Chloroplasts in plant cells show active glassy behavior under low light conditions.

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Plants have developed intricate mechanisms to adapt to changing light conditions. Besides photo- and heliotropism - the differential growth towards light and the diurnal motion with respect to sunlight - chloroplast motion acts as a fast mechanism to change the intracellular structure of leaf cells. While chloroplasts move towards the sides of the plant cell to avoid strong light, they accumulate and spread out into a layer on the bottom of the cell at low light to increase the light absorption efficiency. Although the motion of chloroplasts has been studied for over a century, the collective organelle-motion leading to light adapting self-organized structures remains elusive. Here we study the active motion of chloroplasts under dim light conditions, leading to an accumulation in a densely packed quasi-2D layer. We observe burst-like re-arrangements and show that these dynamics resemble colloidal systems close to the glass transition by tracking individual chloroplasts. Furthermore, we provide a minimal mathematical model to uncover relevant system parameters controlling the stability of the dense configuration of chloroplasts. Our study suggests that the meta-stable caging close to the glass-transition in the chloroplast mono-layer serves a physiological relevance. Chloroplasts remain in a spread-out configuration to increase the light uptake, but can easily fluidize when the activity is increased to efficiently re-arrange the structure towards an avoidance state. Our research opens new questions about the role that dynamical phase transitions could play in self-organized intracellular responses of plant cells towards environmental cues.

keywords: chloroplast photorelocation | active glasses | dense active matter | organelle movement

Plant sensing, consciousness, and movement have astonished scientists and philosophers since ancient times [1]. From Plato, who in *Timaeus* [2] associated plants with a "soul" that lacks "judgment and intelligence", but shares "sensation, pleasure, pain, and desire"; to renaissance scientists like Porta, Bacon, and Hooke who wondered about the mechanism of plant movement, and to seminal work of Charles and Francis Darwin on plant behavior [3]. It is now known that, despite generally being limited in their mobility, plants have evolved many movement mechanisms across both length and time scales to maintain optimal development and adapt to their environment [4, 5]. Examples of these behavioral mechanisms are phototropism (response to light), gravitropism (response to gravity), and thigmotropism (response to touch). Such movements can occur on various time scales [4] and mostly on organismal length scales. However, the response to an environmental cue can also occur on a cellular scale. An example of such responses is the active reaction to light: on large scales, via phototropism, the plant grows and moves towards light [6]. Meanwhile, on a small scale, chloroplasts can rearrange to optimize the photosynthesis and respiration processes [7–9]. Hence, chloroplast motion enables plants to efficiently absorb light in order to generate metabolites as a product of photosynthesis while avoiding damage due to strong light.

The light-induced movement of chloroplasts was first observed in the 19th century [10, 11]. These organelles can change the direction of movement towards or away from light [12, 13] without turning [14, 15], but by photo-receptor dependent re-distribution of a chloroplast-specific membrane protein CHUP1 [16–18]. This protein is essential for correct chloroplast positioning [19, 20] and acts as a nucleation factor [18] for chloroplast-specific short actin filaments [21, 22]. The polymerization of these filaments leads to a net propulsion of the organelle [23]. On the one hand, chloroplasts avoid strong light by fast escape movements [21, 24, 25], which leads to a decrease of potential photo-damage [7, 26, 27]. On the other hand, an accumulation response towards weak light sources [15, 28] leads to an increased absorption efficiency [9] as chloroplasts self-organize into a dense layer on periclinal cell walls [29]. This implies that chloroplast motion simultaneously maximizes photosynthetic performance while minimizing photo-damage.

