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Advancement in the SRO Breeding Program - Families and New Technologies



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Advancement in the SRO Breeding Program –

Families and New Technologies



M. Dove, P. Kube, C. Lind, V. Cumbo, W. O'Connor, M.
Saowaros, A. Elizur, R. Seeto, Z. Gibb, T. Abramov, D. Raftos, E.
Wilkie, SOCo and Southern Cross Shellfish

Future Oysters CRC-P: Accelerated SRO breeding research

- **Project collaborators:**
 - NSW DPI, CSIRO Macquarie University & SOCo
- **Objectives:**
 - More families
 - Increase QX resistance
 - Understand winter mortality
 - Markers for QX resistance (Vivian Cumbo, Macquarie University)





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Accelerated SRO Breeding Research Outcomes

- **Double family production**
 - Improvements to protocols, facilities, husbandry and training
- **One year breeding cycle for QX resistance**
 - doubles the gain
 - Required understanding of the:
 - Genetic architecture of traits
 - SRO reproduction and the QX disease cycle
 - Logistics with respect to growth and condition



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Accelerated SRO Breeding Research Outcomes

Better understanding of winter mortality disease (WM)

- More sites and more tests:
 - Georges and Shoalhaven Rivers
 - Spat and adults
- WM is enigmatic
- No correlation between WM and other traits
- WM is heritable (low to moderate) and can be improved with breeding

Future Oysters CRC-P: New technologies to improve SRO breeding and production

- Project collaborators:

- NSW DPI, University of the Sunshine Coast, University of Newcastle, SOCO, SCS

- Objectives:

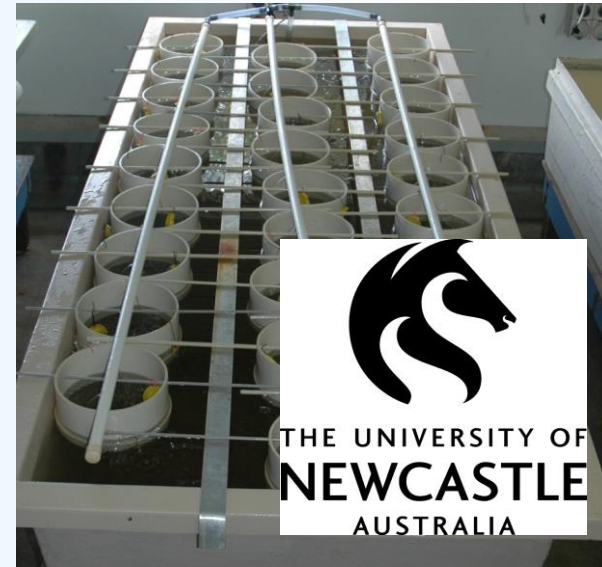
- Increase SRO breeding reliability
- Reduce conditioning time
- SRO Tetraploids



New technologies to improve SRO breeding & production

Improve SRO family success:

- Broodstock conditioning (USC)
 - maturation inducer
- Natural gamete release (USC)
 - spawn inducer
- Preserve gametes when spawning (UoN)
- Tetraploid inductions (Southern Cross Shellfish)



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Preservation of SRO gametes for short term periods.

- Provides a backup in the event of failed breeding
- Cryopreservation not always practical
- Using antioxidants to extend the life span of SRO gametes
- Fine tuning a protocol to allow benchtop storage for 5 days



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AUSTRALIA

In vivo bioassay of buccalin, APGW and tachykinin peptides in the Sydney rock oyster, *Saccostrea glomerata*: gonad conditioning

