Supplementary Information

3 Beneficial cyanosphere heterotrophs accelerate establishment of cyanobacterial biocrust

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6 Supplementary Tables

Laboratory trials on sterile soil substrates							
Treatments	Wetting events						
Treatments	0	3	6	9	12		
M. vaginatus	0.63 ± 0.17	2.59 ± 0.54	3.57 ± 0.36	3.81 ± 0.21	4.11 ± 0.31		
K12	0.53 ± 0.21	2.74 ± 0.21	2.76 ± 0.37	3.16 ± 0.13	3.02 ± 0.28		
O80	0.27 ± 0.18	2.98 ± 0.24	3.49 ± 0.34	6.57 ± 0.94	6.36 ± 0.69		
O64	0.54 ± 0.35	2.23 ± 0.45	3.04 ± 0.62	6.51 ± 0.28	4.97 ± 1.24		
METH4	0.45 ± 0.21	2.05 ± 0.34	3.56 ± 0.61	6.65 ± 1.22	6.36 ± 0.69		
Mixture	0.54 ± 0.21	2.05 ± 0.34	4.62 ± 0.45	7.15 ± 0.95	11.92 ± 1.58		

Table S1. Chlorophyll a areal concentration for laboratory trials of M. vaginatus grown alone and in co-culture with different heterotrophic isolates (Escherichia coli K12, Arthrobacter sp. O80, Bacillus sp. O64, Massilia sp. METH4) or as an equal proportion mixture of cyanosphere heterotrophs (Mixture) for 12 wetting events on sterile soil substrates given in mg Chl a m⁻² $(n=3; \pm \text{ standard deviation})$. Chl *a* values for uninoculated controls were below detection for the duration of experiment and are not shown.

Outdoor trials on hot desert soil substrates								
Treatments	Wetting events							
Treatments	0	3	6	9	12			
Control	0.3 ± 0.04	0.3 ± 0.1	0.3 ± 0.1	0.4 ± 0.1	0.6 ± 0.4			
Heterotrophs	0.3 ± 0.04	$0.3 \pm 0.04 \qquad 0.3 \pm 0.1$		0.4 ± 0.1	0.3 ± 0.1			
Cyanobacteria	4.7 ± 0.4	2.4 ± 0.8	5.3 ± 0.9	10.3 ± 4.9	14.3 ± 6.0			
Consortium	4.3 ± 0.7	$4.0 \pm 1.9 \qquad 20.0 \pm 5.7$		40.0 ± 5.9	37.4 ± 12.1			
Outdoor trials on cold desert soil substrates								
Treatments	Wetting events							
Treatments	0	3	6	9	12			
Control	6.9 ± 0.8	3.8 ± 0.3	7.4 ± 1.8	8.9 ± 1.2	10.7 ± 1.0			
Heterotrophs	7.1 ± 1.1	6.8 ± 1.4	10.3 ± 2.0	18.5 ± 4.7	15.5 ± 2.9			
Cyanobacteria	9.1 ± 1.5	12.9 ± 2.4	21.6 ± 6.0	15.4 ± 1.4	14.2 ± 2.3			
Consortium	9.4 ± 2.0	10.3 ± 2.8	21.9 ± 2.9	22.2 ± 3.9	17.5 ± 3.2			

Table S2. Chlorophyll *a* areal concentrations, given in mg Chl *a* m⁻² (n=5; ± standard deviation),

of unsterilized hot and cold desert soil substrates inoculated with beneficial heterotrophs

23 (Heterotrophs), pioneer cyanobacteria (Cyanobacteria), or with both beneficial heterotrophs and

²⁴ cyanobacteria (Consortium) under field conditions for 12 wetting events.

