A Third Measure-Metastable State in the Dynamics of Spontaneous Shape Change in Healthy Human’s White Cells

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Abstract

Human polymorphonuclear leucocytes, PMN, are highly motile cells with average 12-15 μm diameters and prominent, loboid nuclei. They are produced in the bone marrow, are essential for host defense, and are the most populous of white blood cell types. PMN also participate in acute and chronic inflammatory processes, in the regulation of the immune response, in angiogenesis, and interact with tumors. To accommodate these varied functions, their behavior is adaptive, but still definable in terms of a set of behavioral states. PMN morphodynamics have generally involved a non-equilibrium stationary, spheroid Idling state that transitions to an activated, ellipsoid translocating state in response to chemical signals. These two behavioral shape-states, spheroid and ellipsoid, are generally recognized as making up the vocabulary of a healthy PMN. A third, “random” state has occasionally been reported as associated with disease states. I have observed this third, Treadmilling state, in PMN from healthy subjects, the cells demonstrating metastable dynamical behaviors known to anticipate phase transitions in mathematical, physical, and biological systems. For this study, human PMN were microscopically imaged and analyzed as single living cells. I used a microscope with a novel high aperture, cardioid annular condenser with better than 100 nanometer resolution of simultaneous, mixed dark field and intrinsic fluorescent images to record shape changes in 189 living PMNs. Relative radial roundness, R(t), served as a computable order parameter. Comparison of R(t) series of 10 cells in the Idling and 10 in the Treadmilling state reveals the robustness of the “random” appearing Treadmilling state, and the emergence of behaviors observed in the neighborhood of global state transitions, including increased correlation length and variance (divergence), sudden jumps, mixed phases, bimodality, power spectral scaling and temporal slowing. Wavelet transformation of an R(t) series of an Idling to Treadmilling state change, demonstrated behaviors concomitant with the observed transition.

Introduction

Polymorphonuclear neutrophil granulocytes (PMNs) are the body’s most abundant class of white blood cells. At circulating cell levels of \( \sim 10^{11} \), PMNs make up about 62% of all human white cells [1]. An idling, non-activated neutrophil circulates in the blood for 8 to 12 hours before undergoing apoptosis. When stimulated by inflammatory cytokine signals released by endothelial, mast cells and/or macrophages, PMNs can survive for 24 to 48 hours [2–4].

Once activated, idling PMNs exploit shape change dynamics as they tether to the endothelium of post-capillary venules, making and breaking bonds while rolling along the venule’s endothelial bed of selectins. Following concentration gradients of inflammatory ligands, they wriggle through the vessel wall and migrate to the site of the initiating inflammation. The motions accompanying this migration exploit the PMN’s cytoskeletal, actin polymerization-depolymerization cycle that configures the dynamics of shape change and translation [5,6].

The behavioral dynamics of two of the shape changing metastable states and their associated translational motions are well established. PMNs manifest these metastable shape-motional states in vivo and in vitro. They are: (1) The circular-spherical, Idling state manifesting standing waves and fast and fine random fluctuations in the leucocyte’s apron edge; (2) The activated, ellipsoidal, polarized migrating state, with almost exclusively positive gradient-directed lamelopodia and filopodia formation [7,8].

The complexity of the scenario described above and the necessity of as many as 100 distinct protein/protein interactions to coordinate the actin cytoskeletal apparatus alone, prompts both a phenomenological approach and the consideration of a potentially larger set of shape-motional states which may be transitional, non-stationary and not easily quantified.

In addition to the metastable spherical-round, Idling and elliptical-migratory state, I have found a statistically prominent, third, metastable metastable state in the PMNs from the fresh blood of healthy human subjects. This non-translating, high amplitude, shape changing state has been previously observed and interpreted as a functionally disordered manifestation of immunological and hematological pathology [9–14]. The goals of these studies are to identify, quantitatively characterize and discriminate this third

Citation: Selz KA (2011) A Third Measure-Metastable State in the Dynamics of Spontaneous Shape Change in Healthy Human’s White Cells. PLoS Comput Biol 7(4): e1001117. doi:10.1371/journal.pcbi.1001117

Editor: Michael Shlesinger, Physical Sciences Division, Office of Naval Research, United States of America

Received July 9, 2010; Accepted March 4, 2011; Published April 7, 2011

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Funding: The Fetzer Foundation supported this work. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The author has declared that no competing interests exist.

