

Survival, Growth, and Reproduction of red mangroves (*Rhizophora mangle* L.) in Restoration: Importance and Interaction of Genetic and Environmental factors

Edward Proffitt¹ and Steven E. Travis²

¹ Department of Biological Sciences,
Florida Atlantic University, c/o Harbor Branch
Oceanographic Institute at FAU, Ft. Pierce, FL

² Department of Biology,
University of New England, Biddeford, ME

Foundation Species

➤ Presence forms habitat structure and productivity base for the entire system

- Mangrove trees
- Salt marsh grasses
- Reef-building corals
- Seagrass

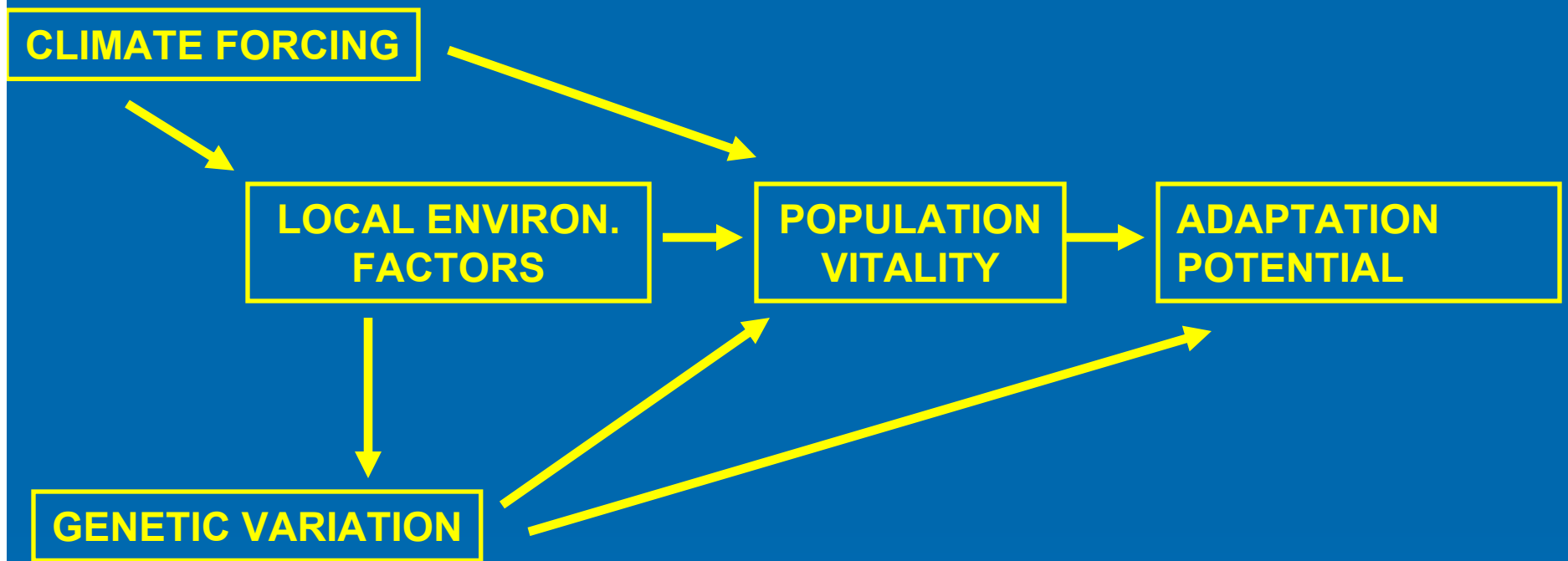
➤ Will interact

- Intermed. Latitudes
- Varying strength

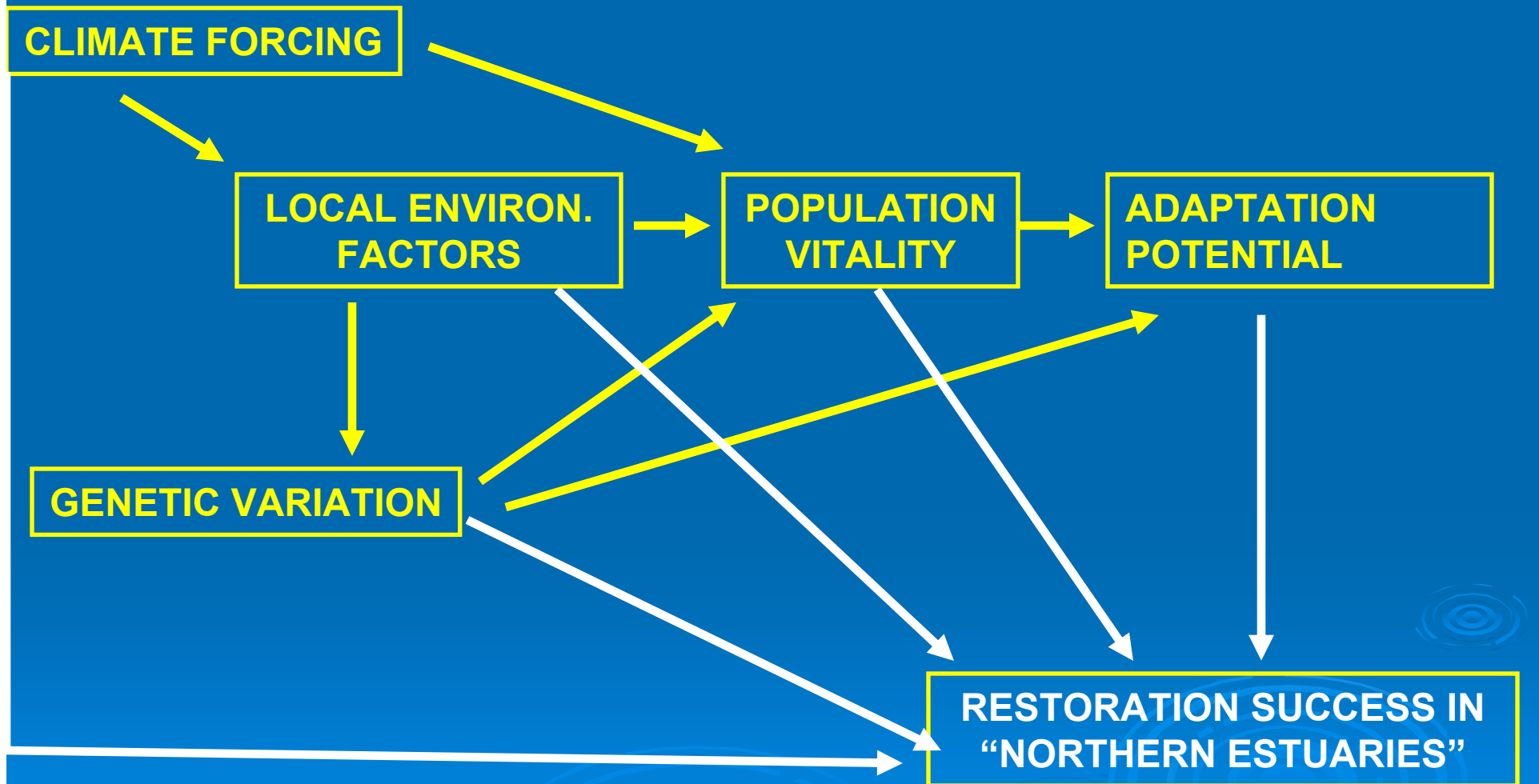
Mangrove “islands” in a salt marsh “matrix”
Merritt Island National Wildlife Refuge



HYPOTHESIZED PATH DIAGRAM: ADAPTATION TO CHANGING CLIMATE



HYPOTHESIZED PATH DIAGRAM: RESTORATION



Field Experiment:

Is there evidence for ecologically important genetic effects in red mangroves?

Address using:

- Full sibling seedling families from individual maternal trees
- Planted in a common garden experiment with an elevation stress gradient
- Evaluate with statistical models and visualize with Norms of Reaction plots

Specific Factors

- **Embayments within Tampa Bay.....5**
- **Donor trees from embayments.....86**
 - **Sibling seedlings from each donor.....40**
- **Islands Planted at Pt Redwing.....5**
- **Elevations w/in each Island.....2**
 - **Low: Less desiccation stress**
 - **High: More stress**

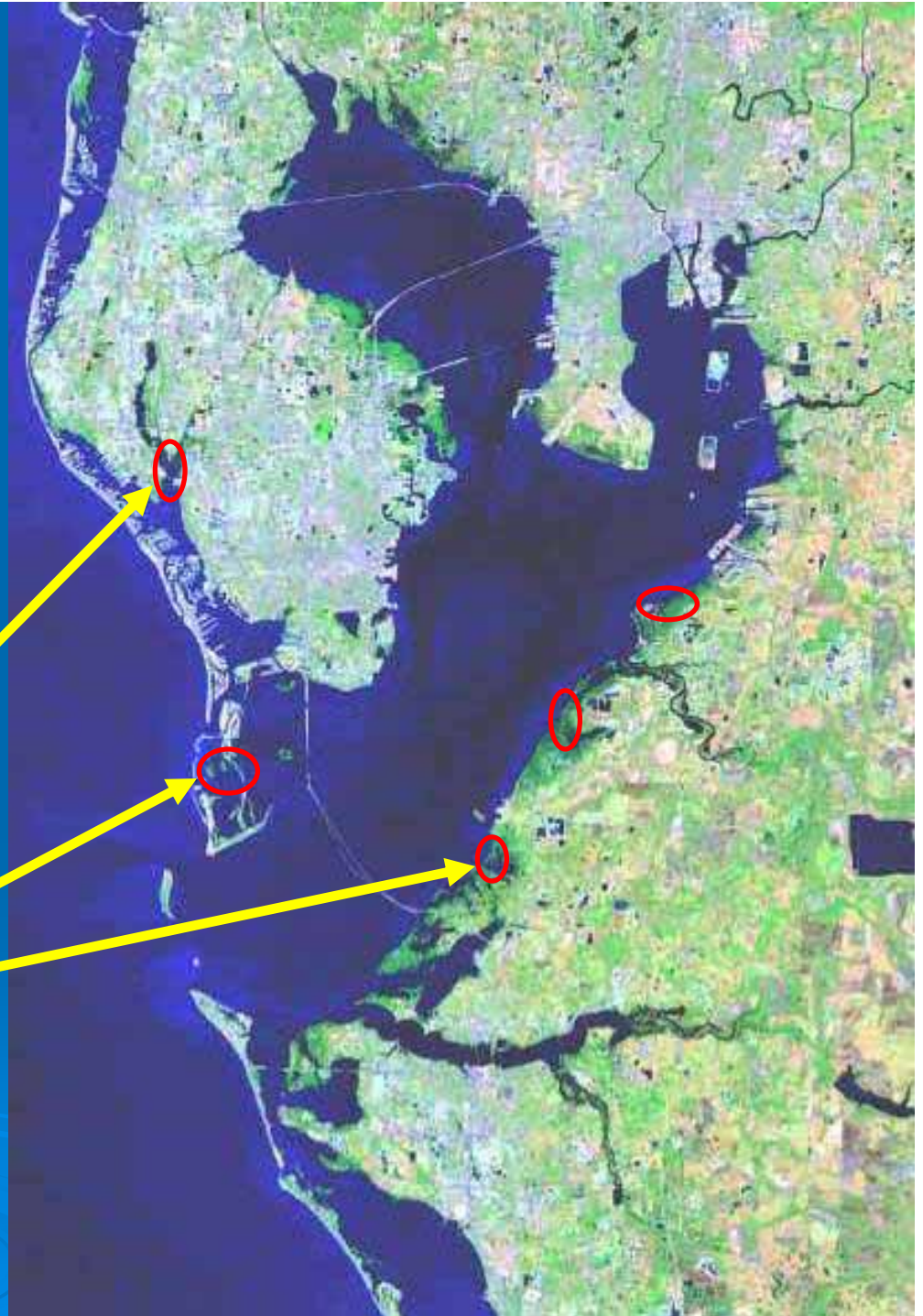


Why use Donor Trees from Different Embayments?

