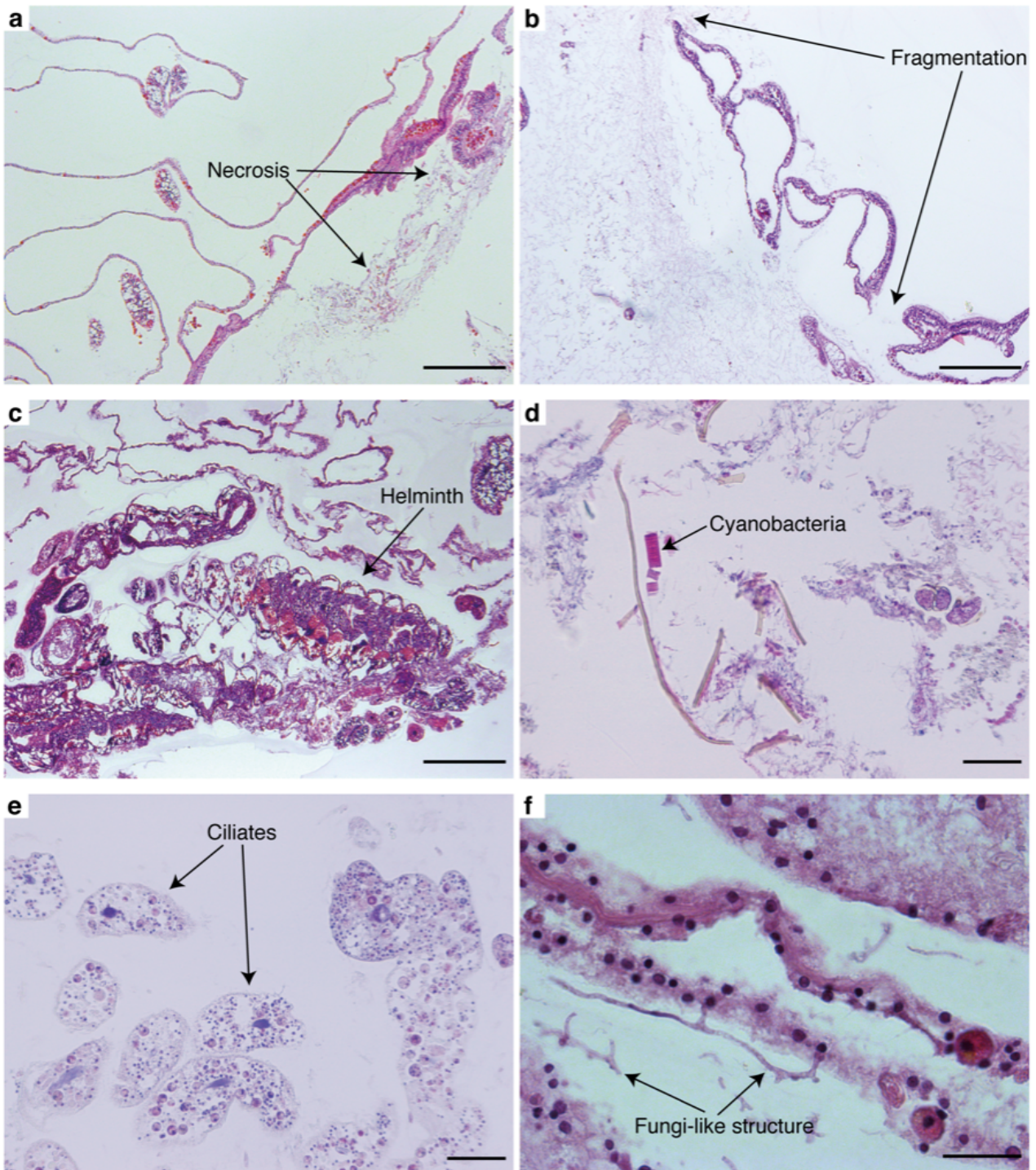
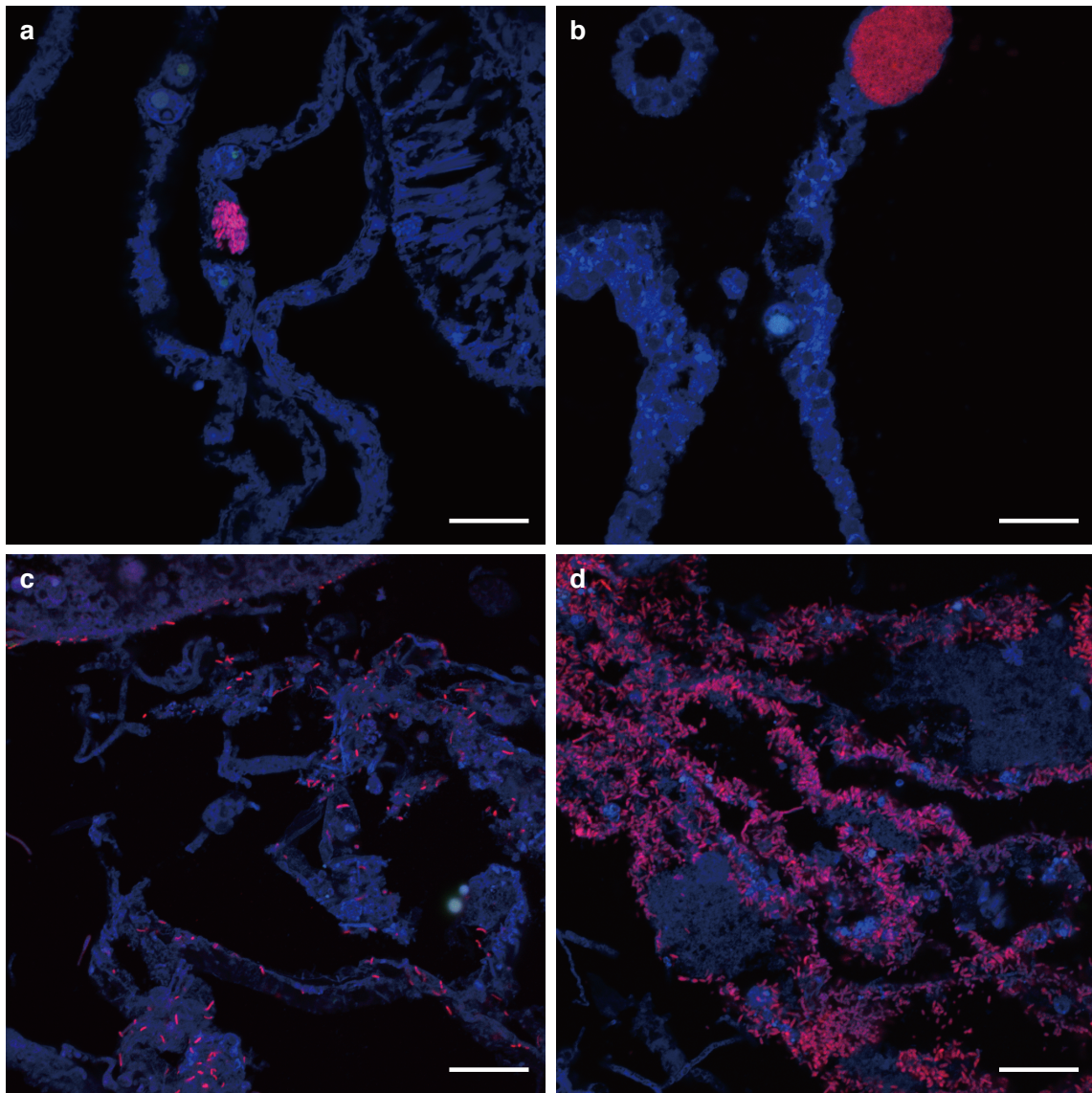


**ENERGY DEPLETION AND OPPORTUNISTIC MICROBIAL COLONISATION IN
WHITE SYNDROME LESIONS FROM CORALS ACROSS THE INDO-PACIFIC**

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Supplementary Figure S1. Histological images of a) tissue necrosis (scale bar = 200 μm); b) fragmentation (scale bar = 200 μm); c) helminths (scale bar = 200 μm); d) cyanobacteria (scale bar = 50 μm); e) ciliates (scale bar = 50 μm); and f) fungi-like structures (scale bar = 20 μm) detected in colonies of WS-diseased *Acropora hyacinthus*.



Supplementary Figure S2. Fluorescence in situ hybridisation micrographs of *Acropora hyacinthus* from a) HH tissue; b) HD tissue, D1 region; c) the lesion front, D2 region; and d) DD tissue, D3 region. Scale bars = 20 μm).

Supplementary Table S1. Comparison of proximate composition of samples types within locations. Values are presented as means \pm SEM. Values in the same row and group (i.e. location) that do not share the same superscripts are significantly different ($p < 0.05$).

