

1 **Electronic Supplementary Material**

2 **Habitat zonation on coral reefs: structural complexity, nutritional**

3 **resources and herbivorous fish distributions**

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7 **S1 Table Associated tables with the main text**

8 **Table 1** Comparison of Bayesian generalised linear mixed effects models used to examine  
9 the relationship between the abundance as well as biomass of roving herbivorous fishes and a  
10 combined complexity metric (PC1), grazing surface area (GSA) and depth. The most  
11 parsimonious model was selected based on the leave-one-out-cross-validation (LOO)  
12 criterion. Shown are the expected log pointwise predictive density differences (ELPD Diff)  
13 between each model relative to the model with the highest ELPD LOO and the standard error  
14 of component-wise differences of ELPD LOO (SE Diff).

Model	Variables	ELPD Diff	SE Diff
Fish abundance	GSA	0.0	0.0
	NULL	-0.8	2.6
	GSA + DEPTH	-1.3	2.4
	DEPTH	-1.8	2.8
	GSA + PC1	-1.9	1.3
	GSA + PC1 + DEPTH	-1.9	2.4
	PC1	-2.4	2.5
	PC1 + DEPTH	-2.8	3.1
Fish biomass	GSA + PC1 + DEPTH	0.0	0.0
	GSA	-0.5	1.7
	GSA + PC1	-0.5	1.0
	GSA + DEPTH	-1.2	1.5
	NULL	-2.2	2.2
	PC1	-2.9	2.2
	DEPTH	-3.6	2.2

	PC1 + DEPTH	-3.6	2.1
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17 **Table 2** Summary of Bayesian hierarchical models used to compare complexity metrics:  
 18 rugosity index, verticality and hard coral cover among habitats along a coral reef depth  
 19 gradient. Bayesian models used 5,000 iterations, 3 chains, a warm-up of 2,500 iterations and  
 20 a thinning interval of 3, with weakly informative priors. SE = standard error, HPDI = high  
 21 posterior density interval. If the HPDIs intersected zero no effect was inferred.

<b>Response variable</b>	<b>Model used</b>	<b>Predictor variable</b>	<b>Estimate</b>	<b>SE</b>	<b>Lower HPDI</b>	<b>Upper HPDI</b>
Rugosity index	Gaussian (identity link)	Intercept	1.670	0.106	1.510	1.850
		Inner flat	-0.534	0.048	-0.625	-0.437
		Mid flat	-0.480	0.049	-0.571	-0.389
		Outer flat	-0.398	0.051	-0.501	-0.299
		Slope	-0.020	0.077	-0.166	0.137
Verticality	Gaussian (identity link)	Intercept	0.724	0.137	0.485	0.952
		Inner flat	-0.486	0.027	-0.536	-0.430
		Mid flat	-0.456	0.028	-0.511	-0.404
		Outer flat	-0.329	0.029	-0.385	-0.274
		Slope	0.087	0.044	-0.002	0.170
Hard coral cover	Beta-binomial (logit link)	Intercept	-0.585	0.403	-1.490	0.209
		Inner flat	-3.000	0.241	-3.460	-2.510
		Mid flat	-2.430	0.233	-2.890	-1.990
		Outer flat	-1.090	0.200	-1.460	-0.710
		Slope	-0.629	0.186	-0.998	-0.276

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28 **Table 3** Summary of pairwise comparisons used to compare complexity metrics: rugosity  
 29 index, verticality and hard coral cover among habitats along a coral reef depth gradient. SE =  
 30 standard error, HPDI = high posterior density interval. P = Probability of a difference  
 31 occurring between contrast levels.

