

Microfluidic assessment of red blood cell mediated microvascular occlusion

Journal:	Lab on a Chip
Manuscript ID	LC-ART-02-2020-000112.R1
Article Type:	Paper
Date Submitted by the Author:	23-Apr-2020
Complete List of Authors:	Man, Yuncheng; Case Western Reserve University Kucukal, Erdem; Case Western Reserve University An, Ran; Case Western Reserve University Watson, Quentin; Case Western Reserve University Bosch, Jürgen; Case Western Reserve University Zimmerman, Peter; Case Western Reserve University Little, Jane; University of North Carolina at Chapel Hill Gurkan, Umut; Case Western Reserve University,

SCHOLARONE[™] Manuscripts

Microfluidic assessment of red blood cell mediated microvascular occlusion

Yuncheng Man^a, Erdem Kucukal^a, Ran An^a, Quentin D. Watson^b, Jürgen Bosch^c, Peter A. Zimmerman^b, Jane A. Little^d, and Umut A. Gurkan^{a,e*}

^a Department of Mechanical and Aerospace Engineering, Case Western Reserve University, Cleveland, OH, USA

^b Center for Global Health and Disease, Department of Pathology, Case Western Reserve University, Cleveland, OH, USA

^c Division of Pediatric Pulmonology, Allergy and Immunology, Case Western Reserve University, Cleveland, OH, USA

^d Department of Hematology and Oncology, School of Medicine, University of North Carolina at Chapel Hill, NC, USA

^e Department of Biomedical Engineering, Case Western Reserve University, Cleveland, OH, USA

*Corresponding Author:

LC-ART-02-2020-000112.R1

Umut A. Gurkan, Ph.D. Warren E. Rupp Associate Professor Case Western Reserve University 10900 Euclid Ave., Cleveland, OH 44106 Phone: +1 (216) 368-6447 Email: <u>umut@case.edu</u> Web: <u>https://engineering.case.edu/emae/faculty/umut_gurkan</u>

ABSTRACT

Abnormal red blood cell (RBC) deformability contributes to hemolysis, thrombophilia, inflammation, and microvascular occlusion in various circulatory diseases. A quantitative and objective assessment of microvascular occlusion mediated by RBCs with abnormal deformability would provide valuable insights into disease pathogenesis and therapeutic strategies. To that end, we present a new functional microfluidic assay, OcclusionChip, which mimics two key architectural features of the capillary bed in the circulatory system. First, the embedded micropillar arrays within the microchannel form gradient microcapillaries, from 20 µm down to 4 µm, which mimic microcapillary networks. These precisely engineered microcapillaries retain RBCs with impaired deformability, such that stiffer RBCs occlude the wider upstream microcapillaries, while less stiff RBCs occlude the finer downstream microcapillaries. Second, the micropillar arrays are coupled with two side passageways, which mimic the arteriovenous anastomoses that act as shunts in the capillary bed. These side microfluidic anastomoses prevent microchannel blockage, and enable versatility and testing of clinical blood samples at near-physiologic hematocrit levels. Further, we define a new generalizable parameter, Occlusion Index (OI), which is an indicative index of RBC deformability and the associated microcapillary occlusion. We demonstrate the promise of OcclusionChip in diverse pathophysiological scenarios that result in impaired RBC deformability, including mercury toxin, storage lesion, end-stage renal disease, malaria, and sickle cell disease (SCD). Hydroxyurea therapy improves RBC deformability and increases fetal hemoglobin (HbF%) in some, but not all, treated patients with SCD. HbF% greater than 8.6% has been shown to improve clinical outcomes in SCD. We show that OI associates with HbF% in 16 subjects with SCD. Subjects with higher HbF levels (HbF>8.6%) displayed significantly lower OI ($0.88\% \pm 0.10\%$, N=6) compared with those with lower HbF levels (HbF \leq 8.6%) who had greater OI (3.18% ± 0.34%, N=10, p<0.001). Moreover, hypoxic OcclusionChip assay revealed a significant correlation between hypoxic OI and subject-specific sickle hemoglobin (HbS) level in SCD. OcclusionChip enables versatile in vitro assessment of microvascular occlusion mediated by RBCs in a wide range of clinical conditions. OI may serve as a new parameter to evaluate the efficacy of treatments improving RBC deformability, including hemoglobin modifying drugs, anti-sickling agents, and genetic therapies.

INTRODUCTION

LC-ART-02-2020-000112.R1

Deformability of red blood cells (RBCs) is critical for continuous blood flow in the microcirculation ^{1, 2}. Healthy RBCs are highly deformable, allowing them to pass through minuscule blood vessels to facilitate oxygen delivery. However, in microcirculatory disorders associated with RBCs, such as in sickle cell disease (SCD) and in malaria, RBC deformability is pathologically altered. In SCD, abnormal RBC deformability is induced by intracellular polymerization of sickle hemoglobin (HbS) under deoxygenated conditions ³. Impaired RBC deformability significantly contributes to the pathophysiological hallmarks of this disorder, including hemolysis, inflammation, microvascular occlusion, and consequent organ failures and stroke ^{4, 5}. In malaria, progressively diminished RBC deformability is caused by membrane-protein network modifications associated with maturing Plasmodium parasites ^{6,7}. Unparasitized RBCs can also lose their deformability in severe malaria 8. Thus, loss of RBC deformability has been associated with severe malaria ^{9, 10}. Consequently, there is a strong impetus to develop robust, functional in vitro RBC deformability assays for standardized and quantitative assessment of disease status and severity. However, sampling large quantities of RBCs in near physiological hematocrit levels is often a limiting factor in developing such assays, since the pathological cells often constitute a small fraction of the entire RBC population. Therefore, a large number of RBCs need to be tested simultaneously to acquire meaningful data, as we demonstrate here using OcclusionChip microfluidic system.

Conventional techniques developed for RBC deformability assessment can be divided into single cell techniques and bulk flow techniques. Single cell techniques, including atomic force microscopy ¹¹, micropipette aspiration ^{12, 13}, and optical tweezers ^{14, 15}, measure cell deformability individually. However, these techniques are typically skill-demanding and time-consuming, which require professional personnel or expensive facilities, and therefore are not capable of providing sufficient blood processing volume throughput to measure a subset of pathological RBCs over a large cell population. Bulk flow techniques, including micropore filtration ¹⁶ and ektacytometry ^{17, 18}, measure the average deformability of bulk cells, but provide obscure information on subpopulations of diseased cells. Recent advancement in microfluidic

technologies have yielded various methods for assessing RBC deformability, based on RBC stretching under shear flow ¹⁹⁻²¹, capillary obstruction ²², wedging in tempered constrictions ^{23, 24}, transit velocity or transit time through constrictions ^{25, 26}, as well as transit pressure through constrictions ²⁷⁻²⁹. While all these methods greatly improved *in vitro* RBC deformability analysis, they remain relatively low-throughput in terms of processing volume, and have significant limitations in testing near-physiological hematocrit blood samples.

Microfluidic filtration has been used to sort cells based on deformability, and has been applied to separating circulating tumor cells from blood ^{30, 31}, and leukocytes from RBCs ³². Studies have employed microspheres to mimic the mechanical filtration of poorly deformable RBCs by the human spleen ^{33, 34}. This method yielded retention rates for pathological RBCs, but provides indirect observation as they pass through the device. Other studies utilized polymer mechanical filtering units (micropillar arrays or capillary networks) achieved by soft lithography to mimic the capillary networks and to assess occlusive events induced by poorly deformable RBCs ³⁵⁻³⁸. While these methods demonstrate improvements and they directly measure capillary obstruction, challenges remain around not being able to process samples at near physiologic hematocrit levels.

In the circulatory system, capillary blood vessel size is not uniform, on the contrary, it gradually changes from the arterioles to microcapillaries and to venules (**Fig. 1A**). Arteriovenous anastomoses are a critical part of the circulatory system providing alternate flow paths around capillary beds ^{39, 40}. Anastomotic passageways help regulate blood flow in the event of constrictions or blockages in smaller microcapillaries. Here, we present a new microfluidic design embodied in OcclusionChip (**Fig. 1B**), which mimics two key features of the capillary bed: a gradient of microcapillary networks in the form of micropillar arrays, and side passageways mimicking the arteriovenous anastomoses (**Fig. 1C**). The embedded micropillar arrays form a gradient of microcapillaries from 20 µm spacing down to 4 µm spacing along the flow direction. In this design, red blood cells (RBCs) with significantly impaired deformability are retained by upstream coarser micropillar arrays, while those with mildly impaired

LC-ART-02-2020-000112.R1

Main Manuscript

deformability are retained by finer downstream micropillar arrays. If the upstream micropillar arrays are fully occluded, blood cells can still flow around and into the downstream micropillar arrays through the anastomotic side passages (**Fig. 1D**). This design allows versatility and processing of near-physiological hematocrit level clinical blood samples and full utilization of the entire microcapillary domain. Leveraging such capabilities of OcclusionChip, we report RBC mediated microcapillary occlusion for a wide spectrum of conditions and disorders, including heavy metal toxin exposure (i.e., mercury), banked blood, and microcirculatory diseases, such as end stage renal disease, malaria, and SCD. We present the design, fabrication, and clinical testing of OcclusionChip for standardized assessment of RBC deformability and associated microvascular occlusion. We introduce a new generalizable parameter, 'Occlusion Index', which may serve as an *in vitro* benchmark of microvascular health, function, and may allow clinical assessment of treatments that target improving RBC deformability.