<i>(mg g sample⁻¹)</i>	WA			PA			GBR		
	DD	HD	HH	DD	HD	HH	DD	HD	HH
Ash	958 \pm 4.67 ^a	942 \pm 5.48 ^b	910 \pm 14.6 ^c	964 \pm 1.61 ^a	945 \pm 9.71 ^{ab}	934 \pm 11 ^b	964 \pm 2.31 ^a	952 \pm 5.72 ^b	949 \pm 2.4 ^b
AFDW	158 \pm 24.9 ^b	285 \pm 34.3 ^a	276 \pm 55.3 ^{ab}	114 \pm 31 ^a	129 \pm 29.6 ^a	223 \pm 54.3 ^a	76.9 \pm 11.5 ^b	113 \pm 15.2 ^a	134 \pm 10.7 ^a
Lipid (<i>mg g AFDW⁻¹</i>)	6.89 \pm 1.93 ^b	17.2 \pm 3.5 ^a	23.6 \pm 2.74 ^a	4.15 \pm 1.17 ^b	7.78 \pm 3.18 ^{ab}	15.3 \pm 4.84 ^a	2.76 \pm 0.44 ^c	5.07 \pm 0.74 ^b	6.79 \pm 0.54 ^a

Supplementary Table S2. Comparison of proximate composition of sample types between locations. Values are presented as means \pm SEM. Values in the same row and group (i.e. sample type) that do not share the same superscripts are significantly different ($p < 0.05$).

<i>(mg g sample⁻¹)</i>	DD			HD			HH		
	GBR	PA	WA	GBR	PA	WA	GBR	PA	WA
Ash	964 \pm 2.31 ^a	964 \pm 1.61 ^{ab}	958 \pm 4.67 ^b	952 \pm 5.72 ^a	945 \pm 9.71 ^a	942 \pm 5.48 ^a	949 \pm 2.4 ^a	934 \pm 11 ^b	910 \pm 14.6 ^b
AFDW	76.9 \pm 11.5 ^b	114 \pm 31 ^{ab}	158 \pm 24.9 ^a	113 \pm 15.2 ^b	129 \pm 29.6 ^b	285 \pm 34.3 ^a	134 \pm 10.7 ^b	223 \pm 54.3 ^{ab}	276 \pm 55.3 ^a
Lipid (<i>mg g AFDW⁻¹</i>)	2.76 \pm 0.44 ^b	4.15 \pm 1.17 ^{ab}	6.89 \pm 1.93 ^a	5.07 \pm 0.74 ^b	7.78 \pm 3.18 ^b	17.2 \pm 3.5 ^a	6.79 \pm 0.54 ^b	15.3 \pm 4.84 ^{ab}	23.6 \pm 2.74 ^a

Supplementary Table S3. Comparison of lipid class composition (mg g lipid⁻¹) between sample types within locations. Values are presented as means \pm SEM. Values in the same row and group (i.e. location) that do not share the same superscripts are significantly different ($p < 0.05$).

Class	WA			Palmyra			GBR		
	DD	HD	HH	DD	HD	HH	DD	HD	HH
WAX	83.6 \pm 13.7 ^a	86.9 \pm 15.9 ^a	114 \pm 75.5 ^a	82 \pm 18.2 ^a	76.4 \pm 13 ^a	62.5 \pm 21.2 ^a	82.2 \pm 7.04 ^b	76.5 \pm 11 ^b	139 \pm 20.9 ^a
TAG	344 \pm 131 ^a	492 \pm 77.7 ^a	462 \pm 169 ^a	97.7 \pm 42.5 ^a	110 \pm 44.9 ^a	325 \pm 150 ^a	56.1 \pm 6.5 ^a	73.9 \pm 21.8 ^a	117 \pm 32.2 ^a
FFA	24 \pm 3.23 ^a	30.8 \pm 4.36 ^a	17.9 \pm 6.13 ^a	26.3 \pm 7.32 ^a	26.6 \pm 7.29 ^a	62.3 \pm 32 ^a	20.4 \pm 3.13 ^b	25.1 \pm 4.47 ^b	38.4 \pm 5.05 ^a
1,2DAG	17.5 \pm 17.5 ^a	0 \pm 0 ^a	9.34 \pm 9.34 ^a	11.5 \pm 11.5 ^a	35.4 \pm 20.4 ^a	7.93 \pm 7.93 ^a	105 \pm 6.91 ^a	45.4 \pm 10.1 ^b	32.3 \pm 11.6 ^b
STEROL	43.6 \pm 6.14 ^a	33.2 \pm 3.44 ^a	44.7 \pm 9.95 ^a	65.9 \pm 17 ^a	61.3 \pm 12.9 ^a	71.5 \pm 38.3 ^a	64.2 \pm 5.49 ^a	66.4 \pm 3.42 ^a	68.2 \pm 4 ^a
AMPL	353 \pm 93.8 ^a	250 \pm 66.5 ^a	229 \pm 108 ^a	446 \pm 103 ^a	469 \pm 33.9 ^a	370 \pm 132 ^a	414 \pm 35.8 ^a	461 \pm 33.9 ^a	380 \pm 37.6 ^a
PE	46.7 \pm 16 ^a	42.6 \pm 7.48 ^a	60.3 \pm 32.2 ^a	104 \pm 29.7 ^a	76.7 \pm 15.2 ^a	60.9 \pm 16.9 ^a	113 \pm 6.64 ^a	90.7 \pm 5.49 ^b	74.1 \pm 6.05 ^b
PSPI	10.4 \pm 10.4 ^a	11.8 \pm 11.8 ^a	5.67 \pm 5.67 ^a	0 \pm 0 ^a	39.1 \pm 27.2 ^a	4.29 \pm 4.29 ^a	0 \pm 0 ^a	27.7 \pm 16.1 ^a	18.7 \pm 9.96 ^a
PC	76.7 \pm 11.4 ^a	53.3 \pm 9.46 ^a	56.9 \pm 16.9 ^a	127 \pm 35.8 ^a	103 \pm 21.9 ^{ab}	35.3 \pm 15.3 ^b	135 \pm 6.83 ^a	133 \pm 6.45 ^a	125 \pm 8.68 ^a
LPC	0 \pm 0	0 \pm 0	0 \pm 0	39.2 \pm 28.9 ^a	3.18 \pm 3.18 ^a	0 \pm 0 ^a	10.8 \pm 7.37 ^a	0 \pm 0 ^a	7.03 \pm 7.03 ^a
ΣSTORAGE	469 \pm 115 ^a	609 \pm 84.1 ^a	604 \pm 161 ^a	218 \pm 45.6 ^a	248 \pm 41.6 ^a	458 \pm 147 ^a	263 \pm 21.2 ^{ab}	221 \pm 35.2 ^b	327 \pm 38 ^a
ΣSTRUCTURAL	531 \pm 115 ^a	391 \pm 84.1 ^a	396 \pm 161 ^a	782 \pm 45.6 ^a	752 \pm 41.6 ^a	542 \pm 147 ^a	737 \pm 21.2 ^{ab}	779 \pm 35.2 ^a	673 \pm 38 ^b
RATIO	1.38 \pm 0.79 ^a	2.01 \pm 0.75 ^a	2.55 \pm 1.36 ^a	0.29 \pm 0.08 ^a	0.34 \pm 0.08 ^a	1.41 \pm 0.72 ^a	0.37 \pm 0.04 ^{ab}	0.32 \pm 0.09 ^b	0.54 \pm 0.11 ^a

Supplementary Table S4. Comparison of lipid class composition (mg g lipid⁻¹) of sample types between locations. Values are presented as means \pm SEM. Values in the same row and group (i.e. sample type) that do not share the same superscripts are significantly different ($p < 0.05$).