<b>Response variable</b>	<b>Contrast</b>	<b>Estimate</b>	<b>Lower HDPI</b>	<b>Upper HDPI</b>	<b>P</b>
Rugosity index	Crest - Inner flat	0.535	0.437	0.625	1.0
	Crest - Mid flat	0.480	0.380	0.571	1.0
	Crest - Outer flat	0.400	0.299	0.501	1.0
	Crest - Slope	0.020	-0.137	0.166	0.60
	Inner flat - Mid flat	-0.054	-0.080	-0.028	0
	Inner flat - Outer flat	-0.135	-0.174	-0.098	0
	Inner flat - Slope	-0.514	-0.634	-0.395	0
	Mid flat - Outer flat	-0.081	-0.120	-0.041	0
	Mid flat - Slope	-0.459	-0.582	-0.344	0
	Outer flat - Slope	-0.379	-0.507	-0.258	0
Verticality	Crest - Inner flat	0.487	0.430	0.536	1.0
	Crest - Mid flat	0.456	0.404	0.511	1.0
	Crest - Outer flat	0.329	0.274	0.385	1.0
	Crest - Slope	-0.087	-0.170	0.002	0.024
	Inner flat - Mid flat	-0.031	-0.049	-0.014	0.0008
	Inner flat - Outer flat	-0.157	-0.183	-0.134	0
	Inner flat - Slope	-0.574	-0.639	-0.503	0
	Mid flat - Outer flat	-0.127	-0.152	-0.103	0
	Mid flat - Slope	-0.543	-0.610	-0.473	0
	Outer flat - Slope	-0.415	-0.484	-0.342	0
Hard coral cover	Crest - Inner flat	20.015	11.423	30.462	1.0
	Crest - Mid flat	11.263	6.886	17.297	1.0
	Crest - Outer flat	2.962	1.961	4.218	0.674
	Crest - Slope	1.881	1.278	2.640	0.021
	Inner flat - Mid flat	0.566	0.303	0.885	0
	Inner flat - Outer flat	0.149	0.088	0.232	0
	Inner flat - Slope	0.094	0.054	0.145	0
	Mid flat - Outer flat	0.264	0.163	0.409	0
	Mid flat - Slope	0.167	0.097	0.248	0
	Outer flat - Slope	0.627	0.401	0.890	0

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35 **Table 4** Summary of Bayesian hierarchical models used to compare complexity metrics:  
 36 refuge density, field-of-view and grazing surface area among habitats along a coral reef depth  
 37 gradient. Bayesian models used 5,000 iterations, 3 chains, a warm-up of 2,500 iterations and  
 38 a thinning interval of 3, with weakly informative priors. SE = standard error, HPDI = high  
 39 posterior density interval. If the HDPIs intersected zero no effect was inferred.

<b>Response variable</b>	<b>Model used</b>	<b>Predictor variable</b>	<b>Estimate</b>	<b>SE</b>	<b>Lower HPDI</b>	<b>Upper HPDI</b>
Refuge density	Negative-binomial (log link)	Intercept	1.880	0.651	0.879	3.390
		Inner flat	-3.440	0.403	-4.240	-2.690
		Mid flat	-2.870	0.309	-3.480	-2.320
		Outer flat	-1.110	0.152	-1.400	-0.816
		Slope	-0.144	0.119	-0.358	0.096
Field-of-view	Gaussian (identity link)	Intercept	0.744	0.040	0.669	0.817
		Inner flat	0.181	0.024	0.136	0.231
		Mid flat	0.163	0.024	0.115	0.210
		Outer flat	0.114	0.024	0.068	0.162
		Slope	-0.022	0.024	-0.069	0.025
Grazing surface area	Beta-binomial (logit link)	Intercept	0.127	1.080	-2.14	2.21
		Inner flat	-2.84	0.248	-3.33	-2.34
		Mid flat	-1.57	0.228	-2.01	-1.11
		Outer flat	0.270	0.211	-0.125	0.691
		Slope	0.143	0.211	-0.270	0.570

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47 **Table 5** Summary of pairwise comparisons used to compare complexity metrics: refuge  
 48 density, field-of-view and grazing surface area among habitats along a coral reef depth  
 49 gradient. SE = standard error, HPDI = high posterior density interval. P = Probability of a  
 50 difference occurring between contrast levels.

