Project Plan SP 2019-031

Do marine reserves adequately represent high diversity cryptobenthic fish assemblages in a changing climate?

Marine Science

Project Core Team

Supervising ScientistShaun WilsonData CustodianShaun Wilson

Site Custodian

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Pending project plan approval

Document endorsements and approvals as of March 18, 2020, 10:17 a.m.

Project Team required
Program Leader required
Directorate required
Biometrician required
Herbarium Curator not required
Animal Ethics Committee required



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Biodiversity and Conservation Science Program

Marine Science

Departmental Service

Service 6: Conserving Habitats, Species and Communities

Project Staff

Role	Person	Time allocation (FTE)
Supervising Scientist	Shaun Wilson	0.15
Research Scientist	Jordan Goetze	0.3
Research Scientist	Alan Kendrick	0.05

Related Science Projects

SP 2012-008 Marine Monitoring Program

Proposed period of the project

July 1, 2019 - June 30, 2022

Relevance and Outcomes

Background

Assemblages of marine fish provide key ecological functions and ecosystem services. Therefore, much effort is spent managing and monitoring fish in Western Australia's network of marine reserves. Monitoring currently focuses on large conspicuous fish, however small bodied species (<5cm) typically represent ~40% of all described species (Brandl et al. 2018). Moreover, many of these cryptobenthic reef fish (CRF) have not been described by science (Allen 2015) and their substantial contribution to current measures of fish diversity (a condition indicator for marine parks) and ecological significance are underestimated (Bellwood et al. 2019).

Short life spans of CRF suggest they are important conduits of energy transfer in marine food webs (Wilson 2004; Depczynski et al. 2007; Goatley and Bellwood 2016) and will respond more rapidly to environmental change and stressors than larger bodied counterparts (Bellwood et al. 2006). This is especially pertinent along the WA coastline, where marine heat waves have had a dramatic impact on coral (Moore et al. 2012), macroalgae (Wernberg et al. 2016) and seagrass (Arias-Ortiz et al. 2018) habitats, yet the implications for the high diversity CRF assemblages is equivocal. Furthermore, the frequency of marine heat waves is increasing (Hughes et al. 2018) and characterising the response of CRF to disturbance related changes in habitat may provide an indication of how larger bodied taxa and related ecological processes will react to climate related stressors over longer time frames (Wilson et al. In review). Similarly, as CRF are prey of many longer-lived species targeted by WA fishers (Farmer and Wilson 2011), shifts in CRF assemblages may be indicative of trophic cascades due to changes in fishing pressure. Indeed, composition of CRF assemblages may be a valuable tool for monitoring predation and how it varies with respect to spatial management of fishing pressure in marine parks (Willis and Anderson 2003).

Standard visual methods of surveying fish are not suitable for monitoring CRF (Ackerman and Bellwood 2000), and whilst sampling with poisons is the preferred technique, this is not always appropriate, requires extensive time under water and is difficult to standardise among samples. Accordingly, this project will investigate an alternate method for assessing CRF assemblages which will involve the collection and analysis of water samples for fish DNA (eDNA), a non-invasive technique with the potential to detect cryptic and rare species with little effort in the field (Pikitch 2018). (Pikitch 2018). Standard poison and visual approaches to assessing fish assemblages



will also be undertaken to compare how measures of CRF assemblage vary among techniques. By collecting data from different habitats and management zones within marine parks it is possible to assess how effectively different techniques detect change in CRF due to spatial variation in environment or management. Results from these surveys will also help identify appropriate indicator species for future monitoring.

The study will initially determine which technique(s) best assess diversity and abundance of CRF in Ningaloo Marine Park. Appropriate techniques will then be used throughout the states network of marine reserves to assess how CRF assemblages vary among habitats and management zones.

Aims

The project will determine appropriate techniques for monitoring CRF, comparing assemblages in fished and protected areas to assess how spatial variation in fishing pressure influences diversity of a KPI ecological value. Thus, the study will provide an indication of marine reserve effectiveness for conserving diversity and a key ecological process. Comparing CRF assemblages in different habitats and levels of habitat quality (e.g. % coral cover) will explore how changes in habitat due to climate related disturbances and pressures may affect fish assemblages. Findings will therefore determine the utility of CRF as indicators of environmental change and adequacy of management zones in marine reserves for capturing fish diversity and important ecological processes. This quantitative approach will provide a basis for assessing the diversity and ecological value of CRF relative to larger bodied fishes throughout the states network of marine reserves.