Class	DD			HD			HH		
	GBR	Palmyra	WA	GBR	Palmyra	WA	GBR	Palmyra	WA
WAX	82.2 \pm 7.04 ^a	82 \pm 18.2 ^a	83.6 \pm 13.7 ^a	76.5 \pm 11 ^a	76.4 \pm 13 ^a	86.9 \pm 15.9 ^a	139 \pm 20.9 ^a	62.5 \pm 21.2 ^b	114 \pm 75.5 ^{ab}
TAG	56.1 \pm 6.5 ^b	97.7 \pm 42.5 ^b	344 \pm 131 ^a	73.9 \pm 21.8 ^b	110 \pm 44.9 ^b	492 \pm 77.7 ^a	117 \pm 32.2 ^b	325 \pm 150 ^{ab}	462 \pm 169 ^a
FFA	20.4 \pm 3.13 ^a	26.3 \pm 7.32 ^a	24 \pm 3.23 ^a	25.1 \pm 4.47 ^a	26.6 \pm 7.29 ^a	30.8 \pm 4.36 ^a	38.4 \pm 5.05 ^a	62.3 \pm 32 ^a	17.9 \pm 6.13 ^b
1,2DAG	105 \pm 6.91 ^a	11.5 \pm 11.5 ^b	17.5 \pm 17.5 ^b	45.4 \pm 10.1 ^a	35.4 \pm 20.4 ^{ab}	0 \pm 0 ^b	32.3 \pm 11.6 ^a	7.93 \pm 7.93 ^a	9.34 \pm 9.34 ^a
STEROL	64.2 \pm 5.49 ^a	65.9 \pm 17 ^a	43.6 \pm 6.14 ^a	66.4 \pm 3.42 ^a	61.3 \pm 12.9 ^a	33.2 \pm 3.44 ^b	68.2 \pm 4 ^a	71.5 \pm 38.3 ^a	44.7 \pm 9.95 ^a
AMPL	414 \pm 35.8 ^a	446 \pm 103 ^a	353 \pm 93.8 ^a	461 \pm 33.9 ^a	469 \pm 33.9 ^a	250 \pm 66.5 ^b	380 \pm 37.6 ^a	370 \pm 132 ^a	229 \pm 108 ^a
PE	113 \pm 6.64 ^a	104 \pm 29.7 ^{ab}	46.7 \pm 16 ^b	90.7 \pm 5.49 ^a	76.7 \pm 15.2 ^a	42.6 \pm 7.48 ^b	74.1 \pm 6.05 ^a	60.9 \pm 16.9 ^a	60.3 \pm 32.2 ^a
PSPI	0 \pm 0 ^a	0 \pm 0 ^a	10.4 \pm 10.4 ^a	27.7 \pm 16.1 ^a	39.1 \pm 27.2 ^a	11.8 \pm 11.8 ^a	18.7 \pm 9.96 ^a	4.29 \pm 4.29 ^a	5.67 \pm 5.67 ^a
PC	135 \pm 6.83 ^a	127 \pm 35.8 ^{ab}	76.7 \pm 11.4 ^b	133 \pm 6.45 ^a	103 \pm 21.9 ^{ab}	53.3 \pm 9.46 ^b	125 \pm 8.68 ^a	35.3 \pm 15.3 ^b	56.9 \pm 16.9 ^b
LPC	10.8 \pm 7.37 ^a	39.2 \pm 28.9 ^a	0 \pm 0 ^a	0 \pm 0 ^a	3.18 \pm 3.18 ^a	0 \pm 0 ^a	7.03 \pm 7.03 ^a	0 \pm 0 ^a	0 \pm 0 ^a
Σ STORAGE	263 \pm 21.2 ^b	218 \pm 45.6 ^b	469 \pm 115 ^a	221 \pm 35.2 ^b	248 \pm 41.6 ^b	609 \pm 84.1 ^a	327 \pm 38 ^a	458 \pm 147 ^a	604 \pm 161 ^a
Σ STRUCTURAL	737 \pm 21.2 ^a	782 \pm 45.6 ^a	531 \pm 115 ^b	779 \pm 35.2 ^a	752 \pm 41.6 ^a	391 \pm 84.1 ^b	673 \pm 38 ^a	542 \pm 147 ^a	396 \pm 161 ^a
RATIO	0.37 \pm 0.04 ^b	0.29 \pm 0.08 ^b	1.38 \pm 0.79 ^a	0.32 \pm 0.09 ^b	0.34 \pm 0.08 ^b	2.01 \pm 0.75 ^a	0.54 \pm 0.11 ^a	1.41 \pm 0.72 ^a	2.55 \pm 1.36 ^a

Supplementary Table S5. Comparison of fatty acid composition (mg g lipid⁻¹) between sample types within locations. Values are presented as means \pm SEM. Values in the same row and group (i.e. location) that do not share the same superscripts are significantly different ($p < 0.05$).

Fatty acid	WA			Palmyra			GBR		
	DD	HD	HH	DD	HD	HH	DD	HD	HH
6:0	0.15 \pm 0.07 ^a	0.06 \pm 0.04 ^a	0.02 \pm 0.02 ^a	0.13 \pm 0.13 ^a	0.16 \pm 0.1 ^a	0.09 \pm 0.05 ^a	0.39 \pm 0.12 ^a	0.14 \pm 0.05 ^a	0.13 \pm 0.03 ^a
8:0	0.01 \pm 0.01 ^a	0.03 \pm 0.02 ^a	0 \pm 0 ^a						
10:0	0.07 \pm 0.03 ^b	0.27 \pm 0.12 ^a	0.14 \pm 0.04 ^{ab}	0 \pm 0 ^a	0.02 \pm 0.02 ^a	0.03 \pm 0.02 ^a	0.01 \pm 0.01 ^b	0.06 \pm 0.03 ^b	2.14 \pm 0.94 ^a
11:0	0.01 \pm 0.01 ^a	0 \pm 0 ^a	0.04 \pm 0.02 ^a	0 \pm 0 ^a	0.02 \pm 0.02 ^a	0.01 \pm 0.01 ^a	0 \pm 0 ^a	0 \pm 0 ^a	0.02 \pm 0.02 ^a
12:0	0.56 \pm 0.16 ^a	0.55 \pm 0.03 ^a	0.77 \pm 0.16 ^a	0.34 \pm 0.2 ^a	0.48 \pm 0.05 ^a	0.32 \pm 0.04 ^a	1.05 \pm 0.26 ^a	0.88 \pm 0.12 ^a	1.04 \pm 0.13 ^a
13:0	0.21 \pm 0.05 ^a	0.16 \pm 0.03 ^a	0.16 \pm 0.09 ^a	0.13 \pm 0.07 ^a	0.32 \pm 0.07 ^a	0.26 \pm 0.11 ^a	0.08 \pm 0.04 ^b	0.25 \pm 0.03 ^a	0.26 \pm 0.07 ^a
14:0	14 \pm 3.95 ^a	19.9 \pm 2.18 ^a	16.6 \pm 4.28 ^a	10.1 \pm 3.86 ^a	14.7 \pm 2.39 ^a	18.6 \pm 5.14 ^a	7.39 \pm 0.72 ^b	8.88 \pm 0.87 ^{ab}	12.8 \pm 2.15 ^a
15:0	0.34 \pm 0.1 ^a	0.22 \pm 0.01 ^a	0.15 \pm 0.03 ^a	0.25 \pm 0.11 ^a	0.31 \pm 0.05 ^a	0.35 \pm 0.17 ^a	0.64 \pm 0.09 ^a	0.4 \pm 0.05 ^b	0.39 \pm 0.04 ^b
16:0	160 \pm 53.4 ^a	221 \pm 45 ^a	201 \pm 53 ^a	97.7 \pm 26.3 ^a	114 \pm 19.9 ^a	165 \pm 44.1 ^a	76.2 \pm 7.47 ^b	82.1 \pm 9.97 ^b	121 \pm 15.6 ^a
17:0	0.28 \pm 0.14 ^a	0.52 \pm 0.14 ^a	0.46 \pm 0.16 ^a	1.83 \pm 1.03 ^a	0.31 \pm 0.09 ^a	0.44 \pm 0.23 ^a	1.68 \pm 0.31 ^a	0.56 \pm 0.12 ^b	0.44 \pm 0.13 ^b
18:0	17.7 \pm 3.61 ^a	23.6 \pm 3.88 ^a	22.3 \pm 3.31 ^a	32.3 \pm 9.23 ^a	25.7 \pm 3.42 ^a	30.3 \pm 5.89 ^a	25.3 \pm 3.71 ^a	28.2 \pm 1.5 ^a	27.9 \pm 1.23 ^a
20:0	3.12 \pm 0.68 ^a	3.7 \pm 0.22 ^a	3.2 \pm 0.42 ^a	3.62 \pm 0.88 ^a	3.54 \pm 0.48 ^a	4.36 \pm 0.94 ^a	3.91 \pm 0.33 ^a	3.66 \pm 0.2 ^a	3.82 \pm 0.19 ^a
21:0	0.48 \pm 0.22 ^a	0.22 \pm 0.06 ^a	0.18 \pm 0.05 ^a	1.16 \pm 0.63 ^a	0.37 \pm 0.04 ^a	0.27 \pm 0.06 ^a	0.99 \pm 0.17 ^a	0.54 \pm 0.06 ^b	0.39 \pm 0.05 ^b
22:0	1.45 \pm 0.33 ^a	1.25 \pm 0.46 ^a	1.16 \pm 0.34 ^a	7.39 \pm 4 ^a	1.95 \pm 0.15 ^a	1.38 \pm 0.52 ^a	5.69 \pm 1.11 ^a	2.8 \pm 0.34 ^b	2.34 \pm 0.43 ^b
24:0	0.64 \pm 0.13 ^a	0.43 \pm 0.11 ^a	0.37 \pm 0.11 ^a	3.26 \pm 1.66 ^a	0.72 \pm 0.12 ^a	0.56 \pm 0.26 ^a	2.81 \pm 0.66 ^a	1.25 \pm 0.2 ^b	0.88 \pm 0.16 ^b
14:1n-5	0.4 \pm 0.19 ^a	0.66 \pm 0.29 ^a	0.41 \pm 0.36 ^a	0.4 \pm 0.27 ^a	0.83 \pm 0.24 ^a	0.26 \pm 0.09 ^a	1.64 \pm 0.6 ^a	1.77 \pm 0.34 ^a	1.43 \pm 0.33 ^a
15:1n-5	0.48 \pm 0.12 ^a	0.68 \pm 0.14 ^a	0.67 \pm 0.1 ^a	0.18 \pm 0.11 ^a	0.56 \pm 0.07 ^a	0.32 \pm 0.15 ^a	1 \pm 0.35 ^a	1.08 \pm 0.23 ^a	1.05 \pm 0.25 ^a
16:1n-7	14.6 \pm 6.47 ^a	19.3 \pm 6.86 ^a	16.6 \pm 5.8 ^a	2.82 \pm 1.08 ^a	4.65 \pm 0.61 ^a	6.29 \pm 1.56 ^a	3.05 \pm 0.25 ^a	3.3 \pm 0.28 ^a	5.78 \pm 1.32 ^a
17:1n-7	23 \pm 6.88 ^a	10.9 \pm 10.9 ^a	3.99 \pm 3.99 ^a	6.88 \pm 5.2 ^a	17.6 \pm 4.67 ^a	19.4 \pm 8.83 ^a	8.62 \pm 0.94 ^a	11.6 \pm 1.9 ^a	17.9 \pm 5.19 ^a
18:1n-7	1.25 \pm 0.3 ^a	1.09 \pm 0.31 ^a	0.98 \pm 0.29 ^a	0.32 \pm 0.11 ^b	0.46 \pm 0.06 ^b	2.41 \pm 1.38 ^a	1.4 \pm 0.27 ^a	2.07 \pm 1.48 ^b	0.61 \pm 0.08 ^b
18:1n-7t	0.03 \pm 0.02 ^b	0.04 \pm 0.04 ^b	0.5 \pm 0.2 ^a	0 \pm 0 ^a	0.07 \pm 0.04 ^a	0.19 \pm 0.12 ^a	0.01 \pm 0.01 ^a	0.05 \pm 0.05 ^a	0.1 \pm 0.06 ^a
18:1n-9	23.6 \pm 9.22 ^a	31 \pm 10.5 ^a	31.3 \pm 10.2 ^a	6.82 \pm 1.31 ^b	6.89 \pm 1.13 ^b	13.8 \pm 2.38 ^a	7.71 \pm 0.55 ^b	13 \pm 5.89 ^b	14.4 \pm 2.71 ^a
18:1n-9t	0.01 \pm 0.01 ^b	0.24 \pm 0.11 ^a	0.51 \pm 0.24 ^a	0 \pm 0 ^b	0.11 \pm 0.09 ^b	1.43 \pm 1.16 ^a	0.11 \pm 0.11 ^a	4.57 \pm 2.98 ^a	0.22 \pm 0.13 ^a