RESULTS

OcclusionChip design and characterization

We have designed the OcclusionChip featuring embedded micropillar arrays forming uniform microcapillaries within each array and overall finer microcapillaries along the flow direction from 20 μ m down to 4 μ m, coupled with two 60- μ m side passages. Microfluidic system design was based on the information available in the literature on human capillary bed and arteriovenous anastomoses ^{39, 40}. OcclusionChip mimics the narrow capillaries, small venules, and arteriovenous anastomoses in the human capillary bed (**Fig. 1A&B**). Characterization of the fabricated PDMS block with micropillar features were achieved by scanning electron microscopy, from which the geometry and dimension of the fabricated PDMS block with microcapillary dimensions were confirmed (**Fig. 1C**). Briefly, the fabricated PDMS micropillars are of 12- μ m height, 20- μ m length, and 10- μ m width. Each micropillar array is 2-mm long and the distance between two successive micropillar arrays is 75 μ m (**Fig. 1C**). Two 60- μ m anastomosis-mimicking side passages were designed to prevent full blockade of the microfluidic channel (**Fig. 1C&D**).

Testing of the micropillar retention mechanism and the function of anastomosis-mimicking side passages was achieved using glutaraldehyde-stiffened RBCs (**Video S1**). RBCs deform and pass through the microcapillaries where no occlusion presented. Such observation confirmed that the micropillar arrays served as a retention mechanism, which is critical for deformability assessment. Further, once severe microcapillary occlusion is induced in the near-inlet portion of the micropillar array (**Fig. 1D** and **Video S1**), local RBC flow is diverted to the nearest anastomotic passage such that full blockade of the microchannel is prevented. The flow environment was characterized via a 2-D numerical simulation performed by a commercially available FEA software package COMSOL 4.3 (Burlington, MA) with the assumption of Newtonian behavior ⁴¹. The boundary conditions at the inlet and outlet were set to pressure inlet and pressure outlet respectively, where the RBC suspension was modeled with a dynamic viscosity and density of 0.001 Pa · s and 993 kg/m^{3 42}. The possible effects of fluid-structure interactions on the flow were neglected in the simulation by treating PDMS structures as rigid bodies. Typical velocity distributions are shown at the micropillar array (**Fig. S1A**), as is estimated maximum velocity and shear rate values across the microchannel (**Fig. S1B**).

Validation testing of OcclusionChip micropillar arrays with rigid fluorescent microbeads

The functional validation of the OcclusionChip was carried out using rigid fluorescent microbeads (Polysciences Inc., Warrington, PA) at 4, 6, or 10-µm diameter. Solutions containing fluorescent microbeads were prepared by mixing the microbead stock solution with BlockAid blocking solution (Thermo Fisher) at a ratio of 1:200 v/v. A visually clear line was observed by the blockage of the 10-µm microbeads at the beginning of the micropillar array with 8-µm gaps (**Fig. S2A**). The same field of view of all the three tests using the three different diameters of microbeads are shown, in which the microbeads were retained within gaps smaller than their outer diameter (**Fig. S2B&C**). This result further confirmed that the micropillar arrays served as a retention mechanism, which is essential to RBC deformability assessment.

Occlusion Index (OI) as a biomarker to assess RBC mediated microcapillary occlusion

To assess the overall RBC mediated microcapillary occlusion in specific samples, we defined a new generalizable parameter 'Occlusion Index' (OI) using the following equation,

$$Occlusion \,Index = \frac{\sum_{i=1}^{n} (c_1 + s \times (i-1)) \times O_i}{\sum_{i=1}^{n} (c_1 + s \times (i-1)) \times N_i} \times 100\% \tag{1}$$

Where

LC-ART-02-2020-000112.R1

- $i = The \ No. \ of \ the \ capillary \ network \ (from \ the \ outlet)$ $c_1 = The \ smallest \ capillary \ dimension$ $s = Step \ change \ of \ capillary \ dimension$ $n = Total \ number \ of \ capillary \ networks \ within \ the \ area \ of \ interest$
- $N_i = Total number of capillaries formed in the ith capillary network$

 O_i = Total number of capillary occlusions induced in the ith capillary network

The OI represents percent occlusion of a microcapillary network, therefore it is generalizable, and can be applied to any microfluidic platform employing capillary networks to assess occlusions due to cellular blockage. We used a microfluidic model employing micropillar arrays assessing RBC deformability in three different scenarios, normal (HbAA) in normoxia (**Fig. S3A**), SCD (homozygous HbSS) in normoxia (**Fig. S3B**), and SCD (homozygous HbSS) in hypoxia (**Fig. S3C**), to demonstrate the generalization of OI. Detailed OI generalization is shown in Supplementary Materials.

OcclusionChip test repeatability

A manual syringe pump was adopted in the experimental process to provide constant pressure source, which significantly improved the cost efficiency of the OcclusionChip assay (**Fig. S4A&B**). To evaluate the repeatability of the experimental process, we tested one normal sample using RBCs from a healthy

donor, and one glutaraldehyde-treated sample using RBCs from another healthy donor. Each sample was run four times, using four different OcclusionChip units, to assess the repeatability of testing. The deformability distribution was highly consistent, where the OIs for the normal sample were 0.19, 0.21, 0.23, and 0.21 (%) (mean \pm standard error of the mean (SEM)=0.21% \pm 0.01%), and the OIs for the glutaraldehyde-treated sample were 3.18, 3.66, 3.36, and 3.54 (%) (mean \pm SEM=3.44% \pm 0.11%). Together, these data show that the variation is minimal, and the test is repeatable.

Occlusion Index to assess chemically treated RBCs with glutaraldehyde, diamide, or mercuric ion

Glutaraldehyde, a non-specific protein cross-linker, and diamide, a thiol group oxidant inducing disulfide cross-linking of band 3, were used to stiffen RBCs in order to mimic impaired RBC deformability, seen in pathological conditions 43, 44. Precisely controlled exposure of RBC to graded concentrations of glutaraldehyde or diamide was utilized to verify OcclusionChip functionality. RBCs from healthy donors exposed to graded concentrations, 0.00% (as control), 0.02%, 0.04%, and 0.08% (w/v), of glutaraldehyde were mixed with untreated normal RBCs to achieve 1% treated cells and were suspended in PBS at 20% hematocrit for microfluidic assessment. RBCs exposed to the same graded concentrations of diamide were directly suspended in PBS for microfluidic assessment. RBCs with impaired deformability were mostly retained at the 4-µm microcapillaries (Fig. 2A), with a smaller number retained at the 6, 8, and 10um microcapillaries (not shown). No appreciable occlusion was observed elsewhere. The retained RBCs were fluorescently labeled with anti-glycophorin A (CD235a, Abcam, Cambridge, MA) antibodies. Results obtained with RBCs from the same blood sample that was exposed to graded concentrations of glutaraldehyde are shown (Fig. 2B, control not shown). Of note, the distribution of retained RBCs within the channel revealed an important aspect of the microfluidic device: it generated distinct patterns in response to the heterogeneous RBC deformability that was induced by various concentrations of glutaraldehyde (Fig. 2C). At the end of the test, numbers of occlusions induced by poorly-deformable RBCs within each micropillar array were quantified (Fig. 2D-F). The number of occlusions induced by RBCs increased as the concentration of glutaraldehyde or diamide increased. The maximum difference in

LC-ART-02-2020-000112.R1

Main Manuscript

total occlusion level was observed with glutaraldehyde treatment at a concentration of 0.08% (w/v), where the rate of increase was more than 16-fold compared to the control. Further, we defined the area of interest as the 4-µm to 10-µm micropillar arrays to compute the OI to assess the overall RBC deformability of individual samples, since typical capillary dimension is 5-10 µm. Our results show that the OI generated by RBCs exposed to glutaraldehyde or diamide increased in a concentration dependent manner (glutaraldehyde: **Fig. 2G**, p<0.001, one-way ANOVA, mean \pm SEM=0.31% \pm 0.03% for control, 1.57% \pm 0.25% for 0.02% glutaraldehyde, 2.17% \pm 0.56% for 0.04% glutaraldehyde, 6.10% \pm 0.74% for 0.08% glutaraldehyde, and diamide: **Fig. 2H**, p=0.01, one-way ANOVA, mean \pm SEM=0.37% \pm 0.07% for control, 0.42% \pm 0.05% for 0.02% diamide, 0.58% \pm 0.09% for 0.04% diamide, and 0.82% \pm 0.11% for 0.08% diamide), which indicates that the defined OI reflect the deformability of the entire RBC population.

