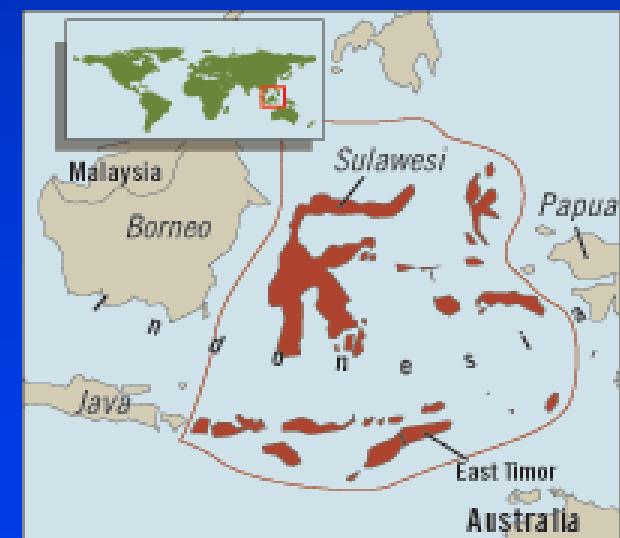


# Putting process on the map: why ecological gradients are important for preserving biodiversity

Tom Smith, Sassan Saatchi, Bob Wayne, Chris Schneider, Catherine Graham, Borja Mila, Wolfgang Buermann and John Pollinger



# Biodiversity Hotspots



# Two limitations of hotspots

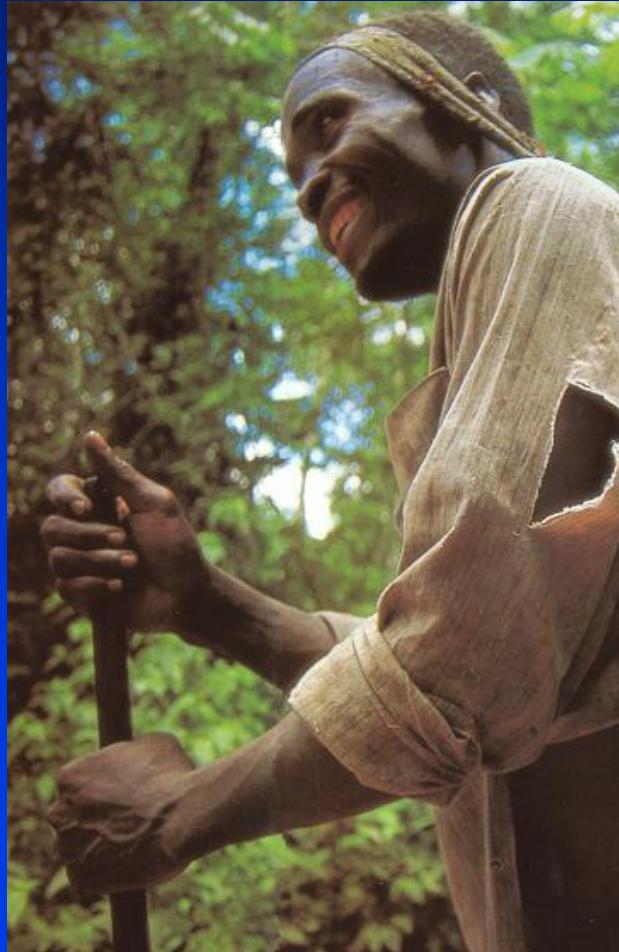
- 1) may fail to capture regions important in generating or maintaining adaptive variation
- 2) may fail to take into account climatic change

# Why preserve gradients?

1) Natural Selection along gradients is a powerful driver of adaptive variation

2) Given climate change preserving them may offer a bet hedging approach

QuickTime™ and a  
TIFF (Uncompressed) decompressor  
are needed to see this picture.

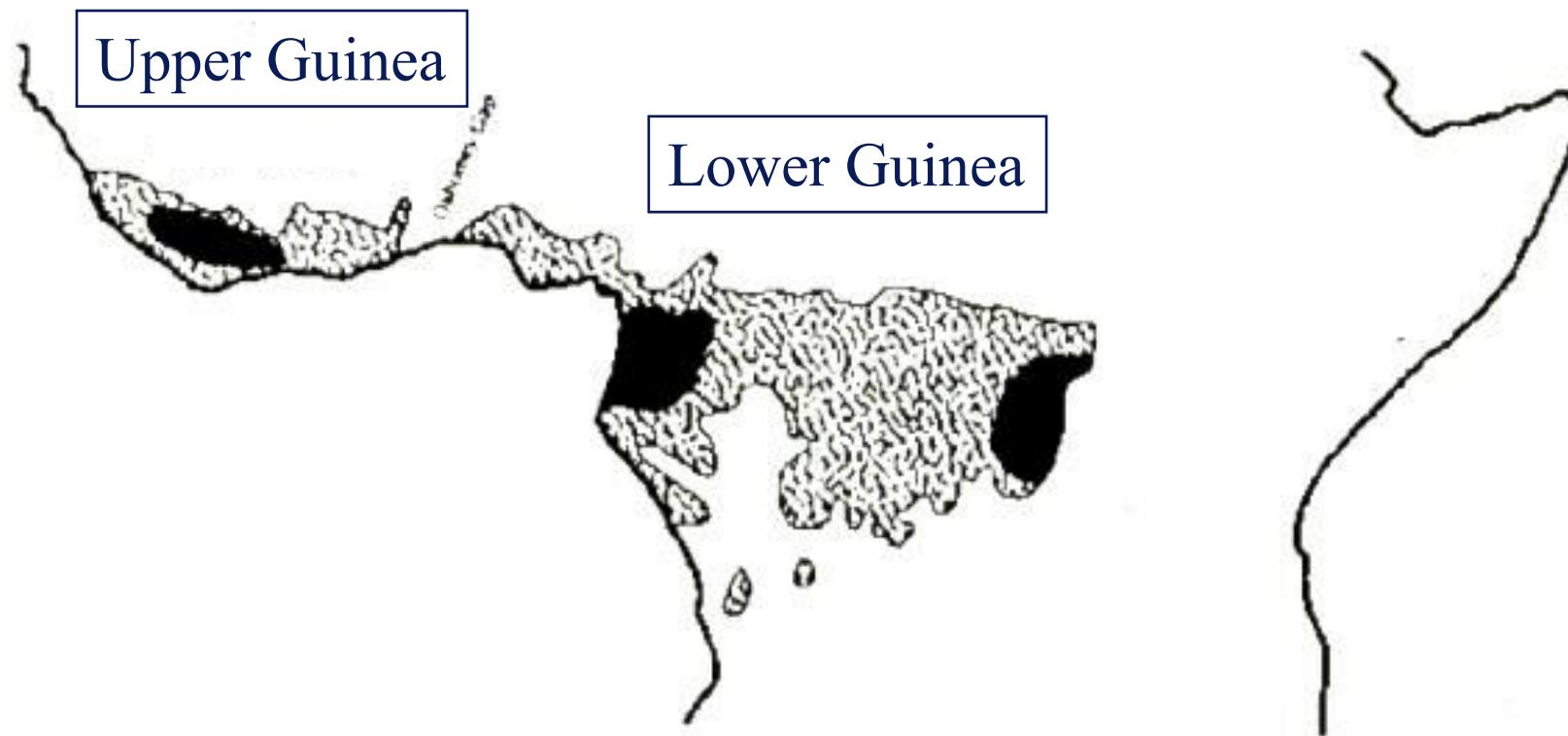


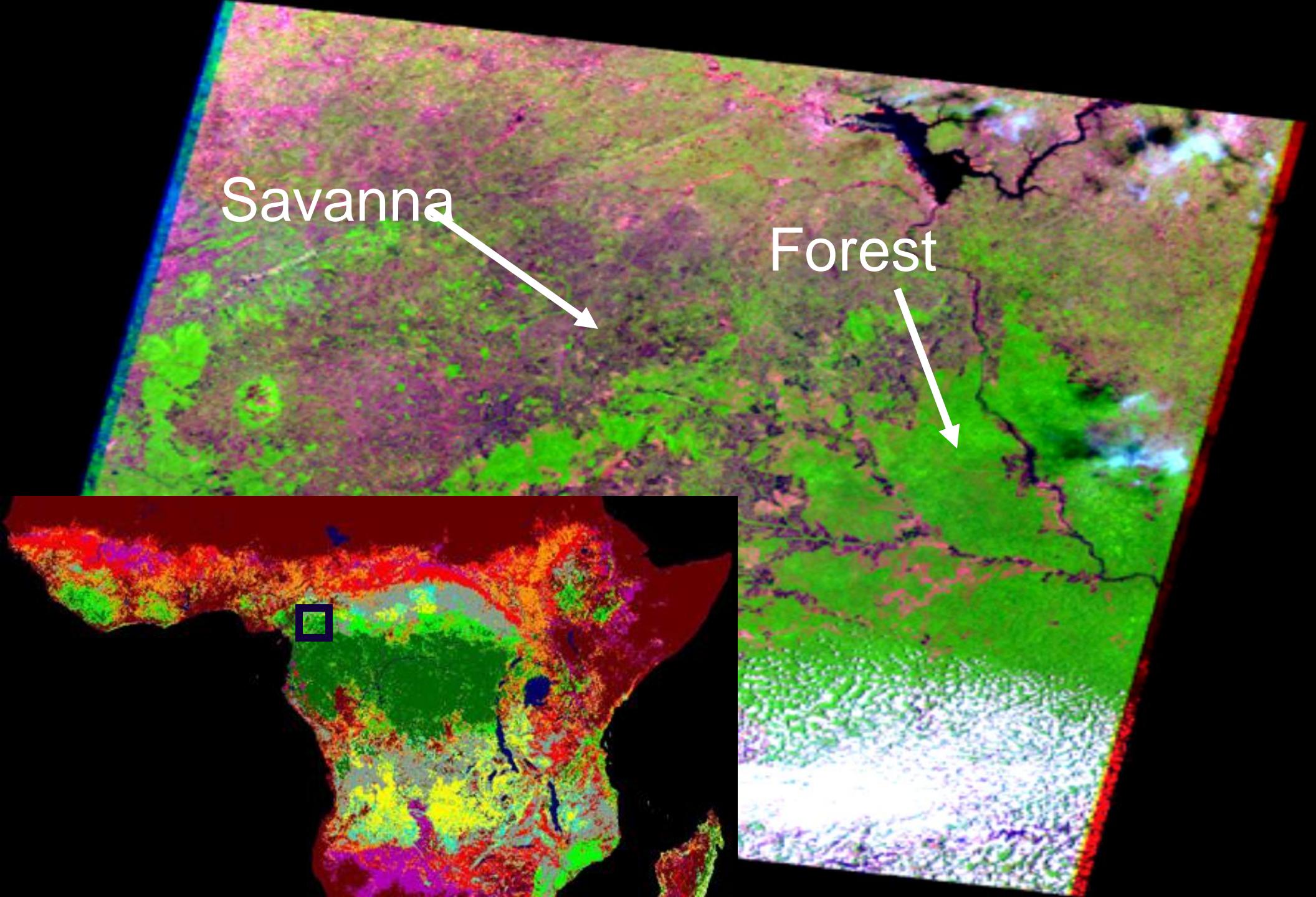
QuickTime™ and a  
TIFF (Uncompressed) decompressor  
are needed to see this picture.