- Embaym. may differ in environmental growing conditions for the maternal trees (that might become maternal effects in the seedlings)

More urban area

Historically contaminated → high mutation rates





Embayments: ○ Regions of Tampa Bay where DONOR trees were located (ie, Maternal Families taken from)

PLANTING LOCATION: □ (5 Islands at Schultz Family Park [Port Redwing])

Maternal Families: Donor Trees (within embayment)

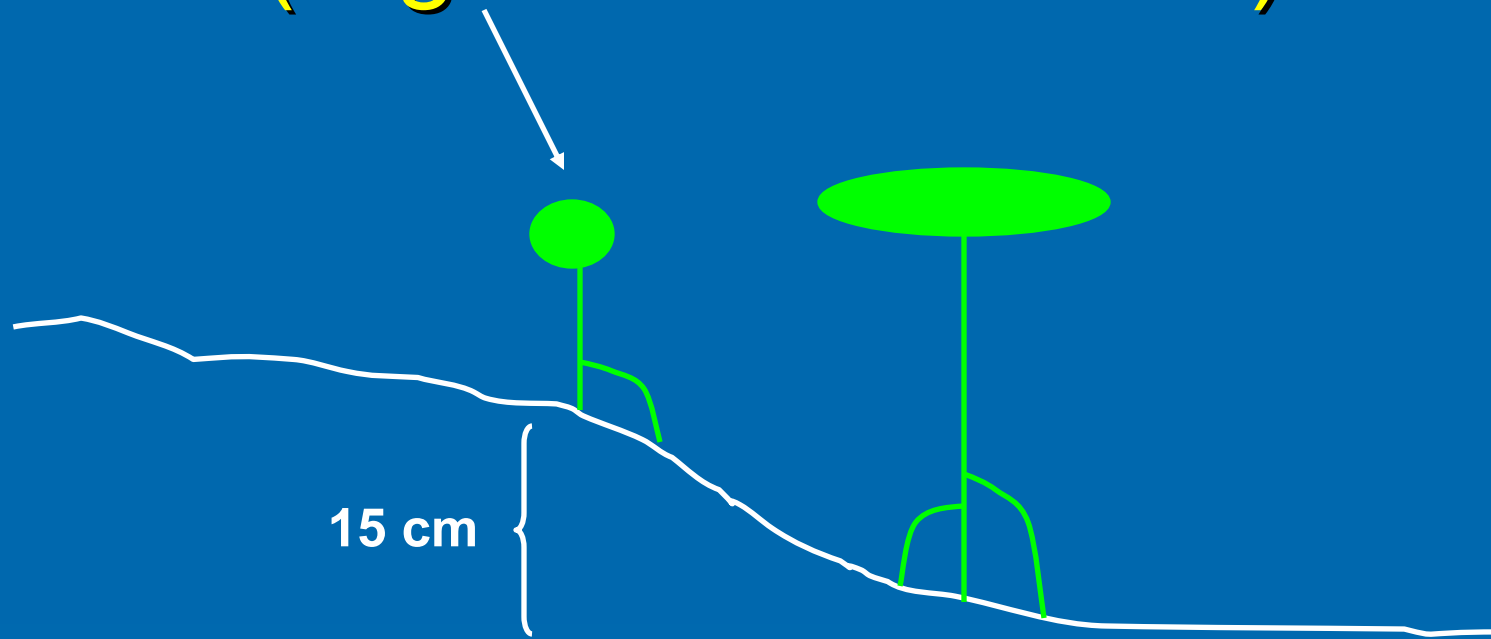
- Seedlings from a donor tree comprise a *sibling cohort* or *Maternal Family*
- Differences among Maternal Families reflects
 - Maternal tree environment that might be incorporated into propagules (e.g., nutrients; stressors like pollutants)
 - Maternal tree genotype

Elevation (**stress**) Effects

- **Planted HIGH & LOW transects on each island separated by 15 cm elevation gradient**



Hypothesized Elevation Effects (High=more stressful)

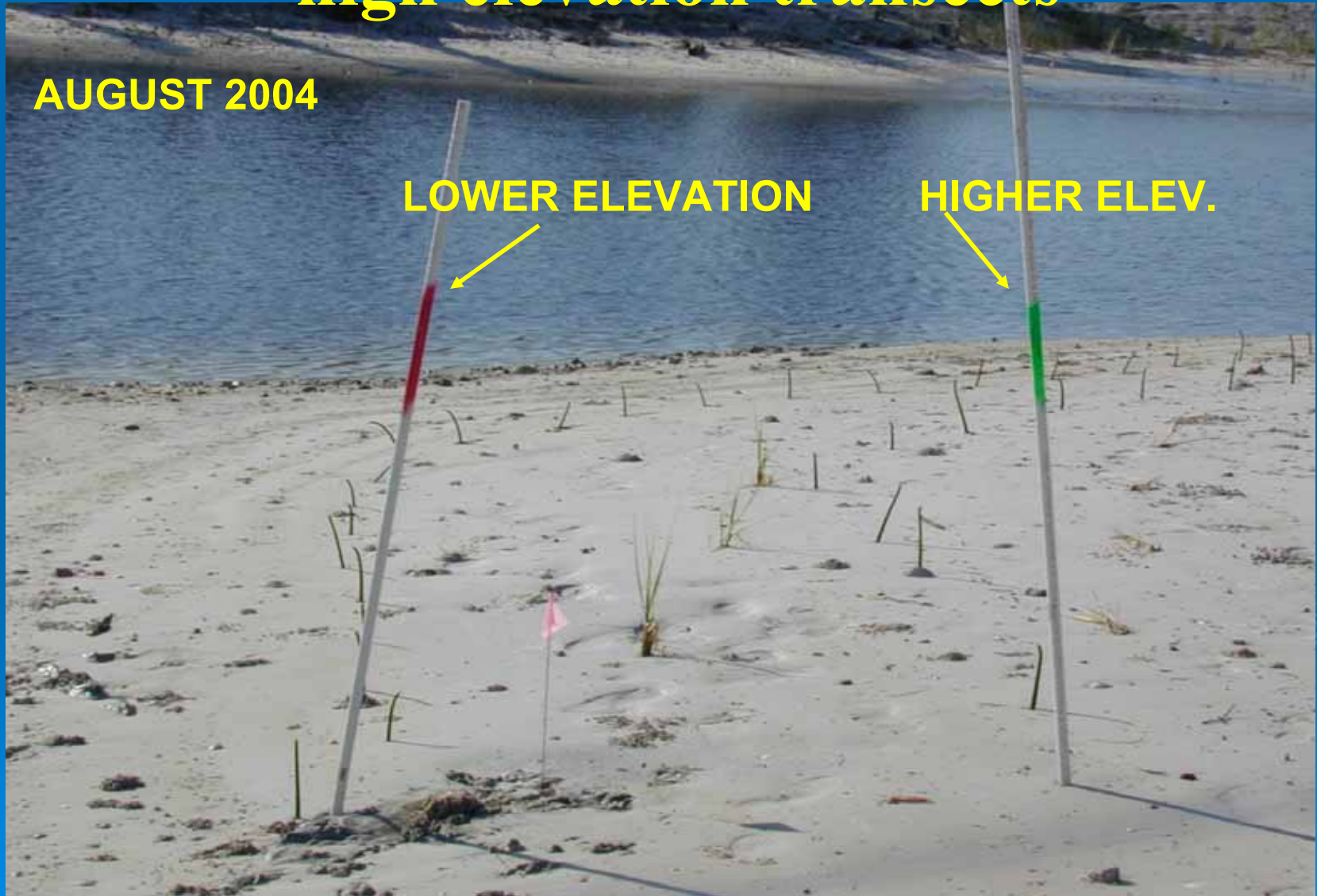


R. mangle experiment showing low & high elevation transects

AUGUST 2004

LOWER ELEVATION

HIGHER ELEV.



