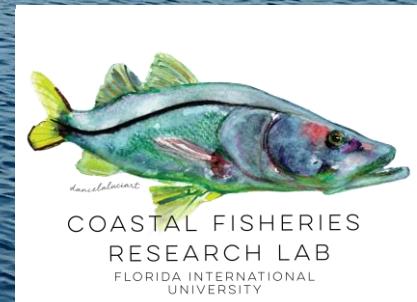
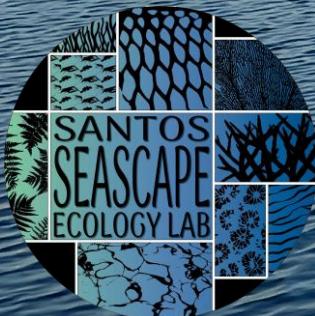


WS01 – Hypervolume modeling: a multivariate tool for seagrass ecosystem assessments

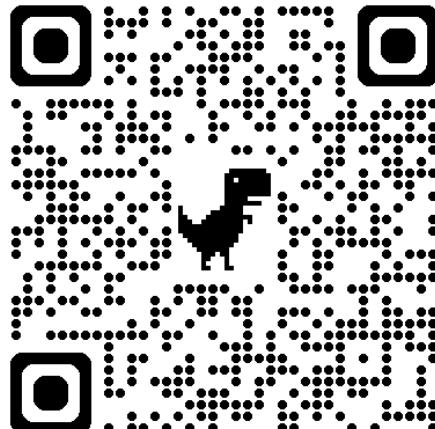
W. Ryan James, Rolando O. Santos, Benjamin Jones,
Jennifer S. Rehage, Gina Badowski, Jonathan Rodemann,
Marianna Coppola

06/21/2024



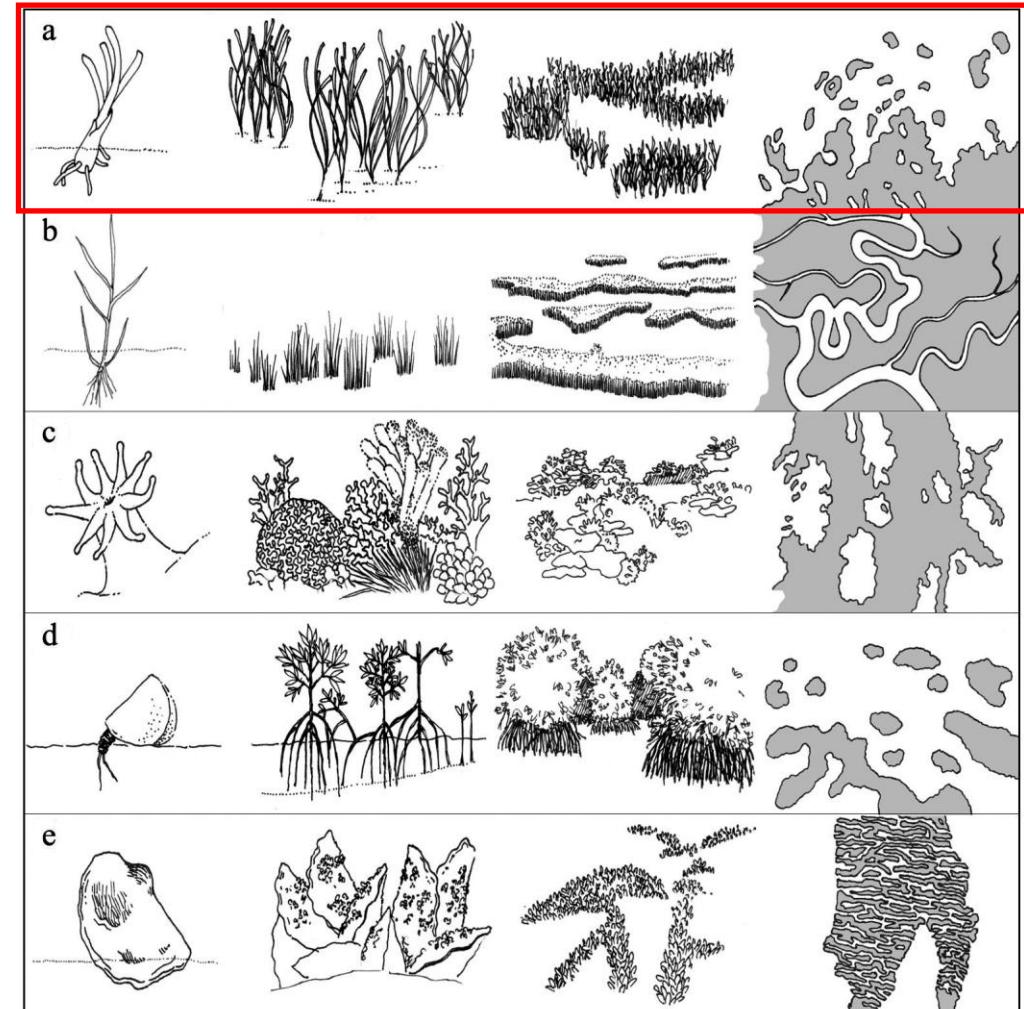
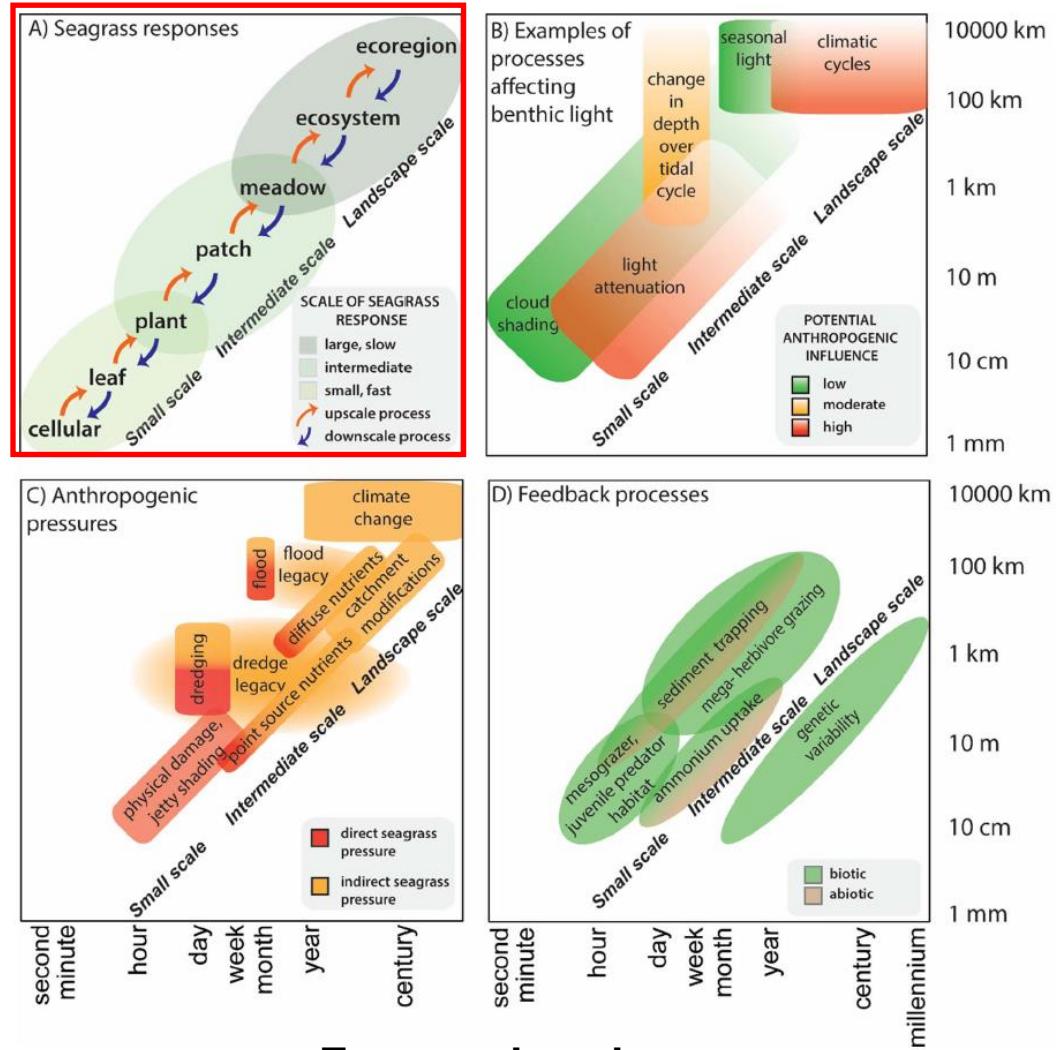
Agenda and materials

- 2:30 – Overview of hypervolumes and metrics
 - 2:50 – Case studies
 - 3:05 – Q & A
 - 3:20 – Code run through
-
- Workshop website:
https://coastalfishscience.github.io/ISBW_HVworkshop/
 - Padlet for questions: <https://padlet.com/badlowski2/hypervolume-modelling-pteo4vn56sk8z2bk>

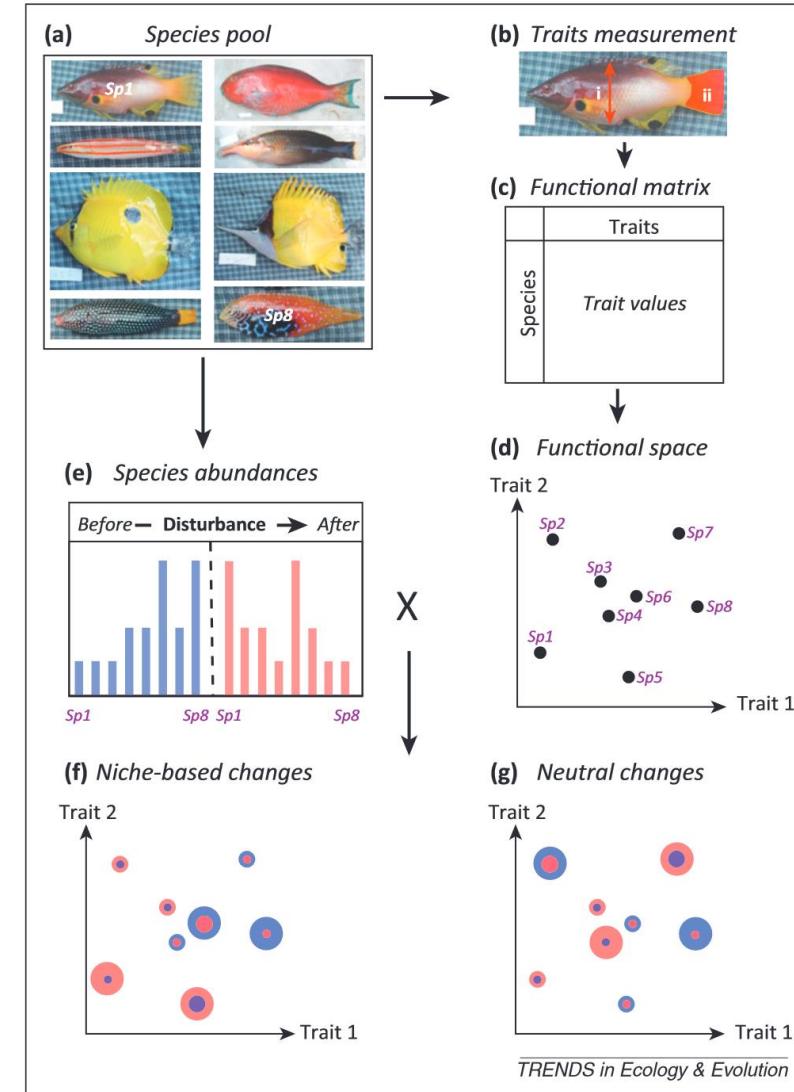
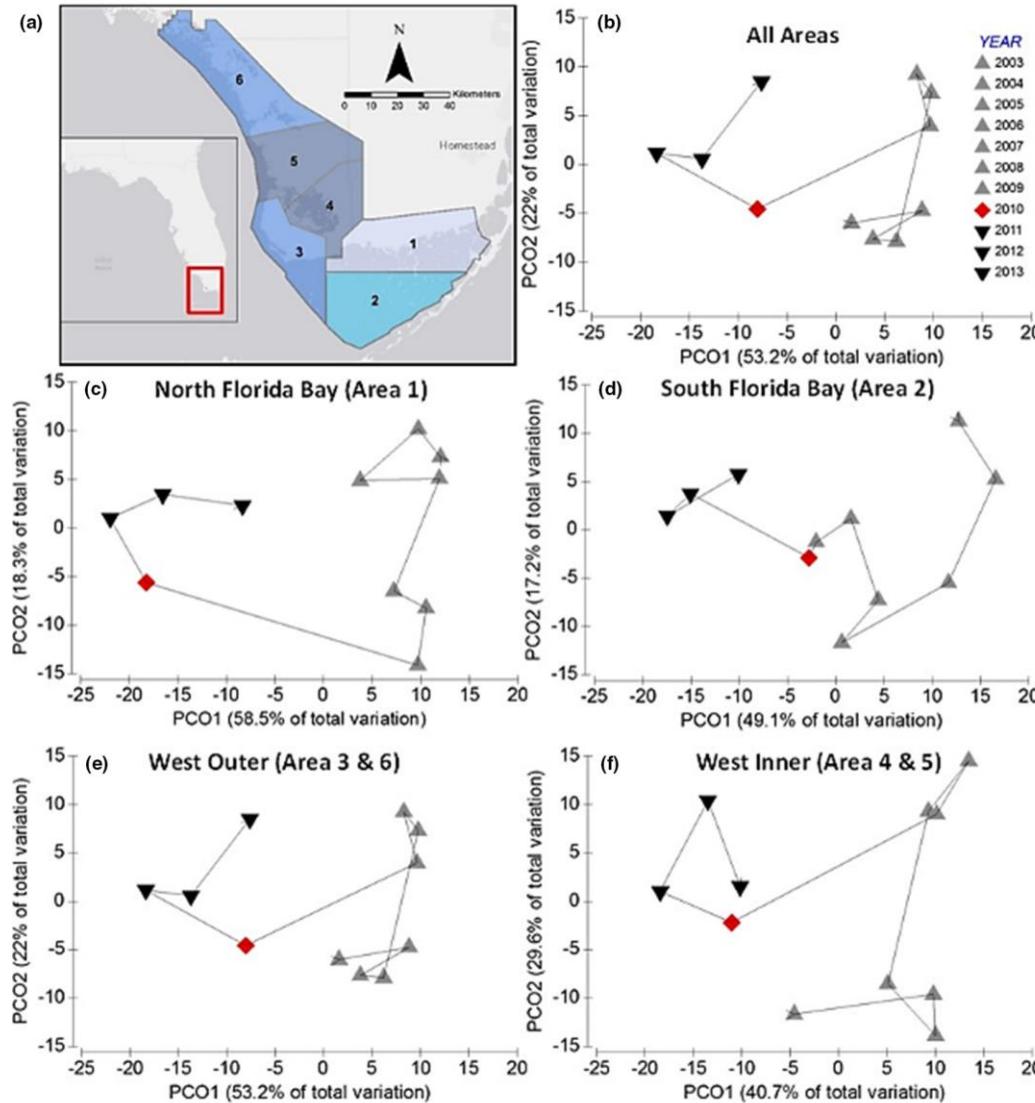


Many ecological questions are multivariate in nature – e.g. multiscale processes and responses

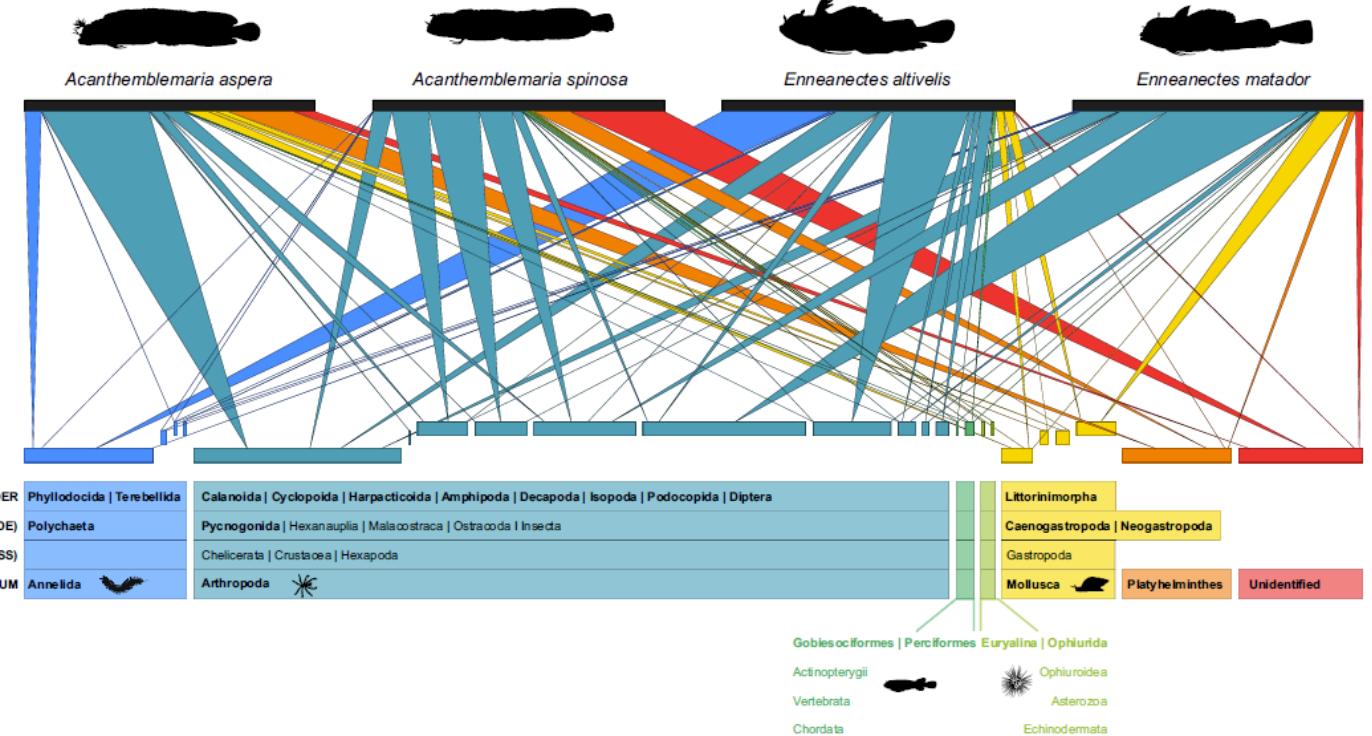
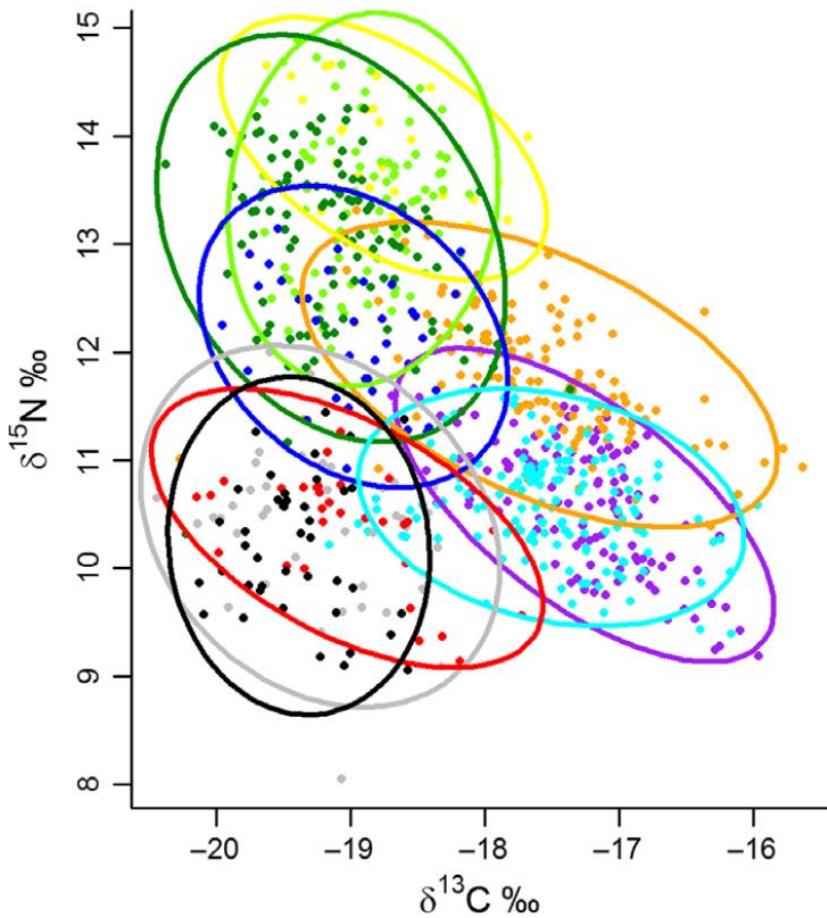
Spatial scales



