# GENETIC SIGNATURES of SELECTION & ASSOCIATION ANALYSIS of the DOMESTICATION EVENT in SOUTH AFRICAN ABALONE, *HALIOTIS MIDAE*

Rhode C., Dale-Kuys R., Vervalle J., Bester- Van der Merwe A., Roodt-Wilding R. Department of Genetics, Stellenbosch University, Private Bag X1, Matieland, 7602 South Africa.

E-mail: clintr@sun.ac.za

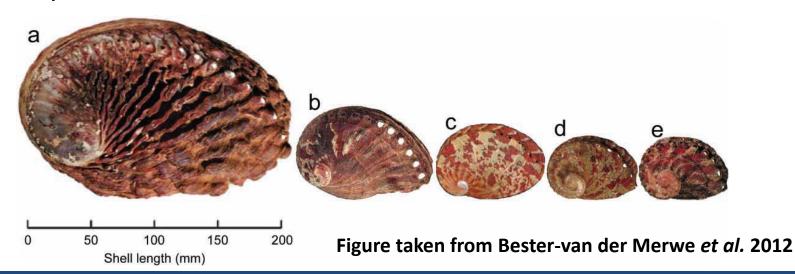






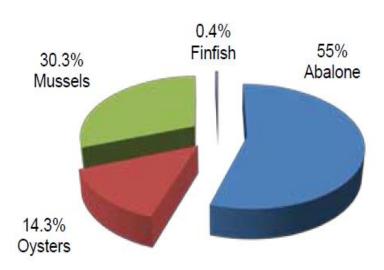
#### Introduction: Abalone

- Gastropod mollusc, genus *Haliotis*, with approximately 56 extant species, globally
- Five endemic species in South Africa
  - a) H. midae; b) H. spadicea; c) H. alfredensis; d) H. parva; e)
     H. queketti

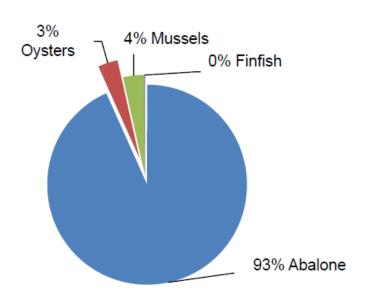


#### Introduction: Abalone

- Haliotis midae, i.e. Perlemoen
  - Local species of commercial importance
  - Fisheries and aquaculture



% Tonnage of total aquaculture production (Figure taken from DAFF, 2012)



% Revenue of total aquaculture production (Figure taken from DAFF, 2012)

## Introduction: Domestication as an Evolutionary Process

#### Selection

- Relaxed natural selection for survival in the wild
- Increased natural selection for adaptation to "humanised environment"
- Increased artificial selection
- Demography
  - Founder effect, reduction in effective population size
    - Greater influence of random drift
    - Increased inbreeding
    - Unequal parental contribution



