Cerebral Blood Flow and Marrow Diffusion Alterations in Children with Sickle Cell Anemia after Bone Marrow Transplantation and Transfusion

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ABSTRACT

BACKGROUND AND PURPOSE: Hematopoietic marrow hyperplasia and hyperperfusion are compensatory mechanisms in sickle cell anemia. We have observed marrow diffusion and arterial spin-labeling perfusion changes in sickle cell anemia following bone marrow transplantation. We aimed to compare arterial spin-labeling perfusion and marrow diffusion/ADC values in patients with sickle cell anemia before and after bone marrow transplantation or transfusion.

MATERIALS AND METHODS: We reviewed brain MRIs from patients with sickle cell anemia obtained during 6 consecutive years at a children's hospital. Quantitative marrow diffusion values were procured from the occipital and sphenoid bones. Pseudocontinuous arterial spin-labeling perfusion values (milliliters/100 g of tissue/min) of MCA, anterior cerebral artery, and posterior cerebral artery territories were determined. Territorial CBF, whole-brain average CBF, and marrow ADC values were compared for changes before and after either bone marrow transplantation or transfusion. Bone marrow transplantation and transfusion groups were compared. Two-tailed paired and unpaired Student t tests were used; P < .05 was considered significant.

RESULTS: Fifty-three examinations from 17 patients with bone marrow transplantation and 29 examinations from 9 patients with transfusion were included. ADC values significantly increased in the sphenoid and occipital marrow following bone marrow transplantation in contrast to patients with transfusion (P > .83). Whole-brain mean CBF significantly decreased following bone marrow transplantation (77.39 \pm 13.78 to 60.39 \pm 13.62 ml/100 g tissue/min; P < .001), without significant change thereafter. CBF did not significantly change following the first (81.11 \pm 12.23 to 80.25 \pm 8.27 ml/100 g tissue/min; P = .47) or subsequent transfusions. There was no significant difference in mean CBF between groups before intervention (P = .22).

CONCLUSIONS: Improved CBF and marrow diffusion eventuate following bone marrow transplantation in children with sickle cell anemia in contrast to transfusion therapy.

ABBREVIATIONS: ACA = anterior cerebral artery; ASL= arterial spin-labeling; BMT = bone marrow transplantation; PCA = posterior cerebral artery; SCA = sickle cell anemia; TI = relaxation time

Sickle cell anemia (SCA) is an inherited hematopoietic disorder, rare in the general US population but more common in African Americans (0.3%; 1/365).¹ The mutated hemoglobin contained in the red blood cells of patients with SCA (HbSS) forms insoluble polymers under low oxygen conditions and deforms the cells, giving them their sickle shape. The "sickle cells" have decreased deformability, rendering them unable to easily

pass through small vessels, and can reduce or occlude blood flow to tissues. Enhanced red blood cell production and blood flow are compensatory mechanisms used by the body in an attempt to maintain homeostasis. Hematopoietic marrow hyperplasia causes marrow signal alterations on MR imaging, including T1 lengthening, T2 shortening, and reduced diffusion.²⁻⁴ Elevated cardiac output, increased arterial blood flow, and vasodilation contribute to cerebral hyperperfusion.⁵ However, end organs remain vulnerable to ischemic injury because these compensatory adjustments are often insufficient. Furthermore, small- and large-vessel vasculopathy can potentiate this rather tenuous situation.⁶ In the brain, injury may take the form of borderzone ischemia, focal small-vessel lacunar infarcts (symptomatic or silent), or large vascular territory infarctions.

Arterial spin-labeling (ASL) perfusion is a noninvasive MR imaging technique that uses flowing blood as an intrinsic tracer to

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assess cerebral blood flow in milliliters/100 g of tissue/min.7-9 Thus, it is an MR imaging technique well-suited to the SCA population because it readily depicts the present state of blood flow to the brain without the need for intravenous gadolinium. In compensated patients with SCA, the combination of hyperperfusion and anemia leads to a generalized increase in ASL brain signal. The signal is then reduced in patients with cerebrovascular disease, including delayed arterial transit (eg, vasculopathy) and hypoperfusion (eg, ischemia or encephalomalacia), or increased in states of hyperperfusion (eg, luxury perfusion or seizure-related).8,10,11 Notable limitations of the ASL technique include a relatively poor signal-to-noise ratio and variables often represented as estimations in calculating the ASL perfusion equation, such as tagging efficiency, postlabel delay, the cortex-blood partition coefficient, and the T1 of blood.8 The latter, however, can be measured directly or calculated with knowledge of the magnetic field strength, hemoglobin, and oxygen saturation.¹²