Chloroplasts in various plants can achieve high surface fractions of around 70% [30, 31] upon the accumulation

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towards dim light. Interestingly, higher numbers of smaller chloroplasts enhances mobility [32], while osmotic expansion of chloroplasts leads to an effective inhibition of the light response [33]. These observations point towards a dependence of the chloroplast mobility on their packing density and their individual activity. It has been shown in other biological systems that packing density and geometry impacts the fluidity and transport properties. Increased densities in confluent tissues can lead to a glass transition [34–36], arresting the cell motion and stabilizing the tissue in a *solid-like* state [37–39]. Active jamming plays a role in morphogenesis of zebrafish embryos [40], stabilizes bronchial epithelium of the airway during development [41], and may be key to understand cancerous tissue [42]. On even smaller scales, the cytoplasm itself features glass-like transport properties in bacteria [43] or human cells [44], indicating that active glassy dynamics occur across scales and are ubiquitous [45].

These observations indicate that jamming- and glass-transitions in biological active matter play a key role in understanding the structural changes and transport in and around cells.

In this study, we investigate the dynamics of chloroplast motion under dim light conditions in the water plant *Elodea Densa* (Fig. 1). We find that the dense, actively driven system shows dynamics similar to systems close to the glass transition, both in their overall statistics and individual particle dynamics. Despite the complexity of the system, we can construct a mathematical model that can re-produce the step size statistics with experimental and physiological parameters. We use this model to test the stability of the state close to the glass transition and show that the chloroplast motion can exhibit more *liquid-like* dynamics when the activity just slightly increases. Our observations suggest that chloroplasts naturally organize close to a critical point in order to reach a large area coverage. Consequently, chloroplasts increase photosynthetic efficiency while maintaining the ability to quickly respond to unfavorable light conditions by actively *fluidizing* the configuration for efficient avoidance movement.

Figure 1: **(a)** *Elodea Densa* waterplant as cultured under controlled environmental conditions. **(b)** Leaves are detached from the plants and **(c)** single epidermal cells are studied. Chloroplasts (arrow) are spread out in a single layer under dim light conditions.

Glassy Features of Chloroplast Motion under dim Light

We track the positions of 1529 chloroplasts under dim light and constant environmental conditions and segment their morphological features (*Materials and Methods*, Supplementary Movie 1). In dim light chloroplasts decrease their mobility [13] and settle onto the periclinal walls in a single layer (Fig. 1c). We measure a two-dimensional average packing fraction of approximately 71%, which is consistent with measurements in land plants [30, 31]. This suggest that chloroplast motion depends on the close-by neighboring chloroplasts. Chloroplasts are anchored to the plasma membrane [46], allowing for mostly two-dimensional motion. Because they can move into all directions without rotating [15], the orientation of the chloroplast is uncoupled from the migration direction. Consequently, an analysis of the two-dimensional center of mass trajectories alone is sufficient to elucidate underlying stochastic processes.

The mean-squared displacement (MSD) (Fig. 2a) reveals a broad spectrum of power law exponents α at small time scales ($\Delta t = 10 - 100$ s), ranging from sub-diffusive ($\alpha < 1$) to super-diffusive ($\alpha > 1$) trajectories (see the histogram in Fig. 2a). At long time scales of $\Delta t \gtrsim 5$ min the MSD approaches overall diffusive scaling ($\alpha = 1$), with approximately exponentially distributed diffusion coefficients (see inset in Fig. 2d) and average diffusion coefficient $D \approx 8 \times 10^{-4}$ μm^2/s. Additionally, the measured velocity auto-correlation (Fig. 2b) reveals that the velocity v of a chloroplast motion de-correlates within several seconds. Together with the measured MSD scaling exponents, the velocity auto-correlation demonstrates a clear picture on the individual dynamics of chloroplasts: they are driven by a stochastic process which has a back-scattering effect, as suggested by the anti-correlation of the velocity after a few seconds (Fig. 2b). A few trajectories show slightly more persistent motion and larger scaling exponents indicating coordinated movement of actively driven chloroplasts, which we will discuss later.