## Introduction: General Observation for South African Abalone

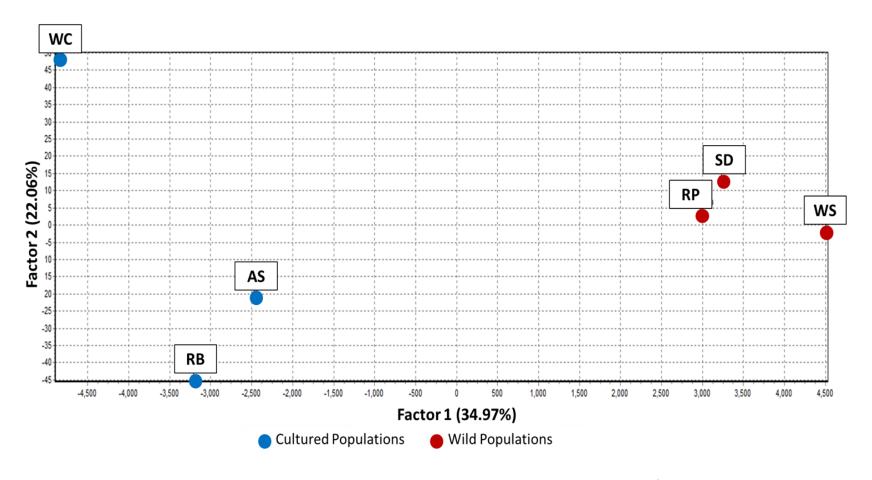
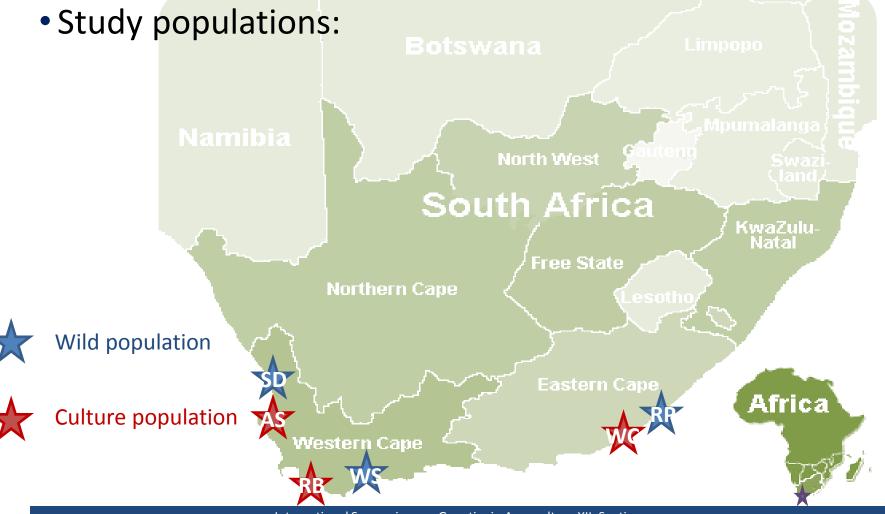


Figure adapted from Rhode et al., 2012

#### Research Question

How is selection shaping the observed patterns of genetic variation in cultured populations of abalone?





Molecular markers

- 150 microsatellite markers
  - Genome-wide distribution
  - On average 10cM interval

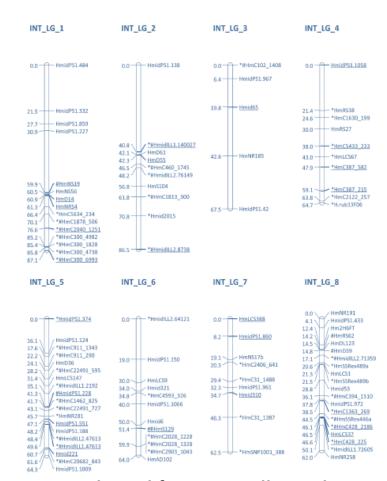
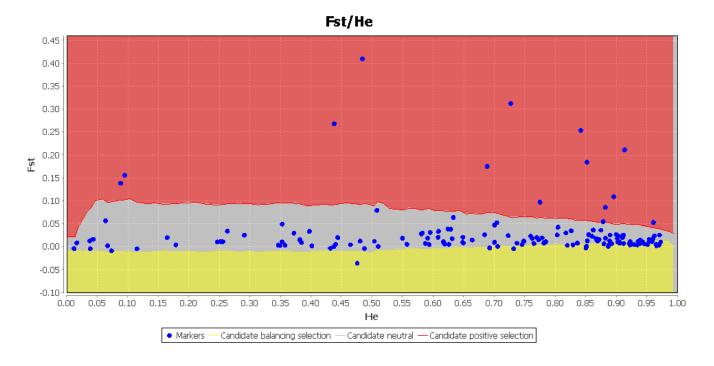


