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# MRI Susceptibility Mapping Suggests Elevated Brain Iron in Sickle Cell Anaemia

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### **Synopsis**

Sickle Cell Anaemia (SCA) is a genetic condition characterized by haemolytic anaemia, cerebral vasculopathy and cognitive impairment. The effect of SCA on brain iron concentrations has not been extensively studied. Brain iron is important in cognitive function and iron overload may accelerate neurodegeneration. Here, susceptibility mapping (QSM) was used to compare brain tissue susceptibility values in 86 SCA patients and 25 healthy controls. Elevated susceptibility was found in the red nucleus of the SCA group versus controls, suggesting increased iron accumulation. In SCA subjects there was no significant effect of silent cerebral infarcts or anaemia severity on brain susceptibility values.

#### Introduction

Sickle Cell Anaemia (SCA) is a genetic condition which affects the development of red blood cells (RBCs). The modified RBCs break down prematurely causing haemolytic anaemia. Cerebral vasculopathy is prevalent, exposing SCA subjects to a high risk of silent cerebral infarcts (SCI) and ischaemic stroke. SCA subjects, including those exhibiting no obvious neurological damage, suffer from cognitive deficits that increase with age<sup>1</sup>.

The accumulation of excess brain iron may contribute to these cognitive deficits observed in SCA. Brain iron is important in cognitive function<sup>2</sup>, and iron accumulation accelerates neurodegeneration in conditions such as Parkinson's<sup>3</sup>. Cerebral vasculopathy is known to trigger iron accumulation<sup>4</sup> and the effect of haemolytic anaemia on brain iron is not well characterized<sup>5</sup>.

Here, we used Susceptibility Mapping (QSM) to compare regional brain iron concentrations between SCA and healthy control (HC) groups. Magnetic susceptibility ( $\chi$ ) values were measured in iron-rich grey matter regions of interest (ROI), where  $\chi$  is directly proportional to iron stored in the form of ferritin macromolecules<sup>6,7</sup>. Within the SCA group, the effects of anaemia severity and vasculopathy on  $\chi$  values were investigated by comparing regional  $\chi$  with blood haematocrit (Hct) and SCI presence respectively.

#### Methods

86 SCA subjects (mean ± standard deviation age: 19.03±10.84 years) and 25 HCs (16.19±5.02 years) were recruited from the Sleep and Asthma Cohort(SAC), and Prevention of Morbidity in Sickle Cell Disease(POMS) clinical studies<sup>8,9</sup>, whose exclusion criteria included chronic blood transfusions.

MRI data were acquired on a 3T Siemens Magnetom Prisma. Susceptibility maps were calculated from multi-echo gradient-recalled-echo (GRE) images. Sequence parameters included: 7 echoes,  $TE_1/\Delta TE/TR:3ms/4ms/38ms$ , 1.15mm isotropic resolution, FOV:180x220x166mm³. A trained radiologist identified SCIs on fluid-attenuated inversion recovery images. Hct was measured in blood sampled during the clinical studies.

QSM Pipeline:  $B_0$  field maps were obtained from a nonlinear fit of the complex GRE images<sup>10</sup> and underwent Laplacian-based unwrapping<sup>11</sup>. Background field removal was performed using Projection onto Dipole Fields<sup>12</sup>. Brain masks were calculated using the FSL Brain Extraction Tool<sup>13</sup>. Field-to- $\chi$  inversion was performed using Tikhonov regularization<sup>14</sup> with regularization parameter  $\alpha$ =0.06, selected using L-Curve methods. Susceptibility values were examined in four iron-rich ROI: Red Nucleus (RN), Caudate Nucleus (CN), Putamen (PT) and Globus Pallidus (GP) (Figure 1). ROI were segmented by coregistration of the Eve QSM atlas<sup>15</sup> to the sixth-echo GRE magnitude image using NiftyReg<sup>16</sup>.

Regression of regional mean  $\chi$  values on log-transformed age was performed on pooled SCA and HC subjects to remove any effect of iron accumulation with age<sup>17</sup>. ANCOVA was applied to examine differences in regional age-corrected  $\chi$  between SCA and HCs, and between SCA subjects with (SCI+) and without SCI (SCI-). In SCA subjects, correlations between the age-corrected ROI mean  $\chi$  values and Hct levels were calculated to assess the effect of anaemia severity on  $\chi$ .

### Results

A significant increase in  $\chi$  with age was observed in all ROI (Figure 2). ANCOVA showed a faster  $\chi$  increase with age in the RN in SCA than in HCs (Figure 3a). The mean age-corrected  $\chi$  in the RN was significantly greater in the SCA group than in the HCs, but there were no significant group differences in any of the other ROI (Figure 3b). There were no significant  $\chi$  differences between the SCI+ and SCI- groups in any ROI (Figure 4). There was no correlation between the mean  $\chi$  values and Hct levels in any of the ROIs (Figure 5).