Primary therapeutic measures in symptomatic patients with SCA include hydroxyurea, red blood cell transfusions, and, in the appropriate population, bone marrow transplantation (BMT), which is curative if successful. Human leukocyte antigenmatched sibling donors offer the best hope for a cure with BMT, with a success rate as high as 98%. ¹³ However, failure can occur in up to 10% and is more common with nonsibling human leukocyte antigen—matched donors and nonmatched donors. ^{14–20} Repeat transplantation would be necessary under these circumstances to achieve success. Imaging markers for transplantation failure would thus be useful adjuncts to guide early treatment decisions.

We have observed changes in brain blood flow and marrow diffusion values following bone marrow transplant in SCA, potentially representing imaging biomarkers of a favorable therapeutic response. The aim of this study was to compare and contrast alterations in ASL perfusion and bone marrow diffusion/ADC values in patients with SCA before and after BMT or initiation of chronic, monthly transfusion therapy.

MATERIALS AND METHODS

This Health Insurance Portability and Accountability Act-compliant retrospective study was performed after institutional review board approval. The brain MR imaging data base from a single academic children's hospital was queried for all examinations performed in patients with sickle cell anemia during a consecutive 6-year period (January 1, 2012, to December 31, 2017) using the terms "sickle cell," "ASL," and "arterial spin-labeling" to build a cohort for retrospective analysis. Each patient's medical record was reviewed to confirm a diagnosis of sickle cell disease, record demographic information, and determine the clinical course of the disease, including time points for transfusions and/or bone marrow transplantation. Each patient's most recent hematocrit and oxygen saturation levels relative to the time of the brain MR imaging were documented to correct for the longitudinal relaxation (T1) value of arterial blood. MR imaging examinations with excessive motion artifacts and technical limitations and without ASL images were excluded. Excessive motion artifacts were determined at the discretion of the board-certified neuroradiologist who reviewed each of the scans (M.T.W.) on the basis of training and experience. Motion artifacts were considered excessive when they limited either the ASL and/or DWI datasets more than a minimal-to-mild degree. Additional exclusion criteria included acute chest syndrome, acute stroke, posterior reversible encephalopathy syndrome, active infection, recent seizure, and vasculopathy based on MRA or transcranial Doppler (when MRA was unavailable). Cerebral hemispheres containing any chronic lesions larger than a lacunar infarct were excluded from analysis; in these cases, the contralateral hemisphere was evaluated in isolation. ASL perfusion images were evaluated qualitatively and quantitatively by a board-certified neuroradiologist with >6 years of clinical experience after board certification who was blinded to the type of therapy/intervention at the time of assessment.

Studies were performed on either a 1.5T (Discovery MR450 or Optima MR450W) or 3T (Discovery MR750) MR imaging scanner (GE Healthcare, Milwaukee, Wisconsin). The following sequences were performed through the brain in each examination: arterial spin-labeling perfusion, sagittal spoiled gradient-echo T1WI, axial T2WI, axial T2 FLAIR, axial DWI, and coronal fatsaturated T2WI. 3D TOF-MRA of the circle of Willis was also acquired concurrently with each brain MR imaging examination.

DWI: Technique, Processing, and Assessment

The bone marrow was evaluated qualitatively on T1, T2, and diffusion-weighted sequences. The T1WI and T2WI were assessed first to ensure lack of substantial marrow hypointensity reflecting transfusion-related iron deposition, which would interfere with the assessment of the DWI due to susceptibility. We excluded from further evaluation those with substantial marrow hypointensity defined as hypointense marrow signal with respect to muscle on all pulse sequences.

DWI acquisition parameters were as follows: TR, 10,000 ms; TE, 83–94 ms; flip angle, 90°; matrix, 128 \times 128 mm; FOV, 24 \times 27 cm; 1–2 excitations; section thickness, 5 mm; spacing, 0. Qualitative marrow diffusion abnormalities were documented using a 3-point scale: 1, within normal limits; 2, mild-to-moderate restricted diffusion; and 3, marked restricted diffusion. Quantitative apparent diffusion coefficient values (in square millimeters/second) were measured using 4-mm circular ROIs placed manually in the marrow cavity of the right greater sphenoid wing and midline occipital bones; these representative sites were selected because the marrow cavity here is reliably thicker than in many other locations of the skull.