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shape-motional PMN state from the other two. I call this third dynamical state, Treadmilling, the word used to also describe a key part of the underlying F.G-actin dynamics [15]. I have observed the Treadmilling state in 12 of the 18 healthy, adult volunteers contributing blood for this study.

A Statistical Concern

We remind ourselves that there are neurobiological limits on the both the resolution afforded by empirically meaningful partitions [16], and the minimal length of a neurobiologically defensible time series [17]. In these studies I must be concerned with how long the cells being observed maintain their anatomical and functional integrity and within that viability limit, how many observations I can make without the vacuous artifice of over-sampling.

The issue more generally is the selection of an appropriate sample size of an intrinsically non-stationary system. Counter-intuitively, it has been shown that under certain conditions of limited information, repeated too-short sample lengths come to be computationally superior globally [18].

In the past I have dealt with this problem by studying repeated time series derived measures yielding populations of not necessarily convergent estimates [19] with, nonetheless, distributional properties of the measures, such that I can estimate each measure’s central and higher moments, range of variation and statistical differences between measures in comparisons of varying observed system state [20]. This is the approach to be taken here.

Methods

Relative Radial Roundness, $R(t)$, as an Order Parameter

It is difficult to find a global quantitative measure on the dynamics of emergent phenomenon with the nice properties of additivity, continuity and differentiability [21,22]. Such a measure has been called an order parameter, named for its use in tracking a system’s dynamics through transitions in the system’s degree of order. In gas-liquid transitions, the order parameter is density [23], while in ferromagnet transitions, for instance, it is net magnetization [23,24]. Perhaps the best known example of an order parameter is relative phase, the Landau-Ginsberg order parameter [23,24], used in the phenomenological description of thermodynamic and superconducting transitions [25].

I have examined the autonomous, time-dependent shape changes of individual Idling and Treadmilling human white cells as real, spatially extended dynamical systems. I use a global measure on the PMN’s relative radial roundness, $R(t)$, as the order parameter. $R(t) = r_1/r_2$ is computed as the ratio of the radius of the cell assumed to be an ideal circle, $r_1 = p/2\pi$ in which $p$ is the sum of the pixels outlining the cell’s perimeter and $r_2 = (A/p)^{0.5}$ using the sum of the pixels within the cell’s silhouette as the cell’s area, $A$. $R(t) = r_1/r_2 = (p/2\pi)(A/p)^{0.5}$, computed at each time step. If both $r_1$ and $r_2$ were derived from an abstract, idealized circle, $R(t) = r_1/r_2 = 1.0$, such that $log[R(t) = r_1/r_2] = 0$, the characteristic lower limit of a generic order parameter. Deviations from this reference characterize changes in state [21-23,25]. My use of the global order parameter, $R(t)$, contrasts with a previous use of an averaged local measure, the power spectral transformation of a sequence of angles resulting from the piece wise linear segmentation of the cell’s circumference [26]. The use of $R(t)$ more closely resembles a differential geometric pattern map [27].

Experimental Procedures

One hundred and eighty-nine PMNs from fifty-three peripheral blood samples were collected from 18 healthy adult volunteers, aged 26 to 72. The blood samples were allowed to sediment gravitationally for 40 minutes at room temperature. A population of PMNs (and other white cells and platelets) were removed from theuffy coat by micropipette and, along with associated plasma, placed within a 12 mm ring painted on a glass slide, forming a ~20–25 μm deep well, compared to the average 5.7 μm vertical space between a plain slide and its cover slip [28]. The 5.7 μm gap of standard slides and coverslips is considerably smaller than the average diameter of PMNs, leading to some mechanical compression of the cell contributing to their activation, and allowing the cell to move along the slide substrate and cover slip simultaneously [28]. The slides used in this study do not suffer from these deficits.

PMN autonomous motions were observed using an Olympus BX41 microscope fitted with CytoViva dark field and fluorescent optical illumination systems, which includes a unique, high-aperture, cardioid annular condenser (www.scitech.com.au). The CytoViva condenser makes it possible to visualize objects of below 100 nm in diameter in real time, and with the cellular samples in an unfixed, living, active state [29,30]. Because PMNs were treated gently, avoiding perturbations of column separation and elution, it became possible to reliably study a PMN continuously for 30 to 60 minutes before the onset of granular clumping, membrane blebbing and other signs of nascent apoptosis [31]. Data collection continued until ten each Idling and Treadmilling cells met the conditions for inclusion in the study. Specifically, only cells that maintained healthy, one state behavior, and did not have contact with any extracellular objects for the entire 30 min recording period were retained for analysis.