20:1n-9	7.81 ± 2.51 ^a	5.45 ± 3.63 ^a	9.8 ± 2.13 ^a	2.12 ± 0.43 ^a	3.05 ± 0.61 ^a	4.48 ± 1.45 ^a	2.93 ± 0.27 ^b	3.82 ± 0.4 ^b	5.57 ± 0.59 ^a
22:1n-9	0.52 ± 0.11 ^a	0.38 ± 0.03 ^a	0.72 ± 0.22 ^a	4.37 ± 1.99 ^a	1.53 ± 0.33 ^{ab}	1.43 ± 0.66 ^b	2.56 ± 0.62 ^a	1.2 ± 0.28 ^b	0.96 ± 0.34 ^b
24:1n-9	0.18 ± 0.12 ^a	0.07 ± 0.02 ^a	0.11 ± 0.05 ^a	0 ± 0 ^a	0.08 ± 0.06 ^a	0.05 ± 0.04 ^a	0.04 ± 0.03 ^a	0.01 ± 0.01 ^a	0.07 ± 0.03 ^a
20:1n-11	0.46 ± 0.25 ^a	0.48 ± 0.27 ^a	0.64 ± 0.22 ^a						
22:1n-11	0.23 ± 0.07 ^a	0.15 ± 0.04 ^a	0.15 ± 0.03 ^a	0.13 ± 0.09 ^a	0.12 ± 0.06 ^a	0.09 ± 0.03 ^a	0.28 ± 0.1 ^a	0.24 ± 0.14 ^a	0.13 ± 0.02 ^a
18:3n-3	0.21 ± 0.05 ^a	0.15 ± 0.04 ^a	0.13 ± 0.02 ^a	0.02 ± 0.02 ^a	0.02 ± 0.02 ^a	0.06 ± 0.04 ^a	0.31 ± 0.12 ^a	0.16 ± 0.09 ^a	0.1 ± 0.03 ^a
18:4n-3	5.14 ± 1.83 ^a	9.04 ± 0.84 ^a	7.84 ± 1.34 ^a	4.71 ± 0.37 ^a	7.99 ± 1.67 ^a	6.14 ± 1.27 ^a	3.8 ± 0.46 ^b	6.43 ± 0.7 ^a	6.66 ± 0.61 ^a
20:4n-3	0.44 ± 0.17 ^a	0.7 ± 0.16 ^a	0.46 ± 0.23 ^a	0 ± 0 ^a	0.12 ± 0.12 ^a	0.43 ± 0.25 ^a	0.08 ± 0.05 ^b	0.22 ± 0.08 ^{ab}	0.42 ± 0.1 ^a
20:5n-3	15.8 ± 3.82 ^a	24.2 ± 2.73 ^a	23.8 ± 0.68 ^a	18.6 ± 3.17 ^a	29 ± 5.85 ^a	19.1 ± 3.8 ^a	13.9 ± 1.28 ^c	22.7 ± 1.24 ^b	26 ± 1.32 ^a
22:3n-3	0 ± 0 ^a	3.29 ± 3.29 ^a	0 ± 0 ^a				0 ± 0 ^a	2.74 ± 1.86 ^a	3.99 ± 2.04 ^a
22:5n-3	4.24 ± 0.96 ^a	5.88 ± 0.22 ^a	5.77 ± 0.61 ^a	8.73 ± 3.38 ^a	7.08 ± 1.24 ^a	4.82 ± 0.69 ^a	5.33 ± 0.65 ^b	6.46 ± 0.6 ^a	7.36 ± 0.57 ^a
22:6n-3	18.5 ± 6.26 ^a	27.5 ± 4.93 ^a	23.8 ± 5.87 ^a	7.83 ± 2.06 ^b	13.4 ± 2.28 ^{ab}	15.1 ± 3.56 ^a	6.18 ± 0.69 ^b	9.55 ± 0.56 ^a	10.3 ± 0.68 ^a
24:5n-3	0 ± 0 ^a	0.1 ± 0.08 ^a	0.01 ± 0.01 ^a	0 ± 0 ^a	0.02 ± 0.02 ^a	0.03 ± 0.02 ^a	0 ± 0 ^a	0 ± 0 ^a	0.03 ± 0.03 ^a
24:6n-3	0.05 ± 0.04 ^a	0.11 ± 0 ^a	0.09 ± 0.02 ^a	0 ± 0 ^b	0.05 ± 0.03 ^{ab}	0.09 ± 0.03 ^a	0.03 ± 0.03 ^a	0 ± 0 ^a	0.02 ± 0.01 ^a
16:2n-4	1.19 ± 0.4 ^a	0.03 ± 0.03 ^a	1.26 ± 0.63 ^a	1.92 ± 0.57 ^a	0.56 ± 0.56 ^{ab}	0.27 ± 0.27 ^b	1.78 ± 0.18 ^a	1.39 ± 0.49 ^{ab}	0.57 ± 0.37 ^b
16:3n-4	1.17 ± 0.73 ^a	3.14 ± 0.72 ^a	2.73 ± 1.24 ^a	0 ± 0 ^b	0.73 ± 0.42 ^{ab}	1.81 ± 0.84 ^a	0.04 ± 0.04 ^b	0.18 ± 0.12 ^{ab}	1.23 ± 0.65 ^a
18:2n-6	7.17 ± 2.28 ^a	9.42 ± 1.51 ^a	7.78 ± 1.75 ^a	2.63 ± 0.58 ^a	2.63 ± 0.41 ^a	3.91 ± 1.07 ^a	2.64 ± 0.22 ^{ab}	2.69 ± 0.24 ^b	3.68 ± 0.48 ^a
18:2n-6t	0.35 ± 0.07 ^a	0.01 ± 0.01 ^c	0.05 ± 0.01 ^b	0 ± 0 ^a	0 ± 0 ^a	0.04 ± 0.02 ^a	0.08 ± 0.06 ^a	0 ± 0 ^a	0.06 ± 0.05 ^a
18:3n-6	29.6 ± 11.5 ^a	43.3 ± 9.99 ^a	37.7 ± 12.2 ^a	8.44 ± 3.03 ^a	14.4 ± 2.6 ^a	19.3 ± 5.71 ^a	7.94 ± 0.89 ^c	10.5 ± 0.75 ^b	18.7 ± 3.05 ^a
20:2n-6	1.71 ± 0.37 ^a	1.85 ± 0.13 ^a	1.74 ± 0.18 ^a	0.89 ± 0.14 ^a	1.22 ± 0.17 ^a	1.18 ± 0.19 ^a	1.01 ± 0.09 ^b	1.32 ± 0.09 ^a	1.18 ± 0.08 ^{ab}
20:3n-6	3.71 ± 1.13 ^a	5.2 ± 1.19 ^a	4.48 ± 1.23 ^a	1.21 ± 0.7 ^a	1.22 ± 0.53 ^a	2.67 ± 1 ^a	0.31 ± 0.17 ^c	0.89 ± 0.19 ^b	1.76 ± 0.32 ^a
20:4n-6	12.8 ± 1.9 ^a	13.5 ± 2.07 ^a	15.1 ± 3.24 ^a	16.1 ± 3.85 ^a	19.2 ± 2.37 ^a	10.8 ± 1.79 ^a	18 ± 1.75 ^b	27.2 ± 1.75 ^a	26.7 ± 1.07 ^a
22:2n-6	0.03 ± 0.03 ^a	0.04 ± 0.02 ^a	0.03 ± 0 ^a	0 ± 0 ^b	0 ± 0 ^b	0.05 ± 0.03 ^a	0.05 ± 0.04 ^a	0.02 ± 0.02 ^a	0.01 ± 0.01 ^a
22:4n-6	6.06 ± 1.01 ^a	4.42 ± 2.31 ^a	8.75 ± 2.17 ^a	7.89 ± 1.68 ^a	9.4 ± 1.23 ^a	6.86 ± 1.37 ^a	7.28 ± 0.92 ^a	10.2 ± 1.63 ^a	8.24 ± 1.88 ^a
22:5n-6	0.14 ± 0.14 ^a	0 ± 0 ^a	0.02 ± 0.02 ^a				0.02 ± 0.02 ^a	0 ± 0 ^a	0.03 ± 0.02 ^a
Total	380 ± 115 ^a	495 ± 81.4 ^a	454 ± 91.2 ^a	261 ± 49.9 ^a	306 ± 40.1 ^a	364 ± 71.2 ^a	224 ± 19.8 ^b	275 ± 18.4 ^a	339 ± 29.2 ^a
ΣSFA	199 ± 60.2 ^a	272 ± 43.1 ^a	246 ± 57.2 ^a	158 ± 36 ^a	163 ± 25.4 ^a	222 ± 55.4 ^a	126 ± 13.6 ^b	130 ± 11.9 ^b	174 ± 19.5 ^a
ΣMUFA	72.6 ± 25.3 ^a	70.5 ± 23.6 ^a	66.4 ± 14.9 ^a	24 ± 6.94 ^b	35.9 ± 7 ^{ab}	50.2 ± 4.82 ^a	29.3 ± 2.37 ^b	42.7 ± 10.6 ^b	48.2 ± 6.19 ^a
ΣPUFA	108 ± 30.3 ^a	152 ± 18 ^a	142 ± 19.7 ^a	78.9 ± 11.3 ^a	107 ± 14.3 ^a	92.7 ± 17.2 ^a	68.8 ± 5.82 ^b	103 ± 3.92 ^a	117 ± 6.42 ^a
ΣTRANS	0.39 ± 0.05 ^a	0.29 ± 0.09 ^a	1.06 ± 0.43 ^a	0 ± 0 ^b	0.18 ± 0.12 ^{ab}	1.66 ± 1.13 ^a	0.2 ± 0.11 ^a	4.61 ± 2.98 ^a	0.38 ± 0.18 ^a

