

Article

Investigation of *Synechocystis* sp. CPCC 534 Motility during Different Stages of the Growth Period in Active Fluids

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Abstract: The motility behavior of suspended microorganisms plays an essential role in the properties of active fluids. Despite the important progress in our understanding of microorganisms' motility in recent years, there are still several open questions about the dynamics of cell motility in active suspensions. Of special interest is the relationship between cell motility and age. In this study, cyanobacterium *Synechocystis* sp. CPCC 534 was used as the model microorganism, and the cell trajectories were tracked for 78 days during the cell growth period. Results showed that the length of cell trajectories had substantially increased from the exponential growth phase to the stationary phase and had declined at the end of the stationary phase. Similar trends were observed for the cells' mean squared displacement (MSD), the time-dependent diffusion coefficient of cell suspensions, and the cell displacement probability density function (PDF). These results suggest that the cellular age of microorganisms has a significant effect on various metrics of cell motility and, therefore, can impact the transport properties of active suspensions.

Keywords: microorganism; cell trajectory; mean squared displacement (MSD); time-dependent diffusion coefficient $D(t)$; displacement probability density function (PDF)



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1. Introduction

Suspensions of photosynthetic microorganisms are often described as “living” or “active” fluids. Active fluids contain a collection of cells, or active particles, suspended in a fluid medium, where the cells can convert chemical nutrients into mechanical energy to drive the flow. In an environment with uniform nutrient distribution, cells move in a random-walk pattern, with high-motility “run” periods separated by low-motility “tumble” periods [1,2]. However, in the presence of gradients in nutrient or light distribution, cells move preferentially in the direction opposite to the gradient (i.e., from low to high availability of nutrients/light). These processes are called chemotaxis and phototaxis for movements driven by nutrients and light, respectively [1,3,4].

Fluid flow, cell motility, microbe biokinetics, and radiative transport are among the important parameters in designing and optimizing photobioreactors used for cell growth in active fluids [5–8]. Among these factors, motility is one of the major ones in providing favorable conditions for microorganisms' nutrient uptake and light harvesting and, therefore, enhancing biomass growth [9–11]. Moreover, motility plays a vital role in the formation of biofilms. In the initial period of biofilm formation, bacteria proceed toward and adhere to wet solid surfaces. This adhesion allows for bacterial surface colonization and creates

protective biofilms. It is essential to understand the motility of the microorganisms to identify the adhesion rate and the subsequent colonization process [12–15].

In a study by Shoemith [16], a simple quantitative method was applied to measure the motility of *Pseudomonas viscosa*, *Bacillus brevis*, and *Escherichia coli* by counting the number of bacteria passing through a small aperture during a certain period of time. The results revealed that bacterial motility is proportional to the concentration and the mean velocity of cells in the suspension. In another study by Kim [17], the impact of motility on the diffusivity of two bacteria: *Pseudomonas aeruginosa*, a motile cell, and *Klebsiella pneumoniae*, a non-motile cell, was investigated through capillary tubes. It was found that the rate of diffusion of motile cells was 1000 times faster, and the diffusion coefficient was 2.3 times higher compared to non-motile bacteria. Arora et al. [18] carried out experiments in an agitated bioreactor to examine the motility behavior of *E. coli* as a function of its growth dynamics. They observed that the microorganisms' motility was reduced by increasing the number of cells in the culture. This was concluded to be a result of a reduction in the "running" phase of the motility pattern. In a recent study by Fadlallah et al. [19], the motility behavior of active fluids (suspensions of *Chlamydomonas reinhardtii* and *Synechocystis* sp.) was studied under different intensities of hydrodynamic shear stress for 15 days. Their results revealed that hydrodynamic shear stress had a considerable effect on the motility of *C. reinhardtii* during the exponential growth period and enhanced the cellular swimming velocity of microorganism cells. On the other hand, *Synechocystis* did not exhibit any specific trend during the growth cycle and showed higher resistance to shear stress. This could be explained by the lack of flagella in *Synechocystis* and/or its small size in comparison to *C. reinhardtii* [19].

In bacterial cultures, once the cells have adapted themselves to the new environment, they move from the lag phase (where they are largely dormant) into an exponential growth phase. The exponential phase continues until the linear growth phase is reached, wherein the cell growth is reduced due to shading or nutrient depletion. At the end of the linear growth phase, bacteria enter the stationary phase in which the number of cell divisions balances with cell death, and the population of microorganisms remains constant in time. As nutrients deplete and waste materials inside the medium rise, the microorganism cells lose the ability to perform metabolic functions, and then the number of dead cells continues to increase and eventually exceeds that of live cells (decline/death phase) [20,21].

To the best of the authors' knowledge, most previous studies have emphasized the motility of various microorganisms for a short time period during the cell's exponential growth phase [19,22]. In this study, the motility of *Synechocystis* sp. CPCC 534 was used as a model organism to study the impacts of cell age on motility over longer periods of time (i.e., up to 78 days), going far beyond the exponential and stationary growth phases. For this purpose, the cell transport parameters such as cell velocity, mean squared displacement (MSD), time-dependent diffusion coefficient $D(t)$, and displacement probability density function (PDF) were calculated [23]. To accomplish this, bacteria trajectories were extracted from the video footage recorded on an inverted video microscope. The average swimming velocity over a given time period was then calculated and correlated with the age of bacteria obtained from the growth curve [19]. These findings have practical implications for biomass and sustainable resource engineering, particularly with regard to designing bioreactors for mobile microorganisms, facilitating the movement of microorganisms in photobioreactors, and regulating the development of biofilms [19,23].

The model microorganism used in this study, *Synechocystis* sp., is a strain of unicellular, non-nitrogen-fixing freshwater cyanobacteria that acquires its energy for movement through photosynthesis [3,24,25]. It is one of the most prevalent microorganisms for studying the fundamental phenomena in bacterial flows for two important reasons: its availability and capability to have both phototrophic growths by oxygenic photosynthesis during light periods and heterotrophic growth by glycolysis and oxidative phosphorylation during dark periods [26–28]. Such bacteria are naturally motile, and they can move toward favorable environmental conditions (such as light, nutrients, and temperature) or move

away from unfavorable ones by using a variety of motility machinery components such as pili [18,29,30].

2. Materials and Methods

2.1. Culture Conditions

Synechocystis sp. CPCC 534 wild type was obtained from the Canadian Phycological Culture Center (Waterloo, ON, Canada). The stock culture was maintained and grown in BG11 medium [31] at 20 ± 1 °C. Suspensions were exposed to a photon flux of $50 \pm 5 \mu\text{mole}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$ with a light cycle of 12/12 h. The initial experimental concentration of *Synechocystis* sp. in the media was adjusted to 1×10^7 cells.mL⁻¹. To provide a homogeneous environment for the cell growth and to prevent cell sedimentation inside the photobioreactor, a magnetic stirrer (VWR Canada) with stirring bars of 7.9 mm diameter and 19.8 mm length at 360 rpm rotation speed was employed. To control environmental conditions, the photobioreactors were placed in a fully controlled incubator (PHCbi) [24,32].