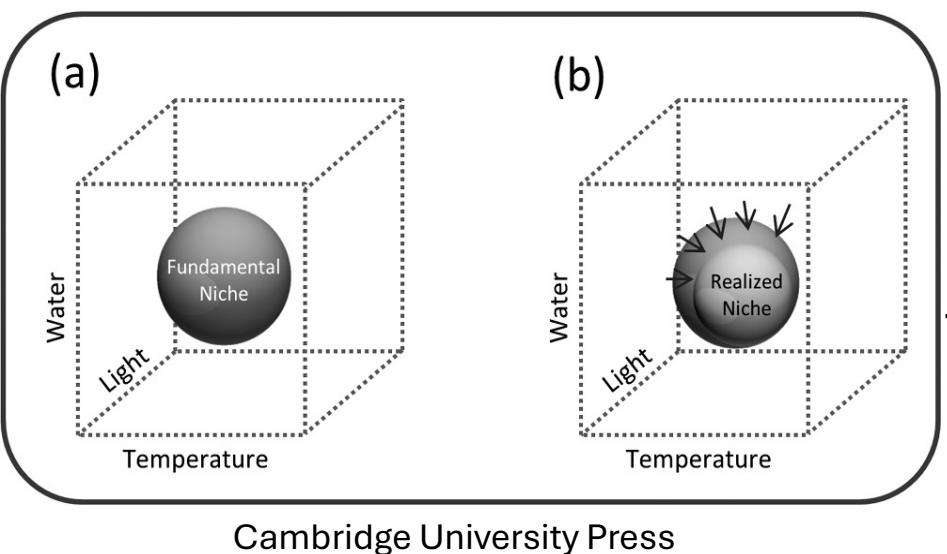
Many ecological questions are multivariate in nature – e.g. taxonomic or trait structure



Many ecological questions are multivariate in nature – e.g. species interactions and trophic dynamics



Adaptation of the Hutchinson's n-dimensional hypervolume to address complex questions



Computational
Power =

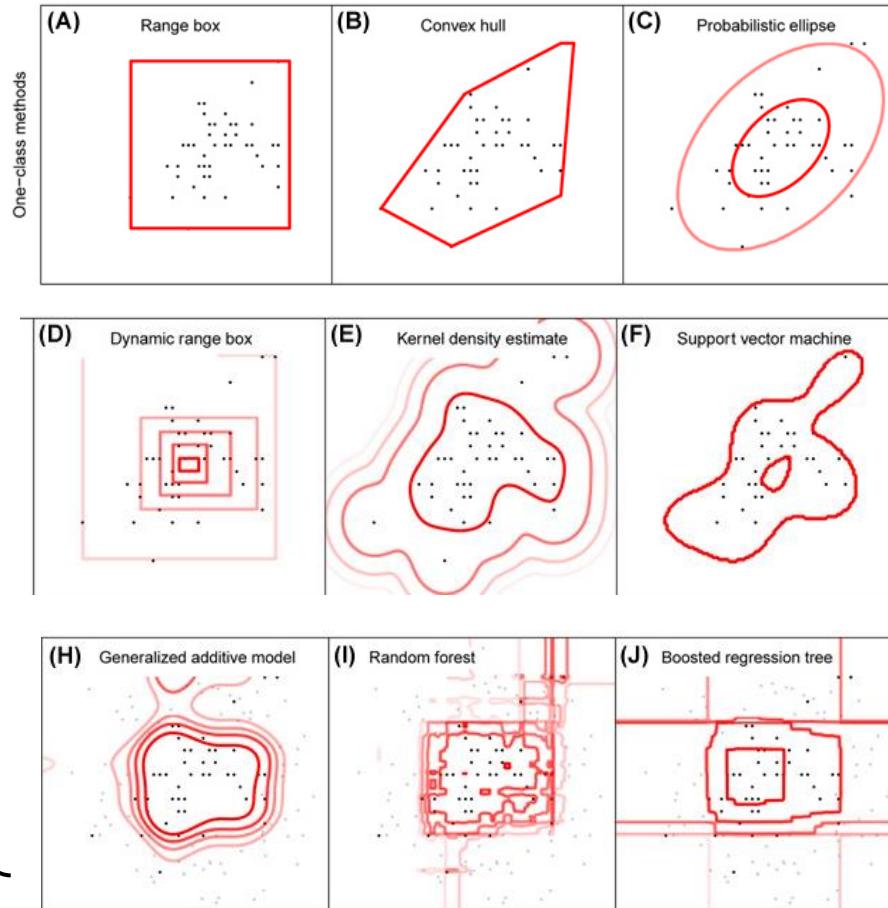
Trait Ecology

Trophic Ecology

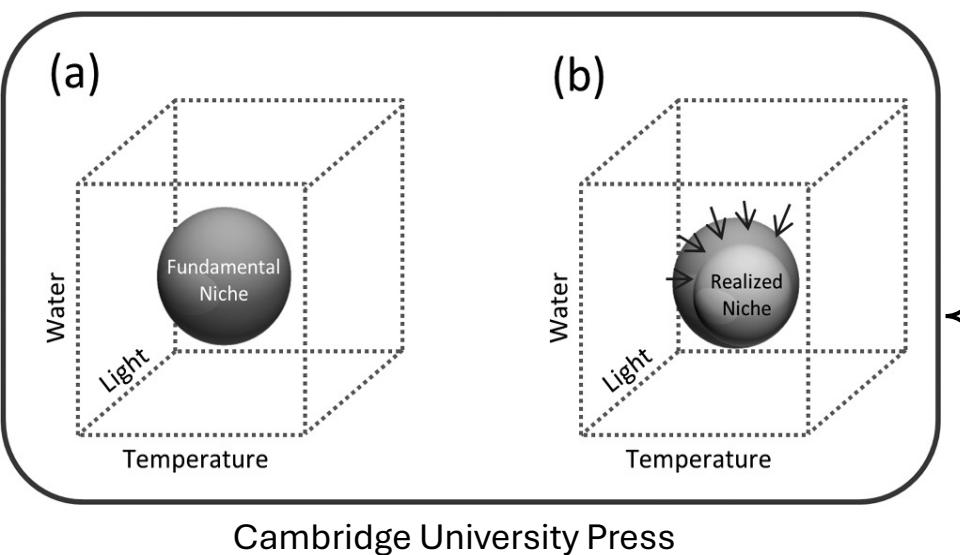
Ecography

Ecosystem

Examples of analytical
approaches



Adaptation of the Hutchinson's n-dimensional hypervolume to address complex questions



Computational
Power =

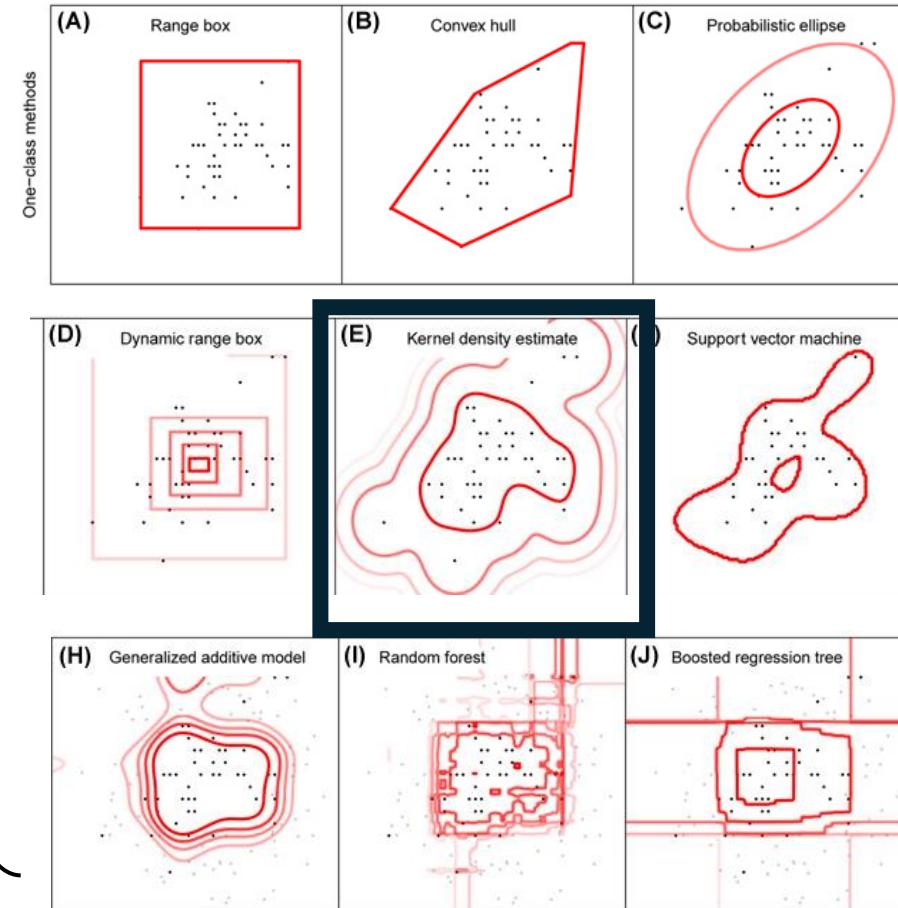
Trait Ecology

Trophic Ecology

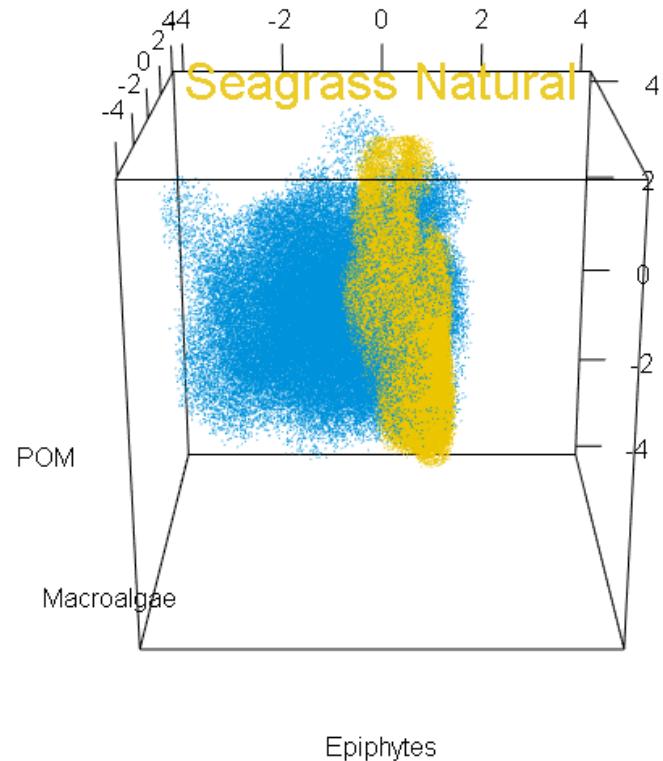
Ecography

Ecosystem

Examples of analytical
approaches

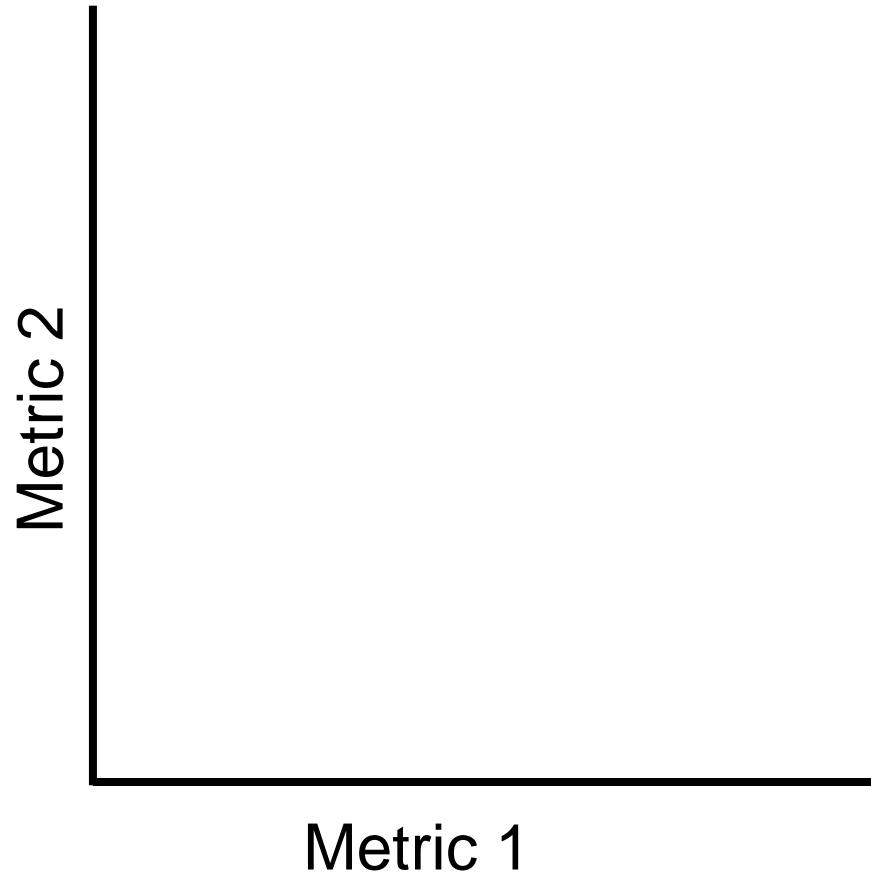


Hypervolumes

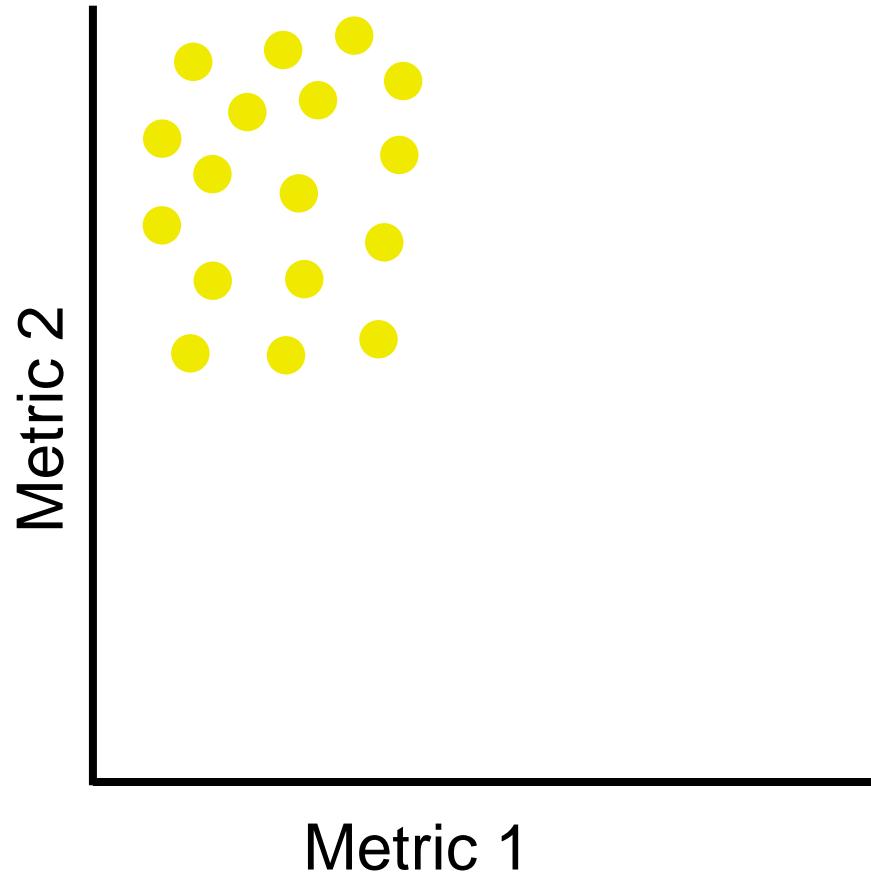


- Hypervolumes (Blonder et al. 2014, 2018)
- Kernel density estimation (but others in package)
- Multivariate tool (n -dimensional)
- Hypervolume metrics useful in ecological analysis

