Current Biology, Volume 33

Supplemental Information

An environmentally induced multicellular

life cycle of a unicellular cyanobacterium

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Figure S1. Growth trajectories and population dynamics of replicate *Cyanothece* sp. populations in batch cultures over time. Related to Figure 1.

(A) Comparison of the growth trajectory in batch cultures in 10 mL BG11 medium with 0 mM and 300 mM NaCl over time. The grey area indicates the time period when filaments were observed in medium with 0 mM NaCl. The shortest generation time in freshwater is $G_{0 mM} = 15.2 h$ (from 72 h to 96 h), while the shortest generation time in the highest salinity is $G_{300 mM} = 17.5 h$ (from 24 h to 48 h).

(B) Population dynamics over the period of 5 days in BG11 *without* added NaCl (populations initiated with 5*10⁵ cells/mL, in 1 mL volume each (24-well plates)).

(C) Population dynamics over the period of 6 days in BG11 *with* added NaCl (300 mM) (populations initiated with $5*10^5$ cells/mL, in 1 mL volume each (24-well plates)).

Error bars represent SDs (of each sub-bar for B) (n = 3).





When diluted, filaments kept growing and increased in length, indicated by the observation of filaments of longer than 16 cells in length and by a significantly higher proportion of 8-celled filaments, in contrast to the original culture, where 24 h later only single cells were observed. Error bars represent standard deviation of each sub-bar (n=3).



Figure S3. Models of the acting substance concentration effect in the disconnecting and connecting compound models. Related to Figure 3.

We consider multiple models of the relationship between acting substance concentration and its effect on the filaments (12 in the disconnecting compound family of models and 8 in the connecting compound family). Disconnecting compound models bring more accurate fit of experimental data than the connecting compound models. The simplest 1-parameteric models yield high regression erros but 3-parameteric models do not bring an advantage over some 2-parametric ones.



Figure S4. Cumulative distribution functions and the minimal regression errors obtained for compound models. Related to Figure 3.

(A) 12 disconnecting compound models can be classified into two groups: models with a good fit having minimal regression errors below 0.17, and models with worse fit, for which the minimal regression error is above 0.21 (can be increased to 0.26 if quadratic model is dropped), see also Table S2. (B) 8 connecting compound models can be classified into two groups: models with a good fit having minimal regression errors around 0.21, and models with worse fits, for which the minimal regression error is above 0.22, see also Table S3. Plots show sample cumulative distribution functions of regression errors from 250 independent optimizations for each model. Dashed lines represent the minimal regression error in each model.

Table S1. The morphology of *Cyanothece* sp. is dependent on the composition of the medium. Related to Figure 2.

Fresh culture medium (BG11) was added to ddH_2O and both supernatants (filament inhibitor: supernatants from cultures inoculated with 5*10⁶ cells/mL starting cell densities, harvested 24 h after inoculation, and filament fragmentor: supernatants from cultures inoculated with 5*10⁵ cells/mL cell densities immediately after filament fragmentation, harvested at 96 h), creating BG11 ratios from 0 – 100 % with 20 % increments. The emergence of the filamentous morphology was recorded after 48 h, starting with single cells of *Cyanothece* sp. in each dilution treatment. "+" represents filament occurrence; "-" represents no filament occurrence. While 20 % of BG11 in ddH₂O provided sufficient nutrients for filament formation, 60-80 % of the BG11 was necessary to dilute the filament fragmentor/inhibitor medium before filaments were observed.

	BG11 ratio						
	100%	80%	60%	40%	20%	0%	
ddH ₂ O	+	+	+	+	+	-	
Filament fragmentor	+	+	+	-	-	-	
Filament inhibitor	+	+	-	-	-	-	

Table S2. Action law E(T) in models used in the toxic and disconnecting compound model families and the minimal regression errors obtained across 250 independent optimizations. Models with the highest quality fitting are highlighted. Related to Figure 3.

Model of the acting	Law of action	The smallest	The smallest				
substance		disconnecting	toxic				
concentration effect		compound	compound				
		regression	regression				
		error	error				
1 parameter models							
Constant	$E(T) = E_0$	0.292	0.590				
Proportional	$E(T) = \alpha T$	0.262	0.732				
2 parameter models							
Linear	$E(T) = \alpha T + E_0$	0.262	0.596				
Step	$E(T) = \begin{cases} 0, & T < T_0 \\ E_0, & T > T_0 \end{cases}$	0.153	0.594				
Fracture	$E(T) = \begin{cases} 0, & T < T_0 \\ \alpha T, & T > T_0 \end{cases}$	0.160	0.721				
Breaking point	$E(T) = \begin{cases} 0, & T < T_0 \\ \alpha(T - T_0), & T > T_0 \end{cases}$	0.170	0.724				
Michaelis-Menten	$E(T) = E_0 \frac{T}{T + T_0}$	0.270	0.598				
Quadratic	$E(T) = \left(\frac{T}{T_0}\right)^2 + E_0$	0.217	0.583				
Top-capped	$E(T) = \begin{cases} \alpha T, & T < T_0 \\ \alpha T_0, & T > T_0 \end{cases}$	0.262	0.593				
Bottom-capped	$E(T) = \begin{cases} \alpha T_0, & T < T_0 \\ \alpha T, & T > T_0 \end{cases}$	0.263	0.613				
3 parameter models							
Sigmoid	$E(T) = \frac{E_0}{1 + e^{-\alpha(T - T_0)}}$	0.152	0.583				
Saturating exponent	$E(T) = E_{\max} - (E_{\max})$	0.269	0.560				
	$-E_{\min})e^{-\alpha I}$						

Table S3. Action law E(T) in models used in the connecting compound models family and the minimal regression errors obtained across 250 independent optimizations. Models with the highest quality fitting are highlighted. Related to Figure 3.

Model of the acting	Law of action	The smallest connecting			
substance		compound regression			
concentration effect		error			
1 parameter models					
Constant	$E(T) = E_0$	0.292			
2 parameter models					
Linear	$E(T) = \alpha(1-T) + E_0$	0.264			
Step	$E(T) = \begin{cases} E_0, & T < T_0 \\ 0, & T > T_0 \end{cases}$	0.208			
Quadratic convex	$E(T) = E_0 + \alpha (1 - T)^2$	0.234			
Quadratic concave	$E(T) = max(0, E_0 - \alpha T^2)$	0.216			
Inverse	$E(T) = \frac{E_0}{1 + \frac{T}{T_0}}$	0.214			
3 parameter models					
Sigmoid	$E(T) = \frac{E_0}{1 + e^{\alpha(T - T_0)}}$	0.213			
Decaying exponent	$E(T) = E_{\max} + (E_{\max} - E_{\min})e^{-\alpha T}$	0.216			