To further investigate the nature of the stochastic process constituting the motion of densely packed chloroplasts, we study the self-part of the van Hove function (step size distribution), $\mathcal{G}_s(\Delta r, \Delta t)$ (Fig. 2c) at various delay times $\Delta t = 10, 100, 1000$ s. Strikingly, the van Hove function deviates from a Gaussian function at small and

Figure 2: Trajectory analysis reveals sub- to super-diffusive transport and local trapping. **(a)** MSD of all trajectories relative to standard deviation. Colormap represents the power-law scaling exponent α fitted on short time scales below 100 s. The exponent is widely distributed but peaks around a diffusive scaling regime ($\bar{\alpha} \approx 0.9$). While a few trajectories exhibit super-diffusive scaling many trajectories show sub-diffusive scaling laws at small times $\Delta t \lesssim 100$ s. Thick lines: ensemble averages over super-diffusive trajectories $\alpha \geq 1.1$ (purple), diffusive trajectories $0.5 < \alpha < 1.1$ (black) and strongly sub-diffusive trajectories $\alpha \leq 0.5$. Notably on large time scales all regimes approach a diffusive scaling. Inset: MSD with units, most trajectories reach one micron displacement after long time ($t > 100$ s) only. Thick lines compare the ensemble average of all trajectories (blue line) to the ensemble averaged MSD of simulations (red line). Purple, black and orange lines correspond to ensemble averages in the main panel. **(b)** Velocity autocorrelation functions (VACF) of trapped (orange) and jumping (black) chloroplasts. Some jumping chloroplast exhibit pronounced temporal correlation, which shifts the VACF to higher values. Anti-correlation as a sign of fast re-orientations is less pronounced in jumping chloroplasts, indicating a directed motion. **(c,d)** Self part of radial van Hove function at different lag-times for experiments (c) and model (d) reveals non-Gaussian displacements on short lag-times (compare with the gray area, representing a Gaussian function). For larger lag-times, the distributions are approaching a Gaussian at small displacements, hence the comparison with a continuous time random walk (CTRW) model [47] (dashed line in (c)) improves. The proposed model (d) accounts for the exponential tails, starting at a critical displacement (vertical line), and the exponential-to-Gaussian crossover at small displacements. Inset: diffusion coefficients of experiments (triangles) and simulation (circles) can be well described by a corrected exponential distribution (solid line).

large displacements in two distinct manners. Firstly, small displacements are governed by a transition from a non-Gaussian to a Gaussian step size distribution for increasing delay times. This feature is commonly found in the dynamics of (fractional) Brownian, non-Gaussian diffusive motion and has been observed in organelle and RNA transport [48, 49]. Secondly, the chloroplast step size distribution exhibits an exponential tail at all time scales. The transition towards the exponential tail happens around $\Delta r \approx 2.3\,\mu$m, corresponding to one average chloroplast radius plus one standard deviation. This displacement is more than twice as large as the inter-chloroplast distance $l \approx 1.026\,\mu$m, which is calculated via the difference of average chloroplast diameter and nearest neighbor distance [50] (SI Fig. 5d,e). Hence, small displacements $\Delta r < l$ correspond to free motion until a chloroplast reaches l. Larger displacements $l \leq \Delta r < 2.3\,\mu$m are likely connected to the interaction of multiple chloroplasts, while displacements exceeding $2.3\,\mu$m show a sudden transition towards an exponential step size distribution. Such a heavy tail has been previously observed in other systems on various scales, such as the bacterial cytoplasm [43], particle transport in human cells [44], motion of cells in confluent tissues [34], and non-active systems like super-cooled liquids [51] and colloidal glasses [52]. Using a continuous-time random walk (CTRW) model it was shown that exponential tails of the van Hove function display an universal feature of particle motion close to the glass transition [47, 53], which can be fitted to our experimental results (dotted line in Fig. 2c). However, this model does not fully account for the non-Gaussian center of the observed distribution, as we will discuss in more detail later.

With a high packing fraction of about $\phi \approx 71\,\%$ (see *Materials and Methods*) the amorphous chloroplast monolayer bears a noticeable structural resemblance with colloidal glasses, which may explain the key features of chloroplast motion. The direct study of individual trajectories in colloidal glasses has led to the conclusion that the stretched exponential step size distribution arises as a result of heterogeneous dynamics of slow and fast particles [52]. This heterogeneity was found in active systems and also constitutes a hallmark for glasses [45]. Hence, our observation of the exponentially tailed step size distribution indicates the existence of dynamical heterogeneity underlying chloroplast motion.