Planting 5 Different Islands

- Effect of seedling environment over a spatial scale of 10s of m



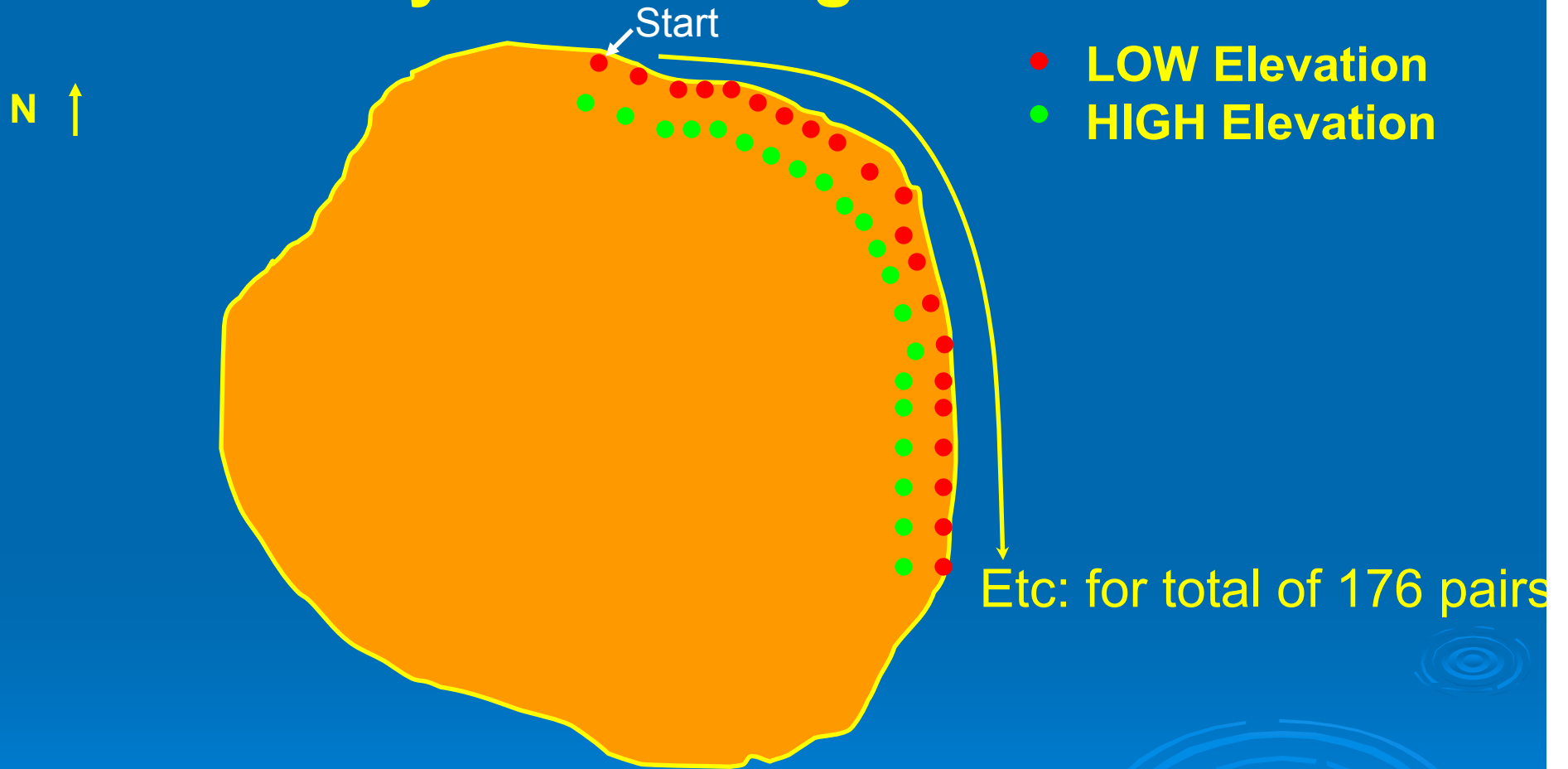
The Islands Planted

August 2006
Photo



Photo courtesy of B. Henningsen, SWFWMD

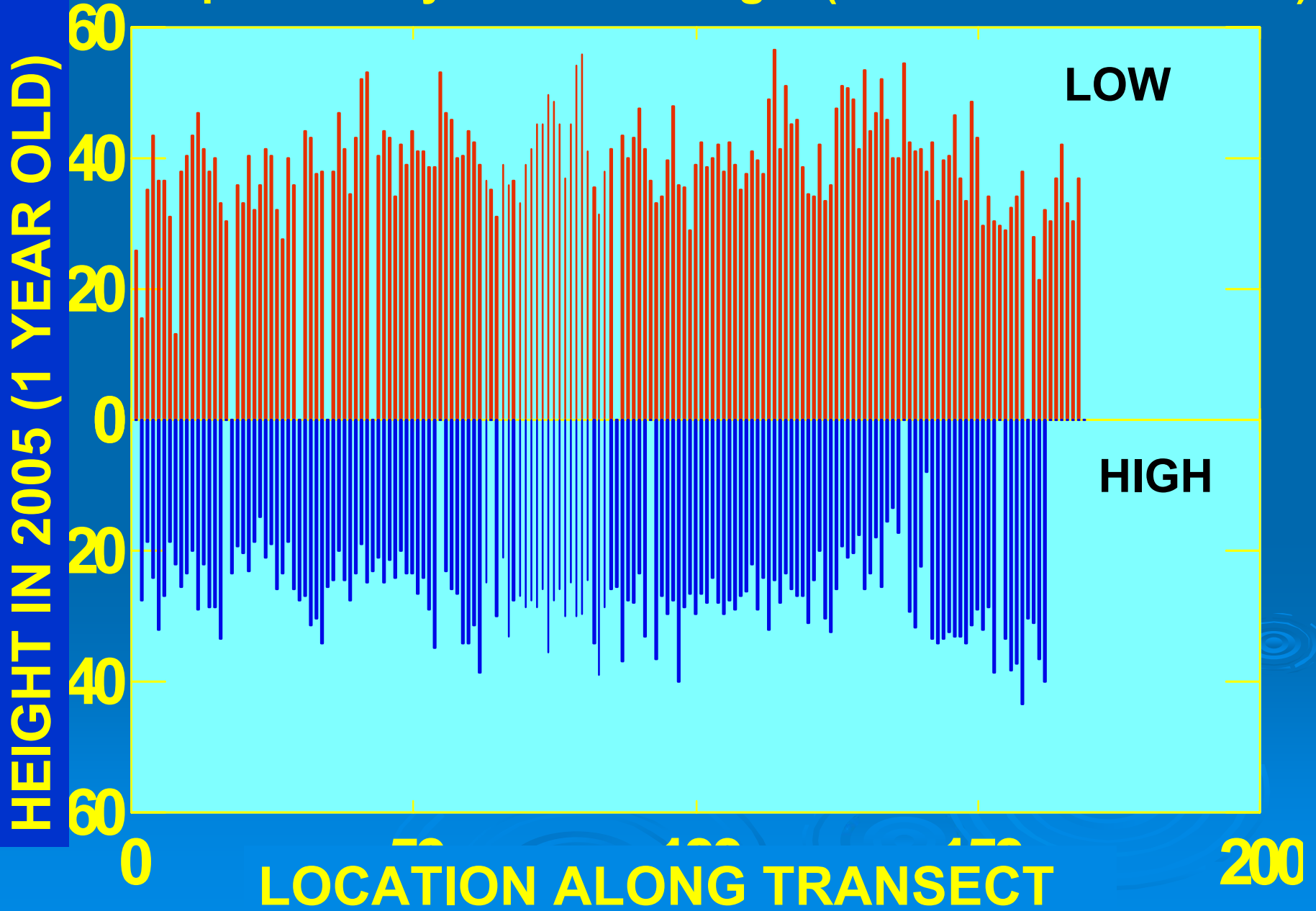
Smallest Spatial Scale: different planting units every 0.5 m along Elev. transects



Island 4: Illustration of the LOW & HIGH elevation transects

ISLAND 4: Heights along transects

- Notice peak/valley "runs" in height (local environ. effects)



Propagule Size

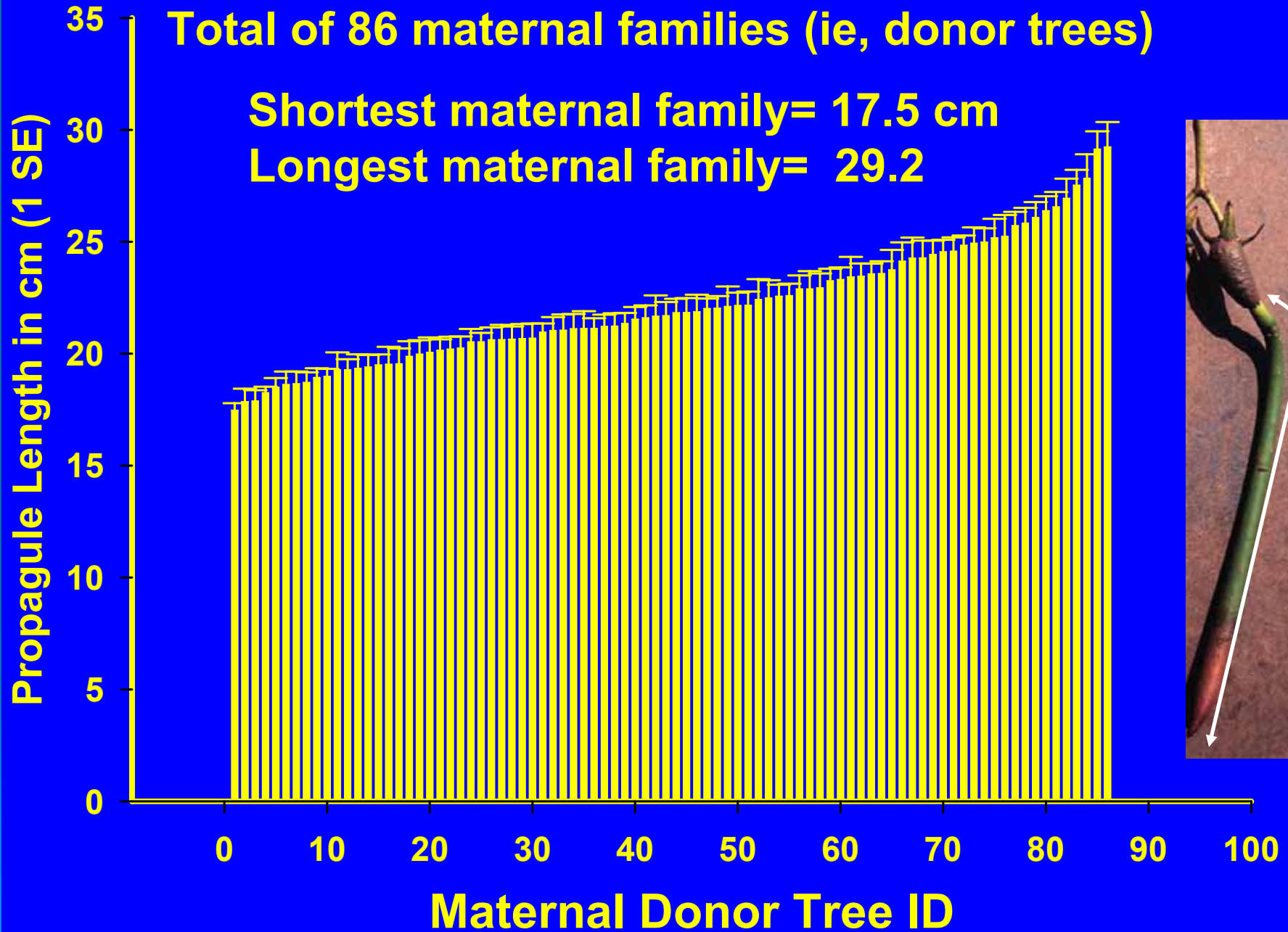
- **Covariate in analyses.**
- **May reflect degrees of maternal stores**

Rhizophora propagule length (mean & SE by donor tree)

Total of 86 maternal families (ie, donor trees)

Shortest maternal family= 17.5 cm

Longest maternal family= 29.2



Determining Genetic Effects (from maternal families)

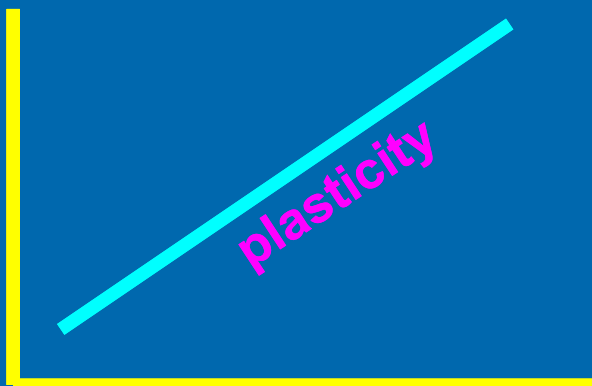
- **This Experiment: Norms of Reaction for different maternal families of seedlings over the elevation stress gradient**
- **Next Experiment: confirm using F_1 offspring from these seedlings**

How separate genetic, maternal, plastic, and environmental effects?

➤ Field Experiment:

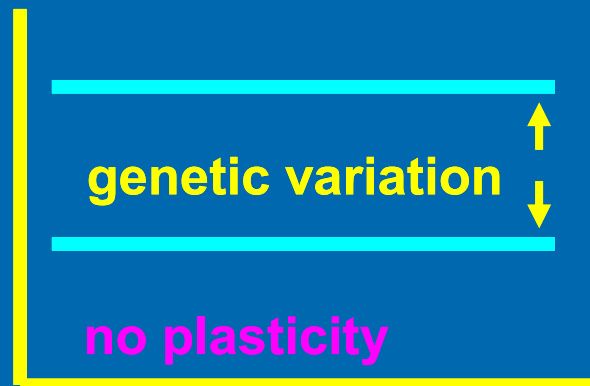
- Norms of reaction over an environmental gradient (Elevation) of a series of seedling sibling families (ie, from different donor trees)
- Assess presumed maternal effects by:
 - Use propagule size as a covariate
- Determine local environmental effects –
 - Diff's among maternal tree environments (embayments within Tampa Bay)
 - Diff's. among islands planted
 - Diff's along transects on each island planted

Phenotype



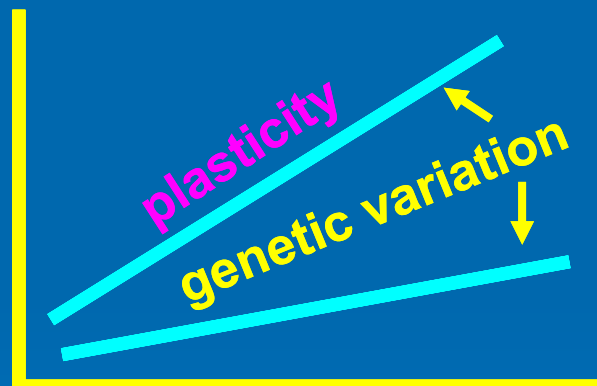
Environment

Phenotype



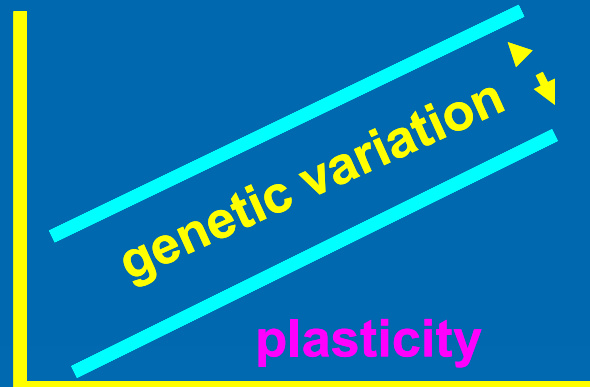
Environment

Phenotype



Environment

Phenotype



Environment

NORMS OF REACTION

RESULTS



Experimental Design: Reminder



Statistical Effects (fixed)

- **ELEVATION** : (2 Levels)
- **ISLAND PLANTED**: (5 Levels)
- Maternal Tree Home Embayment or **EMBAYMENT**: (5 Levels)
- **MATERNAL FAMILY**: (86 Levels)

PROPAGULE LENGTH: Covariate

Total Propagules Planted = 1,685

Response Variables

- Survival
- Plant Height (annually for 3 years)
- Trunk Diameter (annual for 3 years)
- Annual Incremental *Growth* in Height
- Number of Stems
- Canopy Area (based on diameters of major & minor axes of “ellipse”)
- Ratio of Sq. Rt(Canopy Area) : Height
- Reproductive output (propagules produced at year 3)