# Present distribution of rainforest (stipple) and postulated locations of refuges (black)

(from Mayr and O'Hara 1986)





Savanna

Forest

N  
↑

## UPPER GUINEA

## LOWER GUINEA

Dahomey  
Gap

16○

17●

15●

13●

9●

11●

14●

10●

8●

7●

12●

6○

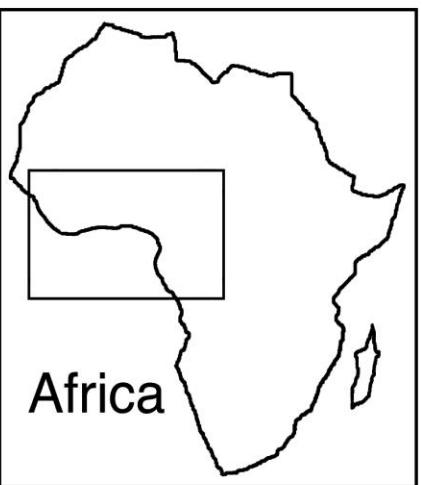
4●

1●

2●

3○

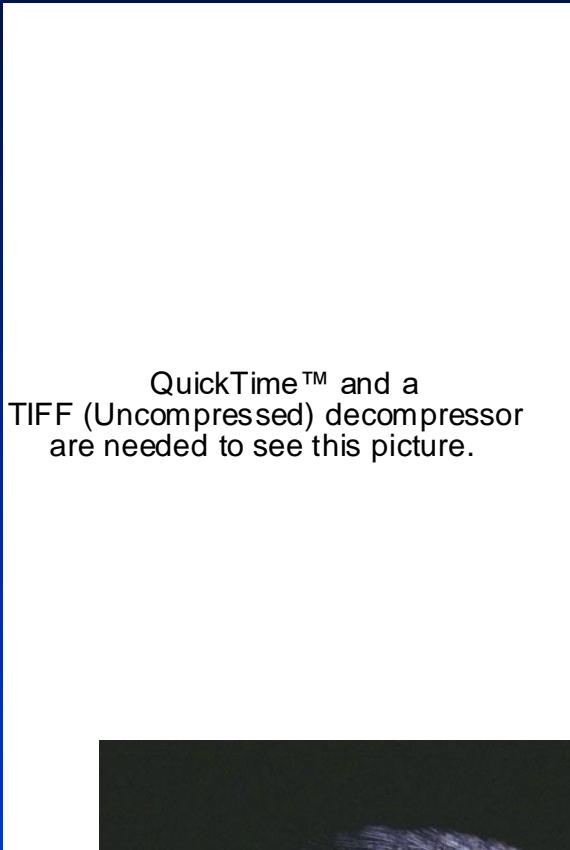
5●

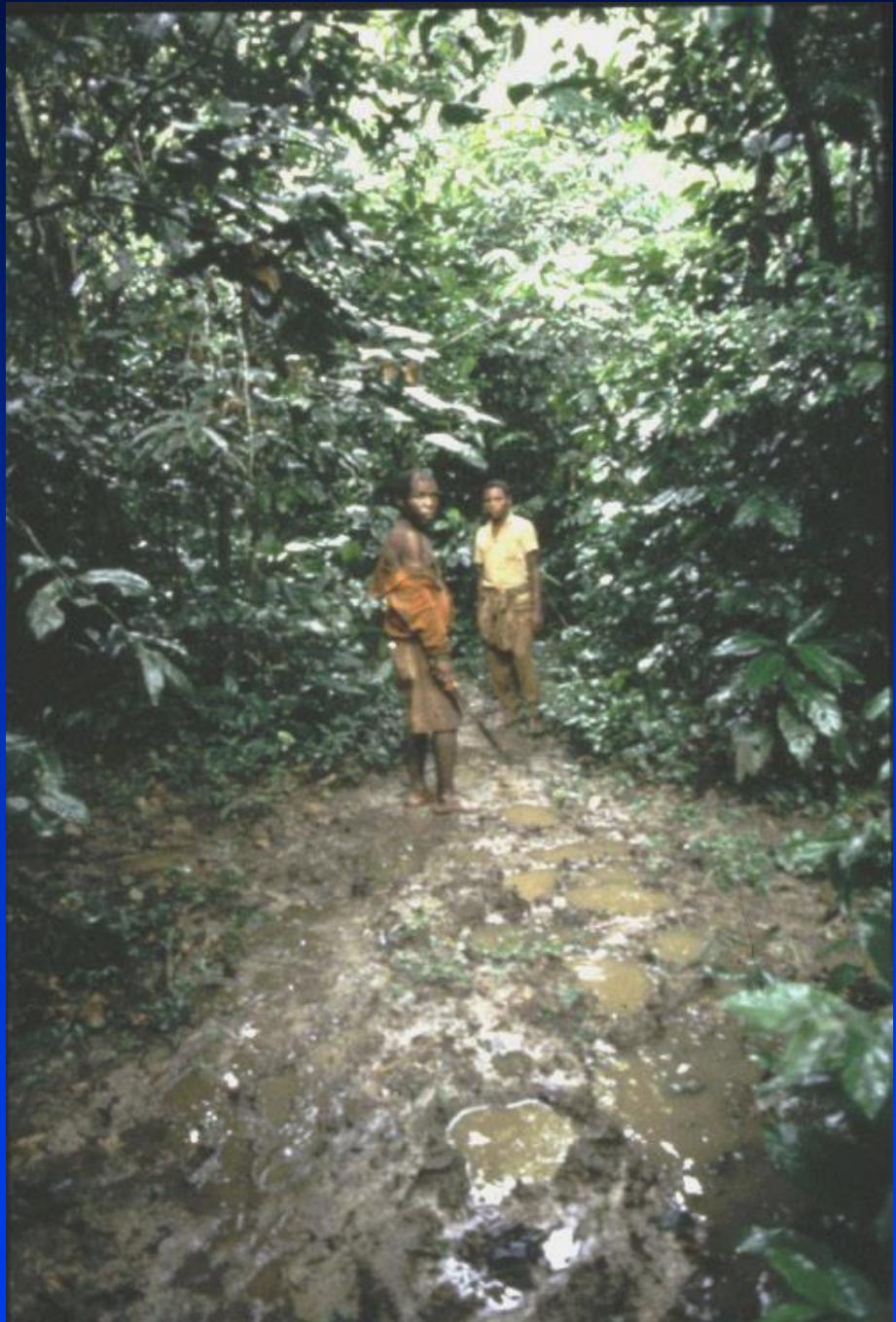


- [Savanna pattern] Savanna
- [Ecotone pattern] Ecotone
- [Forest pattern] Forest
- [Black square] Refuge

0 300 600 900 1200 miles







QuickTime™ and a  
TIFF (Uncompressed) decompressor  
are needed to see this picture.

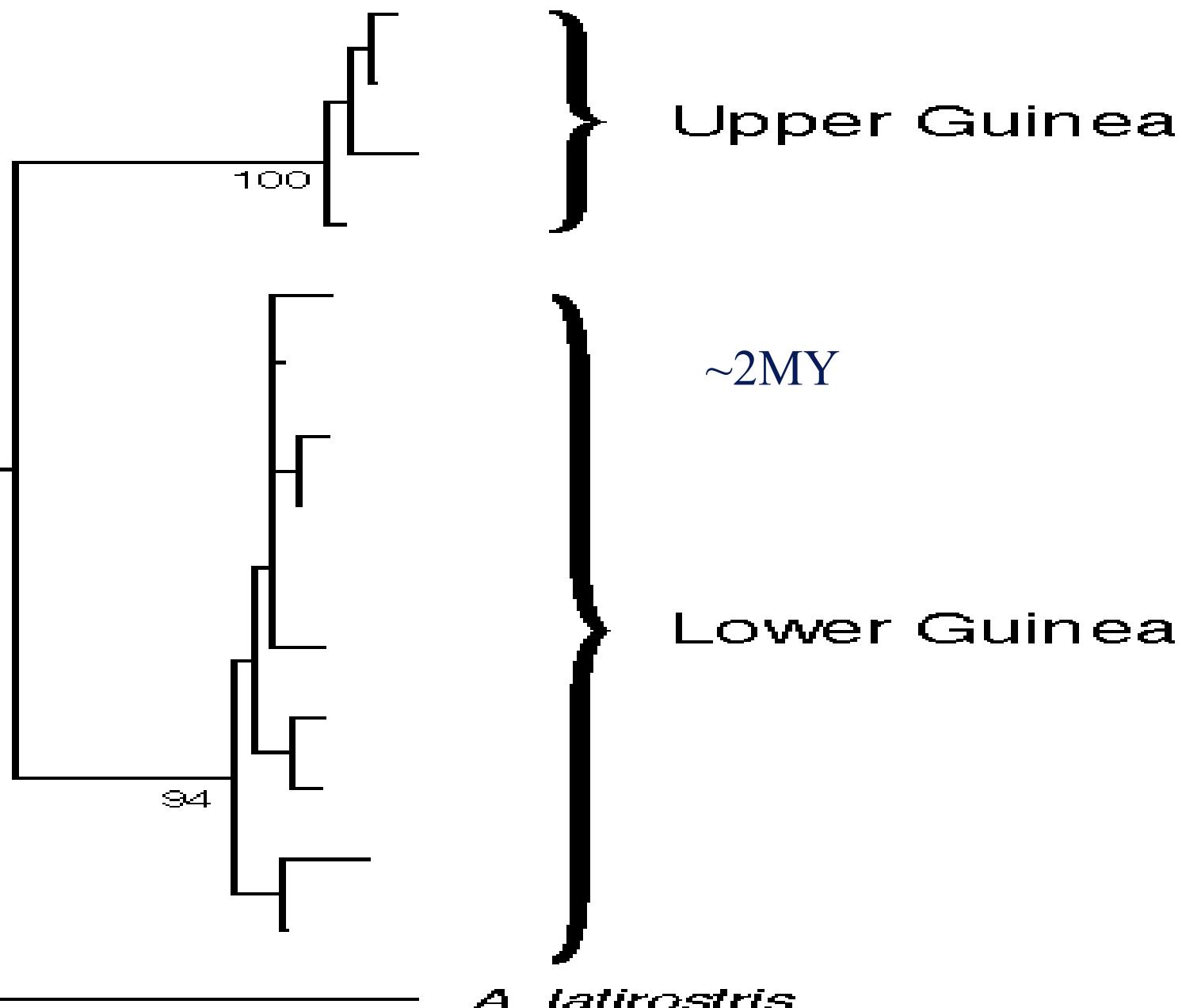




Little Greenbul  
*Andropadus*  
*virens*



*Andropadus virens*



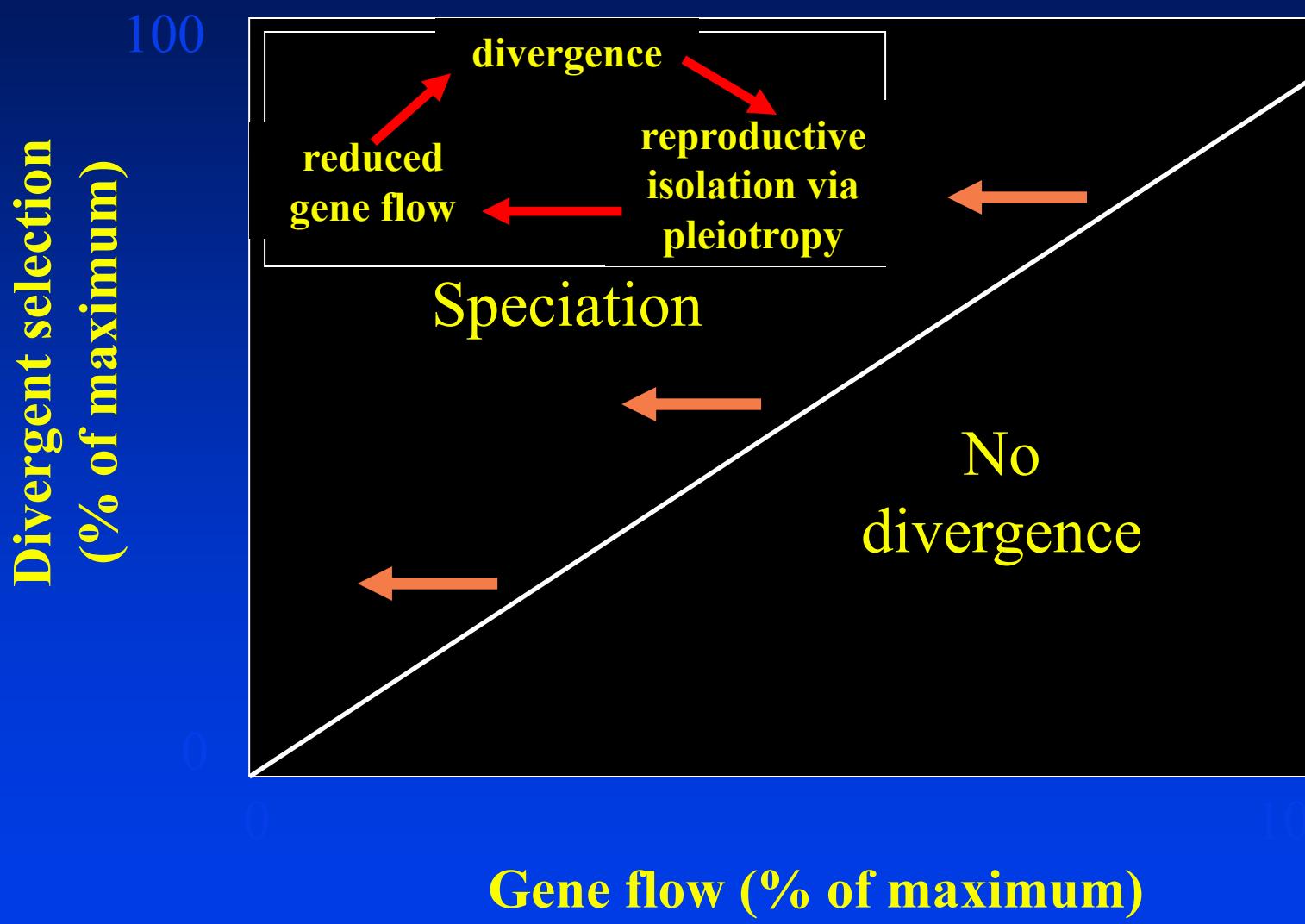
2.0%  
sequence  
divergence

Two way MANOVA between region (upper and lower guinea)  
and habitat (forest vs. ecotone)