Dynamic Heterogeneity due to the Co-existence of locally entrapped and re-arranging Chloroplasts

The aforementioned hypothesis requires analysis of individual chloroplast trajectories. We find qualitatively different trajectories. While many chloroplasts remain caged by their neighbors (slow particles), a small fraction (88 out of 1529) can escape their local confinement and move for longer distances until they get trapped again as a jump-like motion (fast particles) (Fig. 3a,e). The dynamic heterogeneity is studied by disentangling the fast and slow dynamics of the system, based on the chloroplasts maximal displacement d_{max} during the experimental

Figure 3: Intermittent re-arrangements of chloroplasts. **(a)** Time series of chloroplast motion during a bursting event. Trajectories are color-coded by their final distance d. While most chloroplasts remain at their initial location, a small region shows enhanced activity (white box). **(b)** Time series of average velocity (gray) and the 95th percentile velocity (black) of the cell in (a). The gray region around the peak corresponds to the time series in (a) and (c). The large increase of the 95th percentile indicates that the single event is an extreme of the velocity distribution. **(c)** Velocity and distance d from initial position for trajectories which undergo strong re-arrangement (a, white box). Correlated motion of the sudden re-location becomes apparent. Colors for comparison of trajectories in (c) and (d). **(d)** Close-up of trajectories in (a), white box. Chloroplasts show random trapped motion, followed by transient collective motion, and subsequent fluctuations. Scale bar: $10\,\mu m$. Dark color saturation indicates high speed. **(e)** Histogram of maximal displacements with threshold value (dotted line, see *Materials and Methods*). Inset: Distance from initial position for all chloroplasts exceeding the distance threshold showing similar pattern: sudden re-arrangements and subsequent plateau at approximately discrete positions of a few chloroplast diameters. Trajectories from (c) and (d) are marked in respective colors.

time (Fig. 3e) [52, 54] and their maximal speeds (see *Materials and Methods*). We uncover large regions of trapped particles and smaller regions of more mobile chloroplasts (Fig. 3a, SI Fig. 5 a) by marking trajectories by their total displacement from their initial position. Fast chloroplasts re-arrange within a short period of time and arrest the dynamics afterwards. The intermittent bursts of chloroplast re-locations can be detected in the tails of the speed distribution (Fig. 3b), as represented by the 95th percentile, while the average speed does not exhibit strong changes. This confirms that re-arrangements are connected to exceptionally high speeds of chloroplast motion, which transiently occur in a small fraction of all chloroplasts in a cell.

Besides the spatial co-localization of chloroplast re-locations, we find that their motion can be temporally correlated and follows a string-like profile. Individual chloroplasts can break out of their local cage and push their neighbors with them (Fig. 3c,d, SI Fig. 5a-c). Additionally, fast chloroplasts have longer velocity auto-correlation times, suggesting an increase in the persistent motion, while slow particles oppose their motion on smaller time scales (Fig. 2b). The movement of all re-arranging chloroplasts follows a similar scheme: a sudden chloroplast displacement at elevated velocities plateaus at a distance of only a few chloroplast diameters (Fig. 3e inset). This suggests that they are locally entrapped again. Similar sudden and hardly predictable re-arrangements have been found in simulations of active Ornstein-Uhlenberg particles [55], sheared granular media [56] and in epithelial tissues [34]. Interestingly, the string-like cooperative motion of particles in a close-to glassy state were observed in super-cooled liquids [57, 58], simulations of Lennart-Jones mixtures [59] and granular beads [60].

The striking similarity to many systems close to the glass transition implies that chloroplast motion in this dense single-layered configuration obeys similar physical constraints and mechanisms. In the following, we will build a mathematical formulation based on physiological parameters that models the co-existence of trapped particles and regions of higher mobility in the chloroplast dynamics. This approach allows us to study the effect of activity and crowding on the chloroplast dynamics. Our goal is to infer the stability around the critical state of the chloroplast mono-layer.