Figure adapted from Vervalle et al., 2012

 Frequentist (LOSITAN) and Bayesian (BayeScan) F<sub>st</sub>outlier analysis

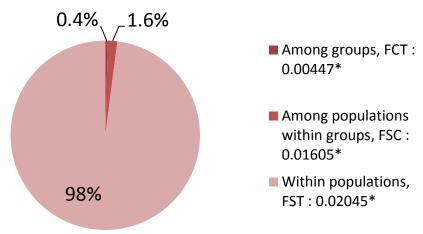


#### % Loci under selection

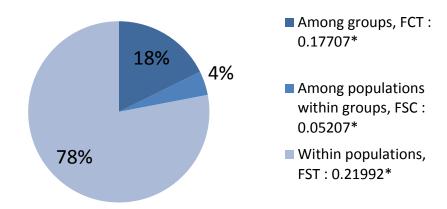
	Across cultured populations	Across wild populations	Across all populations
Directional selection	2-4	3-13	~9
Balancing selection	4-12	2-3	6-18

Variance explained due to directional selection





### AMOVA based on loci under directional selection (\*P < 0.05)



Growth rate currently selected trait

 Artificial phenotypic selection causative to molecular signature of selection?

Correlation of genetic variants with growth?



Commercial cohort of 661 F1 animals phenotyped

- Top and bottom ±15% selected for genotyping
  - 13 loci under directional selection

 Parentage analysis to correct for family size bias (final cohort: 80)

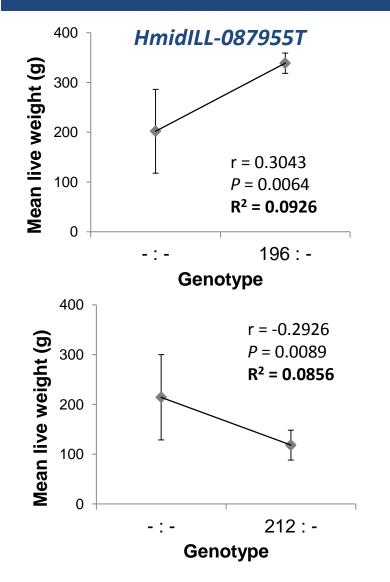
- Association tests:
  - Case/Control analysis
  - Quantitative analysis

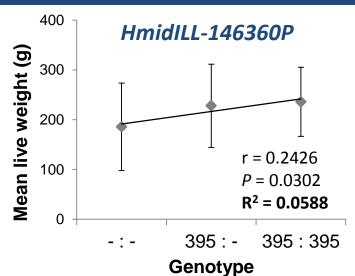
- Simple correlation analysis
  - Allelic correlations with size

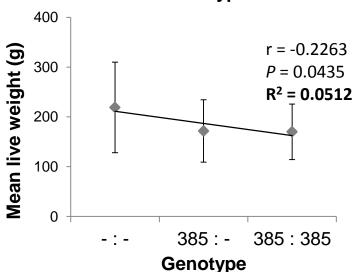
BLAST – putative functional analysis

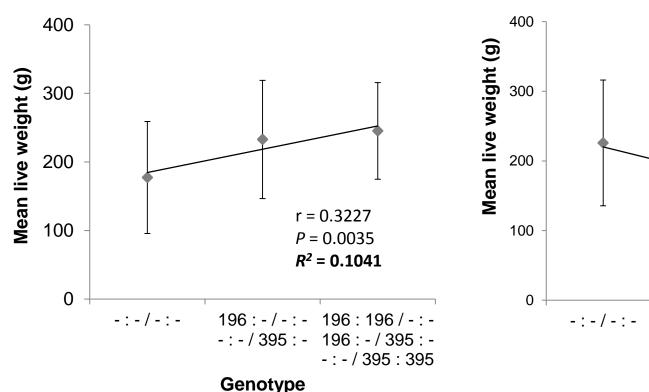
**Locus-by-locus AMOVA: Large vs Small** 