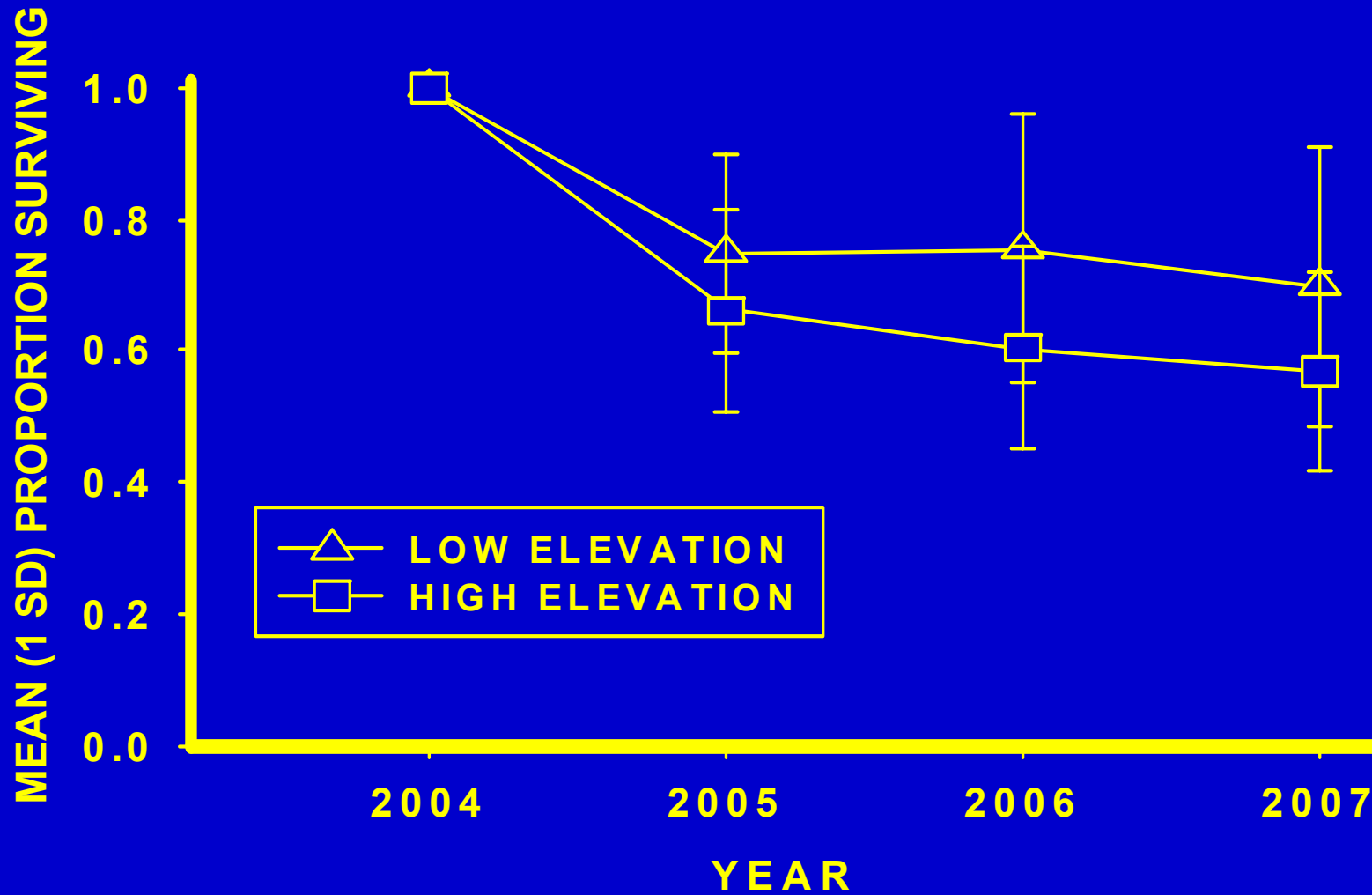
Main Research Hypotheses

- Does plant performance differ with *seedling maternal family* ?
- Do seedlings from a maternal family respond differently to *Low & High elevation*?

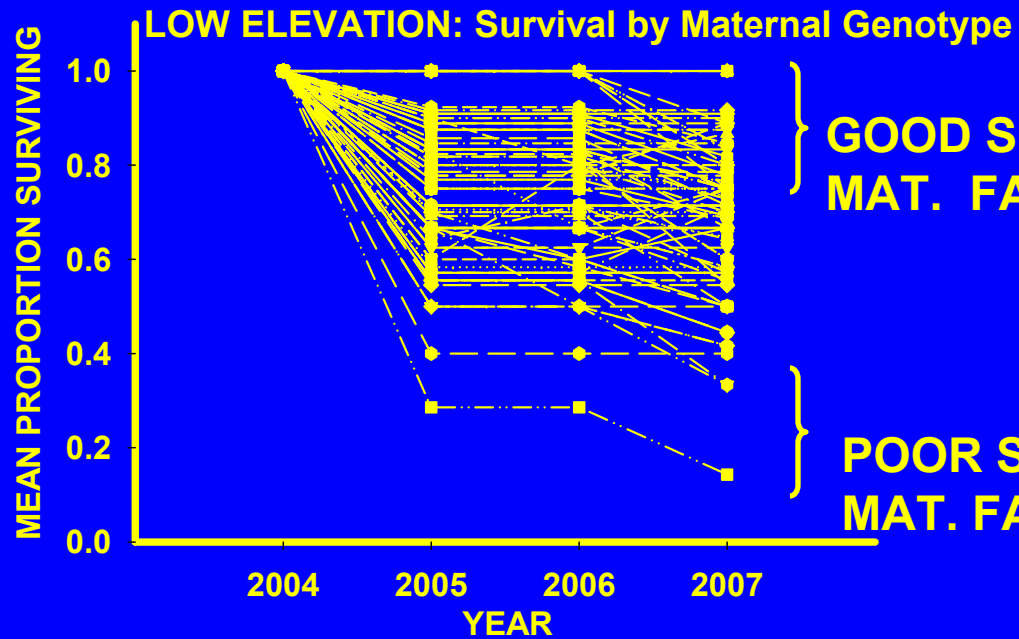
SEEDLING SURVIVAL



SURVIVAL AT LOW & HIGH ELEVATIONS

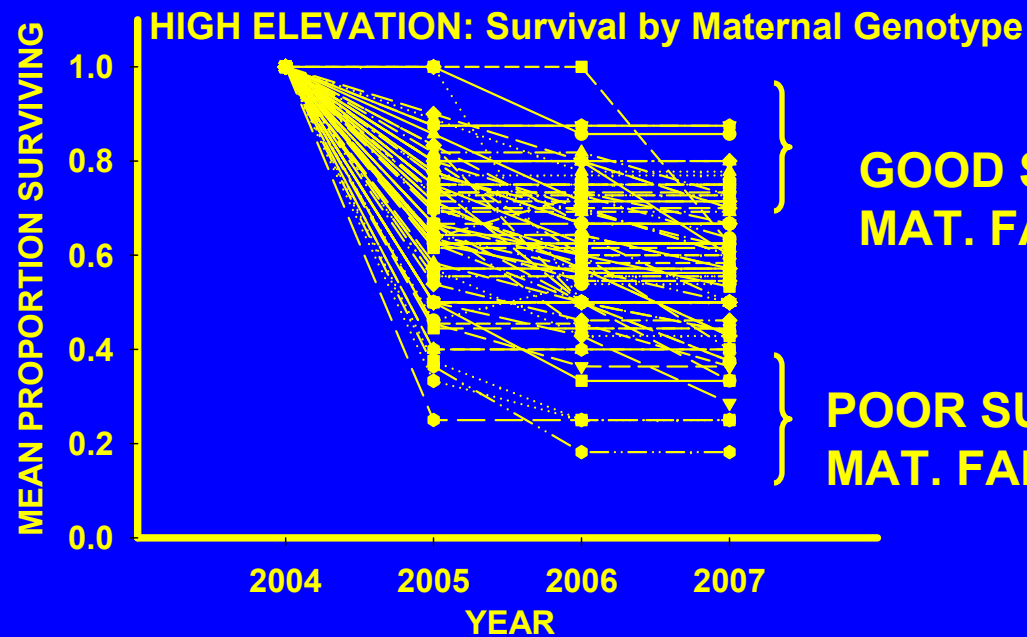


SURVIVAL BY MATERNAL FAMILY



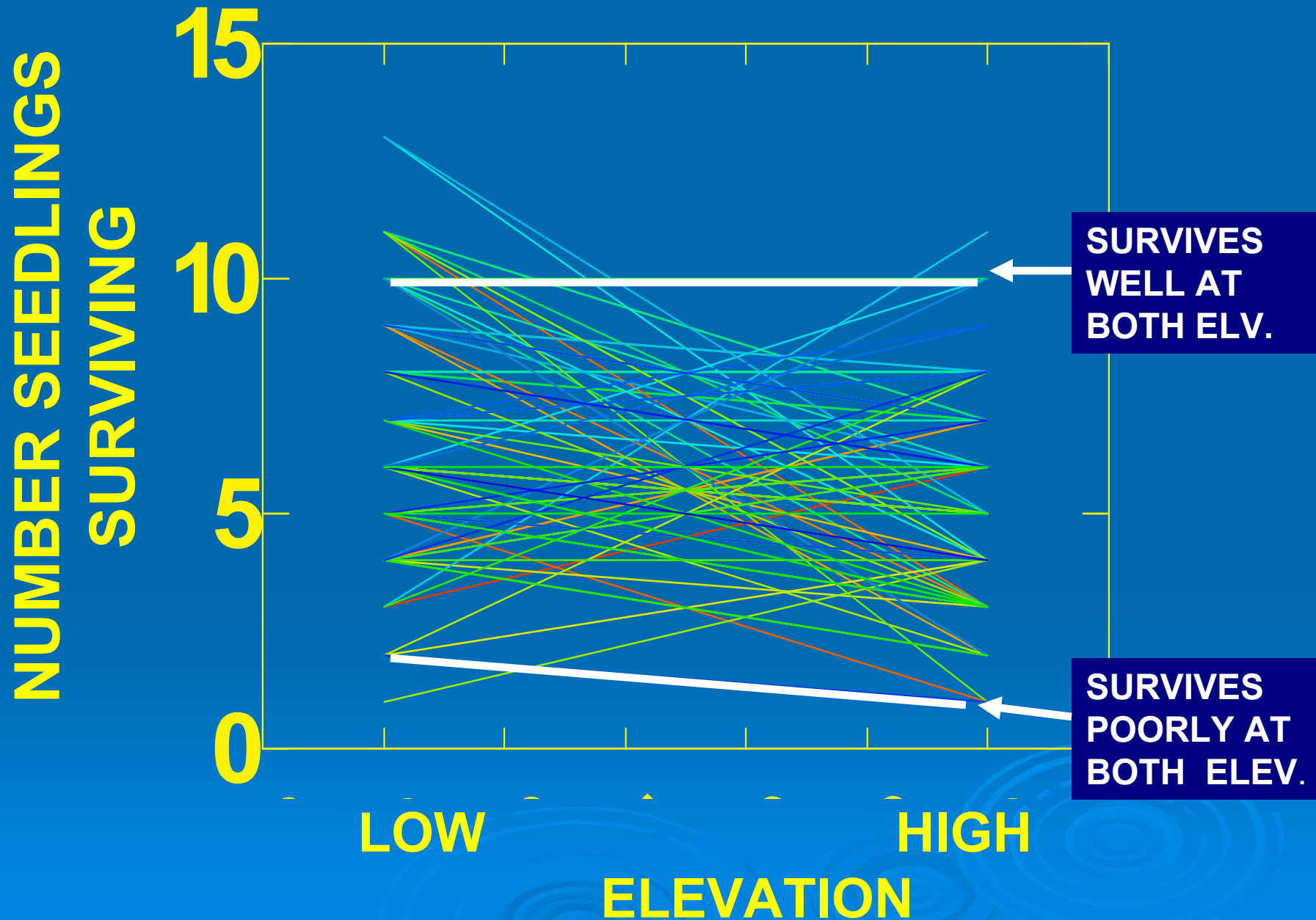
SURVIVAL

LOW ELEV.