A Threshold-Based Jump Diffusion Model reveals that the dim light adapted Chloroplast mono-layer is close to a Glass-transition

We propose a stochastic model of chloroplasts to elucidate the impact of critical length-scales and active diffusion of the system. In this model discrete jumps are triggered by a threshold-based mechanism (Fig. 4a). We consider a particle in a harmonic potential $V(x) = -\frac{k}{2}x^2$ that can undertake a jump of finite velocity v upon exceeding

a critical distance x_c from the potential minimum position $a(t)$. This leads us to a Langevin-equation:

$$\dot{x} = \begin{cases} -\frac{x}{\theta} + \sqrt{2D(t)}\xi(t), & x - a(t) < x_c \\ -\frac{x_c}{\theta} \pm v + \sqrt{2D(t)}\xi(t), & x - a(t) \geq x_c \end{cases} \tag{1}$$

where $\xi(t)$ is a Gaussian white noise $\langle\xi\rangle = 0$, $\langle\xi(t)\xi(t')\rangle = \delta(t - t')$. The effective spring constant k emerges from the confinement of the neighboring particles [61, 62]. Together with a damping pre-factor γ it defines a correlation time $\theta = \frac{\gamma}{k}$. After every jump the center position of the potential a is translated by an equal amount to maintain the particle inside the harmonic potential. Based on the fluctuating and disordered nature of the chloroplast environment, we furthermore assume that the diffusion coefficient $D(t)$ is described by the square of an Ornstein-Uhlenbeck process [63, 64]:

$$D(t) = Y(t)^2, \quad \dot{Y}(t) = -\frac{Y}{\tau} + \sigma\eta(t). \tag{2}$$

In the equation above, τ is a relaxation time scale, and σ is the noise amplitude of a Gaussian white noise $\eta(t)$. Equation (2) models the fluctuating heterogeneity of the environment and leads to temporal correlations of the diffusion coefficient. We support this assumption by observing that this "diffusing diffusivity" model [63] predicts an exponential distribution of diffusion coefficients [64] consistent with measurements of the long term diffusion coefficient (Fig. 2d, inset). This allows us to use the average long-term (active) diffusion coefficient $\langle D\rangle = \sigma^2\tau = 8 \times 10^{-4}\ \frac{\mu m^2}{s}$ to estimate parameters for the model. Other parameters are inferred from particle trajectories and physiological parameters (*Materials and Methods*). A typical trajectory of this model displays diffusive dynamics with time-correlated variance and sudden jumps of discrete size (Fig. 4b). The resemblance to the experiments is also reflected in the van Hove function (Fig. 2c).

The discrete-sized jumps can account for the exponential tails, while the non-Gaussian to Gaussian transition for increasing timescales and small displacements in the van Hove function can be well described by the diffusing diffusivity model. This model has been used previously to describe active processes in cytoskeletal dynamics [65] and the transport of RNA [49] or colloidal beads on membranes [48].

Note that our model neglects the subdiffusive dynamics at small time scales (Fig. 2 (a)) by construction [63, 64]. Together with the damping parameter and the critical length scale, the effective diffusivity $D = \sigma^2\tau$ steers the caging dynamics, as observed in the MSD of the trajectories (Fig. 4c). At a small noise amplitude σ the system is strongly caged, as chloroplasts are drastically less probable to reach a high enough displacement to jump. If, however, σ is increased by

Figure 4: Simulation results. **(a)** Sketch of the mechanistic model. A Brownian particle is connected to a spring and a jump-element which undergoes sudden extension or compression if a critical displacement is reached. **(b)** Example trajectory shows sudden jumping events before being trapped again. **(c)** Ensemble-averaged MSD from low to large steady-state diffusion coefficients D shows a transition from trapped to free motion. Parameters obtained from our data suggest that the system is close to a transition towards unconfined motion. Inset: variability of trajectories shows dynamics ranging from caged to jumping motion, dotted line: ensemble average MSD.