$ \begin{array}{ c c c c c c } \hline Cyanosphere Isolate \\ \hline Treatment \\ \hline \hline Cyanosphere Isolate \\ \hline \hline O64 \\ \hline O80 \\ \hline METH \\ \hline \hline Control \\ - \\ \hline - \\ \hline Control \\ - \\ \hline - \\ \hline - \\ \hline Control \\ \hline - \\ \hline - \\ \hline - \\ \hline Control \\ \hline - \\ \hline - \\ \hline - \\ \hline - \\ \hline Control \\ \hline - \\ \hline - \\ \hline - \\ \hline - \\ \hline Control \\ \hline - \\ \hline Control \\ \hline - \\ \hline Control \\ \hline - \hline$		Hot Desert				Cold D	Desert	
O64O80METH4O64O80METH4ControlControl-++++Heterotroph+++++++-Heterotroph-++++CyanobacteriaCyanobacteria-++++Consortium+++++-Consortium+++++Table S3. Presence of sequences matching those of the cyanosphere heterotrophs (<i>Bacillus</i> sp.O64, Arthrobacter sp. O80, Massilia sp. METH4) in final bacterial community for outdoor triafrom hot and cold desert locations (n=3). Crosses (+,++,+++) represent number of samples ineach treatment with detected cyanosphere isolate sequences, while dashes (-) represent their	Treatment	Cyanosphere Isolate			T ()	Cyanosphere Isolate		
Heterotroph++++-Heterotroph-+++++++CyanobacteriaCyanobacteria-++++-Consortium++++++++-Consortium++++++Table S3. Presence of sequences matching those of the cyanosphere heterotrophs (<i>Bacillus</i> sp.O64, Arthrobacter sp. O80, Massilia sp. METH4) in final bacterial community for outdoor triafrom hot and cold desert locations ($n=3$). Crosses (+,++,+++) represent number of samples ineach treatment with detected cyanosphere isolate sequences, while dashes (-) represent their		O64	080	METH4	- Treatment	O64	080	METH4
CyanobacteriaCyanobacteria-+++Consortium++++++-Consortium++++Table S3. Presence of sequences matching those of the cyanosphere heterotrophs (<i>Bacillus</i> sp.O64, Arthrobacter sp. O80, Massilia sp. METH4) in final bacterial community for outdoor triafrom hot and cold desert locations ($n=3$). Crosses (+,++,+++) represent number of samples ineach treatment with detected cyanosphere isolate sequences, while dashes (-) represent their	Control	-	-	-	Control	-	+++	-
Consortium++++++-Consortium+++++Table S3. Presence of sequences matching those of the cyanosphere heterotrophs (<i>Bacillus</i> sp.O64, <i>Arthrobacter</i> sp. O80, <i>Massilia</i> sp. METH4) in final bacterial community for outdoor triatfrom hot and cold desert locations ($n=3$). Crosses (+,++,+++) represent number of samples ineach treatment with detected cyanosphere isolate sequences, while dashes (-) represent their	Heterotroph	+++	+++	-	Heterotroph	-	+++	++
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O64, <i>Arthrobacter</i> sp. O80, <i>Massilia</i> sp. METH4) in final bacterial community for outdoor triated from hot and cold desert locations ($n=3$). Crosses (+,++,+++) represent number of samples in each treatment with detected cyanosphere isolate sequences, while dashes (-) represent their	Consortium	+++	+++	-	Consortium	+	+++	+
		-		-			•	
absence.	each treatment	with detect	ted cyanos	phere isolate	e sequences, wł	ile dashes	(-) represe	nt their
	absence.							

57 Supplementary Figures

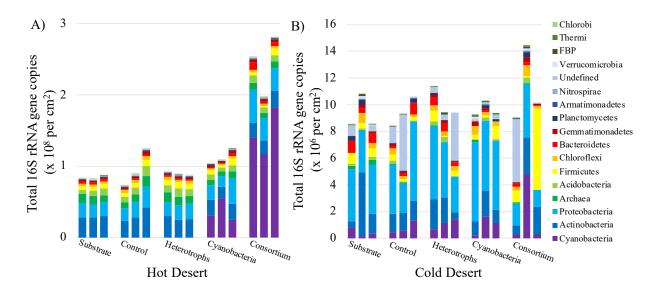




Figure S1. Bacterial abundance and community structure of original hot and cold desert soil
substrates (Substrate) and of unsterilized hot and cold desert soil substrates inoculated with
beneficial heterotrophs (Heterotrophs), *Microcoleus* spp. (Cyanobacteria), or with both beneficial
heterotrophs and cyanobacteria (Consortium) at phylum-level as determined by high-throughput
16S rRNA gene analysis coupled to qPCR. Treatments and uninoculated control (Control) were
subjected to 12 wetting events under field conditions.

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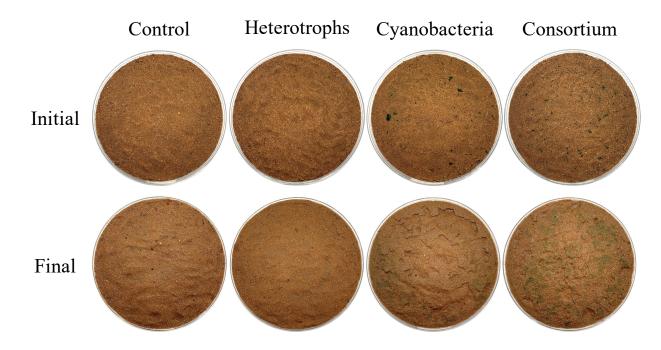


Figure S2. Visual aspect of unsterilized hot desert soil substrates inoculated with beneficial heterotrophs (Heterotrophs), Microcoleus spp. (Cyanobacteria), or both beneficial heterotrophs and cyanobacteria (Consortium), as well as uninoculated controls (Control) after incubation for 12 wetting events. Visible formation of biocrust can be seen at the final timepoint of the Cyanobacteria and Consortium treatments.

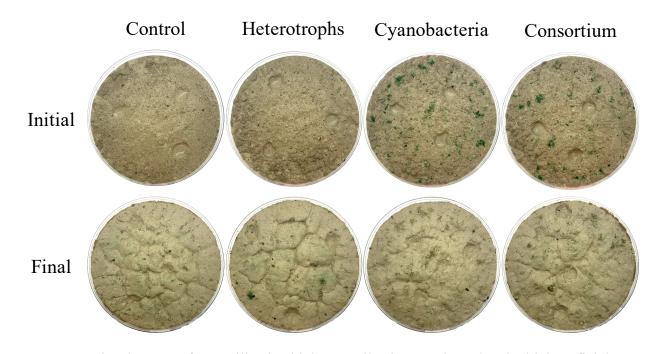


Figure S3. Visual aspect of unsterilized cold desert soil substrates inoculated with beneficial
heterotrophs (Heterotrophs), *Microcoleus* spp. (Cyanobacteria), or with both beneficial
heterotrophs and cyanobacteria (Consortium), as well as uninoculated controls (Control) after
incubation for 12 wetting events. Visible green coloring on Cyanobacteria and Consortium
treatments at initial timepoint is due to phototrophic biomass additions.