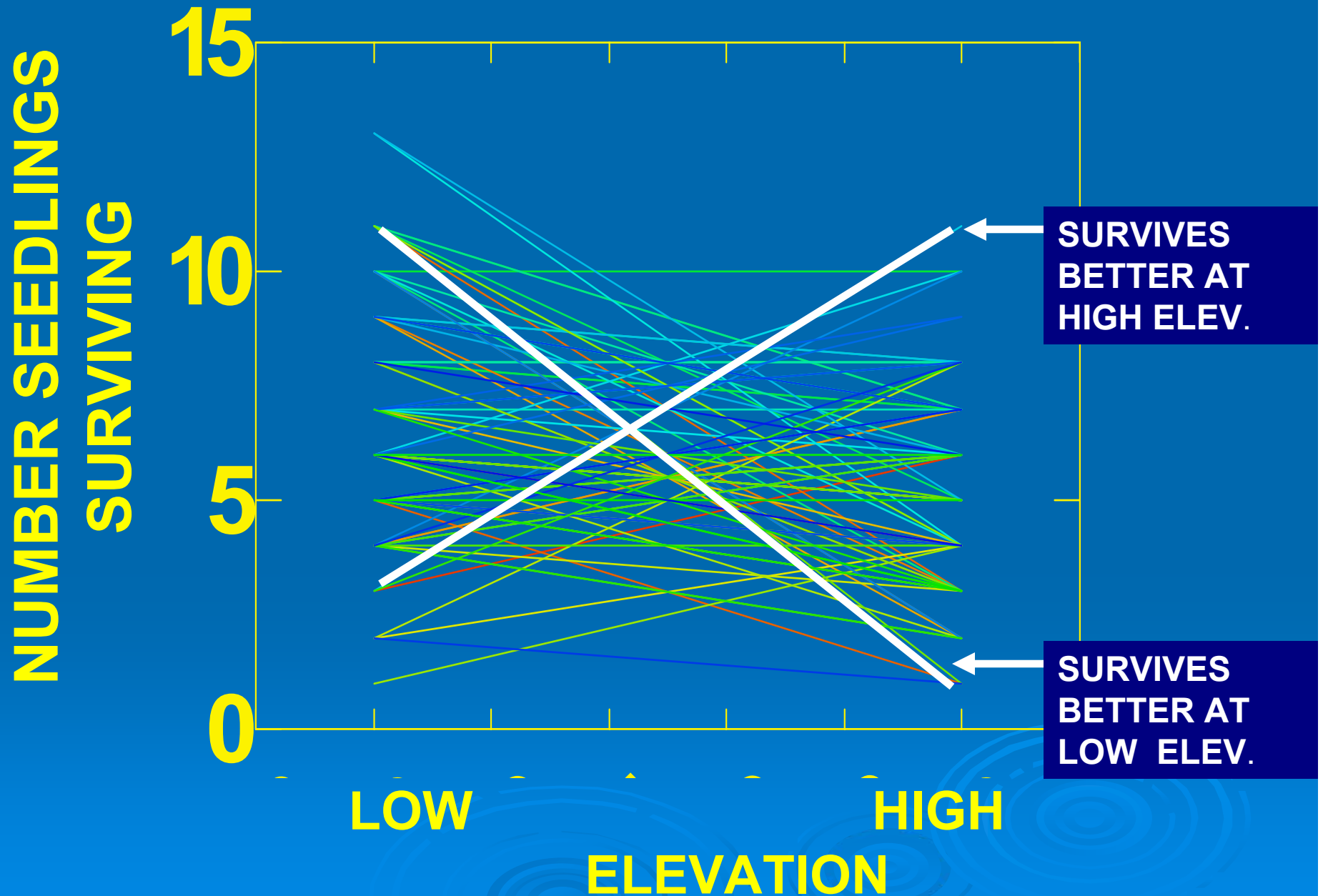


HIGH ELEV.

NORMS OF REACTION: 86 FAMILIES (3 YR)



NORMS OF REACTION: 86 FAMILIES (3 YR)



SUMMARY: NORMS OF REACTION

- Plasticity exists within maternal families (with environmental stress)
 - Independent of propagule size
- Genetic differences among maternal families
 - Assumption that aspects of survival are heritable

Logit modeling Results : Survival at 3 years

- Response Variable (modeled alive = 1, dead = 0),
 $p < 0.0005$, McFadden's Rho Squared = 0.29,
- Significant Explanatory Variables were (best model selected by AIC):
 - Elevation – Odds of surviving 3.1x greater at LOW Elv. When maternal family not considered
 - Maternal family – Odds ratios ranged from
 - 3.2 : 1 (greater survival at LOW elv. For one seedling family)
 - 0.06:1 (greater surv. At HIGH elv. For another seedling family)
 - Island Planted (no sig. effect of “local” seedling environment)
 - Propagule size (no sig. effect)

Growth in Height



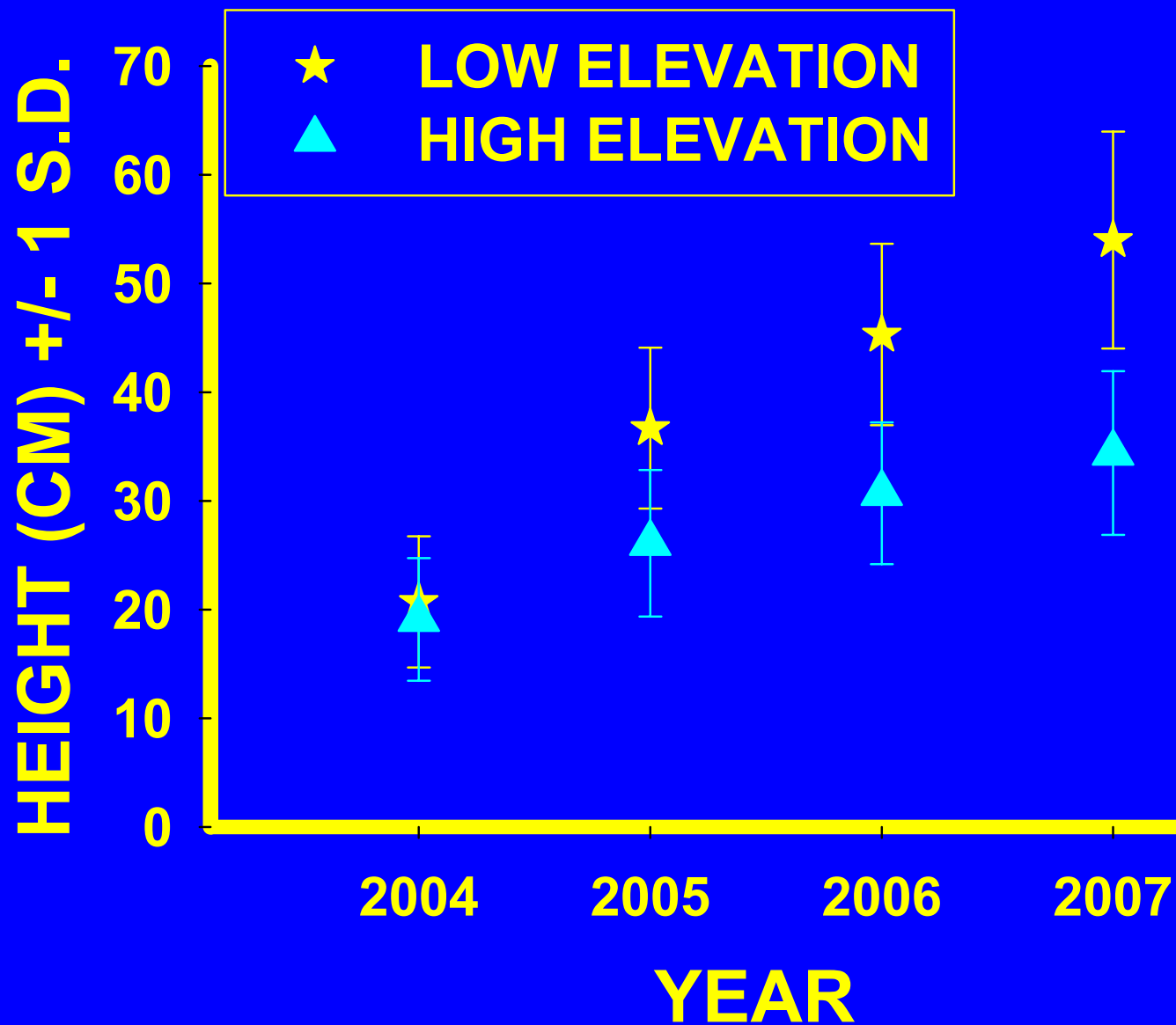
HEIGHT (2005): 1 year



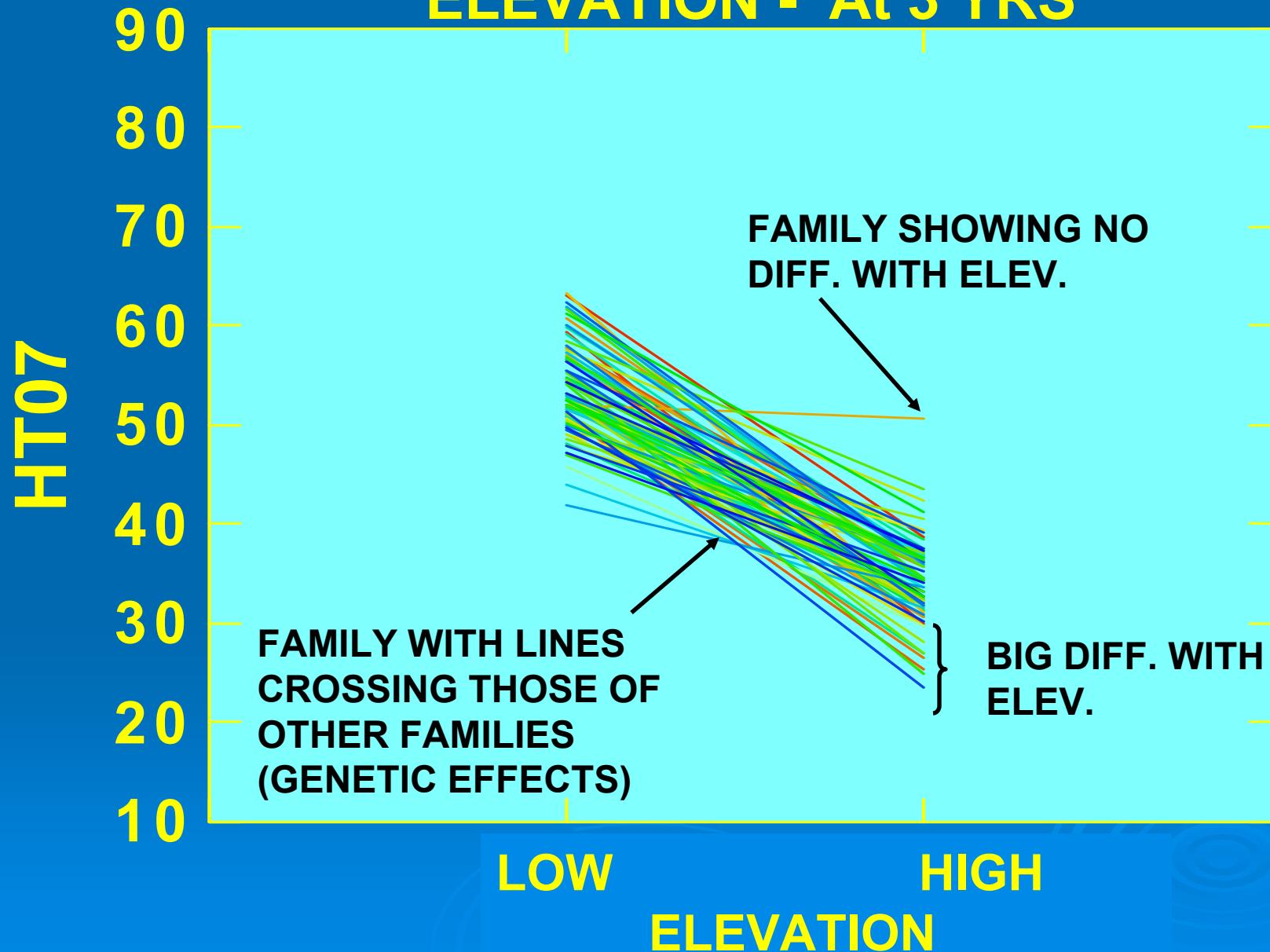
2007 (3 yrs old)



GROWTH IN HEIGHT SINCE EXPERIMENT INITIATION IN 2004



HEIGHTS BY MATERNAL FAMILY & ELEVATION - At 3 YRS



HEIGHT (Repeated-Measures ANOVA over 3 years)