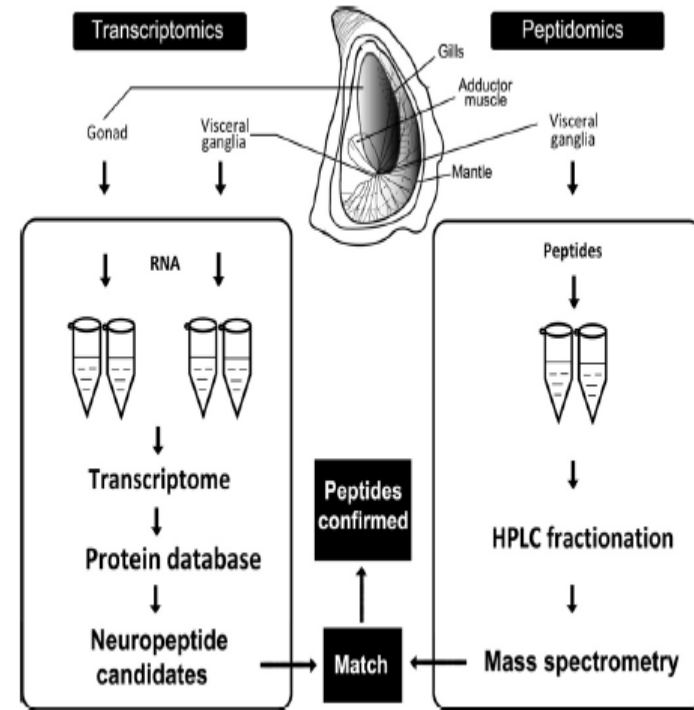
Introduction

Peptides 82 (2016) 109–119

Contents lists available at ScienceDirect

Peptides

journal homepage: www.elsevier.com/locate/peptides



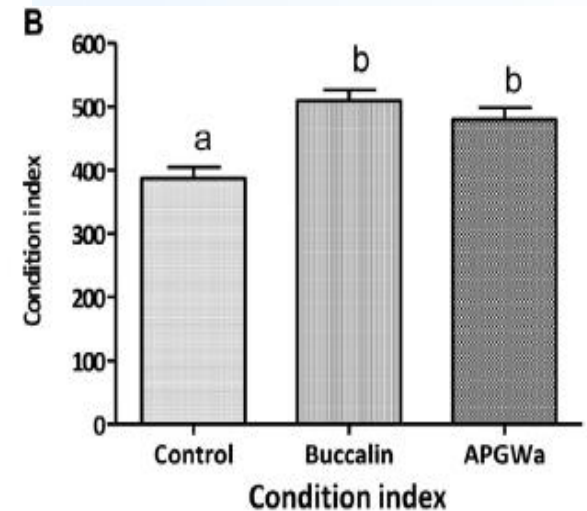
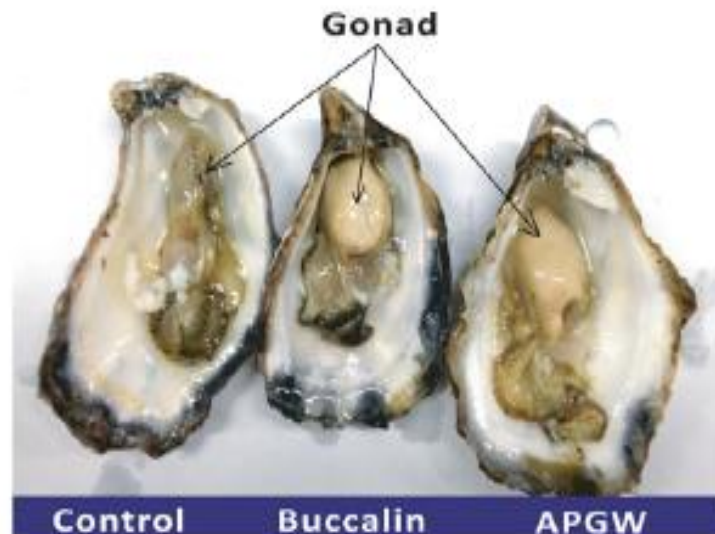
Reproductive neuropeptides that stimulate spawning in the Sydney Rock Oyster (*Saccostrea glomerata*)

Vu Van In^{a,b}, Nikoleta Ntalamagka^a, Wayne O'Connor^{a,c}, Tianfang Wang^a, Daniel Powell^a, Scott F. Cummins^a, Abigail Elizur^{a,*}

