## Microbial community changes correlate with impaired host fitness of *Aurelia aurita* after environmental challenge

Nicola Diamavil#	, Cynthia M. Chibani	1 (:	COHEMA 1. + Level	Name (NA)	. D
Nicole Pinnow+,".	. Cynthia iyi. Chibani	+. Simon (	Juliert*, T. and	Nancy Welland	ı-Brauer+',"'

- <sup>1</sup> Kiel University, General Microbiology, Am Botanischen Garten 1-9, 24118 Kiel, Germany
- # Joint first authors
- <sup>†</sup> Current address: Sysmex Inostics GmbH, Falkenried 88, 20251 Hamburg, Germany
- \* Correspondence: nweiland@ifam.uni-kiel.de; Tel.: , 49-431-880-1649 (Germany)

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**Tab. S1: Pairwise result of Log-rank test for survival.** Log-rank test was used with the survival library (<a href="https://cran.r-project.org/web/packages/survival/index.html">https://cran.r-project.org/web/packages/survival/index.html</a>). Kaplan Meier test produced % of survival rates, chi-squares, and calculated the listed p-values for significance (< 0.05). Significance levels \*, < 0.05; \*\*, < 0.01; ns, not significant.

## fitness trait: survival

comparison		p-value	significance	
	native normal	sterile normal	0.300	ns
	native normal	native high salinity	0.250	ns
	native normal	native high temperature	0.003	**
	native normal	native future ocean	0.400	ns
	sterile normal	sterile high salinity	0.020	*
	sterile normal	sterile high temperature	0.010	*
	sterile normal	sterile future ocean	0.200	ns
	native high salinity	sterile high salinity	0.030	*
	native high temperature	sterile high temperature	0.600	ns
	native future ocean	sterile future ocean	0.200	ns

**Tabs. S2-S6: PERMANOVA tests.** Multivariate statistics were conducted using R for each fitness trait and the treatments growth, feeding, budding, segmentation, and ephyrae release for native and sterile conditions. PERMANOVA p-values are depicted with their corresponding significance levels \*, < 0.05; \*\*, < 0.01; \*\*\*,< 0.001; ns, not significant. PERMANOVA settings included Permutation test for adonis under reduced model, free Permutation, and 9999 No. of permutations adonis2.

Tab. S2 fitness trait: growth

comparison		p-value	significance	
	native normal	sterile normal	0.001	***
	native normal	native high salinity	0.001	***
	native normal	native high temperature	0.001	***
	native normal	native future ocean	0.001	***
	sterile normal	sterile high salinity	0.001	***
	sterile normal	sterile high temperature	0.001	***
	sterile normal	sterile future ocean	0.002	**
	native high salinity	sterile high salinity	0.001	***
	native high temperature	sterile high temperature	0.012	*
	native future ocean	sterile future ocean	0.042	*

Tab. S3 fitness trait: feeding

comparison			p-value	significance
	native normal	sterile normal	0.896	ns
	native normal	native high salinity	0.030	*
	native normal	native high temperature	0.036	*
	native normal	native future ocean	0.592	ns
	sterile normal	sterile high salinity	0.010	**
	sterile normal	sterile high temperature	0.010	**
	sterile normal	sterile future ocean	0.016	*
	native high salinity	sterile high salinity	0.596	ns
	native high temperature	sterile high temperature	0.498	ns
	native future ocean	sterile future ocean	0.110	ns

Tab. S4 fitness trait: budding

comparison		p-value	significance
native normal	sterile normal	0.001	***
native normal	native high salinity	0.001	***
native normal	native high temperature	0.001	***
native normal	native future ocean	0.001	***
sterile normal	sterile high salinity	0.001	***

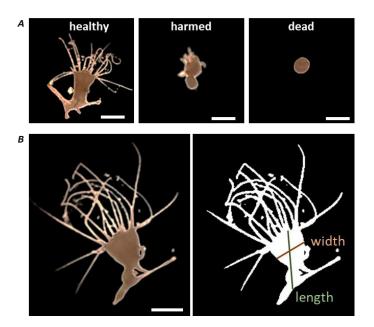
sterile normal	sterile high temperature	0.001	***
sterile normal	sterile future ocean	0.001	***
native high salinity	sterile high salinity	0.775	ns
native high temperature	sterile high temperature	0.001	***
native future ocean	sterile future ocean	0.010	**

Tab. S5 fitness trait: segmentation

comparison		p-value	significance	
	native normal	sterile normal	0.001	***
	native normal	native high salinity	0.001	***
	native normal	native high temperature	0.001	***
	native normal	native future ocean	0.001	***
	sterile normal	sterile high salinity	0.001	***
	sterile normal	sterile high temperature	0.001	***
	sterile normal	sterile future ocean	0.001	***
	native high salinity	sterile high salinity	0.002	**
	native high temperature	sterile high temperature	0.001	***
	native future ocean	sterile future ocean	0.007	**

Tab. S6 fitness trait: ephyrae release

comparison		p-value	significance	
	native normal	sterile normal	0.001	***
	native normal	native high salinity	0.001	***
	native normal	native high temperature	0.001	***
	native normal	native future ocean	0.001	***
	sterile normal	sterile high salinity	0.002	**
	sterile normal	sterile high temperature	0.001	***
	sterile normal	sterile future ocean	0.001	***
	native high salinity	sterile high salinity	0.003	**
	native high temperature	sterile high temperature	0.897	ns
	native future ocean	sterile future ocean	0.002	**



**Fig. S1:** Monitoring polyp fitness traits survival and growth. (*A*) Polyps categorized as healthy, harmed, and dead based on their phenotypical appearance. Healthy and harmed are alive polyps. (*B*) Example original photograph of an alive polyp (left panel) and its black/white projection (right panel) showing the axes for length and width measurement of the calyx.