Expected outcome

The project will determine appropriate techniques for monitoring CRF, comparing assemblages in fished and unfished areas to assess how spatial variation in fishing pressure influences diversity of a KPI ecological value. Thus, the study will provide an indication of marine reserve effectiveness for conserving diversity and a key ecological process. Comparing CRF assemblages in different habitats and levels of habitat quality (e.g. % coral cover) will explore how changes in habitat due to climate related disturbances and pressures may affect fish assemblages. Findings will therefore determine the utility of CRF as indicators of environmental change and adequacy of management zones in marine reserves for capturing fish diversity and important ecological processes. This quantitative approach will provide a basis for assessing the diversity and ecological value of CRF relative to larger bodied fishes throughout the states network of marine reserves.

Knowledge transfer

The project will develop indicators and methodologies that can improve the monitoring of fish assemblages in the states network of marine reserves by the marine science program. DBCA researchers will learn methods for eDNA sampling, which has shown great potential as a rapid and non-destructive monitoring tool for a broad range of marine life. Improved understanding of the diversity and habitat associations of fishes will help with conservation planning. Findings from the project will be disseminated within DBCA via information sheets, meetings and presentations. As results will also be of interest to international researchers and conservationists, they will be published in peer reviewed journals, presented at conferences and circulated through social media.

Tasks and Milestones

Milestone/task	When
Collect samples with icthyodcide, UVC, eDNA	March 2020
ID and analyse samples	March-August 2020
Analyse of data, write up results of method comparison for publication	August 2020- 2021
Analyse data, write up results on effects of no-take areas	Feb 2021-July 2021
Use appropriate method to sample across different habitats	Feb 2022-Feb 2023
Analyse samples	Feb 2022-July 2023
Write up results comparing samples spatially	Jul 2023-Dec 2023



References

Ackerman JL, Bellwood DR (2000) Reef fish assemblages: a re-evaluation using enclosed rotenone stations. Mar Ecol-Prog Ser 206:227–237

Allen GR (2015) Review of Indo-Pacific coral reef fish systematics: 1980 to 2014. Ichthyol Res 62:2-8

Arias-Ortiz A, Serrano O, Masqué P, Lavery PS, Mueller U, Kendrick GA, Rozaimi M, Esteban A, Fourqurean JW, Marbà N (2018) A marine heatwave drives massive losses from the world's largest seagrass carbon stocks. Nat Clim Change 8:338

Bellwood DR, Hoey AS, Ackerman JL, Depczynski M (2006) Coral bleaching, reef fish community phase shifts and the resilience of coral reefs. Glob Change Biol 12:1587–1594

Bellwood DR, Streit RP, Brandl SJ, Tebbett SB (2019) The meaning of the term 'function' in ecology: A coral reef perspective. Funct Ecol 0: +

Berry TE, Osterrieder SK, Murray DC, Coghlan ML, Richardson AJ, Grealy AK, Stat M, Bejder L, Bunce M. (2017). DNA metabarcoding for diet analysis and biodiversity, a case study using the endangered Australian sea lion (*Neophoca cinerea*). Ecol Evol, 7, 5435-5453.

Brandl SJ, Goatley CH, Bellwood DR, Tornabene L (2018) The hidden half: ecology and evolution of cryptobenthic fishes on coral reefs. Biol Rev

Deagle BE, Tollit DJ. (2007). Quantitative analysis of prey DNA in pinniped faeces, potential to estimate diet composition? Conserv Genet, 8, 743-747.

Depczynski M, Bellwood DR (2004) Microhabitat utilisation patterns in cryptobenthic coral reef fish communities. Mar Biol 145:455–463

Depczynski M, Fulton CJ, Marnane MJ, Bellwood DR (2007) Life history patterns shape energy allocation among fishes on coral reefs. Oecologia 153:111–120

Farmer BM, Wilson SK (2011) Diet of finfish targeted by fishers in North West Australia and the implications for trophic cascades. Environ Biol Fishes 91:71–85

Fisher R, Wilson SK, Sin TM, Lee AC, Langlois TJ (2018) A simple function for full-subsets multiple regression in ecology with R. Ecol Evol 8:6104–6113

Fulton CJ, Abesamis RA, Berkström C, Depczynski M, Graham NAJ, Holmes TH, Kulbicki M, Noble MM, Radford BT, Tano S, Tinkler P, Wernberg T, Wilson SK (2019) Form and function of tropical macroalgal reefs in the Anthropocene. Funct Ecol 0:

Goatley CHR, Bellwood DR (2016) Body size and mortality rates in coral reef fishes: a three-phase relationship. Proc R Soc B 283:20161858