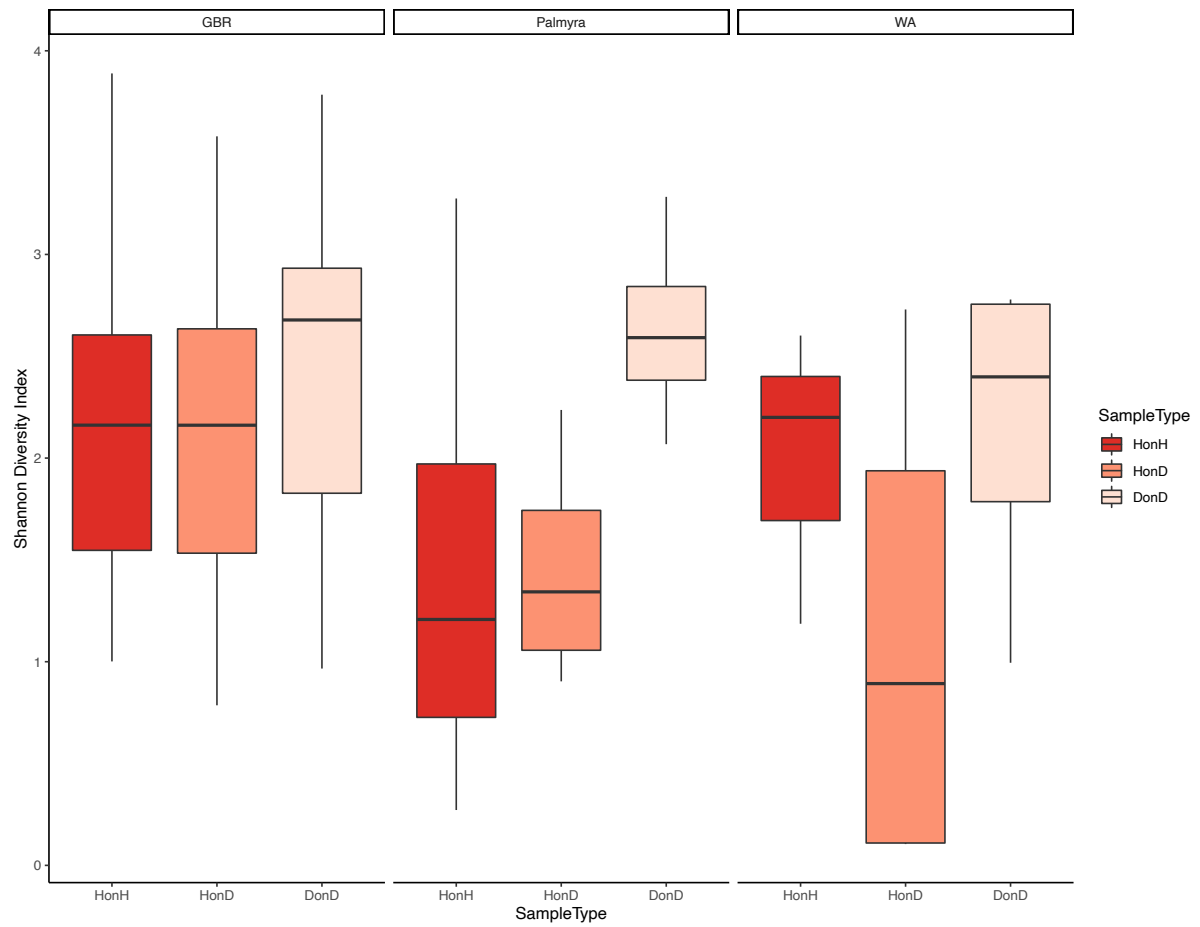
Σ n-3 PUFA	44.4 ± 12.7 ^a	70.9 ± 8.06 ^a	61.9 ± 7.4 ^a	39.9 ± 6.65 ^a	57.6 ± 10.3 ^a	45.7 ± 9.06 ^a	29.7 ± 2.74 ^b	48.2 ± 2.42 ^a	54.8 ± 3.58 ^a
Σ n-3 LC PUFA	39.1 ± 11 ^a	61.7 ± 8.09 ^a	54 ± 6.05 ^a	35.1 ± 6.31 ^a	49.6 ± 8.68 ^a	39.5 ± 7.83 ^a	25.6 ± 2.23 ^b	41.6 ± 2.16 ^a	48.1 ± 3.27 ^a
Σ n-6 PUFA	61.5 ± 16.7 ^a	77.7 ± 11 ^a	75.7 ± 10.6 ^a	37.1 ± 5.73 ^a	48.1 ± 4.37 ^a	44.8 ± 7.47 ^a	37.3 ± 3.21 ^b	52.8 ± 2.86 ^a	60.4 ± 3.84 ^a
Σ n-6 LC PUFA	22.7 ± 3.83 ^a	23.1 ± 0.61 ^a	28.4 ± 4.33 ^a	25.2 ± 5.5 ^a	29.8 ± 3.32 ^a	20.4 ± 2.5 ^a	25.6 ± 2.54 ^b	38.3 ± 2.75 ^a	36.8 ± 2.11 ^a
EPA:DHA	1.01 ± 0.16 ^a	0.94 ± 0.17 ^a	1.18 ± 0.36 ^a	2.81 ± 0.66 ^a	2.21 ± 0.3 ^{ab}	1.36 ± 0.23 ^b	2.38 ± 0.23 ^a	2.4 ± 0.1 ^a	2.57 ± 0.11 ^a
EPA:ARA	1.19 ± 0.2 ^a	1.83 ± 0.18 ^a	1.76 ± 0.43 ^a	1.28 ± 0.19 ^a	1.51 ± 0.23 ^a	1.83 ± 0.28 ^a	0.79 ± 0.04 ^b	0.86 ± 0.06 ^{ab}	0.98 ± 0.05 ^a
n-3:n-6	0.71 ± 0.02 ^b	0.93 ± 0.09 ^a	0.82 ± 0.02 ^a	1.12 ± 0.16 ^a	1.17 ± 0.12 ^a	1.01 ± 0.03 ^a	0.81 ± 0.04 ^a	0.94 ± 0.06 ^a	0.93 ± 0.07 ^a