ASL: Technique, Processing, and Assessment

Each subject underwent pseudocontinuous ASL with background suppression and a segmented 3D fast spin-echo readout in accordance with current consensus guidelines⁷ using the following scan parameters: flip angle, 111°–155°; inversion time/postlabel delay, 1025–1525 ms; labeling duration, 1500–2000 ms; TR, 4300–4600 ms; TE, 11 ms; ETL (echo train length), 1; NEX, 2–3; matrix, 512 × 8; FOV, 24–55 × 24–29 cm; 32–35 slices; 3- to 4-mm slice thickness; 3- to 4-mm spacing; total sequence acquisition time, 4–5 minutes. Anatomic landmarks were used to select the labeling plane just below the inferior margin of the cerebellum. ASL post-processing was performed on the MR imaging scanner. Gray-scale ASL perfusion images and pseudocolor images were generated and displayed with age-dependent scaling based on the

Table 1: Transplant group—hemoglobinopathy subtype, transplant data, and MR imaging data analyzed

								ASL (Right	ASL (Left
Pt	Diagnosis	Source	Donor	HLA Match	Scans (No.)	Age (yr)	DWI (No.)	Cerebrum) (No.)	Cerebrum) (No.)
1	Hbss	BM	R	8/8	2	18,19	2	2	2
2	Hbss	BM	R	8/8	4	15,16,16,18	4	4	4
3	HbS-B0Th	PBSC	R, H	4/8	2	12,13	2	2	2
4	HbSC	BM	R	8/8	5	13,14,14,16,17	5	5	5
5	Hbss	BM	R	8/8	4	7,10,11	3	3	3
6	Hbss	BM	R	8/8	2	7,9	2	2	2
7	HbS-B0Th	BM	R	8/8	2	10,12	2	2	2
8	Hbss	BM	R	8/8	3	11,11,12	3	2	2
9	Hbss	BM	R	8/8	4	5,7,8,9	4	4	4
10	Hbss	UCB	R	5/6	4	8,8,9,10	4	0	4
11	Hbss	BM	R	8/8	3	8,9,10	3	3	3
12	Hbss	PBSC	R, H	3/6	3	4,5,5	3	0	3
13	Hbss	BM	U	8/8	5	18,18,18,19,20	5	3	3
14	Hbss	BM	R	8/8	2	1,2	2	2	2
15	Hbss	BM	R	10/10	3	3,3,3	2	3	3
16	HbSS	BM	R	8/8	2	15,16	2	2	2
17	Hbss	UCB	U	6/6	3	3,3,4	3	3	3

Note:—Pt indicates patient; BM, bone marrow; PBSC, peripheral blood stem cell; UCB, umbilical cord blood; R, related; H, haploidentical; U, unrelated; HbSS, homozygous hemoglobin S; HbSC, sickle cell hemoglobin C; HbS-B0Th, hemoglobin S-β-thalassemia disease; HLA, human leukocyte antigen.

Table 2: Chronic transfusion group—hemoglobinopathy subtype and MR imaging data analyzed^a

	,					
Pt	Diagnosis	Scans (No.)	Age (yr)	DWI (No.)	ASL (Right Cerebrum) (No.)	ASL (Left Cerebrum) (No.)
1	Hbss	3	7,9,11	0	3	0
2	Hbss	6	0,1,2,5,6	3	1	5
3	Hbss	2	7,8	0	2	2
4	Hbss	2	8,11	2	0	2
5	Hbss	6	2,3,4,5,6,7	0	6	6
6	Hbss	2	4,6	1	2	2
7	Hbss	2	6,8	1	2	2
8	Hbss	3	3,4,5	1	3	3
9	Hbss	3	9,10,11	0	3	3

Note:—Pt indicates patient; HbSS, homozygous hemoglobin S.

magnitude of normal cerebral blood flow from the neonatal-to-early childhood period (younger than 1 year: 0–60 mL/100 g of tissue/min; older than 1 year: 0–100 mL/100 g of tissue/min).²¹