2.2. Experimental Protocol

Experiments were carried out in triplicate and were performed in 125 mL Pyrex flask photobioreactors. A Spectronic 200E spectrophotometer was used to measure the optical density (OD) of *Synechocystis* sp. CPCC 534 suspensions daily at 750 nm wavelength as a proxy for cell growth. The wavelength of maximal absorption for chlorophyll-a was found to be at 670 nm and for chlorophyll-b at 652 nm. To calculate the reference values, the intensity of the transmitted light at 750 nm was measured. The absorption at 750 nm results only from chlorophyll-free structures in the cells. A calibration curve was used to calculate the number of *Synechocystis* sp. cells per milliliter in a sample solution. Therefore, a dilution series of *Synechocystis* sp. CPCC 534, cultivated at the same environmental growth conditions, were prepared, and the optical density and the cell density (number of cells.mL⁻¹) were measured using the Spectronic 200E spectrophotometer (Thermo Fisher Scientific, Waltham, CA, USA) and a Hauser Scientific hemocytometer (Thermo Fisher Scientific, Waltham, CA, USA), respectively [24,32].

2.3. Calculation of *Synechocystis* sp. Growth Rate (r) and Doubling Time (k)

The growth rate (r) of the *Synechocystis* sp. cells was estimated through the logarithmic growth phase by Equation (1) [33].

$$r = \frac{\ln N_t - \ln N_0}{\Delta t} \quad (1)$$

where N_0 is the number of cells at the beginning of an observation period, N_t is the cell population at time t , and Δt is the duration of the observation period ($t_t - t_0$).

The growth rate (r) is used to evaluate the number of cell divisions per day, known as doubling per day (k), by Equation (2) [33].

$$k = \frac{r}{\ln 2} = \frac{r}{0.6931} \quad (2)$$

2.4. Video Microscopy and Image Acquisition

An aliquot of 40 μL of suspension sample was placed in the cavity of a concave slide, covered with a glass coverslip, and finally sealed with grease to form a micro slide. To observe and record cell displacement in the closed micro slide, it was placed with its glass coverslip side on the stage of an inverted optical Olympus IX83 video microscope equipped with a 40X magnification lens and a CCD camera. The bacteria settled on the flat side of the micro slide after about 30 min. Then the cells' movement in the sample was recorded in "vsi" format for 50 min at a rate of one frame per second. The resolution of the CCD camera was 325 nm per pixel [1,23].

2.5. Analysis of Cell Dynamics

The recorded video footage was post-processed with Fiji ImageJ software (1.53c, Wayne Rasband) to acquire binary images. The binary video footage in “tif” format was then used for particle tracking [34] using MATLAB Version: 9.11.0.2207237 (R2021b) (Natick, MA, USA). Cell tracking was carried out in three steps: first, all cells’ x and y positions were extracted from the recorded video footage, frame by frame, to determine cell trajectories. Then, cell motility was estimated from the average cell velocity by eliminating the non-motile cells (or other debris) in the calculations. Finally, the mean squared displacement (MSD), diffusion coefficient, and displacement probability density function (PDF) were calculated to evaluate the dynamics of *Synechocystis* sp. CPCC 534 [35].

2.6. Statistical Analysis

For the statistical analysis, the “statsmodels” package in Python was used with 95% confidence intervals, and $p < 0.05$ was evaluated as significant [36].

3. Results and Discussion

3.1. Calibration Curve

The calibration curve was established by plotting the OD versus the number of *Synechocystis* sp. CPCC 534 cells, as is shown in Figure 1. The concentration of *Synechocystis* sp. CPCC 534 in all subsequent experiments was determined using the equation derived from the calibration curve (Figure 1) [24].

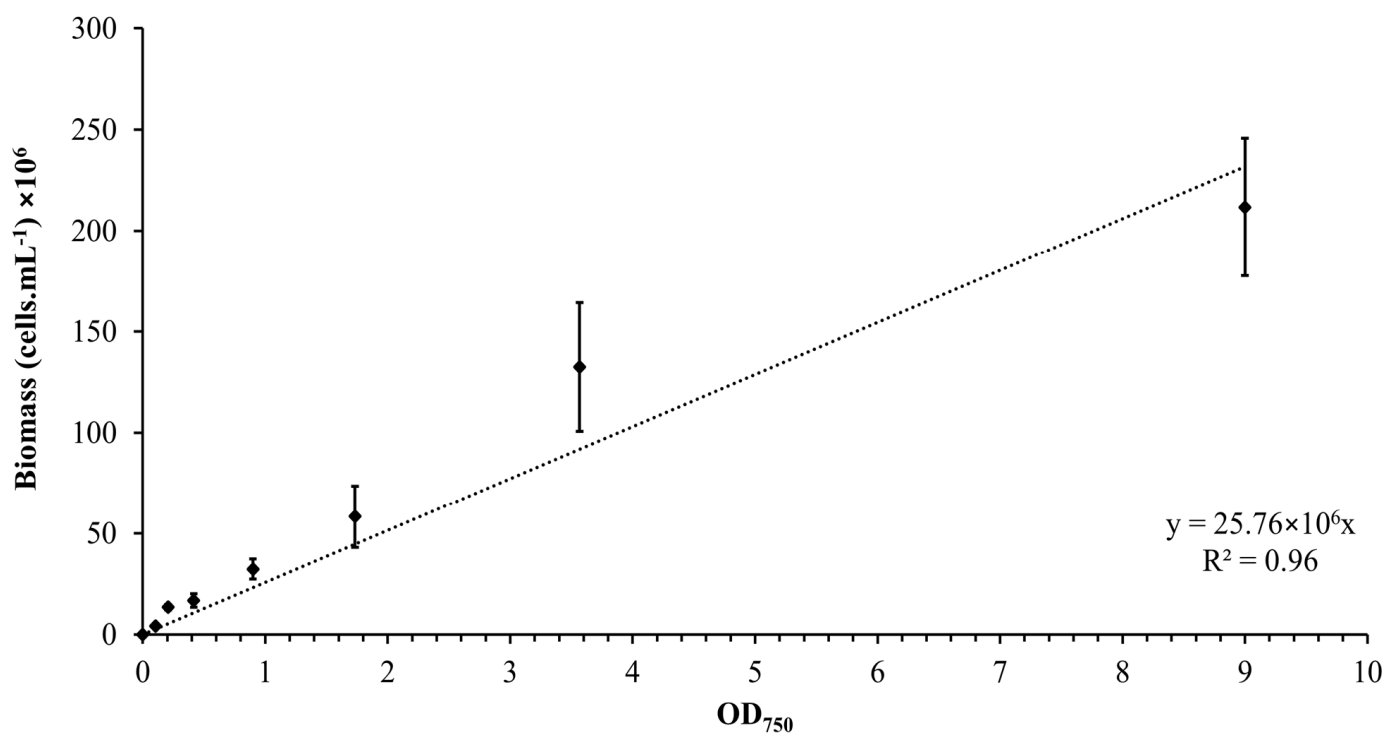


Figure 1. Calibration curve relating the OD₇₅₀ to the biomass concentration for *Synechocystis* sp. CPCC 534 suspensions ($n = 3 \pm \text{SD}$).