$20 - 40\%$, the jumping dynamics are more pronounced, suggesting that chloroplasts are close to a critical point and can easily transition into a *liquid-like* state. In our model this transition is, to a great extent, governed by the rate of jump events undertaken by the particle. We define the jump time distribution in our model by construction as the first passage time distribution of an Ornstein-Uhlenbeck process at long timescales, at which the fluctuating diffusion coefficient approaches $D = \sigma^2\tau$. The first passage time distribution approaches a series of exponential functions [66] and the mean first passage time is controlled by a dimensionless quantity $x_c/\sqrt{D\theta}$ [67]. This suggests that a similar transition is also possible when the critical length-scale x_c increases, while D is held constant (SI Fig. 7), which has been observed in studies of plant cells with varying chloroplast numbers and areas [32]. Hence, despite the simplified description of the cage effect and jumping dynamics, this toy model is sufficient to predict that, not only the activity σ, but also the critical size x_c can easily drive the system out of the caged dynamics.

Conclusion

Plants have developed multiple adaptation mechanisms to changing light conditions [27]. One of these mechanisms is orchestrated via independent movements of disk-like chloroplasts towards or away from light. The motion of densely packed chloroplasts adapted to low light conditions exhibits a striking similarity to the caged dynamics of super-cooled liquids or colloidal suspensions close to glass transition. We observe actively driven organelles confined in a single layer at high packing fraction. Intermittent coordinated motion of single or

multiple chloroplasts leads to a heavy-tailed step size distribution caused by the heterogeneous nature of the environment. Furthermore, we uncover dynamical features similar to systems close to the glass transition, such as dynamically heterogeneous regions, sudden chloroplast displacements and even string-like coordinated movements. This renders chloroplasts a novel biological system exhibiting glassy behavior, while maintaining an unaltered disk-like shape, contrary to other biological active glasses like epithelial cell sheets [34–37, 40–42].

We compare the observed dynamics to a one-dimensional model, whose parameters are obtained from experiments and physiological values. The statistics of this model are in agreement with our data. This indicates that burst-like chloroplast jumps could be triggered upon exceeding a threshold, similar to the cage length in glasses [50]. By construction, the waiting times between the discrete jump events in this model are identical to the first-passage time of an Ornstein-Uhlenbeck process with diffusing diffusivity, which is approximately exponentially distributed for large enough boundaries and times [66] (SI Fig. 7c). This indicates that the underlying statistics of this model are similar to those of the continuous time random walk (CTRW) model proposed by Chaudhuri et al. [47, 53], which is used to describe a universal step size distribution function in various systems close to the glass transition. However, this CTRW model needs the mean waiting time between jump events as a fit parameter, which is difficult to infer from limited data directly. Our model enabled us to convert a waiting time between jump events into a length scale x_c, which can be more easily extracted. Our model calculations with varied parameters suggest that chloroplasts operate at a meta-stable point and can easily perturb the mono-layer structure by slight enhancements of their activity, which could explain the fast adaptation mechanism towards strong light. Additionally, these simulations hint towards a cage effect inhibiting chloroplast motion when the chloroplast size is changed, as has been observed before [32, 33].

We interpret our results as follows: chloroplasts accumulate into a single layer in response to weak light in order to achieve a high surface coverage and therefore maximize photosynthetic efficiency [9]. Since we know that chloroplasts avoid highly intense light [68], they must maintain the ability to undertake this avoidance motion efficiently without being hindered by other chloroplasts. Therefore, the ability to increase the activity and a sufficiently loose packing of chloroplasts are necessary. Our study shows that chloroplasts in dim light conditions resemble a system close to the glass transition. However, the transitions towards and away from this state are yet to be discovered. How chloroplasts spread on the periclinal cell walls as a result of an accumulation response under dim light remains a subject for further research, since a direct observation of this process was not possible. Further studies on collective escape from this low-light adapted state towards an avoidance configuration will shine a light on the various dynamic phase transitions this system can undergo, which were also found in active glasses [55, 69]. Additionally, the impact of confinement due to the cell walls on the transitions between dark and light-adapted states remains a matter for further investigation.