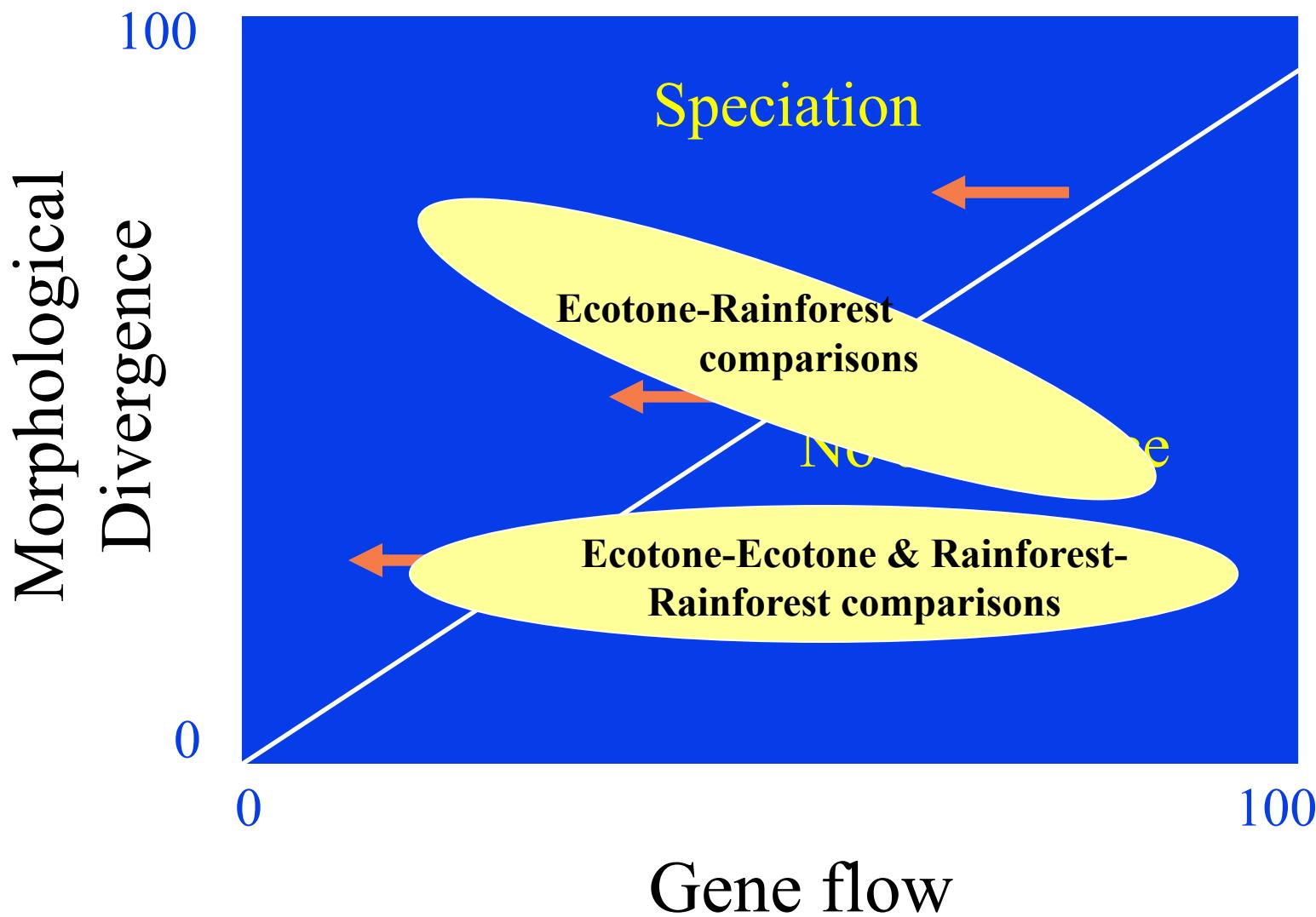
Adult Males

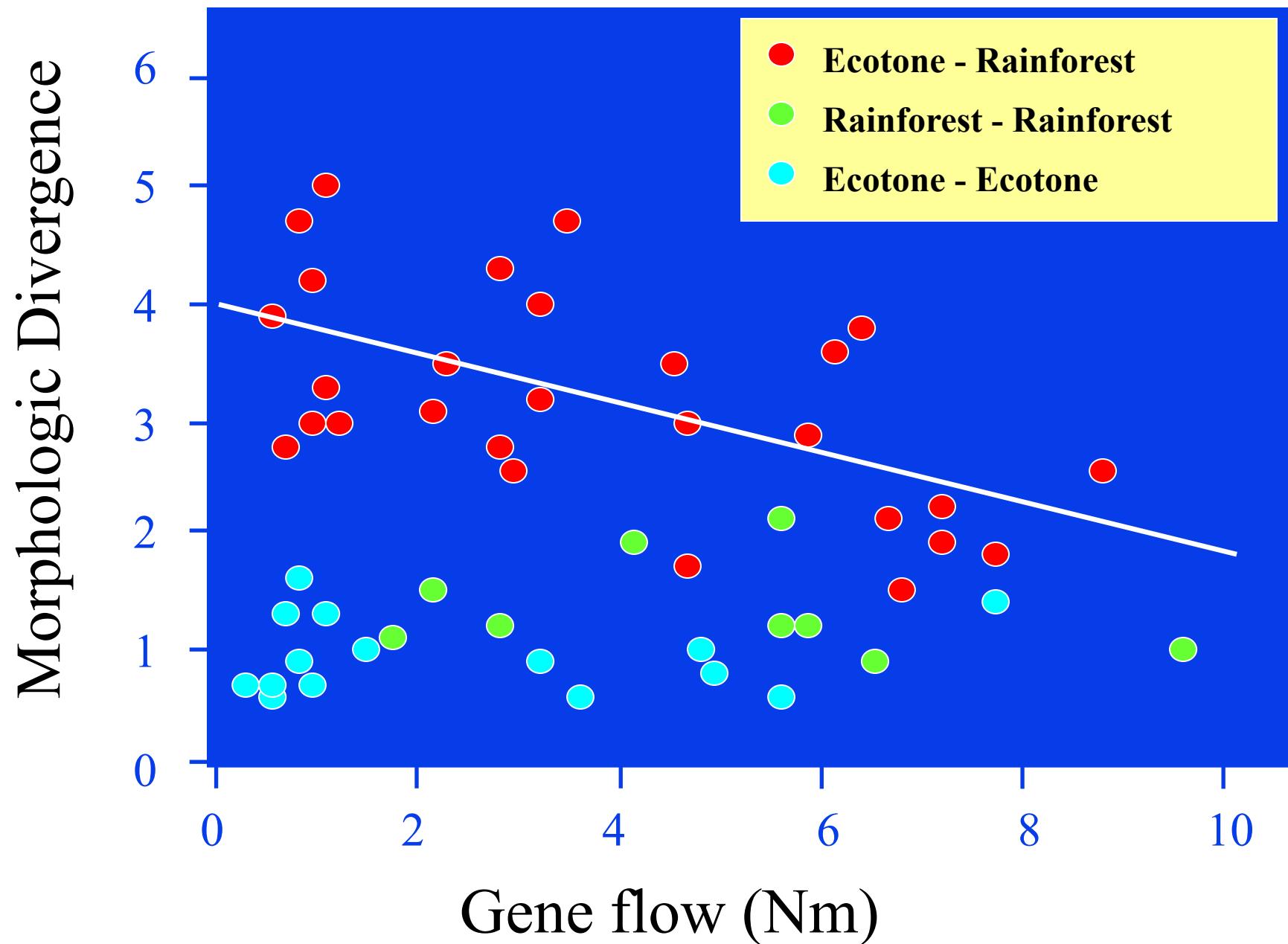
Character	Whole Model	Habitat	Region
	<i>P</i>	<i>P</i>	<i>P</i>
Mass (g)	<b>0.0001</b>	<b>0.0001</b>	<b>0.01</b>
Wing length (mm)	<b>0.0001</b>	<b>0.0001</b>	0.35
Tail length	<b>0.001</b>	<b>0.0001</b>	0.41
Tarsus length	<b>0.0001</b>	<b>0.0001</b>	0.29
Upper mandible l.	<b>0.01</b>	0.49	<b>0.002</b>
Bill depth	<b>0.06</b>	<b>0.022</b>	0.64

# Divergence-with-gene-flow model of speciation



# Divergence-with-gene-flow in little greenbuls





(Smith et al. Science 1997)

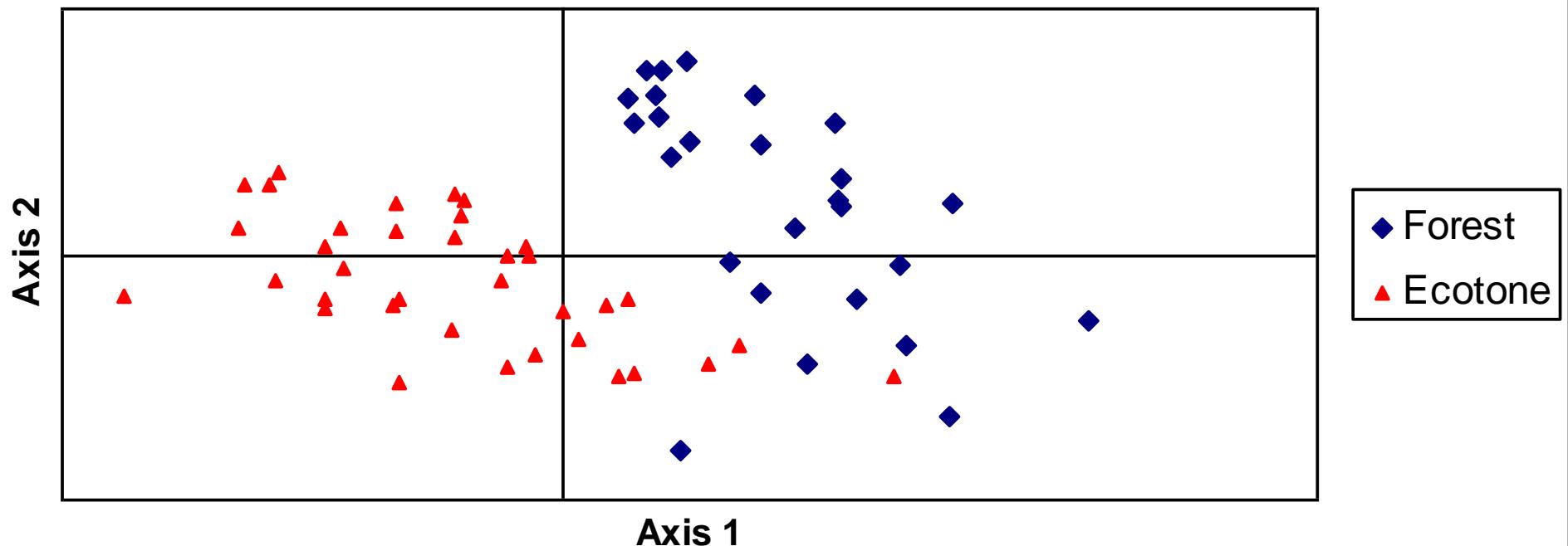
# Evidence of morphological divergence and divergence with gene flow

What is the evidence for genetic  
divergence and reproductive isolation?