3.2. Growth Curve

The growth curve, Figure 2, represents the variation of live *Synechocystis* sp. CPCC 534 cells number during a 78-day growth period. In the present culture, *Synechocystis* sp. cells immediately moved to the exponential (log) growth phase and reached the linear phase after eight days. The transition from the exponential to linear phase is evident from the slope change in the growth curve [35].

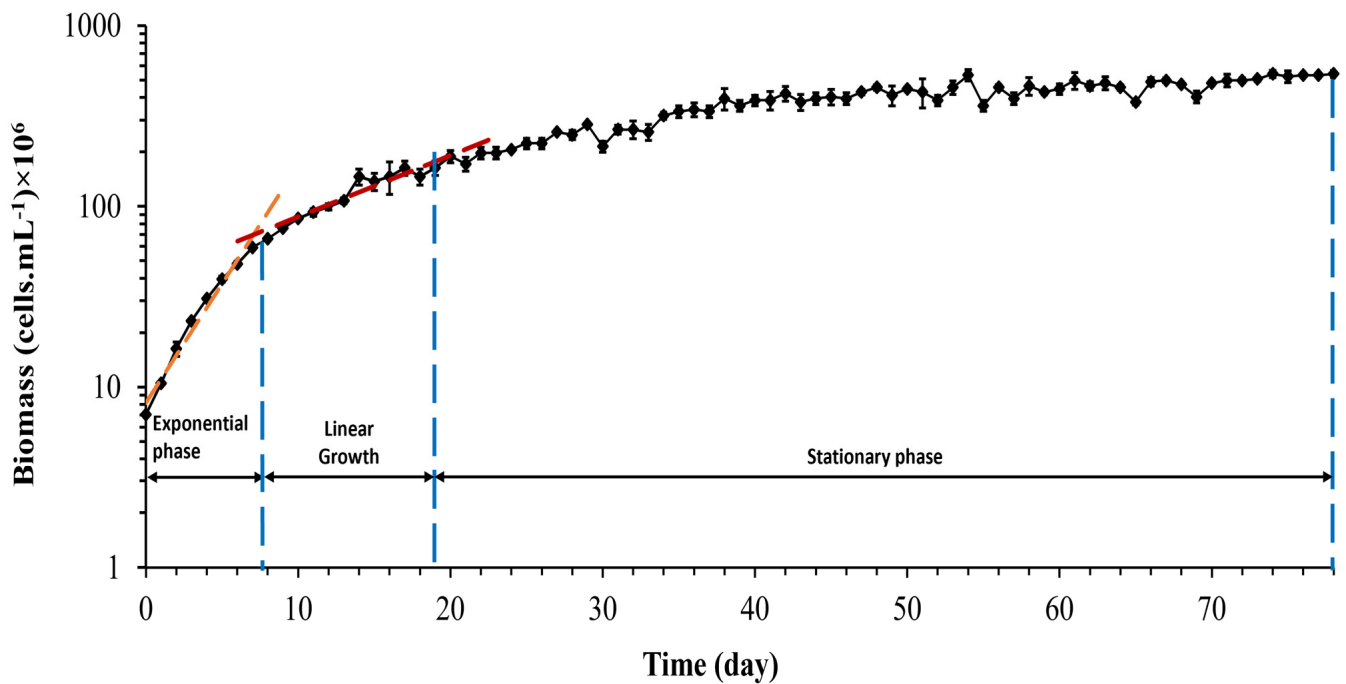


Figure 2. Growth curve of *Synechocystis* sp. CPCC 534 ($n = 3 \pm \text{SD}$). The blue dashed lines show the end of each growth phase in this experiment.

The growth rate and doubling per day calculated during the exponential growth phase were $0.3 \pm 0.01 \text{ d}^{-1}$ and $0.43 \pm 0.015 \text{ d}^{-1}$, respectively. The growth rate equation for both the exponential phase and linear phase, respectively, are:

$$\ln(C_{biomass}) = \ln(8 \times 10^6) + 0.3d \quad (3)$$

$$\ln(C_{biomass}) = \ln(4 \times 10^7) + 0.08d \quad (4)$$

where $C_{biomass}$ is the biomass concentration, and d is the time in days.

3.3. Cell Motion

For cell tracking, the concentration of microorganisms in the suspension was set to $10^7 \text{ cells.mL}^{-1}$ and video microscope images were recorded for 50 min. Figures 3–5 show the trajectories of the population of *Synechocystis* sp. CPCC 534 at (Figure 3) the beginning (on the first day) of the exponential growth phase, (Figure 4) the end of the exponential phase, and (Figure 5) the end of the stationary phase (at day 1, day 17, and day 78, respectively). The duration of all trajectories displayed in Figures 3–5 is 50 min. The results show the length of cell trajectories had increased noticeably during the exponential phase compared to the beginning of the exponential growth phase and the end of the stationary phase. In Figures 3 and 5, the cell trajectories mainly consist of “tumbling” movement and extend around 10 pixels in space, while in Figure 4, cells perform more “run” movements, and the trajectories extend over 80 pixels in space. This observation implies that cell motility improved with the cells’ age. The decline of cell motility at the end of the stationary phase could be due to nutrient depletion in the PBR and/or the fact that most cells had reached the end of their cell life cycle, Figure 5 [5,32].