MATERIALS AND METHODS

Preparation and Imaging

Elodea Densa was kept in an aquatic culture. Ambient light conditions were applied 1 h before image acquisition. For imaging, a low-light adapted leaf was detached and placed between two glass slides. We study the lower cell layer of the two-layered leaf tissue [70], as these are more strongly involved in photosynthesis [71]. The upper epidermal cell layer is important to enhance gas exchange due to reduced gas diffusion in water [72]. Bright field microscopy was performed with a `Nikon TI2` microscope using a halogen light source and a red-light 620 nm cut-on wavelength filter at low light intensities of $7.2\,\mathrm{W/cm^2}$ in combination with a `Photometrics BSI Express` sCMOS camera with high quantum yield to enable imaging every 1 s with a $63\times$ water-immersion object ive (NA= 1.2) and pixel resolution of $0.1\,\mu\mathrm{m/px}$.

Image Processing

Image time series were background subtracted using a difference-of-Gaussian method with variances $\sigma_1 = 2\,\mathrm{px}$ and $\sigma_2 = 37\,\mathrm{px}$ and subsequently inverted (SI Fig. 6). The resulting image shows white circular spots, reminiscent of nuclei training data for the StarDist versatile model [73]. Using this pre-trained StarDist network we segmented the images and detect center of mass positions of the particles. The segmented objects were analysed for their area and aspect ratio. Small objects of an area of 250 px, corresponding to an equivalent diameter of less than $1.78\,\mu\mathrm{m}$, were removed. Additionally, elongated objects with aspect ratios of above 3.5 or below 0.3 were excluded to avoid mis-detections of cell walls. We hand-segmented the surrounding cell wall to discriminate chloroplasts in different cells (SI Fig. 5b). The average packing fraction was calculated by the sum of all areas of segmented chloroplast relative to the area of the segmented cells.

Trajectory Analysis

The center-of-mass positions of all segmented chloroplasts were linked to trajectories using a linear assignment problem solver *trackpy* [74]. Spurious trajectories of a duration smaller than $400\,\mathrm{s}$ were not considered in the analysis. Trajectories were classified as *fast* when their maximal displacement lies significantly (one standard-deviation) above the mean particle size (see dotted line in Fig. 3c, SI Fig. 5d) [52, 54], and whose maximal velocity lies above $0.65\,\mu\mathrm{m/s}$. Particle velocities were calculated with the first derivative of the third-order Savitzky-Golay filtered data, with a kernel length of $11\,\mathrm{s}$. The self-part of the van Hove function and the mean squared displacements were calculated with trackpy [74], whilst the MSD scaling exponents α were extracted using a local fitting scheme [75] between $\Delta t = 10\,\mathrm{s}$ and $100\,\mathrm{s}$. As we track the center of mass of the segmented particles, we have two contributions to the error of the tracking algorithm: one from the limited pixel-size of $\sigma_{px} = 0.11\,\mu\mathrm{m}$ and area fluctuations of the mass. As the mask can differ from image to image, we take the area-fluctuations as a proxy to estimate a detection error: $\sigma \approx \sqrt{\frac{\sigma_A}{\langle A \rangle}}\sigma_{px} \approx 0.07\,\mu\mathrm{m}$, with the mean $\langle A \rangle$ and the standard deviation σ_A of the areas obtained by temporal and ensemble averaging over all trajectories.