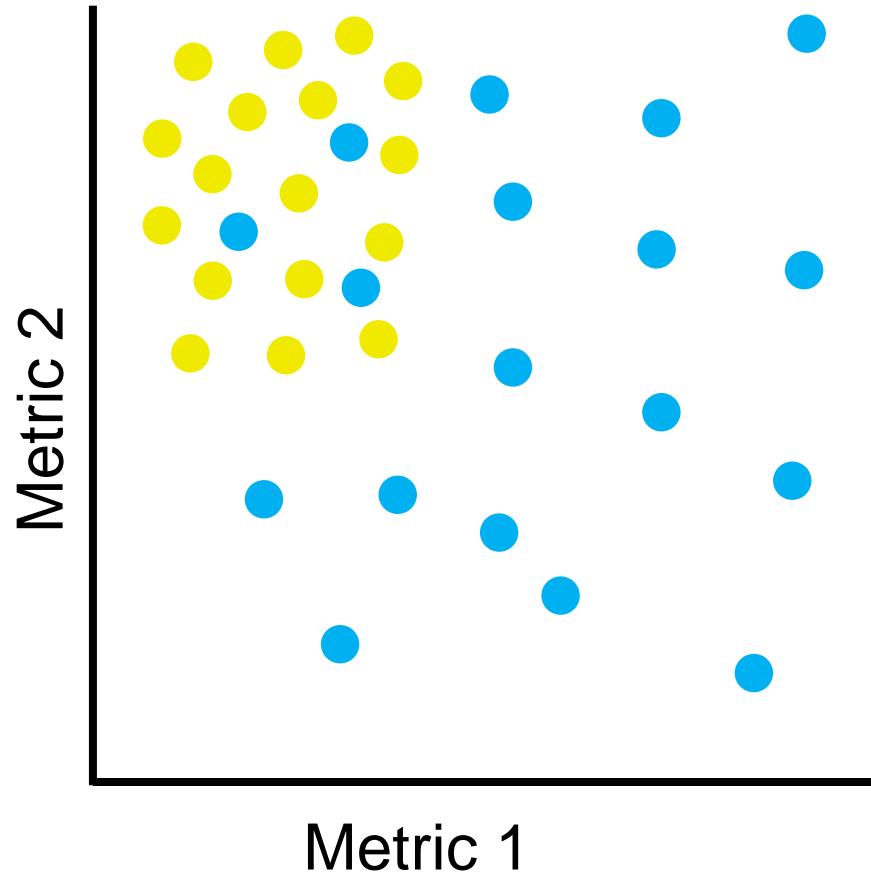
Building Hypervolumes



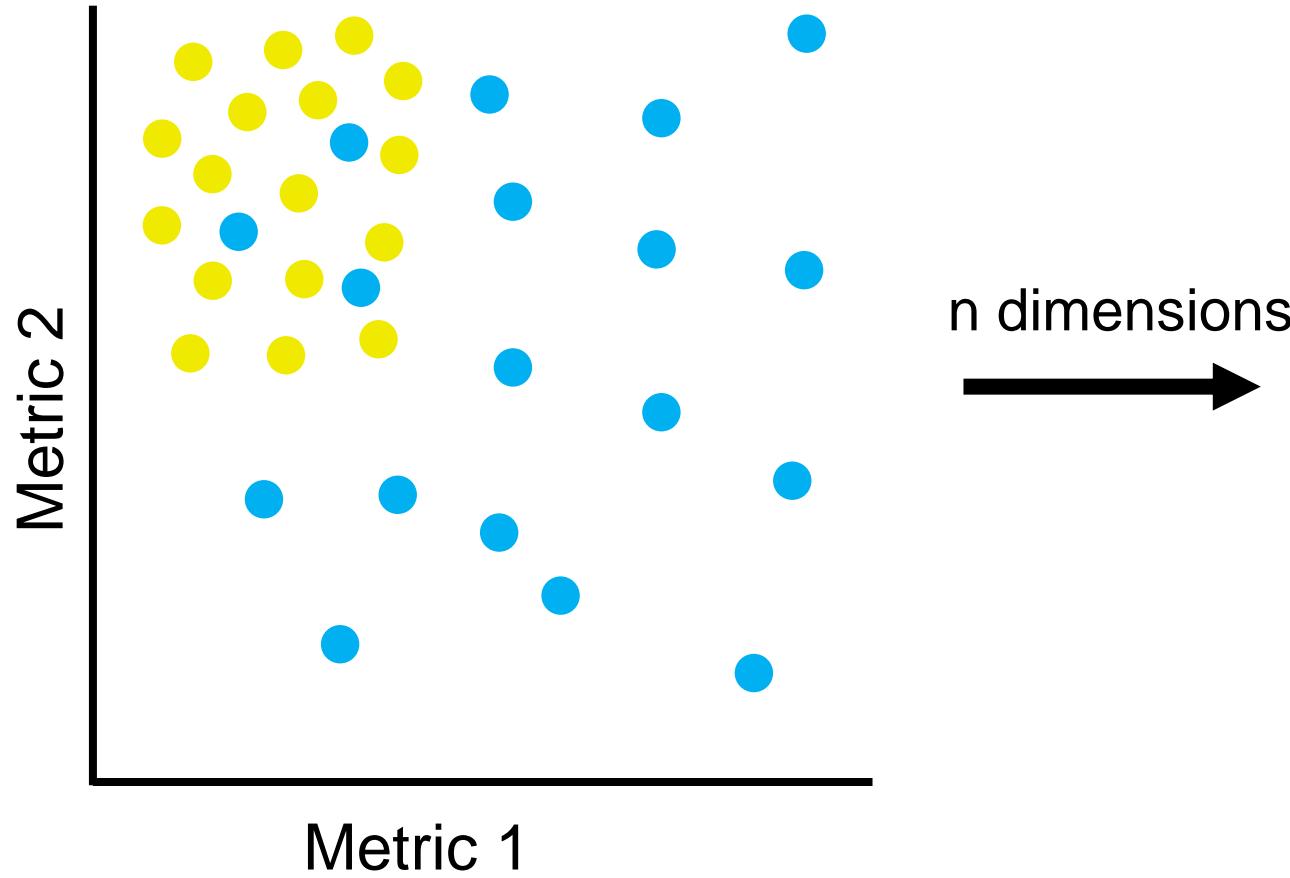
Building Hypervolumes



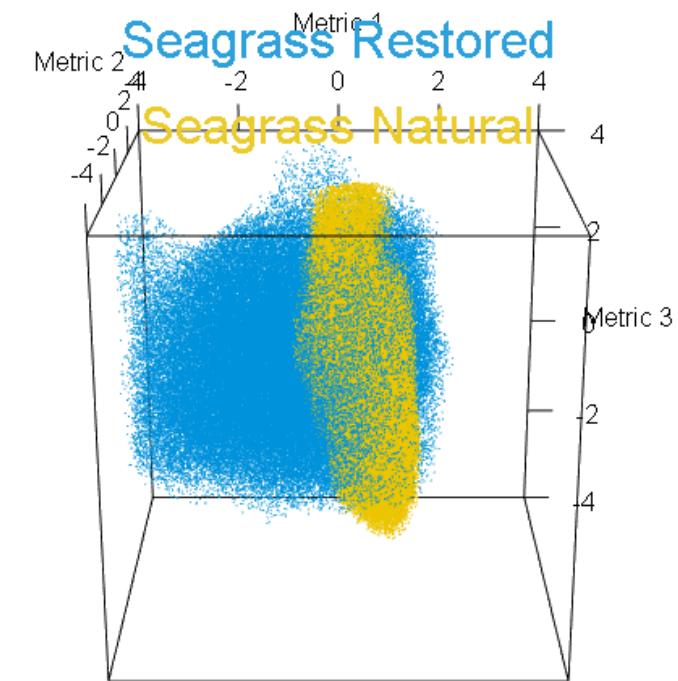
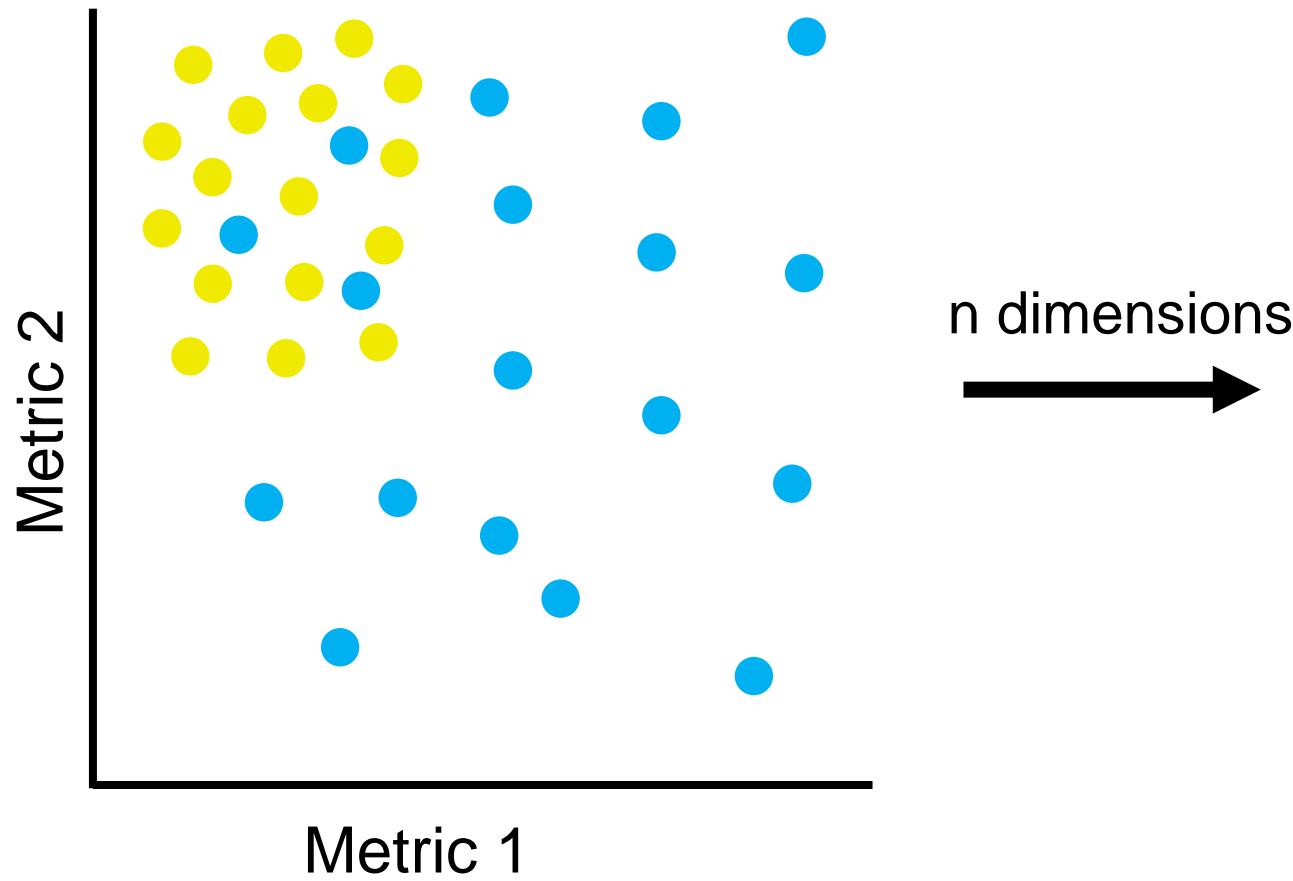
Building Hypervolumes



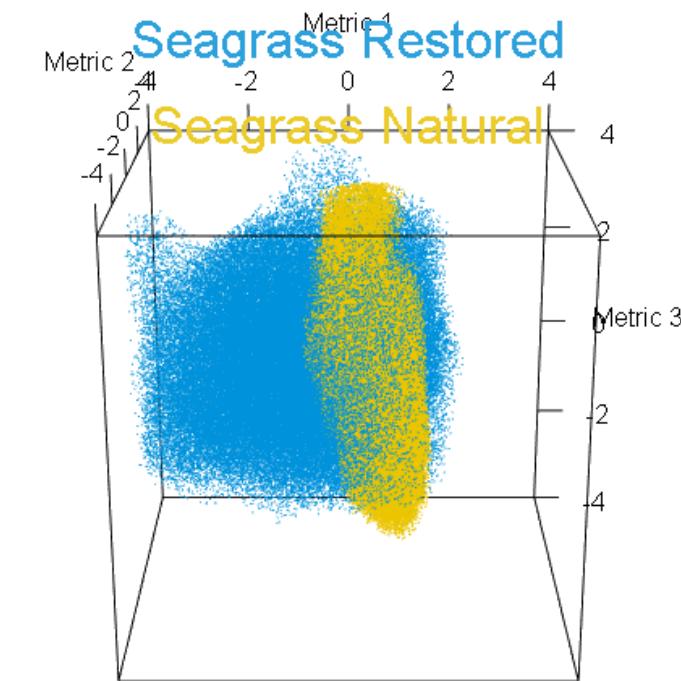
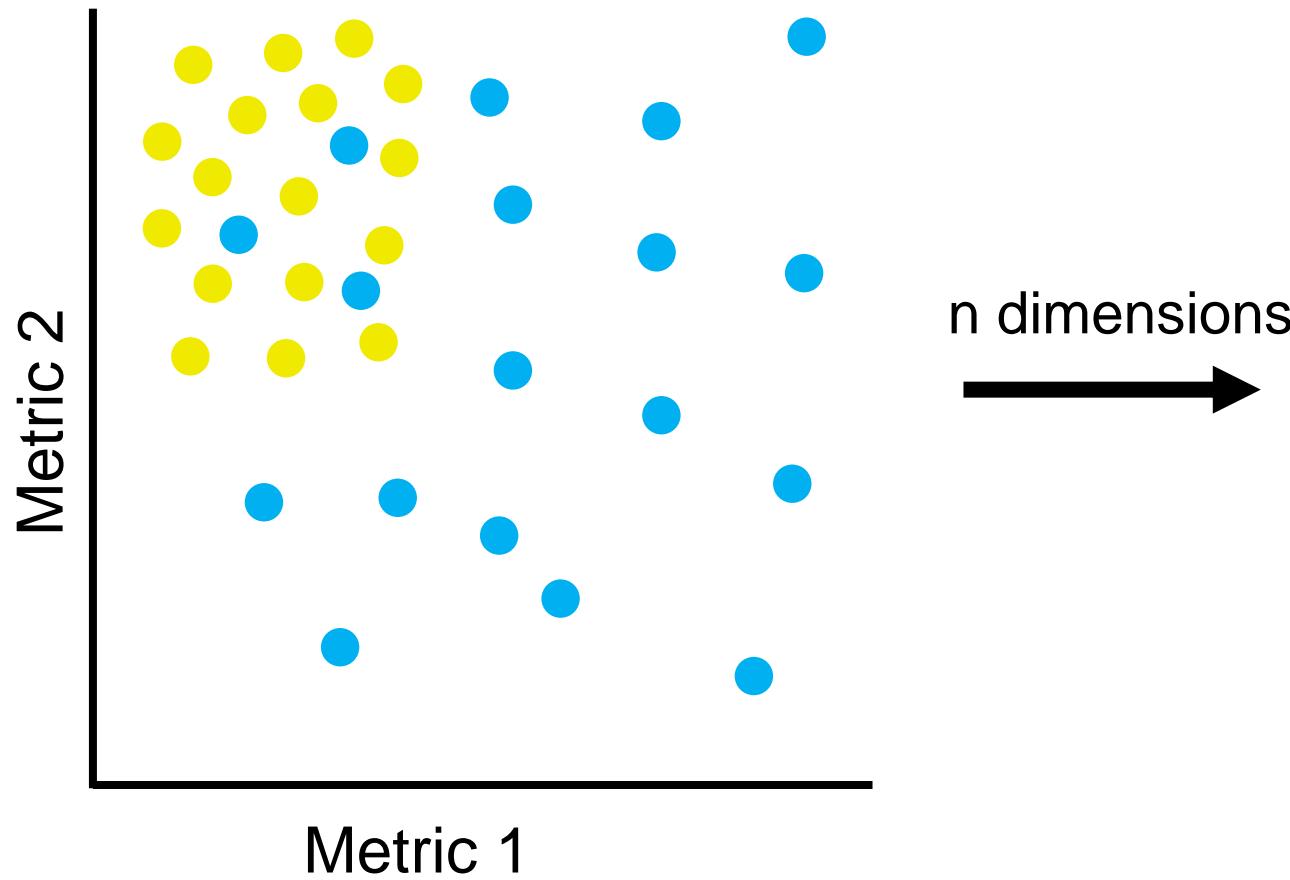
Building Hypervolumes



Building Hypervolumes

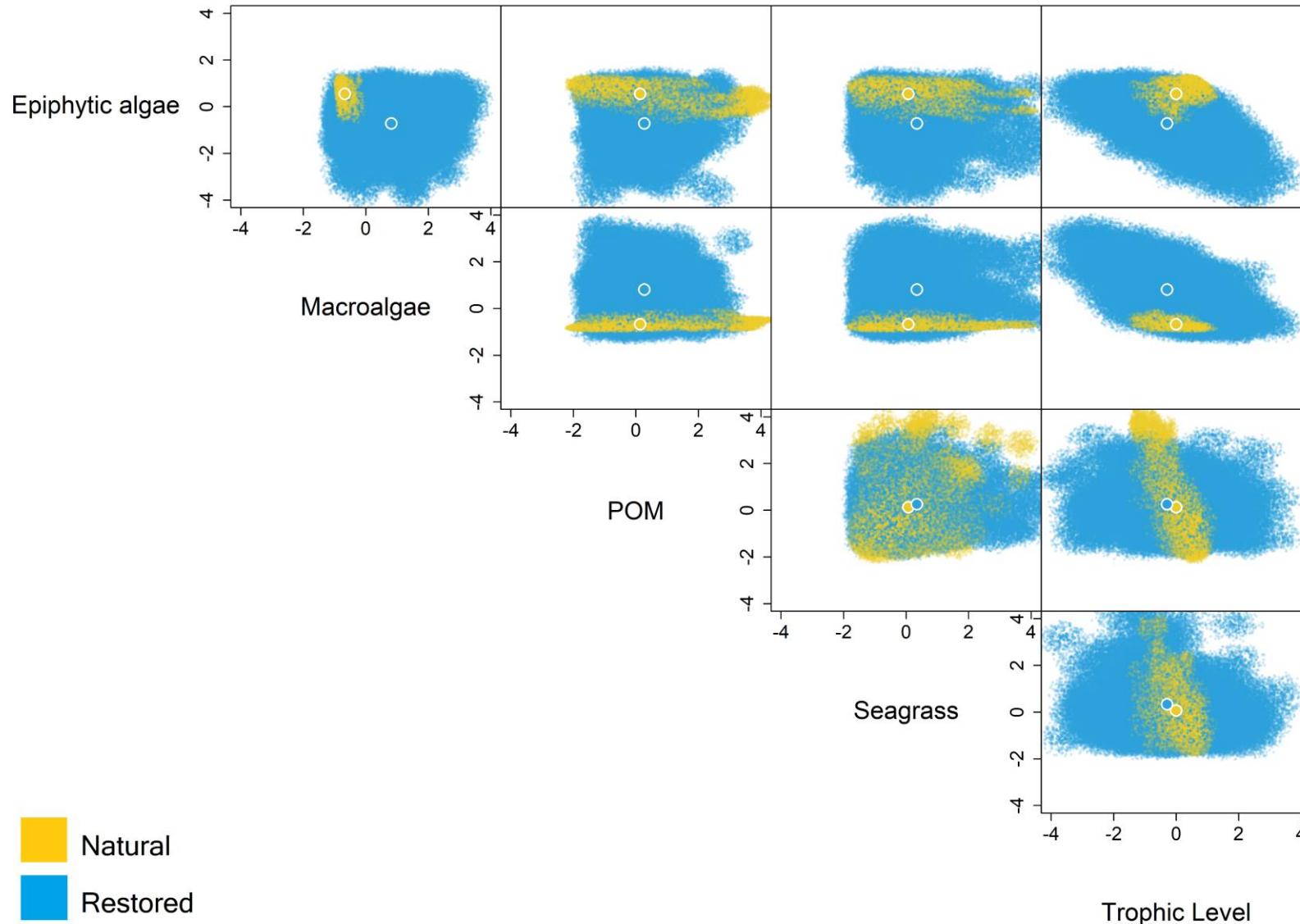


Building Hypervolumes

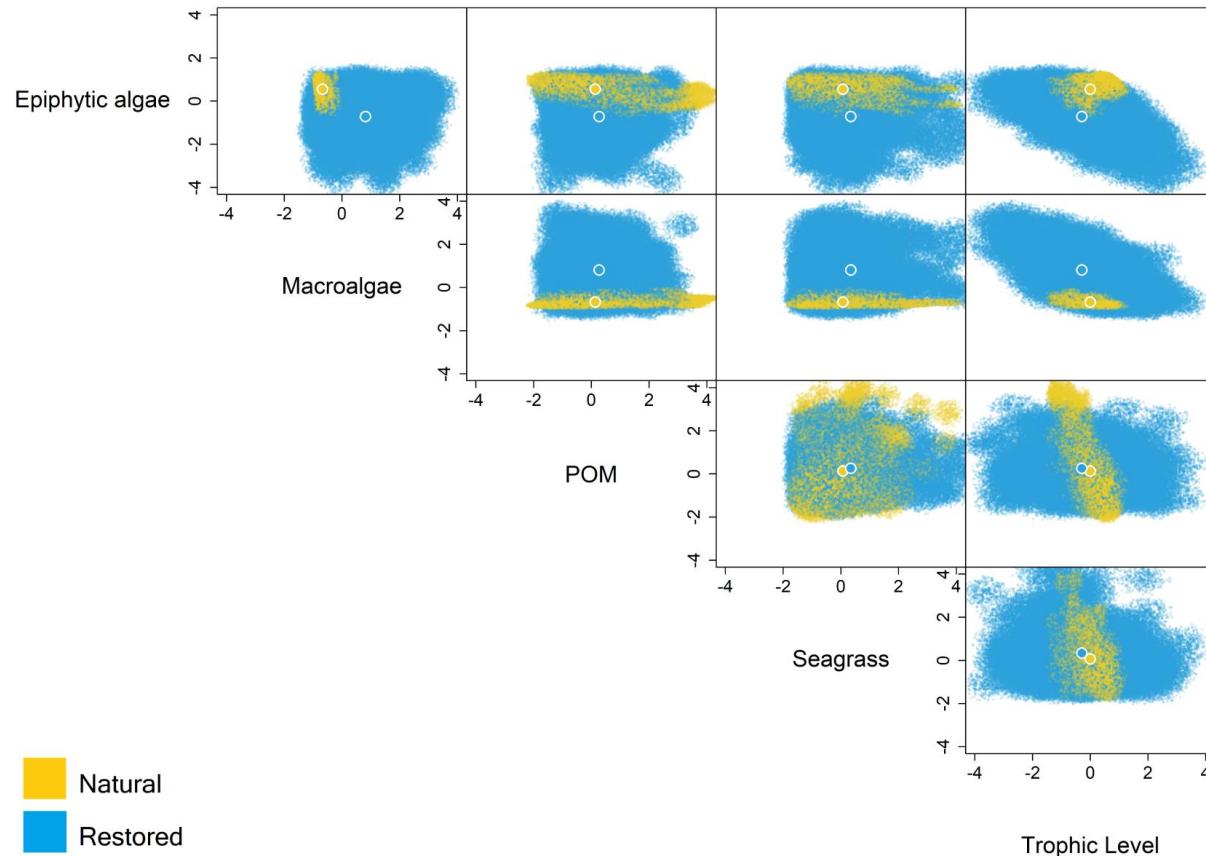


Size based on variability

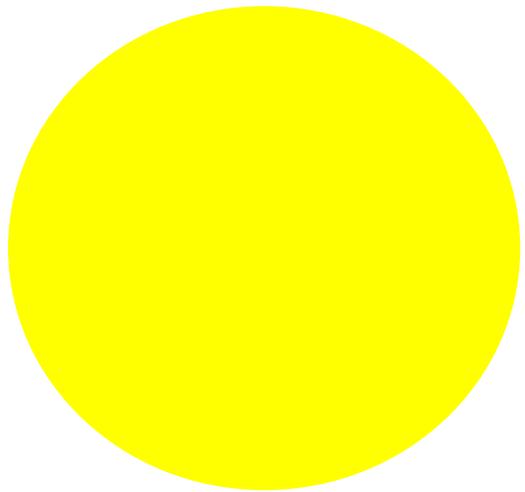
Hypervolume biplot



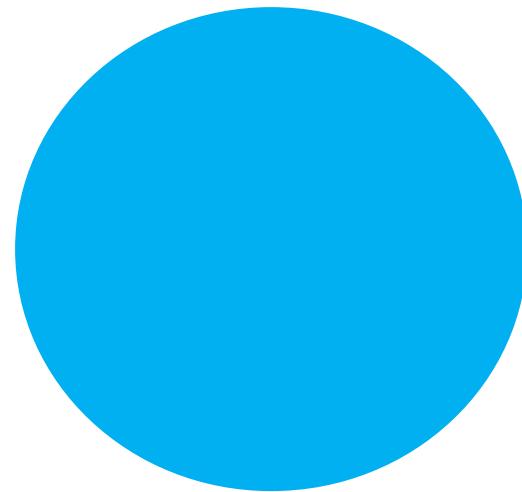
Hypervolume biplot



Hypervolume metrics



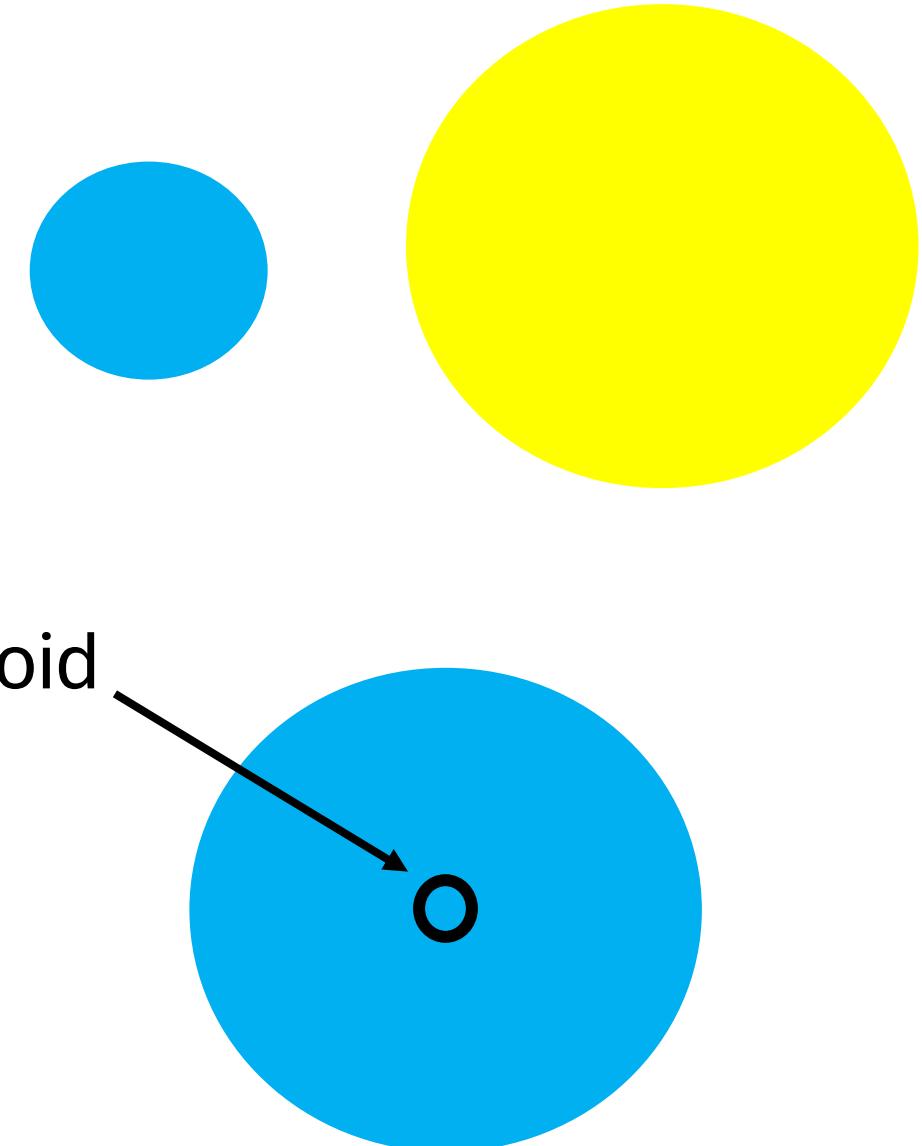
HV1



HV2

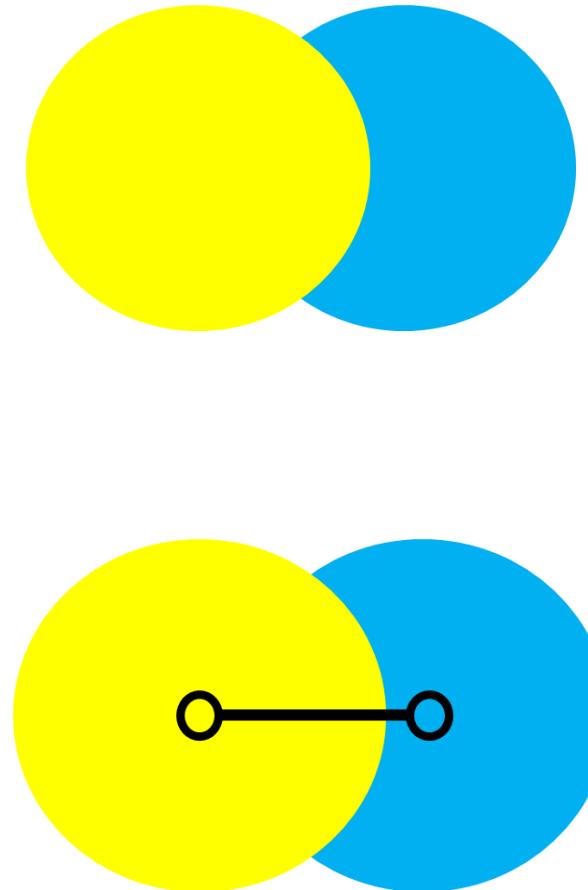
Hypervolume metrics – single

- Hypervolume size (volume) – total space occupied by a hypervolume
 - Variation of input data
 - Trophic niche size
 - Trait diversity
 - Spatial variability
- Centroid – center point of hypervolume
 - Mean conditions / location in hypervolume space
 - Community weighted mean

