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Spontaneous Neutrophil Migration Patterns during Sepsis after Major Burns

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Abstract

Finely tuned to respond quickly to infections, neutrophils have amazing abilities to migrate fast and efficiently towards sites of infection and inflammation. Although neutrophils ability to migrate is perturbed in patients after major burns, no correlations have yet been demonstrated between altered migration and higher rate of infections and sepsis in these patients when compared to healthy individuals. To probe if such correlations exist, we designed microfluidic devices to quantify the neutrophil migration phenotype with high precision. Inside these devices, moving neutrophils are confined in channels smaller than the neutrophils and forced to make directional decisions at bifurcations and around posts. We employed these devices to quantify neutrophil migration across 18 independent parameters in 74 blood samples from 13 patients with major burns and 3 healthy subjects. Blinded, retrospective analysis of clinical data and neutrophil migration parameters revealed that neutrophils isolated from blood samples collected during sepsis migrate spontaneously inside the microfluidic channels. The spontaneous neutrophil migration is a unique phenotype, typical for patients with major burns during sepsis and often observed one or two days before the diagnosis of sepsis is confirmed. The spontaneous neutrophil migration phenotype is rare in patients with major burns in the absence of sepsis, and is not encountered in healthy individuals. Our findings warrant further studies of neutrophils and their utility for early diagnosing and monitoring sepsis in patients after major burns.
Introduction

Today, sepsis is the leading cause of death after major burn injury \[1, 2\] and a major cause of death in the intensive care units, with a mortality rate of about 30% \[3\]. Identifying sepsis early is critical, considering that for every 6 hours delay in the diagnosis of sepsis, survival decreases by 10% \[4\]. However, identifying sepsis in burn patients is challenging, because it is masked by the systemic inflammation response syndrome (SIRS), which occurs in almost all patients with major burns \[5\]. Diagnosis may also be delayed by the time required for microbiological confirmation of infection. Although advanced mass spectrometry tools in microbiology labs can accelerate the identification of infections in blood, they still require at least 12–24 hours of blood culture before the bacteria reach levels that can be detected \[6, 7\]. In the absence of sepsis, inappropriate use of antibiotics can have long term consequences for patients by interfering with the normal microbiome, facilitating the development of multidrug resistant bacterial strains, and increasing the cost of hospitalization \[8\]. Contributing to the challenge of diagnosing sepsis is the fact that the pathophysiology of sepsis is not well understood and there is no reliable marker for sepsis. For example, C-reactive protein (CRP) is a marker of the inflammatory response post-burn and recombinant-CRP has been shown to help treating sepsis \[9\]. However, the levels of CRP fail to predict infection or sepsis in patients with major burns \[10\]. Neutrophil CD64 expression can be markedly up-regulated at the onset of bacterial infections. However, recent studies have found that CD64 also increases after major trauma and SIRS \[11\] or sterile insult after major surgery \[12\]. Numerous other molecules have been considered as potential markers, including IL1, IL6, procalcitonin, and reactive oxygen species. However, none has yet been proven effective in the clinic \[13–15\].

One opportunity for detecting sepsis in patients with major burns arises from the study of innate immune responses, particularly those of neutrophils, the major white blood cell population, and first responders to tissue injury. Among the features of the complex neutrophil phenotype, chemotaxis could be an appropriate integrator and sensitive measure for neutrophil function. Neutrophils can integrate a broad range of chemokine and metabolic changes triggered by the burn injury \[16\]. Chemotaxis is altered well before other important neutrophil functions are affected, including phagocytosis \[17\] and the production of reactive oxygen species \[18\], cytokines \[19\] or release of lytic enzymes \[20\]. However, until now, neutrophil chemotaxis changes measured in burn patients have been nonspecific and no features specific to sepsis have been identified. Earlier efforts to characterize neutrophil chemotaxis in sepsis, using traditional cell migration tools, such as the transwell assay \[21\], have been hampered by intrinsic limitations of these technologies, such as the instability of the chemical gradients, and lack of single cell resolution. First employed for studies of cell migration a decade ago \[22\], microfluidic assays enabled very precise control of the shape and stability of chemical gradients during neutrophil chemotaxis \[23–27\]. However, biological issues related to the intrinsic variability of directionality and speed of neutrophils
moving on flat surfaces limited the precision of the motility phenotype measurements in these devices \[28, 29\]. Recently, microfluidic devices that confine the moving cells into micro-channels enabled us to decouple the measurement of speed from directionality and persistence and to independently quantify these parameters with high precision \[30, 31\].