Arterial territory–specific quantitative ASL data in milliliters/100 g of tissue/min were procured using an AW workstation with Func-Tool software (GE Healthcare). The T1 value of arterial blood was calculated in each patient using patient-specific hematocrit levels, oxygen saturation at or around the time of imaging (median, 2 days; range, 0–71 days), and the magnetic field strength according to previously published guidelines. Then, 6 distinct manual ROIs were drawn around the cerebral cortex and subcortical white matter at the level of the basal ganglia to approximate each of the following vascular territories: anterior cerebral artery (right and left), middle cerebral artery (right and left), and posterior cerebral artery (right and left), using a previously established technique. Cerebral blood flow was calculated from the following equation:

$$CBF = \frac{\lambda (1 - e^{\frac{-t_{\text{sat}}}{T_{\text{lg}}}})}{2\alpha T_{\text{lb}} (1 - e^{\frac{-\tau}{T_{\text{lb}}}})} \frac{PW}{PD} e^{\frac{-PLD}{T_{\text{lb}}}}$$

where λ indicates the cortex-blood partition coefficient (0.9, assumed); t_{sat} , time of saturation before imaging (2000 ms); T_{1g} , T1

of gray matter (1200 ms, assumed); α , labeling efficiency (0.8, assumed); $T_{1\text{b}}$, T1 of the blood (calculated for each patient using the hematocrit, oxygen saturation, and magnetic field strength based on Hales et al¹²); τ , labeling duration (1500 ms); PW, perfusion weighted signal intensity; PD, proton density signal intensity; and PLD, postlabeling delay time. Mean whole-brain perfusion was estimated in each examination by averaging CBF values from all 6 evaluated territories. Gray-scale and corresponding postprocessed pseudocolor data were also evaluated together qual-

itatively. Qualitative cerebral signal magnitude was recorded using a 3-point scale: 1, hypoperfusion; 2, within normal limits; and 3, hyperperfusion.

Statistical Methods

The normal Gaussian distribution of the data mean was determined and confirmed by a Shapiro-Wilk test. Subsequently, 2-tailed paired t tests were used separately in each group of patients with SCA before and after BMT and transfusion, respectively. Two-tailed unpaired t tests were then performed to evaluate differences in the mean cerebral blood flow and marrow diffusion values between the posttransplantation and posttransfusion arms. The Wilcoxon signed rank test was used to evaluate paired samples of qualitative ordinal data. Observed power was calculated on the basis of the postintervention whole-brain CBF mean and SD differences between groups. $P \le .05$ was considered significant.

RESULTS

Fifty-three brain MR imaging examinations from 17 patients with BMT (mean, 3; range, 2–5; mean length of follow-up, 1.9 ± 0.9 years; range, 0.7–4.2 years) (9 male, 8 female) and 29 brain MR

a SCA subtype, transplant information, total MRI scans, number of DWIs, and number of ASL sequences analyzed (right and left cerebral hemispheres) from 17 patients with transplants.

^a SCA subtype, total MRI scans, number of DWIs, and number of ASL sequences analyzed (right and left cerebral hemispheres) from 9 patients with transfusion.

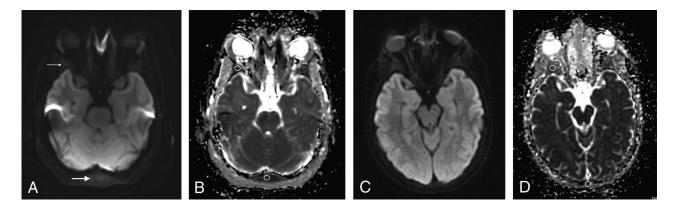


FIG 1. A 13-year-old girl with sickle cell disease. Pre-bone marrow transplantation axial DWI (A) (TR/TE = 10,000/83 ms, slice thickness = 5 mm) through the greater sphenoid wing (thin arrow) and occipital bone (thick arrow) and corresponding quantitative ADC map with ROIs (B) placed demonstrating reduced diffusion of the bone marrow. Follow-up MR imaging at 14 years of age from the same patient post-bone marrow transplantation with axial DWI (C) (TR/TE = 10,000/79.8 ms, slice thickness = 5 mm) at the same level as A and corresponding quantitative ADC map (D) demonstrating facilitated diffusion with respect to the pretransplantation scan.

