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Supplementary file

Methods

CMR protocol

CMR studies were performed on a 3.0 T (Philips Achieva) equipped with a 32-channel receiver coil.

Patients were asked to avoid caffeine for 24 hours prior to having the CMR performed.

Cine images

Scout images were used to plan cine images in the vertical long axis, pseudo short axis and horizontal long axis (balanced steady state free precession [bSSFP] acquisition). Cine image stack were acquired covering the entire heart. Image acquisition parameters for bSSFP as follows: TR 2.6 ms, TE 1.3 ms, flip angle 40°, spatial resolution $2.0 \times 1.63 \times 8 \text{ mm}^3$, 30 cardiac phases (1, 2).

T1 maps acquisition

Native and post contrast T1 mapping were planned in a single short axis slice at mid LV level using an ECG-triggered modified Look-Locker inversion (MOLLI) using the '3 of 5' approach, with 3x R-R interval recovery epochs, voxel size $1.7 \times 2.14 \times 10 \text{ mm}^3$ Trigger delay at end-diastole, flip angle 35°, FOV 320 – 420 mm (3, 4); Post-contrast T1 mapping was acquired 15 minutes after contrast administration.

First pass myocardial perfusion

Stress first-pass myocardial perfusion was acquired after intravenous adenosine administration (140 mcg/kg/min for three minutes) under continuous ECG monitoring and assessment for adequate haemodynamic response using a Gadolinium-DTPA (diethylene triamine pentaacetic acid (0.075 mmol/kg) bolus) for both stress and rest first-pass myocardial perfusion. A spoiled turbo gradient echo with a $5 \times$ k-t Broad-use Linear Acquisition Speed-up Technique (BLAST), 11 training profiles, $1.31 \times 1.32 \times 10 \text{ mm}^3$ acquired resolution, pre-pulse delay of 100 ms, acquisition shot of 123 ms/slice, flip angle 20°, TR/TE 3.0/1.42 ms and three short axis slices was used (5).

Late gadolinium enhancement

Late gadolinium enhancement (LGE) images were acquired between 10 and 15 minutes following contrast administration using inversion recovery-prepared T1-weighted gradient echo. The optimal inversion time to null signal from normal myocardium was determined using a Look-Locker approach. 10 to 12 short axis slices were acquired, with further slices in the vertical and horizontal long axis orientations, or phase-swapped if indicated. Typical parameters for LGE images were as follows: TE 2.0 ms, TR 3.5 ms, flip angle 25°, acquired spatial resolution 1.54 × 1.76 × 10 mm.³. Alternate heart beat acquisitions by navigator was used for poor breath holders.

CMR analysis

Image analysis was performed using cvi42 (v 5.6.5, Circle Cardiovascular Imaging Inc., Calgary, Canada) in accordance to recognised reporting standards(6) and using established and validated protocols.(7) LGE and myocardial perfusion were assessed by 2 blinded expert readers.

For LV volume and function, LV contours were drawn manually at both end diastole and end systole on the LV short axis SSFP cine stack.(6) Papillary muscles were included in the LV blood pool.

Native and post-contrast myocardial T1 was measured by delineating a region of interest in the mid interventricular septum(8) and in the LV blood pool. Care was taken to avoid partial-volume effects from neighbouring tissue or blood pool when delineating the region of interest (ROI). The following formula was used to calculate extracellular volume (ECV), where R1 is 1/T1, *myo pre* and *myo post* are the pre and post-contrast myocardial T1 values and *blood pre* and *blood post* are the pre-contrast and post-contrast blood pool T1 values.(9)

$$ECV = (1 - hematocrit) \times \frac{R1_{myo\ post} - R1_{myo\ pre}}{R1_{blood\ post} - R1_{blood\ pre}}$$

LGE was reported according to the 16 segment American Heart Association (AHA) model (10). The five-standard deviation (5SD) method was used for LGE scar mass quantification (11). Two separated

ROIs, one representing the region with LGE and the other representing remote myocardium, were delineated on the LGE short axis images to obtain the scar mass.

Perfusion was firstly assessed visually by comparing the rest and stress perfusion images using the 16 segment AHA model.(10)