^a Centre of Genetics, Ecology and Physiology, Univ

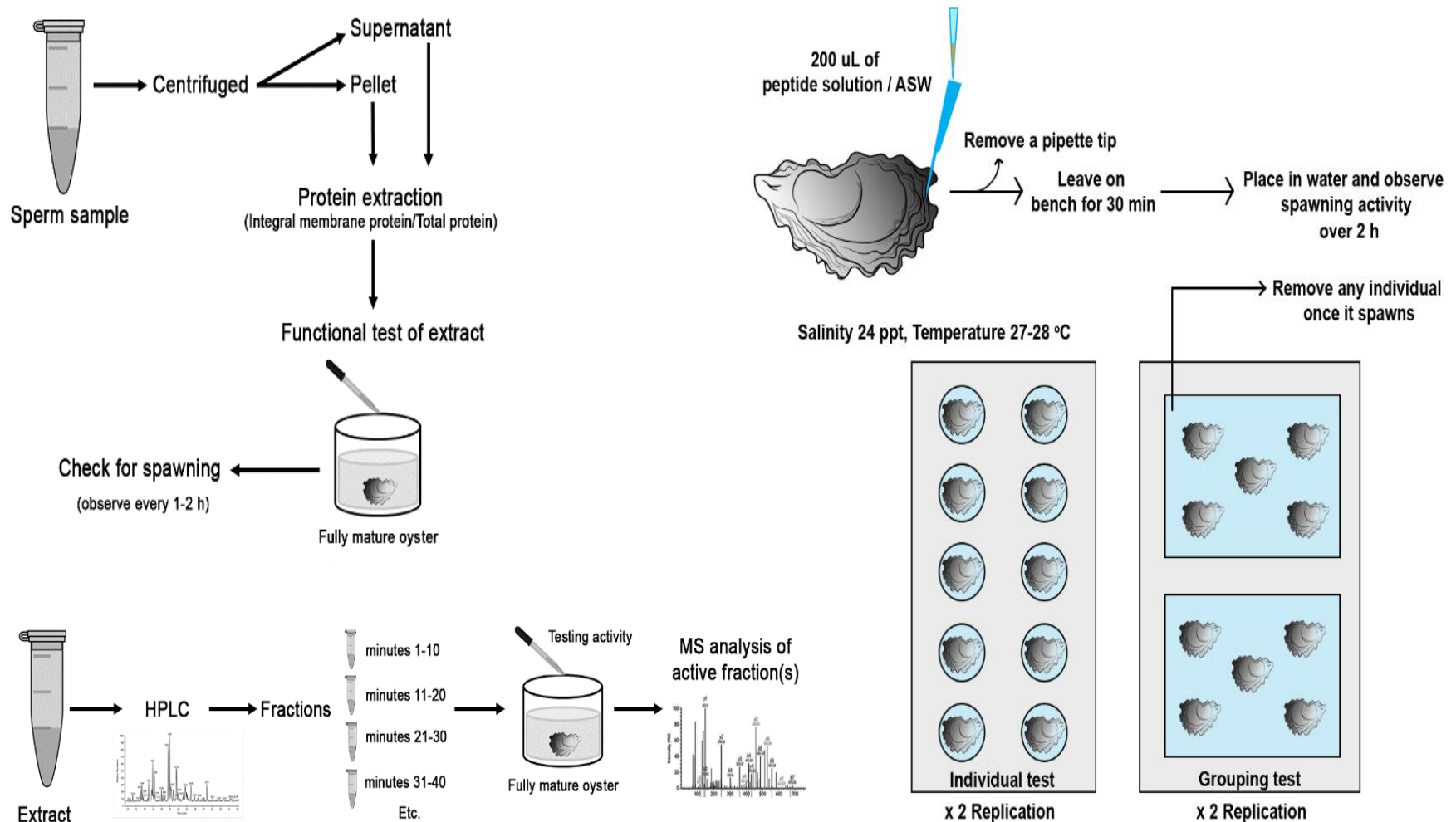
^b Northern National Broodstock Center for Maricu

^c Industry and Investment NSW, Department of Pr



Identification of spawning inducing pheromones in the sperm of the Sydney rock oyster, *Saccostrea glomerata*

Aim: To identify the spawning inducing pheromones (SIPs) in the sperm of the Sydney rock



The Future



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- SRO BP now operational
- Further development:
 - improve **family production success**
 - Incorporate:
 - **WM resistance**
 - **Marker assisted selection** for QX resistance
- Breeding program responsive to unidentified future problems and needs
- Breeding program that can ensure long term sustainability of this industry



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AIM

To develop a marker assisted selection program that can help provide commercial quantities of disease resistant oysters to the Australian oyster industry



Experiments

1. Compile a list of genetic markers that could indicate disease resistance
2. Create a method to non-lethally sample oysters
3. Experimentally test markers to determine which indicate disease resistance



Transcriptomic Differences: Gene Expression Markers

Phenotypic differences between disease resistant and susceptible oysters at the molecular level

qPCR is a fast and cost-effective molecular method for screening large numbers of animals