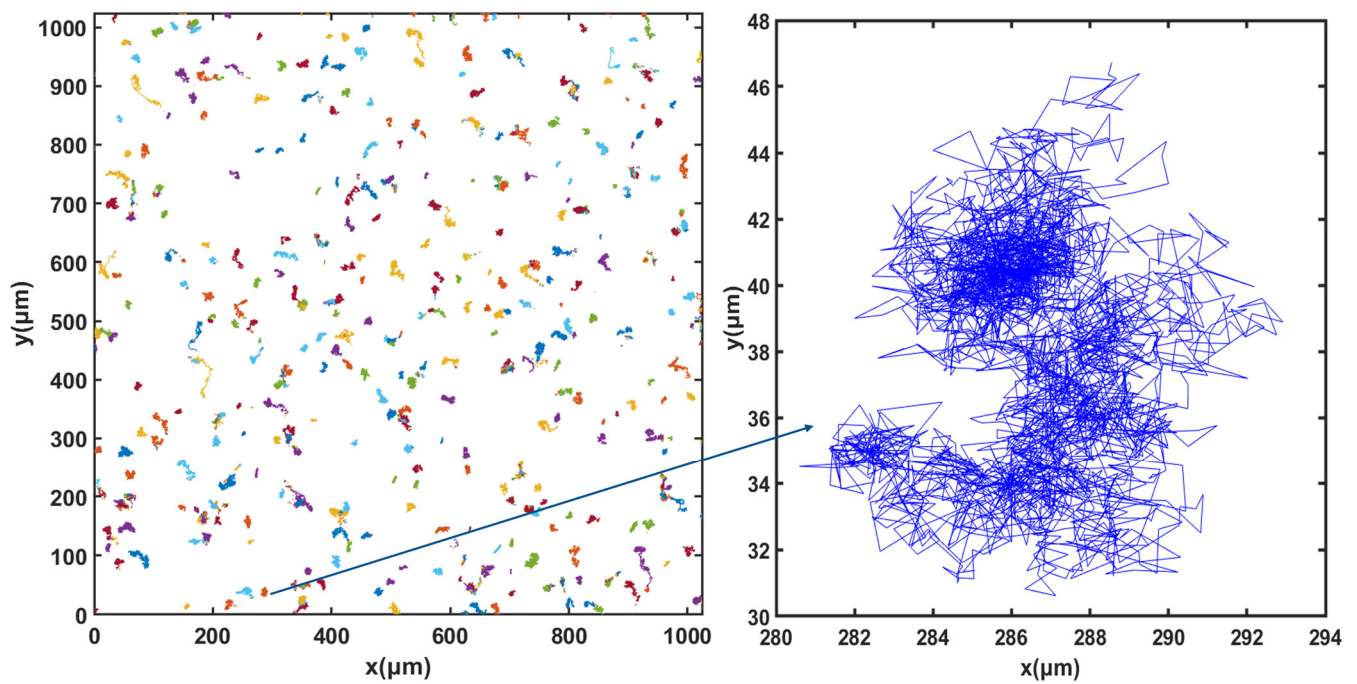


Figure 3. Extracted trajectories of *Synechocystis* cells from the video microscope images, recorded over 50 min, for the first day of the exponential phase, day 1.

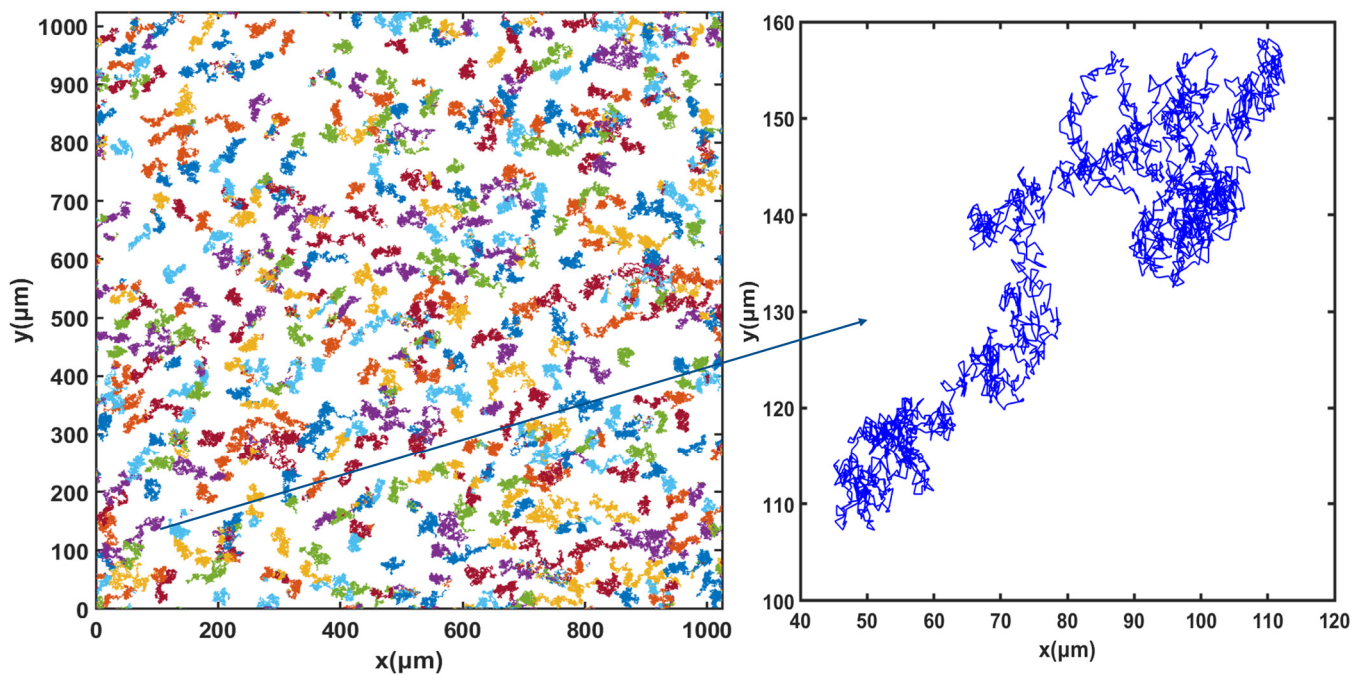


Figure 4. Extracted trajectories of *Synechocystis* cells from the video microscope images, recorded over 50 min, for the end of the exponential phase, day 17.