Table 3: Pre- and posttransplantation ASL perfusion changes^a

	Pretransplantation								Posttransplantation						
Pt	MCA (R)	MCA (L)	ACA (R)	ACA (L)	PCA (R)	PCA (L)	Avg	MCA (R)	MCA (L)	ACA (R)	ACA (L)	PCA (R)	PCA (L)	Avg	P Value
1	71	70	68	73	55	60	66	78	79	69	75	58	68	71	.02 ^b
2	92	82	88	85	96	89	89	74	78	70	73	61	65	70	.008 ^c
3	82	80	79	83	92	90	84	60	63	62	64	64	64	63	<.001 ^c
4	66	63	64	60	50	51	59	34	35	37	38	28	30	34	$< .001^{c}$
5	100	80	93	85	97	83	90	62	61	62	61	63	54	61	$< .001^{c}$
6	90	89	96	93	87	91	91	81	77	74	70	71	73	74	<.001 ^c
7	68	65	68	63	63	59	64	69	62	37	59	51	57	56	.14
8	76	72	73	67	73	73	72	59	54	56	52	48	46	53	<.001°
9	84	65	77	64	86	77	76	97	87	86	84	79	71	84	.16
10	NA	57	NA	56	NA	55	56	NA	45	NA	48	NA	40	44	.03°
11	73	81	76	77	72	74	76	78	73	75	74	71	74	74	.47
12	89	NA	77	NA	100	NA	95	82	NA	72	NA	87	NA	80	.07
13	63	66	59	68	66	37	60	61	60	67	55	58	67	61	.82
14	85	81	79	76	77	74	79	77	79	65	70	62	63	69	.006 ^c
15	68	71	67	72	63	64	68	64	65	63	62	53	50	60	.004 ^c
16	76	73	76	71	75	71	74	66	66	57	58	45	47	57	.005°
17	100	99	101	94	91	94	97	40	40	37	37	35	37	38	$< .001^{c}$
Т	81	76	79	75	78	72	77	66	63	60	60	57	55	60	<.001 ^c

Note:—Pt indicates patient; R, right; L, left; Avg, average/mean whole-brain CBF; NA, not applicable/excluded; T, mean of each arterial territory.

imaging examinations from 9 patient with transfusion (mean, 3; range, 2–6; mean length of follow-up, 3 ± 1.4 years; range, 1.1– 5.6 years) (3 male, 6 female) met the inclusion criteria (Tables 1 and 2). Two patients with transfusion were excluded; one lacked ASL, and the other had a severe vasculopathy with multiple infarctions and excessive motion artifacts. Three follow-up examinations after BMT were excluded from 2 different patients due to posterior reversible encephalopathy syndrome. One hemisphere was excluded from ASL analysis due to infarctions or vasculopathy in 4 cases from 2 patients in the transplant group and in 5 cases from 3 patients in the transfusion group; however, there was no significant difference between pre- and posttransfusion and preand posttransplantation CBF values between the subgroups that underwent unilateral and bilateral cerebral analysis. At the time of the initial MR imaging, the mean age of the BMT group was 9.8 \pm 5 years (range, 1–18 years), while the mean age of the transfusion group was 6.3 ± 4 years (range, 0–9 years) (P = .13). Transfusions were initiated 20 days before the follow-up MR imaging on average (range, 4-60 days). The mean time between BMT and the first follow-up MR imaging was 281 ± 170 days (range, 3-875 days). No patients who underwent BMT had graft failure.

Marrow ADC Values

ADC values significantly increased in the marrow of the greater sphenoid wing $(0.97 \pm 0.34 \text{ to } 1.59 \pm 0.55 \times 10^{-3} \text{ mm}^2/\text{s}$; P = .025) and occipital bone $(1.11 \pm 0.27 \text{ to } 1.44 \pm 0.51 \times 10^{-3} \text{ mm}^2/\text{s}$; P = .043) following bone marrow transplantation (Fig 1). Qualitatively, ADC sequences also demonstrated visually brighter marrow following BMT (the mean score decreased from 3 to 1), representing more facilitated diffusion (P = .01428) (Fig 1). These changes were sustained across time on follow-up examinations, with no significant difference in the interval mean ADC values (P > .05). There was no significant difference in sphenoid wing and occipital marrow mean ADC values be-

^a Arterial territory—specific and mean whole-brain CBF values (in milliliters/100 g of tissue/min) from 17 patients with BMT prior to (left columns) and at the first scan after (right columns) transplantation. In the interest of space and simplicity, CBF values were rounded to the nearest whole number for this display.

^b Significant increase in mean whole-brain CBF.

^c Significant.

tween patients pre- and posttransfusion (P > .83); however, only 2 patients could be evaluated posttransfusion due to MR imaging evidence of marrow iron deposition. Post-BMT and posttransfusion

marrow ADC values were significantly different from one another (P = .0323). There was no significant difference between marrow ADC values obtained at 1.5T and 3T in either group (P > .05).