Mercury is a toxic heavy metal that binds to RBC membranes with high affinity and results in RBC hemolysis ^{45, 46}. Mercuric ions induce membrane conformational changes and impair RBC deformability ⁴⁷. In this study, we interrogated the effect of short-term exposure of mercuric ion on RBC deformability, at both low (5 μ M Hg²⁺) and high levels (50 μ M Hg²⁺), as the range of the mercury level *in vivo* has been reported at 16,000 μ g/L (58 μ M) to 11 μ g/L (0.055 μ M) in blood ⁴⁸. Hg²⁺-exposed RBC samples were perfused through the OcclusionChip, and enhanced retention of poorly deformable RBCs was observed at the 4- μ m microcapillaries for all Hg²⁺-treated samples (**Fig. 2F**), suggesting moderate impairment in RBC deformability following short-term mercuric ion exposure. The OI generated by Hg²⁺-exposed RBCs increased in a mercuric ion concentration dependent manner (**Fig. 2I**, p=0.007, Kruskal-Wallis, median \pm SEM=0.29% \pm 0.03% for control, 0.50% \pm 0.08% for 5 μ M, and 1.00% \pm 0.15% for 50 μ M mercury). Scanning electron microscopy results confirm that mercury induced morphologic changes in the RBCs, including the loss of the characteristic biconcave shape and the formation of spiked cell membrane (**Fig. S5**). Detailed sample processing for scanning electron microscopy is summarized in Supplementary Materials. Taken together, our results demonstrate that short-term mercuric ion exposure induces RBC

morphology change and moderately reduces RBC deformability, which can be assessed using OcclusionChip.

Occlusion Index to assess impaired RBC deformability induced by *Plasmodium falciparum* infection, blood storage lesion, or end-stage renal disease

To demonstrate the versatility and clinical relevance of OcclusionChip, we tested RBCs from a stored blood unit, exposed to *Plasmodium falciparum* infection, from subjects with end-stage renal disease, or from healthy donors. Typical retained Hoechst-labeled *Plasmodium falciparum*-infected RBCs (*pf*-iRBCs) within the microchannel are shown (**Fig. S6**). We observed that the *pf*-iRBCs were preferentially retained by various sizes of microcapillaries, which generated a significantly higher level of OIs, compared to normal, non-infected RBCs (**Fig. 3**, N=5, malaria: mean \pm SEM=4.09% \pm 1.06%, and normal: N=5, 0.15% \pm 0.03%, p=0.012, Mann-Whitney). In addition, the OI of one blood sample stored for up to 6 weeks were closely monitored, from which we observed the OI increased as the storage progressed, indicating the progressively decreased deformability of the stored RBCs (**Fig. 3**, mean=0.27% for fresh, 0.39% for week 1-2, 0.50% for week 3-4, and 0.55% for week 5-6). The progressively reduced deformability of stored RBCs is also reflected by the hemolysis level, glucose consumption, and lactate production (**Fig. S7A-D**). Furthermore, RBCs from hemodialyzed samples were assessed, from which we observed a small change of OI due to the retention of poorly deformable RBCs primarily within the 4-µm micropillar array, suggesting a moderate reduction of RBC deformability in subjects with end-stage renal disease. (**Fig. 3**, N=2, mean \pm SEM=0.43% \pm 0.01%).

Occlusion Index to assess pathologically impaired RBC deformability in SCD

To further demonstrate the clinical and pathophysiological relevance of OcclusionChip, we assessed RBCs from 16 adult subjects with homozygous (HbSS) SCD (**Fig. 3**). The close-up view of microcapillaries in OcclusionChip show that, the occlusions of 4-µm microcapillaries were induced by a morphologically heterogeneous RBC population from subjects with SCD, including disc-shaped

LC-ART-02-2020-000112.R1

Main Manuscript

(indicated in orange zone), mildly sickled (indicated in rose zone), and highly sickled (indicated in red zone) cell morphologies within the same field of view (**Fig. 4A**). Notably, HbSS RBCs from subjects with SCD appeared to stagnate around the front surface of the micropillars due to the local flow environment (the dead zone showed in **Fig. S1A**). The numbers of occlusions induced by HbSS RBCs across the device were quantified (**Fig. 4B**). OI of HbSS RBCs from subjects with SCD was significantly higher compared to normal RBCs (HbAA) from healthy donors (**Fig. 4C**, HbSS: N=16, mean \pm SEM=2.32% \pm 0.36%, and HbAA: N=5, 0.15% \pm 0.03%, p=0.001, Mann-Whitney).

Occlusion Index associates with fetal hemoglobin level (HbF%) in subjects with SCD

HbF% level in blood has been shown to associate with clinical outcomes in SCD and it is inducible via Hydroxyurea therapy ⁵⁰, although the treatment response is heterogeneous among patients ^{51, 52}. HbF level greater than 8.6% has been considered as a clinically therapeutic threshold in SCD ⁵³. Therefore, we investigated the association between OI measured by OcclusionChip and the subject-specific HbF% in the 16 adult subjects with SCD (**Fig. 5**). We observed that subjects with higher HbF levels (>8.6%, N=6) had significantly lower OIs (**Fig. 5A**, N=6, mean \pm SEM=0.88% \pm 0.10%), compared to those with lower HbF levels (<8.6%) who had significantly higher OIs (N=10, mean \pm SEM=3.18% \pm 0.34%) as measured by OcclusionChip (p<0.001, one-way ANOVA). Among the 11 subjects tested who were on Hydroxyurea therapy, 5 subjects had HbF% less than 8.6% and lower OIs (**Fig. 5B**, triangles) compared with subjects who were on Hydroxyurea with HbF greater than 8.6%.

Hypoxic Occlusion Index associates with sickle hemoglobin level (HbS%) in subjects with SCD

Since hypoxia regulates hemoglobin desaturation and therefore the pathophysiological process in SCD, we created a deoxygenation system to couple OcclusionChip to study the behavior of RBCs in postcapillary networks in SCD (**Fig. 6A**). We utilized a custom gas exchanger to deoxygenate the blood flow

developed in one of our previous studies (**Fig. 6B**), which desaturates hemoglobin in the sample to an approximate SpO₂ of 83% ⁵⁴. We designed the custom gas exchange tubing and the gas exchange microchannel to stabilize the oxygen level within the microchannel as the gas permeability of PDMS was recognized (**Fig. 6C**). The detailed fabrication process is presented in Supplementary Materials. Validation of oxygen diffusion was performed by introducing an oxygen-sensitive luminescence probe tris (4,7-diphenyl-1,10-phenanthroline) ruthenium (II) dichloride complex (Santa Cruz Biotechnology, Dallas, TX) (1 mg/ml in PBS) into the microchannel (**Fig. 58**). Molecular oxygen can quench the luminescence of this probe; thus the dissolved oxygen level can be measured by the luminescent intensity ⁵⁵. In our experiment, a controlled gas flow (95% N₂ plus 5% CO₂) was allowed at 0 s and the luminescence was monitored under illumination at 488 nm. Gas exchange was completed at approximately 420 s as the luminescent intensity saturated (**Fig. 6D**).

RBCs from 7 subjects with SCD (4 homozygous HbSS, and 3 compound-heterozygous HbSC subjects, all termed as HbS-carrying) and 3 healthy donors were tested under ambient or hypoxic conditions. Although it has been reported that the sickling of HbS-carrying RBCs depends on the blood oxygen level ⁵⁶, here we observed heterogeneous levels of cell sickling and changes in morphology under identical degrees of hypoxia (**Fig. 6E** and **Video S2**). As expected, HbAA RBCs showed no change in morphology under hypoxic conditions (not shown). We tested the effect of hypoxia on the OI of HbS-carrying RBCs and HbAA RBCs. The sample perfusion duration was decreased to 2 minutes in order to prevent saturation in hypoxic OcclusionChip. Our scanning electron microscopy results show that when oxygen level is reduced, HbS-carrying RBCs undergoing sickling were unable to pass through the microcapillaries, thereby inducing enhanced microchannel occlusion and resistance to blood flow (**Fig. 6F**). Detailed sample processing for scanning electron microscopy is summarized in Supplementary Materials. The OI was significantly greater in hypoxic sickle RBCs compared to sickle RBCs examined under ambient conditions. As expected, the OI of HbAA RBCs did not change under hypoxia (**Fig. 6G**, HbS-carrying RBCs: N=7, mean \pm SEM=0.25% \pm 0.03% vs 57.31% \pm 4.79%, and HbAA: N=3, 0.05% \pm

LC-ART-02-2020-000112.R1

Main Manuscript

0.01% vs $0.05\% \pm 0.01\%$ for normoxia vs. hypoxia). Profiles of occlusions induced by HbAA RBCs and HbS-carrying RBCs in hypoxic conditions are shown, from which full saturation of the 4-µm and 6-µm micropillar arrays were observed for all HbS-carrying RBC samples (**Fig. 6H**). A statistically significant correlation between the OI and HbS% of the subjects was observed (**Fig. 6I**, PCC=0.78, p=0.04, N=7). Of note, we observed recovery of sickled HbS-carrying RBCs (**Video S3**) as well as clearance of retained HbS-carrying RBCs within the microchannel (**Video S4**) by stopping the controlled gas flow, which demonstrates that the intracellular HbS polymerization and RBC sickling due to hypoxia is reversible, and OcclusionChip is capable of detecting this effect.

