

Genética Spring

GENÉTICA SPRING'S (GS) BREEDING PROGRAM



Benchmark Holdings plc



Benchmark's aquaculture business

INTRODUCTION





SUMMARY

- **2**000:
 - Benchmark Holdings plc
- **2**004:
 - Fish Vet Group
- **2**014:
 - SalmoBreed
 - Stofnfiskur
- **2**015:
 - Akvaforsk Genetics
 - Spring Genetics
- **2**015:
 - INVE
- **2**016:
 - Genética Spring CENIACUA





BENCHMARK AQUACULTURE **BUSINESS Operations** in 27 countries R&D facilities and farms Diagnostic laboratories Commercial services Manufacturing/production



BENCHMARK'S AQUACULTURE PLATFORM



Shaping Sustainable Aquaculture



The Fish Site

SERVICES INCL.

R&D & trials facilities



BENCHMARK GENETICS





- CENIACUA
 - P. vannamei breeding program
 - Established in 1997
 - For the Colombia farming industry
- BENCHMARK
 - Acquired the shrimp breeding programs from CENIACUA in 2016
 - Genética Spring is formed
 - Developing breeding lines for the global shrimp industry









P. vannamei diseases today!

WSSV

The most significant viral pathogen

AHPND ("EMS")

The most significant bacterial disease

EHP

"New kid on the block"



SOME COSTS OF SHRIMP DISEASES ACROSS ASIA



WSSV, AHPND ("EMS") AND EHP

LOST OF PRODUCTIO ALONE*

- Thailand
 - - US\$ 7 billion from 2011 to 2016
 - EHP
 - US\$180 million in 2016
- Vietnam 2015
 - **WSSV**
 - US\$ 11 million
 - - US\$ 26 million
- * Feed sales losses are estimated to be 30% of production losses!





AHPND IN THAILAND P. vannamei



In the absence of AHPND and assuming that the peak production of 2011 would have been maintained, then the AHPND outbreak has cost Thailand an estimated US\$7 billon to date.



Genética Spring

GENÉTICA SPRING'S (GS) BREEDING PROGRAM

A TALE OF TWO BREEDING LINES



A TALE OF TWO BREEDING LINES

Atlantic breeding program

- 1997: Between and within-family selection
- Selected for:
 - Resistance to TSV, NHP and Vibriosis
 - General pond survival and growth
- Base Line

Pacific breeding program

- 2008: Mass selection
- Selected for:
 - WSSV resistance and general pond survival
- WSSV R Line





Atlantic line – Base line.

Complete pedigree information

- Origin from eight (8) different countries, both wild & farmed
- Actual inbreeding around 9-10%
- Between and within family selection
- 100 families per batch (full and half sib scheme)
- Two batches for year
- Current preparing the 14th generation



BASE LINE DESIGN

BREEDING NUCLEUS





PACIFIC LINE – WSSV R LINE

Mass selection

- Origin from:
 - Base Line + four (4) lines from Ecuador + one (1) wild population
- High selection pressure
 - <0.001% (1 in 10,000)</pre>
- One batch per year
- Selected for WSSV resistance and pond survival
- Current preparing the 9th generation



WSSV R LINE DESIGN

F₀ Generation 2008

- Challenge test PL40
- Broodstock raising
- Reproduction

F₈Generation 2016

- Challenge test PL40
- Challenge test lab
- Farm stocking
- Broodstock raising
- Reproduction

F₁ Generation 2009

- Challenge test PL40
- Challenge test lab
- Farm stocking
- Broodstock raising
- Reproduction



SHRIMP FIT FOR THEIR ENVIRONMENT

- The Shrimp sector will continue to be repeatedly exposed to new epidemics while depending on broodstock poorly adapted to local conditions.
- To break this vicious cycle:

SAFE

Use of clean animals ROBUST

Use of SPR robust animals

ADAPTED

Select breeders adapted to local conditions



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| Pathogen/test | PCR | RT-PCR | Histopathology |
|---|-----|--------|----------------|
| White Spot Syndrome Virus (WSSV) | ✓ | | ✓ |
| Taura Syndrome Virus (TSV) | | × | ✓ |
| Yellow-Head Virus (YHV) | | ~ | ~ |
| Infectiuos Myonecrosis Virus (IMNV) | | ~ | ~ |
| Nodavirus de P vannamei | | ~ | ~ |
| IHHNV | | | ~ |
| AHPND | ~ | | ✓ |
| Necrotizing Hepatopancreatitis (NHP) | ~ | | ✓ |
| BP – MBV - HPV | | | ✓ |
| EHP | ~ | | ✓ |
| Gregarines – Nematodes - Haplosporidians | | | ✓ |
| Bacterial infections | ~ | | ✓ |



SURVEILLANCE ANALYSIS (# of samples tested)





ARIZONA. TUCSON ARIZONA AQUACULTURE PATHOLOGY LABORATORY School of Animal & Comparative Biomedical Sciences Bldg. #90, Room 102 1117 E. Lowell Street, Tucson, Arizona 85721-0090 Phone: 520-621-4438; Email: aquapath@ag.arizona.edu



April 6, 2017

Genetica Spring SAS Andres Suarez Calle 32 # 8-33 Oficina 215 Cartagena Bolivar, Colombia

E-mail: asuarez@ceniacua.org

Case: 17-163

Dear Mr. Suarez:

The PCR and RT-PCR tests you requested for the detection of WSSV, IHHNV, TSV, YHV, IMNV, AHPND, NHP-B and EHP have been completed. The 57 vials of *P. vannamei* arrived on March 28th, 2017 in good condition. The samples were collected from Punta Canoas Breeding Facility in 2015, 2016, and 2017. Representative samples from the vials (approx. 30mg) were collected for DNA and RNA extractions. WSSV, IHHNV, TSV, YHV, IMNV, AHPND, NHP-B and EHP were not detected in the samples tested. The testing was completed on April 5th, 2017. A summary of the tests and results is provided on the following page.