Supplementary Table S6. Comparison of fatty acid composition (mg g lipid⁻¹) of sample types between locations. Values are presented as means ± SEM. Values in the same row and group (i.e. sample type) that do not share the same superscripts are significantly different ($p < 0.05$).

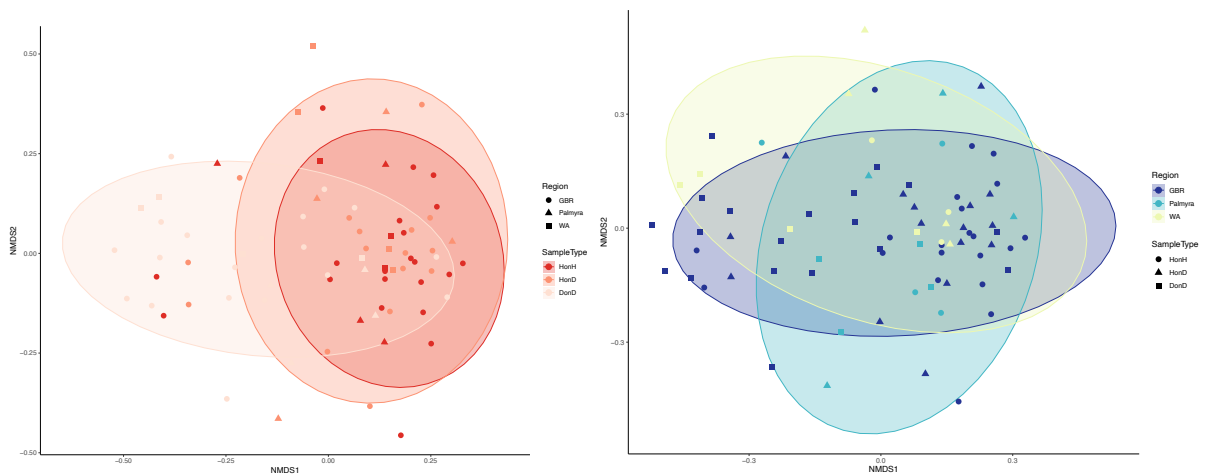
Fatty acid	HH			HD			DD		
	GBR	Palmyra	WA	GBR	Palmyra	WA	GBR	Palmyra	WA
6:0	0.13 ± 0.03 ^a	0.09 ± 0.05 ^a	0.02 ± 0.02 ^a	0.14 ± 0.05 ^a	0.16 ± 0.1 ^a	0.06 ± 0.04 ^a	0.39 ± 0.12 ^a	0.13 ± 0.13 ^a	0.15 ± 0.07 ^a
8:0				0 ± 0 ^b	0 ± 0 ^b	0.03 ± 0.02 ^a	0 ± 0 ^a	0 ± 0 ^a	0.01 ± 0.01 ^a
10:0	2.14 ± 0.94 ^a	0.03 ± 0.02 ^a	0.14 ± 0.04 ^a	0.06 ± 0.03 ^b	0.02 ± 0.02 ^b	0.27 ± 0.12 ^a	0.01 ± 0.01 ^b	0 ± 0 ^b	0.07 ± 0.03 ^a
11:0	0.02 ± 0.02 ^a	0.01 ± 0.01 ^a	0.04 ± 0.02 ^a	0 ± 0 ^a	0.02 ± 0.02 ^a	0 ± 0 ^a	0 ± 0 ^a	0 ± 0 ^a	0.01 ± 0.01 ^a
12:0	1.04 ± 0.13 ^a	0.32 ± 0.04 ^b	0.77 ± 0.16 ^a	0.88 ± 0.12 ^a	0.48 ± 0.05 ^b	0.55 ± 0.03 ^{ab}	1.05 ± 0.26 ^a	0.34 ± 0.2 ^a	0.56 ± 0.16 ^a
13:0	0.26 ± 0.07 ^a	0.26 ± 0.11 ^a	0.16 ± 0.09 ^a	0.25 ± 0.03 ^{ab}	0.32 ± 0.07 ^a	0.16 ± 0.03 ^b	0.08 ± 0.04 ^a	0.13 ± 0.07 ^a	0.21 ± 0.05 ^a
14:0	12.8 ± 2.15 ^a	18.6 ± 5.14 ^a	16.6 ± 4.28 ^a	8.88 ± 0.87 ^b	14.7 ± 2.39 ^a	19.9 ± 2.18 ^a	7.39 ± 0.72 ^a	10.1 ± 3.86 ^a	14 ± 3.95 ^a
15:0	0.39 ± 0.04 ^a	0.35 ± 0.17 ^{ab}	0.15 ± 0.03 ^b	0.4 ± 0.05 ^a	0.31 ± 0.05 ^a	0.22 ± 0.01 ^a	0.64 ± 0.09 ^a	0.25 ± 0.11 ^b	0.34 ± 0.1 ^{ab}
16:0	121 ± 15.6 ^a	165 ± 44.1 ^a	201 ± 53 ^a	82.1 ± 9.97 ^b	114 ± 19.9 ^{ab}	221 ± 45 ^a	76.2 ± 7.47 ^a	97.7 ± 26.3 ^a	160 ± 53.4 ^a
17:0	0.44 ± 0.13 ^a	0.44 ± 0.23 ^a	0.46 ± 0.16 ^a	0.56 ± 0.12 ^a	0.31 ± 0.09 ^a	0.52 ± 0.14 ^a	1.68 ± 0.31 ^a	1.83 ± 1.03 ^{ab}	0.28 ± 0.14 ^b
18:0	27.9 ± 1.23 ^a	30.3 ± 5.89 ^a	22.3 ± 3.31 ^a	28.2 ± 1.5 ^a	25.7 ± 3.42 ^a	23.6 ± 3.88 ^a	25.3 ± 3.71 ^a	32.3 ± 9.23 ^a	17.7 ± 3.61 ^a
20:0	3.82 ± 0.19 ^a	4.36 ± 0.94 ^a	3.2 ± 0.42 ^a	3.66 ± 0.2 ^a	3.54 ± 0.48 ^a	3.7 ± 0.22 ^a	3.91 ± 0.33 ^a	3.62 ± 0.88 ^a	3.12 ± 0.68 ^a
21:0	0.39 ± 0.05 ^a	0.27 ± 0.06 ^{ab}	0.18 ± 0.05 ^b	0.54 ± 0.06 ^a	0.37 ± 0.04 ^{ab}	0.22 ± 0.06 ^b	0.99 ± 0.17 ^a	1.16 ± 0.63 ^a	0.48 ± 0.22 ^a
22:0	2.34 ± 0.43 ^a	1.38 ± 0.52 ^a	1.16 ± 0.34 ^a	2.8 ± 0.34 ^a	1.95 ± 0.15 ^{ab}	1.25 ± 0.46 ^b	5.69 ± 1.11 ^a	7.39 ± 4 ^a	1.45 ± 0.33 ^b
24:0	0.88 ± 0.16 ^a	0.56 ± 0.26 ^a	0.37 ± 0.11 ^a	1.25 ± 0.2 ^a	0.72 ± 0.12 ^{ab}	0.43 ± 0.11 ^b	2.81 ± 0.66 ^a	3.26 ± 1.66 ^a	0.64 ± 0.13 ^b