Simulations

Our model (equations (1), (2)) was integrated using a stochastic Euler-Maruyama scheme with a time stamp $dt = 0.1\,\mathrm{s}$ for 10^6 time steps. Simulation parameters were retrieved from trajectories. We choose the $1/e$ decay of the positional auto-correlation function (see SI Fig. 5f) to estimate the elastic relaxation time $\theta = 1063\,\mathrm{s}$. Although the function does not follow an exponential decay it gives a reasonable estimate for the correlation time. Additionally, we inferred the noise amplitude σ of the diffusing diffusivity from the measurements of the long-time diffusion coefficients of the system as $D = \sigma^2\tau$. Here we set the correlation time $\tau = 90\,\mathrm{s}$ which is well within the range of actin turnover times [76]. The jump velocity $v \approx 0.6\,\mu\mathrm{m/s}$ was estimated from the average maximal velocity of jumping chloroplasts. The critical length scale $x_c = 2.3\,\mu\mathrm{m}$ is the point at which the van Hove function transitions to an exponential (Fig. 2c). This critical length scale is close to the average particle radius plus one standard deviation (SI Fig. 5c). For parameter sweeps, x_c^{-1} and σ were changed from $20\,\%$ to $180\,\%$ of their experimental value, and simulations were repeated 200 times for each parameter. Simulation data was analysed exactly the same as trajectories (see above). The continuous time random walk model (Fig. 2c, dashed line) was implemented as outlined in [44] with parameters $\tau_1 = 3000\,\mathrm{s}$, $\tau_2 = 100\,\mathrm{s}$, $d = 0.6\,\mu\mathrm{m}$, $l = 1.2\,\mu\mathrm{m}$.

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SUPPLEMENTARY INFORMATION

Supplementary Movies

Supplementary movies 1 and 2 can be found under this link: https://drive.google.com/drive/folders/1ezeK4Iy6htm5bvcWWkMW2JPsMTKHkisz?usp=sharing

SI Movie 1: Time lapse (60 min) of background subtracted chloroplasts, false colors. Microscopy under low light assures that the system remains in a dim-light adapted state.

SI Movie 2: Time lapse (60 min) of background subtracted chloroplasts. Close up of a re-arranging cell. Individual chloroplasts in a region re-arrange within short time, while most other chloroplasts remain entrapped between their neighbors.

Figure 5: Extended datafigure. **(a)** All trajectories, colored by the total distance shows localized fast displacements d_{tot}. **(b)** Chloroplasts at time point $t = 0$ colored by their maximal velocity during re-arrangements. Correlates with distance in (a). Gray mask: segmented cells. **(c)** Nearest-neighbor network (gray lines) of chloroplasts at $t = 0$ with corresponding average nearest neighbor distance d_{NN} by colormap. **(d)** Histogram of particle radii R with the critical x_c (dotted line) corresponding to the mean radius increased by one standard deviation. **(e)** Radial density function of the particles around the average density ρ, with a nearest neighbor peak around two particle radii (dotted line) plus their standard deviation (gray). The nearest neighbor shell size can be estimated from an average until the first minimum [50]. **(f)** The autocorrelation function of the position for all particles (gray) and the ensemble average (orange). The $1/e$ decay is located at $\Delta t = 1063$ s (dotted line).

Figure 6: Image processing workflow: brightfield images (left) are background subtracted using a difference-of-Gaussian method. The subtracted image is segmented into individual regions using a StarDist neural network. Each region can be analyzed for its area, aspect ratio and center of mass position. This procedure is done for every frame of the image time series. The positions are assigned to trajectories using trackpy[74] (right).

Figure 7: Parameter variations in the jump model. **(a)** Mean squared displacement, normalized by the displacement at $\Delta t = 0$ for different critical length scales $x_c* = x_{c_0}/x_c$ (solid lines), where $x_{c_0} = 2.3\,\mu$m is the experimental value. Additionally, a comparable parameter sweep is performed by changing the activity $\sigma \sim \sqrt{D}$ (dotted lines) with relative diffusion coefficients $D* = D/D_0$, where $D_0 = 8 \times 10^{-4}\,\mu\text{m}^2/\text{s}$ is the experimental value. **(b)** Waiting time distributions for same parameters as in panel a (data points shown for x_c^*) with double-exponential fits for x_c^* (solid lines) and D^* (dashed lines). **(c)** Example trajectories for different x_c^* (top) and D^* (bottom), for larger D^* or x_c^{-1}, more jump events lead to larger displacements resembling a transition towards a more *liquid* state.