Idling PMNs are characterized by their near spheroid shape (quasi-circular in two dimensions). In this state, the microscopically visible autonomous motions are limited to standing waves on the cell surface and low amplitude fluctuations of the cell’s microvillus border. In contrast, Treadmilling PMNs demonstrate large and irregular changes in cell shape. Multiple transient, often simultaneously appearing, pseudopodia and lamelopodia emerge from the cell surface, oriented apparently randomly and without significant movement of the PMNs center of mass. The cell’s movement was less than 1.5 times the maximum diameter of the cell over the typical ~30 minute recording sessions. Figure 1
portrays binary color coded, characteristic silhouettes of the round Idling, Treadmilling and elliptically polarized/translocating shape-states of PMNs. Only videos micrographs of mature, segmented neutrophils that did not have contact with any cells or extracellular entities and remained visibly healthy over the thirty minutes of observation, in addition to manifesting stable state behavior were retained for further analysis.

Images were collected every 2 sec for 30 min using an Optronics Microfire 1200×1600 CCD array camera [30] resulting in a 900 point \( R(t) \) series of high resolution images per PMN. The slowness of the cell shape changing motions led to the finding that more frequent sampling within the time limit of cellular integrity was obviously redundant. In the geometric computations, each primary image was used to produce two binary, \( 0,1 \), digital daughter images: an area map of \( A \), and perimeter map of \( p \). The 0,1 coding of the pixels of the two daughter images were converted into binary arrays and used compute the \( R(t) \) time series.

Computation of the Measures on \( R(t) \)

In light of the above discussion of biological constraints on sample length and the intrinsic non-stationarity of the PMNs shape motion series, statistical distributions of often individually non-convergent measures made on each of the cells, serve as the basis for comparisons of Idling and Treadmilling states. Statistical evaluations are then made on populations of possibly incomplete measures, not on the raw observations. Rules of thumb concerning sample length requirements for any particular measure [32] though easily attainable in physical and computational systems, often ignore the intrinsic series limits and non-stationarity of real, behaving, biological systems. In addition to the use of the distribution of each particular kind of measure, I study an aggregate of several, often incomplete measures, each reflecting different aspects of the shape-motional dynamics of PMNs.

On the \( R(t) \) of each cell, I study: (1) The central moments of the \( R(t) \) distribution, the mean \( S_1 \) and standard deviation, \( S_2 \), as well as

### Table 1. Measure averages for cells in each state group.

| Measure  | Idling (n = 10) | Treadmilling (n = 10) | \( t(df); \rho \) |
|----------|-----------------|-----------------------|------------------|
| Mean     | 2.935           | 2.932                 | \( t_{18} = 0.004; \rho < 0.4983 \) |
| SD       | 0.1023          | 0.3882                | \( t_{18} = 4.816; \rho < 0.0001 \) |
| Skew     | 0.2993          | 0.7273                | \( t_{18} = 2.419; \rho < 0.0132 \) |
| Kurtosis | 0.7229          | 1.165                 | \( t_{18} = 0.652; \rho = 0.2612 \) |
| \( \lambda_1 \) | 0.6242       | 0.5066                | \( t_{18} = 2.592; \rho < 0.0090 \) |
| \( \alpha \) | -0.4561        | -1.0290               | \( t_{18} = 10.600; \rho < 0.0001 \) |

Table 1 reports the results of measures made on the \( R(t) \) series of ten Idling PMN and ten Treadmilling PMN, all of which had remained healthy and in a single behavioral state for the 30 min recording period and made no contact with extracellular bodies during that time.

doi:10.1371/journal.pcbi.1001117.t001
the skewness, $S_3$, indicating the asymmetry of the density distribution of $R(t)$, estimated using the third moment, $m_3$, divided by the cubed root of the variance squared, $S_3 = m_3 / \text{var}^{3/2}$. The kurtosis, $S_4$, of $R(t)$ is computed using the relation, $S_4 = m_4 / \text{variance}^2 - 3$ [33]; (2) An estimate of the $R(t)$'s orbital divergence, its sensitivity to initial conditions, in a three dimensional embedding space, was computed using a generic algorithm for the leading Lyapounov exponent, $\Lambda_1$ [34]; (3) Differences in a hierarchical scaling property of $R(t)$, by computing the scaling exponent $\alpha$ derived from its power (frequency)spectrum, as the slope of the middle third of the linear best fit of the log power-log frequency relation [35]; (4) An example of the time dependence of the scaling of $R(t)$ was estimated from a Morlet continuous wavelet transformation using standard algorithms [36–38].