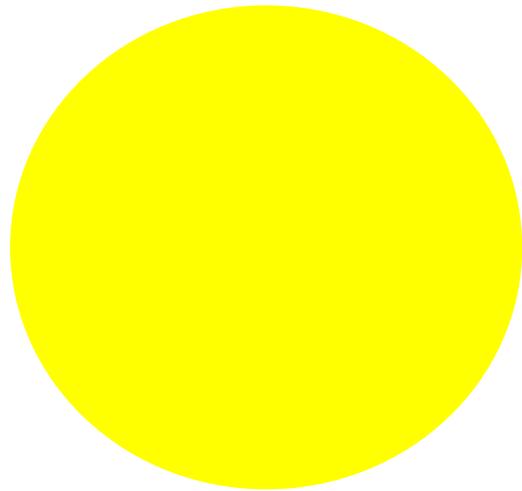


Hypervolume metrics – comparison

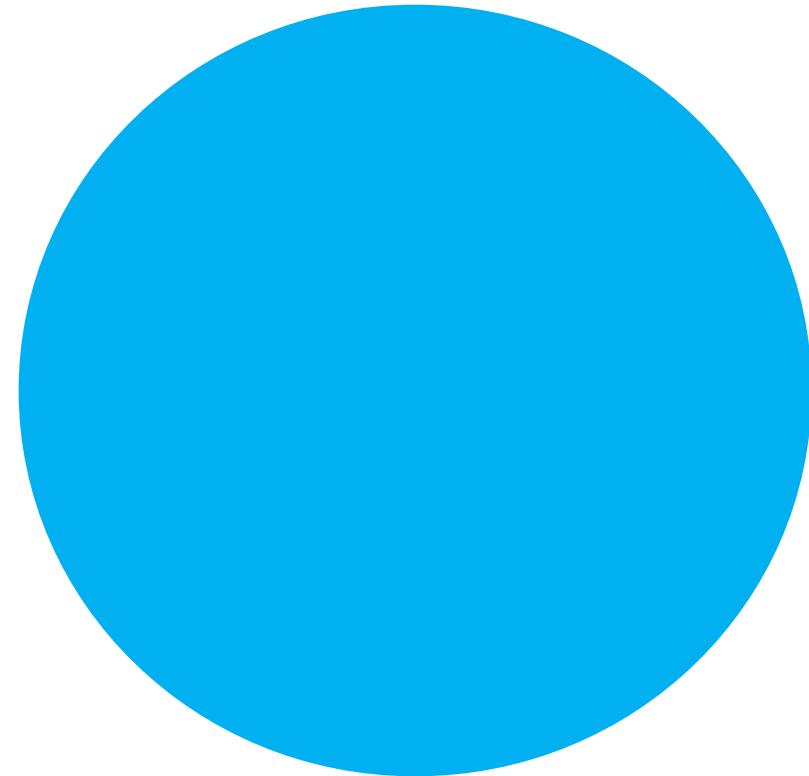
- Hypervolume overlap – amount of space two hypervolumes share relative to overall space of two hypervolumes
 - Similarity
 - Niche overlap/packing/resource partitioning
 - Trait similarity/packing
- Centroid distance – Euclidean distance between two hypervolume centroids
 - mean conditions / location in hypervolume space
 - Stability
 - Niche/trait shift



Hypervolume size

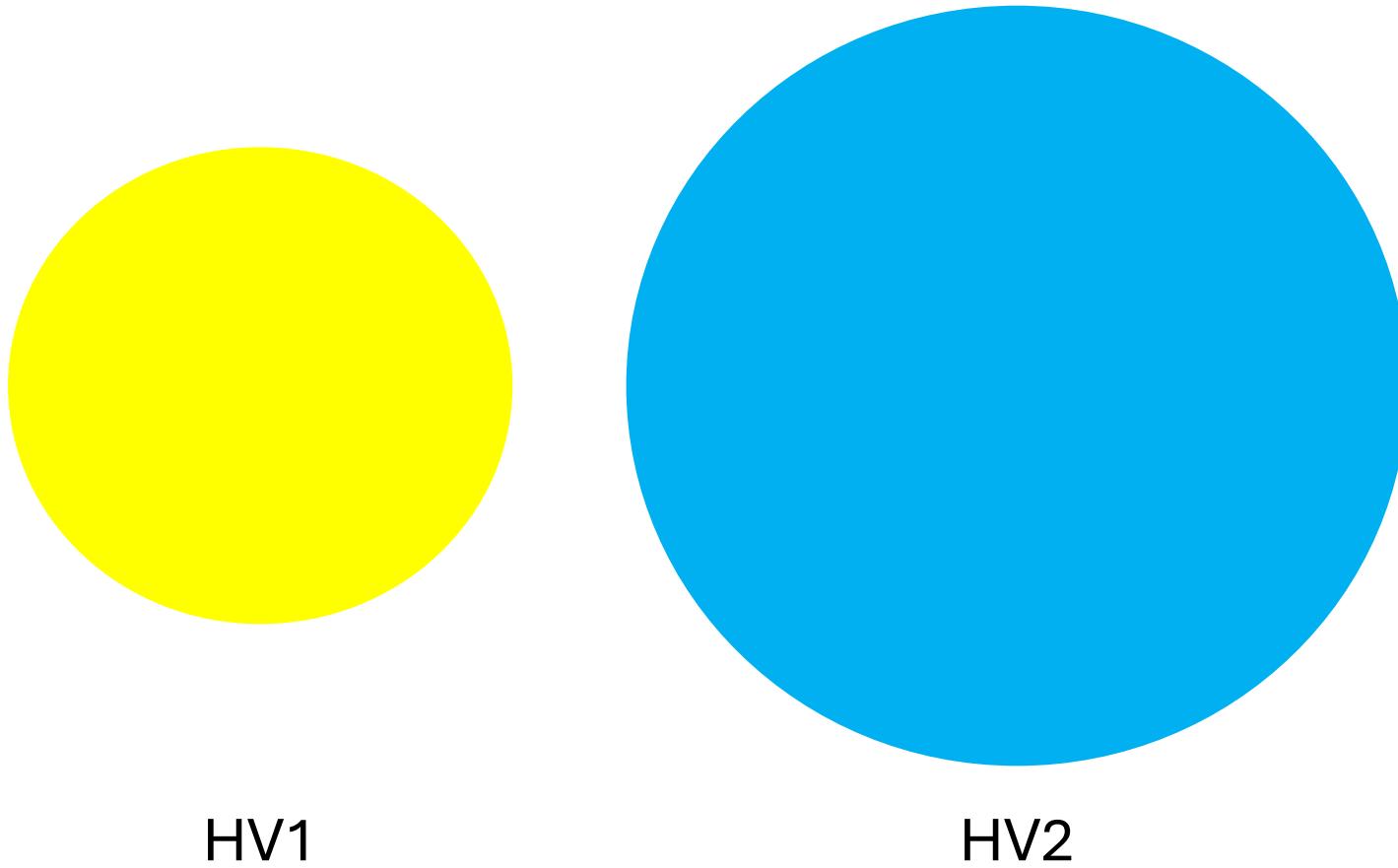


HV1



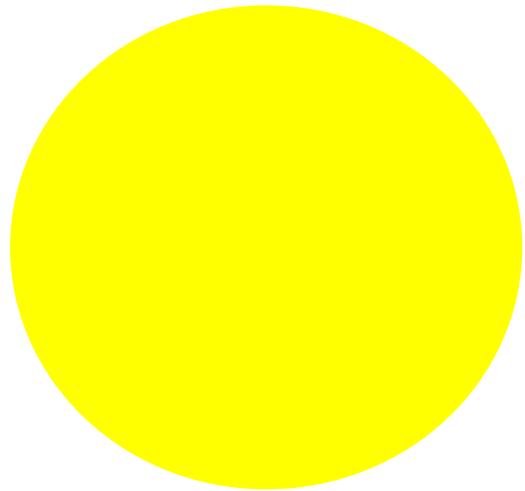
HV2

Hypervolume size

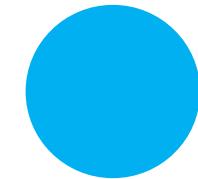


HV1 variation < HV2 variation

Hypervolume size

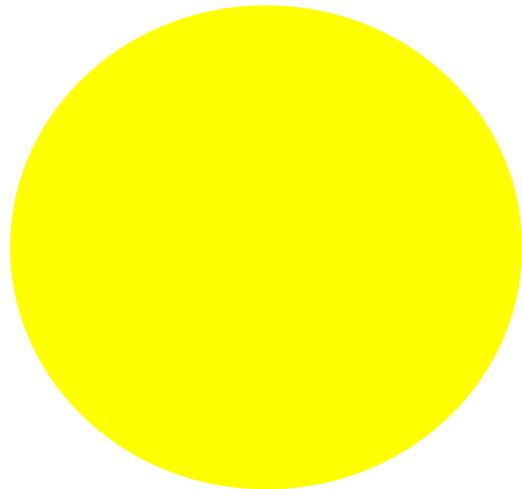


HV1

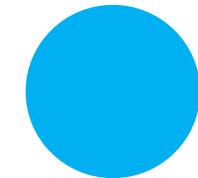


HV2

Hypervolume size



HV1

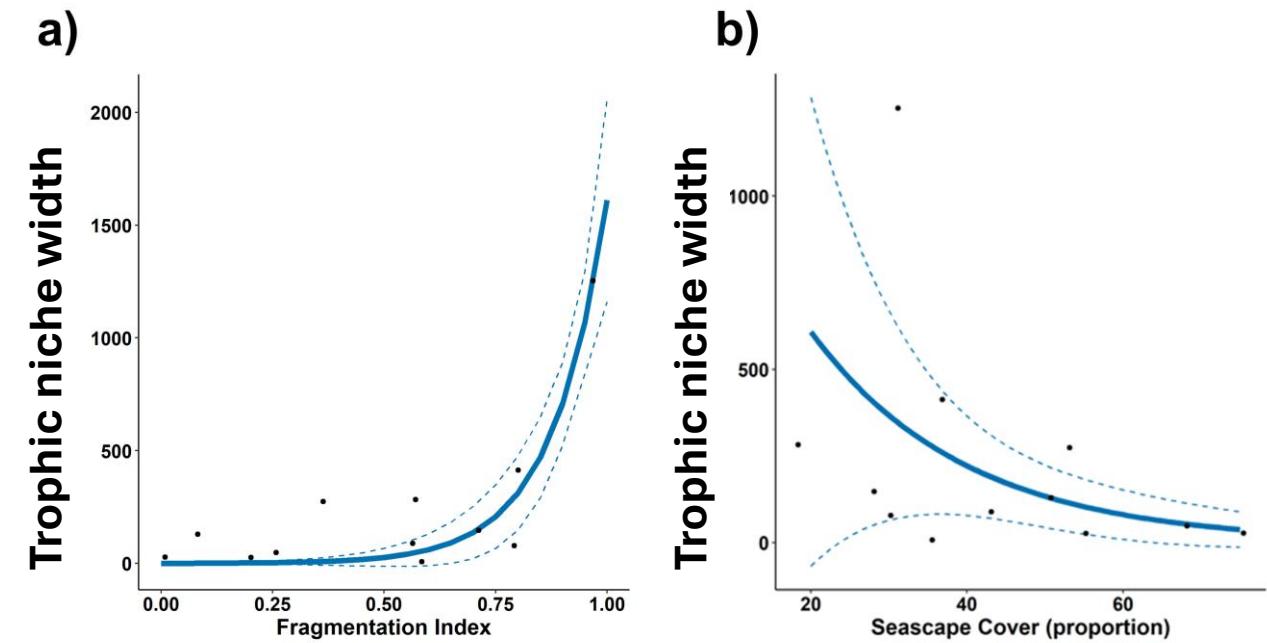
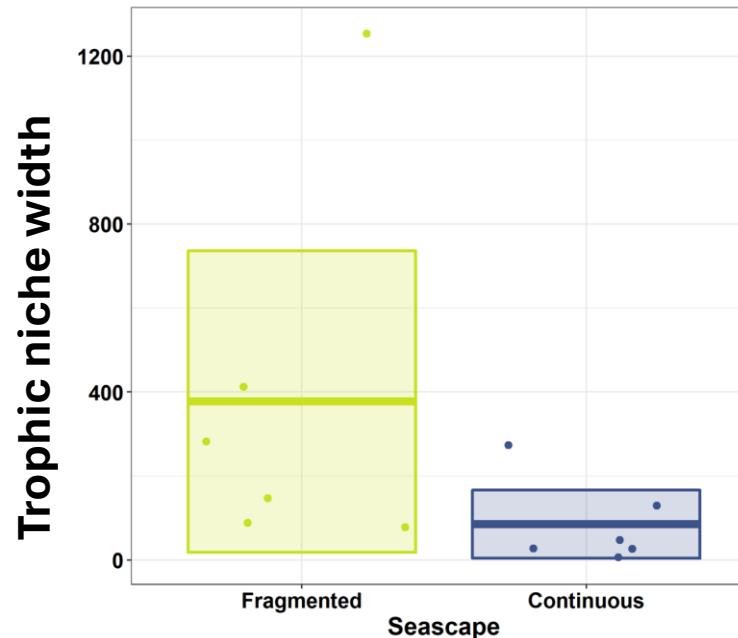


HV2

HV1 variation < HV2 variation

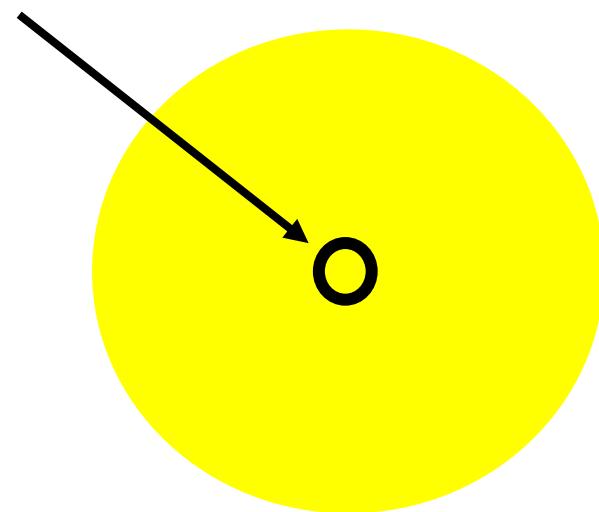
Hypervolume size to understand trophic niche width (Santos et al. 2022)

- Change in trophic niche based on seascape context
- Hypervolumes based on resource use and trophic level of pinfish