Graham NAJ, Jennings S, MacNeil MA, Mouillot D, Wilson SK (2015) Predicting climate-driven regime shifts versus rebound potential in coral reefs. Nature 518:94–97

Hughes TP (1994) Catastrophes, phase shifts, and large-scale degradation of a Caribbean coral reef. Science 265:1547–1551

Hughes TP, Anderson KD, Connolly SR, Heron SF, Kerry JT, Lough JM, Baird AH, Baum JK, Berumen ML, Bridge TC (2018) Spatial and temporal patterns of mass bleaching of corals in the Anthropocene. Science 359:80–83

Hughes TP, Graham NAJ, Jackson JBC, Mumby PJ, Steneck RS (2010) Rising to the challenge of sustaining coral reef resilience. Trends Ecol Evol 25:633–642

Koziol A, Stat M, Simpson T, Jarman S, DiBattista JD, Harvey ES, Marnane M, McDonald J, Bunce M (2018) Environmental DNA metabarcoding studies are critically affected by substrate selection. Mol Ecol Resour

Moore JAY, Bellchambers LM, Depczynski MR, Evans RD, Evans SN, Field SN, Friedman KJ, Gilmour JP, Holmes TH, Middlebrook R, Radford BT, Ridgway T, Shedrawi G, Taylor H, Thomson DP, Wilson SK (2012) Unprecedented Mass Bleaching and Loss of Coral across 12° of Latitude in Western Australia in 2010–11. PLoS ONE 7:e51807

Mouillot D, Graham NAJ, Villéger S, Mason NWH, Bellwood DR (2013) A functional approach reveals community responses to disturbances. Trends Ecol Evol 28:167–177

Pikitch EK (2018) A tool for finding rare marine species. Science 360:1180-1182

Rodemann JR, Brandl SJ (2017) Consumption pressure in coastal marine environments decreases with latitude and in artificial vs. natural habitats. Mar Ecol Prog Ser 574:167–179

Stat, M, Huggett MJ, Bernasconi R, DiBattista JD, Berry TE, Newman SJ, Harvey ES, Bunce, M. (2017). Ecosystem biomonitoring with eDNA, metabarcoding across the tree of life in a tropical marine environment. Scientific Reports, 7, 12240.

Warwick RM, Clarke KR (1998) Taxonomic distinctness and environmental assessment. J Appl Ecol 35:532–543



Wernberg T, Bennett S, Babcock RC, Bettignies T de, Cure K, Depczynski M, Dufois F, Fromont J, Fulton CJ, Hovey RK, Harvey ES, Holmes TH, Kendrick GA, Radford B, Santana-Garcon J, Saunders BJ, Smale DA, Thomsen MS, Tuckett CA, Tuya F, Vanderklift MA, Wilson S (2016) Climate-driven regime shift of a temperate marine ecosystem. Science 353:169–172

Willis T, Anderson M (2003) Structure of cryptic reef fish assemblages: relationships with habitat characteristics and predator density. Mar Ecol Prog Ser 257:209–221

Wilson SK (2004) Growth, mortality and turnover rates of a small detritivorous fish. Mar Ecol Prog Ser 284:253–259

Wilson SK, Fisher R, Pratchett MS (2013) Differential use of shelter holes by sympatric species of blennies (Blennidae). Mar Biol 160:2405–2411

Wilson SK, Graham NAJ, Polunin NVC (2007) Appraisal of visual assessments of habitat complexity and benthic composition on coral reefs. Mar Biol 151:1069–1076

Wilson SK, Robinson JP, Chong-Seng K, Robinson J, Graham NAJ (In review) Boom and bust of keystone structure on coral reefs. Coral Reefs

Study design

Methodology

Method/management comparisons

The composition of CRF assemblages will be assessed by; icthyocide collections, underwater visual surveys and collection of water for eDNA. CRF assemblages will be sampled at six sites inside sanctuary zones and six sites within recreational zones in the Ningaloo lagoon. Within each of the sites there will be 4-5 replicate survey areas over rubble dominated benthos. This type of benthos is common at Ningaloo, and typically has high diversity and abundance of CRF (Depczynski and Bellwood 2004).

Underwater visual census (UVC), will be undertaken using 4-5 point counts (5m radius) over the site that will also be used for ichtyocide sampling. UVC will also occur over larger 5x50 transects to assess the abundance of more mobile cryptic species and the abundance of potential predators. Following visual assessment, 4-5 small plots (~0.5m²) will be enclosed using a fine mesh and fish within the enclosed area will be collected using clove oil (Depczynski and Bellwood 2004). All CRF collected will be euthanised and preserved for identification.