Visualizing Phase Space Behavior of $R(t)$

To visualize the phase space behavior I used relatively denoised, three dimensional Broomhead-King, B/K, eigenfunction, $\Psi$, embedding of the $R(t)$'s. To do this, I computed and plotted $\Psi_1$, $\Psi_2$, and $\Psi_3$ with respect to each other [39,40]. Each $R(t)$ series

Figure 2. Shows the eigenfunction space embedding (see text) for four representative idling cells (left column) and four Treadmilling cells (right column).

doi:10.1371/journal.pcbi.1001117.g002
generated an M-lagged data matrix on which an MxM Hermitean autocovariance, \( C_M \), matrix was computed, with \( M = 8 \), a typical correlation decay interval. \( C_M \) was then decomposed into its eigenvalue-ordered eigenvectors. The eigenvectors associated with the three largest eigenvalues were each composed with the original \( R(t) \) series to form B/K eigenfunctions \( \Psi_1, \Psi_2, \) and \( \Psi_3 \). These formed the axes of the B/K eigenspace reconstruction. Because \( R(t) \) behavior attributable to the lower, excluded, eigenvectors accounts for the trivial, “noise” component of the variance, the resulting eigenfunction space embedding (each successive point being a triple) is relatively denoised compared with the more commonly used phase delay space construction [41].

Another graphical representation of the orbital behavior of \( R(t) \) is its two dimensional, \( i = \tau_1, j = \tau_2 \), Recurrence Plot, \( RP[R(t)]_{ij} \), introduced by Eckmann [42]. Graphical representations of \( RP[R(t)]_{ij} \) are two dimensional lattices, each point computed as \( RP_{ij} = \Theta(\varepsilon \| \mathbf{x}_i - \mathbf{x}_j \|) \), \( i,j = 1...N \), where \( \mathbf{x}_i \) in \( \mathbb{R}^2 \) represents the location of the orbit in phase space at time \( i \). \( \varepsilon \) is the static distance defining the “closeness” threshold, and \( \Theta \) is the Heavyside function. The resulting binary series, each point \( \varepsilon \)-close or not to the previous value, is coded in black and white. Here, a standard time delay three dimensional embedding was used, with delay \( \tau = 1 \) [43]. If \( \mathbf{x}_i \) falls within the distance \( \varepsilon \) of \( \mathbf{x}_j \), \( \mathbf{x}_i \) is considered to be a recurrence of \( \mathbf{x}_j \), otherwise not. Clustering in \( RP_{ij} \) has been used to discriminate among three characteristic patterns of intermittency [44].

**Results**

There were highly significant differences between the measures \( S_2, S_3, \Lambda_1, \) and \( \alpha \) that were made on the \( R(t) \) series of the PMNs in the *Idling* versus the *Treadmilling* state, see Table 1. No significant differences were found between the two distributions of \( S_1 \) or \( S_4 \).

The qualitative differences in the shape-motional patterns implied by the statistically significant differences in the measures in Table 1 are consistent with behavior that was observed microscopically in the two pre-polarized states: (1) The small, stochastically wavy border fluctuations in cell shape of the generic *Idling* PMNs; (2) A range of large, simultaneously multiscale motions in cell shape variations of \( R(t) \) in the *Treadmilling* state. For examples, compared with *Idling*, the increase in asymmetric amplitude in *Treadmilling* is reflected in increases in \( S_2 \) and \( S_3 \), and the increase in shape-motional order in *Treadmilling* is seen in the statistically significant decrease in \( \Lambda_1 \), the leading Lyapounov index of expansive, orbital mixing [45]. The larger, smoother, more correlated shape motions of the *Treadmilling* state are seen in statistically significant increases in \( \alpha \) in the *Treadmilling* versus *Idling* states. Without a significant difference in the means of \( R(t) \), the variational measures make the discrimination between *Idling* and *Treadmilling* states.