DISCUSSION

In OcclusionChip, embedded micropillar arrays are coupled with side passageways mimicking arteriovenous anastomoses in the capillary bed. Other capillary microfluidic systems that we are aware of lack these side passages around capillary openings 57, 58. This design feature enables versatility for a wide variety of clinical and pathological conditions (Fig. 3), such that if the upstream micropillar arrays are blocked in a severe case, blood cells can still reach downstream arrays through the microfluidic anastomoses (Fig. 1D). OcclusionChip design allows processing of clinical blood samples near physiological hematocrit levels (20% hematocrit), while other microcapillary based microfluidic systems typically accommodate low hematocrit levels (e.g., 1% hematocrit in recent studies ^{26, 59}). Nearphysiological hematocrit blood sample processing of OcclusionChip was vital for testing a wide variety of clinical blood samples presented in this work (Fig. 3). OcclusionChip design allows more than three orders of magnitude greater throughput in processing volume (3,000,000 cells per minute) compared to other microfluidic designs that we are aware of (e.g., 600 cells per minute in a recently published study ⁵⁹, or ~ 0.5 million cells per hour as reported in another microfluidic system ²⁹). Typical blood sample flow rate in OcclusionChip was measured to be approximately 1.5 µL/minute at near-physiological (20%) hematocrit level, which translates to processing of 3,000,000 cells per minute (with estimated 2 million cells/µL of blood at 20% hematocrit 60). OcclusionChip design enables a wide dynamic OI range.

LC-ART-02-2020-000112.R1

Main Manuscript

Previous microfluidic systems utilized single size constrictions by design, and therefore, if the constrictions are too large or too small, the system is not sufficiently sensitive to interrogate cells with mildly or severely compromised deformability. For example, if the constrictions are too narrow, such systems are swiftly saturated by cells with severely compromised deformability, which significantly reduces the dynamic detection range. In OcclusionChip, stiff RBCs are retained by upstream, coarser micropillar arrays, while those less stiff flow through the side anastomoses and are interrogated and/or retained by finer downstream micropillar arrays. These characteristics contribute to the wide dynamic range of OI measurement of OcclusionChip.

To verify the retention mechanism and demonstrate the physiological relevance of the OcclusionChip, we utilized precisely controlled exposure of normal RBCs to glutaraldehyde or diamide, which are widely used to verify new RBC deformability measurement techniques. We further interrogated abnormal RBC deformability that is induced by exposure to mercuric ion. We found that short-term mercuric ion exposure altered RBC deformability in a concentration dependent manner (**Fig. 2I**, **Fig. 3**), which is in accordance with early *in vitro* studies on the hemolytic effect of inorganic mercuric ion on RBCs compromising membrane fragility and cell deformability ^{47, 61}.

Envisioning a universal RBC deformability assessment tool for all circulatory diseases, we further used the OI to assess the abnormal deformability of *in vitro* malaria-infected RBCs, and RBCs from patients with end-stage renal disease (**Fig. 3**). Early *ex vivo* studies have revealed the impaired deformability of malaria infected RBCs, which is stage dependent, decreasing as intra-erythrocytic parasites mature ^{33, 62, 63}. In full agreement with this, we observed enhanced occlusions within multiple micropillar arrays induced by malaria infected RBCs, with a markedly increased OI compared to normal RBCs (**Fig. 3**, **Fig. S6B**). It has been shown that RBC lifetime is significantly reduced in patients with end-stage renal disease due to uremic toxins, mechanical and osmotic trauma ^{64, 65}. Lipidic and phospholipidic compositions on cell membrane are significantly altered ⁶⁶ in these patients. However, it is still not clear why the uremic state leads to impaired RBC deformability and reduced lifetime in circulation ⁶⁷. Previous studies using

LC-ART-02-2020-000112.R1

Main Manuscript

membrane filters demonstrated that RBCs from patients with end-stage renal disease required more time to pass through and thus they were less deformable compared to normal RBCs ⁶⁸⁻⁷⁰. In full agreement, we observed that RBCs from subjects on hemodialysis presented mildly elevated OIs, suggesting that RBCs from individuals with end-stage renal disease display mildly impaired deformability.

SCD is one of the common inherited blood disorders worldwide, which results in abnormal intracellular hemoglobin polymerization and reduced RBC deformability ^{71, 72}. As expected, RBCs from subjects with homozygous SCD (HbSS) resulted in significantly greater OIs compared with RBCs from healthy donors (HbAA) (Fig. 4C). Assessing RBC deformability is of great importance in SCD, since earlier studies revealed a significant association between the incidence of microcirculatory vaso-occlusive events and impaired cellular deformability 73, 74. Hydroxyurea therapy is a widely utilized Food and Drug Administration (FDA)-approved, pharmaceutical treatment for SCD, which stimulates HbF production, reducing sickling and improving RBC deformability ^{50, 75}. Hydroxyurea increases blood HbF% in some, but not all, treated patients with SCD ^{51, 52}, hence close monitoring of clinical outcomes is essential ⁷⁶. It has been reported that high levels of HbF (>8.6%) associates with improved clinical outcomes and survival in SCD ⁵³. In line with these findings, we observed that OI associates with HbF% in subjects with SCD (Fig. 5A). Furthermore, subjects with higher HbF levels (>8.6%) displayed significantly lower RBC mediated microcapillary occlusion in vitro as indicated by lower OIs (Fig. 5B) compared to those with lower HbF levels (≤8.6%). Importantly, among the SCD subjects tested, 5 out of 11 subjects who were on Hydroxyurea therapy, displayed HbF levels lower than 8.6% and higher OIs (Fig. 5B). The remaining 6 subjects on Hydroxyurea displayed HbF levels greater than 8.6% and lower OIs (Fig. 5B). These findings suggest that OcclusionChip and OI may serve as an *in vitro* therapeutic benchmark in SCD to help identify subjects who may not sufficiently respond to Hydroxyurea with an increased HbF% to a therapeutically effective level. Taken together, the presented OcclusionChip platform, together with other clinical laboratory tests, would be clinically useful in assessing RBC associated microvascular occlusion in SCD.

LC-ART-02-2020-000112.R1

Main Manuscript

Recent studies showed that RBCs become less deformable after storage ⁷⁷⁻⁷⁹. In accordance, our results indicated that during the 6-week storage period, the OI of the stored RBCs increased modestly over time (**Fig. 3**). Transfusion therapy is one of the main treatments utilized in SCD, in which stored RBCs from healthy donors are exchanged with patient's own RBCs to lower the sickle RBC fraction in the circulatory system. The quality of the stored blood units may significantly impact the clinical outcomes of transfusion therapy, especially in anemic hematological disorders such as SCD ⁸⁰. A quantitative assessment of RBC mediated microcapillary occlusion, as described here, would ensure additional quality control for stored blood. Improved quality control in transfusion medicine may considerably enhance outcomes by reducing the frequency of needed transfusions and thereby alleviating the incidence of transfusion-related complications, such as allergic reaction, infection, iron overload, and transfusion therapy present significantly higher RBC adhesion to heme-activated endothelium compared to subjects with SCD who are not on transfusion therapy ⁸⁵. Results presented here suggest that HbF level stays less than 8.6% (as expected), while OI remains higher in 5 subjects with SCD compared with subjects on Hydroxyurea therapy with HbF% greater than 8.6% (**Fig. 5B**).