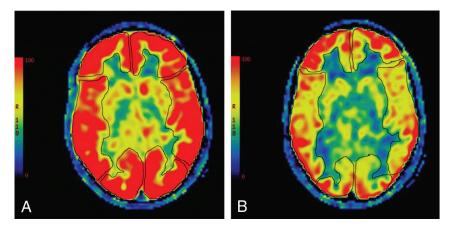


FIG 2. An 11-year-old boy with sickle cell disease. Pre-bone marrow transplantation, axial pseudocolor ASL (A) depicts manual ROIs drawn to interrogate CBF from the bilateral ACA, MCA, and PCA territories. Diffusely increased signal is present throughout the cerebrum, consistent with elevated cerebral blood flow. Follow-up MR imaging at 12 years of age from the same patient, post-bone marrow transplantation, with axial pseudocolor ASL (B) depicting persistent-but-improved/decreased hyperperfusion.

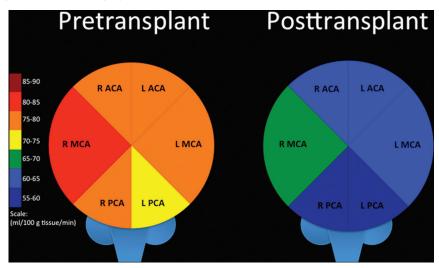


FIG 3. Comparison of arterial territory–specific quantitative mean CBF in patients with SCA before BMT (*left*) and at the first scan after BMT (*right*). R indicates right; L, left.

Cerebral Blood Flow

Preintervention mean whole-brain perfusion for both groups combined was 78.43 ± 11.18 mL/100 g of tissue/min. CBF values obtained at 1.5T were not significantly different from those obtained at 3T in either group (P > .05). There was no significant difference in mean quantitative cerebral blood flow between the BMT and transfusion groups before intervention (P = .22). Mean whole-brain cerebral blood flow was significantly lower after BMT compared with transfusion (P = .0004). The observed power was strong at 99.6%.

Whole-brain mean quantitative cerebral blood flow significantly decreased following BMT, from 77.39 \pm 13.78 to $60.39 \pm 13.62 \text{ mL/}100 \text{ g tissue/min (first)}$ scan; P = .00004), and did not significantly change thereafter (Table 3 and Figs 2 and 3). The mean CBF significantly decreased in all bilateral vascular territories (anterior cerebral artery [ACA], MCA, posterior cerebral artery [PCA]) on the first scan following BMT, and most of these significant differences were sustained across time. On an individual level, most patients (11/17) had a significant drop in whole-brain CBF on the first scan following BMT. Half of the patients who did not experience significantly decreased CBF on the first scan were reimaged at ≥ 1 later time point; CBF ultimately decreased significantly in two-thirds of these patients. The first posttransplantation scan was performed at 0.8 ± 6 years on average

Table 4: Pre- and postinitiation of chronic transfusion therapy ASL perfusion changes^a

		-	Pretra	ansfusion	ı			Posttransfusion							
Pt	MCA (R)	MCA (L)	ACA (R)	ACA (L)	PCA (R)	PCA (L)	Avg	MCA (R)	MCA (L)	ACA (R)	ACA (L)	PCA (R)	PCA (L)	Avg	P Value
1	96	NA	78	NA	93	NA	89	95	NA	94	NA	87	NA	92	.69
2	79	71	64	65	62	73	69	NA	87	NA	84	NA	80	84	.06
3	103	91	102	98	111	107	102	75	78	76	76	73	75	76	$<$.001 b
4	NA	89	NA	80	NA	80	83	NA	75	NA	82	NA	76	78	.37
5	90	96	81	93	82	88	88	93	96	95	101	91	93	95	.02 ^c
6	84	102	88	100	78	83	90	79	96	81	86	78	76	83	.02 ^b
7	98	93	88	90	92	88	92	81	84	74	85	74	83	81	.005 ^b
8	70	68	70	65	65	64	67	74	79	64	71	60	60	68	.74
9	70	70	64	66	63	71	67	69	74	77	85	73	80	87	.02 ^c
Т	86	78	79	82	81	82	81	81	84	80	80	77	78	80	.47

Note:—Pt indicates patient; R, right; L, left; Avg, average/mean whole-brain CBF; NA, not applicable/excluded; T, mean of each arterial territory.