Here, we performed comprehensive measurements of neutrophil migration in blood samples from adult patients after major burns (>20% total body surface area). We monitored the changes in neutrophil migration using microfluidic devices for up to one month after injury, across 18 independent parameters, in 74 blood samples from burn patients and healthy subjects. Blinded, retrospective analysis determined the presence of sepsis in the group of patients in this study. By studying the correlations between neutrophil migration and sepsis, we identified one unanticipated neutrophil migration phenotype that was highly specific for sepsis. We found that spontaneous migration of neutrophils in the absence of a chemoattractant associated with sepsis could begin before sepsis is diagnosed, and ceased after sepsis was effectively treated. These results could have practical implications for the diagnosis and monitoring of sepsis in patients after major burns, as well as warrant further studies to better understand the roles of neutrophils during sepsis.

**Materials and Methods**

**Design and fabrication of the microfluidic devices**

The design of the device enables one to generate chemical gradients in two simple steps performed at the beginning of the experiment, after which the device does not require any user intervention or external syringe pumps to operate. The device is first primed with chemoattractant and then washed with a buffer solution, based on principles described in detail before \[32\]. A gradient forms between an array of sources of chemoattractant and a large buffer reservoir, connected through long channels with small cross section. An array of orthogonal side-channels (6 μm width, 3 μm height), primed with the chemoattractant, serve as source and a central channel (500 μm width, 50 μm height), washed with media, serves as a sink. The array of channels incorporates straight channels for measuring the speed of neutrophil migration, as well as channels with several types of mazes (centered rectangle, off-centered rectangle, and posts) to quantify neutrophil directionality and persistence during migration.

The microfluidic devices were manufactured using standard techniques. Two layers of photoresist (SU8, Microchem, Newton, MA), the first one 3 μm thin and the second one 50 μm thick, were patterned on one silicon wafer by sequentially employing two photolithography masks and processing cycles according to the instructions from the manufacturer. The wafer with patterned photoresist was used as a mold to produce PDMS (Polydimethylsiloxane, Fisher Scientific, Fair Lawn, NJ) parts, which were thenbonded irreversibly to standard glass slides (1 × 3 inches, Fisher).
Blood samples
Blood samples of 3 mL were drawn from patients with large thermal injuries admitted to the Massachusetts General Hospital (MGH). Enrollment criteria included burns covering at least 20% of total body surface area, age between 18 and 81, and we enrolled both male and female, proportionally distributed among ethnic groups. The first sample was obtained within 72 hours after burn injury, and two more samples were drawn at 48 hour intervals afterward (phase 1). Samples were then obtained weekly for up to two months during treatment (phase 2). During a suspected septic event, three additional samples were drawn at 48 hour intervals (phase 3). No samples were collected within 24 hours after an operative procedure.

Ethics Statement
All patient samples were obtained with written informed consent and through procedures approved by the Institutional Review Board at the Massachusetts General Hospital (2008-P-002123).

Neutrophil isolation
Using sterile technique, neutrophils were isolated from whole blood by density gradient separation using HetaSep for 15 min. at room temperature. Neutrophils were then purified using the EasySep Human Neutrophil Kit (STEMCELL Technologies Inc. Vancouver, Canada) following the manufacturer’s protocol. The purity of neutrophils was assessed to be >98% using a Cell Counter. The final aliquot of neutrophils was re-suspended in 20–50 μL of 1 × HBSS +0.2% HSA before loading the neutrophils into the chemotaxis device. Samples were processed within one hour after each blood draw, and were maintained at 37°C.