Marker Selection

Literature search

Genes that were differentially expressed between selectively bred and wild type oysters under ambient conditions

79 genes identified, and 66 chosen for initial analysis



Differential expression of genes encoding anti-oxidant enzymes in Sydney rock oysters, *Saccostrea glomerata* (Gould) selected for disease resistance
Timothy J. Green^{a,*}, Tom J. Dixon^{b,c}, Emilie Devic^a, Robert D. Adlard^{d,e}, Andrew C. Barnes^{a,f}

Received: 31 March 2017 | Revised: 28 July 2017 | Accepted: 7 August 2017
DOI: 10.1111/mec.14333

ORIGINAL ARTICLE

WILEY **MOLECULAR ECOLOGY**

Transcriptomic profiling of adaptive responses to ocean acidification

Priscila Goncalves^{1,2} | David B. Jones^{1,2} | Emma L. Thompson^{1,2,3} |
Laura M. Parker³ | Pauline M. Ross³ | David A. Raftos^{1,2}

MOLECULAR ECOLOGY

Molecular Ecology (2016) 25, 4836–4849

doi: 10.1111/mec.13808

Rapid transcriptional acclimation following transgenerational exposure of oysters to ocean acidification

PRISCILA GONCALVES,^{††} KELLI ANDERSON,^{††} EMMA L. THOMPSON,^{††} AROON MELWANI,^{††} LAURA M. PARKER,[‡] PAULINE M. ROSS[‡] and DAVID A. RAFTOS^{*,††}

^{*}Department of Biological Sciences, Macquarie University, Sydney, NSW 2109, Australia, [†]Sydney Institute of Marine Science, Chowder Bay, NSW 2088, Australia, [‡]School of Biological Sciences, University of Sydney, Sydney, NSW 2006, Australia

Vol. 599: 115–127, 2018
<https://doi.org/10.3354/meps12455>

MARINE ECOLOGY PROGRESS SERIES
Mar Ecol Prog Ser

Published February 23

Intraspecific differences in the transcriptional stress response of two populations of Sydney rock oyster increase with rising temperatures

Dominic McAfee^{1,2,*}, Vivian R. Cumbo¹, Melanie J. Bishop¹, David A. Raftos¹

Experiment 1: Non-lethal Sampling of Oysters

1. Low oyster mortality
2. Remain reproductively active after sampling
3. Enough tissue sampled to obtain for genetic analysis of 66 genes

Known disease resistant and disease susceptible family lines from 2015 year class

Susceptible	Survival
2015042	4.0%
2015032	5.3%
2015038	12.0%
2015020	14.6%
2015040	16.0%

Resistant	Survival
2015006	51.3%
2015007	67.0%
2015025	68.3%
2015022	71.0%
2015008	76.0%

40 oysters from each family line acclimated for 2 weeks in 750L tanks

Treatments:

- Anesthetized + tissue sampled
- Anesthetized only
- Seawater control

Oysters placed in experimental trays. Air exposed overnight



Anesthetized oysters mantle tissue biopsied and placed into RNA*later*



10 oysters per family line sampled

1hr

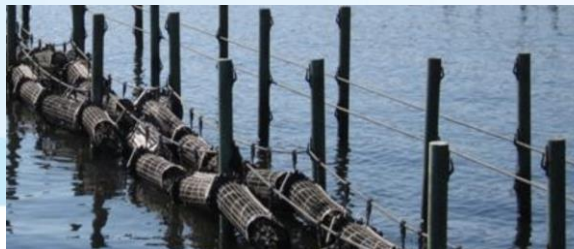
Relaxant bath of $MgCl_2$ water added 16hrs later
(50g/L; salinity 32.5, Temp 21°C)