Hypoxia and hemoglobin desaturation play an essential role in regulating pathophysiological processes in SCD ^{93, 94}. Hypoxia induces intracellular polymerization of sickle hemoglobin, sickling of RBCs, and impairment of RBC deformability and adhesiveness ⁹⁵. Therefore, a functional *in vitro* assay for RBC deformability and microcapillary occlusion under hypoxia may significantly benefit the development and assessment of novel therapeutic strategies that directly target the hemoglobin content of the RBCs, such as anti-sickling drugs and genetic therapies in SCD ⁹⁶⁻⁹⁸. To that end, we coupled the OcclusionChip assay with hypoxia control, and assessed RBC mediated microcapillary occlusion under hypoxia in subjects with SCD (**Fig. 6A-F**). Using hypoxic OcclusionChip assay we revealed a significant association between RBC mediated microcapillary occlusion *in vitro* (**Fig. 6G&H**), the resulting hypoxic OI, and the subject-specific HbS levels in SCD (**Fig. 61**, Pearson Correlation Coefficient: 0.78, p: 0.04). The finding suggests

LC-ART-02-2020-000112.R1

Main Manuscript

that hypoxic OI would be essentially helpful in assessing therapies that target sickle hemoglobin polymerization to improve RBC deformability ^{98, 99} or genetic therapies that aim at reducing the sickle hemoglobin content in blood along with improving RBC qualities in hemoglobin disorders ^{97, 100, 101}. For example, Aes-103 (5-hydroxymethylfurfural) is a putative anti-sickling agent that is in clinical trials for potential treatment of SCD ¹⁰², and therefore we would expect to observe a difference in terms of OI when testing sickle RBCs before and after Aes-103 treatment under hypoxia. On the other hand, a recent study reported that after receiving gene therapy, the fraction of sickle RBCs under hypoxia significantly decreased ¹⁰⁰. Consequently, the OI of RBCs from individuals with SCD measured under hypoxia is expected to drop after receiving gene therapy, which could be considered as a functional assessment of the therapy outcome.

In conclusion, we present a functional microfluidic assay, the OcclusionChip, for assessing microvascular occlusion mediated by RBCs. The significant aspects of this functional assay are anastomosis-mimicking side passageways, the ability to process clinical blood samples at near-physiologic hematocrit, and the versatility of the assay for its potential use in various clinical conditions and diseases. We validated the OcclusionChip assay initially by testing control samples and glutaraldehyde or diamide-treated RBCs. Clinical relevance and versatility of the assay was demonstrated by testing toxic mercuric ion-exposed RBCs, stored blood, subjects with end-stage renal disease, malaria-infected RBCs, and subjects with SCD. Importantly, in subjects with SCD, we show that the OI, an indicative index of RBC mediated microcapillary occlusionChip assay helped us reveal a significant correlation between hypoxic OI and subject-specific HbS% in SCD. Taken together, these findings suggest that OcclusionChip functional microfluidic assay and the OI can be utilized for quantitative assessment of microvascular occlusion mediated by RBCs. OcclusionChip and OI would be applicable to a wide range of clinical conditions and in assessment of novel targeted and curative therapies that improve RBC quality, such as anti-sickling drugs and gene therapies in SCD.

MATERIALS AND METHODS

OcclusionChip fabrication and assembly

The OcclusionChip was fabricated based on standard lithography techniques (Fig. S9). A negative template on a 3-inch silicon wafer (University Wafers, Boston, MA) was initially fabricated through photolithography. Briefly, a layer of negative photoresist SU8-2010 (Thermo Fisher Scientific, Waltham, MA) was spin-coated on the wafer at a thickness of 12 µm. After being soft-baked at 95°C for 4 min, the wafer was exposed to UV light with alignment to selectively cure the photoresist. Following the postexposure bake, in which the wafer was baked in the same condition as soft-bake, the wafer was developed in a photoresist solvent propylene glycol monomethyl ether acetate (PGMEA, Sigma Aldrich, St. Louis, MO), and hard-baked at 110°C overnight. A 2-hour surface passivation using trichloro (1H, 1H, 2H, 2Hperfluorooctyl) saline (Sigma Aldrich, St. Louis, MO) was performed under vacuum to facilitate the separation of the molded polymer from the master wafer. Next, a polydimethylsiloxane (PDMS, Thermo Fisher Scientific, Waltham, MA) pre-polymer was mixed with the curing agent at a ratio of 10:1 (v/v) and degassed in a desiccator to remove any air bubbles. The mixture was poured over the master wafer and cured at 80 °C overnight. Two 0.5 mm-diameter holes were punched as the inlet and the outlet after the PDMS block was separated from the master wafer. Excessive saline was removed by sonicating with isopropanol for 5 min. Tubing was assembled after the PDMS block was bonded to a microscope glass slide (Microscopy Sciences) through surface modification under oxygen-plasma treatment. The fabricated microchannel was rinsed in 100% ethanol and phosphate-buffered saline (PBS), and incubated with 2%

bovine serum albumin (BSA, ProSpec-Tany TechnoGene Ltd, East Brunswick, NJ). Prior to introducing blood samples, the microchannel was rinsed with PBS to remove excessive BSA.

Blood sample collection and preparation

All experiments were performed in accordance with the study protocol approved by the Institutional Review Board (IRB) committee at University Hospitals Cleveland Medical Center (IRB 05-14-07C). Study participants were fully informed regarding the purposes of the study and consent was obtained. Blood samples from healthy donors (HbAA) and SCD subjects (homozygous HbSS and heterozygous HbSC) were anticoagulated with ethylenediaminetetraacetic acid (EDTA) and stored at 4°C. Healthy RBCs were treated with glutaraldehyde, Diamide, and mercuric ion according to protocols detailed in Supplementary Materials. Pf-iRBCs were cultured at 4% hematocrit under standard conditions in RPMI 1640 according to published protocols ¹⁰³. Parasitized RBCs were labeled with Hoechst 33342 for 30 minutes in room temperature. The mean parasitemia of the five malaria samples tested was 4.55 ± 0.29 % (± SEM). Stored blood samples were obtained from Hemanext (Avon, MA). Briefly, packed RBCs were prepared from whole blood anticoagulated with citrate-phosphate-double dextrose (CP2D) anticoagulant after centrifugation and removal of the plasma. The RBCs were re-suspended in AS3 red cell storage solution and stored in a conventional storage bag for 42 days at 4°C. A total of seven blood samples were tested over a six-week period. These samples were shipped to us individually over time, which originated from the same unit of stored blood. Lactate production and glucose consumption were analyzed to determine the metabolic activities of the stored RBCs. Hemolysis levels were well below FDA guidelines of 1%, maximum allowable limit for red cell product for transfusion. Blood samples from patients on hemodialysis were obtained from the clinics at University Hospitals, with IRB approval. HbF% and HbS% levels in whole blood samples were determined using HPLC (VARIANT[™] II, Bio-Rad Laboratories, Inc., Hercules, California).

Microfluidic assessment of blood samples

Whole blood samples were centrifuged at 500 x g for 5 min at 4° C, plasma was removed, and the RBCs were washed twice in PBS, and re-suspended at a 20% hematocrit value. The OcclusionChip was placed on an Olympus IX83 inverted motorized microscope stage for high-resolution imaging. The RBC samples were perfused through the microchannel under constant pressure for 20 min. Non-retained RBCs were rinsed away by injecting PBS into the microchannel under the same pressure. Negligible contamination by platelets and white blood cells was observed by visual inspection. Phase contrast, bright field, and fluorescent images were recorded at 20X using the Olympus Cell Sense live imaging software (excitation/emission wavelength, 488/505-580 for Green Fluorescent Protein, GFP). Recorded images were post-processed using Adobe Photoshop (San Jose, CA), and the number of occlusions in each micropillar array was counted manually. When a microcapillary formed by two neighbored micropillars was obstructed by an RBC, it was counted as one occlusion. Typically, it takes less than 1 hour to complete the microfluidic processing, imaging, and image analysis for testing one clinical blood sample. Oxygen tension was controlled in hypoxia experiments using a custom designed gas exchange setup as described in Supplementary Materials. Scanning electron microscopy was performed on the micropillars and RBCs as described in Supplementary Materials. OcclusionChip microfluidic system was designed to be single use and disposable, to prevent any cross-contamination between the samples tested. Each data point in this manuscript was generated with a single use OcclusionChip.

Statistical analyses

The data are reported as mean \pm SEM in this study. A test of normality was performed. One-way ANOVA was used for normally distributed data with Tukey's post hoc analysis for more than two groups. Non-parametric Mann-Whitney U-test or Kruskal-Wallis H tests with Dunn's post hoc analysis for comparing more than two groups for non-normally distributed data. Statistical significance was defined with p-value less than 0.05 (p < 0.05). Statistical analyses were performed using Minitab 19 (Minitab, Inc., State College, PA) and Matlab (Mathworks, MA, USA).

ACKNOWLEDGMENTS

The authors greatly thank Grace Gongaware from Cleveland Institute of Art for crafting the scientific illustration. The authors also acknowledge Dr. Thomas Hostetter for providing blood samples from hemodialysis patients, and Dr. Samuel Coker from Hemanext for providing the stored blood samples. The authors acknowledge with gratitude the contributions of patients and clinicians at Seidman Cancer Center (University Hospitals, Cleveland). UAG would like to thank the Case Western Reserve University, University Center for Innovation in Teaching and Education (UCITE) for the Glennan Fellowship, which supports the scientific art program and the art student internship in CASE-BML.

FUNDING

This work was supported by National Science Foundation (NSF) CAREER Award 1552782, National Heart, Lung, and Blood Institute (NHLBI) R01HL133574, OT2HL152643, U01HL117659, and T32HL134622.