^a Arterial territory—specific and mean whole-brain CBF values (in milliliters/100 g of tissue/min) from 9 patients with transfusion prior to (left columns) and at the first scan after (right columns) transfusion initiation. In the interest in space and simplicity, CBF values were rounded to the nearest whole number for this display.
^b Significant.

^c Significant increase in mean whole-brain CBF.

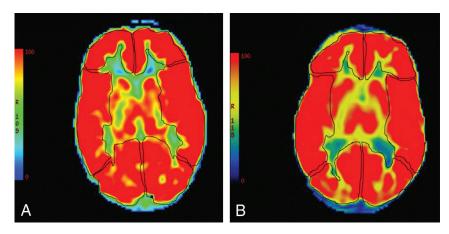


FIG 4. A 31-month-old girl with sickle cell disease. Pretransfusion axial pseudocolor ASL (A) depicts manual ROIs drawn to interrogate CBF from the bilateral ACA, MCA, and PCA territories. Diffusely increased signal is present throughout the cerebrum, consistent with elevated cerebral blood flow. Follow-up MR imaging at 36 months of age from the same patient at the same level posttransfusion with axial pseudocolor ASL (B) depicting persistent hyperperfusion without significant change.

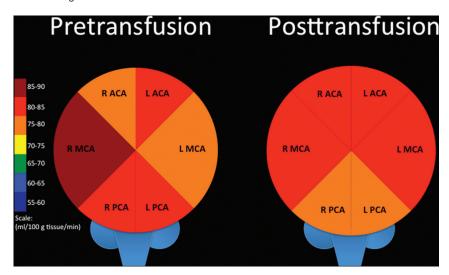


FIG 5. Comparison of arterial territory–specific quantitative mean CBF in patients with SCA before initial transfusion (*left*) and at the first scan after transfusion initiation (*right*). R indicates right; L, left.

after BMT. There was no significant difference in the time interval between transplant and the first posttransplantation brain MR imaging between patients who did and did not experience decreased CBF (P=.5). One of the 6 patients without significant CBF alteration had an unrelated BMT without myeloablative preconditioning.

Whole-brain mean perfusion values did not significantly change following the first transfusion (81.11 \pm 12.23 to 80.25 \pm 8.27 mL/100 g tissue/min; P=.47) or across time after subsequent transfusions (Table 4 and Figs 4 and 5). Whole-brain CBF did not significantly decline in 6 of the 9 patients and actually significantly increased in 2 of these patients. A minority of patients (3/9) did experience significantly decreased whole-brain CBF after transfusion; unfortunately, none of these patients had further follow-up imaging to determine the duration of the change.

DISCUSSION

Marrow diffusion and cerebral perfusion significantly improved in patients with sickle cell anemia following bone marrow transplantation and did not change significantly after transfusion alone. Because marrow ADC values negatively correlate and CBF positively correlates with the severity of anemia, these may be useful surrogate imaging biomarkers for therapeutic success.²⁻⁴,12

Apparent diffusion coefficient values of bone marrow are inversely proportional to the degree of hematopoietic constituents. Red marrow hyperplasia causes T1 prolongation, T2 shortening, and reduced diffusion.²⁻⁴ Proliferative hematopoietic marrow changes in SCA are an attempt to compensate for the diminished oxygen-carrying capacity of the sickled erythrocytes and to replenish red blood cells prematurely removed from circulation. All patients in our cohort had evidence of reduced diffusion in the sphenoid and occipital bone marrow before intervention. Although SCAassociated quantitative ADC values in the skull have not been previously assessed in the literature to our knowledge, the significant increase in ADC and facilitated diffusion following BMT raises the possibility that marrow diffusion is reduced at baseline compared with age-expected values. These increased marrow ADC values were sustained across time on follow-up examinations after BMT, suggesting diminished or resolved hematopoietic marrow proliferation. No significant change in marrow ADC values was found in patients with SSA pre- to posttransfusion. However, interpretation was confounded because transfusion causes increased bone marrow T1 pro-

longation and T2 shortening due to iron deposition. ^{2,3} Thus, we excluded examinations with qualitative marrow hypointensity from marrow DWI assessment. Unfortunately, in the transfusion group, this exclusion left only 2 patients for analysis.