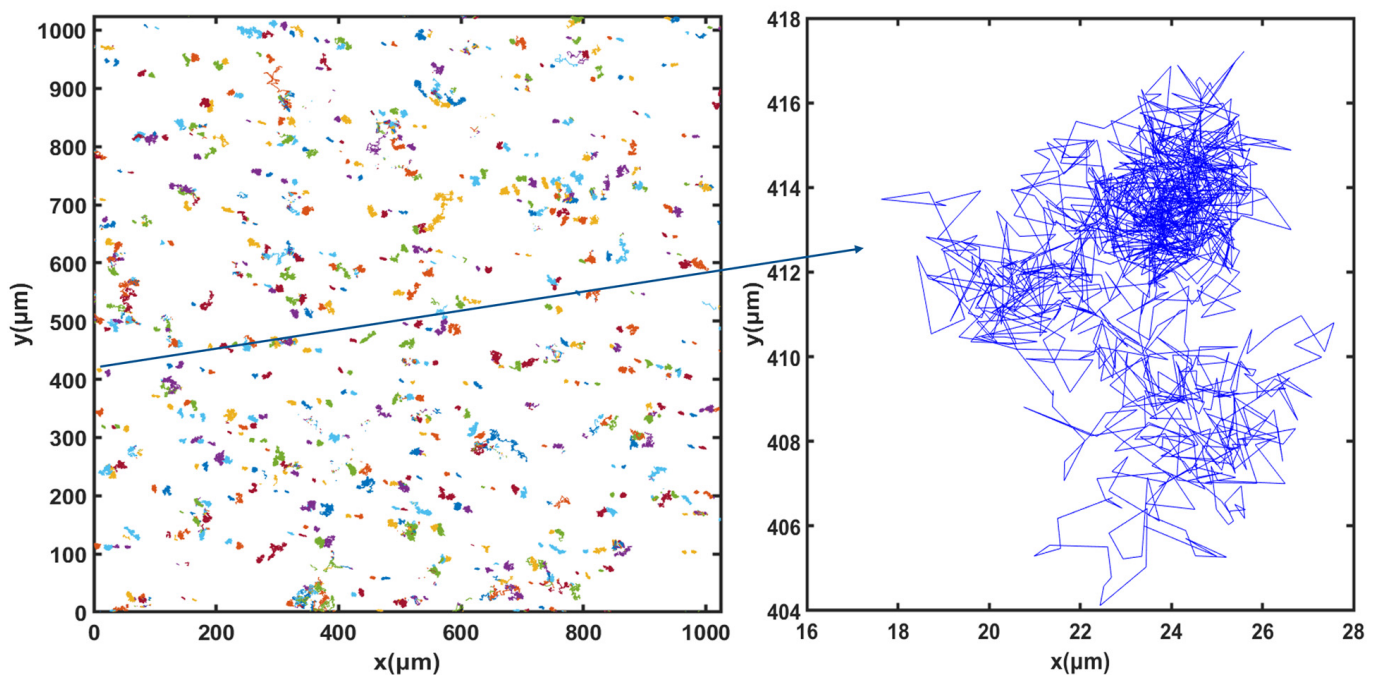


Figure 5. Extracted trajectories of *Synechocystis* cells from the video microscope images, recorded over 50 min, for the end of the stationary phase, day 78.

3.4. Evolution of Cell Motility during *Synechocystis* Growth

The instantaneous velocity of a single cell was calculated as the displacement between two consecutive frames multiplied by the acquisition framerate (one frame per second). The instantaneous velocity for each cell was then averaged over the experimental time period to determine the average cell velocity, V_p . Finally, the ensemble average velocity was calculated as follows:

$$V_m = \frac{1}{n_p} \sum_{p=1}^{n_p} V_p \quad (5)$$

where n_p is the number of motile *Synechocystis* cells present in the recorded video [19]. The measurement time for each experimental run was 50 min, and cell concentrations for all experiments were constant (10^7 cells.mL⁻¹). In Figure 6, the average velocity of *Synechocystis* sp. cell is plotted as a function of time for 78 days of observation. The growth curve corresponding to the same observation period is also superposed on the average velocity curve. It is observed that the cells' average velocity increases with time during the exponential and linear growth periods and starts to decline monotonously from the beginning of the stationary growth phase.

This observation suggests cell motility was directly correlated with cell growth and cell aging attenuated cell motility. Of special interest is the observation in the stationary growth region where the motility of the cells decays with time. The question arises whether the aged cells give birth to fewer motile cells or whether other factors, such as self-shading or nutrient depletion, caused this reduction in cell motility. We could not address this question due to experimental limitations, but it would be an interesting topic for future study.

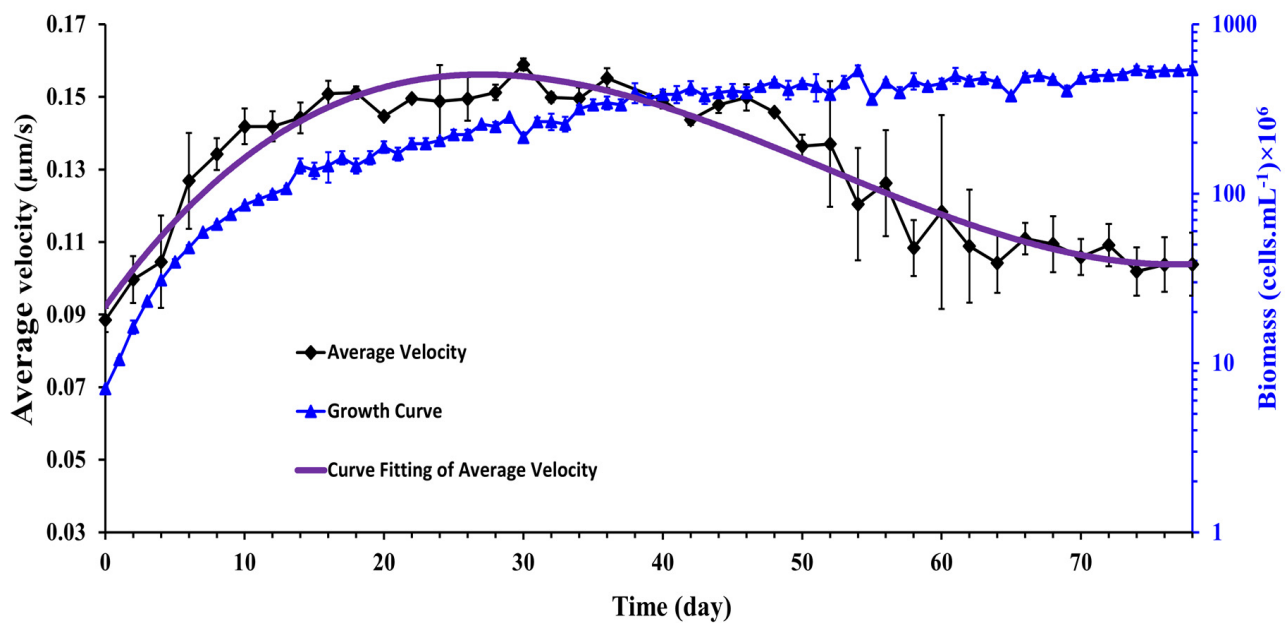


Figure 6. Time history of the average velocity of *Synechocystis* cells. The purple line was fitted to a non-linear third-order polynomial model. ($n = 3 \pm \text{SD}$).

