

Appendix D
Supporting Data
2011 SBOO Stations
Macrobenthic Communities

Appendix D.1

SBOO two-way crossed ANOSIM results for benthic infauna (A=sediment type, B=depth stratum).

Global Test: Factor A

Tests for differences between depth strata (across all sediment types)

Sample statistic (Global R):	0.616
Significance level of sample statistic:	0.01%
Number of permutations:	9999
Number of permuted statistics greater than or equal to Global R:	0

Pairwise Tests: Factor A

Tests for pairwise differences between individual depths across all sediment types: r values (p values)

	Sand with coarse	Sand with fines	Coarse with sand	Sand with coarse and fines
Sand	0.305 (0.002)	0.433 (0.0001)	0.955 (0.01)	0.376 (0.015)
Sand with coarse		0.755 (0.0001)	0.836 (0.048)	0.964 (0.048)
Sand with fines			1 (0.008)	0.167 (0.300)
Coarse with sand				no test

Global Test: Factor B

Tests for differences between sediment types (across all depth strata)

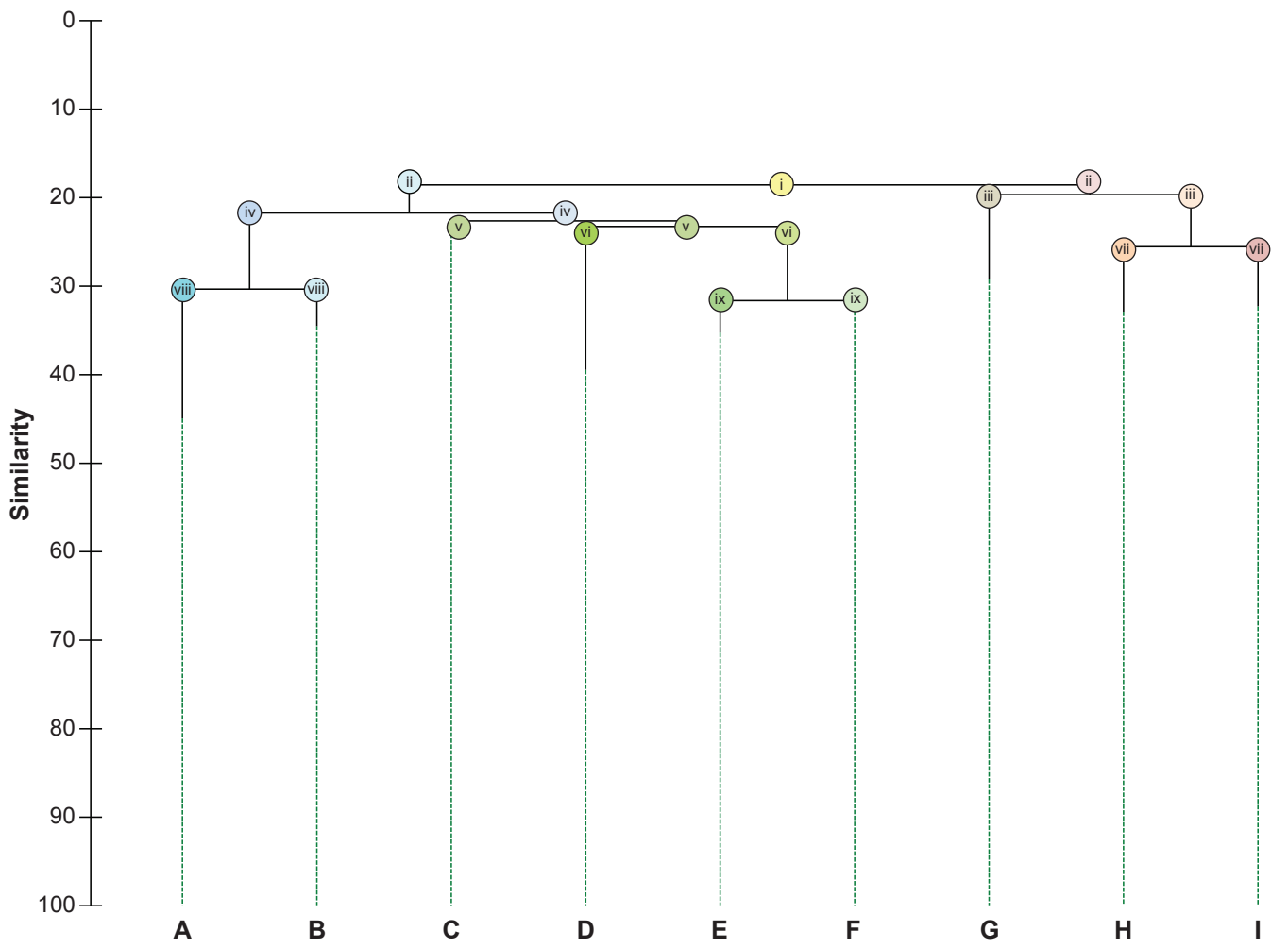
Sample statistic (Global R):	0.539
Significance level of sample statistic:	0.01%
Number of permutations:	9999
Number of permuted statistics greater than or equal to Global R:	0