Neutrophil chemotaxis measurements
Chemokine concentration gradients are maintained for at least 3 hours in the axial direction of the channels by diffusion between the dead-end channels (primed with chemokine at the start of the experiment and functioning as source) and the neutrophil loading channel (washed with buffer when neutrophil suspension is loaded, the and functioning as a sink). Approximately 15 minutes before neutrophils were loaded, the microfluidic device was primed with the chemokine formyl-methionyl-leucyl-phenylalanine (fMLP - 100 nM), leukotriene B4 (LTB4 - 100 nM), or no chemoattractant, serum-free Hank’s Balanced Salt Solution (HBSS) with 0.2% Human serum albumine (HSA). All conditions included extracellular matrix protein fibronectin [100 μM]. The side channels are closed at the distal end, devices are primed with chemoattractant and a linear gradient is formed along the length of the channel when the central well is washed with media. For this, a 1 mL syringe filled with the solution of chemoattractant and fibronectin was connected to one port of the device, and the outlet port was
blocked with a hemostat. By applying pressure to the syringe, the solution was instilled into the device, and the displaced air diffused out through the PDMS. Neutrophils were then infused into the device and allowed to settle in the main channel by clamping the ports of the device. Neutrophil migration in the direction of chemoattractant started immediately, and was recorded using and time-lapse imaging on a Biostation IM at 37°C and 80% humidity. Images of the moving neutrophils are acquired automatically every 2 minutes for up to 2 hours, from at least 30 distinct locations on each microfluidic device (minimum 10 per condition).

The analysis of neutrophil migration was performed by an independent researcher with no information about the status of the patients. All migrating neutrophils were manually tracked in each sample, and velocities were calculated using Image J (NIH). Because the moving neutrophils are mechanically confined in the 6 μm width, 3 μm height channels, neutrophil migration paths are constrained and cell trajectories are uncomplicated to measure.

**Sepsis criteria**

For the purpose of this study, sepsis was established blindly and retrospectively, by a physician not involved in the care of the patients in this study. Sepsis was defined based on the presence simultaneously of documented infection and SIRS, following the guidelines from the American College of Chest Physicians (ACCP), Society of Critical Care Medicine (SCCM) and the European Society of Intensive Care Medicine (ESICM) [5]. SIRS was established based on two or more of the following criteria: temperature above 38°C or below 36°C, tachycardia (heart rate above 90 beats per minute), tachypnea (respiratory rate faster than 20 breaths per minute), and white blood count above $12 \times 10^9/L$ or below $4 \times 10^9/L$ or more than 10% immature (band) forms. Infections were established based on documented positive cultures of blood samples, intra-operative deep wound specimens, deep tracheal aspirate, or urine samples. Patients who had positively documented cultures without signs of SIRS were not considered septic. Septic patients were considered no longer septic when cultures converted into sterility after antimicrobial treatment. Antimicrobial treatment was not considered as an associated factor for diagnosis of infection due to frequent empirical antimicrobial regimens for common SIRS conditions in burn patients. Thus, our criteria for sepsis were stricter than the current guidelines that consider suspected infections [5].

**Statistical Analysis**

STATA software (StataCorp., College Station, TX) was utilized to apply a logistic model to the entirety of our neutrophil functional data set and determine whether a patient’s septic status could be predicted by their neutrophil phenotypic function. Exploratory analysis determined 3 variables, all quantifying the spontaneous migration in the absence of chemoattractant, to significantly
correlate to a septic diagnosis in the clinic. These three variables included: number of neutrophils migrating in the absence of chemoattractant (NMC_N), oscillatory migration in the absence of chemoattractant (OM_N) and the total length of migration in the channels in the absence of chemoattractant (TL_N). A neutrophil activation score in the absence of chemoattractant (NAS_N) was defined as NMC_N*(TL_N + OM_N). A logistic model (Generalized Estimating Equations (GEE)) was used to determine statistical significance of the NAS_N variable as a clinical biomarker for sepsis. The GEE took into account the patient’s repeated NAS_N measurements over time and did not require normality of the predictor variable. To test if the mean of the activation state during infection is different than that of the infection-free state, we used a random intercept linear mixed effects model (repeated measures of ANOVA). To compare the parameters of neutrophil migration between healthy controls and burn patients we used a t-test and difference s were considered significant for p<0.05.