A similar pattern of results was observed by Fadlallah et al. [19] where they cultured *C. reinhardtii* at 22 ± 1 °C with a photon flux of 70 ± 5 $\mu\text{mole}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$ and a stirring rate of 360 rpm for 15 days. Their results showed the mean velocity of *C. reinhardtii* improved noticeably during the exponential phase while the average velocity oscillated randomly around a constant value in the stationary phase. In this experiment, however, the authors did not observe the decay of cell motility, perhaps because the experiments were not followed a long time enough in the stationary growth phase to verify the motility decay as was conducted in the present work.

These velocity data were fitted to a non-linear third-order polynomial model in the form:

$$V_m = a_1d^3 + a_2d^2 + a_3\cdot d + a_4 \quad (6)$$

The model parameters were determined using the “statsmodels” package in Python. The parameter values and their 95% confidence intervals (given in brackets) are:

- $a_1 = 8.36 \times 10^{-7}$ (6.44×10^{-7} to 1.03×10^{-6})
- $a_2 = -1.31 \times 10^{-4}$ (-1.54×10^{-4} to -1.09×10^{-4})
- $a_3 = 5.30 \times 10^{-3}$ (4.53×10^{-3} to 6.06×10^{-3})
- $a_4 = 9.25 \times 10^{-2}$ (8.57×10^{-2} to 9.92×10^{-2})

All parameters were found to be statistically significant ($p < 0.05$). It should be noted that Equation (6) is obtained specifically for *Synechocystis* cells under the given growth conditions. Its generality needs to be studied in future work.

3.5. Dynamics of Cell Motion during Growth

The mean squared displacement (MSD) as a function of experimental time is used to describe the dynamics and the nature of cell motion. After determining cell trajectories, the MSD was computed to analyze the cell transport parameters, such as cell diffusion in its surrounding environment. Figure 7 shows the mean squared displacement of *Synechocystis* sp. CPCC 534 increases linearly with time over a 3000 s observation time, confirming Fickian diffusion behavior. A similar pattern was also found by Vourc’h et al. [23], who studied the surface diffusion of *Synechocystis* sp. PCC 6803 during the early stages of cell contact with a solid surface in diluted suspension. The maximum value of the MSD was estimated from Figure 7 and plotted in Figure 8 as a function of time. Due to the

improvement in the cell motility, the MSD_{max} increased linearly during the exponential and linear growth phases, followed by an almost constant value up to day 44. Then the MSD_{max} declined progressively in the stationary range and roughly leveled off until the end of this period.

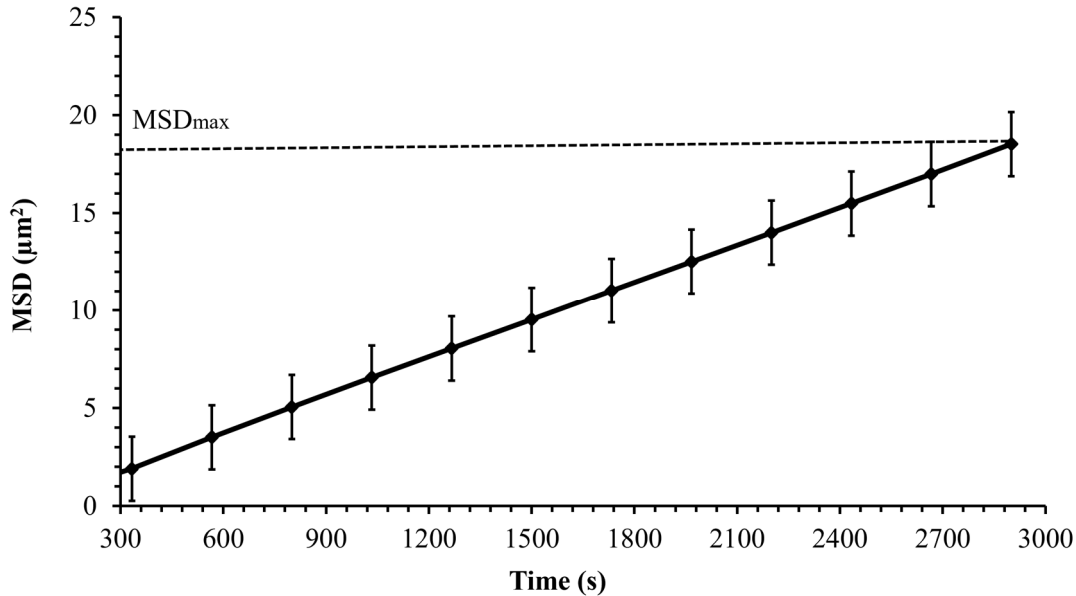


Figure 7. The MSD variation of *Synechocystis* sp. CPCC 534 versus time for a single experiment during 50 min observation period ($n = 3 \pm SD$). Data were fitted to a non-linear third-order polynomial model.

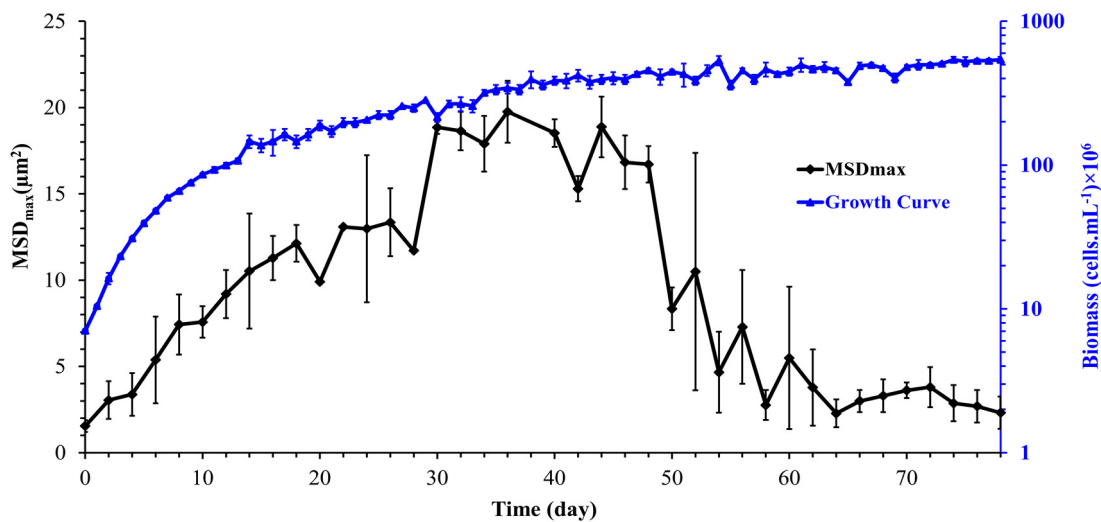


Figure 8. Variation of the maximum mean squared displacement (MSD_{max}) with time during *Synechocystis* sp. CPCC 534 growth ($n = 3 \pm SD$).

