

## Bletecsin

# Development of a Nonlinear K-Law Spectral Signature Index to Classify Basophilic Inclusion Bodies of the White Spot Syndrome Virus

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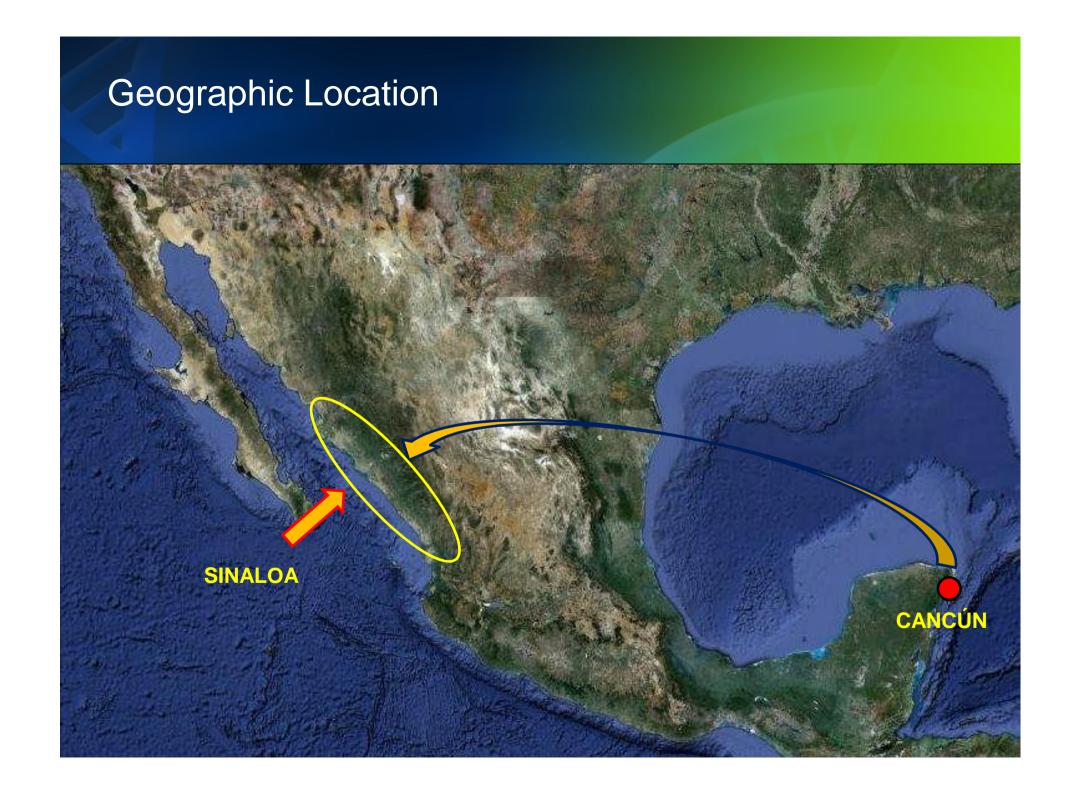
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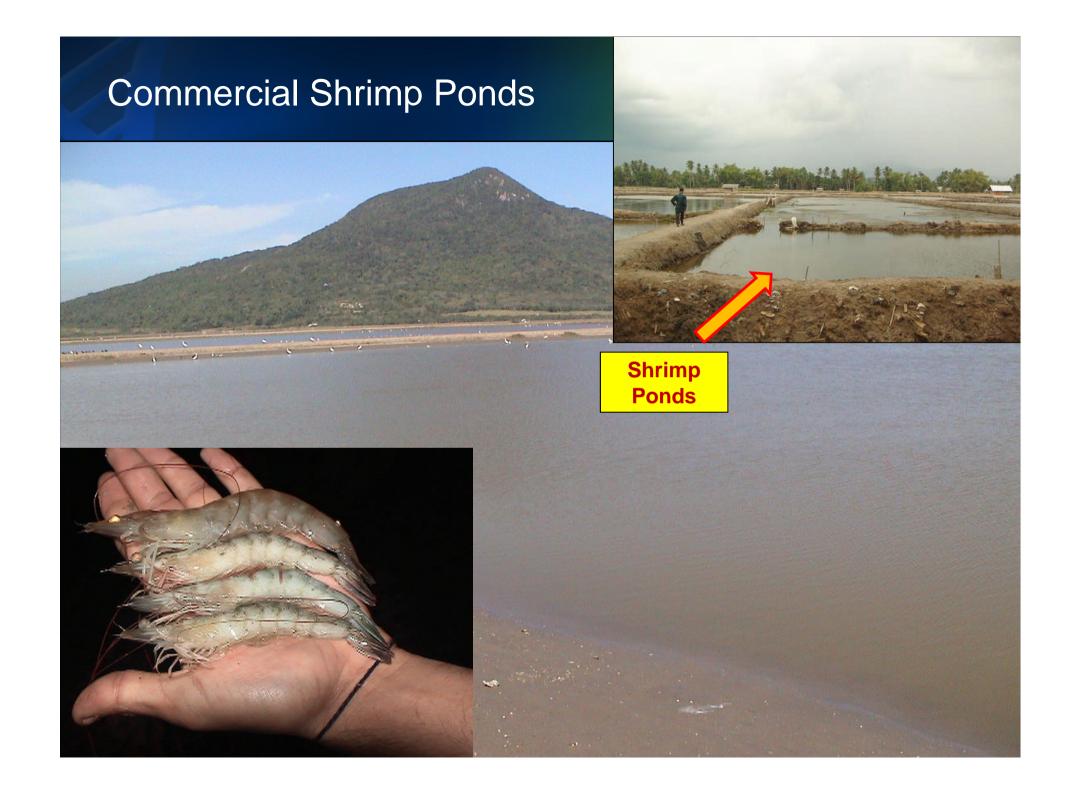
# Shrimp Producers