We hope that this information will be helpful to you. A hard copy of this report will be mailed to you. If there are any questions regarding this case, please feel free to contact us.

UAZ Policy on certification: This report provides our findings on the samples submitted to our laboratory for examination, health status evaluation, disease diagnosis, or pathogen detection. It is our policy and intent to perform the most appropriate assay(s) for the determination of the health/pathogen status of all specimens submitted to our laboratory. However, this report in no way constitutes a stock or facility "certification" or a "certificate" of health/pathogen status for the sample(s) tested or for the stocks, or facility, from which the sample(s) were derived.

PCR: disclaimer: This report provides our findings on the samples submitted to our laboratory for pathogen detection. The PCR assay used by this laboratory for the detection of shrimp pathogens is a research tool. The results should be considered as experimental and tentative. Whenever possible, PCR results should be confirmed by alternative assay. This report in no way constitutes a stock or facility "certification" or a "certificate" of health/pathogen status for the sample(s) tested or for the stocks, or facility, from which the sample(s) were derived. The **UAZ Aquaculture Pathology Lab** is a OIE (Office International des Epizooties or the Organization or World Animal Health Organization) Reference Laboratory for White Spot Disease, Taura Syndrome, Infectious Hypodermal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, International Syndrome, Infectious Mypodermal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Tetrahedral Baculovirus, Interious Syndrome, Infectious Hypodermal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Tetrahedral Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Tetrahedral Baculovirus, Tetrahedral Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Tetrahedral Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Baculovirus, Tetrahedral Baculovirus, Infectious Hypodernal and Hematopoietic Necrosis, Spherical Bacu

Sincerely yours,

Irina Maskaykina Research Specialist

Assun Mumae Ita

Arun K. Dhar, Ph.D. Associate Professor Aquaculture Pathology Laboratory Director





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GENÉTICA SPRING POPULATIONS ARE SPR FOR:





Genética Spring





ADAPTED

- Distinct lines:
 - Selected for superior performance under specific conditions.
- Our dissemination protocols:
 - Select lines best suited to our clients conditions.
 - Local adaptation in the multiplication phase.





DIVERSE GENETIC BASE

"GENE SUPERMARKET"

Countries – Original Gene Pool: Colombia¹ Costa Rica² Ecuador² Hawaii¹ Panama² Peru² El Salvador² ✓Venezuela¹

¹Domesticate stocks; ²Wild stocks





GENOTYPE BY ENVIRONMENT INTEREACTION

GxE interactions are relevant in *P.vannamei* culture

DOES ONE SIZE FIT ALL?





LINES UNDER DEVELOPMENT

- Resistant Lines:
 - WSSV Genomic Selection
 - AHPND Genomic Selection
 - General survival
- Maturation:
 - No ablation
 - Dry feed 100%





LINES UNDER DEVELOPMENT

Growth Lines:

- Early growth (60 DOC)
- Late growth (100 DOC)
- Lined outdoor ponds (100 PL/m2)

Low salinity (<5ppt)</p>





GENÉTICA SPRING FACILITIES

Punta Canoas: SPF & SPR

- NBC
 - Family selection
- MC
 - Commercial breeders

Pacific Coast: SPR

- NBC
- Testing facility

Bogotá:

Single pathogen challenge testing





GENÉTICA SPRING INFRASTRUCTURE

Punta Canoas – Biosecure Facility

- Physically isolated from commercial farms
- Indoor NBC
- Covered MC
- Fenced perimeter
- No visitors
- Testing infrastructure
 - Earth ponds
 - Lined intensive ponds
 - Indoor/outdoor tanks
- Lab infrastructure
 - PCR; Histology; Bacteriology













BASE LINE x WSSV R LINE



LINE AND FAMILY CONCEPT

Line (or strain):

- Group of families with one or more common characteristic such as:
 - WSSV resistant
 - Fast growing
- Family:
 - Shrimps originated from one individual spawn
 - One male crossed with one female





MATINGS DESIGN



40 families total

- 10 families from each line
 - Two pure lines
 - Two hybrids lines

- Full sib and half sib families.
 - The same male inseminated one female from each line.



RESULTS

FAMILY SURVIVAL FOR A 30 DAYS WSSV CHALLENGE TEST MAY 2017



Families



FAMILY SURVIVAL FOR A 30 DAYS WSSV CHALLENGE TEST



Families



Families

FAMILY GROWTH AT 100 DOC



AVERAGE WITHIN EACH LINE FOR: SURVIVAL FOR WSSV CHALLENGE TEST GROWTH FOR 100 DOC (PL-10)





CUTTING EDGE TECHNOLOGIES

Pit Tags:

- Individual monitoring of breeders candidates:
 - From growth selection to maturation
- Parental assignment:
 - 192 SNPs panel
 - Replacing elastomer tagging
 - Common environment
- Genomic Selection
 - Training data: Trial Base line x WSSV R line
 - 2378 SNPs: different in the two lines



Technologies

from Salmon



Other major farmed species



Acknowledgments

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- Dr. Andy Shinn FVG
 For the scary disease data
- Morten Rye AFGC
 - For the genomic data





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Portal Oficial de Turismo do Rio Grande do Norte