Repeated Measures Analysis of Variance Between-Subjects Effects

| Source | DF | Type III SS | Mean Square | F Value | Pr > F |
|---------------|-----|-------------|-------------|---------|--------|
| Embayment | 4 | 103.3598 | 25.8399 | 0.23 | 0.9236 |
| Elevation | 1 | 169264.1010 | 169264.1010 | 1483.66 | <.0001 |
| Island Plntd | 4 | 17010.2821 | 4252.5705 | 37.28 | <.0001 |
| MatFam(Embay) | 81 | 14224.8712 | 175.6157 | 1.54 | 0.0022 |
| Propag. Len. | 1 | 6890.1448 | 6890.1448 | 60.39 | <.0001 |
| Error | 951 | 108495.1138 | 114.0853 | | |

All significant EXCEPT environmental influences among *embayments-within-Tampa Bay* (large spatial scale; locations of maternal trees)

Height con't: effects over time

Within subjects over time (**Univariate effects**)

| Source | DF | MS | F Value | Pr > F | G - G |
|-------------------|------|------------|---------|--------|--------|
| height | 2 | 1225.22 | 73.33 | <.0001 | <.0001 |
| height*Embaym | 8 | 19.85263 | 1.19 | 0.3022 | 0.3096 |
| height*Elev. | 2 | 5127.79451 | 306.89 | <.0001 | <.0001 |
| height*ISLAND | 8 | 70.26781 | 4.21 | <.0001 | 0.0003 |
| ht*MatFam(Emby) | 162 | 20.00410 | 1.20 | 0.0519 | 0.0773 |
| height*prop Leng. | 2 | 162.39419 | 9.72 | <.0001 | 0.0003 |
| Error(height) | 1902 | 16.70889 | | | |

The effects on height of:

elevation
island planted
propagule length

} *Changed over time*

The effects on height of:

Maternal family

} *Constant over time*

Height: Independent Variables Effect Sizes

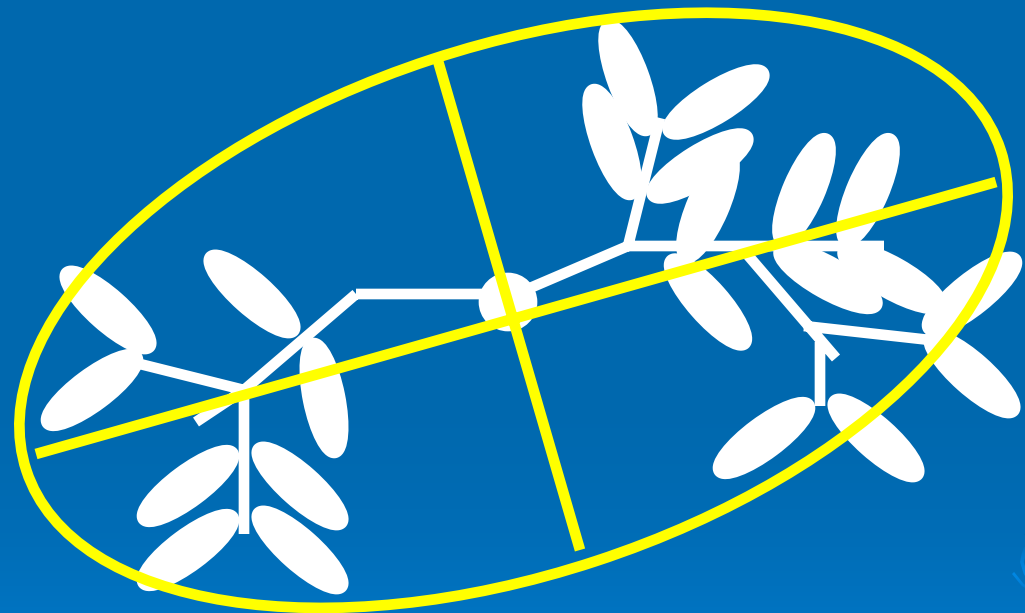
- Calculate at year 3 (cumulative effect over 3 years)

HEIGHT AT 3 YRS (2007) ANOVA & EFFECT SIZES

| <i>SOURCE</i> | <i>SIG.</i> | <i>EFFECT SIZE</i> |
|---------------------------|-------------|--------------------|
| MATERNAL TREE EMBAYMENT | NS | |
| ELEVATION (LOW or HIGH) | 0.0005 | 9.94 |
| PLANTING ISLAND (1...5) | 0.0005 | +3.99 TO -3.47 |
| MATERNAL FAMILY | 0.011 | +6.40 TO -7.14 |
| PROPAGULE LENGTH (COVAR.) | 0.0005 | 0.64 |
| ELEV. X ISLAND | 0.0005 | +3.46 TO -1.99 |

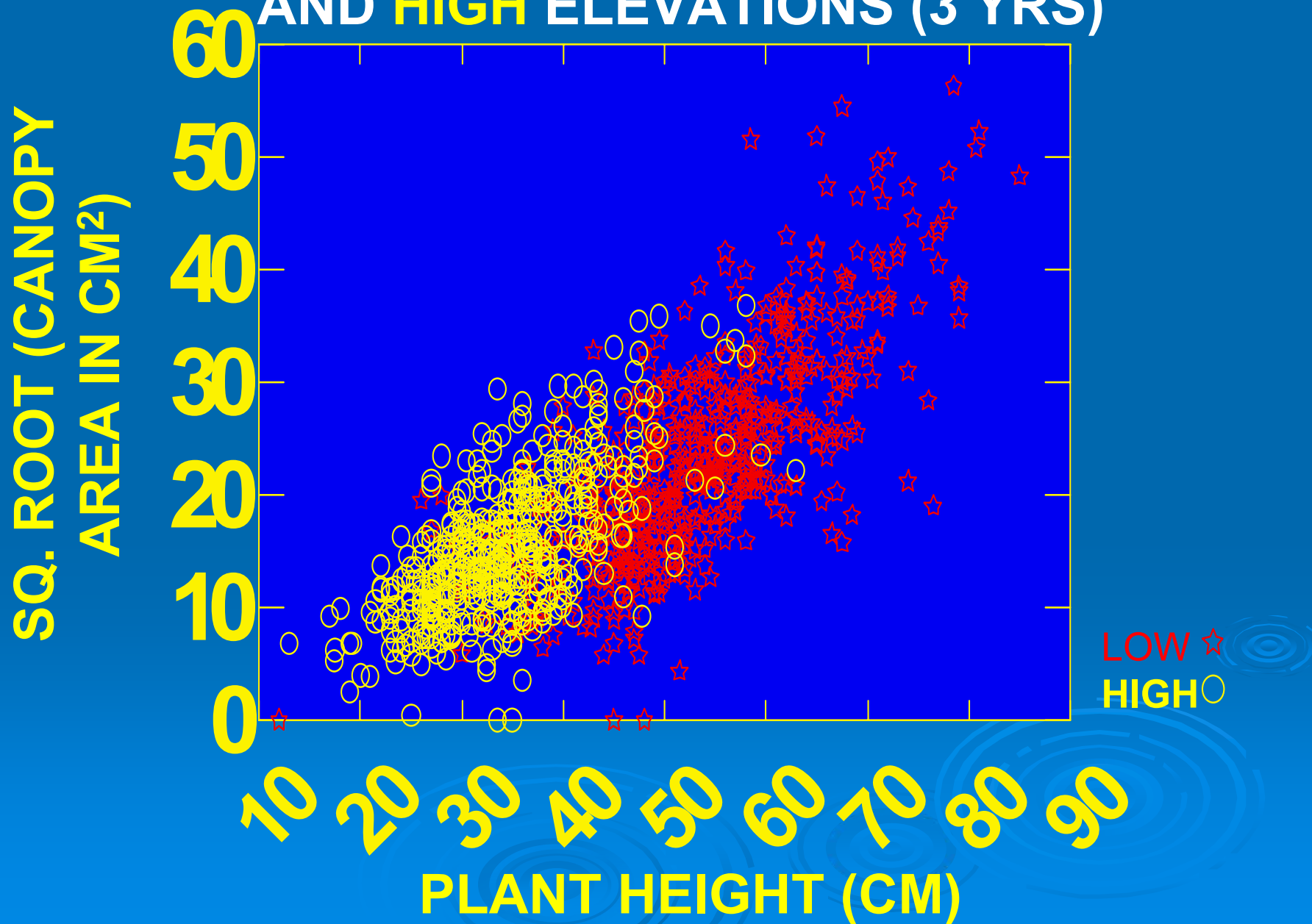
CANOPY AREA (AT 3 YEARS)

ILLUSTRATION FROM TOP VIEW



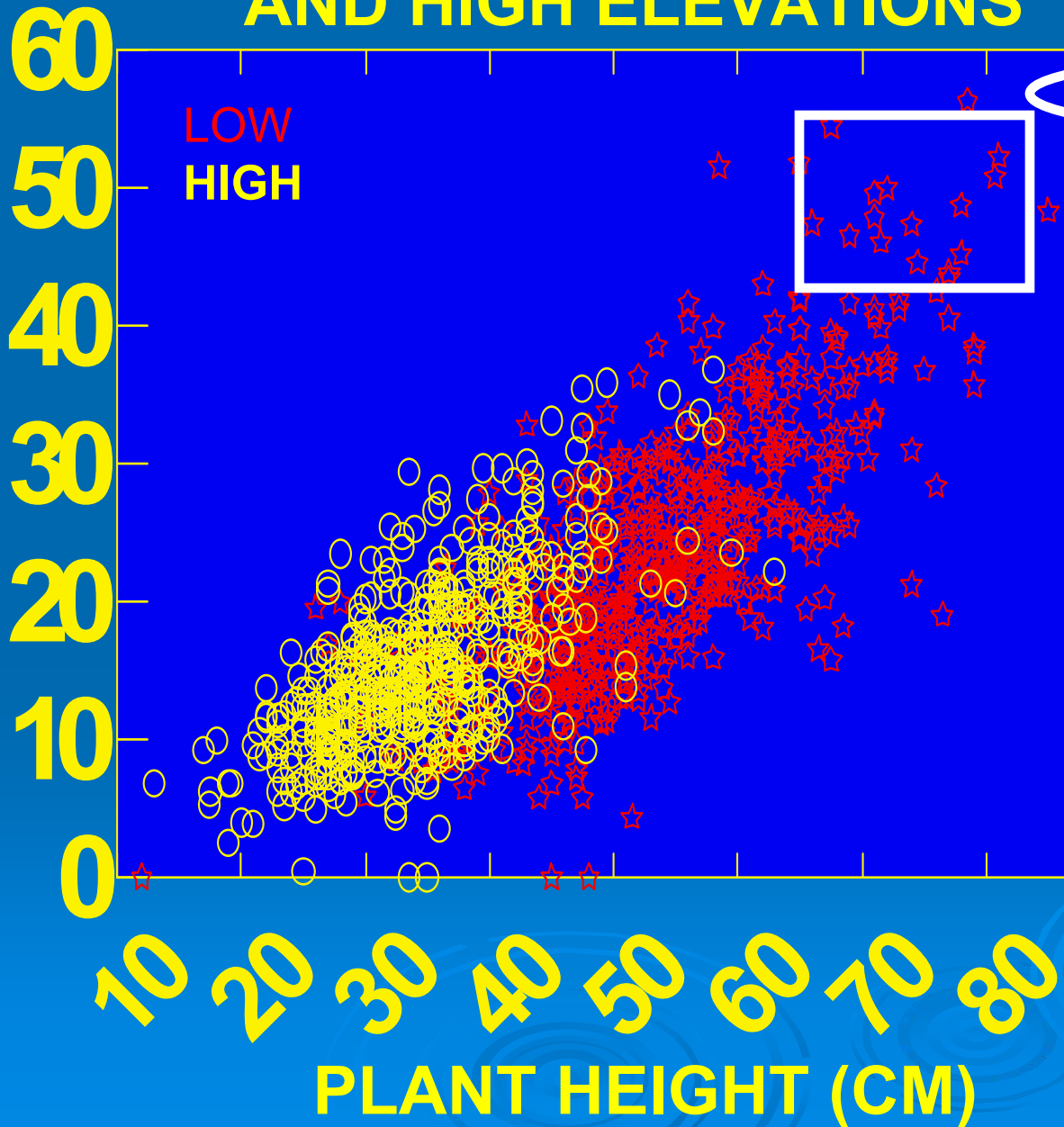
- MEASURE LENGTH OF MAJOR & MINOR AXES
- ESTIMATE AREA FROM EQUATION FOR ELIPSE

CANOPY AREA VS HEIGHT AT **LOW** AND **HIGH** ELEVATIONS (3 YRS)