# Principal Coordinates Analysis of AFLP Markers

75 individuals  
4 Forest sites; 5 Ecotone sites  
210 markers (c9 + c13)

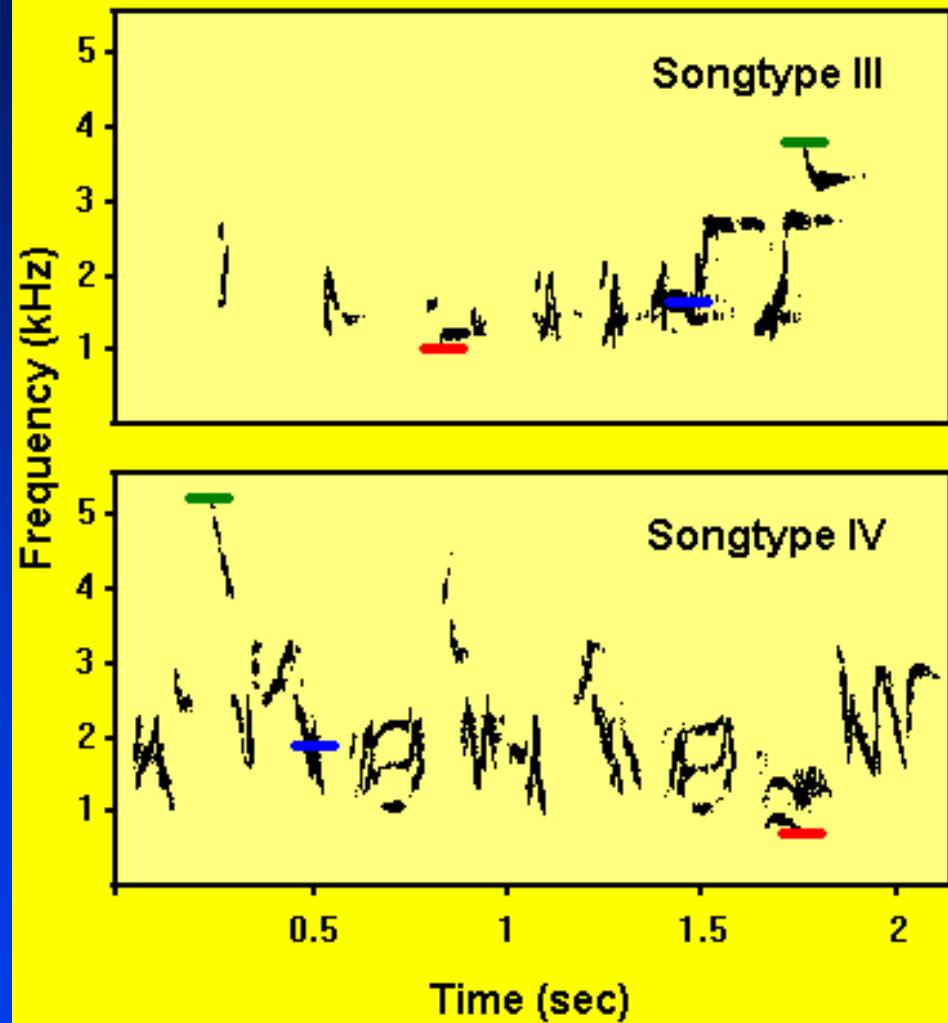
Principal Coordinates (1 vs 2)



# Breeding activity of *Andropadus virens* by habitat and period

<u>Habitat</u>	<u>June-July</u>	<u>Jan-Feb</u>
ECOTONE	-	+
FOREST	+	-

## Acoustic Divergence



	Rainforest	Ecotone	F	P
$F_{\text{max}}$	3669	3385	44.65	<0.001
$F_{\text{peak}}$	2069	2093	0.002	NS
$F_{\text{min}}$	1009	1036	19.67	<0.001
Delrate	7.8	6.6	49.27	<0.001
$F_{\text{max}}$	5086	5206	4.10	NS
$F_{\text{peak}}$	2398	2318	4.48	NS
$F_{\text{min}}$	904	957	37.18	<0.001
Delrate	8.6	8.6	0.27	NS

\*\*  
\*\*  
\*\*  
\*\*

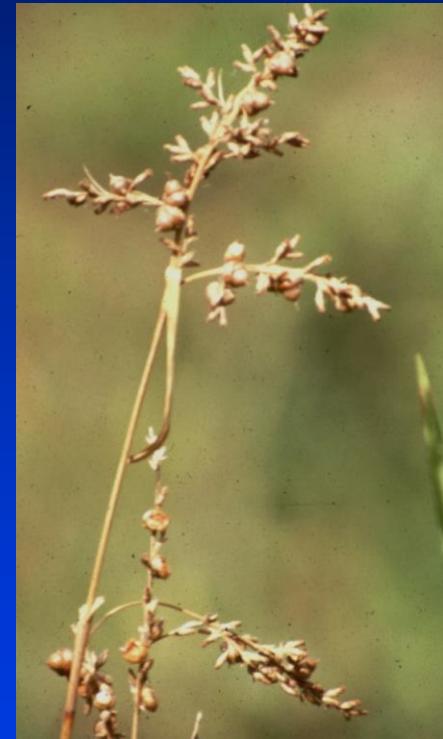
# Evidence for reproductive divergence?

## Prezygotic isolating barriers

- ✓ 1) Habitat      forest vs. ecotone
- ✓ 2) Temporal differ in breeding time
- ✓ 3) Behavioral      differences in song

# Agents of Natural Selection?

- Food?
- Disease?
- Habitat structure?
- Predation?



# Black-bellied Seedcracker *Pyrenestes ostrinus*

Large and small morph in the forest





Morph  
specializing  
small |  
large |

## SEED HARDNESS

Species	Maximum Fracture Force (N)	Diameter (mm)
	Mean $\pm$ SE	Mean $\pm$ SE
Soft sedge ( <i>Scleria goossensisii</i> )	$12.3 \pm 2.0$	$2.92 \pm 0.44$
Hard sedge ( <i>S. verrucosa</i> )	$146.9 \pm 2.0$	$3.26 \pm 0.02$





## Third bill morph in ecotone

$F_{st} = 0.005$  (ns)  
between mega and  
other morphs

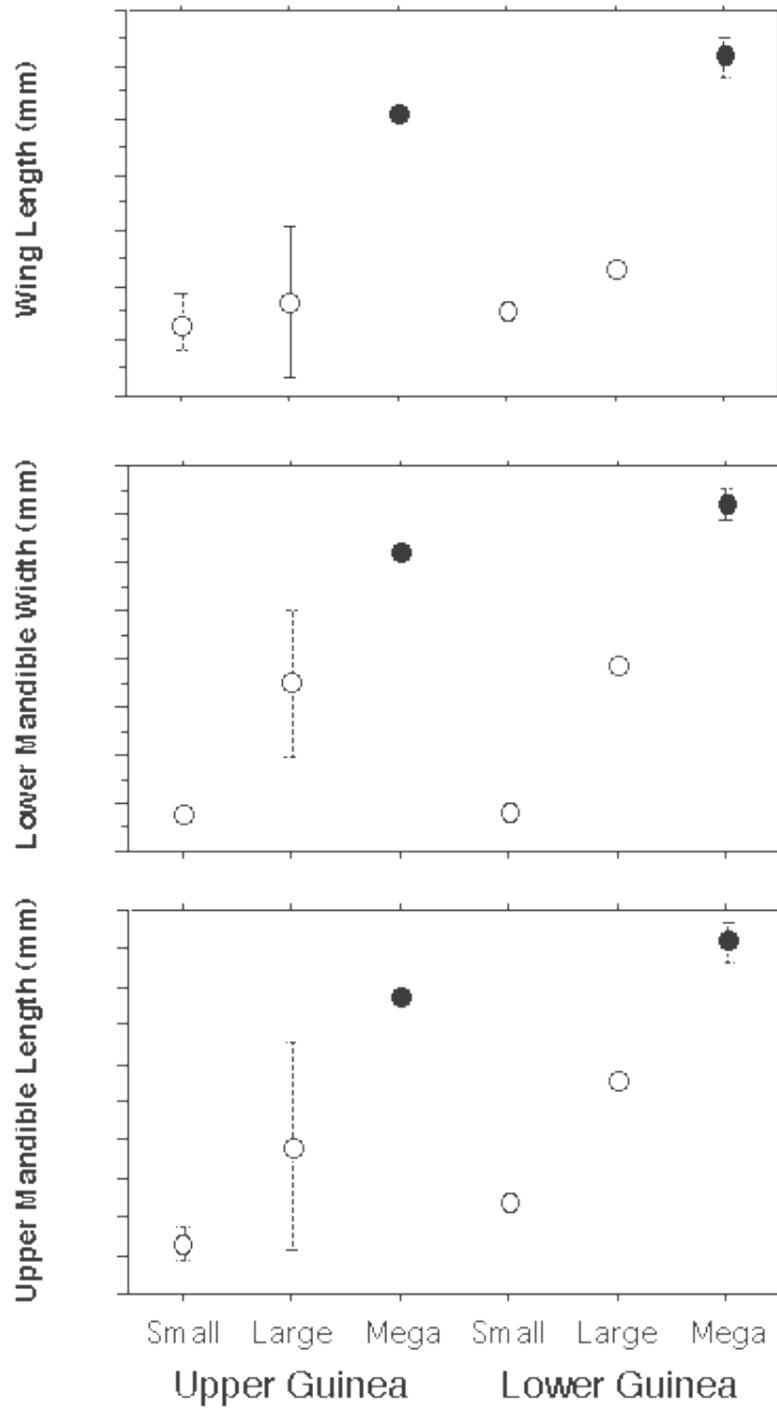
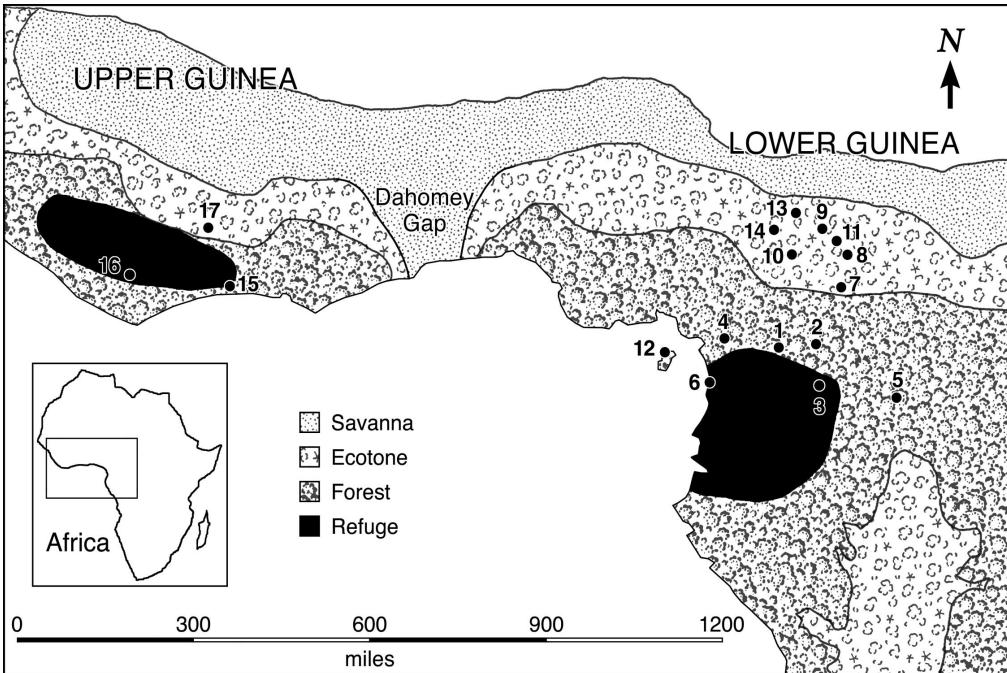


## SEED HARDNESS

	Species	Maximum Fracture Force (N)	Diameter (mm)
Morph		Mean $\pm$ SE	Mean $\pm$ SE
small	Soft sedge ( <i>Scleria goossensii</i> )	$12.3 \pm 2.0$	$2.92 \pm 0.44$
large	Hard sedge ( <i>S. verrucosa</i> )	$146.9 \pm 2.0$	$3.26 \pm 0.02$
mega	Ecotone Hard Sedge ( <i>S. racemosa</i> )	$299.5 \pm 11.9$ ECOTONE ONLY	$4.12 \pm 0.32$



# Pattern of morphological variation between refugia is similar



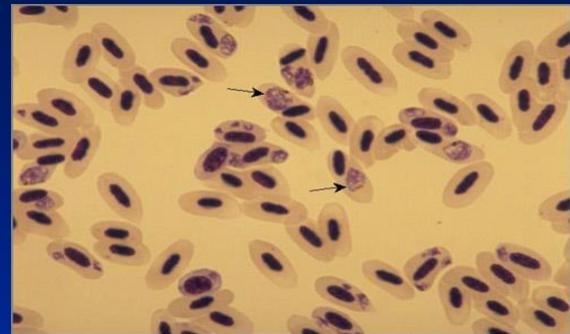
# Agents of Natural Selection?

- Food?
- Disease?