Consistent with the differences in behavior described by direct observation and the aggregate of measures (see Table 1), Figure 2 portrays the previously described \( \{ \Psi_1, \Psi_2, \Psi_3 \} \) B/K eigenfunctions embedding of four representative *Idling* cells (left column) and four *Treadmilling* cells (right column). The phase portraits of the *Idling* cells reflect symmetric, small, random fluctuations around a near stationary state. *Treadmilling* cells manifest larger, more irregular, asymmetric phase space motions which occupy almost an order of magnitude larger volume than that by the *Idling* state.

Another geometric, graphical treatment of the cell’s shape motional behavior is displayed in Figure 3. We see the recurrence...
plots, $R[t]_{i,j}$ of the four representative PMNs in the *Idling* state (top row) and four in the *Treadmilling* state (bottom row), in which $\varepsilon = 1$ for all plots. The $R[t]_{i,j}$ of the *Idling* cells demonstrate more homogeneous temporal distributions of returns typical of more random data with shorter correlation lengths/relaxation times. The square patches of only lightly increased density overlaid on the more uniform surround are consistent with both the visualized small amplitude oscillations in $R(t)$ in the *Idling* state and with the statistical results reported in Table 1. The $R[t]_{i,j}$ of cells in the *Treadmilling* state are, as expected, less homogeneously distributed, manifesting clustering in the return times across multiple times scales, as well as apparent discontinuous changes in their phase space patterns. For example, short interval "bursting" interleaved with low amplitude, long interval behavior is seen in the *Treadmilling* cells' $R[t]_{i,j}$. *Treadmilling* PMNs $R[t]_{i,j}$ portraits are consistent with recurrence patterns of intermittency [44,46].

Four of the seven order parameter measures demonstrated statistically significant differences between the *Idling* and *Treadmilling* PMNs, Table 1. While observing and recording the real-time behavior of 189 PMNs, I witnessed many cells transitioning from one state to another among my three defined behavioral regimes. Data series including such transitions were plagued by the same complications as were the single state series (e.g., cell-cell interactions, apoptotic behavior) in addition to too short times in one or more behavioral state to allow any analysis. I was finally

Figure 4. Shows a “close-up” of a PMN during a phase transition. The upper panel shows the $R(t)$ series, while the lower panel depicts the continuous 1D Morlet wavelet moduli for that time interval (x-axis). Scale of the basis function increases up the y-axis. The colormap passes through the visible spectrum; blues representing low Morlet moduli amplitudes to high valued reds. The cell is initially *Idling*, and begins *Treadmilling* at data point $t = 283$. Note that anticipatory slow high amplitude fluctuations begin to appear at about point $t = 198$. doi:10.1371/journal.pcbi.1001117.g004
able to make sufficient observations portraying a single PMN shape motion transformation in real time. Figure 4 is a Morlet wavelet graph, in continuous time along the x-axis, and scale (~wavelength) along the Y axis. Figure 4 contributes evidence for a continuous transition in shape motion state, here from Idling to Treadmilling. Table 2 lists measures before and after this single cell transition. Note that the direction and approximate magnitude of change resemble those of the population of statistically significant values in Table 1.

Discussion

There are established physiological mechanisms and behavior that are consistent with both our qualitative microscopic observations and quantitative aggregate measure descriptions. PMNs are known to oscillate on multiple time and space scales, from 7 sec, 70 sec, and 250 sec membrane potential fluctuations [47] and 25 sec calcium flux oscillations [48], to the ~8 sec bound/unbound actin oscillations [49], to 21.6 sec and 230 sec glycolytic cycles producing NAD(P)H oscillations [47], and 10 sec bound/unbound actin oscillations [49], to 21.6 sec and 230 sec pericellular proteolysis fluctuations [48], among many others. The $R(t)$ series in this study evidenced scaling, board hand power spectra with multiple resonances [50]. It is likely some reported modes contribute to the cell shape fluctuations directly and others contribute to the emergence of other dynamical patterns. The slowest Fourier mode in $S_n [R(t)]$ of the Idling state had an average 8.457 minutes oscillation, whereas that of the slowest $S_n [R(t)]$ of the Treadmilling state averaged a 4.201 minutes oscillation. It is interesting that these characteristic times correspond roughly to the results of studies of the characteristic remodeling times composed of actin filament diffusion, polymerization and then turnover coordinated with cellular migratory motions [51,52]. It appears that the transition from Idling into the intermittent Treadmilling regime occurs as the Idling state loses some of its dynamical structural stability, and its shape motion scenario becomes driven by several quasiperiodic, multi-periodic metabolic and physiological cellular oscillator mechanisms [53,54].