Quantitative analysis of the perfusion data was performed to generate estimates of myocardial blood flow (MBF) at stress and rest as previously described. (12) Myocardial perfusion reserve (MPR) values were calculated by dividing the stress by the rest MBF. Breathing motion in the dynamic series was corrected using an automated registration algorithm (Circle Cardiovascular Imaging, Calgary, AB, Canada). Regions of interest defining the myocardium and a region within the left ventricular blood pool were then used to generate signal versus time curves. Signal values were converted to contrast agent concentrations using the pre-contrast T_1 measurements and the equation for the imaging sequence derived from the Bloch equations.(13, 14) The calibration factor S_0 was derived from the pre-contrast image signal (SI_{pre}) and pre-contrast T_1 (T_{1_pre}) as follows:

$$S_0 = \frac{SI_{pre}}{f(T_{1_pre})}$$

Where $f(...)$ denotes the equation describing the imaging pulse sequence.(13) The T_1 at each time point $T_1(t)$ was then obtained from the signal intensity $SI(t)$:

$$T_1(t) = \min \left\{ \left(S_0 \cdot f(T_1(t)) - SI(t) \right)^2 \right\}$$

The contrast agent concentration, $C(t)$, was then obtained as follows:

$$C(t) = \frac{\frac{1}{T_1(t)} - \frac{1}{T_{1_pre}}}{r_1}$$

Where r_1 is the contrast agent relaxivity.

The concentration versus time curves for the blood pool and myocardium were then used to estimate myocardial blood flow using model independent deconvolution.(14, 15) The following equation was minimized to find the flow weighted response function $R_f(t)$:

$$\min \left\{ \|A \cdot R_f(t) - C_m(t)\|^2 - \lambda^2 \|L \cdot R_f(t)\|^2 \right\}$$

A is the convolution matrix operator calculated from the arterial concentration curve(16) (4), C_m is the myocardial concentration versus time curve, L is the identity matrix and λ , is a coefficient that determines the degree to which the solution is forced to be smooth by the side constraint $\|LR_f\|^2$. The optimal value for λ was determined using the L-curve method. (14, 17) The MBF was taken as the maximum value of the flow weighted response function $R_f(t)$.

Results

One hundred and twenty-three SSc patients were screened for the study, out of whom twenty-nine were excluded as they did not meet the inclusion criteria or refused the CMR. Ninety-three patients were recruited but only eighty-three had CMR data available. Reasons for absent data is illustrated in figure 1. Of the eighty-three patients with CMR data, function/volume assessment was available for all patients, LGE-CMR and T1 native data was available for 80 patients, ECV% for 78 patients, perfusion data was available for 61 patients.

Figure S1. Flowchart showing patient selection, recruitment and feasibility

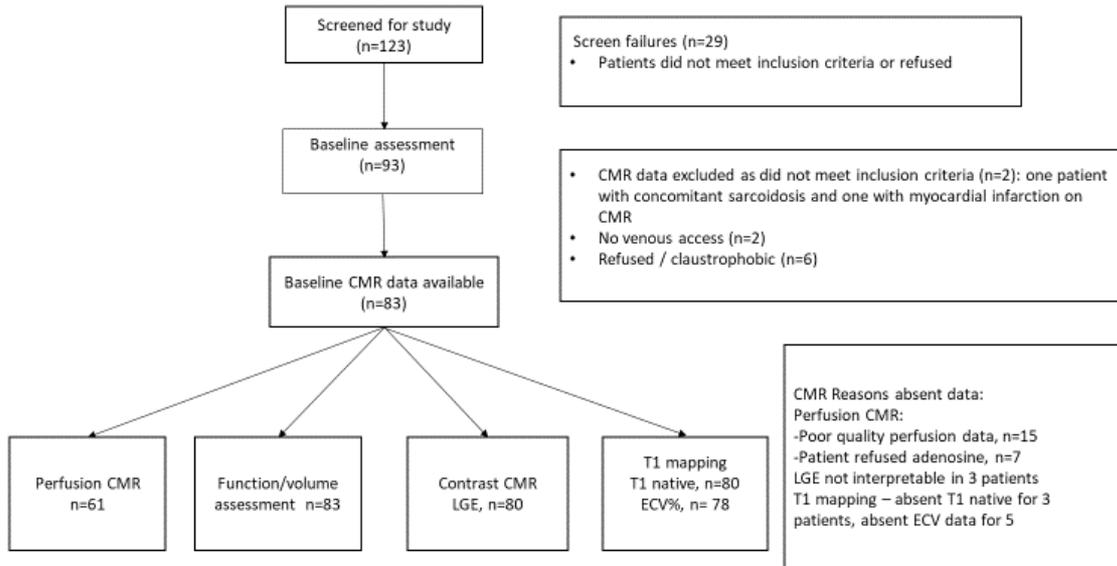


Table S1. Standard of care cardio-pulmonary profile

Standard of care cardio-pulmonary profile (% predicted value)	
Forced vital capacity %, mean (SD)	101 (21)
Total lung capacity %, mean (SD)	92 (15)
DLCO %, mean (SD)	62 (14)
DLCO/VA %, mean (SD)	80 (15)
Cardiovascular risk profile, n%	
Dyslipidaemia	3 (4)
Hypertension	8 (10)
Smoking	7 (8)
Family history of CVD	8 (10)
Patients with any CV risk factors, n%	22 (27)
1 cv risk factor	17 (21%)
2 cv risk factors	5 (6%)
Systolic BP, mean (SD)	116 (16)
Diastolic BP, mean (SD)	67 (12)
BMI, median (IQR)	26 (20,31)
Cardiac serum biomarkers	
NT-proBNP, median (IQR)	92 (48,143)
Hs-TnI, median (IQR)	4 (3,8)
CK, median (IQR)	73 (63,105)