# Blood-borne parasites

*Plasmodium relictum*



*Haemoproteus columbae*

## Vectors

mosquito



lousefly



midge

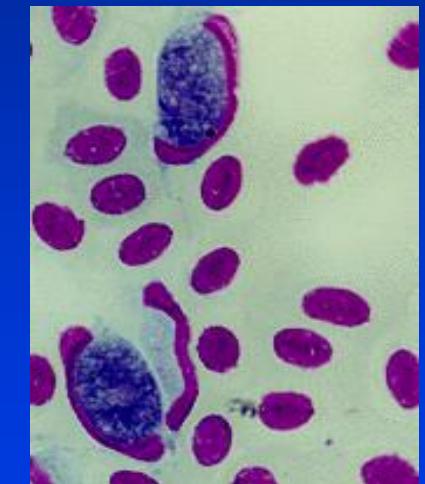
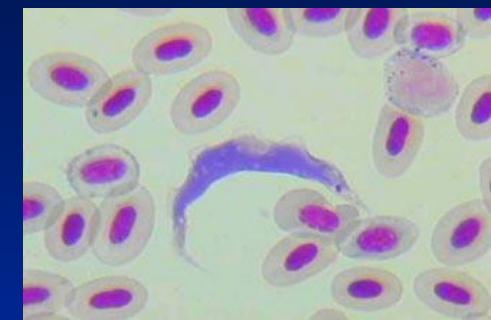


blackfly



## Host

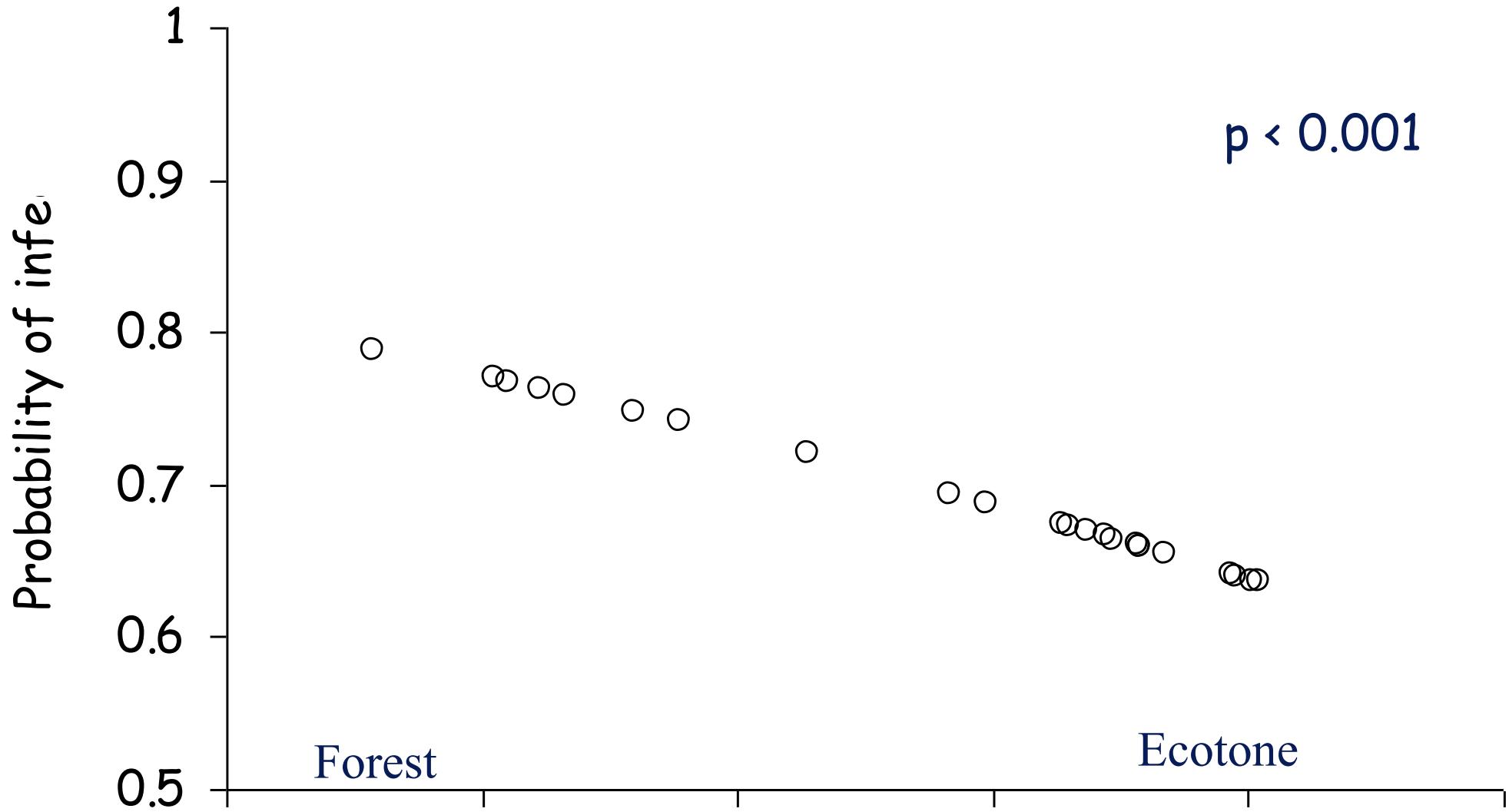
*Trypanosoma avium*



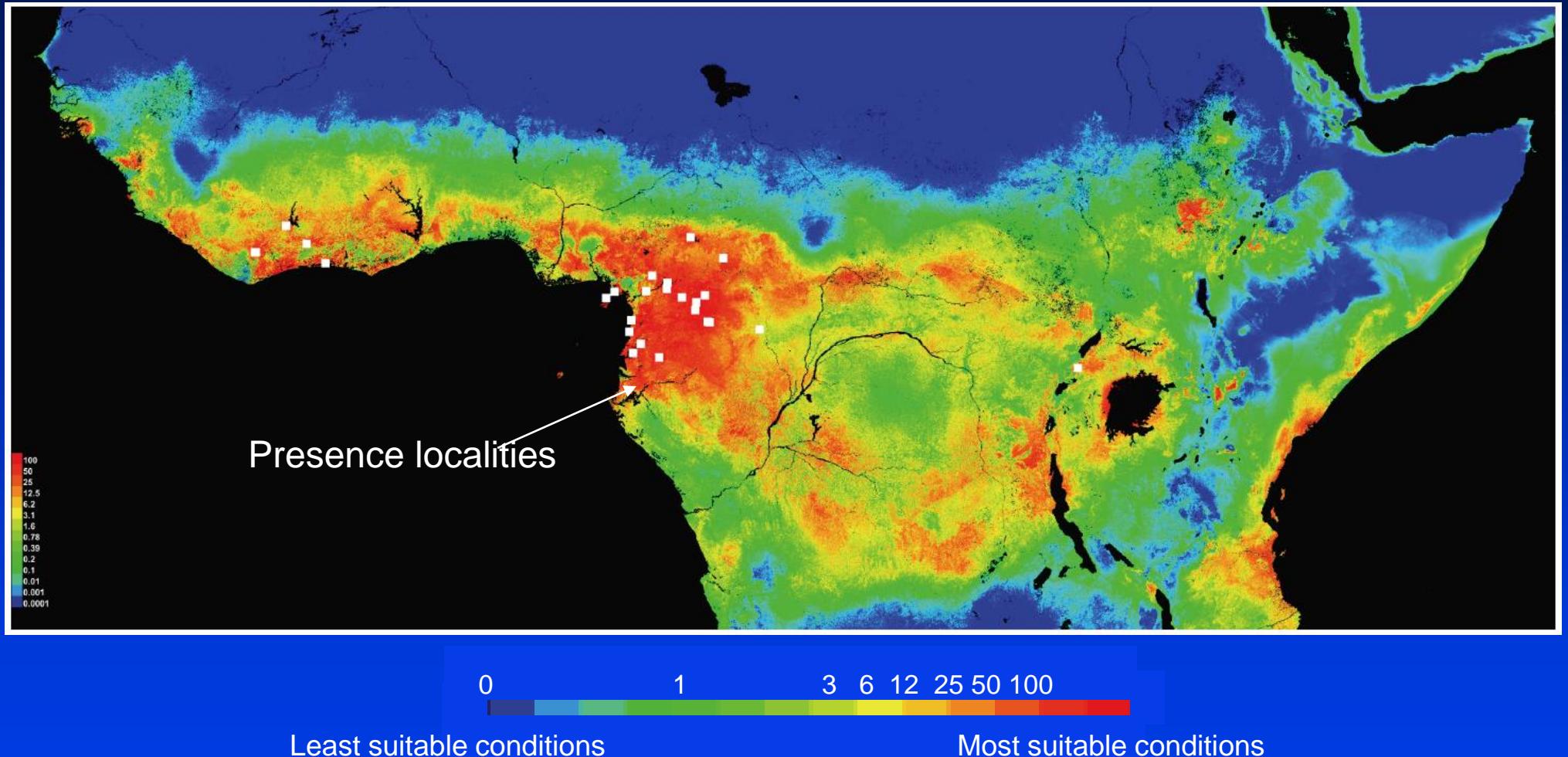
*Leucocytozoon smithi*

# Infection with *Plasmodium* &/or *Haemoproteus*

QuickTime™ and a  
TIFF decompressor  
are needed to see this picture.



# Africa - Maxent Predictions for Avian Malaria with prevalence\*



\* Presence localities weighted by relative prevalence

# Moving beyond simple surrogates of biodiversity...

## Putting Process on the Map

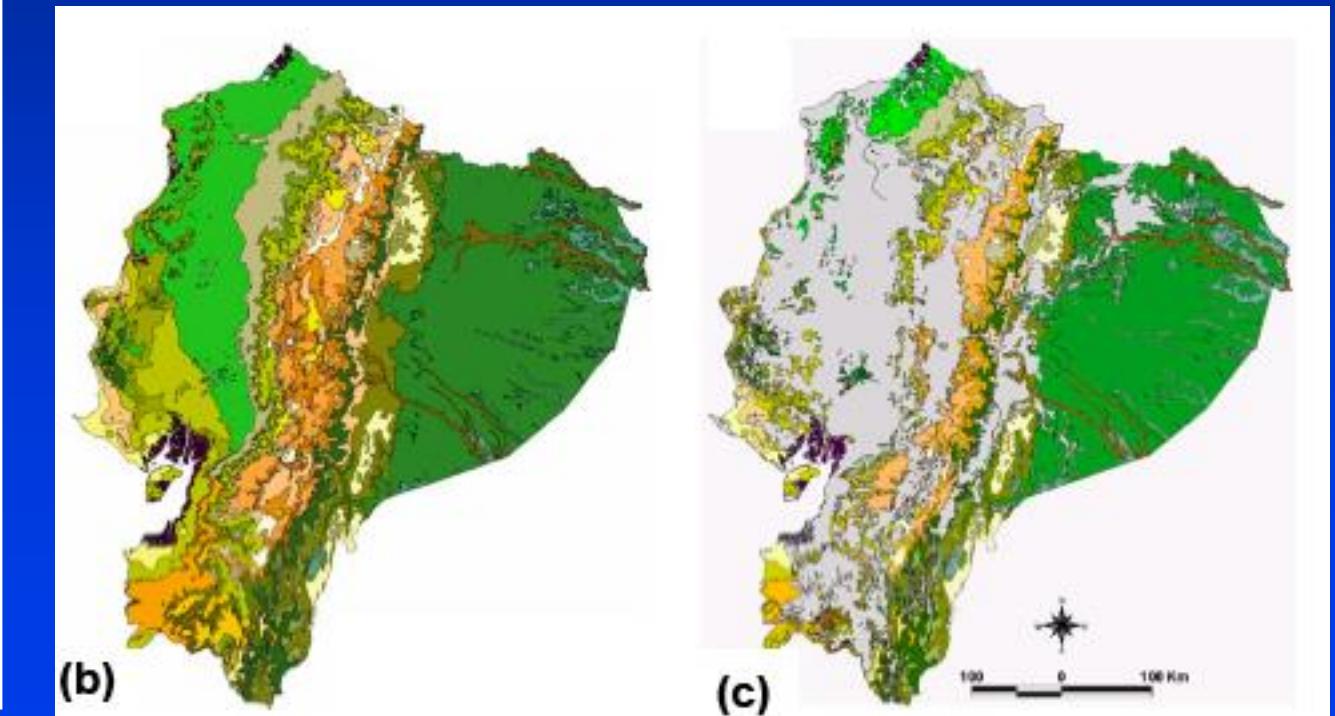
The steps:

- 1) Assess biodiversity pattern
- 2) Assess biodiversity process
- 3) Predict the influence of climate change
- 4) Combine 1-3 and levels of threat to identify critically important regions

# Ecuador

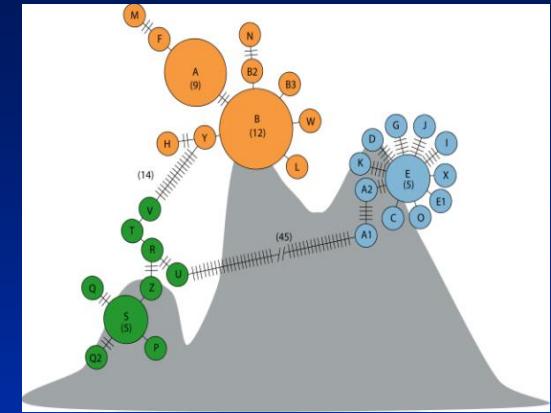


QuickTime™ and a  
TIFF (Uncompressed) decompressor  
are needed to see this picture.