Oysters placed back into 750L tanks for 1 week

Survival monitored @ 1 week

1 week



Oysters transferred on DPI lease

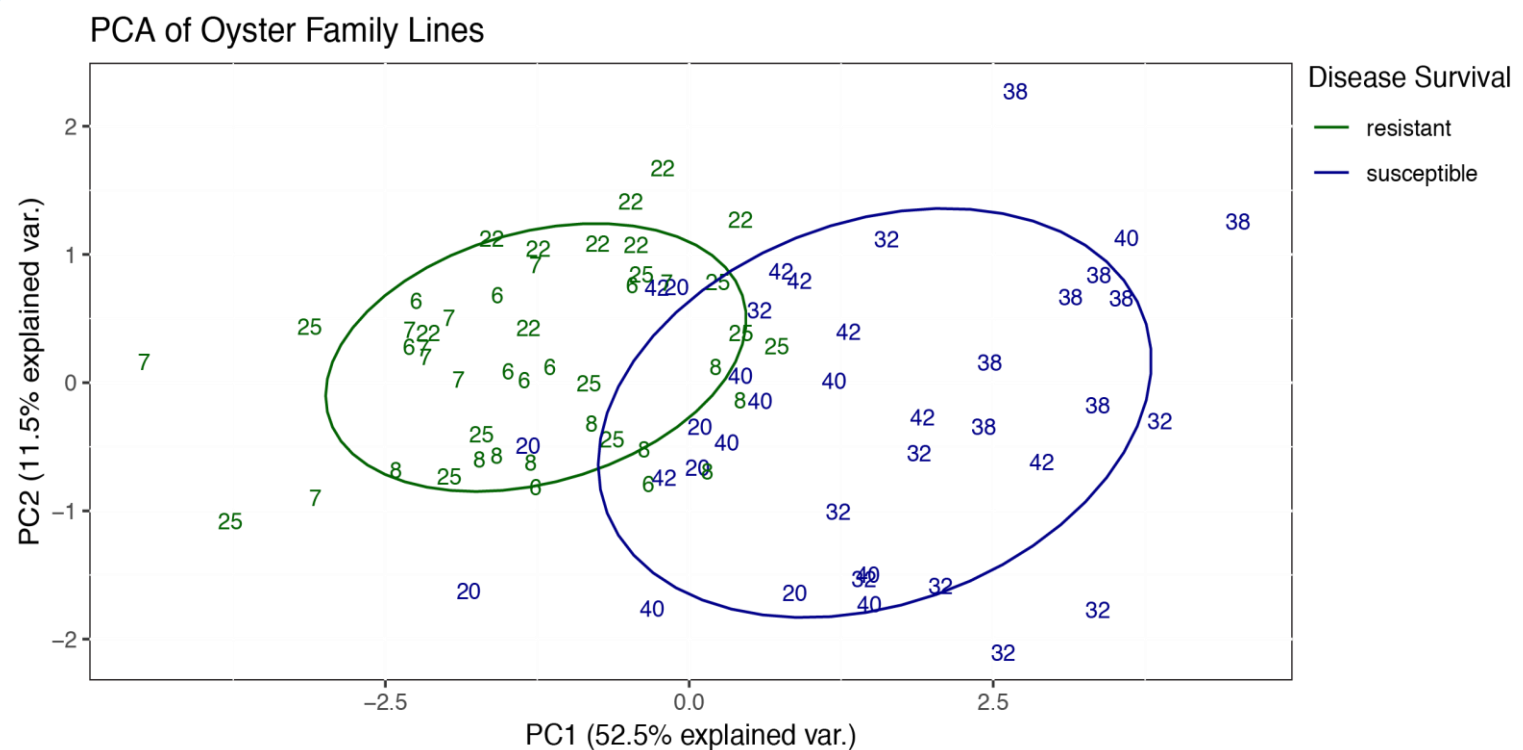
Survival monitored @ 49 days post exposure

Experiment 1: Non-lethal Sampling of Oysters

1. Low oyster mortality - No mortality detected
2. Remain reproductively active after sampling - Yes
3. Enough tissue sampled to obtain for genetic analysis of 66 genes - Yes

Experiment 2: Markers for Disease Resistance

- Molecular methods carried out
- Statistically analyzed results and found a combination of 7 gene that separate disease resistant and susceptible oysters



Conclusions

Successful development of a method to sample oyster tissue for genetic analysis that does not affect their reproduction cycle or survival

7 gene expression markers that could be incorporated into the breeding program to accelerate disease resistant selection

Developed a fast and cost-effective method for screening large numbers of oysters



Current limitations

Additional validation steps required:

- Integration of these genetic data into the EBV model
- Consider additional costs associated with screening large numbers of oysters
- Optimal time to screen animals (to get maximum discrimination between susceptible and resistant oysters and have data available for breeding decisions)
- Management of tagged broodstock prior to spawning
- Determine if we can screen sufficient numbers to account for the poor single pair mating success rate



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