As listed in Table 1 and Table 2, a comparison of Idling with Treadmilling PMNs reveals significantly different $R(t)$ order parameter dynamics. Projected to a two dimensional plane (Figure 1), one sees an associated difference in the underlying planar geometry, with the Idling PMNs manifesting one centroids in their circularity, and the Treadmilling PMNs with two point defined, barycentric ellipses.

Many characteristics of the changes in measures in the distinct single state observations and in the computable, real-time transition from Idling to Treadmilling suggest the typical signs of a phase transition [21–23,25]. These included: (1) Increasing amplitude of $R(t)$ variability seen in the $S_2$ and $S_3$ of the cell shape fluctuations; (2) Decreased leading $\Lambda_1$ becoming less positive in the direction of zero, shadowing the leading eigenvalue of the unknown underlying partial differential equation; (3) An increase in the log-log power spectral scaling index, $\alpha$, reflecting a “less white” spectral pattern of $R(t)$ fluctuations, also consistent with slowing; (4) The Morlet wavelet transformation of a continuous time $R(t)$, evidenced anticipatory, high amplitude slowing and a mixed phase regimes in the neighborhood of a real-time PMN shape fluctuation transition. The eigenfunction space embedding of the sequence of triples, $\{P_1, P_2, P_3\}$, demonstrated directly the space-time morphogenetic transformation undergone by $R(t)$ in the Treadmilling state with reference to that of the Idling cell state. Recurrence plots, $R_{ij}$ depicted increased phase space clustering consistent with the more hierarchical, intermittent dynamics of the Treadmilling PMNs in contrast with the more randomly distributed and metrically transitive space of the Idling $R_{ij}$. It should be noted that the action spaces of less uniform intermittency and those of more uniform transitivity reflect common metastable alternatives in the dynamics of some biological sciences [40].

Finally, I have spent hundreds of hours microscopically tracking 109 individual PMN cells in the hopes of answering these questions about state and state transitions. While only one such transition was recorded with sufficient observations in both the Idling and Treadmilling states to allow statistical analyses, many transitions were observed. I have seen Idling cells transition to Treadmilling, and Treadmilling cells ball up and Idle (although with slightly ragged apsorns). I have also observed numerous instances of Idling cells polarizing and Translocating until they reach some point at which point they Idle again. The only transitions that were not observed were from the Treadmilling to the polarized, single lamelopod, Translocating state or vice versa. In either case the cells ball-up briefly into an Idling appearance before changing again. See Table 3.

Acknowledgments

I wish to thank Prof. Arnold Mandell and Dr. Paul Gailey for their helpful comments and suggestions.

Author Contributions

Conceived and designed the experiments: KAS. Performed the experiments: KAS. Analyzed the data: KAS. Contributed reagents/materials/analysis tools: KAS. Wrote the paper: KAS. Designed and wrote computer programs used in data transformation and analysis: KAS.

Table 2. Measures made on a neutrophil before and after idling to treadmill state change.

|                  | Idling               | Treadmilling          |
|------------------|----------------------|-----------------------|
| Mean             | 2.0969               | 2.1614                |
| SD               | 0.0348               | 0.1620                |
| Skew             | -0.1942              | 2.0076                |
| Kurtosis         | 2.8552               | 8.0831                |
| $\lambda_1$      | 0.6520               | 0.498                 |
| $\alpha$         | -0.5681              | 1.0486                |

Table 2 reports the results of the same measures listed in Table 1, this time made on the $R(t)$ series of a single PMN first in an Idling state and then in a Treadmilling state. This PMN also remained healthy for the 30 min recording period and made no contact with extracellular bodies during that time. doi:10.1371/journal.pcbi.1001117.t002

Table 3. Observed cell state transitions.

| From       | To       | Idling | Treadmilling | Translocating |
|------------|----------|--------|--------------|--------------|
| Idling     | X        | X      | X            |              |
| Treadmilling| X        | X      |              |              |
| Translocating| X      |        | X            |              |

doi:10.1371/journal.pcbi.1001117.t003
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