# Point locality, genetic and phynotypic data

- Birds
- Bats
- Frogs
- Small mammals
- Primates\*
- Trees\*

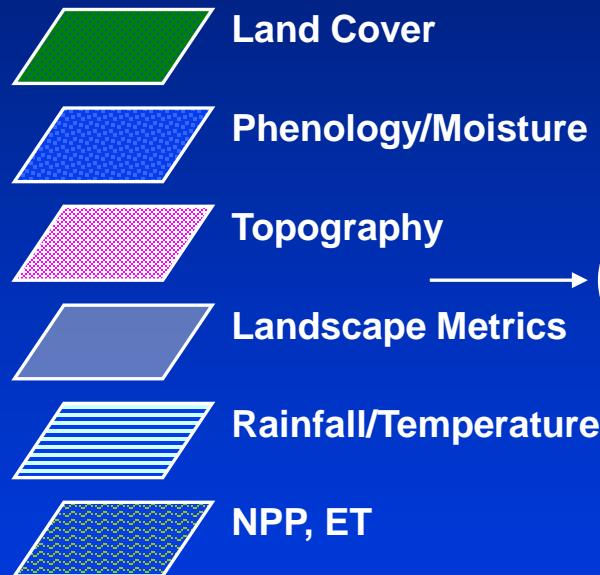


\*distribution data only

White-plumed antbird  
(*Pithys albifrons*)

# Spatial Analysis Model

## Remote Sensing & Environmental Variable Layers



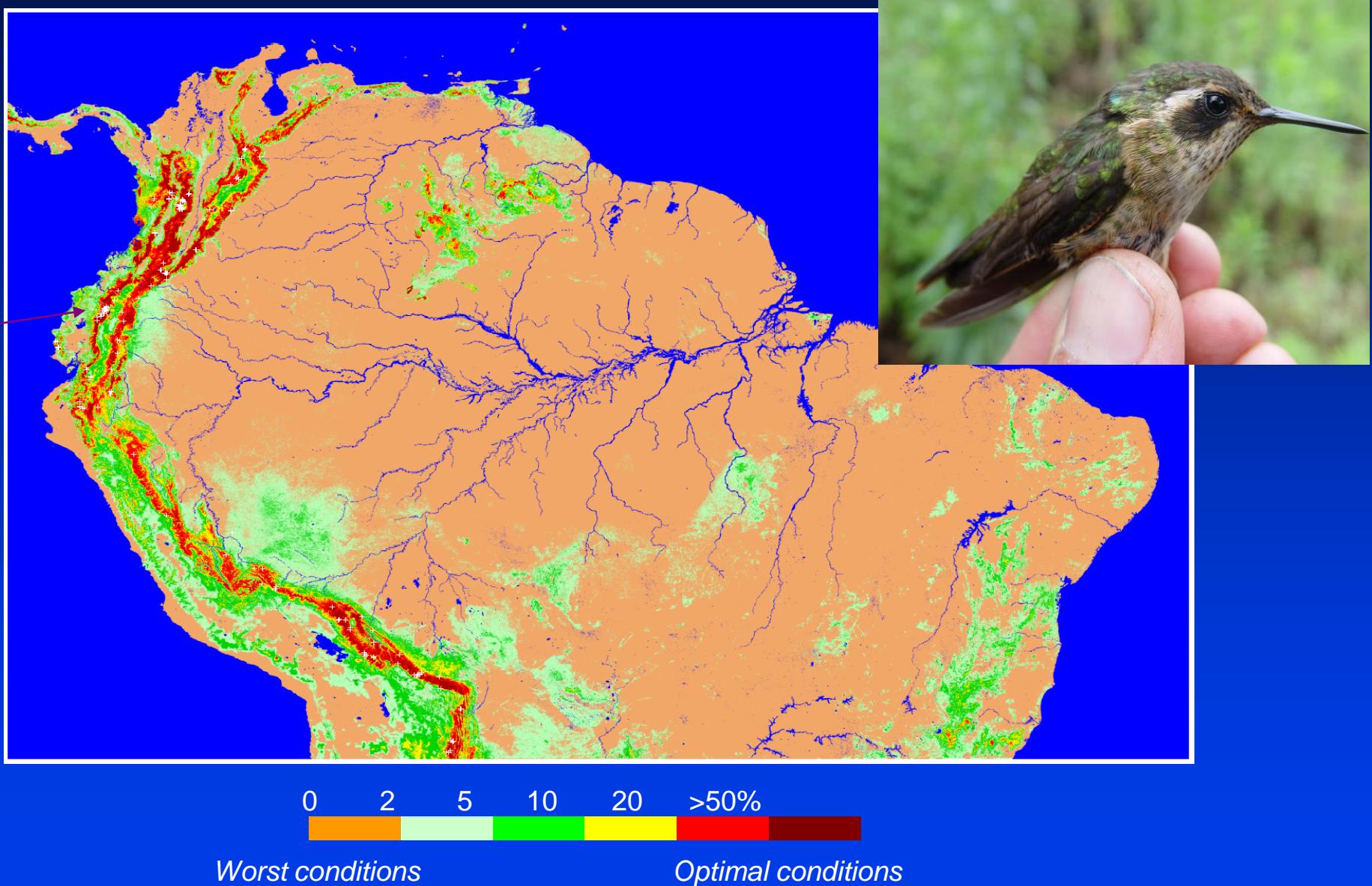
Biodiversity  
Spatial  
Analysis

Current &  
Predicted  
Distribution of  
Species  
Diversity

Species Point Locality Data

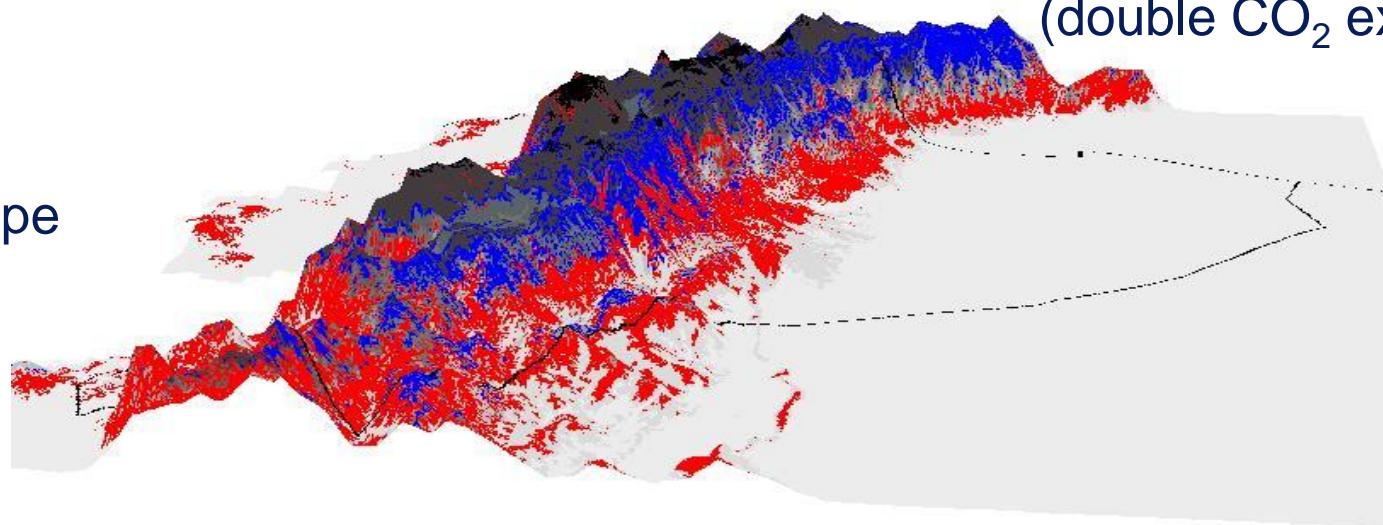
# Predicted Range of the Speckled Hummingbird

*Point  
Locations*



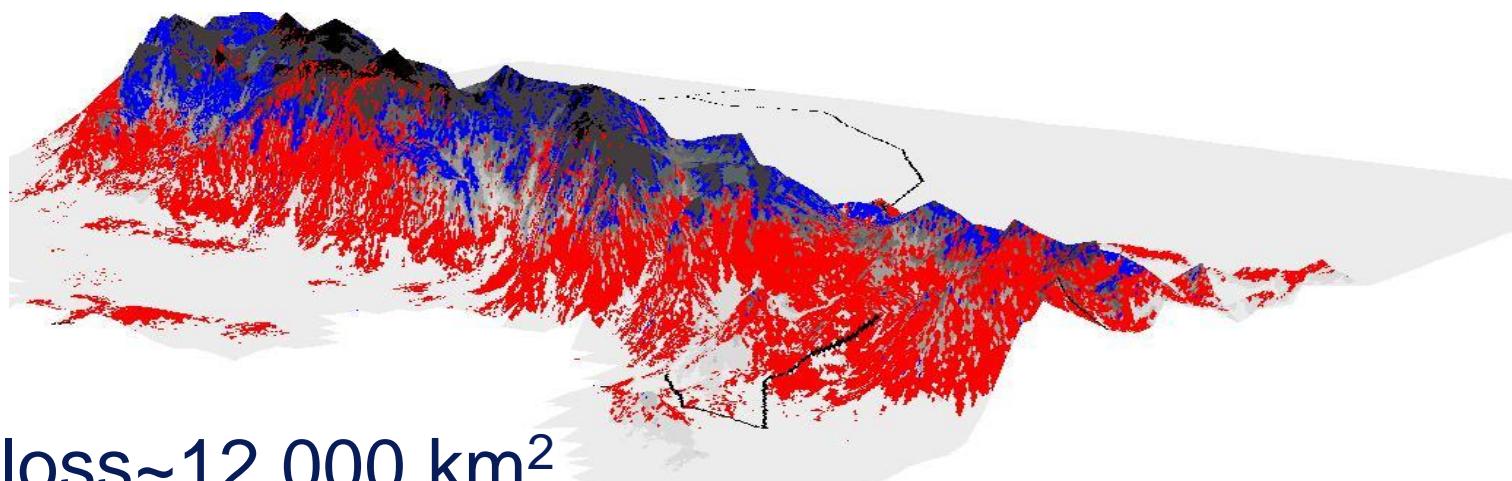
# Net loss of speckled hummingbird habitat with climate warming (double CO<sub>2</sub> experiment)

East Slope  
Andes



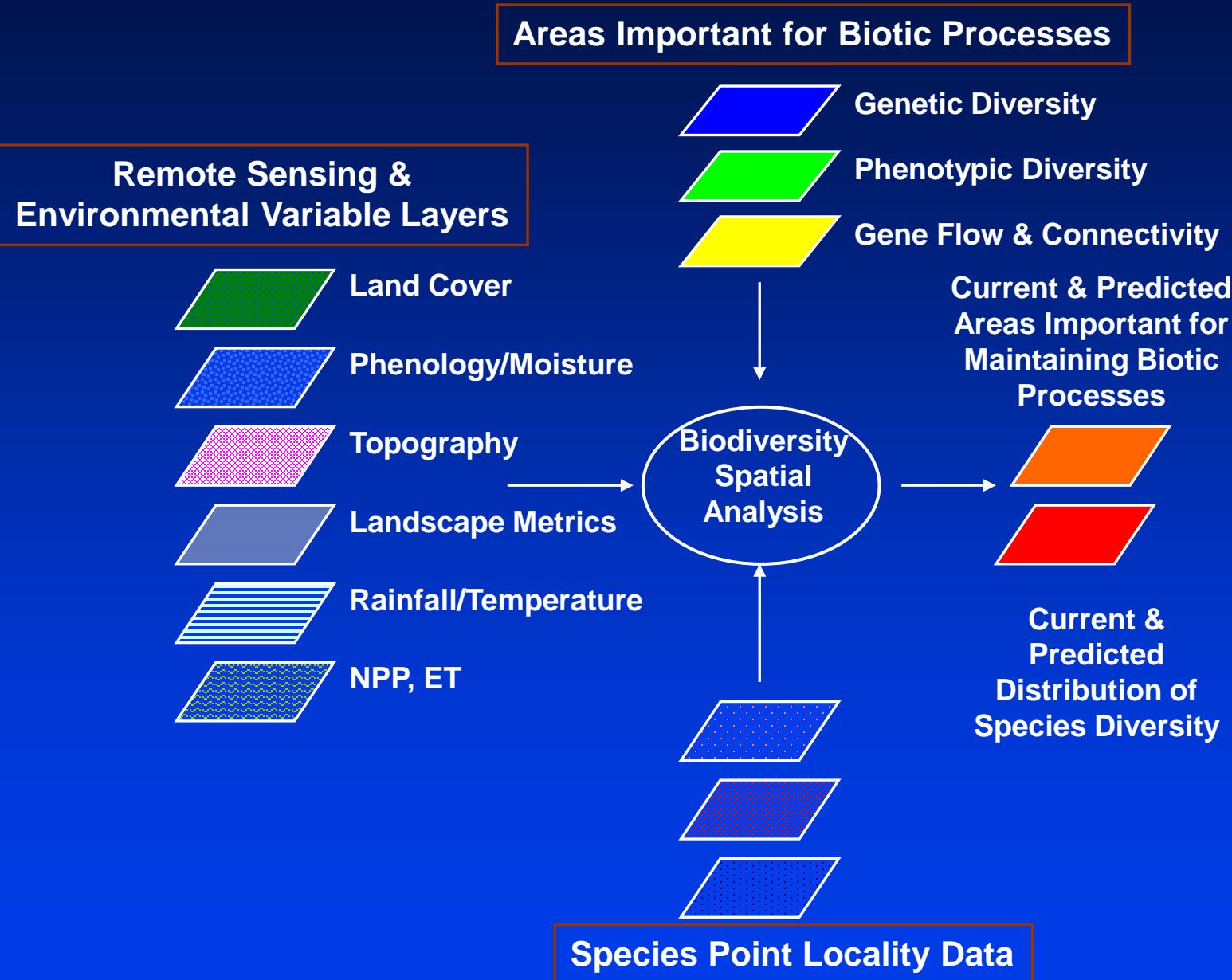
■ Areas becoming less suitable ■ Areas becoming more suitable