CANOPY AREA VS HEIGHT AT LOW AND HIGH ELEVATIONS

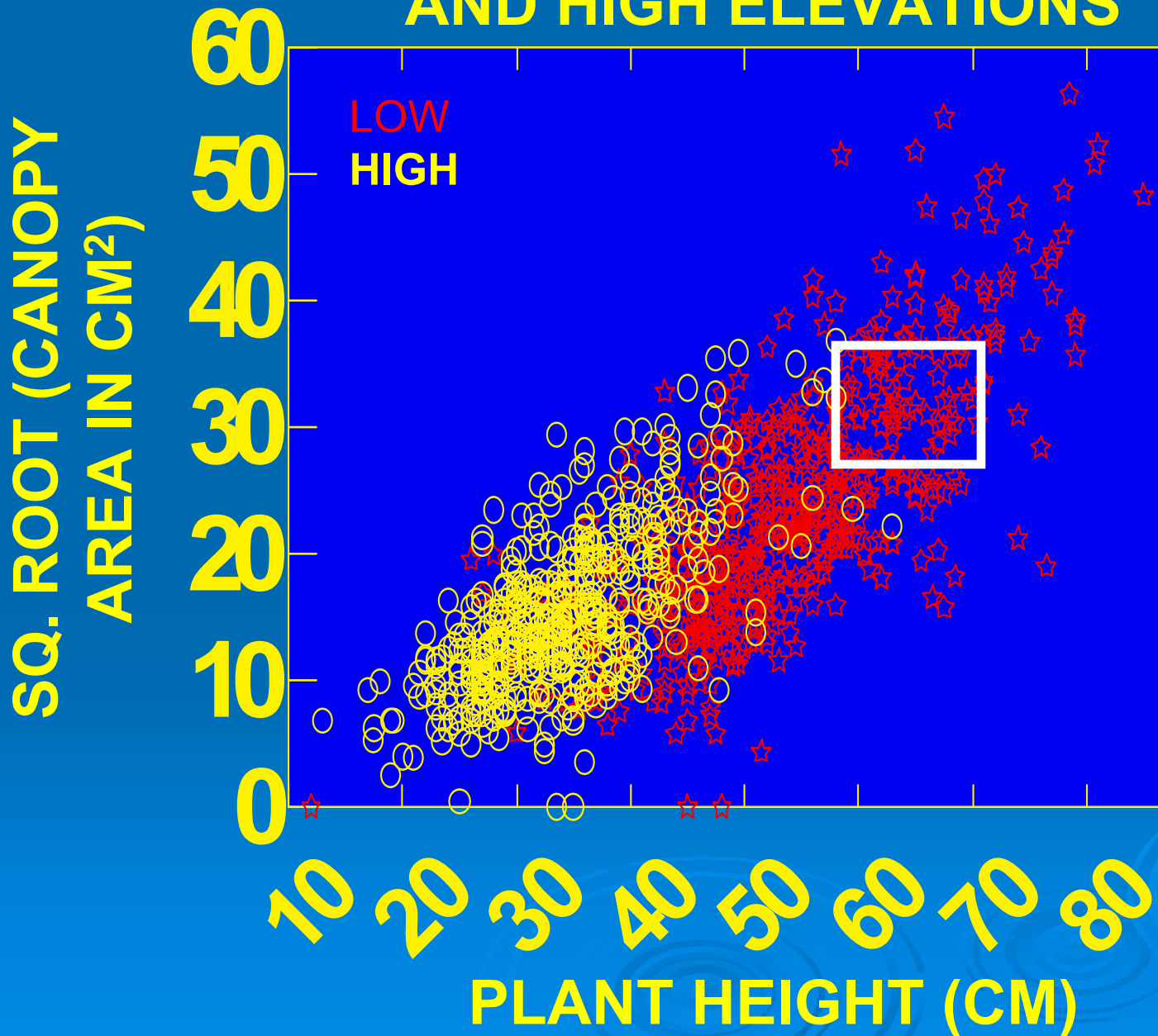
SQ. ROOT (CANOPY AREA IN CM²)



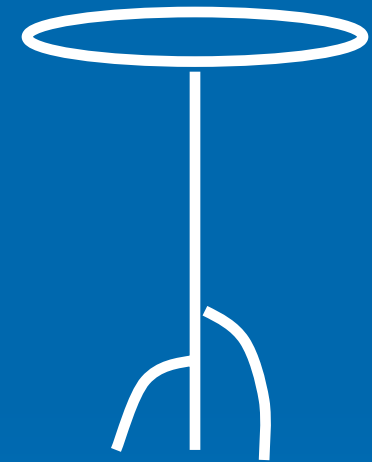
LOW ELV.



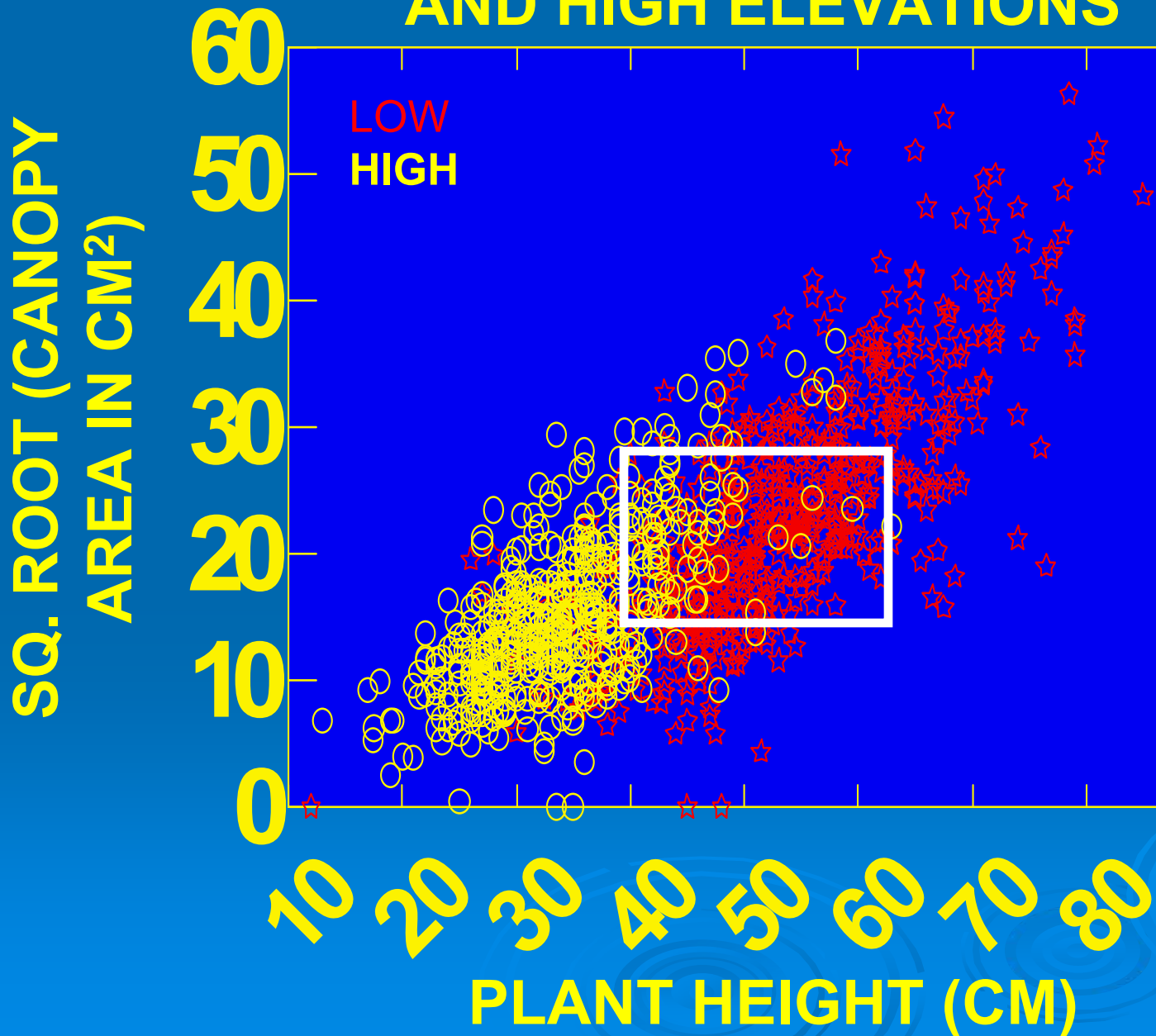
CANOPY AREA VS HEIGHT AT LOW AND HIGH ELEVATIONS



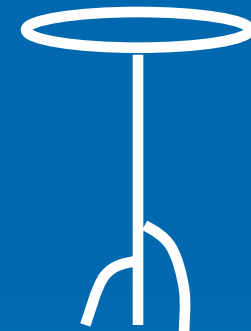
LOW ELV.



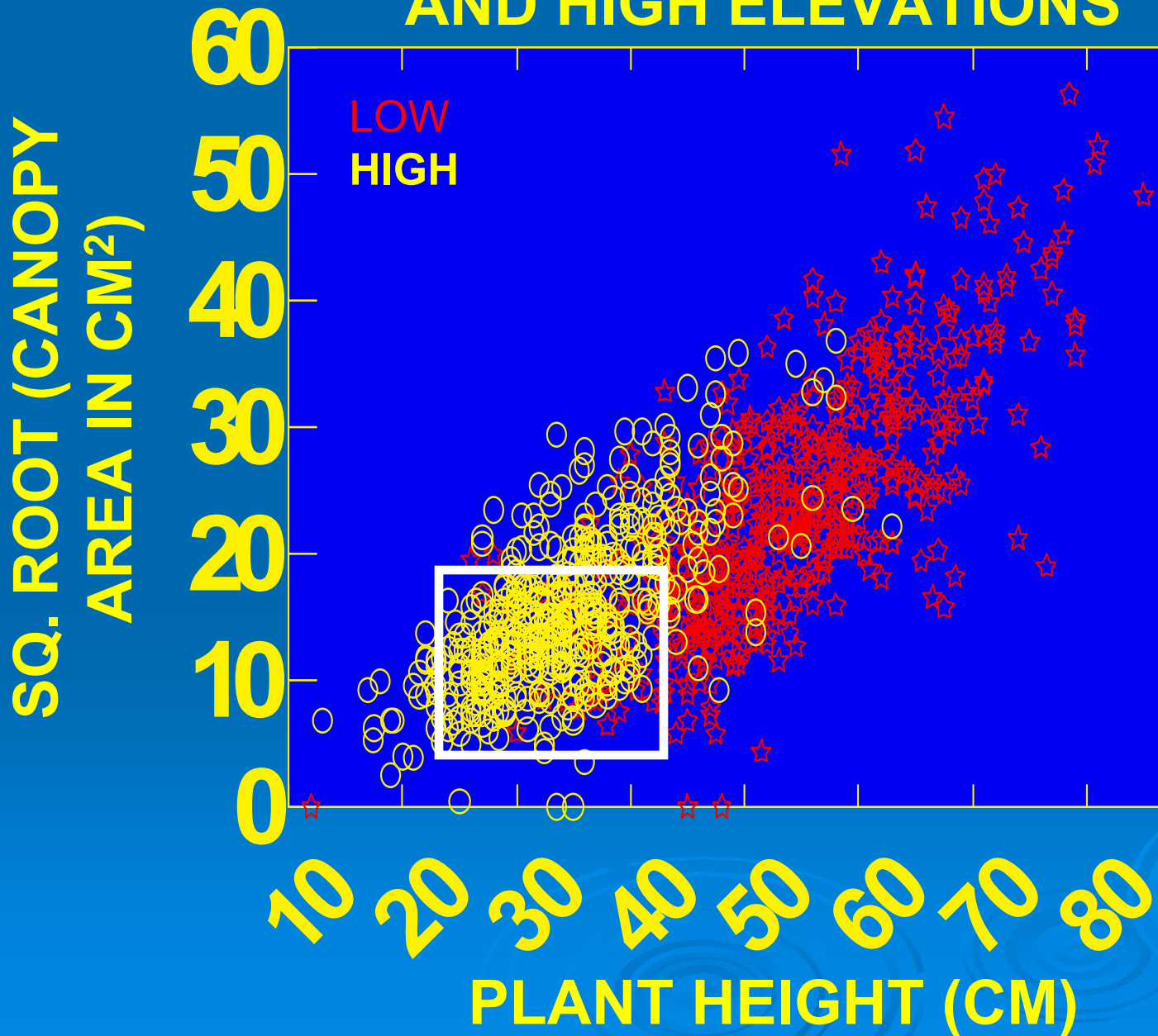
CANOPY AREA VS HEIGHT AT LOW AND HIGH ELEVATIONS



LOW & HIGH ELV.