Results

Microfluidic platform for probing neutrophil migration phenotype

The goal of this study was to measure changes of the neutrophil migration phenotype in patients with major burns and sepsis and to identify specific changes associated exclusively with sepsis. The study was enabled by microfluidic devices designed for precision measurements of the neutrophil migration phenotype. Two critical features contributing to the precision of the measurements are the confinement of moving neutrophils in channels with cross-sections smaller than the neutrophil size, and the placement of bifurcations and post-obstacles along the channels. In the straight section of the channels, we can measure neutrophil migration speed and persistence with high precision. At the posts and bifurcations, we force the moving neutrophils to make binary directional decisions, which are easy to quantify and compare.

Neutrophils separated from patients’ blood (Fig. 1A – steps 1 & 2) are loaded inside the device and their migration responses tested in 3 distinct conditions which include serum-free Hank’s Balanced Salt Solution (HBSS) and gradients of formyl-methionyl-leucyl-phenylalanine (fMLP) and leukotriene B4 (LTB4) (Fig. 1A – step 3). The migration of neutrophils, recorded using automated microscopy, is analyzed for 6 parameters in each of the three conditions (Fig. 1A – step 4). While the neutrophils from healthy volunteers are able to migrate efficiently around the posts in the channel (S1 Movie), patient neutrophils often get stuck on the post, lose directionality, and exit the device (Fig. 1A – step 5, S2 Movie). All healthy neutrophils are able to pick the shortest path and follow the steeper gradient in the offset rectangle, whereas a large fraction of patient neutrophils loop inside the rectangle, unable to correctly follow the direction of the chemoattractant gradient. We then compared the neutrophil migration parameters to identify markers and scores to distinguish between burn patients with and without sepsis (Fig. 1A – steps 6 & 7). The analysis of neutrophil
migration phenotype can be completed in less than 4 hours after the blood samples become available. Blood samples were collected every two days for the first 5 days in the hospital (phase 1) and once a week for three more weeks (phase 2). Additional samples were collected every two days for one week whenever the clinical team requested blood cultures from burn patients with fever and leukocytosis, and/or thrombocytosis (phase 3 - Fig. 1B).
Neutrophil migration phenotype in patients with major burns

We measured eighteen independent parameters of neutrophil migration in 74 samples from 13 patients with major burns, during treatment at Massachusetts General Hospital (Table 1) and 3 healthy volunteers. We validated first the device and compared it with previous studies [32], by measuring neutrophil migration in three healthy volunteers. The measured differences in migration speed between patients with major burns and healthy subjects are in agreement with previous observations [16, 32–39]. We also measured differences in the parameters of directionality and persistence between the neutrophils from patients with major burns and healthy controls. These differences between neutrophils from burn patients and healthy controls are summarized in Table 2 and a heat map (Fig. 2B). Neutrophils from patients with major burns migrate in smaller numbers (MC), at slower average velocity (v), and are less directional (D). Fewer neutrophils reach the end of the channels in response to the two chemoattractants tested (T, TL). For the majority of samples, these parameters were on average at least two standard deviations below healthy donors.

Neutrophil spontaneous migration is characteristic for sepsis

We compared the migration parameters of neutrophils from burn patients based on the status of the patient at the time of sample collection. Of the 71 blood samples from burn patients analyzed, 21 were collected from patients during sepsis, 29 during SIRS, and 21 from patients with no signs of SIRS. We observed that a significant numbers of neutrophils from patients during sepsis advanced inside the channels in the absence of a chemoattractant (S3 Movie). To better characterize this unique phenotype, we recorded the number of neutrophil migrating spontaneously (MCN) and the fraction of neutrophils moving spontaneously and crossing back and forth the longitudinal-middle of the channels (OMN – Fig. 2C). However, on a two dimensional plot of oscillatory migration and migration count, the neutrophils samples from patients during sepsis often overlap with those in the absence of sepsis (Fig. 3A). Similarly, significant overlap between sepsis and non-sepsis samples was present when we analyzed the average distance that neutrophils advanced inside the channels length (TLN) vs. the oscillatory migration. Other parameters of neutrophil migration in the absence of chemoattractant did not correlate with sepsis, alone or in combinations (Fig. 3B). Moreover, no linear combination of the 18 parameters appeared to be effective in differentiating between sepsis and no-sepsis.