<b>Response variable</b>	<b>Contrast</b>	<b>Estimate</b>	<b>Lower HDPI</b>	<b>Upper HDPI</b>	<b>P</b>
Refuge density	Crest - Inner flat	30.551	11.710	65.228	1.0
	Crest - Mid flat	17.556	8.844	30.353	1.0
	Crest - Outer flat	3.033	2.180	3.958	0.765
	Crest - Slope	1.157	0.906	1.427	0
	Inner flat - Mid flat	0.575	0.161	1.257	0.0005
	Inner flat - Outer flat	0.100	0.031	0.186	0
	Inner flat - Slope	0.038	0.012	0.068	0
	Mid flat - Outer flat	0.175	0.078	0.291	0
	Mid flat - Slope	0.066	0.032	0.109	0
	Outer flat - Slope	0.381	0.271	0.502	0
Field-of-view	Crest - Inner flat	-0.181	-0.231	-0.136	0
	Crest - Mid flat	-0.163	-0.210	-0.115	0
	Crest - Outer flat	-0.114	-0.162	-0.068	0
	Crest - Slope	0.022	-0.025	0.069	0.83
	Inner flat - Mid flat	0.017	-0.030	0.064	0.78
	Inner flat - Outer flat	0.067	0.018	0.112	0.99
	Inner flat - Slope	0.203	0.156	0.250	1.0
	Mid flat - Outer flat	0.049	0.003	0.097	0.98
	Mid flat - Slope	0.186	0.136	0.231	1.0
	Outer flat - Slope	0.137	0.090	0.184	1.0
Grazing surface area	Crest - Inner flat	17.094	9.303	26.491	1.0
	Crest - Mid flat	4.775	2.799	7.044	0.99
	Crest - Outer flat	0.763	0.493	1.120	0
	Crest - Slope	0.867	0.533	1.251	0
	Inner flat - Mid flat	0.279	0.156	0.441	0
	Inner flat - Outer flat	0.045	0.024	0.070	0
	Inner flat - Slope	0.051	0.028	0.079	0
	Mid flat - Outer flat	0.159	0.095	0.238	0
	Mid flat - Slope	0.181	0.105	0.267	0
	Outer flat - Slope	1.140	0.703	1.643	0

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54 **Table 6** The roving herbivorous fish species observed during visual censuses in Pioneer bay,  
 55 Orpheus Island.

Family	Taxon name
Acanthuridae	<i>Acanthurus</i> spp.
Labridae	<i>Chlorurus bleekeri</i>
Labridae	<i>Chlorurus microrhinos</i>
Labridae	<i>Chlorurus spilurus</i>
Labridae	<i>Hipposcarus longiceps</i>
Labridae	<i>Scarus altipinnis</i>
Labridae	<i>Scarus flavipectoralis</i>
Labridae	<i>Scarus rivulatus</i>
Labridae	<i>Scarus schlegeli</i>
Labridae	<i>Scarus</i> spp.
Siganidae	<i>Siganus doliatus</i>
Siganidae	<i>Siganus lineatus</i>
Siganidae	<i>Siganus</i> spp.

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66 **Table 7** Summary of Bayesian hierarchical models used to examine if nominal herbivorous  
67 fish abundance and biomass varied among habitats along a coral reef depth gradient.  
68 Bayesian models used 5,000 iterations, 3 chains, a warm-up of 2,500 iterations and a thinning  
69 interval of 3, with weakly informative priors. SE = standard error, HPDI = high posterior  
70 density interval. If the HDPIs intersected zero no effect was inferred.

<b>Response variable</b>	<b>Model used</b>	<b>Predictor variable</b>	<b>Estimate</b>	<b>SE</b>	<b>Lower HPDI</b>	<b>Upper HPDI</b>
Abundance	Negative Binomial (log link)	Intercept	-2.090	0.640	-3.300	-0.828
		Inner flat	-1.270	0.556	-2.410	-0.229
		Mid flat	-1.830	0.558	-2.910	-0.714
		Outer flat	1.020	0.538	0.029	2.110
		Slope	-0.810	0.558	-1.930	0.307
Biomass	Gamma (log link)	Intercept	3.13	1.05	1.11	5.13
		Inner flat	-2.69	0.677	-4.02	-1.40
		Mid flat	-5.00	0.683	-6.31	-3.59
		Outer flat	0.017	0.670	-1.26	-1.36
		Slope	-0.847	0.694	-2.10	0.546

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79   **Table 8** Summary of pairwise comparisons used to compare nominal herbivorous fish  
 80   abundance and biomass among habitats along a coral reef depth gradient. SE = standard error,  
 81   HPDI = high posterior density interval. P = Probability of a difference occurring between the  
 82   contrast levels.