To assess the details of the temporal evolution of the motility during an observation period, we divide the 50 min duration of the recording into 12 temporal windows centered around time t . We denote δ , the time elapsed between two successive values of t . Inside each window centered on t , we compute $MSD(t, \Delta)$ as the MSD computed for a time interval Δ as:

$$MSD(t, \Delta) = \frac{1}{N_t} \sum_{i=1}^{N_t} \frac{1}{\delta - \Delta} \sum_{t'=t-\frac{\delta}{2}+\frac{\Delta}{2}}^{t+\frac{\delta}{2}-\frac{\Delta}{2}} \left[X_i \left(t + \frac{\Delta}{2} \right) - X_i \left(t - \frac{\Delta}{2} \right) \right]^2 \quad (7)$$

where N_t is the number of trajectories recorded on the temporal window.

For most applications, one would need to know the behavior of the diffusion coefficient as a function of time. As $MSD(t, \Delta)$ is proportional to Δ inside each temporal window centered around t , this enables the derivation of a time-dependent diffusion coefficient $D(t)$ such as:

$$D(t) = \frac{MSD(t, \Delta)}{4\Delta} \quad (8)$$

Figure 9 shows a slow decrease in the diffusion coefficient with time reaching a constant value asymptotically after 3000 s. A time-dependent diffusion coefficient was also observed in [23] and was related to the secretion of EPS by *Synechocystis* cells. As was shown in the previous section, the *Synechocystis* motility is age-dependent, and therefore, it is expected that the diffusion coefficient is age-dependent as well. Thus, the instantaneous diffusion coefficient becomes concomitantly time and age-dependent, a fact that makes its dynamics quite complex. However, the long-term diffusion coefficient is of special interest in most practical applications. The long-term diffusion coefficient obtained from the experiments carried out for each point on the growth curve is plotted in Figure 10 for the whole growth period of *Synechocystis* sp. CPCC 534. Based on Figure 10, it followed the same pattern as the MSD_{max} profile, confirming that cell motility dominates the diffusion dynamics of *Synechocystis* sp. CPCC 534.

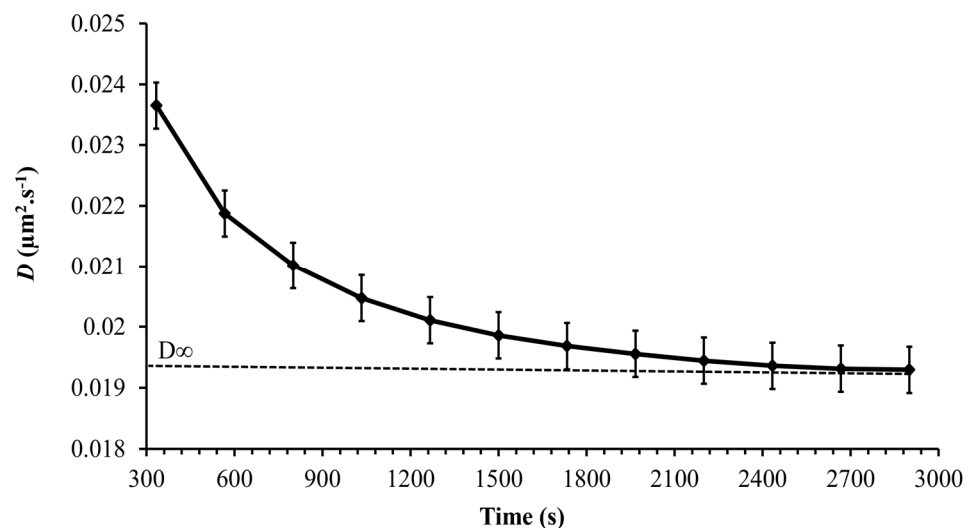


Figure 9. Variation, during 50 min observation time, of the time-dependent diffusion coefficient of *Synechocystis* sp. CPCC 534 ($n = 3 \pm SD$).

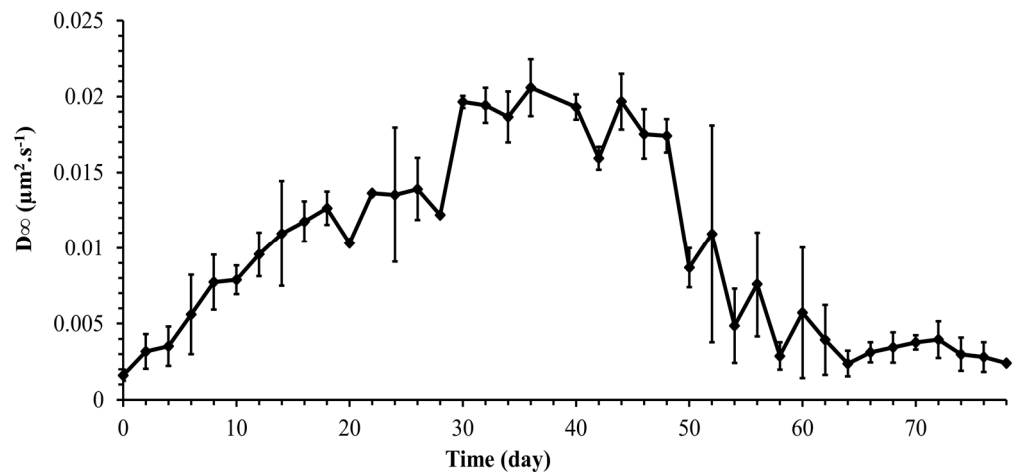


Figure 10. Variation of long-term diffusion coefficient (D_{∞}) during the growth period of *Synechocystis* sp. CPCC 534 ($n = 3 \pm SD$).

3.6. Probability Density Function (PDF)

In active suspensions, the probability density function (PDF) of cell displacements is usually non-Gaussian; or is a combination of a Gaussian distribution for long time intervals and an exponential distribution for short time intervals [37]. Figure 11 plots the PDFs along the x-axis for various time intervals at (a) the beginning of the exponential growth phase, (b) the end of the exponential growth phase, and (c) the end of the stationary phase. The PDF distributions are symmetrical and zero-centered and show a smooth transition from exponential to a Gaussian distribution with passing time. The central part of the PDF relates to the low motility (tumble) part of cell trajectories, while the tails indicate the higher cell motility (run) parts [38]. Figure 11a shows that the PDF consists of two parts: a sharp central part and wide tails. The central part of the PDFs for each observation time corresponds to the non-diffusive behavior of *Synechocystis* sp. CPCC 534 (tumble), and their maximum values decline with the passage of time as the motility of cells increases. The tail of each displacement distribution broadened as time passed, implying that the diffusive behavior of cyanobacteria tends to be Gaussian at longer times. The difference between Figure 11a,b indicates that the tail of each displacement distribution had been expanded noticeably from the beginning of the exponential growth phase to the end of this phase, signifying that cell motility had increased as cells matured. However, the PDF was remarkably narrowed at the end of the stationary phase, Figure 11c, when cells aged, and their motility declined and was dominated by tumbling motion.