To better characterize the unique phenotype of spontaneous migration of neutrophils, we took into account the observation that these neutrophils either migrate long distances inside the channels or display oscillatory migration patterns. Thus, we defined a nonlinear neutrophil activation score in the absence of chemoattractant (NASN), which captures both behaviors in the same equation: NASN = MCN * (TLN + OMN). We analyzed the changes in NASN during time after major burns and observed that the score remained low for the duration of hospital stay in patients with no sepsis (Fig. 3C, N=28 samples from 6 patients).
In contrast, in patients with sepsis, we observed transient increases of the NAS\textsubscript{N} score that corresponded with the time of sepsis (Fig. 3D, N=43 samples from 7 patients). The increases of NAS\textsubscript{N} were usually transient during the first week after burn, and appeared to be more prolonged later in the disease (after week 3). Moreover, the score usually decreased after the start of antibiotic treatment and, in some of the patients that developed sepsis, the score increased days before sepsis was diagnosed (Fig. 3D).

A logistic model of generalized estimating equations (GEE) indicated that the NAS\textsubscript{N} score is statistically significant correlated with sepsis and differentiates between septic patients and those with systemic inflammatory response syndrome (SIRS - P=0.038) and no complications (P=0.021, Fig. 4A). The score increased significantly at least 3 days before clinical diagnostic of sepsis (P=0.036). Moreover, the NAS\textsubscript{N} decreased 3-fold after effective antibiotic treatment was started, compared to the first draw after sepsis diagnosis (Fig. 4B). A random intercept linear mixed effects model (repeated measures of ANOVA) revealed that the mean of the NAS\textsubscript{N} during infection was different than that of the infection-free state (p=0.077).

To further probe the potential utility of analyzing the spontaneous neutrophil migration patterns that act as a biomarker for sepsis in burn patients, we analyzed the receiver operating characteristic (ROC) for various cut-off levels of NAS\textsubscript{N}. Using the NAS\textsubscript{N} on the day of the sepsis the area under the curve was 0.74, at the threshold of good performance. Moreover, the performance of the NAS\textsubscript{N} could be improved further by considering the average of two successive samples. The area under the curve increases to 0.80, the best sensitivity to 80%, and specificity to 77% (Fig. 4C). This analysis suggests that the NAS\textsubscript{N} has the potential, when monitored frequently, to become useful adjunct to current clinical protocols to identify and monitor sepsis in patients after major burns.

**Discussion**

The microfluidic platform presented in this study enabled us to characterize the migration of human neutrophils from burn patients, at single-cell resolution and

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total number of patients</td>
<td>13</td>
</tr>
<tr>
<td>Total blood samples</td>
<td>71</td>
</tr>
<tr>
<td>Average size burn (TBSA)</td>
<td>39.3%</td>
</tr>
<tr>
<td>Patients with sepsis during hospital treatment</td>
<td>7</td>
</tr>
<tr>
<td>Patients with SIRS during hospital treatment</td>
<td>12</td>
</tr>
<tr>
<td>Patients with infection</td>
<td>7</td>
</tr>
<tr>
<td>Patients treated with antibiotics</td>
<td>13</td>
</tr>
<tr>
<td>Mortality</td>
<td>1</td>
</tr>
</tbody>
</table>

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with high precision, across 18 parameters. In addition to confirming that significant differences in neutrophil migration phenotype exist between burn patients and healthy [40], we uncovered a new neutrophil migration phenotype, the spontaneous migration in the absence of chemoattractants, which is highly correlated with the occurrence of sepsis in the burn patients. The spontaneous migration of neutrophils is a unique phenotype, which has not been observed before. The spontaneous migration takes place in the absence of any chemoattractants and is thus distinct from chemokinesis, which requires continuous exposure to chemoattractant. The spontaneous migration shows long

### Table 2. Summary of neutrophil phenotypic migration parameters to an array of chemoattractants (no chemoattractant, fMLP, and LTB₄).