<b>Response variable</b>	<b>Contrast</b>	<b>Estimate</b>	<b>Lower HDPI</b>	<b>Upper HDPI</b>	<b>P</b>
Abundance	Crest - Inner flat	3.562	0.879	9.074	0.99
	Crest - Mid flat	6.301	1.189	14.785	1.00
	Crest - Outer flat	0.362	0.080	0.884	0.03
	Crest - Slope	2.284	0.502	5.659	0.93
	Inner flat - Mid flat	1.769	0.383	4.427	0.85
	Inner flat - Outer flat	0.100	0.025	0.257	0.00
	Inner flat - Slope	0.642	0.095	1.603	0.21
	Mid flat - Outer flat	0.057	0.013	0.139	0.00
	Mid flat - Slope	0.361	0.061	0.942	0.04
	Outer flat - Slope	6.228	1.355	16.236	0.99
Biomass	Crest - Inner flat	14.70	1.52	44.66	1.00
	Crest - Mid flat	149.00	20.10	455.95	1.00
	Crest - Outer flat	0.99	0.13	2.95	0.49
	Crest - Slope	2.31	0.38	7.21	0.89
	Inner flat - Mid flat	10.10	1.41	31.39	1.00
	Inner flat - Outer flat	0.07	0.01	0.20	0.003
	Inner flat - Slope	0.16	0.02	0.50	0.01
	Mid flat - Outer flat	0.01	0.00	0.02	0.003
	Mid flat - Slope	0.02	0.00	0.05	0
	Outer flat - Slope	2.40	0.26	7.42	0.90

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88 **Table 9** Summary of Bayesian hierarchical models used to explore the relationship between  
 89 nominal herbivorous fish abundance as well as biomass with grazing surface area (GSA).  
 90 Bayesian models used 5,000 iterations, 3 chains, a warm-up of 2,500 iterations and a thinning  
 91 interval of 3, with weakly informative priors. SE = standard error, HPDI = high posterior  
 92 density interval. If the HDPIs intersected zero no effect was inferred.

<b>Response variable</b>	<b>Model used</b>	<b>Predictor variable</b>	<b>Estimate</b>	<b>SE</b>	<b>Lower HPDI</b>	<b>Upper HPDI</b>
Herbivore abundance	Gamma (log link)	Intercept	-3.47	0.710	-4.77	-1.97
		GSA	3.01	1.360	4.91	5.81

  

Herbivore biomass	Gamma (log link)	Intercept	-4.88	0.866	-6.42	-3.07
		GSA	4.77	1.86	1.18	8.21

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94 **Table 10** List of water visibility measurements recorded with a secchi disk at the  
 95 approximate location of each survey site. Measurements were recorded during high tide  
 96 before the beginning of each survey.

<b>Date</b>	<b>Survey site</b>	<b>Visibility (m)</b>
27/04/2018	1	3.5
28/04/2018	2	3.7
26/04/2018	3	4.4
02/05/2018	1	4.2
30/04/2018	2	3.1
01/05/2018	3	2.8
03/05/2018	1	4.7
02/05/2018	2	4.4
03/05/2018	3	3.9

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99      **Table 11** List of average depth measurements for each habitat (n = 3) within each site (n = 3).