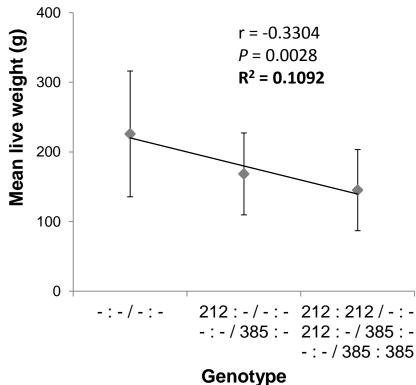
Marker	SSDa	Df	Variance	% Variation	F <sub>st</sub>	<b>G</b> " <sub>st</sub>
HmidILL-087955T	0.3091	1	0.0004	0.1179	0.0064	0.0051
HmLCS5M	0.1293	1	0.0000	0.0000	0.0024	-0.0330
HmLCS48M	0.1453	1	0.0000	0.0000	0.0036	-0.0200
Hm-NS18M	0.5574	1	0.0055	1.4459	0.0117*	0.0355*
HmidILL-146360P	1.4740	1	0.0133	3.2478	0.0232*	0.1569*
HmidILL-064192P	0.0264	1	0.0000	0.0000	0.0005	-0.0313
HmNR106D	0.3132	1	0.0000	0.0000	0.0026	-0.0577
HmidPS1.559	0.2919	1	0.0020	1.8547	0.0238*	0.0395*











HmidILL-087955T & HmidILL-146360P

#### HmidILL-087955T :

- 3' UTR of retinol dehydrogenase 14a gene
- Activation of retinol (vitamin A) to retinoic acid
- Associated with development

#### HmidILL-146360P :

- 3' UTR of malectin gene
- N-linked glycosylation / carbohydrate binding
- Carbohydrate metabolism / transmembrane / signalling

#### Conclusions

- Rapid divergence between wild progenitor and cultured populations
  - Role of selection
  - Artificial vs Natural selection

- Application in marker assisted breeding
  - Confirm in independent and larger cohorts
  - Larger genome project in future

### Acknowledgements



















Aquaculture 356-357 (2012) 235-242 Marine Genomics 10 (2013) 27-36



Contents lists available at SciVerse ScienceDirect

#### Aquaculture

journal homepage: www.elsevier.com/locate/aqua-online





Contents lists available at SciVerse ScienceDirect

#### Marine Genomics

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



A population genetic analysis of abalone domestication events in South Africa: Implications for the management of the abalone resource

Clint Rhode \*, Juli-ann Hepple, Suzaan Jansen, Tanja Davis, Jessica Vervalle, Aletta Elizabeth Bester-van der Merwe, Rouvay Roodt-Wilding

Department of Genetics, Stellenbosch University, Private bag X1, Matieland, 7602, South Africa

Detection of molecular signatures of selection at microsatellite loci in the South African abalone (*Haliotis midae*) using a population genomic approach

Clint Rhode \*, Jessica Vervalle, Aletta E. Bester-van der Merwe, Rouvay Roodt-Wilding

Department of Genetics, Stellenbosch University, Private Bag X1, Matieland, 7602, South Africa



Immunogenetics, Molecular Genetics and Functional Genomics



SHORT COMMUNICATION

doi: 10.1111/age.12142

Comparison of population genetic estimates amongst wild, F1 and F2 cultured abalone (*Haliotis midae*)

Clint Rhode, Simo N. Maduna, Rouvay Roodt-Wilding and Aletta E. Bester-van der Merwe Department of Genetics, Stellenbosch University, Private Bag X1, Matieland, 7602 South Africa.

Journal of Shellfish Research, Vol. 32, No. 1, 89-103, 2013.

#### INTEGRATED LINKAGE MAP OF HALIOTIS MIDAE LINNAEUS BASED ON MICROSATELLITE AND SNP MARKERS

JESSICA VERVALLE, JULI-ANN HEPPLE, SUZAAN JANSEN, JANA DU PLESSIS, PEIZHENG WANG, CLINT RHODE AND ROUVAY ROODT-WILDING\*

Molecular Aquatic Research Group, Department of Genetics, Stellenbosch University, Private Bag XI, Matieland, 7602, South Africa

#### Thank You!

#### **Questions?**