<table>
<thead>
<tr>
<th>Neutrophil Chemotaxis Parameter</th>
<th>Abbr.</th>
<th>Definition</th>
<th>Chemo-attractant</th>
<th>Healthy Donor (Avg. +/- Std. Dev.)</th>
<th>Burn Patient (Avg. +/- Std. Dev.)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Migration Count</td>
<td>MC</td>
<td>The total number of neutrophils that migrated from the central cell loading channel into the side migration channel.</td>
<td>fMLP</td>
<td>110.5 +/- 25.8</td>
<td>62.7 +/- 86.6</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>139 +/- 49.6</td>
<td>73.0 +/- 76.4</td>
</tr>
<tr>
<td>Average Velocity</td>
<td>V</td>
<td>The average velocity of neutrophil chemotaxis in straight channels or the linear portion of the channels past the mazes [μm/min]</td>
<td>fMLP</td>
<td>17.5 +/- 2.1</td>
<td>9.5 +/- 7.6</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>23.5 +/- 5.3 •</td>
<td>10.0 +/- 7.5 •</td>
</tr>
<tr>
<td>Directionality</td>
<td>D</td>
<td>The fraction of neutrophils that navigate through the mazes along the shortest route, without turning back at the bifurcations or being delayed at posts.</td>
<td>fMLP</td>
<td>0.78 +/- 0.11</td>
<td>0.48 +/- 0.35</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>0.87 +/- 0.08</td>
<td>0.63 +/- 0.35</td>
</tr>
<tr>
<td>Average Migration Length</td>
<td>TL</td>
<td>The average distance that neutrophils migrate along the straight channel [μm].</td>
<td>fMLP</td>
<td>425.6 +/- 45.1 •</td>
<td>29.3 +/- 10.6 •</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>469.2 +/- 22.4 •</td>
<td>31.1 +/- 12.9 •</td>
</tr>
<tr>
<td>Oscillatory Migration</td>
<td>OM</td>
<td>The percentage of neutrophils moving inside the channels that cross-back and forth over the mid-line (x=250 μm) of the straight channel.</td>
<td>fMLP</td>
<td>2.3 +/- 5.7</td>
<td>6.8 +/- 10.4</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>4.2 +/- 10.2</td>
<td>2.5 +/- 5.4</td>
</tr>
<tr>
<td>Through</td>
<td>T</td>
<td>The percentage of neutrophils that migrate persistently the entire length of the straight channels in the field of view (x=500 μm).</td>
<td>fMLP</td>
<td>88 +/- 6.3 •</td>
<td>14.0 +/- 21.4 •</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>LTB₄</td>
<td>93 +/- 8 •</td>
<td>31.2 +/- 33.9 •</td>
</tr>
</tbody>
</table>

Averages +/- standard deviations are shown for all parameters for both healthy donors and burn patients. • indicates average parameters that are significantly different between healthy donors and burn patients, (p<0.05)

doi:10.1371/journal.pone.0114509.t002
strides of persistent migration, however, it is different from chemotaxis because these strides occur in the absence of any chemical gradient to trigger the persistent migration. Spontaneous migration is also unlike retrotaxis, when neutrophils move persistently and independently against chemical gradients, but only following chemotaxis [41].

The spontaneous neutrophil migration phenotype could not be captured by existing chemotaxis assays. Conventional assays, such as the transwell assay [21], do not have single cell resolution. They only measure average migration, and will miss entirely a small population of cells with spontaneous and oscillatory