### **Discussion and Conclusions**

The greater mean age-corrected susceptibility and faster susceptibility increase with age suggests increased iron accumulation in the red nucleus of subjects with sickle cell anaemia compared to healthy controls. The underlying mechanism remains unclear as the increased  $\chi$  was not correlated with Hct (a putative measure of anaemia severity) or SCI presence (a putative measure of vasculopathy) in the SCA group. A more direct measure of vasculopathy than SCI (e.g. transcranial Doppler or MR angiography) together with measures of tissue compromise (e.g. diffusion tensor imaging) and hypoxic exposure (e.g. oxygen saturation) could be acquired in future to investigate any relationship with iron accumulation. Furthermore, it would be interesting to use QSM to examine iron accumulation outside these basal ganglia ROIs, closer to cortical areas affected by SCI. The increased iron accumulation observed in SCA may affect cognitive function. We are currently awaiting cognitive results from the clinical studies, and future work will include comparing  $\chi$  values and cognitive performance.

### **Acknowledgements**

This study was supported by the EPSRC-funded UCL Centre for Doctoral Training in Medical Imaging (EP/L016478/1) and the Department of Health's NIHR-funded Biomedical Research Centre at Great Ormond Street Hospitals.

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## **Figures**

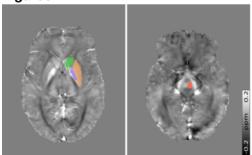


Figure 1: Unilateral segmented ROIs overlaid on the susceptibility maps of a representative healthy control subject. Caudate Nucleus (Green), Putamen (Orange) Globus Pallidus (Purple) and Red Nucleus (Red).

Correlation between	Mean Regional χ	Values and Log-Transformed Age
for the Combined SC	A and HC groups	
Region of Interest	Cor.Coef	p-value
Caudate Nucleus	0.48	9.5e-08***
Putamen	0.78	<2.2e-16***
Globus Pallidus	0.79	<2.2e-16***
Red Nucleus	0.81	<2.2e-16***

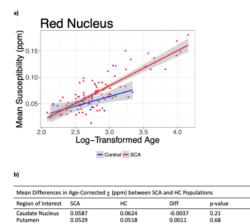
Figure 2: Table showing the results of correlation analysis on mean regional susceptibility values against the log-transformed age at the time of MRI. Highly significant correlations (p<0.001) were observed in all of the regions of interest considered. \*\*\* Indicates p-value significance at the 0.001 level.

Globus Pallidus

Red Nucleus

0.1342

0.0714



0.1340

0.0002

0.0124

0.96

Figure 3: Regression plot a) showing the mean susceptibility values in the red nucleus as a function of the log transformed age. A greater rate of susceptibility increase with age can be observed in the subjects with sickle cell anaemia (red) compared to the healthy controls (blue). The grey shaded areas indicate the uncertainty in the fit of the linear regression models. Table b) showing differences in the regional mean age-corrected susceptibility values between the sickle cell anaemia and healthy control groups. \*Indicates p-values significance at the 0.05 level.

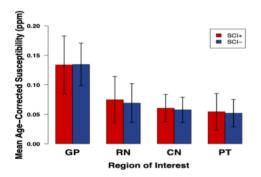
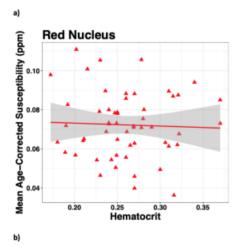


Figure 4: Bar charts illustrating an absence of significant differences in susceptibility between SCA subjects with (SCI+) and without (SCI-) silent cerebral infarcts. Mean age-corrected susceptibility values are plotted in each of the four regions. Standard deviations of the means are shown as error bars. There were no significant differences in any of the ROIs considered.



Regional Mean Age-Corrected $\chi$ Values and Hct Correlation in SCA			
Region of Interest	Cor.Coef	p-value	
Caudate Nucleus	-0.14	0.31	
Putamen	-0.07	0.61	
Globus Pallidus	-0.099	0.47	
Red Nucleus	-0.04	0.77	

Figure 5: Mean age-corrected susceptibility values in the red nucleus plotted as a function of haematocrit level in the SCA subjects a). The grey shaded area indicates the uncertainty in the fit of the linear regression model. Table b) showing the results of correlation analysis on regional mean age-corrected susceptibility values against blood haematocrit. There was no correlation observed between the susceptibility values and Hct in the red nucleus, and all other regions considered (not plotted).

Proc. Intl. Soc. Mag. Reson. Med. 27 (2019)