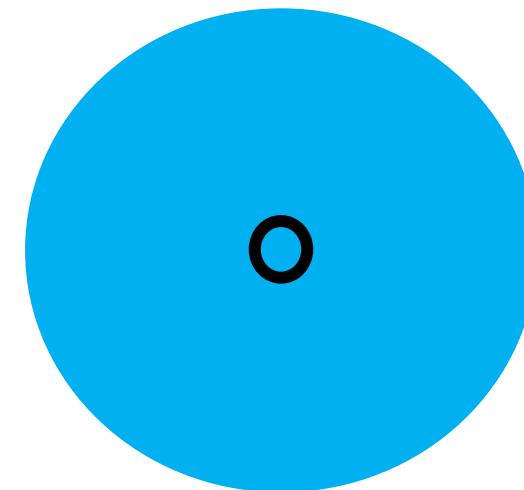


Centroid distance

Centroid

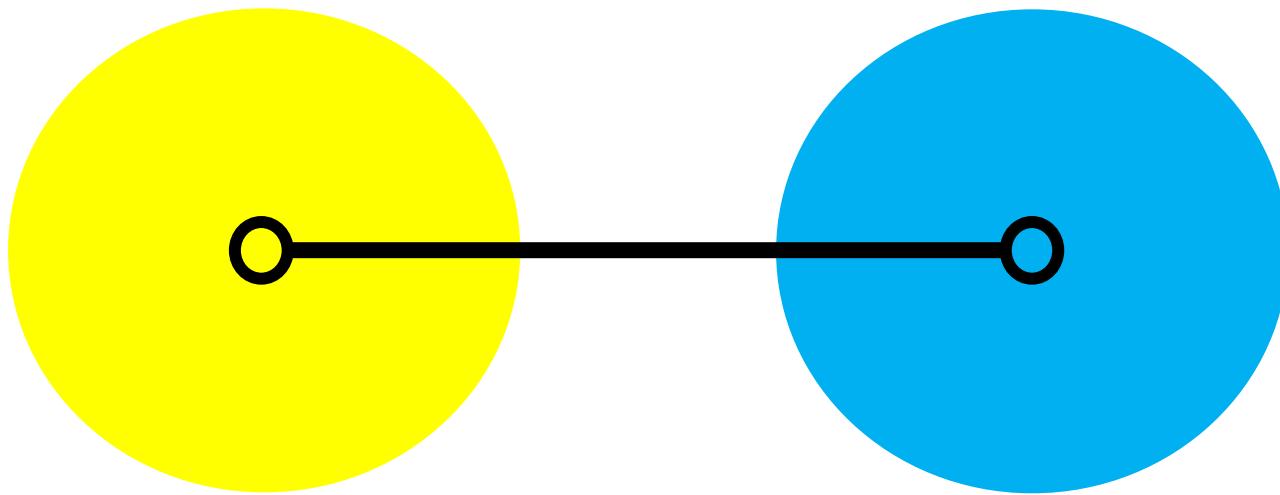


HV1



HV2

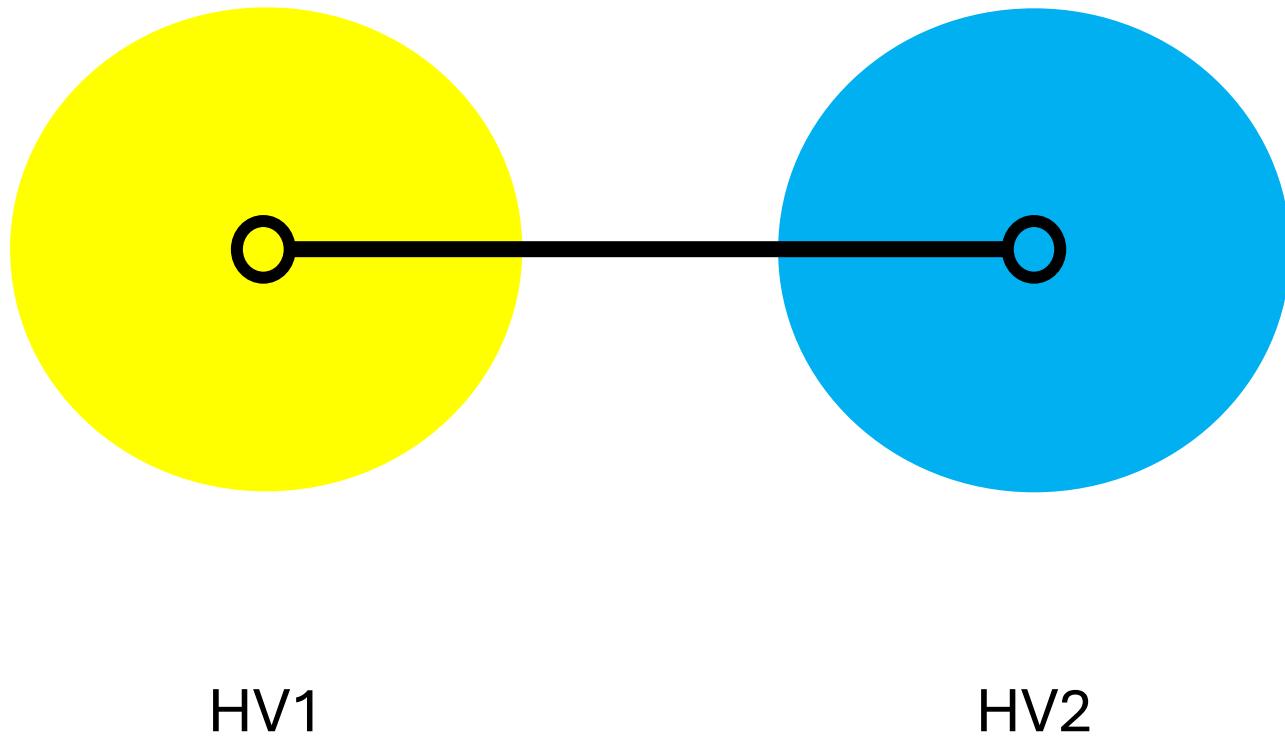
Centroid distance



HV1

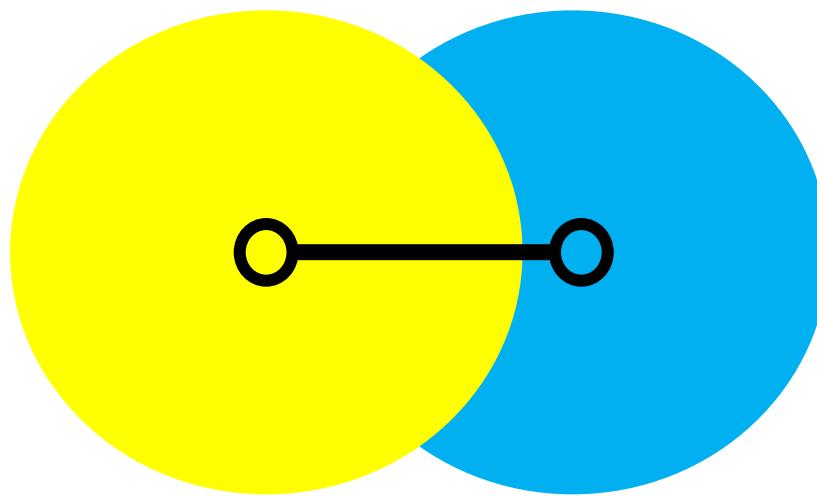
HV2

Centroid distance



HV1 and HV2 big difference in mean conditions

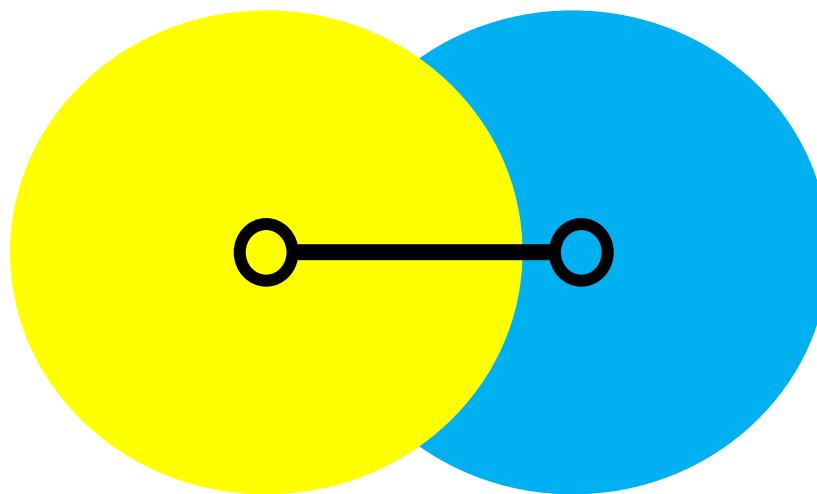
Centroid distance



HV1

HV2

Centroid distance



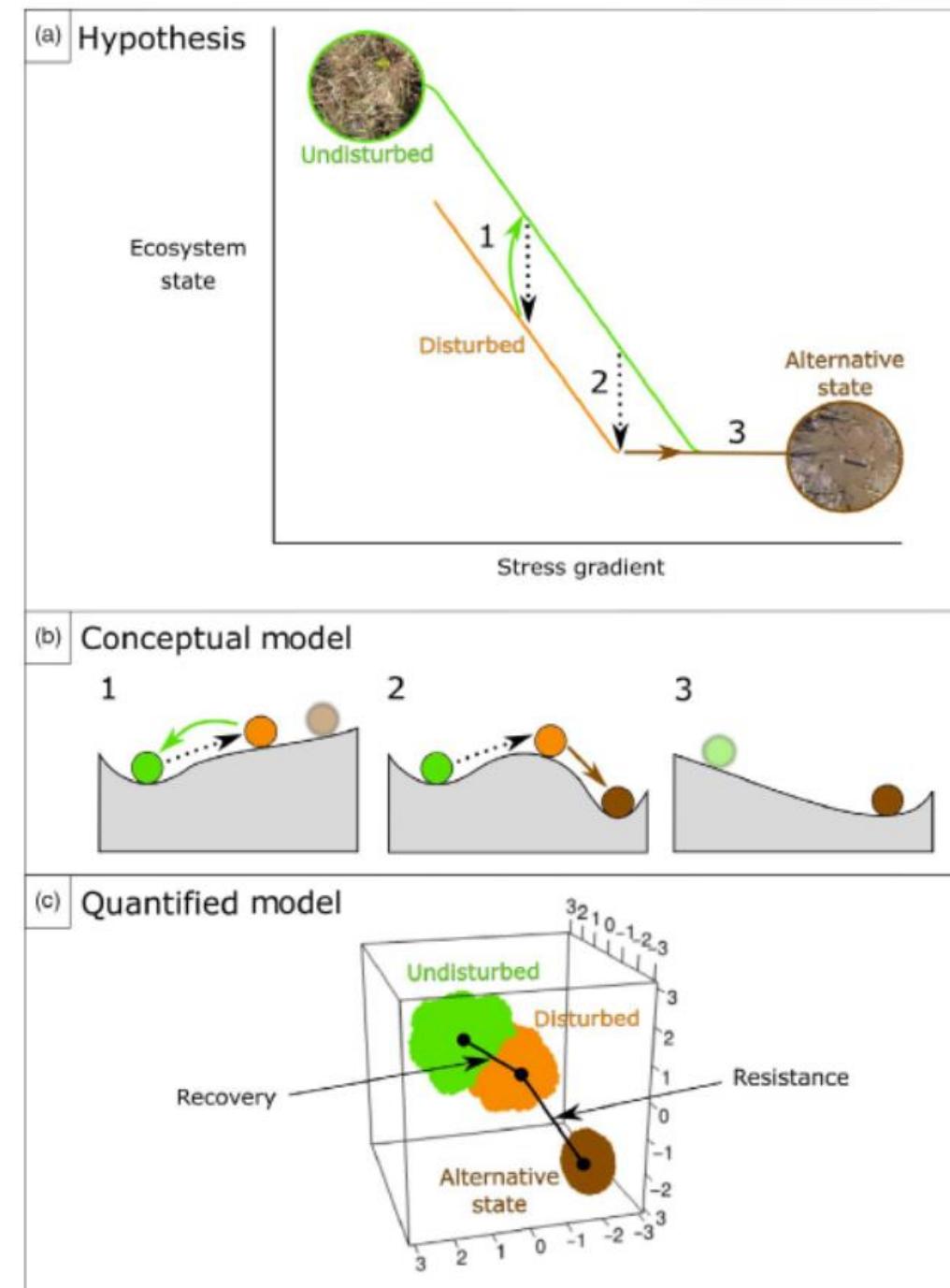
HV1

HV2

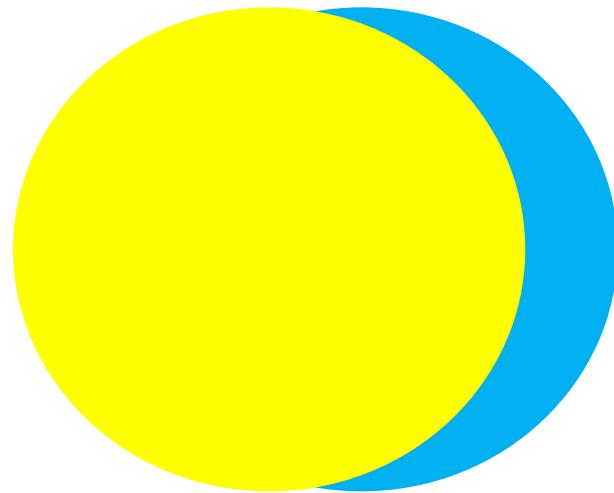
HV1 and HV2 small difference in mean conditions

Centroid distance to understand ecosystem resilience

- Jones et al. 2020 *Journal of Ecology*
- Hypervolumes to quantify state and recovery of marshes along stress gradient (flooding) to disturbance
- Recovery – centroid distance from disturbed to undisturbed
- Resistance – centroid distance from disturbed to alternate state



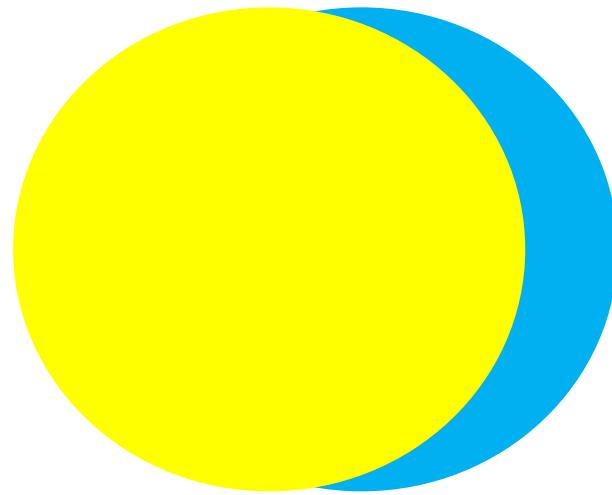
Hypervolume overlap



HV1

HV2

Hypervolume overlap

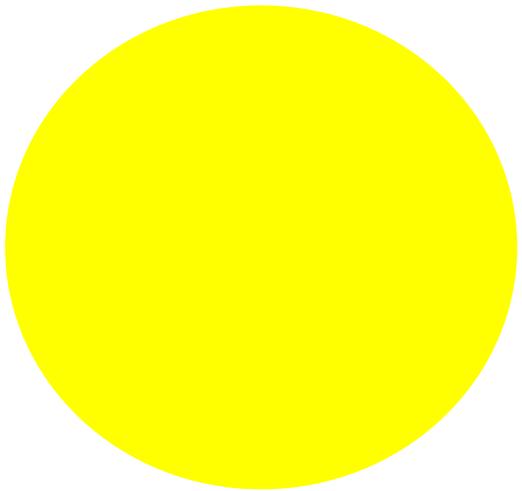


HV1

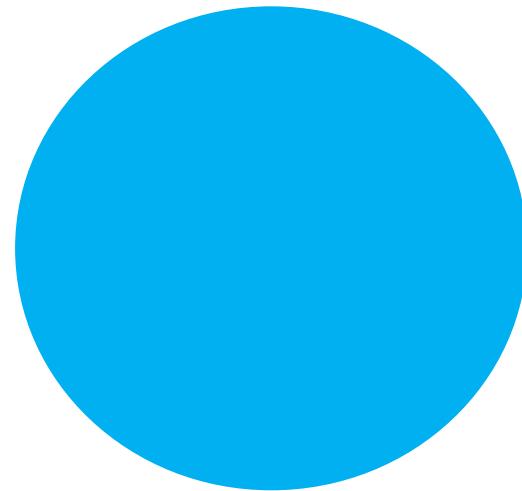
HV2

HV1 and HV2 have high similarity

Hypervolume overlap

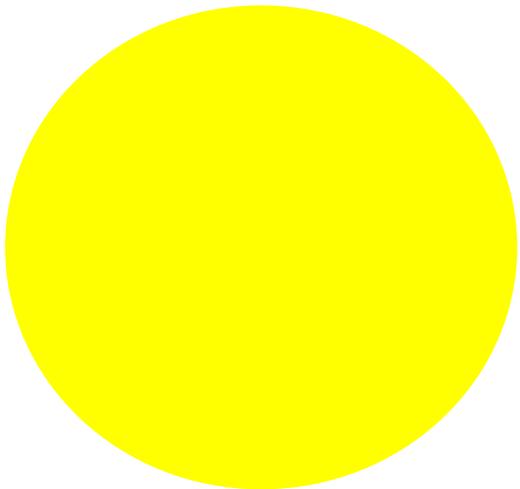


HV1

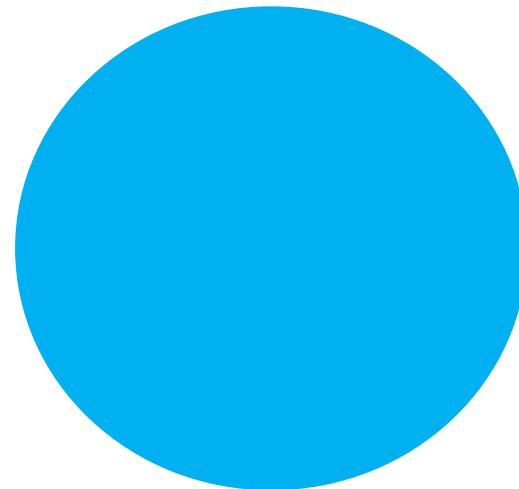


HV2

Hypervolume overlap



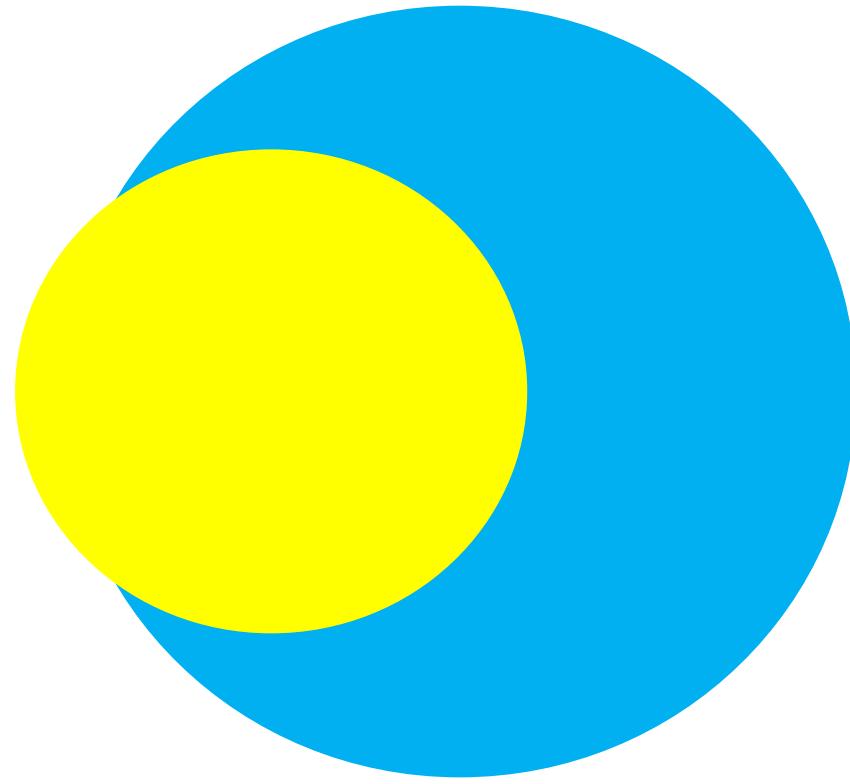
HV1



HV2

HV1 and HV2 have low similarity

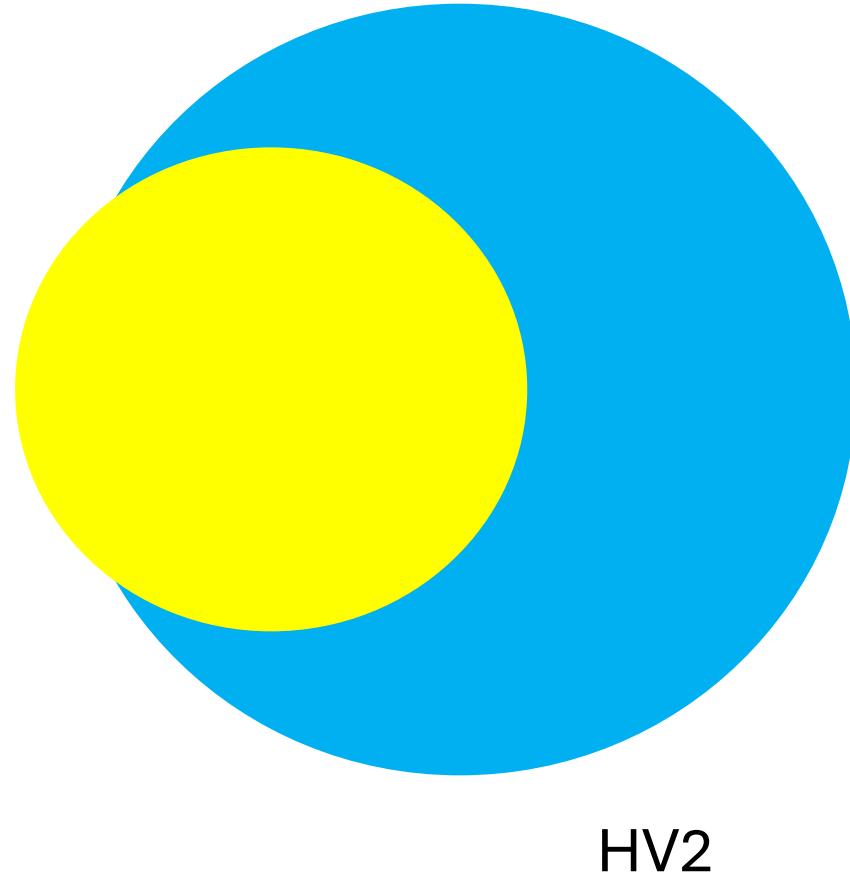
Hypervolume overlap



HV1

HV2

Hypervolume overlap

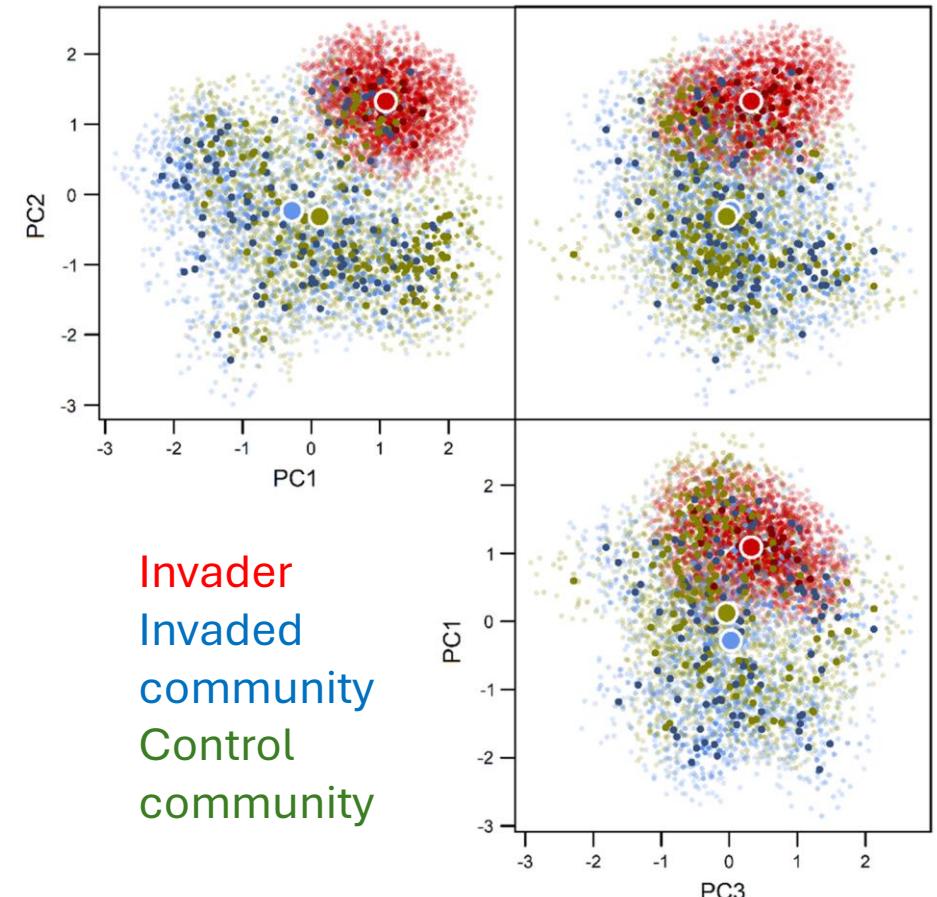


HV1 not unique and HV2 high % unique, so have low similarity

Overlap to understand trait similarity of invasive plant (Helsen et al. 2016)