14:1n-5	1.43 ± 0.33 ^a	0.26 ± 0.09 ^b	0.41 ± 0.36 ^b	1.77 ± 0.34 ^a	0.83 ± 0.24 ^a	0.66 ± 0.29 ^a	1.64 ± 0.6 ^a	0.4 ± 0.27 ^a	0.4 ± 0.19 ^a
15:1n-5	1.05 ± 0.25 ^a	0.32 ± 0.15 ^a	0.67 ± 0.1 ^a	1.08 ± 0.23 ^a	0.56 ± 0.07 ^a	0.68 ± 0.14 ^a	1 ± 0.35 ^a	0.18 ± 0.11 ^a	0.48 ± 0.12 ^a
16:1n-7	5.78 ± 1.32 ^a	6.29 ± 1.56 ^a	16.6 ± 5.8 ^a	3.3 ± 0.28 ^b	4.65 ± 0.61 ^a	19.3 ± 6.86 ^a	3.05 ± 0.25 ^b	2.82 ± 1.08 ^b	14.6 ± 6.47 ^a
17:1n-7	17.9 ± 5.19 ^a	19.4 ± 8.83 ^a	3.99 ± 3.99 ^a	11.6 ± 1.9 ^a	17.6 ± 4.67 ^a	10.9 ± 10.9 ^a	8.62 ± 0.94 ^{ab}	6.88 ± 5.2 ^b	23 ± 6.88 ^a
18:1n-7	0.61 ± 0.08 ^a	2.41 ± 1.38 ^a	0.98 ± 0.29 ^a	2.07 ± 1.48 ^a	0.46 ± 0.06 ^a	1.09 ± 0.31 ^a	1.4 ± 0.27 ^a	0.32 ± 0.11 ^b	1.25 ± 0.3 ^a
18:1n-7t	0.1 ± 0.06 ^b	0.19 ± 0.12 ^{ab}	0.5 ± 0.2 ^a	0.05 ± 0.05 ^a	0.07 ± 0.04 ^a	0.04 ± 0.04 ^a	0.01 ± 0.01 ^a	0 ± 0 ^a	0.03 ± 0.02 ^a
18:1n-9	14.4 ± 2.71 ^a	13.8 ± 2.38 ^a	31.3 ± 10.2 ^a	13 ± 5.89 ^b	6.89 ± 1.13 ^b	31 ± 10.5 ^a	7.71 ± 0.55 ^a	6.82 ± 1.31 ^a	23.6 ± 9.22 ^a
18:1n-9t	0.22 ± 0.13 ^a	1.43 ± 1.16 ^a	0.51 ± 0.24 ^a	4.57 ± 2.98 ^a	0.11 ± 0.09 ^a	0.24 ± 0.11 ^a	0.11 ± 0.11 ^a	0 ± 0 ^a	0.01 ± 0.01 ^a
20:1n-9	5.57 ± 0.59 ^{ab}	4.48 ± 1.45 ^b	9.8 ± 2.13 ^a	3.82 ± 0.4 ^a	3.05 ± 0.61 ^a	5.45 ± 3.63 ^a	2.93 ± 0.27 ^{ab}	2.12 ± 0.43 ^b	7.81 ± 2.51 ^a
22:1n-9	0.96 ± 0.34 ^a	1.43 ± 0.66 ^a	0.72 ± 0.22 ^a	1.2 ± 0.28 ^a	1.53 ± 0.33 ^a	0.38 ± 0.03 ^b	2.56 ± 0.62 ^a	4.37 ± 1.99 ^a	0.52 ± 0.11 ^b
24:1n-9	0.07 ± 0.03 ^a	0.05 ± 0.04 ^a	0.11 ± 0.05 ^a	0.01 ± 0.01 ^b	0.08 ± 0.06 ^{ab}	0.07 ± 0.02 ^a	0.04 ± 0.03 ^b	0 ± 0 ^b	0.18 ± 0.12 ^a
20:1n-11	0 ± 0 ^b	0 ± 0 ^b	0.64 ± 0.22 ^a	0 ± 0 ^b	0 ± 0 ^b	0.48 ± 0.27 ^a	0 ± 0 ^b	0 ± 0 ^b	0.46 ± 0.25 ^a
22:1n-11	0.13 ± 0.02 ^a	0.09 ± 0.03 ^a	0.15 ± 0.03 ^a	0.24 ± 0.14 ^a	0.12 ± 0.06 ^a	0.15 ± 0.04 ^a	0.28 ± 0.1 ^a	0.13 ± 0.09 ^a	0.23 ± 0.07 ^a
18:3n-3	0.1 ± 0.03 ^a	0.06 ± 0.04 ^a	0.13 ± 0.02 ^a	0.16 ± 0.09 ^a	0.02 ± 0.02 ^a	0.15 ± 0.04 ^a	0.31 ± 0.12 ^a	0.02 ± 0.02 ^{ab}	0.21 ± 0.05 ^a
18:4n-3	6.66 ± 0.61 ^a	6.14 ± 1.27 ^a	7.84 ± 1.34 ^a	6.43 ± 0.7 ^a	7.99 ± 1.67 ^a	9.04 ± 0.84 ^a	3.8 ± 0.46 ^a	4.71 ± 0.37 ^a	5.14 ± 1.83 ^a
20:4n-3	0.42 ± 0.1 ^a	0.43 ± 0.25 ^a	0.46 ± 0.23 ^a	0.22 ± 0.08 ^b	0.12 ± 0.12 ^b	0.7 ± 0.16 ^a	0.08 ± 0.05 ^b	0 ± 0 ^b	0.44 ± 0.17 ^a
20:5n-3	26 ± 1.32 ^a	19.1 ± 3.8 ^a	23.8 ± 0.68 ^a	22.7 ± 1.24 ^a	29 ± 5.85 ^a	24.2 ± 2.73 ^a	13.9 ± 1.28 ^a	18.6 ± 3.17 ^a	15.8 ± 3.82 ^a
22:3n-3	3.99 ± 2.04 ^a	0 ± 0 ^a	0 ± 0 ^a	2.74 ± 1.86 ^a	0 ± 0 ^a	3.29 ± 3.29 ^a			
22:5n-3	7.36 ± 0.57 ^a	4.82 ± 0.69 ^b	5.77 ± 0.61 ^{ab}	6.46 ± 0.6 ^a	7.08 ± 1.24 ^a	5.88 ± 0.22 ^a	5.33 ± 0.65 ^a	8.73 ± 3.38 ^a	4.24 ± 0.96 ^a
22:6n-3	10.3 ± 0.68 ^b	15.1 ± 3.56 ^{ab}	23.8 ± 5.87 ^a	9.55 ± 0.56 ^b	13.4 ± 2.28 ^{ab}	27.5 ± 4.93 ^a	6.18 ± 0.69 ^a	7.83 ± 2.06 ^a	18.5 ± 6.26 ^a
24:5n-3	0.03 ± 0.03 ^a	0.03 ± 0.02 ^a	0.01 ± 0.01 ^a	0 ± 0 ^b	0.02 ± 0.02 ^{ab}	0.1 ± 0.08 ^a			
24:6n-3	0.02 ± 0.01 ^b	0.09 ± 0.03 ^a	0.09 ± 0.02 ^a	0 ± 0 ^c	0.05 ± 0.03 ^b	0.11 ± 0 ^a	0.03 ± 0.03 ^a	0 ± 0 ^a	0.05 ± 0.04 ^a
16:2n-4	0.57 ± 0.37 ^a	0.27 ± 0.27 ^a	1.26 ± 0.63 ^a	1.39 ± 0.49 ^a	0.56 ± 0.56 ^a	0.03 ± 0.03 ^a	1.78 ± 0.18 ^a	1.92 ± 0.57 ^a	1.19 ± 0.4 ^a
16:3n-4	1.23 ± 0.65 ^a	1.81 ± 0.84 ^a	2.73 ± 1.24 ^a	0.18 ± 0.12 ^b	0.73 ± 0.42 ^b	3.14 ± 0.72 ^a	0.04 ± 0.04 ^b	0 ± 0 ^b	1.17 ± 0.73 ^a
18:2n-6	3.68 ± 0.48 ^b	3.91 ± 1.07 ^{ab}	7.78 ± 1.75 ^a	2.69 ± 0.24 ^b	2.63 ± 0.41 ^b	9.42 ± 1.51 ^a	2.64 ± 0.22 ^a	2.63 ± 0.58 ^a	7.17 ± 2.28 ^a
18:2n-6t	0.06 ± 0.05 ^a	0.04 ± 0.02 ^a	0.05 ± 0.01 ^a	0 ± 0 ^b	0 ± 0 ^b	0.01 ± 0.01 ^a	0.08 ± 0.06 ^b	0 ± 0 ^b	0.35 ± 0.07 ^a
18:3n-6	18.7 ± 3.05 ^a	19.3 ± 5.71 ^a	37.7 ± 12.2 ^a	10.5 ± 0.75 ^b	14.4 ± 2.6 ^{ab}	43.3 ± 9.99 ^a	7.94 ± 0.89 ^a	8.44 ± 3.03 ^a	29.6 ± 11.5 ^a
20:2n-6	1.18 ± 0.08 ^b	1.18 ± 0.19 ^b	1.74 ± 0.18 ^a	1.32 ± 0.09 ^b	1.22 ± 0.17 ^b	1.85 ± 0.13 ^a	1.01 ± 0.09 ^a	0.89 ± 0.14 ^a	1.71 ± 0.37 ^a
20:3n-6	1.76 ± 0.32 ^a	2.67 ± 1 ^a	4.48 ± 1.23 ^a	0.89 ± 0.19 ^b	1.22 ± 0.53 ^b	5.2 ± 1.19 ^a	0.31 ± 0.17 ^b	1.21 ± 0.7 ^b	3.71 ± 1.13 ^a
20:4n-6	26.7 ± 1.07 ^a	10.8 ± 1.79 ^b	15.1 ± 3.24 ^b	27.2 ± 1.75 ^a	19.2 ± 2.37 ^b	13.5 ± 2.07 ^b	18 ± 1.75 ^a	16.1 ± 3.85 ^a	12.8 ± 1.9 ^a