100     All measurements were recorded during high tide.

<b>Site</b>	<b>Habitat</b>	<b>Depth (mean, m)</b>
1	Slope	4.12
2	Slope	3.93
3	Slope	5.00
1	Crest	2.31
2	Crest	1.97
3	Crest	1.91
1	Outer Flat	1.31
2	Outer Flat	1.64
3	Outer Flat	1.52
1	Mid Flat	1.20
2	Mid Flat	1.32
3	Mid Flat	1.34
1	Inner Flat	0.98
2	Inner Flat	1.16
3	Inner Flat	1.01

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110   **Table 12** List of average habitat exposure times for each habitat (n = 3) within each site (n =  
111   3). Measurements used tide data for the Lucinda (offshore) region (18° 53' S, 146° 33' E)  
112   and calculated the percentage of time per year that each reef habitat was covered by < 30 cm  
113   of water, thereby excluding roving herbivorous fishes from grazing.

Site	Habitat	Exposure (mean, %)
1	Slope	0
2	Slope	0
3	Slope	0
1	Crest	0
2	Crest	0.03
3	Crest	0.14
1	Outer Flat	5.00
2	Outer Flat	1.04
3	Outer Flat	1.91
1	Mid Flat	7.72
2	Mid Flat	4.71
3	Mid Flat	4.30
1	Inner Flat	15.24
2	Inner Flat	8.89
3	Inner Flat	14.21

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123 **Table 13** Parameters used in PhotoScan software.

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<i>Photo alignment parameters</i>	
Accuracy	High
Key Point limit	40,000
Tie Point limit	4,000
<i>Optimization</i>	
Camera accuracy	10 m
Marker accuracy	0.005 m
Scale bar accuracy	0.001 m
Projection accuracy	0.1 pix
Tie point accuracy	1 pix
Fit f	Enabled
Fit aspect	Disabled
Fit cx, cy	Enabled
Fit skew	Disabled
Fit k1, k2, k3	Enabled
Fit p1, p2	Enabled
Fit k4	Disabled
<i>Dense point cloud</i>	
Quality	Low
Depth filtering	Aggressive
<i>Mesh</i>	
Surface type	Arbitrary
Source data	Dense cloud
Face count	High
Interpolation	Enabled

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128 **Table 14** Summary of priors used in Bayesian hierarchical models used to compare  
 129 complexity metrics and nominal herbivore abundance and biomass among habitats along a  
 130 coral reef depth gradient. Dpar = distributional parameter.

Response	Prior	Class	Dpar
<i>Rugosity</i>			
	student_t(3,1,10)	Intercept	
	student_t(3,0,10)	Intercept	sigma
	student_t(3,0,10)	sd	
<i>Verticality</i>			
	student_t(3,0,10)	Intercept	
	student_t(3,0,10)	Intercept	sigma
	student_t(3,0,10)	sd	
<i>Field of view</i>			
	student_t(3,1,10)	Intercept	
	student_t(3,0,10)	sd	
	student_t(3,0,10)	sigma	
<i>Refuges</i>			
	student_t(3,1,10)	Intercept	
	student_t(3,0,10)	sd	
	gamma(0.01,0.01)	shape	
<i>Grazing surface area</i>			
	student_t(3,0,10)	Intercept	
	gamma(0.01,0.01)	phi	
	student_t(3,0,10)	shape	
<i>Hard coral cover</i>			
	student_t(3,0,10)	Intercept	
	gamma(0.01,0.01)	phi	
	student_t(3,0,10)	shape	
<i>Herbivore abundance</i>			
	student_t(3,2,10)	Intercept	
	student_t(3,0,10)	sd	
	gamma(0.01,0.01)	shape	
<i>Herbivore biomass</i>			
	student_t(3,2,10)	Intercept	
	student_t(3,0,10)	sd	
	gamma(0.01,0.01)	shape	

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133   **Table 15** Summary of priors used in Bayesian hierarchical models used to explore the  
134   relationship between nominal herbivorous fish abundance and biomass with grazing surface  
135   area (GSA).

Response	Prior	Class
<i>Herbivore abundance</i>		
	student_t(3,-3,10)	Intercept
	student_t(3,0,10)	sd
	gamma(0.01,0.01)	shape
<i>Herbivore biomass</i>		
	student_t(3,-4,10)	Intercept
	student_t(3,0,10)	sd
	gamma(0.01,0.01)	shape

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