AUTHOR CONTRIBUTIONS

YM and UAG conceived the idea. YM and EK performed the simulations. YM conducted the experiments, acquired and analyzed data, prepared figures, and wrote the manuscript. YM, EK, RA, JB, PAZ, JAL, and UAG discussed and interpreted data. YM, EK, RA, JB, PAZ, JAL, and UAG reviewed and edited the manuscript. QDW, JB, and PAZ provided cultured malaria samples, and JAL provided blood samples from subjects at University Hospitals Cleveland Medical Center Adult Sickle Cell Clinic.

CONFLICT OF INTERESTS

The authors declare no conflict interest.

REFERENCES

- 1. N. Mohandas and P. G. Gallagher, *Blood*, 2008, **112**, 3939-3948.
- 2. S. Chien, *Microvascular research*, 1985, **29**, 129-151.
- 3. D. C. Rees, T. N. Williams and M. T. Gladwin, *Lancet*, 2010, **376**, 2018-2031.
- G. A. Barabino, M. O. Platt and D. K. Kaul, *Annual review of biomedical engineering*, 2010, 12, 345-367.
- Y. Alapan, A. Fraiwan, E. Kucukal, M. N. Hasan, R. Ung, M. Kim, I. Odame, J. A. Little and U. A. Gurkan, *Expert review of medical devices*, 2016, 13, 1073-1093.
- 6. C. A. Moxon, G. E. Grau and A. G. Craig, *British journal of haematology*, 2011, **154**, 670-679.
- 7. A. M. Dondorp, E. Pongponratn and N. J. White, *Acta tropica*, 2004, **89**, 309-317.
- A. M. Dondorp, B. J. Angus, M. R. Hardeman, K. T. Chotivanich, K. Silamut, R.
 Ruangveerayuth, P. A. Kager, N. J. White and J. Vreeken, *Am J Trop Med Hyg*, 1997, 57, 507-511.
- 9. C. A. Moxon, G. E. Grau and A. G. Craig, *British journal of haematology*, 2011, **154**, 670-679.
- J. Rey, P. A. Buffet, L. Ciceron, G. Milon, O. Mercereau-Puijalon and I. Safeukui, *Sci Rep*, 2014, 4, 3767-3767.
- I. Dulinska, M. Targosz, W. Strojny, M. Lekka, P. Czuba, W. Balwierz and M. Szymonski, Journal of biochemical and biophysical methods, 2006, 66, 1-11.
- 12. R. M. Hochmuth, Journal of biomechanics, 2000, 33, 15-22.
- 13. R. P. Hebbel, A. Leung and N. Mohandas, *Blood*, 1990, **76**, 1015-1020.

- 14. M. Dao, C. T. Lim and S. Suresh, J Mech Phys Solids, 2003, 51, 2259-2280.
- 15. S. Henon, G. Lenormand, A. Richert and F. Gallet, *Biophys J*, 1999, **76**, 1145-1151.
- H. L. Reid, A. J. Barnes, P. J. Lock, J. A. Dormandy and T. L. Dormandy, *Journal of clinical pathology*, 1976, 29, 855-858.
- N. L. Parrow, P. C. Violet, H. Tu, J. Nichols, C. A. Pittman, C. Fitzhugh, R. E. Fleming, N. Mohandas, J. F. Tisdale and M. Levine, *Journal of visualized experiments : JoVE*, 2018, DOI: 10.3791/56910.
- O. K. Baskurt, M. R. Hardeman, M. Uyuklu, P. Ulker, M. Cengiz, N. Nemeth, S. Shin, T. Alexy and H. J. Meiselman, *Biorheology*, 2009, 46, 251-264.
- 19. Y. Zheng, M. A. Cachia, J. Ge, Z. S. Xu, C. Wang and Y. Sun, *Lab Chip*, 2015, 15, 3138-3146.
- 20. Y. Katsumoto, K. Tatsumi, T. Doi and K. Nakabe, Int J Heat Fluid Fl, 2010, 31, 985-995.
- 21. Y. Zheng, J. Nguyen, C. Wang and Y. Sun, *Lab Chip*, 2013, **13**, 3275-3283.
- J. P. Shelby, J. White, K. Ganesan, P. K. Rathod and D. T. Chiu, *P Natl Acad Sci USA*, 2003, 100, 14618-14622.
- A. T. Santoso, X. Y. Deng, J. H. Lee, K. Matthews, S. P. Duffy, E. Islamzada, S. M. McFaul, M. E. Myrand-Lapierre and H. S. Ma, *Lab Chip*, 2015, 15, 4451-4460.
- M. A. Lizarralde Iragorri, S. El Hoss, V. Brousse, S. D. Lefevre, M. Dussiot, T. Xu, A. R. Ferreira, Y. Lamarre, A. C. Silva Pinto, S. Kashima, C. Lapoumeroulie, D. T. Covas, C. Le Van Kim, Y. Colin, J. Elion, O. Francais, B. Le Pioufle and W. El Nemer, *Lab Chip*, 2018, 18, 2975-2984.

- LC-ART-02-2020-000112.R1
- A. Adamo, A. Sharei, L. Adamo, B. Lee, S. Mao and K. F. Jensen, *Anal Chem*, 2012, 84, 6438-6443.
- J. M. Sosa, N. D. Nielsen, S. M. Vignes, T. G. Chen and S. S. Shevkoplyas, *Clin Hemorheol Micro*, 2014, 57, 275-289.
- 27. Q. Guo, S. J. Reiling, P. Rohrbach and H. S. Ma, *Lab Chip*, 2012, **12**, 1143-1150.
- M. E. Myrand-Lapierre, X. Y. Deng, R. R. Ang, K. Matthews, A. T. Santoso and H. S. Ma, *Lab Chip*, 2015, 15, 159-167.
- Q. Guo, S. P. Duffy, K. Matthews, X. Deng, A. T. Santoso, E. Islamzada and H. Ma, *Lab Chip*, 2016, 16, 645-654.
- X. Qin, S. Park, S. P. Duffy, K. Matthews, R. R. Ang, T. Todenhofer, H. Abdi, A. Azad, J. Bazov,
 K. N. Chi, P. C. Black and H. S. Ma, *Lab Chip*, 2015, 15, 2278-2286.
- G. Vona, A. Sabile, M. Louha, V. Sitruk, S. Romana, K. Schutze, F. Capron, D. Franco, M.
 Pazzagli, M. Vekemans, B. Lacour, C. Brechot and P. Paterlini-Brechot, *Am J Pathol*, 2000, 156, 57-63.
- 32. Q. Guo, S. P. Duffy, K. Matthews, E. Islamzada and H. S. Ma, *Sci Rep*, 2017, 7.
- G. Deplaine, I. Safeukui, F. Jeddi, F. Lacoste, V. Brousse, S. Perrot, S. Biligui, M. Guillotte, C.
 Guitton, S. Dokmak, B. Aussilhou, A. Sauvanet, D. C. Hatem, F. Paye, M. Thellier, D. Mazier, G.
 Milon, N. Mohandas, O. Mercereau-Puijalon, P. H. David and P. A. Buffet, *Blood*, 2011, 117, E88-E95.

- S. A. S. Diakite, P. A. Ndour, V. Brousse, F. Gay, C. Roussel, S. Biligui, M. Dussiot, V. Prendki,
 T. M. Lopera-Mesa, K. Traore, D. Konate, S. Doumbia, J. Cros, S. Dokmak, R. M. Fairhurst, M.
 Diakite and P. A. Buffet, *Malaria J*, 2016, 15.
- J. Picot, P. A. Ndour, S. D. Lefevre, W. El Nemer, H. Tawfik, J. Galimand, L. Da Costa, J. A. Ribeil, M. de Montalembert, V. Brousse, B. Le Pioufle, P. Buffet, C. Le Van Kim and O. Francais, *American journal of hematology*, 2015, **90**, 339-345.
- E. Du, M. Diez-Silva, G. J. Kato, M. Dao and S. Suresh, *P Natl Acad Sci USA*, 2015, **112**, 1422-1427.
- J. C. Ciciliano, R. Abbaspour, J. Woodall, C. Wu, M. S. Bakir and W. A. Lam, *Lab Chip*, 2017, DOI: 10.1039/c7lc00720e.
- 38. S. S. Shevkoplyas, T. Yoshida, S. C. Gifford and M. W. Bitensky, *Lab Chip*, 2006, 6, 914-920.
- 39. A. G. Hudetz, *Microcirculation*, 1997, 4, 233-252.
- 40. J. P. Delaney, *Am J Physiol*, 1969, **216**, 1556-1561.
- 41. J. P. W. Baaijens, A. A. Vansteenhoven and J. D. Janssen, *Biorheology*, 1993, **30**, 63-74.
- 42. E. Kucukal, J. A. Little and U. A. Gurkan, *Integr Biol-Uk*, 2018, **10**, 194-206.
- 43. A. M. Forsyth, J. D. Wan, W. D. Ristenpart and H. A. Stone, *Microvascular research*, 2010, **80**, 37-43.
- 44. A. Pantaleo, E. Ferru, G. Giribaldi, F. Mannu, F. Carta, A. Matte, L. De Franceschi and F. Turrini, *Biochem J*, 2009, **418**, 359-367.