Fig. 2. Neutrophil migration parameters in burn patients. (A) Heat-map showing the results of 18 neutrophil migratory phenotype measurements in 74 blood samples from 13 burn patients and three healthy volunteers. Specific measurements are explained in detail in Table 2, including averages for each parameter. Each parameter was measured in three conditions: in the absence of chemoattractant, and in the presence of fMLP and LTB4 chemoattractant gradients. Bright green illustrates 2 S.D. below healthy donor averages, light green illustrates 1 S.D. below healthy donor averages, bright red illustrates 2 S.D. above healthy donor averages, and light red illustrates 1 S.D. above healthy donor averages. Below the heat map, a color coded bar represents the status of the blood donor at the time of the draw. Red illustrates sepsis, green illustrates SIRS, and black illustrates no SIRS status. The day post burn is indicated above the bar, the patient identifier is presented below the bar. Healthy donors, burn patients with sepsis, and burn patients without sepsis are grouped by the accolades. (B) Kymograph showing an example of oscillatory migration of a neutrophil from a septic burn-patient, in the absence of chemoattractant. The time interval between successive frames is 2 minutes. Vertical scale bar is 100 μm.

doi:10.1371/journal.pone.0114509.g002
migration phenotype. Other assays, recently reviewed in [25, 42] lack the necessary precision to characterize complex neutrophil migration patterns like those during spontaneous cell migration. Most of microfluidic devices where neutrophils migrate on one-dimensional surfaces [22, 26, 29], could not decouple the oscillatory migration from changes in direction, and thus do not have the necessary precision to distinguish the new phenotype. Moreover, the new microfluidic platform is robust, easy to use, and requires no external pumps or other accessories besides the microscope.

Overall, sensitivity and specificity values for our assay are only slightly better than other markers for sepsis in burn patients [43, 44]. However, it is important to
note that, enabled by technology, we could observe the changes in neutrophil migration phenotype, up to two days before sepsis. This unique capability of our assay, to predict the occurrence of sepsis, may eventually enable early interventions or preventive treatments in the context of burns. It also points to a potential critical role that neutrophils play in the earliest stages leading to sepsis. The spontaneous neutrophil migration phenotype may be an integrator of multiple inputs (cytokines, chemokines, leukotrienes, extracellular matrix context, and microbial products), which allows neutrophils to prioritize these signals for their contextual importance [45] during tissue injury [46] and infections [47]. Until now, such stimuli could only be measured individually e.g. CD64 [11],
IL-6 [48], serum macrophage migration inhibitory factor [49], interleukin-1 receptors [50], and neutrophil gelatinase-associated lipocalin [51] and integrated into comprehensive scores afterwards by artificial models. Using the neutrophil phenotype as a marker has the advantage of natural adaptation to changes and compensation for variability in the levels of signals among individuals.

Additional benefits from the use of microfluidic technologies to monitor neutrophil function may eventually emerge, besides the better precision for identifying sepsis. Correcting the functionality of neutrophils in patients could become as an important objective among sepsis prevention measures. Supporting this possibility, one recent study showed that treatment using lipid mediators of inflammation resolution could help restore normal neutrophil migration phenotype and improve the survival after burns in an animal model [39]. A reliable infection marker in the context of burns and treatments to correct neutrophil functionality could discourage the preventive use of antibiotics and reduce the chances of selecting for antibiotic resistant bacterial strains, which are common in burn units [52]. Further studies of neutrophil phenotype in larger populations of patients with sepsis after major burns could lead to better understanding of the roles that neutrophils play before and during sepsis and facilitate the design of new tools for the diagnosis, monitoring, and prevention of sepsis in patients with major burns.

Supporting Information

S1 Movie. Neutrophils from healthy volunteers migrate fast and efficiently around the posts in the channel.
doi:10.1371/journal.pone.0114509.s001 (AVI)

S2 Movie. Neutrophils from patients with major burn injury display defective migration. Patient neutrophils often get stuck on the posts, lose directionality, and reverse the direction of migration to exit the migration channel.
doi:10.1371/journal.pone.0114509.s002 (AVI)

S3 Movie. A significant numbers of neutrophils from patients during sepsis advanced inside the channels in the absence of a chemoattractant.
doi:10.1371/journal.pone.0114509.s003 (AVI)

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Author Contributions

Conceived and designed the experiments: CNJ RGT SPF DI. Performed the experiments: CNJ MM LD AA. Analyzed the data: CNJ MM LD AA AI BAH HSW
RGT SPF DI. Contributed reagents/materials/analysis tools: CNJ LD AI BAH HSW. Wrote the paper: CNJ HSW RGT DI.

References