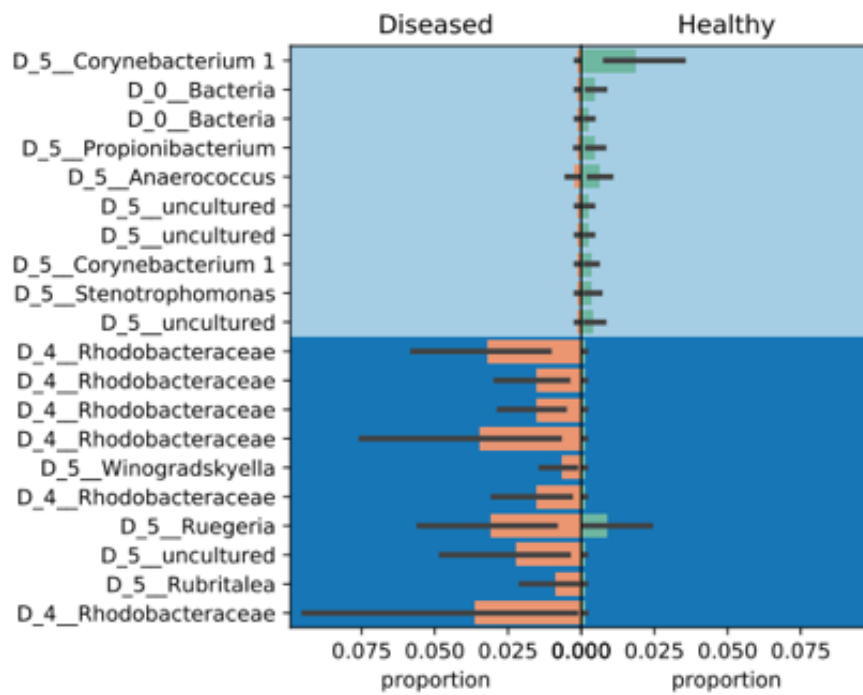
22:2n-6	0.01 ± 0.01 ^b	0.05 ± 0.03 ^a	0.03 ± 0 ^a	0.02 ± 0.02 ^a	0 ± 0 ^a	0.04 ± 0.02 ^a	0.05 ± 0.04 ^a	0 ± 0 ^a	0.03 ± 0.03 ^a
22:4n-6	8.24 ± 1.88 ^a	6.86 ± 1.37 ^a	8.75 ± 2.17 ^a	10.2 ± 1.63 ^a	9.4 ± 1.23 ^{ab}	4.42 ± 2.31 ^b	7.28 ± 0.92 ^a	7.89 ± 1.68 ^a	6.06 ± 1.01 ^a
22:5n-6	0.03 ± 0.02 ^a	0 ± 0 ^a	0.02 ± 0.02 ^a				0.02 ± 0.02 ^b	0 ± 0 ^b	0.14 ± 0.14 ^a
Total	339 ± 29.2 ^a	364 ± 71.2 ^a	454 ± 91.2 ^a	275 ± 18.4 ^b	306 ± 40.1 ^{ab}	495 ± 81.4 ^a	224 ± 19.8 ^a	261 ± 49.9 ^a	380 ± 115 ^a
ΣSFA	174 ± 19.5 ^a	222 ± 55.4 ^a	246 ± 57.2 ^a	130 ± 11.9 ^b	163 ± 25.4 ^{ab}	272 ± 43.1 ^a	126 ± 13.6 ^a	158 ± 36 ^a	199 ± 60.2 ^a
ΣMUFA	48.2 ± 6.19 ^a	50.2 ± 4.82 ^a	66.4 ± 14.9 ^a	42.7 ± 10.6 ^a	35.9 ± 7 ^a	70.5 ± 23.6 ^a	29.3 ± 2.37 ^a	24 ± 6.94 ^a	72.6 ± 25.3 ^a
ΣPUFA	117 ± 6.42 ^{ab}	92.7 ± 17.2 ^b	142 ± 19.7 ^a	103 ± 3.92 ^b	107 ± 14.3 ^{ab}	152 ± 18 ^a	68.8 ± 5.82 ^a	78.9 ± 11.3 ^a	108 ± 30.3 ^a
ΣTRANS	0.38 ± 0.18 ^a	1.66 ± 1.13 ^a	1.06 ± 0.43 ^a	4.61 ± 2.98 ^a	0.18 ± 0.12 ^a	0.29 ± 0.09 ^a	0.2 ± 0.11 ^b	0 ± 0 ^b	0.39 ± 0.05 ^a
Σn-3 PUFA	54.8 ± 3.58 ^a	45.7 ± 9.06 ^a	61.9 ± 7.4 ^a	48.2 ± 2.42 ^b	57.6 ± 10.3 ^{ab}	70.9 ± 8.06 ^a	29.7 ± 2.74 ^a	39.9 ± 6.65 ^a	44.4 ± 12.7 ^a
Σn-3 LC PUFA	48.1 ± 3.27 ^a	39.5 ± 7.83 ^a	54 ± 6.05 ^a	41.6 ± 2.16 ^b	49.6 ± 8.68 ^{ab}	61.7 ± 8.09 ^a	25.6 ± 2.23 ^a	35.1 ± 6.31 ^a	39.1 ± 11 ^a
Σn-6 PUFA	60.4 ± 3.84 ^{ab}	44.8 ± 7.47 ^b	75.7 ± 10.6 ^a	52.8 ± 2.86 ^b	48.1 ± 4.37 ^b	77.7 ± 11 ^a	37.3 ± 3.21 ^a	37.1 ± 5.73 ^a	61.5 ± 16.7 ^a
Σn-6 LC PUFA	36.8 ± 2.11 ^a	20.4 ± 2.5 ^b	28.4 ± 4.33 ^b	38.3 ± 2.75 ^a	29.8 ± 3.32 ^{ab}	23.1 ± 0.61 ^b	25.6 ± 2.54 ^a	25.2 ± 5.5 ^a	22.7 ± 3.83 ^a
EPA:DHA	2.57 ± 0.11 ^a	1.36 ± 0.23 ^b	1.18 ± 0.36 ^b	2.4 ± 0.1 ^a	2.21 ± 0.3 ^a	0.94 ± 0.17 ^b	2.38 ± 0.23 ^a	2.81 ± 0.66 ^a	1.01 ± 0.16 ^b
EPA:ARA	0.98 ± 0.05 ^b	1.83 ± 0.28 ^a	1.76 ± 0.43 ^a	0.86 ± 0.06 ^b	1.51 ± 0.23 ^a	1.83 ± 0.18 ^a	0.79 ± 0.04 ^b	1.28 ± 0.19 ^a	1.19 ± 0.2 ^a
n-3:n-6	0.93 ± 0.07 ^a	1.01 ± 0.03 ^a	0.82 ± 0.02 ^a	0.94 ± 0.06 ^a	1.17 ± 0.12 ^a	0.93 ± 0.09 ^a	0.81 ± 0.04 ^{ab}	1.12 ± 0.16 ^a	0.71 ± 0.02 ^b



Supplementary Figure S3. Shannon diversity index for microbial communities associated with healthy and diseased colonies of *A. hyacinthus*.



Supplementary Figure S4. nMDS of microbial community composition with ellipses representing 90% confidence intervals for A) health states and B) locations.



Supplementary Figure S5. Correlation clustering identified sequence variants with significantly different relative abundance in diseased and healthy tissue.