To assess fish presence using environmental DNA, 4-5 replicate 1L water samples will be collected from the same depth at each site. Water samples will be filtered using 0.45tm membranes on the day of collection and filters frozen at -20°C. Water has been chosen as the medium for DNA collection as it tends to contain more fish DNA than other substrates (Koziol et al. 2018). DNA extractions, including negative field controls, will be conducted in a dedicated DNA laboratory using automated sample preparation (QIAGEN QIAcube, Venlo, Netherlands). A single step metabarcoding approach using oligonucleotides targeting fish taxa will be used to amplify fish DNA (Stat et al. 2017) and will include both negative and positive laboratory controls. Resulting samples will be pooled (equimolar) into a library and sequenced using an Illunina Miseq platform (Illumina; San Diego, USA). All data generated through sequencing will be filtered through a series of quality control steps prior to taxonomic assignment (Stat et al. 2017).

Variation in CRF across habitats

CRF assemblages will be assessed across a range of habitats, from healthy coral reefs, with high structural complexity to reefs dominated by canopy forming macroalgae and low levels of hard structural complexity. The extremes of this range represent two prominent and ecologically important habitat types in tropical shallow waters (Fulton et al. 2019), as well as alternate stable states that evolve due to differing exposure to disturbance and anthropogenic stressors (Hughes et al. 2010). In particular, coral dominated reefs are highly susceptible to heat stress and extensive bleaching of corals following heat stress, or gradual erosion of reef resilience due to fishing, disease and cyclones, can result in phase shifts to macroalgal dominated reefs (Hughes 1994; Graham et al. 2015). Understanding how CRF assemblages vary across this range of habitats will therefore give an indication of how fish assemblages may change due to climate and human related pressures. Identifying species unique to different habitat types will also be the basis of developing suitable indicators for monitoring the condition of CRF relative to environmental pressures. All samples will be collected using the technique identified from method comparisons as being most appropriate for collecting CRF. The habitat present within each site will be assessed using line intercept transects and structural complexity will be recorded using a combination of seascape and small scale measures relevant to fish of different body sizes (Wilson et al. 2007, 2013).

Statistical analyses



1. i) Method/management comparison

Data from each sample will be analysed to determine the number of species present and indices of; species, taxonomic (Warwick and Clarke 1998) and functional diversity (Mouillot et al. 2013) calculated for all replicates. How these measures of diversity differ with respect to the sampling methods (UVC, eDNA, ichthyocides), and management zones (sanctuary, general use) will be investigated using generalised additive mixed models. Method and management zones will be included in models as fixed factors and sites will be random factors. All combinations of variables will be considered in analyses and the best model will be identified using AIC and model weights (Burnham and Anderson 2010; Fisher et al. 2018). Differences in CRF assemblage composition among methods, and management zones will be examined using permutational analysis of variance.

1. ii) Variation in CRF across habitats

The influence of habitat on CRF assemblages will be assessed using a combination of GAMM (Generalised additive mixed models), DistlM (Distance based linear modelling) and PCO (Principal Coordinates Analysis). The number of individuals, species, taxonomic and functional diversity will be calculated for each replicate and used as response variables in GAMM, whilst habitat measures (e.g. coral cover, macroalgal cover, complexity) will be co-variates and site a random factor. All combinations of variables will be considered using a full subsets approach (Fisher et al 2018) and the best model for explaining variation in community measures will be based on the lowest AIC value and model weights (Burnham and Anderson 2010). Differences in CRF community composition will be examined using DistlM, with habitat measures used as predictor variables and findings viewed in multivariate space using PCO.

Biometrician's Endorsement

required

Data management

No. specimens

Herbarium Curator's Endorsement

not required

Animal Ethics Committee's Endorsement

required

Data management

The raw data will be stored on MSP T drive with a full back-up stored on an external hard drive. Additionally, a page will be created on the DBCA Confluence site that will provide links to the data sets (csv), R scripts, and output figures as appropriate. The metadata reports for each field trip will be uploaded onto Confluence under a newly created project linked to the SPP number.

Data custodian: Marine Science Program, Shaun Wilson

Budget

Consolidated Funds

Source	Year 1	Year 2	Year 3
FTE Scientist	0.5	0.5	0.5
FTE Technical	0.05	0.05	0.05



Source	Year 1	Year 2	Year 3
Equipment	12,500	12,000	
Vehicle			
Travel	3,500	3,500	5,000
Other			
Total	16,000	15,500	7,000

External Funds

Source	Year 1	Year 2	Year 3
Salaries, Wages, Overtime			
Overheads			
Equipment			
Vehicle			
Travel			
Other			
Total			