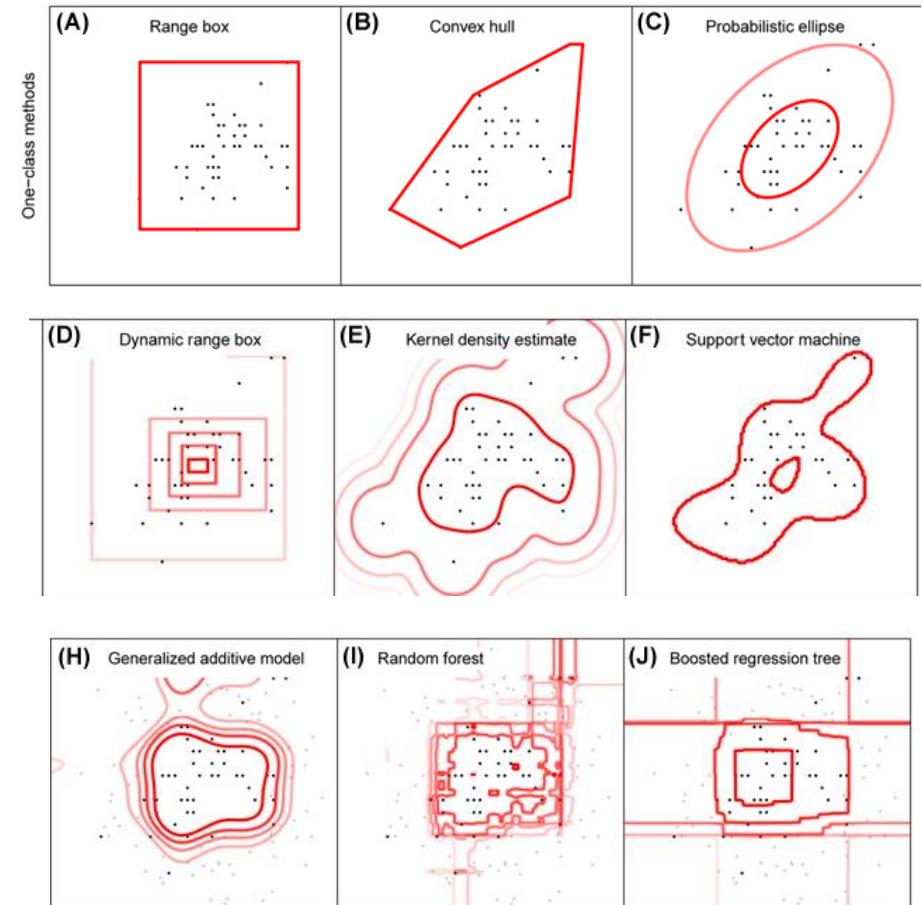
- PC on traits to reduce # of axes
- Test function distinctive hypothesis
 - Invader unique part of hv
 - Community shift native to minimize overlap

	Centroid distance (SD)	Jaccard similarity	Unique volume fraction (%)
<i>Impatiens glandulifera</i>			
NAT _C – NAT _I	0.251 (0.201–0.302)	0.529 (0.513–0.544)	27.4 (25.7–29.2) – 33.9 (32.0–36.2)
	0.217 (0.123–0.314)	0.543 (0.514–0.571)	25.0 (22.5–27.5) – 33.7 (29.9–37.2)
NAT _C – NAT _T	0.115 (0.067–0.174)	0.748 (0.724–0.770)	15.2 (12.7–18.2) – 13.6 (11.3–16.5)
NAT _C – INV	2.152 (2.121–2.185)	0.029 (0.023–0.033)	96.5 (96.0–97.2) – 85.7 (83.5–88.4)
	2.223 (2.165–2.285)	0.030 (0.022–0.037)	96.5 (95.7–97.4) – 83.4 (79.5–88.0)
NAT _I – INV	2.046 (2.026–2.063)	0.051 (0.047–0.054)	94.1 (93.7–94.6) – 73.3 (71.6–75.7)
	2.168 (2.109–2.212)	0.043 (0.034–0.051)	95.1 (94.3–96.1) – 73.8 (68.9–79.4)
NAT _T – INV	2.124 (2.104–2.141)	0.036 (0.032–0.039)	95.6 (95.3–96.2) – 82.4 (81.1–84.6)
<i>Rosa rugosa</i>			
NAT _C – NAT _I	0.703 (0.657–0.758)	0.461 (0.441–0.477)	28.3 (26.3–30.1) – 43.6 (41.3–46.0)
	0.471 (0.326–0.633)	0.496 (0.460–0.538)	27.8 (23.7–32.6) – 38.6 (33.8–42.9)
NAT _C – NAT _T	0.238 (0.172–0.286)	0.781 (0.752–0.809)	9.9 (7.8–12.4) – 14.6 (11.7–18.3)
NAT _C – INV	1.978 (1.928–2.015)	0.052 (0.047–0.057)	94.1 (93.6–94.8) – 68.1 (65.5–71.2)
	2.112 (2.027–2.212)	0.042 (0.033–0.048)	95.5 (94.8–96.4) – 66.7 (61.8–72.9)
NAT _I – INV	2.361 (2.342–2.379)	0.024 (0.021–0.025)	97.3 (97.2–97.7) – 81.7 (80.3–84.1)
	2.400 (2.311–2.487)	0.018 (0.012–0.023)	98.0 (97.5–98.6) – 82.9 (78.5–88.3)
NAT _T – INV	2.096 (2.067–2.113)	0.049 (0.047–0.051)	94.5 (94.3–94.7) – 68.5 (67.3–69.7)



Multivariate approaches

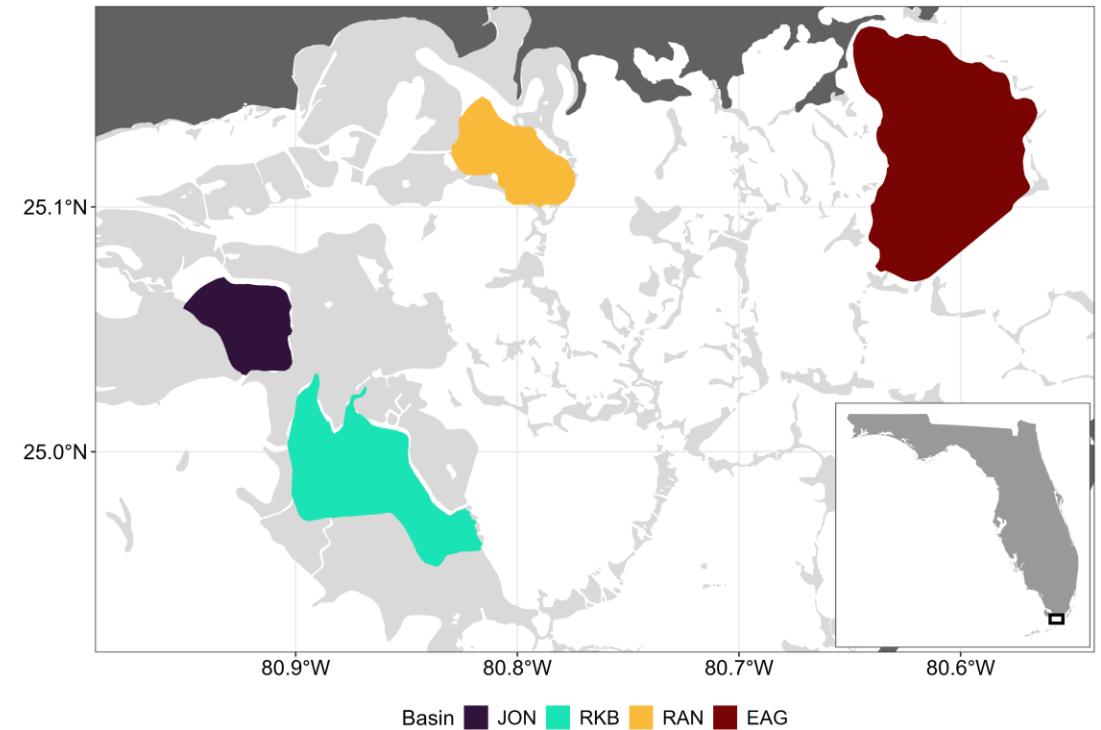
- Based on data and question
- Take advantage of variation
 - Allow for holes in data distribution
- Blonder et al. 2014 *GEB*
- Blonder et al. 2017 *Ecography*



Blonder et al. 2017. Ecography

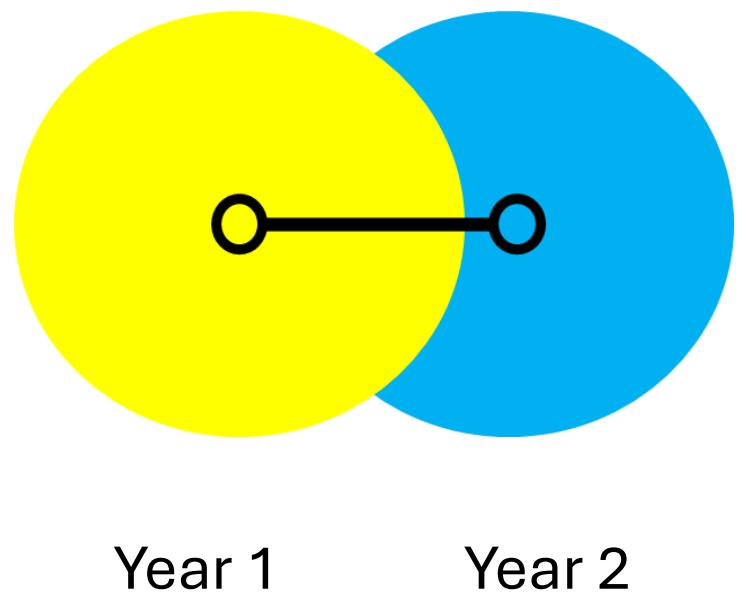
Case study 1 – Centroid distance SAV community stability

- Temporal stability of SAV communities in Florida Bay
- Monitoring data ~ 30 sites per basin
- HV – Percent cover of *Thalassia testudinum*, *Halodule wrightii*, *Syringodium filiforme*, total macroalgae, and total drift algae and seagrass richness
- Centroid distance across time (17 years)

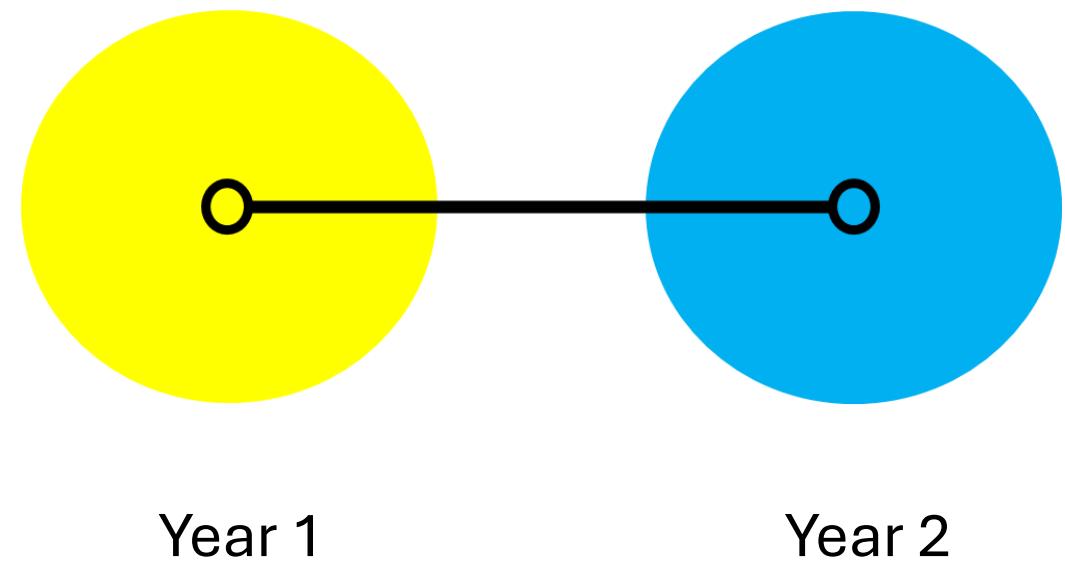


Stability – change in mean conditions

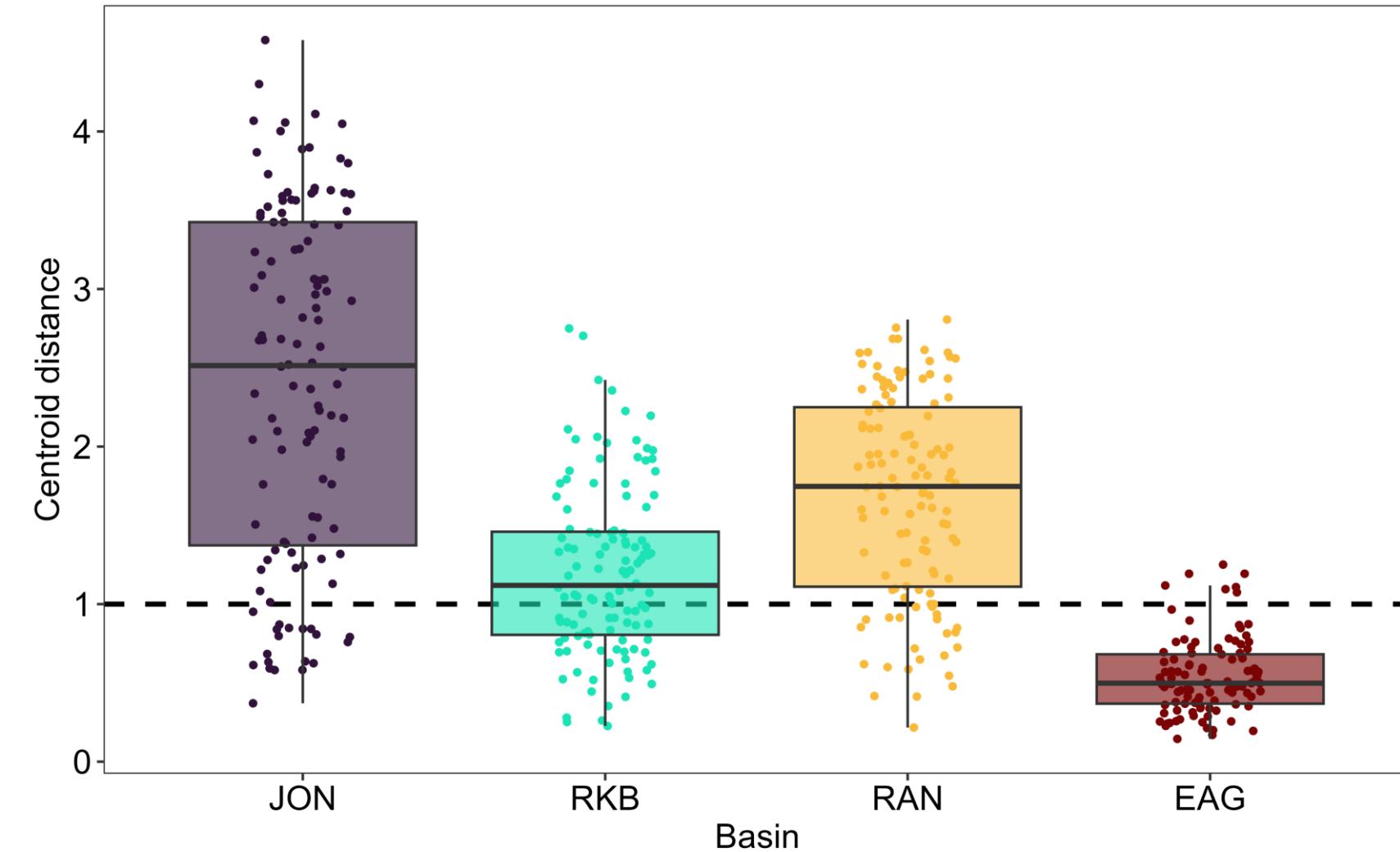
High temporal stability



Low temporal stability

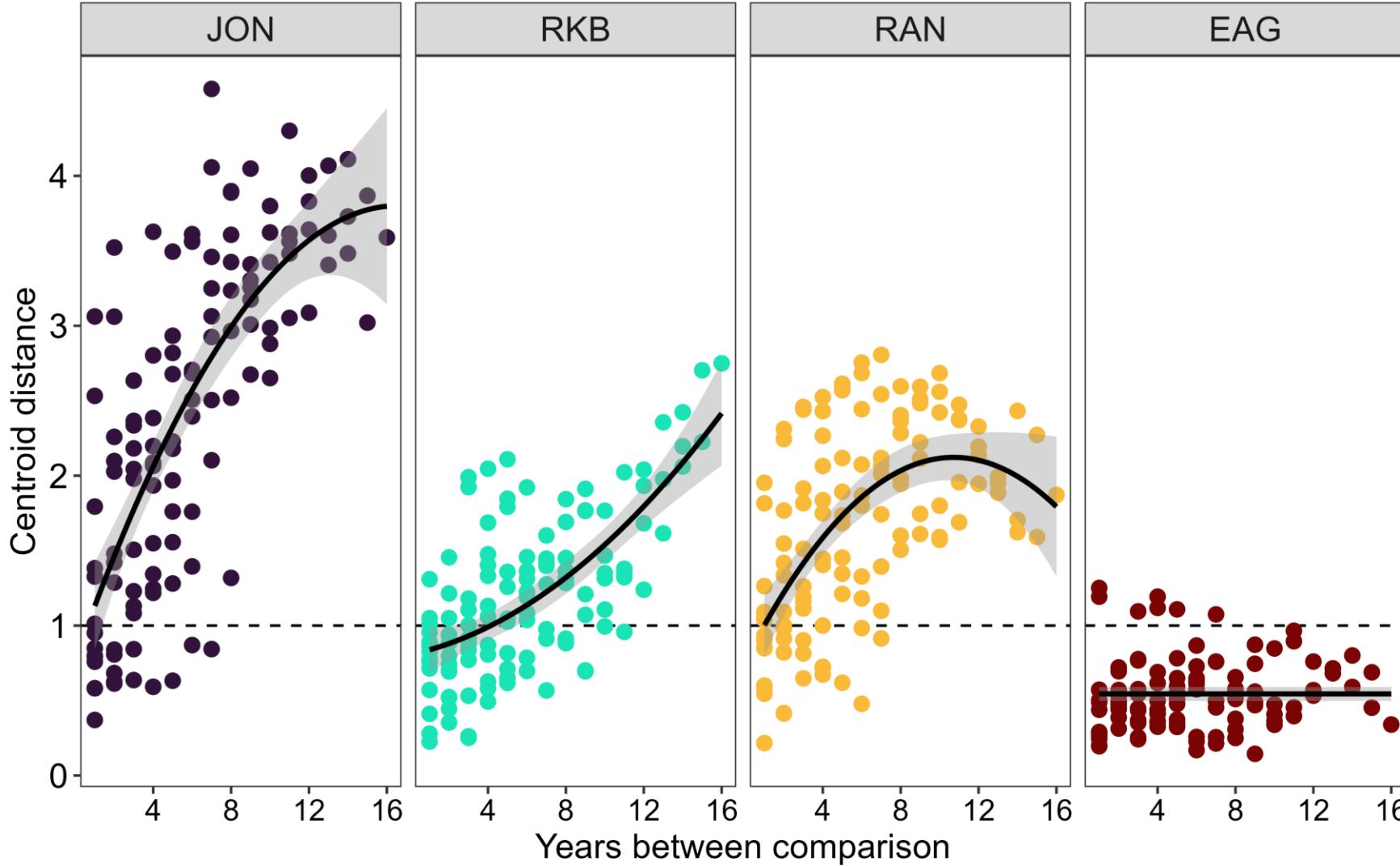


Temporal stability in mean conditions



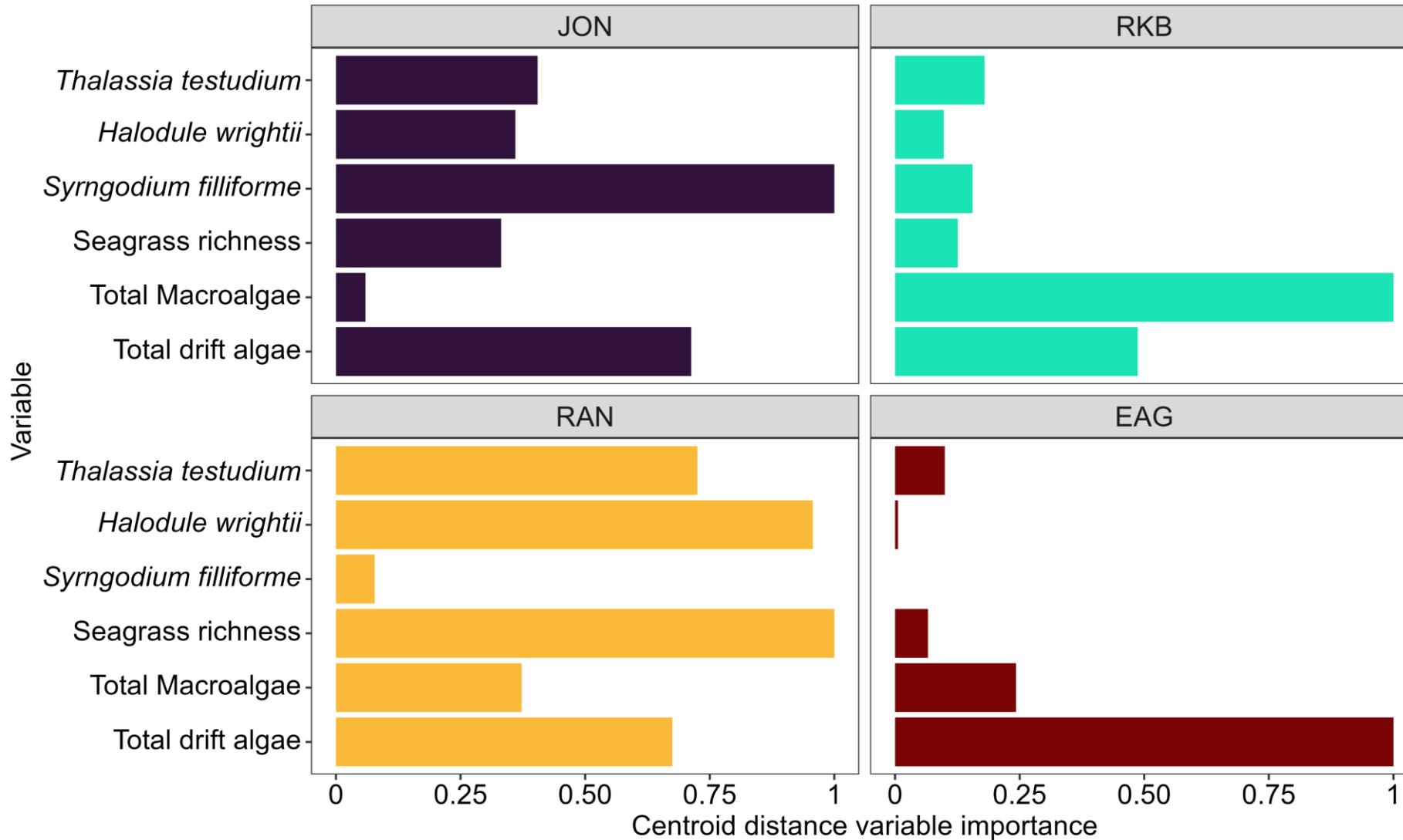
- Variation in stability over across basins