West Slope  
Andes



Habitat loss~12,000 km<sup>2</sup>

# Spatial Analysis Model

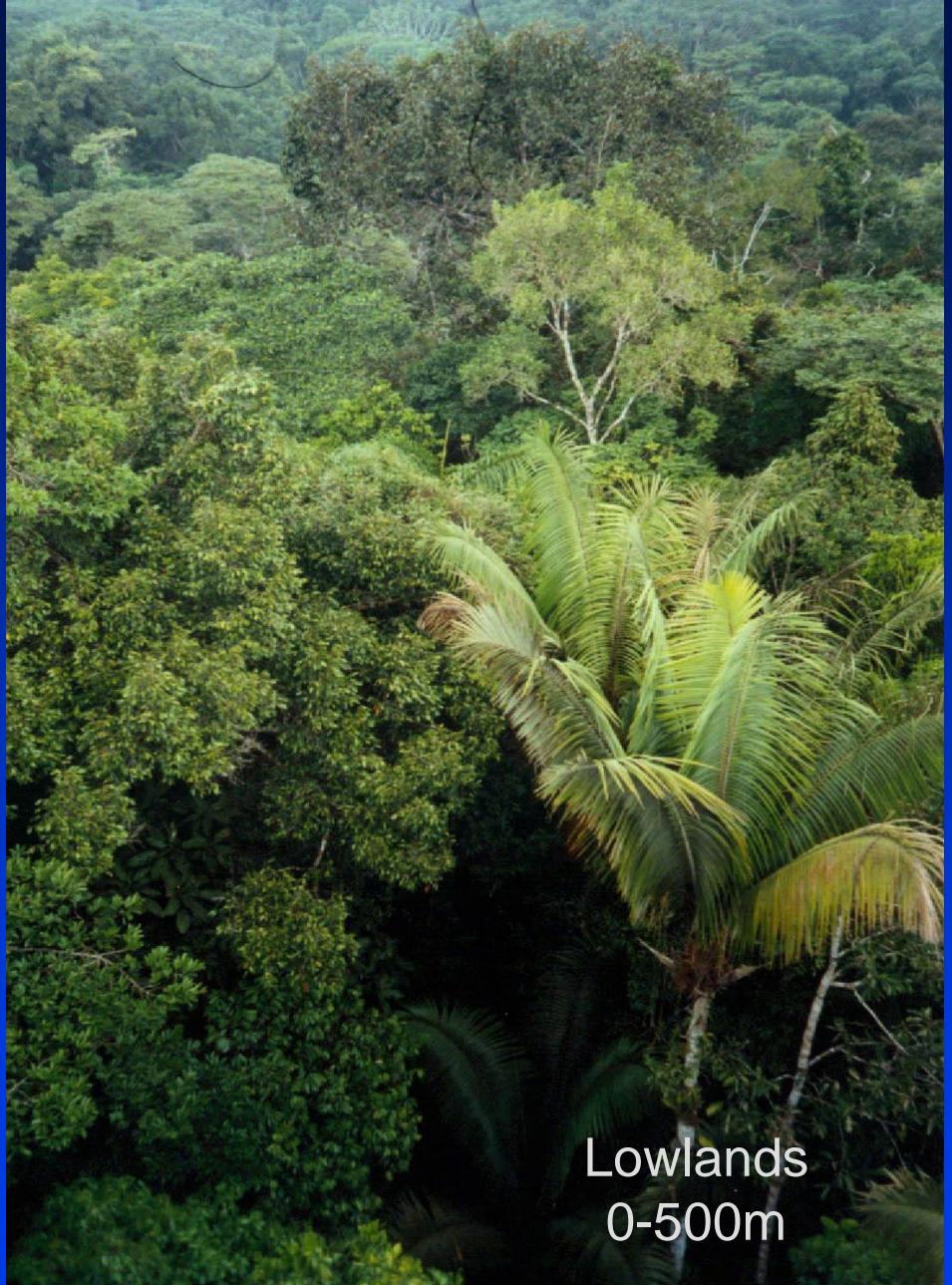


# Evidence for differentiation along ecological gradients in the Andes

Wedge-billed woodcreeper  
(*Glyphorynchus spirurus*)

N=196  
29 sites





Lowlands  
0-500m



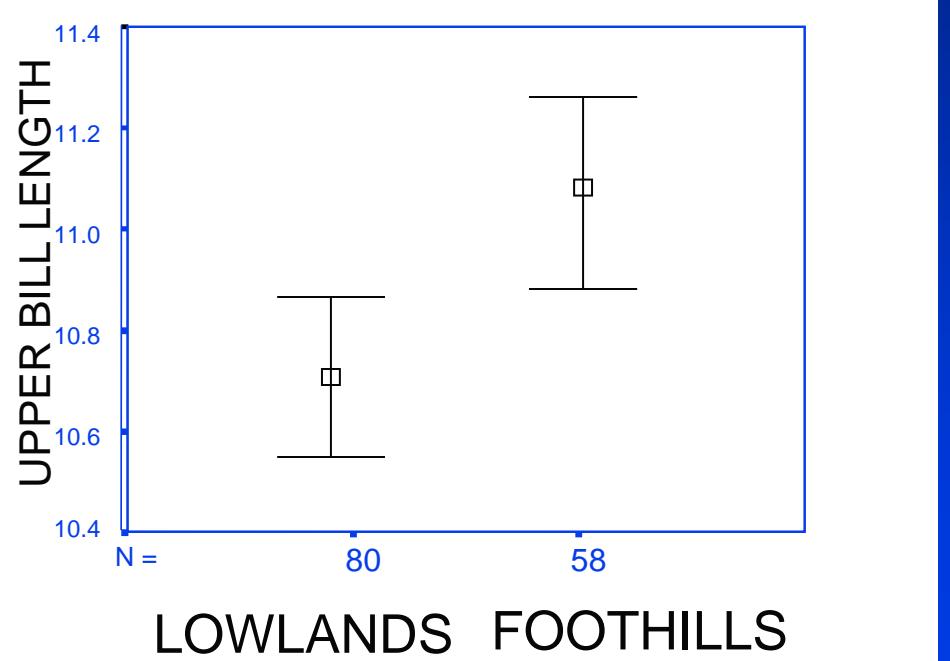
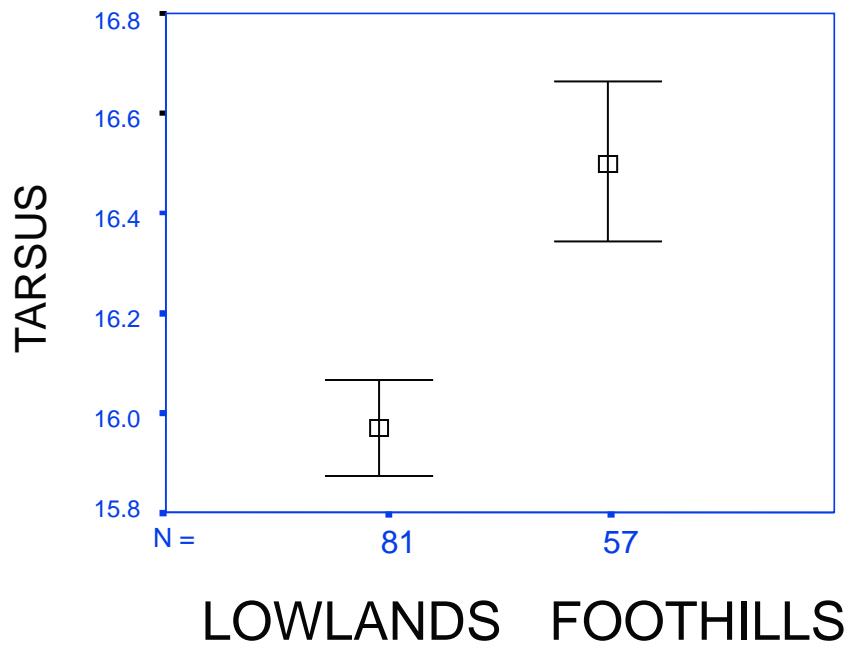
Foothills  
800-1500m



