

Background





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* Preliminary results, subject to revision and not for citation

Created coastal marshes have equivalent diversity to natural coastal marshes: A case study from Louisiana

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Study sit

Figure 3. Among-site rarefied diversities for individual *plots/samples (α; x-axis) among-samples (β; insets),* and total site (*y*; *y*-axis) for all taxonomic groups.

 α -diversity (S_N

taxonomic groups. ----->

cross-taxonomic	analysis of ma	arsh biodiversit	У У
	Samp	oling	
Таха	Taxonomic resolution	Sampling method	Replicates
Fish &	Species	Minnow trap	27
Macroinvertebrates	Species	Trawling	8
Plants	Species	Quadrats	15
Spiders	Morpho type	Sweeps	1
Microbes	Order	Core (surface)	4
	Order	Core (8-10 cm depth)	4
Macroinfauna	Species/Genus	Core	10

Reference marshes



PS7



Figure 4. Abundance (mean ± 1SD) among created (LHA & LHB; green symbols) and reference marshes (LHC, WPH01, WPH02, & PS07) for all



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	chang
	•• LHA •• sSBR
500 450 400	Surface Microbes
350 300 500	1 2 Below-Surface Microbes
450 400 350 300	
10.0 7.5	1 2 Plants
5.0 2.5 S	
25 20 20 15 10	Macroinfauna
Spec	1 2 4 8 Spiders
16 12 8	On-marsh nekton
4	1 2 4 8 16 Off-marsh
15 10	nekton
5	1 2 4
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estion: How does species richness ge across scales amongst groups?



Takeaways Crossed streams: Scale-dependence in all taxonomic groups



Figure 5. Sample-based rarefactions (left column) maintaining spatial configuration and (middle column) breaking spatial relationships. (right column) Individual-based rarefactions based on sampling individuals randomly.

stion: Do the effects of aggregation, ity, and SAD differ among marshes?



Conclusions

ed marshes were not notably different than ce marshes

ed marshes harbor similar levels of biodiversity as e marshes: Important tools for the maintenance toration of coastal biodiversity

al marshes are not equivalent. Different building plans may lead to different biodiversity patterns