Transitions in mean conditions/state



- JON – shift with stabilization/recovery
- RKB – shift
- RAN – recovery
- EAG – stable

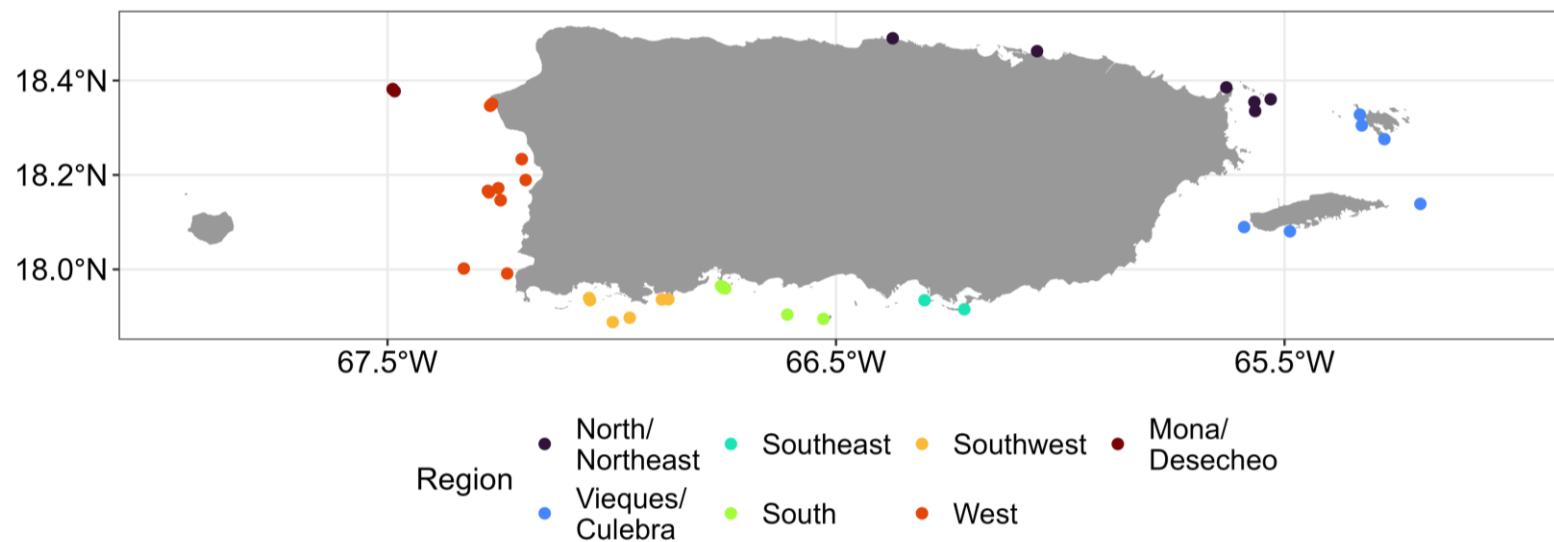
Variables contributing to state change



- Spatial variation in the variable contributing most to the centroid distance between years

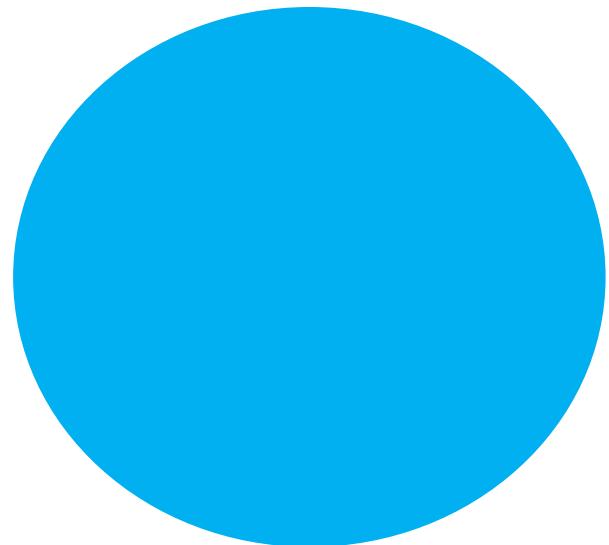
Case study 2 – Hypervolume size Coral trait diversity

- How does trait diversity vary across regions
- Benthic monitoring of coral communities combined with literature trait values
- HV – community weighted (using percent cover) for 5 functional traits based on random percent cover
- Hypervolume size for trait diversity



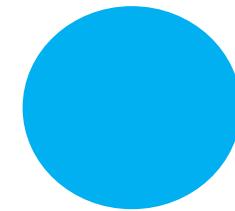
Trait diversity – variation in traits across community

High diversity

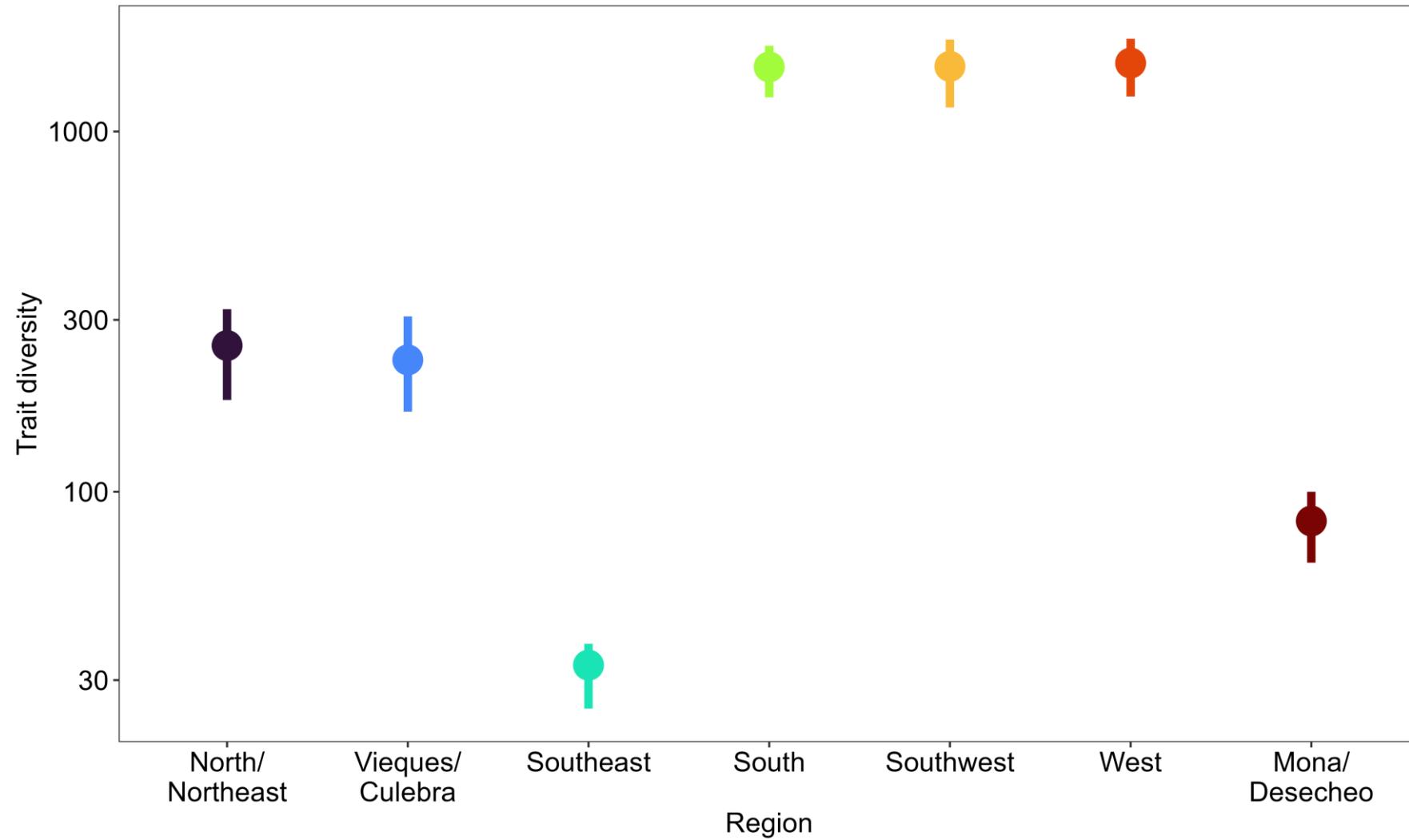
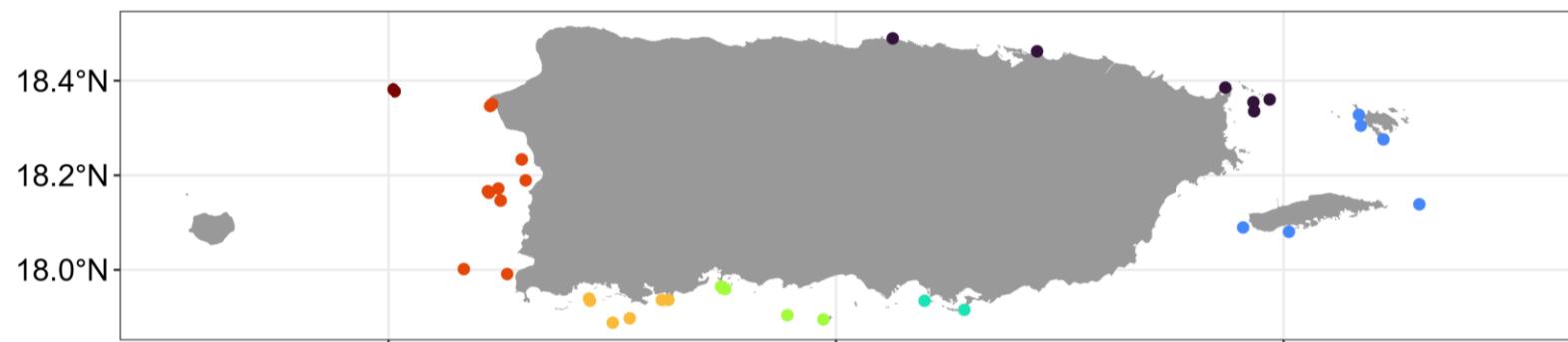


Region

Low diversity

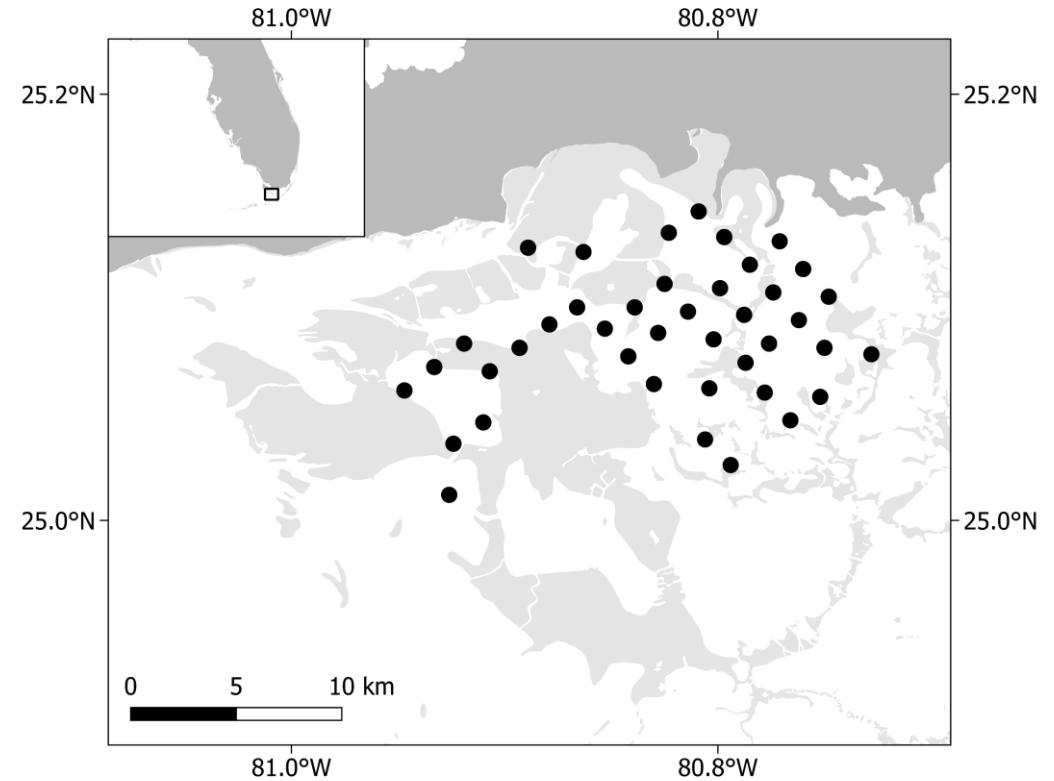


Region

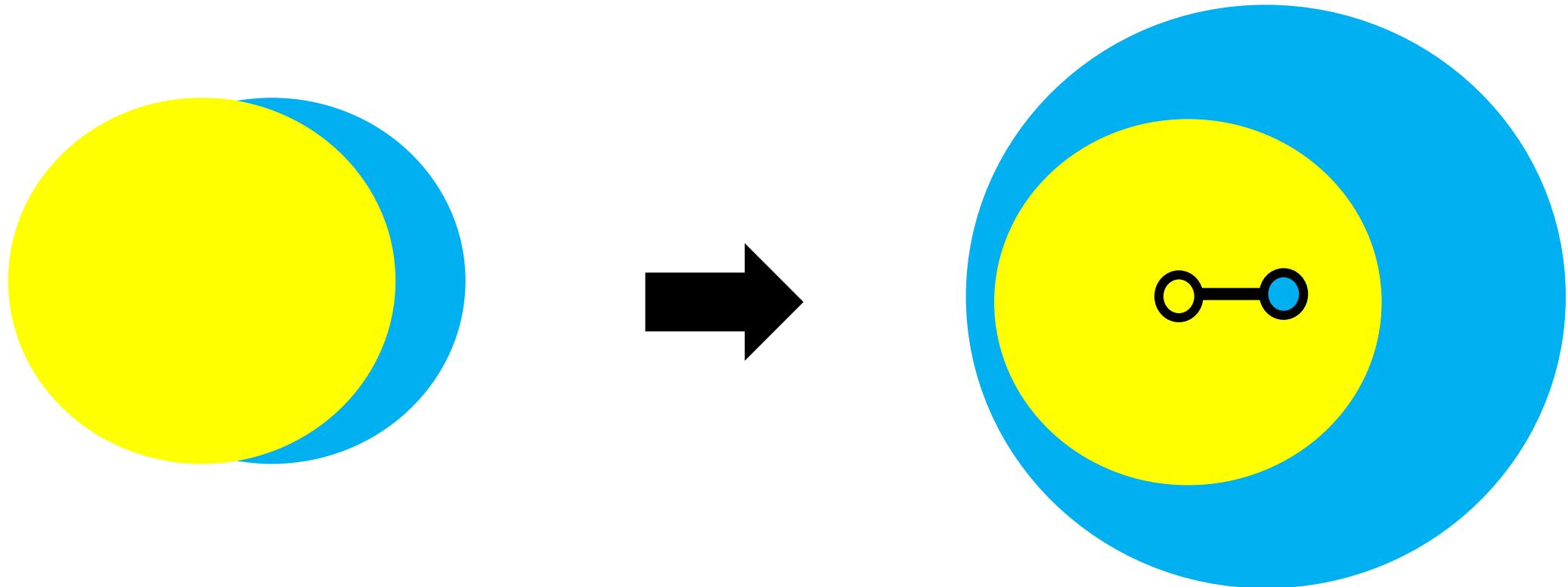


Case study 3 – overlap Trophic niche seagrass consumers

- Seasonal variation in trophic niche dynamics 7 seagrass consumers
- Axes – basal resource use (seagrass, epiphytes, algae, mangrove)
- HV – each species and season based on random points from mean and sd of resource use
- Overlap, size, and centroid distance

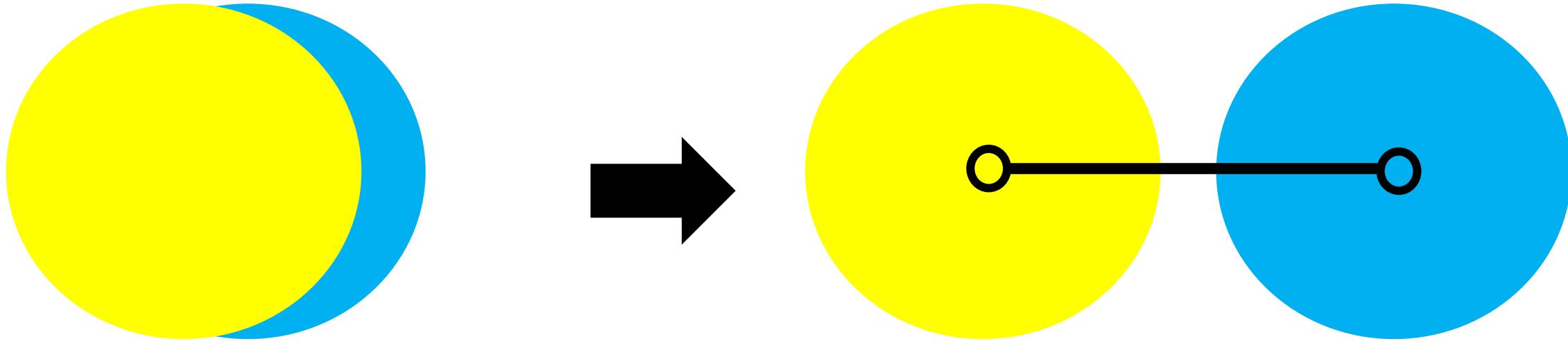


Niche expansion/contraction



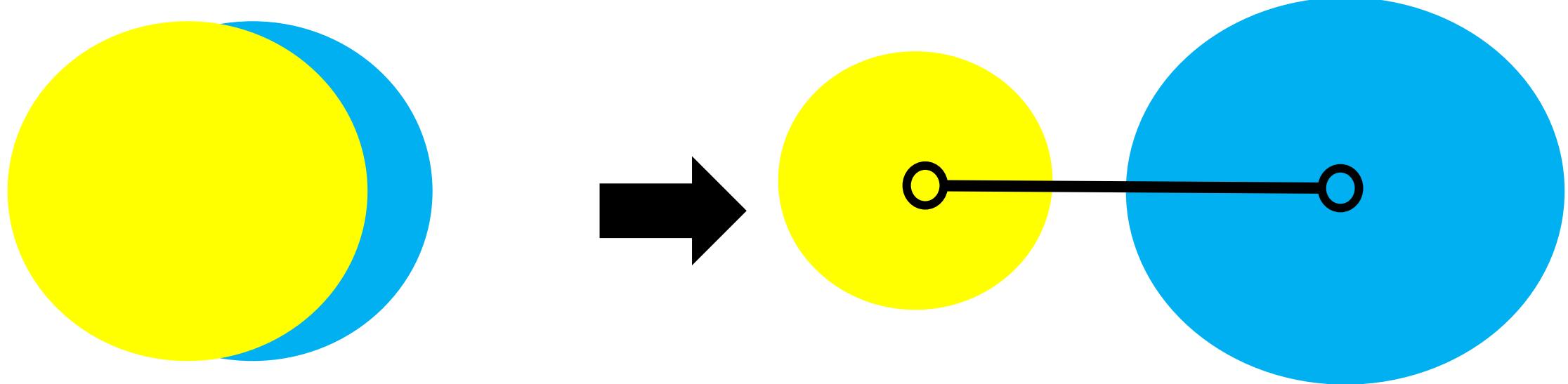
- Difference in size
- Small centroid distance
- Low overlap
- Niche unique high for hv1, low for hv2