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Appendix D.2

Delineation of cluster groups (see Figure 5.4) by species exclusivity (i.e., species that occur solely in each supported clade versus species that occur in multiple non-related clades). Roman numerals and colored circles in dendrogram (below) correspond to numbers and colors delineating each SIMPROF-supported split featured in the appendix (following pages). Inner=inner shelf (5–30 m), mid=mid-shelf (30–120 m). S=sand, Sc=sand with coarse, Sf=sand with fines, Scf=sand with coarse and fines, Cs=coarse with sand. CG = cluster group.

CG	# grabs (n)			Depth	Depth			Sed.	Fines			depth/sediment exceptions
	invert	sed.	nearfield		mean	min	max		mean	min	max	
A	4	2	0	mid	55	55	55	Scf	22.3	21.3	23.3	—
B	5	3	0	mid	59	55	60	S	9.0	8.6	9.7	—
C	4	2	0	inner	20	19	21	S	9.0	8.8	9.3	—
D	2	1	0	inner	18	18	18	S	5.4	5.4	5.4	—
E	35	16	11	inner	30	28	38	Sf	13.5	0.0	29.4	Sf/mid = 1, S/inner = 1, S/mid = 1, Sc/inner = 1
F	15	7	0	inner	19	19	21	S	15.8	7.9	33.8	Sf = 2
G	3	3	0	inner	26	19	38	Cs	9.2	0.5	25.6	Sf/mid = 1
H	14	6	0	mid	47	38	55	Sc	4.3	0.0	10.8	Sf = 1, S = 2
I	26	12	5	inner	28	18	36	varied	1.7	0.0	4.8	S/mid = 4, Sc/inner = 4, S/inner = 2, Sc/mid = 1, Cs/inner = 1



Appendix D.2 *continued*

(i.) Species occurring in all cluster groups

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Aricidea (Acmira) catherinae</i>	3.3	1.4	0.3	0.5	2.9	0.5	0.3	0.07	0.04
<i>Leptochelia dubia</i>	9.8	2.6	0.8	1.0	1.9	0.6	1.0	3.5	0.27
Lineidae	2.5	1.6	0.8	0.5	2.5	1.5	1.3	0.79	1.58
<i>Lumbrineris lingulata</i>	8.3	4.0	0.3	0.5	0.5	0.5	11.3	1	0.38
<i>Paranemertes californica</i>	0.5	0.2	0.5	0.5	1.0	0.8	1.0	0.57	0.04
<i>Prionospio (Prionospio) jubata</i>	15.5	5.0	2.0	1.5	13.0	2.3	2.3	2	3.08
<i>Scoloplos armiger</i> Cmplx	1.0	7.8	0.5	3.0	4.0	1.1	0.7	1.29	5.69
<i>Spiophanes berkeleyorum</i>	2.5	0.2	0.3	1.0	4.1	1.3	0.3	0.29	0.31
<i>Spiophanes norrisi</i>	57.0	14.2	1.3	11.0	43.7	21.9	61.7	42.64	196.27
<i>Tiron biocellata</i>	1.0	0.2	1.3	2.0	0.6	0.3	0.3	0.14	0.04

(ii.) Species delineating the separation of cluster groups A, B, C, D, E, and F from cluster groups G, H, and I (18.57% similarity)