The ASL signal magnitude is influenced by a number of factors in addition to blood flow, namely the spin-lattice relaxation time (T1) of arterial blood, blood flow velocity, labeling efficiency, and the brain-blood partition coefficient of water. The latter variable is difficult to accurately measure and is therefore typically estimated at 0.9 mL/g for averaged brain gray and white matter. Labeling efficiency is inversely related to blood flow velocity and requires additional measurements for calculation, such as with phase-contrast imaging. However, the T1 of blood can and should be adjusted on a patient-to-patient basis, especially if there are known deviations from normal hematocrit. Anemia will prolong the T1 value of blood and cause spuriously increased ASL signal that, all other factors being equal, would be interpreted as hyper-

perfusion. Arterial T1 values can be measured directly, but specific sequences are required at the time of MR imaging. ^{23–25} Alternatively, as used in our study, the T1 value can be calculated after determining the hematocrit, $\rm O_2$ saturation, and magnetic field strength. ¹²

Cerebral perfusion decreases during childhood in normal states. 26 It has been shown that anemia positively correlates with cerebral blood flow and negatively correlates with cerebrovascular reserve and that these relationships are strongly associated with anemia severity in children with SCA.⁵ In untreated patients with asymptomatic sickle cell anemia, cerebral blood flow is diffusely elevated relative to aged-matched healthy subjects without regard for age, manifested on ASL as diffusely increased brain signal.⁶ Before intervention, we estimated the mean whole-brain CBF in our SCA cohort (transfusion and BMT groups combined) to be elevated at 78 \pm 11 mL/100 g of tissue/min (64 \pm 14 mL/100 g of tissue/min is considered normal).²⁷ Previous literature reported quantitative whole-brain CBF values in patients with SCA to be in a similar range (83–87 \pm 24 mL/100 g of tissue/min), though different ASL techniques were used and anemia-related blood T1 value adjustments were not made in that study, possibly accounting for slightly higher reported values.^{6,28}

Therapeutic measures in SCA include blood transfusions and allogenic bone marrow transplantation. The improved O2 delivery conferred by transfused normal red blood cells enhances cerebrovascular reserve, decreases the oxygen extraction fraction, and decreases CBF.^{5,23} Guilliams et al²³ reported a significant mean CBF drop following transfusion, from 88 to 82 mL/100 g of tissue/ min. A minority of patients in our cohort also experienced decreased CBF posttransfusion; however, the mean whole-brain CBF encompassing the entire group did not change significantly (from 81 to 80 mL/100 g of tissue/min). It is unclear whether this finding could relate to a smaller sample size and/or differences in techniques among studies. In contrast, whole-brain CBF significantly decreased after BMT (from 76 to 63 mL/100 g of tissue/ min) and was sustained across time on follow-up MR imaging. This change occurred as early as 3 days after transplantation and, therefore, could be a neuroimaging marker heralding favorable engraftment. Engraftment can be observed clinically from 1 to 4 weeks posttransplantation with the definition based on a sustained rebound in the absolute neutrophil count of $\geq 500/\mu L$ and platelet count of \geq 20,000/ μ L.²⁹

Several study limitations should be noted. The sample size is relatively small due to the rarity of SCA-related BMT, the high prevalence of brain and vascular pathology factoring into exclusions, and the novelty of ASL. The posttransfusion bone marrow diffusion data were particularly sparse because iron deposition precluded accurate interpretation in most cases. In the few cases that required single cerebral hemisphere exclusion from analysis, contralateral cerebral ASL perfusion data could have been unreliable to some extent; however, there was no significant difference between pre- and posttransfusion and pre- and posttransplantation CBF values between the subgroups that underwent unilateral and bilateral cerebral analysis to suggest that unilateral cerebral hemisphere exclusion from analysis had any bearing on the results. Future studies with a larger number of subjects will be useful to confirm our findings. Increased blood velocity due to chronic

anemia is a known cause of decreased labeling efficiency using ASL, which may have resulted in underestimation of absolute CBF.^{30,31} Because this was a retrospective study, we did not have corresponding phase data available to approximate labeling efficiency in each case. However, relative changes across time and between groups would be unaffected. In addition, we used a single rather than a multicompartmental model for quantitative ASL calculation, and we did not account for venous outflow effects in SSA.^{24,30}

CONCLUSIONS

Improved cerebral blood flow and marrow diffusion ensue following bone marrow transplantation in children with sickle cell anemia. We found no significant alteration in CBF or marrow diffusion in patients with SCA who underwent transfusion therapy, though the sample size is a limitation. ASL perfusion and quantitative marrow diffusion values are useful posttherapeutic imaging biomarkers.

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