Niche shift



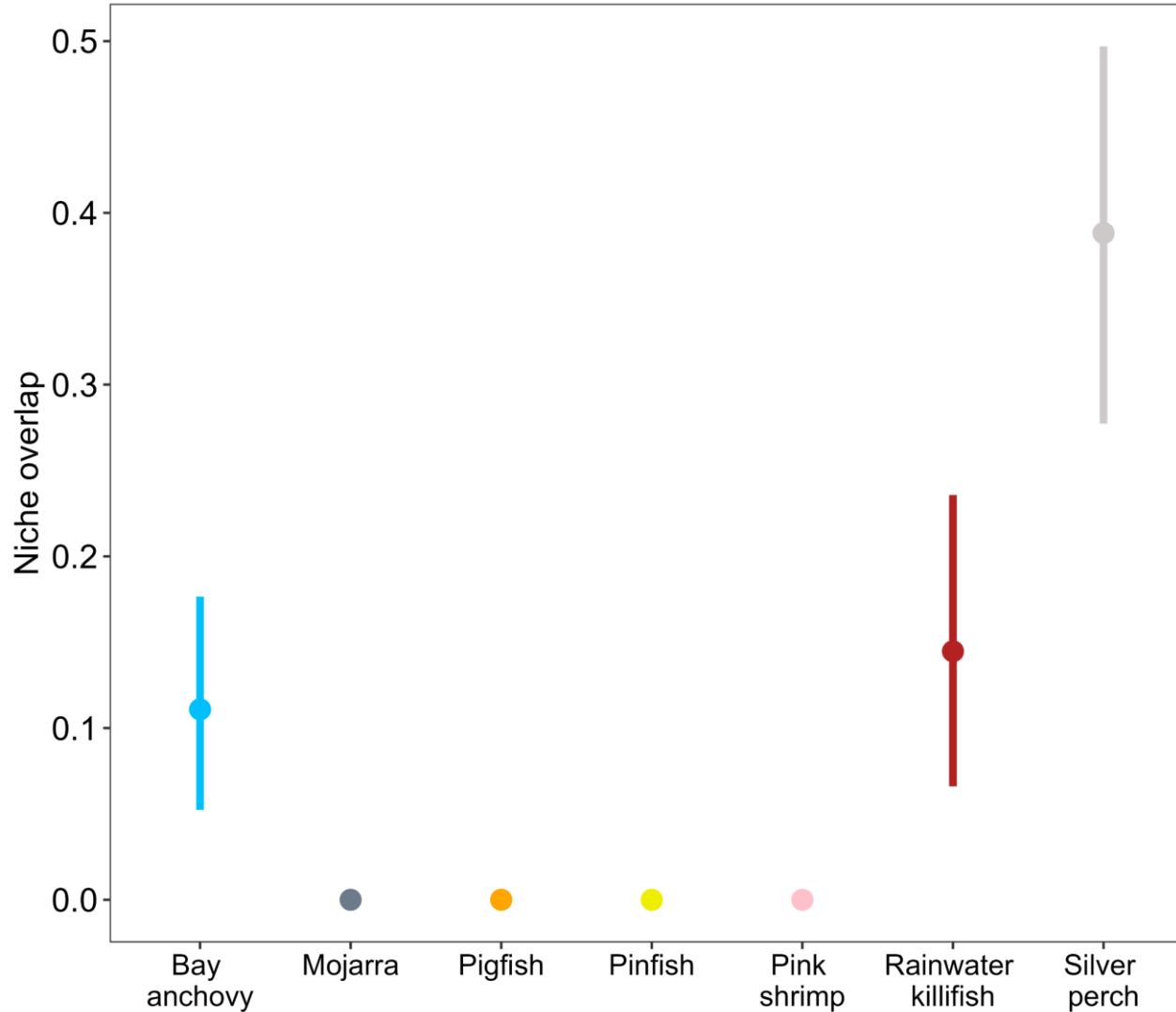
- No difference in size
- Large centroid distance
- Low overlap
- Niche unique high for hv1 and hv2

Niche expansion/contraction and niche shift



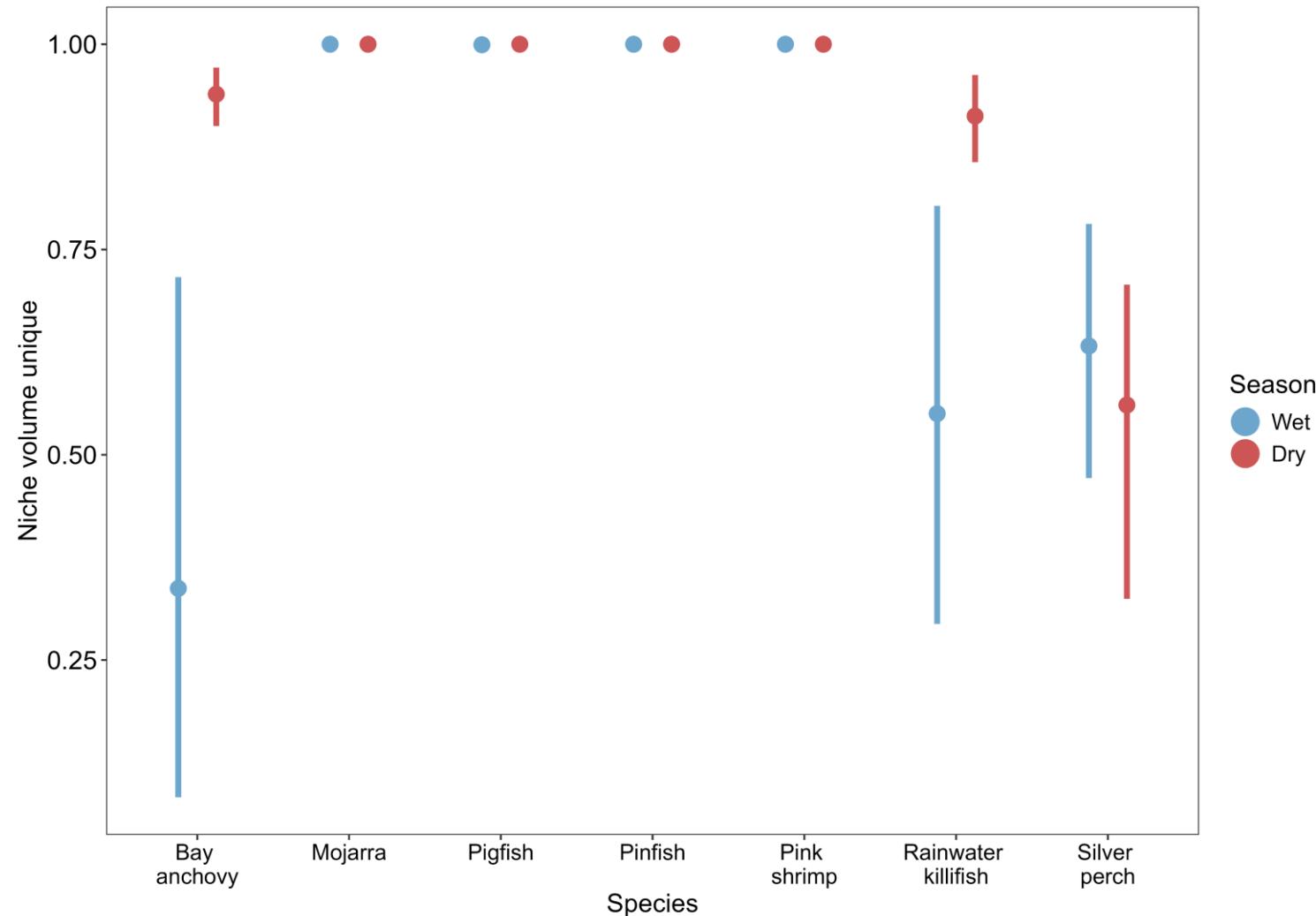
- Difference in size
- Large centroid distance
- Low overlap
- Niche unique high for hv1 and hv2

Niche similarity between seasons



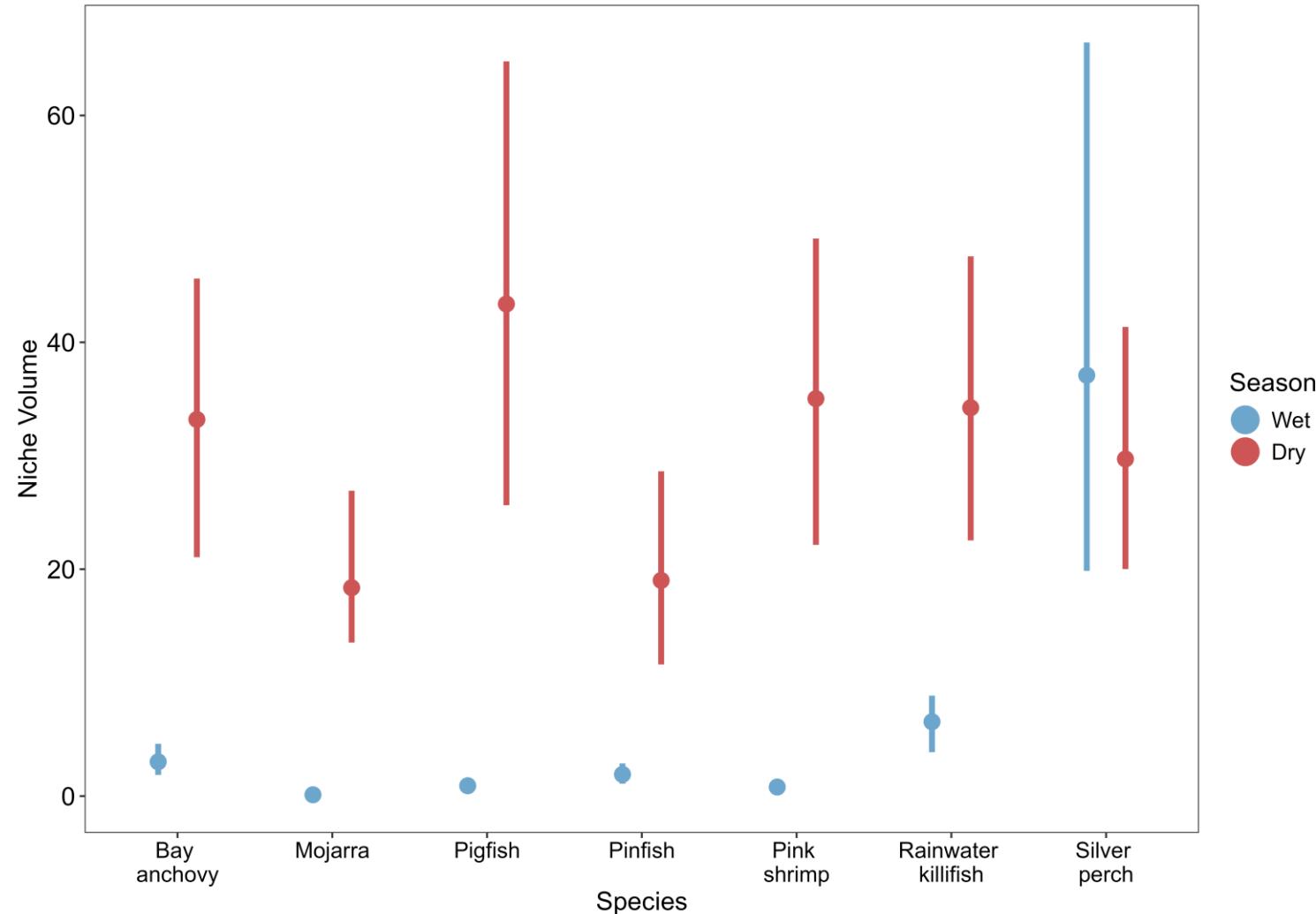
- No overlap for mojarra, pigfish, pinfish, and pink shrimp
- Intermediate/low overlap for bay anchovy and rainwater killifish
- Higher overlap for silver perch

Unique niche for each season



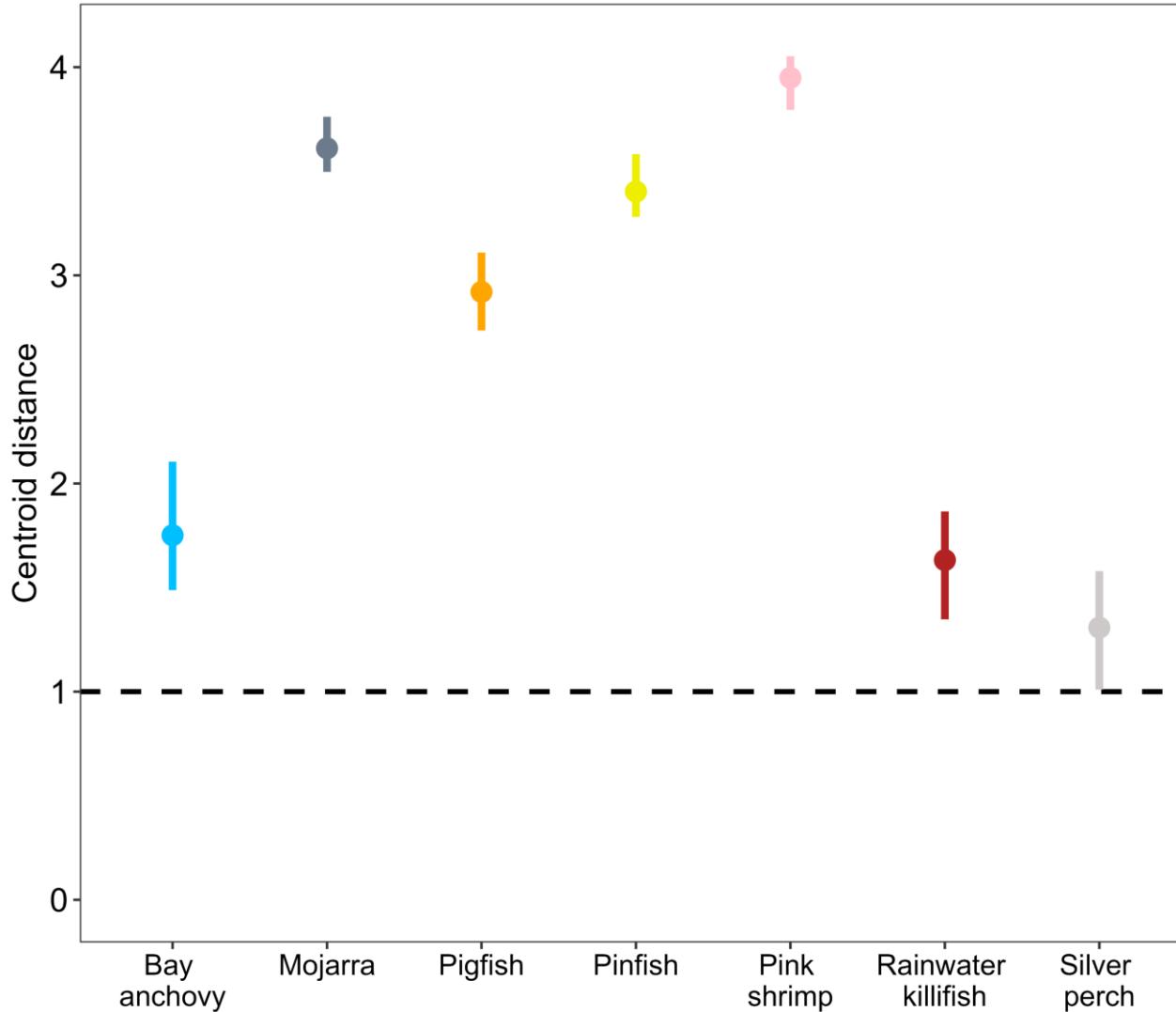
- 100% unique for mojarra, pigfish, pinfish, and pink shrimp in both seasons
- High unique for dry and low for wet for bay anchovy and rainwater killifish
- Low unique for silver perch in both seasons

Changes in trophic niche size



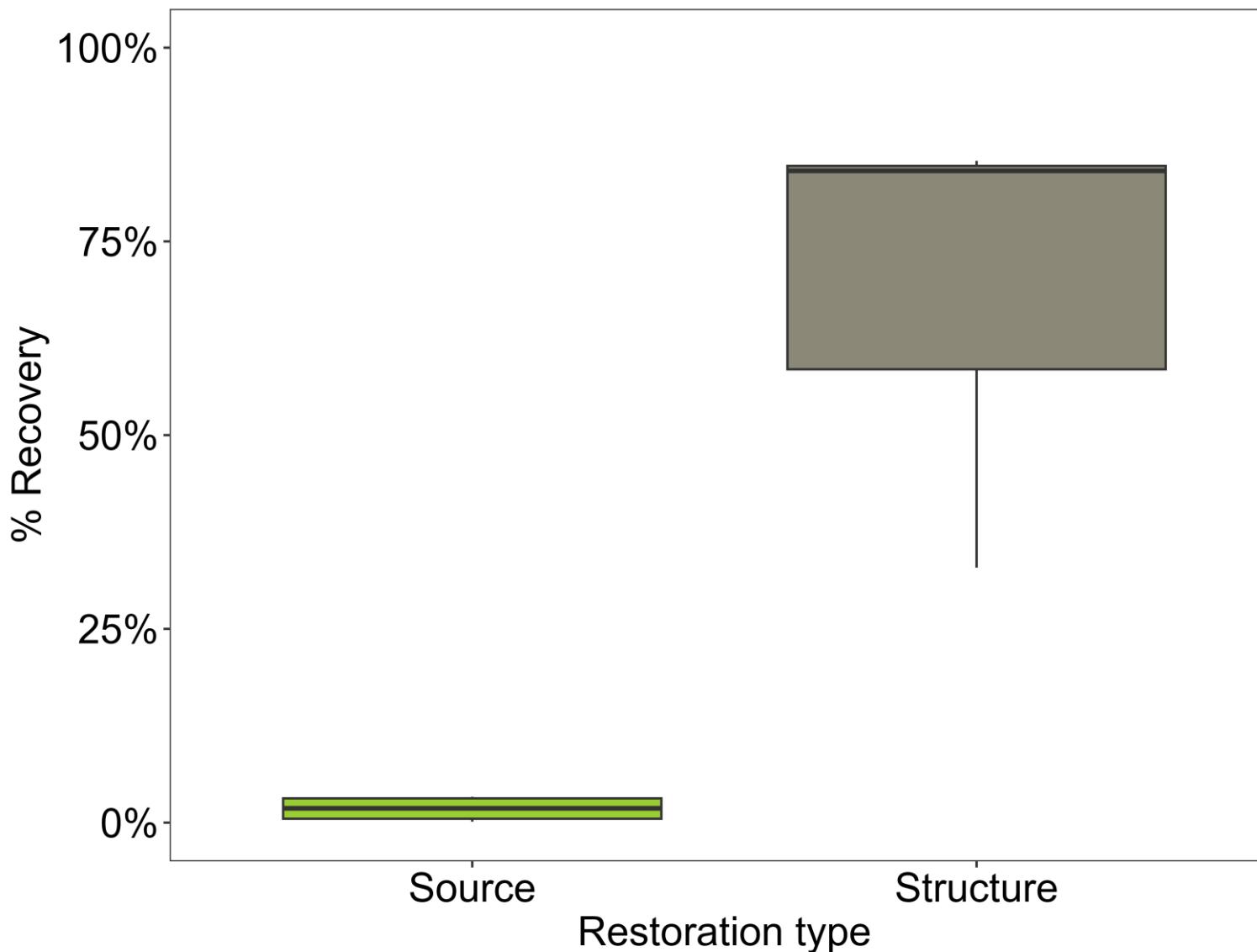
- Larger in dry for all species except
- Silver perch no difference between seasons

Changes in mean resource use



- Large shifts for mojarra, pigfish, pinfish, and pink shrimp
- Smaller shift for bay anchovy, rainwater killifish, and silver perch

Overlap to quantify restoration recovery (James et al. 2020)



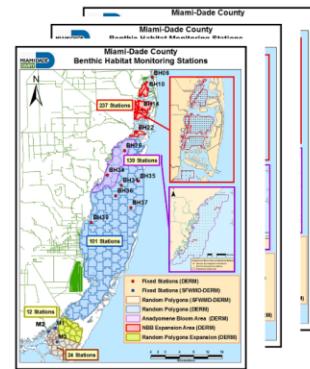
- Recovery slower in restorations of biogenic habitat than abiotic habitat structure
- Recovery of ecosystem function unrelated to age of restoration

Seagrass state

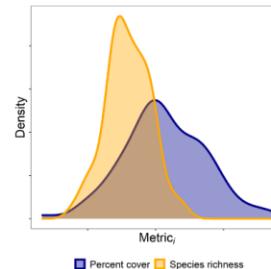


Objective 1 – Quantify Seagrass State and Stability

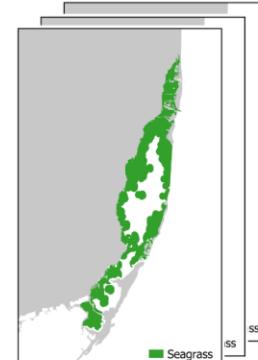
Monitoring Data



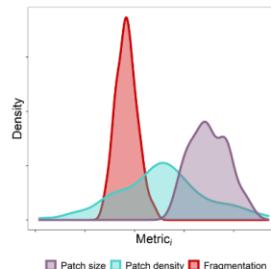
SAV metrics



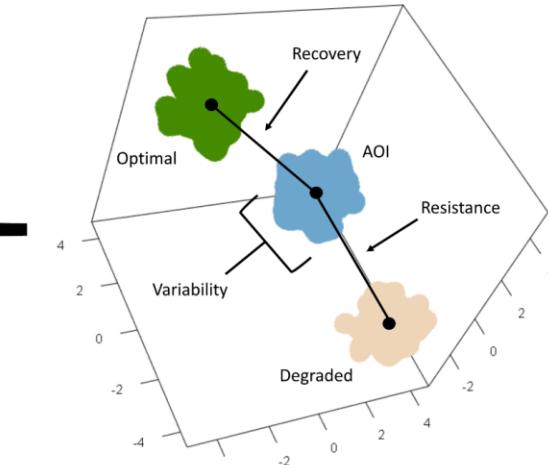
Remote sensing



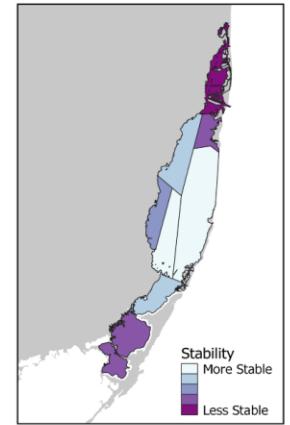
Seascape metrics



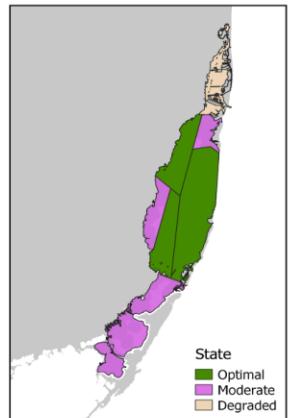
Hypervolumes



Stability



State



@santos_seascape_ecology



wjames@fiu.edu, rsantosc@fiu.edu