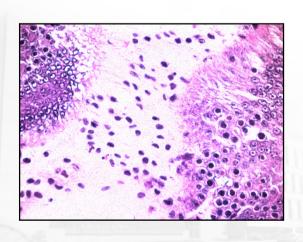


#### **Shrimp Producers Location**

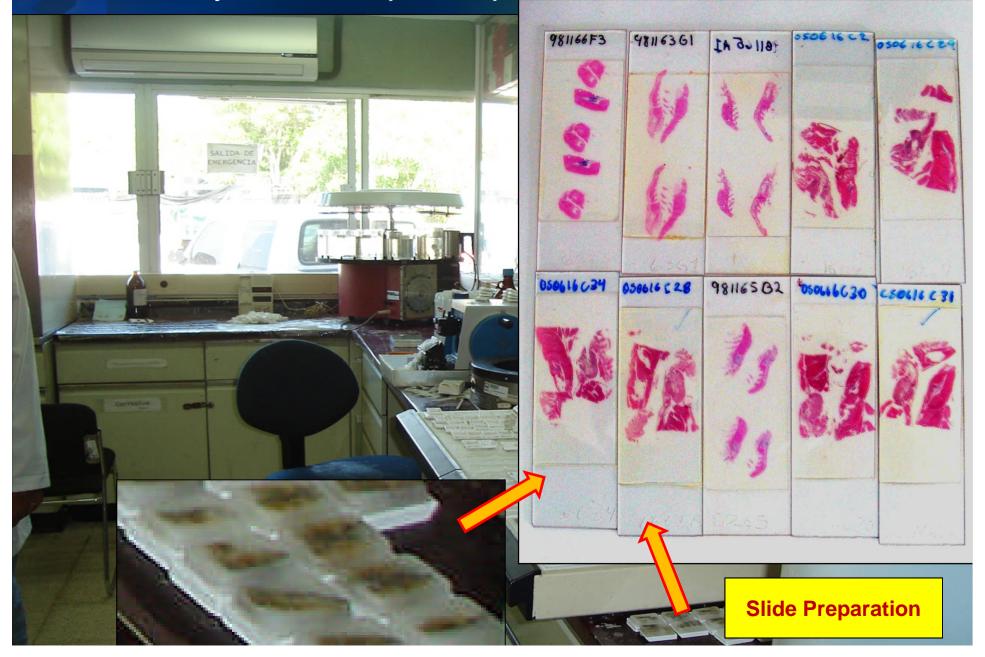




# Histology



#### Laboratory for Shrimp Samples



# Shrimp Samples Ready to be Analyzed f (x,y) is obtained

# Mathematical Foundation

$$\mathscr{F}[f(t)] = F(\omega) = \int_{-\infty}^{\infty} f(t)e^{-j\omega t}dt$$

# Non Linear K-Law Filter Mathematical Foundation

Fourier Based Filter General Form:

$$I(u,v) = |I(u,v)|^{k} \exp[-i\phi(u,v)]$$

K = 1:

$$I(u,v) = |I(u,v)|^{k=1} \exp\left[-i\phi(u,v)\right] = \Im\left(I(x,y)\right)$$

K = 0:

$$I(u,v) = \exp[-i\phi(u,v)] \rightarrow Phase \ only \ filter$$

0 < K < 1

$$I(u,v)_{k-Law} = |I(u,v)|^{0 < k < 1} \exp\left[-i\phi(u,v)\right]$$

#### Spectral Signature (SSF)

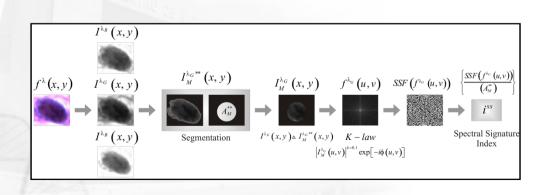
#### **Spectral Signature**

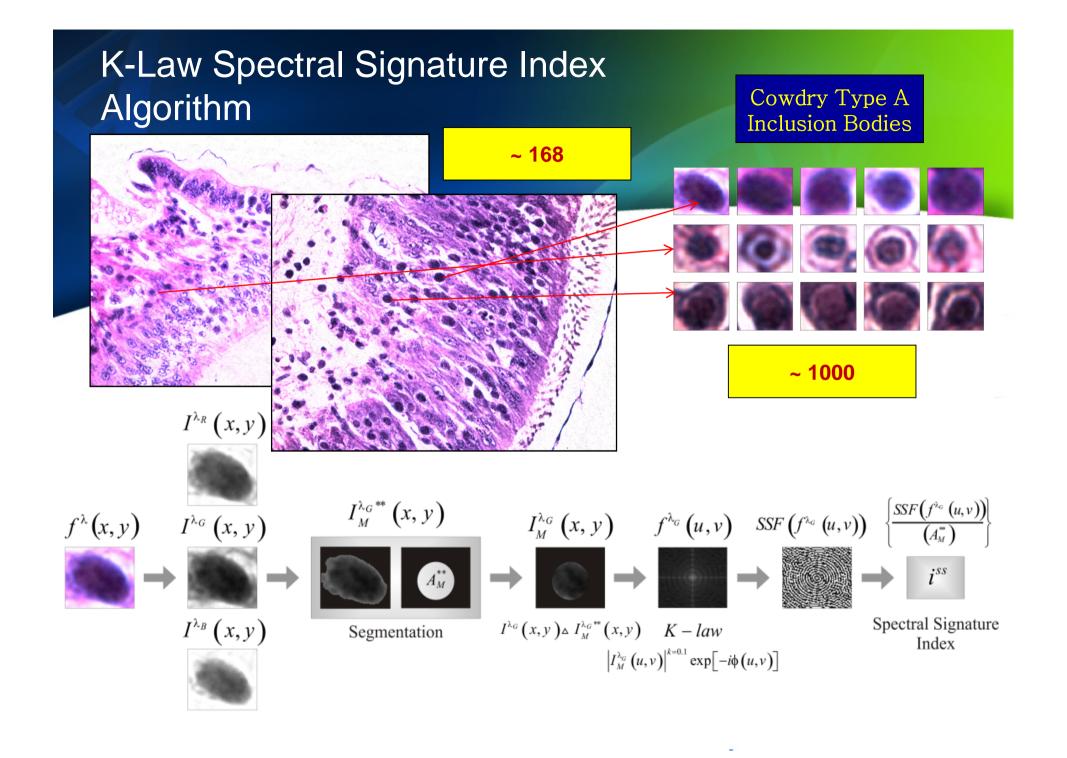
$$SSF\left(f_{w}^{\lambda_{G}}\left(u,v\right)_{k}\right) = \begin{cases} 1, & if \ \operatorname{Re}\left(f_{w}^{\lambda_{G}}\left(u,v\right)_{k}\right) > 0\\ 0, & otherwise \end{cases}$$

Spectral Signature Index

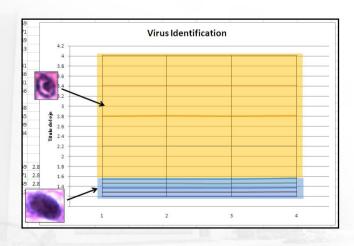
$$i_{k}^{ss} = \left\{ \frac{SSF\left(f_{w}^{\lambda_{G}}\left(u,v\right)_{k}\right)}{\left(A_{M}^{**}\right)_{w}} \middle| \left(u,v\right) \in \square \right\}$$

# Spectral Signature Algorithm



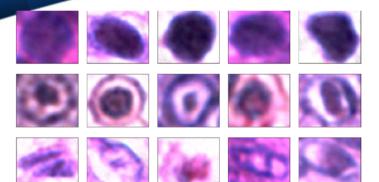


# Results



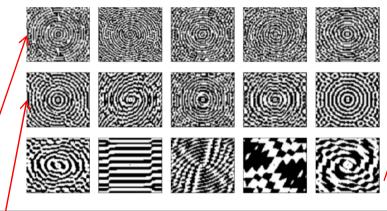
#### Classification Results

Cowdry Type A Inclusion Bodies Bank



100
Representative
CAs

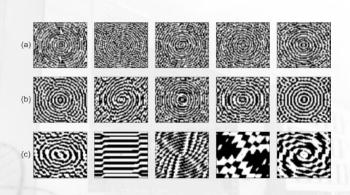
## Spectral Signature



WSSV Group	Signature Index Statistical Behavior				
	$\frac{-}{x_{i^{ss}}}$	$\sigma_{_{i^{zz}}}$	1SE	2.SE	
/ I	1.3748	0.4817	0.0852	0.1703	
II	2.6069	1.8533	0.4953	0.9906	
IIIª	159.4229	352.5394	94.2201	188.4402	
ΙV <sup>b</sup>	1.7498	1.2362	0.1823	0.3645	

- a. Non-infected tissue group particles;
- b. Groups I and II analyzed together

# Conclusion



#### Conclusion

A new algorithm based on the analysis of frequencies found in the green channel and processed by K-Law Fourier filter was developed with an efficient performance (< 0.5s by CAs).

Results show that inclusion bodies are well defined in a numerical fringe;

 $(1.3853 \le i^{ss} \le 2.1143)$ 

This algorithm is easy to be implemented inside an automatic system to find if is present the WSSV.

The SSI can be extrapolated to work on a new kind of histology tissues.

# Thank you !!!