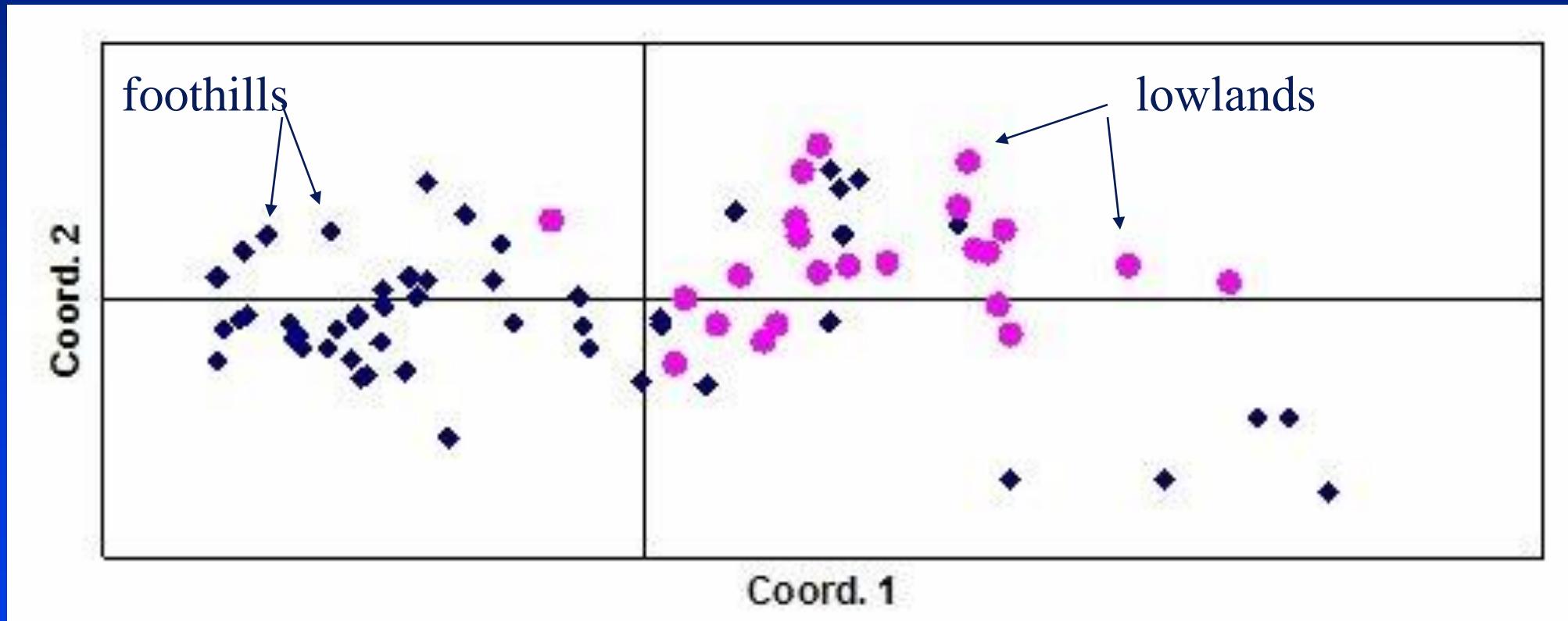
# Morphological analysis

## Tarsus length

## Bill length



# Principal coordinates analysis of 74 individuals from the eastern slope of the Andes using 126 AFLP markers



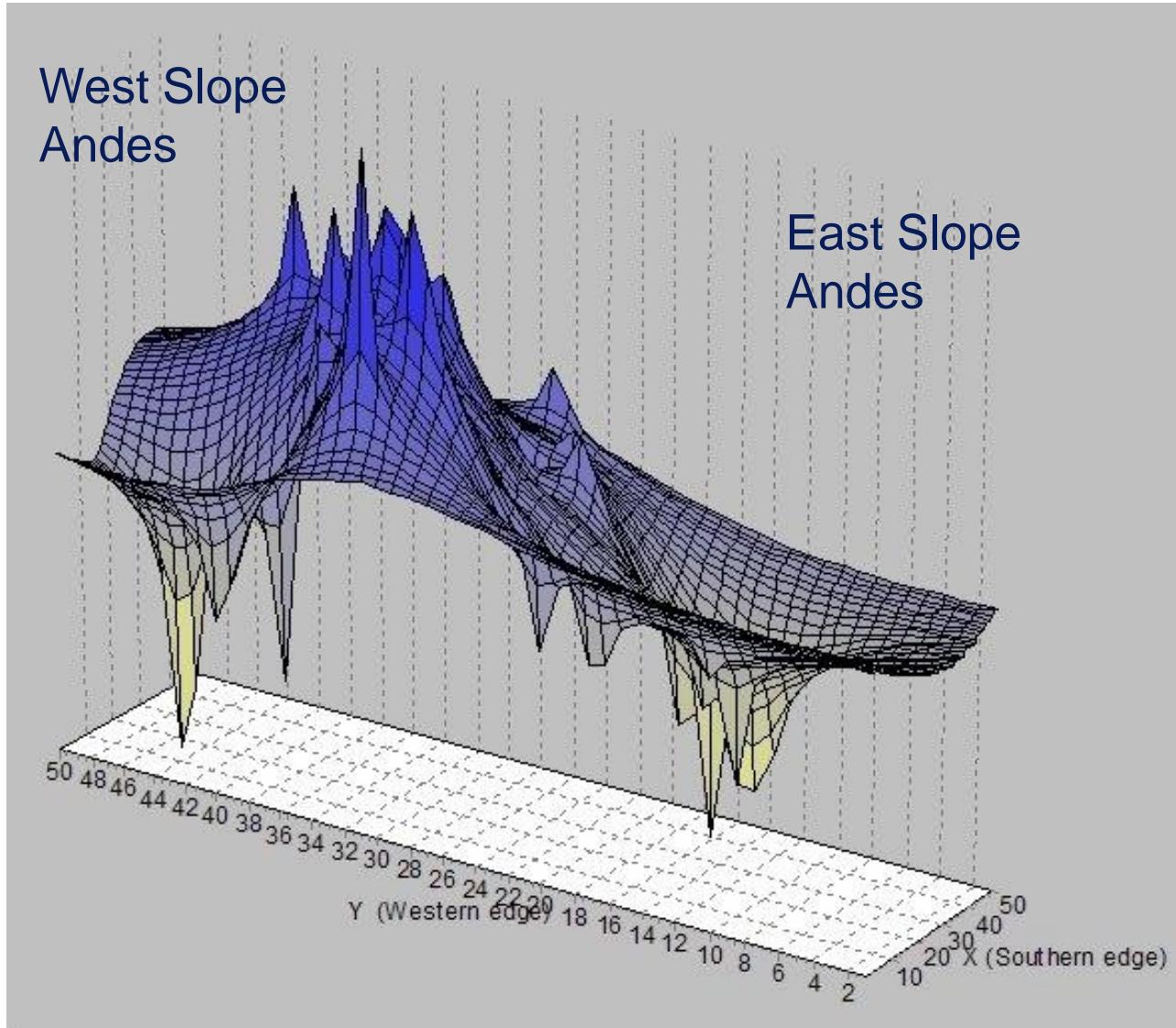


Lowlands

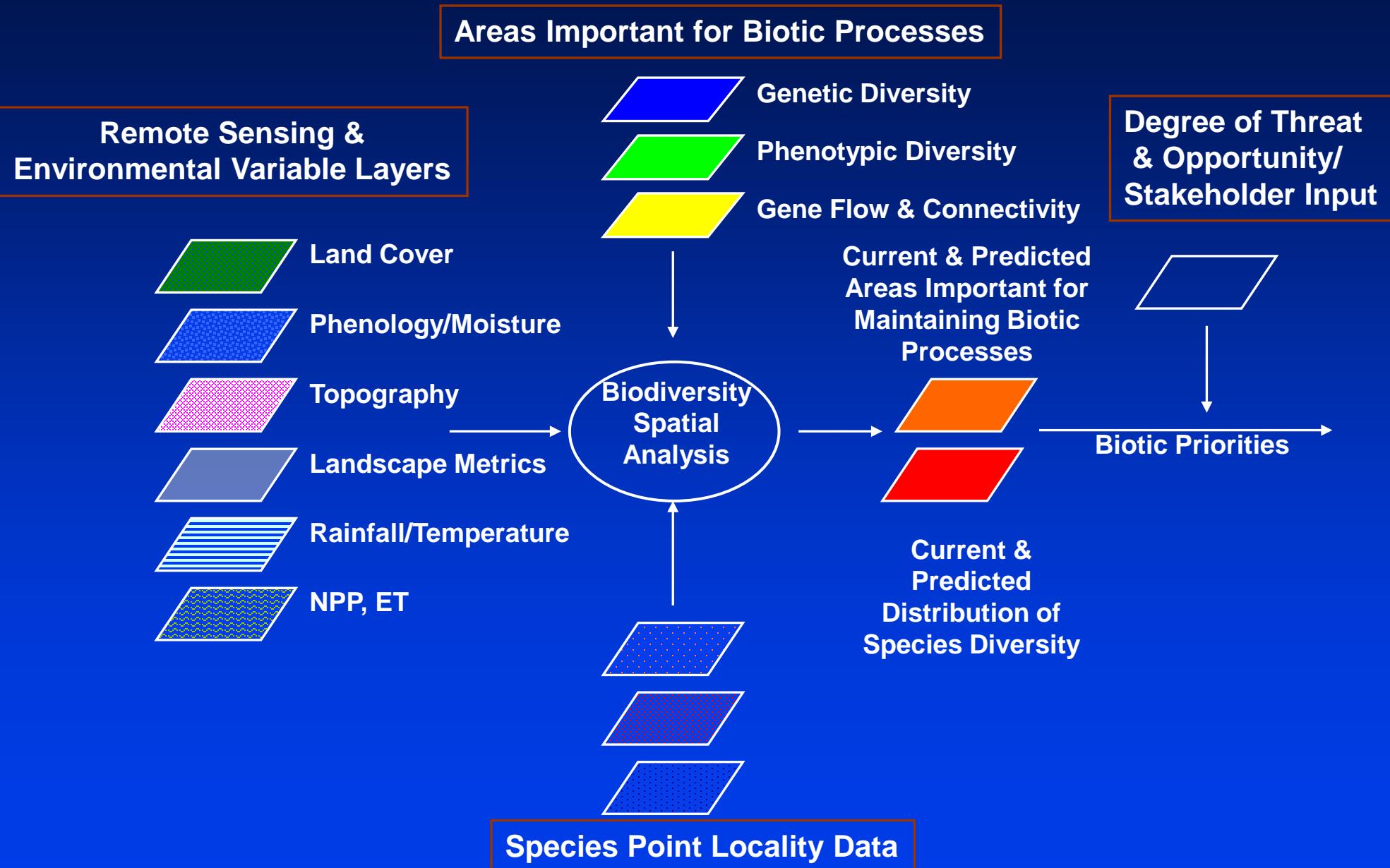


Foothills

# Genetic landscape for wedge-billed woodcreeper populations in the Andes of Ecuador



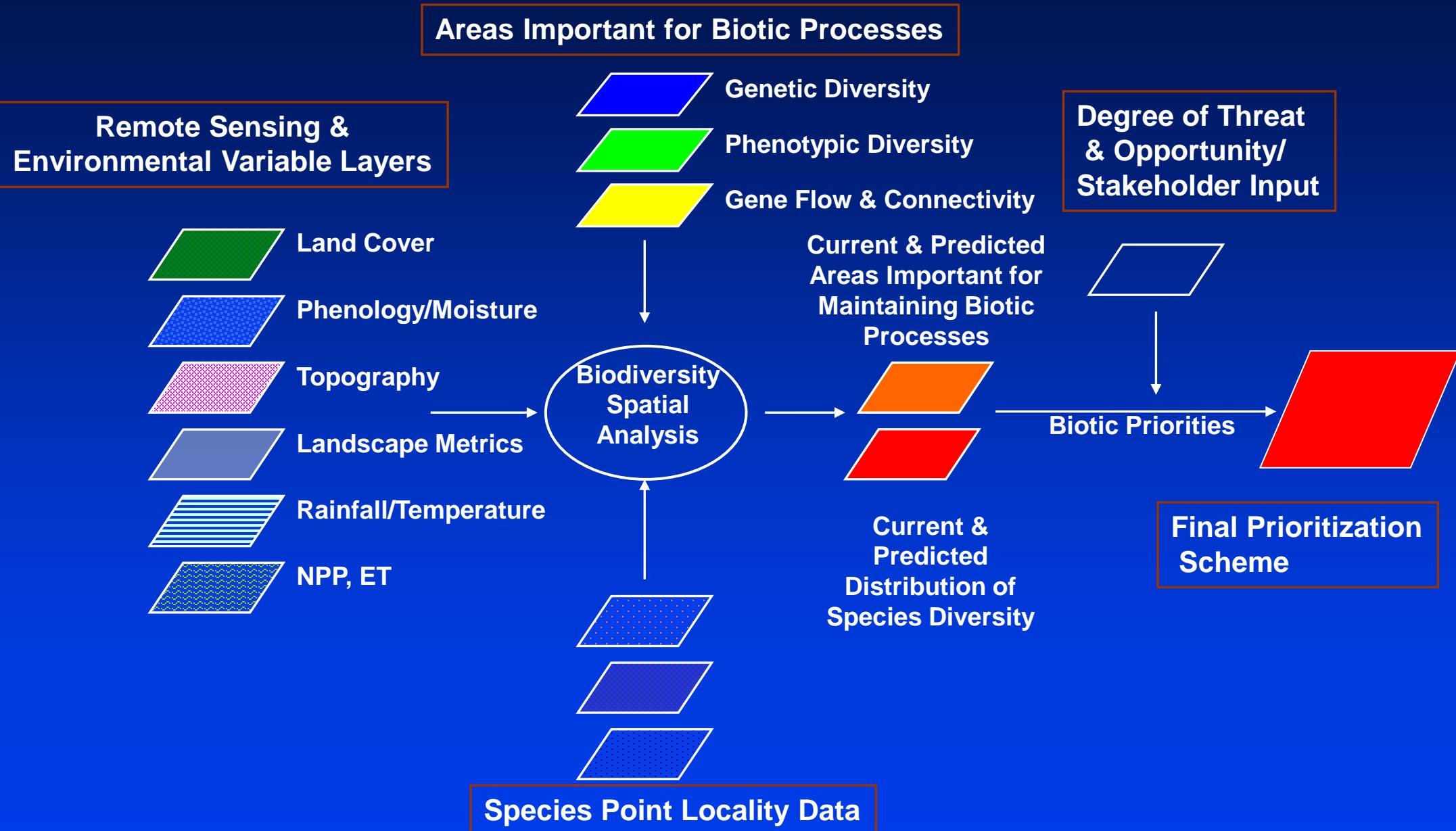
# Spatial Analysis Model



# Threats and opportunities and stakeholder input



# Spatial Analysis Model



# SUMMARY

- 1) Gradients can be important drivers of adaptive genetic variation
- 2) Conserving gradients may offer a bet hedging approach for conserving adaptive variation in the face of climate change
- 3) Integrating biodiversity pattern, process and degree of threat may offer a valuable approach for prioritizing regions for conservation

# Collaborators

Camille Bonneau  
Wolfgang Buermann  
Ryan Calsbeek  
Jaime Chavez  
Catherine Graham  
Alex Kirschel  
Borja Mila  
Ana Paula

John Pollinger  
Cassia Prates  
Jorge Velesquez  
Sassan Saatchi  
Chris Schneider  
Ravinder Sehgal  
Hans Slabbekoorn  
Bob Wayne

# Support

NASA  
NSF  
National Geographic  
Society

QuickTime™ and a  
TIFF (Uncompressed) decompressor  
are needed to see this picture.



BOSTON UNIVERSITY