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Typosyllis</i> sp SD1	0.0	0.0	0.0	0.0	0.0	0.0	9.3	4.43	0.27
<i>Micranellum crebricinctum</i>	0.0	0.0	0.0	0.0	0.0	0.0	2.7	0.29	0.38
<i>Cnemidocarpa rhizopus</i>	0.0	0.0	0.0	0.0	0.0	0.0	2.3	2	0.38
<i>Sipunculus nudus</i>	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.07	0.19

(iii.) Species delineating the separation of cluster group G from cluster groups H and I (19.67% similarity)

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Aricidea (Acmira) cerrutii</i>	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.79	0.15
<i>Clymenella</i> sp A	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.36	0.31
<i>Aphelochaeta</i> sp SD5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.29	0.42
<i>Clymenella complanata</i>	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.29	0.08
additional 5 taxa (<0.15)	0.0	0.0	0.0	0.0	0.0	0.0	0.0	x	x
<i>Saccocirrus</i> sp	0.0	0.0	0.0	0.0	0.0	0.0	76.0	0.0	0.0
<i>Pareurythoe californica</i>	0.0	0.0	0.0	0.0	0.0	0.0	27.7	0.0	0.0
<i>Eulalia</i> sp SD1	0.0	0.0	0.0	0.0	0.0	0.0	6.0	0.0	0.0
Leptoplanidae	0.0	0.0	0.0	0.0	0.0	0.0	2.0	0.0	0.0
<i>Typosyllis</i> sp SD6	0.0	0.0	0.0	0.0	0.0	0.0	1.7	0.0	0.0
<i>Dorvillea (Schistomeringos)</i> sp	0.0	0.0	0.0	0.0	0.0	0.0	1.3	0.0	0.0
<i>Microphthalmus</i> sp	0.0	0.0	0.0	0.0	0.0	0.0	1.3	0.0	0.0
<i>Polygordius</i> sp	0.0	0.0	0.0	0.0	0.0	0.0	1.3	0.0	0.0
<i>Rhabdocoela</i> sp A	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0
additional 11 taxa (<0.68)	0.0	0.0	0.0	0.0	0.0	0.0	x	0.0	0.0

(iv.) Species delineating the separation of cluster groups A and B from cluster groups C, D, E, and F (21.73% similarity)

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Aphelochaeta williamsae</i>	5.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Euchone incolor</i>	2.8	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Aphelochaeta</i> sp SD13	2.3	1.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Proclea</i> sp A	2.0	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Eclysippe trilobata</i>	0.5	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Eusyllis habei</i>	0.5	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Chiridota</i> sp	0.5	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0
additional 5 taxa (<0.25)	x	x	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Neotrypaea</i> sp	0.0	0.0	0.3	2.0	0.1	0.1	0.0	0.0	0.0

[illegible]

Appendix D.2 *continued*

(viii.) Species delineating the separation of cluster groups A and B (30.35% similarity)

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Aphelochaeta tigrina</i>	4.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Clymenura gracilis</i>	4.8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Streblosoma</i> sp SD1	3.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Nuculana hamata</i>	1.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Gymnonereis crosslandi</i>	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Photis bifurcata</i>	1.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Cardiomya pectinata</i>	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Cirratulus</i> sp	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Dougaloplus</i> sp A	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Eulalia californiensis</i>	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Thyasira flexuosa</i>	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
additional 39 taxa (≤ 0.75)	x	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Aricidea</i> (<i>Allia</i>) <i>antennata</i>	0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Polycirrus</i> sp SD3	0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<i>Tritella pilimana</i>	0.0	1.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0
additional 21 taxa (≤ 0.80)	0.0	x	0.0	0.0	0.0	0.0	0.0	0.0	0.0

(ix.) Species delineating the separation of cluster groups E and F (31.66% similarity)

Cluster groups	A	B	C	D	E	F	G	H	I
<i>Caprella penantis</i>	0.0	0.0	0.0	0.0	0.6	0.0	0.0	0.0	0.0
additional 74 taxa (< 0.25)	0.0	0.0	0.0	0.0	x	0.0	0.0	0.0	0.0
<i>Astyris gausapata</i>	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0
additional 14 taxa (< 0.25)	0.0	0.0	0.0	0.0	0.0	x	0.0	0.0	0.0