BP, blood pressure; BMI, body mass index; CK, creatine kinase; CV, cardiovascular; CVD, cardiovascular disease; DLCO, diffusing capacity of the lungs for carbon monoxide; DLCO/VA, DLCO adjusted for volume; FVC, forced vital capacity; Hs-TnI, high-sensitivity troponin I; NT-proBNP, N-terminal pro brain natriuretic peptide; TLC, total lung capacity

Right ventricular (RV) dimension and function in SSc patients

RV parameters of SSc patients were within normal range, with a mean (SD) RVEF of 58 (7)%, RVED/body surface area (BSA) of 79 (18), RVESV/BSA of 35 (12) and RVSV/BSA of 46 (8) (ml/m²).

Association between native T1 and clinical SSc phenotype

T1 native in SSc patients versus HC did not reach statistical significance, after applying Bonferroni correction. The association between T1 native and clinical phenotype is detailed in Table S2. T1 native associated with MRSS and had a negative association with dcSSc.

Table S2. Logistic and linear regression to predict the relationship between SSc clinical phenotype and native T1

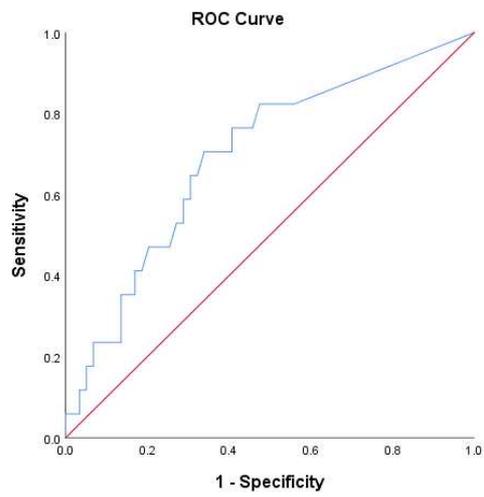
Variable	T1 native			
	Linear univariate analysis		Multivariate analysis, R ² =0.238	
	Beta	p value	Beta	p value
Male gender	-0.89	0.431	-0.194	0.129
Age	0.122	0.281		
Presence of CV risk factors	-0.081	0.475	-0.146	0.181
Disease duration	-0.076	0.508		
Presence of ILD	-0.151	0.181		
DLCO/VA	-0.054	0.641		
MRSS	0.249	0.026*	0.457	0.001*
DcSSc	-0.132	0.241	-0.312	0.015*
Digital ulcers	0.204	0.070	0.149	0.177
CRP	0.107	0.346	0.163	0.170
DMARD treatment	0.063	0.577		
ACE inhibitor treatment	-0.209	0.063		

* p<0.05; ** p<0.001

ACE, angiotensin converting enzyme; CV, cardiovascular; CRP, C-reactive protein; dcSSc, diffuse cutaneous systemic sclerosis; DLCO/VA, DLCO adjusted for volume; DMARD, disease modifying antirheumatic drugs; ECV, extracellular volume; ILD, interstitial lung disease; LGE, late gadolinium enhancement; MPR, myocardial perfusion reserve; MRSS, modified Rodnan skin score; OR odds ratio

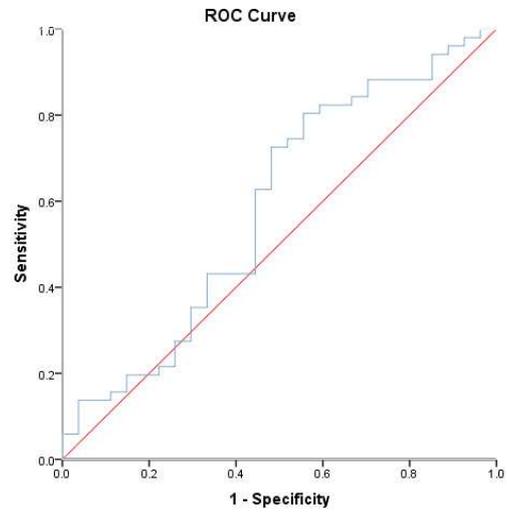
Figure S2A. ROC curve of hs-TnI for predicting LGE. S2B ROC curve of NT-proBNP for predicting ECV (ECV <29; ECV ≥29)

S2A



AUC (95%CI): 0.695(0.55, 0.837); p=0.015

S2B



AUC (95%CI) =0.586 (0.447; 0.726);p=0.213

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