- LC-ART-02-2020-000112.R1
- L. Zolla, G. Lupidi, A. Bellelli and G. Amiconi, *Biochimica et biophysica acta*, 1997, **1328**, 273-280.
- 46. L. Zolla, G. Lupidi and G. Amiconi, *Toxicology in vitro : an international journal published in association with BIBRA*, 1994, **8**, 483-490.
- 47. H. C. Mel and T. A. Reed, *Cell Biophys*, 1981, **3**, 233-250.
- 48. K. L. Nuttall, Ann Clin Lab Sci, 2004, **34**, 235-250.
- A. Turhan, L. A. Weiss, N. Mohandas, B. S. Coller and P. S. Frenette, *P Natl Acad Sci USA*, 2002, 99, 3047-3051.
- 50. S. Charache, M. L. Terrin, R. D. Moore, G. J. Dover, F. B. Barton, S. V. Eckert, R. P. McMahon and D. R. Bonds, *The New England journal of medicine*, 1995, **332**, 1317-1322.
- 51. M. H. Steinberg, British journal of haematology, 2005, **129**, 465-481.
- 52. A. R. Chand, H. Xu, L. G. Wells, B. Clair, C. Neunert, A. E. Spellman, L. J. Clay, K. Natrajan and A. Kutlar, *Journal*, 2014.
- 53. O. S. Platt, D. J. Brambilla, W. F. Rosse, P. F. Milner, O. Castro, M. H. Steinberg and P. P. Klug, *The New England journal of medicine*, 1994, **330**, 1639-1644.
- 54. M. Kim, Y. Alapan, A. Adhikari, J. A. Little and U. A. Gurkan, *Microcirculation*, 2017, 24.
- 55. P. Hartmann, M. J. Leiner and M. E. Lippitsch, *Anal Chem*, 1995, **67**, 88-93.
- 56. D. K. Kaul and H. Xue, *Blood*, 1991, 77, 1353-1361.
- 57. J. Kim, H. Lee and S. Shin, *Journal of Cellular Biotechnology*, 2015, 1, 63-79.

- 58. G. Tomaiuolo and S. Guido, J Vasc Res, 2015, 52, 15-15.
- 59. E. Islamzada, K. Matthews, Q. Guo, A. T. Santoso, S. P. Duffy, M. D. Scott and H. Ma, *Lab Chip*, 2019, DOI: 10.1039/c9lc01058k.
- 60. H. E. Judy and N. B. Price, *Journal of the American Medical Association*, 1958, **167**, 563-566.
- 61. E. Kerek, M. Hassanin and E. J. Prenner, *Bba-Biomembranes*, 2018, **1860**, 710-717.
- I. Safeukui, J. M. Correas, V. Brousse, D. Hirt, G. Deplaine, S. Mule, M. Lesurtel, N. Goasguen,
 A. Sauvanet, A. Couvelard, S. Kerneis, H. Khun, I. Vigan-Womas, C. Ottone, T. J. Molina, J. M. Treluyer, O. Mercereau-Puijalon, G. Milon, P. H. David and P. A. Buffet, *Blood*, 2008, 112, 2520-2528.
- R. Suwanarusk, B. M. Cooke, A. M. Dondorp, K. Silamut, J. Sattabongkot, N. J. White and R. Udomsangpetch, *The Journal of infectious diseases*, 2004, 189, 190-194.
- N. Sotirakopoulos, T. Tsitsios, M. Stambolidou, G. Athanasiou, M. Peiou, V. Kokkinou and K. Mavromatidis, *Renal Failure*, 2004, 26, 179-183.
- 65. N. Z. Ertan, S. Bozfakioglu, E. Ugurel, M. Sinan and O. Yalcin, *Plos One*, 2017, **12**, e0171371.
- M. Dessi, A. Noce, P. Bertucci, G. Noce, S. Rizza, A. De Stefano, S. M. di Villahermosa, S. Bernardini, A. De Lorenzo and N. Di Daniele, *Lipids Health Dis*, 2014, 13.
- 67. S. Forman, M. Bischel and P. Hochstein, Ann Intern Med, 1973, 79, 841-843.
- Y. Kikuchi, T. Koyama, Y. Koyama, S. Tozawa, T. Arai, M. Horimoto and Y. Kakiuchi, *Nephron*, 1982, 30, 8-14.

- LC-ART-02-2020-000112.R1
- C. D. Brown, H. S. Ghali, Z. H. Zhao, L. L. Thomas and E. A. Friedman, *Kidney Int*, 2005, 67, 2066-2067.
- P. Zachee, M. A. Boogaerts, R. L. Lins, V. Denbroeck and M. E. Debroe, *Clin Nephrol*, 1986, 25, 315-315.
- 71. F. B. Piel, M. H. Steinberg and D. C. Rees, *The New England journal of medicine*, 2017, 377, 305.
- Y. Man, U. Goreke, E. Kucukal, A. Hill, R. An, S. Liu, A. Bode, A. Solis-Fuentes, L. V. Nayak, J.
 A. Little and U. A. Gurkan, *Blood Cells Mol Dis*, 2020, 83, 102424.
- W. M. Lande, D. L. Andrews, M. R. Clark, N. V. Braham, D. M. Black, S. H. Embury and W. C. Mentzer, *Blood*, 1988, **72**, 2056-2059.
- S. K. Ballas, J. Larner, E. D. Smith, S. Surrey, E. Schwartz and E. F. Rappaport, *Blood*, 1988, 72, 1216-1223.
- E. Nader, M. Grau, R. Fort, B. Collins, G. Cannas, A. Gauthier, K. Walpurgis, C. Martin, W.
 Bloch, S. Poutrel, A. Hot, C. Renoux, M. Thevis, P. Joly, M. Romana, N. Guillot and P. Connes, *Nitric Oxide*, 2018, 81, 28-35.
- 76. A. Marahatta and R. E. Ware, *Blood Cells Mol Dis*, 2017, **67**, 135-142.
- E. Bennett-Guerrero, T. H. Veldman, A. Doctor, M. J. Telen, T. L. Ortel, T. S. Reid, M. A.
 Mulherin, H. Zhu, R. D. Buck, R. M. Califf and T. J. McMahon, *Proceedings of the National Academy of Sciences of the United States of America*, 2007, **104**, 17063-17068.
- S. Henkelman, M. J. Dijkstra-Tiekstra, J. de Wildt-Eggen, R. Graaff, G. Rakhorst and W. van Oeveren, *Transfusion*, 2010, 50, 941-948.

- S. M. Frank, B. Abazyan, M. Ono, C. W. Hogue, D. B. Cohen, D. E. Berkowitz, P. M. Ness and V. M. Barodka, *Anesth Analg*, 2013, **116**, 975-981.
- M. García-Roa, M. Del Carmen Vicente-Ayuso, A. M. Bobes, A. C. Pedraza, A. González-Fernández, M. P. Martín, I. Sáez, J. Seghatchian and L. Gutiérrez, *Blood Transfus*, 2017, 15, 222-231.
- R. E. Domen and G. A. Hoeltge, *Archives of pathology & laboratory medicine*, 2003, **127**, 316-320.
- 82. S. J. Wagner, *Vox sanguinis*, 2005, **88**, 60.
- P. Harmatz, E. Butensky, K. Quirolo, R. Williams, L. Ferrell, T. Moyer, D. Golden, L. Neumayr and E. Vichinsky, *Blood*, 2000, 96, 76-79.
- 84. P. G. Firth, Y. Tsuruta, Y. Kamath, W. H. Dzik, C. S. Ogilvy and R. A. Peterfreund, *Canadian journal of anaesthesia = Journal canadien d'anesthesie*, 2003, **50**, 895-899.
- E. Kucukal, A. Ilich, N. S. Key, J. A. Little and U. A. Gurkan, *American journal of hematology*, 2018, DOI: 10.1002/ajh.25159.
- 86. L. S. Lessin, J. Kurantsinmills and H. B. Weems, *Blood Cells*, 1977, **3**, 241-262.
- 87. M. J. Messer and J. W. Harris, *J Lab Clin Med*, 1970, **76**, 537-&.
- 88. T. Itoh, S. Chien and S. Usami, *Blood*, 1995, **85**, 2245-2253.
- 89. G. B. Nash, C. S. Johnson and H. J. Meiselman, *Blood*, 1986, 67, 110-118.
- 90. L. A. Verduzco and D. G. Nathan, *Blood*, 2009, **114**, 5117-5125.