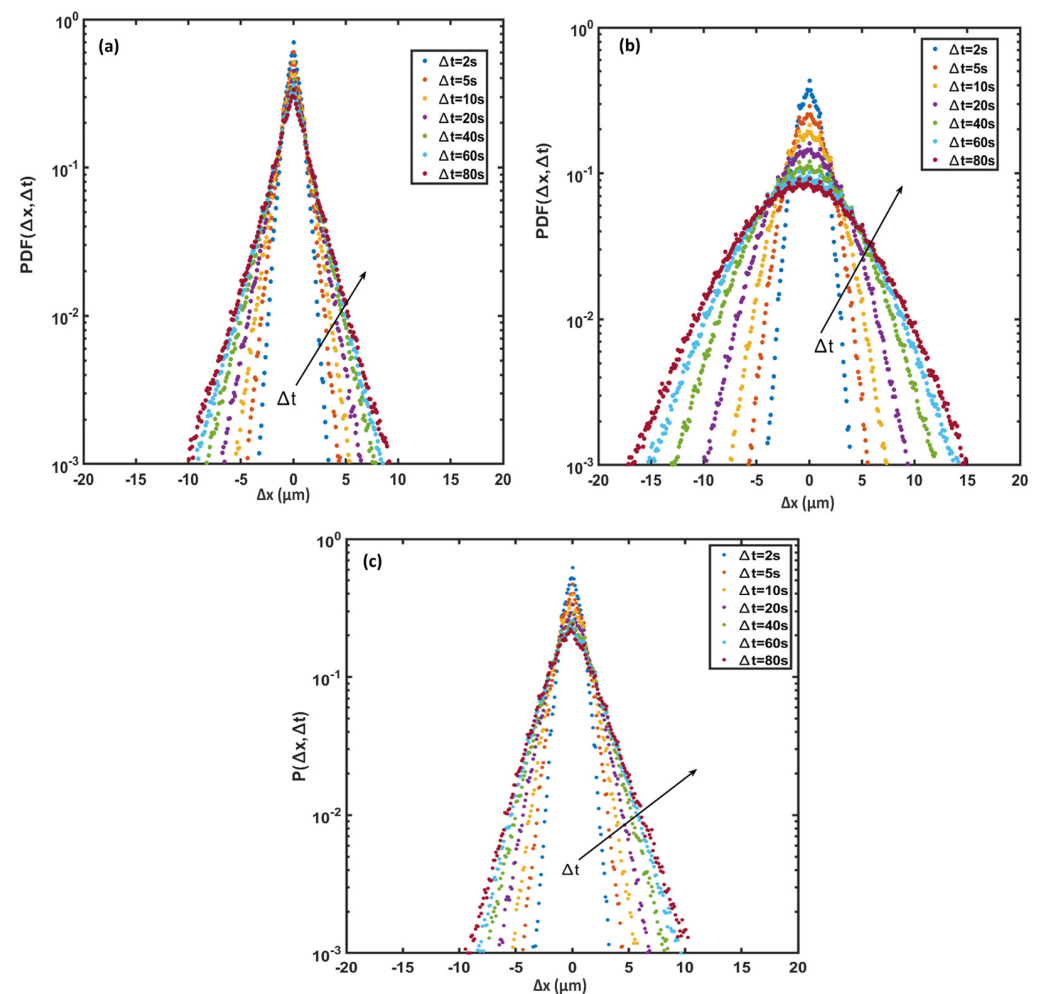


Figure 11. Displacement probability density function (PDF) along the x-axis for *Synechocystis* sp. CPCC 534 cell trajectories (a) beginning of the exponential growth phase, (b) end of the exponential growth phase, and (c) end of the stationary phase. Arrows show increasing time intervals.

4. Conclusions

The motility of wild-type *Synechocystis* sp. CPCC 534 was investigated during its complete growth cycle for 78 days. This rather long observation time, new in the literature, was aimed at investigating the correlation between the cell growth phase and cell motility evolution during the whole growth cycle, from the exponential to the decay phase.

The results revealed a direct correlation between cell motility and the cell growth phase: cell motility increased with time during the exponential linear growth phases. It declined all through the stationary phase before reaching a constant value asymptotically. The mean squared distance traveled by cells at different stages of growth followed a trend similar to that of the cell motility time-variation, as was expected.

The instantaneous diffusion coefficient of *Synechocystis* at all growth states decreased with time and asymptotically reached a constant value. This type of time dependence for diffusion coefficient has already been observed [23] and has been shown experimentally to be due to the secretion of exopolysaccharides by *Synechocystis* cells [5]. However, the present work revealed that the long-term diffusion coefficient is also directly correlated with the growth state of the cells and follows the same time evolution as that of the MSD.

The PDF of the cell displacement, on the other hand, confirmed the close correlation between the cell motility and its growth state; at the beginning of the exponential phase, the PDFs were of exponential type, revealing the non-diffusive behavior of the cells. The tail of each displacement distribution then broadened as time passed, implying the diffusive behavior of *Synechocystis* cells, and tended to Gaussian at longer times. The results of this study find their application in the prediction of biofilm formation on photobioreactor walls.

Author Contributions: Conceptualization, H.P.; Methodology, Z.S., M.M.A., T.V., C.T.D. and H.P.; Software, Z.S., T.V. and C.T.D.; Validation, M.M.A., C.T.D., T.V. and H.P.; Formal Analysis, Z.S., M.M.A., T.V., C.T.D. and H.P.; Investigations, Z.S. and M.M.A.; Resources, C.T.D. and H.P.; Data Curation, Z.S.; Writing—Original Draft, Z.S.; Review and Editing, M.M.A., T.V., H.P. and C.T.D.; Supervision, M.M.A., C.T.D. and H.P.; Project Administration, M.M.A. and H.P.; Funding Acquisition, H.P. and C.T.D. All authors have read and agreed to the published version of the manuscript.

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Data Availability Statement: There is no supporting data for this paper.

Conflicts of Interest: The authors declare no conflict of interest.

Abbreviations

d	Day
D	Diffusion coefficient
EPS	Exopolysaccharide
k	Doubling per day
MSD	Mean squared displacement
OD	Optical density
PBR	Photobioreactor
PDF	Probability density function
r	Growth rate
V_m	Average velocity

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