there was no difference between HF groups in  $f_pD^*$ , there is a trend for reduced fpD\*, PC Flux and perfusion with severity of disease. In addition structural changes are reflected by the increase in T<sub>1</sub>, though these are not reflected in D alone. Renal dysfunction in heart failure is mediated by decreased renal perfusion. Further, prolonged T<sub>1</sub> reflecting chronic structural renal changes/congestion might be the primary culprit in the pathophysiology of the chronic cardiorenal syndrome. These structural changes appear to be associated with classical cardiovascular risk factors.

References: [1] Ronco et al. Blood Purif 2009; 27:114-126, [2] Gardener et al. MRM 2010; 63:1627-36, [3] Buxton et al. MRM 1998; 40:383-96