- LC-ART-02-2020-000112.R1
- M. Kim, Y. Alapan, A. Adhikari, J. A. Little and U. A. Gurkan, *Microcirculation*, 2017, 24, 10.1111/micc.12374.
- C. Yuan, E. Quinn, E. Kucukal, S. Kapoor, U. A. Gurkan and J. A. Little, *Blood Cells Mol Dis*, 2019, **79**, 102350-102350.
- C. Yuan, E. Quinn, E. Kucukal, S. Kapoor, U. A. Gurkan and J. A. Little, *Blood Cells Mol Dis*, 2019, **79**, 102350.
- 94. S. J. Rotz, M. Ann O'riordan, C. Kim, N. Langer, C. Cruz, R. Schilz, C. Lance and J. A. Little, *American journal of hematology*, 2016, **91**, E355-356.
- 95. M. Kim, Y. Alapan, A. Adhikari, J. A. Little and U. A. Gurkan, *Microcirculation*, 2017, 24.
- 96. M. J. Telen, P. Malik and G. M. Vercellotti, *Nat Rev Drug Discov*, 2019, 18, 139-158.
- 97. Z. Romero, M. DeWitt and M. C. Walters, *Expert Opin Biol Ther*, 2018, 18, 1123-1136.
- 98. F. A. Ferrone, Semin Hematol, 2018, 55, 53-59.
- 99. K. Dufu, M. Patel, D. Oksenberg and P. Cabrales, *Clin Hemorheol Microcirc*, 2018, **70**, 95-105.
- J. A. Ribeil, S. Hacein-Bey-Abina, E. Payen, A. Magnani, M. Semeraro, E. Magrin, L. Caccavelli,
 B. Neven, P. Bourget, W. El Nemer, P. Bartolucci, L. Weber, H. Puy, J. F. Meritet, D. Grevent, Y.
 Beuzard, S. Chretien, T. Lefebvre, R. W. Ross, O. Negre, G. Veres, L. Sandler, S. Soni, M. de
 Montalembert, S. Blanche, P. Leboulch and M. Cavazzana, *The New England journal of medicine*, 2017, **376**, 848-855.
- 101. S. H. Orkin and D. E. Bauer, Annu Rev Med, 2019, 70, 257-271.

- G. J. Kato, M. P. Lawrence, L. G. Mendelsohn, R. Saiyed, X. Wang, A. K. Conrey, J. M. Starling,G. Grimes, J. G. Taylor and J. McKew, *Journal*, 2013.
- 103. N. Willie, R. K. Mehlotra, R. E. Howes, T. A. Rakotomanga, S. Ramboarina, A. C. Ratsimbasoa and P. A. Zimmerman, *Am J Trop Med Hyg*, 2018, **98**, 1683-1691.

FIGURES



Figure 1. OcclusionChip design mimics the capillary bed with arteriovenous anastomoses. (A) A schematic representation of the human capillary bed consisting of arterioles, microcapillaries, venules and arteriovenous anastomoses. (B) Macro view of the OcclusionChip with a clinical blood sample at near physiological hematocrit flowing through the microchannel. Scale bar represents a length of 2 mm. (C) OcclusionChip design features nine micropillar arrays with openings ranging from 20 μ m down to 4 μ m, with anastomosis-mimicking side passages that are 60 μ m wide. Inset: scanning electron microscopy images showing the organization of micropillar arrays and side microfluidic anastomosis, with dimensions in micrometers. Scale bars represent a length of 100 μ m and 10 μ m, respectively. (D) Anastomosis side flow takes place around occluded microcapillaries (i.e., 4 μ m array in this panel). Scale bar represents a length of 50 μ m.



Figure 2. Assessment of microvascular occlusion mediated by chemically treated RBCs in the OcclusionChip. (A) A bright field and EGFP overlay image showing microcapillary occlusions induced by poorly deformable RBCs. (B) (Top to bottom) Overview of retained RBCs stiffened by glutaraldehyde with the concentration of 0.02%, 0.04% and 0.08% (w/v) (control not shown) in the microfluidic device. Scale bars represent a length of 500 μ m. Arrows indicate flow direction. Blue dash zones indicate each micropillar array. (C) Schematics of the patterns of the retained RBCs stiffened by graded glutaraldehyde. Profiles of occlusions within the four micropillar arrays generated by RBCs exposed to graded

LC-ART-02-2020-000112.R1

concentrations of **(D)** glutaraldehyde, **(E)** diamide, and **(F)** mercuric ion, and the accordingly computed Occlusion Index generated by RBCs exposed to graded concentrations of **(G)** glutaraldehyde, **(H)** diamide, and **(I)** mercuric ion are shown (N=4 in each group). Error bars represent standard error of the mean.



Figure 3. Versatility of OcclusionChip revealed by the spectrum of Occlusion Indices (OIs) of RBCs from various pathological conditions. These OIs were generated by abnormal RBCs in various conditions including non-specific protein cross-linking (Glutaraldehyde), cytoskeletal-specific stiffening (Diamide), heavy metal toxin (HgCl₂), storage lesion, and circulatory diseases (renal failure or end-stage renal disease, malaria, sickle cell disease: SCD). Stars (*) at the top of the bars indicate statistically significant difference (*: p<0.05; **: p<0.01; *: p<0.001). These results indicate the versatility of OcclusionChip assay for a wide variety of clinical and pathological conditions that impact RBC deformability and function.

LC-ART-02-2020-000112.R1



Figure 4. Assessment of microvascular occlusion mediated by pathologically abnormal RBCs from subjects with SCD in the OcclusionChip. (A) Representative fluorescent image of retained HbSS RBCs across the microfluidic device. Typical morphologies of retained HbSS RBCs within microcapillaries are shown in subsequent insets. Scale bar represents a length of 500 μm, 50 μm, and 10 μm, respectively. Arrow indicates flow direction. (B) Profiles of numbers of microcapillary occlusions induced by normal (HbAA) RBCs and SCD (HbSS) RBCs (5 HbAA and 16 HbSS) in each arrays. (C) The Occlusion Index (OI) of HbSS RBCs is significantly higher compared to that of HbAA RBCs (p=0.001, Mann-Whitney).



Figure 5. Red blood cell mediated microcapillary occlusion *in vitro* and the resultant Occlusion Index (OI) associate with fetal hemoglobin level (HbF%) in subjects with sickle cell disease (SCD). (A) HbF% has been shown to associate with improved clinical outcomes in SCD, where HbF>8.6% is considered as a therapeutic threshold. OI associates with blood HbF level in SCD, where subjects with higher HbF levels (HbF>8.6%) displayed significantly lower OIs (mean \pm SEM = 0.88% \pm 0.10%, N=6), compared with those with lower HbF levels (HbF≤8.6%) (mean \pm SEM = 3.18% \pm 0.34%, N=10, p<0.001, one-way ANOVA). Error bars represent standard error of the mean (SEM). (B) Hydroxyurea therapy is known to improve RBC deformability and to increase HbF% in some but not all patients with SCD. Among the 11 subjects tested who were on Hydroxyurea therapy (circles), 5 subjects presented lower HbF levels (HbF≤8.6%) and higher OIs (red circles), while the other 6 subjects presented higher HbF levels (HbF>8.6%) and lower OIs (black circles). All 5 subjects on Transfusion therapy displayed HbF levels less than 8.6% and higher OIs (red triangles) compared with subjects who were on Hydroxyurea with HbF greater than 8.6%. These results suggest that OI would be useful in assessing the clinical outcomes and treatment efficacy in SCD, along with other important clinical parameters, such as HbF%.



Figure 6. Hypoxia-mediated HbS-carrying RBC sickling and microcapillary occlusion in the OcclusionChip. (A) Macroscale view of the OcclusionChip integrated with the custom designed gas exchange tubing and the gas exchange microchannel. (B) Schematic of axisymmetric cross section of the

gas exchange tubing showing oxygen diffusion. Arrows indicate the flow direction. **(C)** Schematic of the fabrication of the gas exchange microchannel. Cross-sectional view shows the oxygen level stabilization within the microchannel. **(D)** Normalized gray-level intensity measured near the outlet of the microchannel when a solution containing the oxygen-sensitive luminescence probe was flushed at 0 s. **(E)** Identification of hypoxia-mediated cell sickling of HbS-carrying RBCs from two frames of Video S2. **(F)** Scanning electron microscopy images showing hypoxia-enhanced microcapillary occlusion, which confirm that the deformability of HbS-carrying RBCs is mediated by hypoxia. **(G)** The Occlusion Index (OI) of HbS-carrying RBCs increased in response to hypoxia while that of HbAA RBCs remained hypoxia non-responsive. **(H)** Profiles of microcapillary occlusions induced by HbS-carrying RBCs within the microchannel under normoxic or hypoxic conditions. **(I)** A strong positive correlation (PCC=0.7636, p=0.046, N=7) was observed between the OI and the subject-specific HbS levels under hypoxia.

Table of Contents Entry



A gradient of microcapillary networks and microfluidic anastomoses enable standardized quantitative assessment of red blood cell mediated microvascular occlusion.