CANOPY AREA VS HEIGHT AT LOW AND HIGH ELEVATIONS



CANOPY AREA VS HEIGHT AT LOW AND HIGH ELEVATIONS (3 YRS)

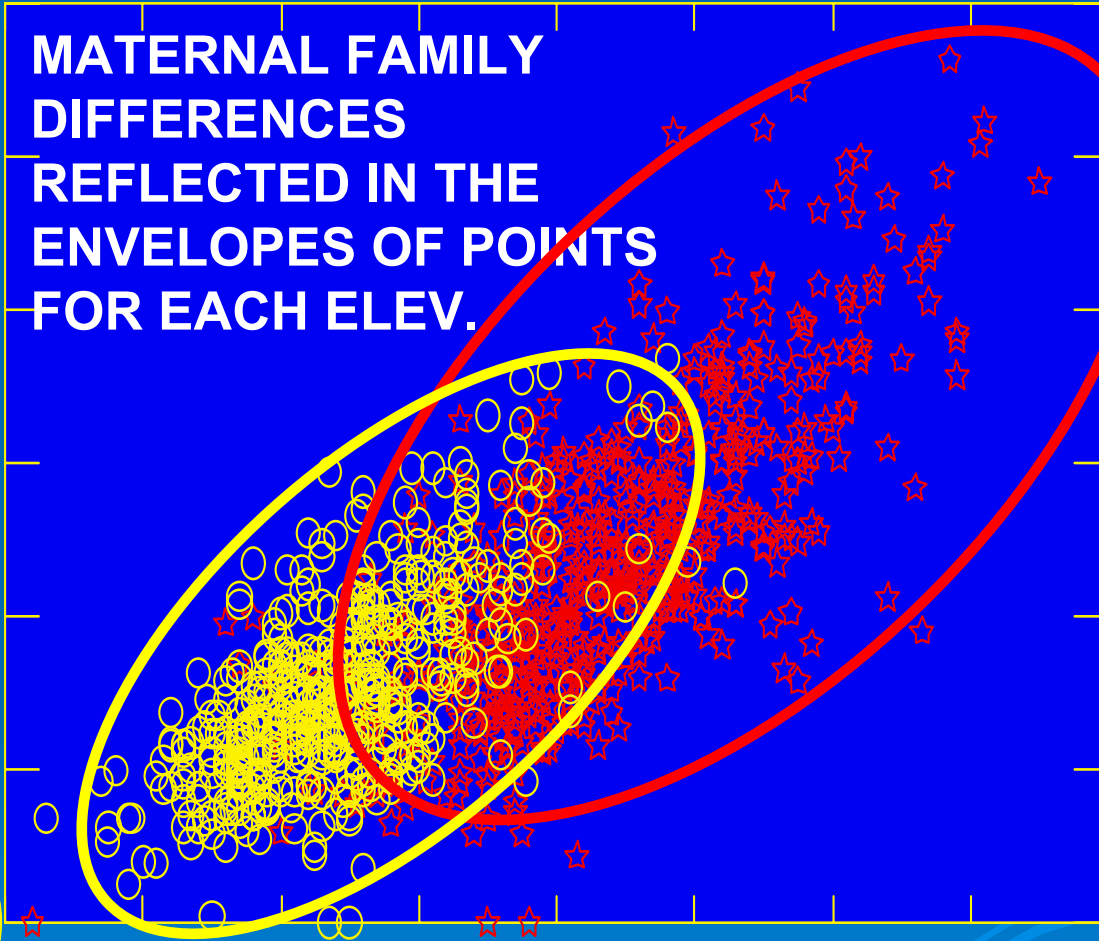
SQ. ROOT (CANOPY AREA IN CM²)

60
50
40
30
20
10
0

MATERNAL FAMILY DIFFERENCES REFLECTED IN THE ENVELOPES OF POINTS FOR EACH ELEV.

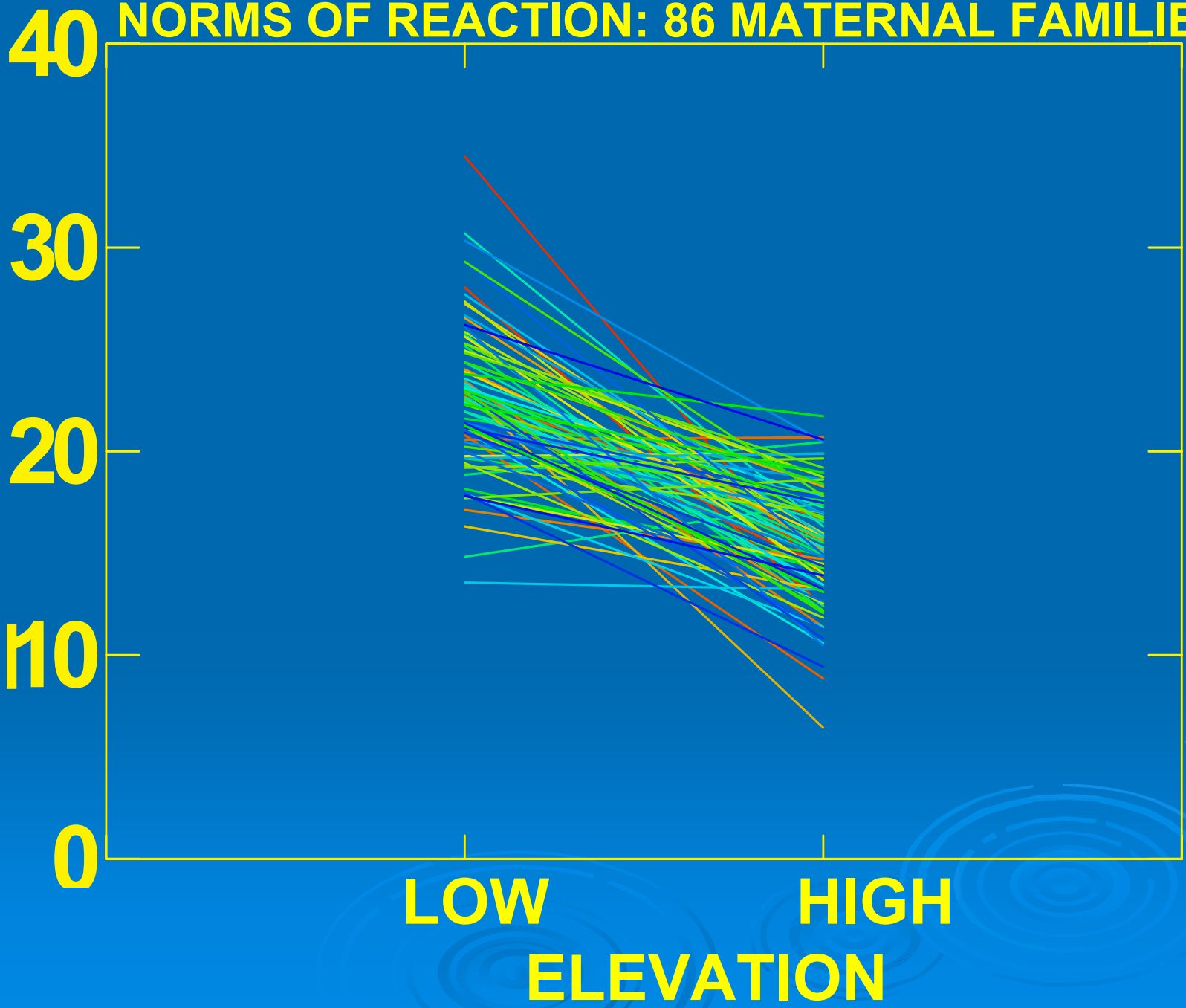
10 20 30 40 50 60 70 80 90
PLANT HEIGHT (CM)

LOW ☆
HIGH ○



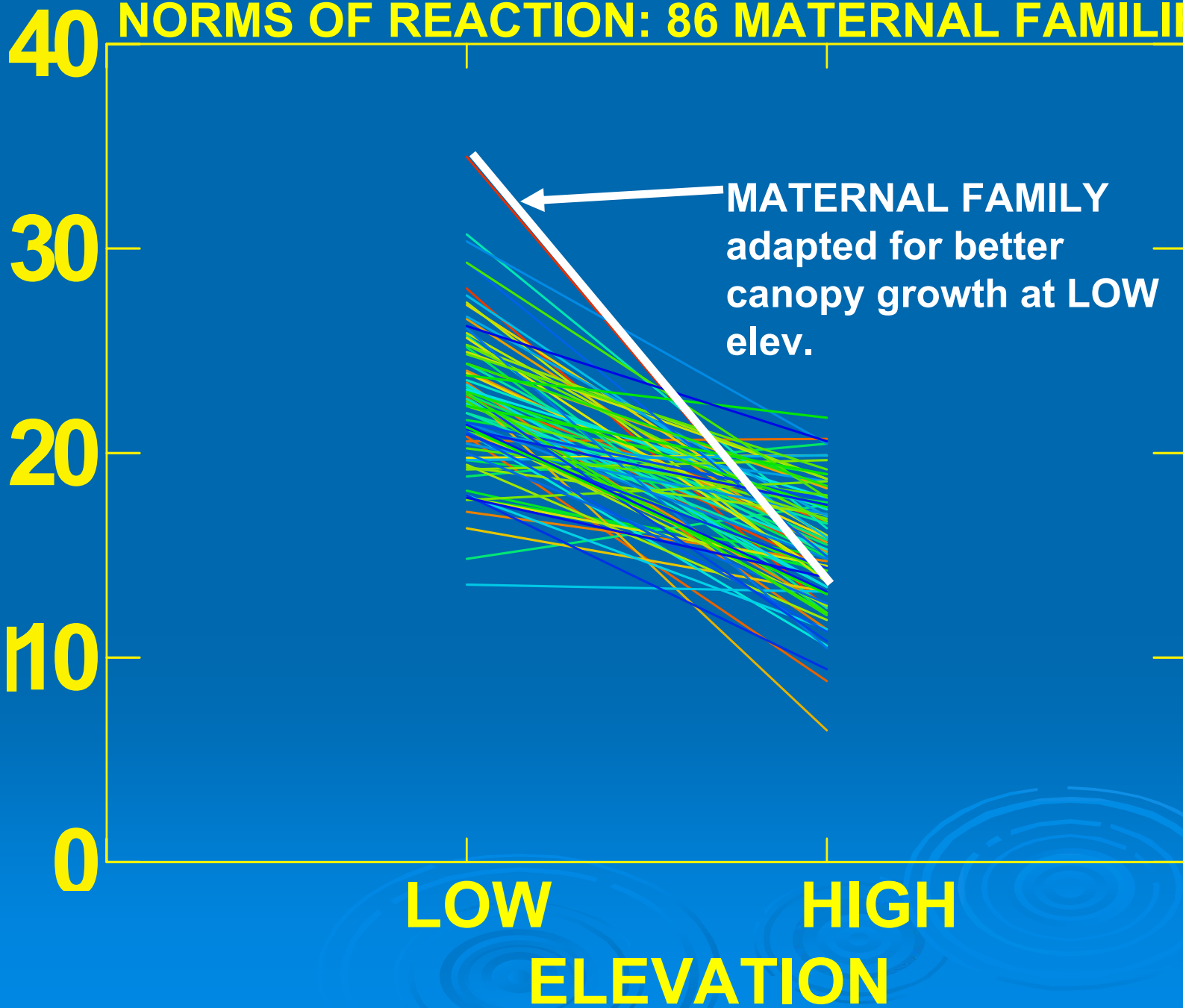
SQ RT(CANOPY AREA IN CM²)

NORMS OF REACTION: 86 MATERNAL FAMILIES



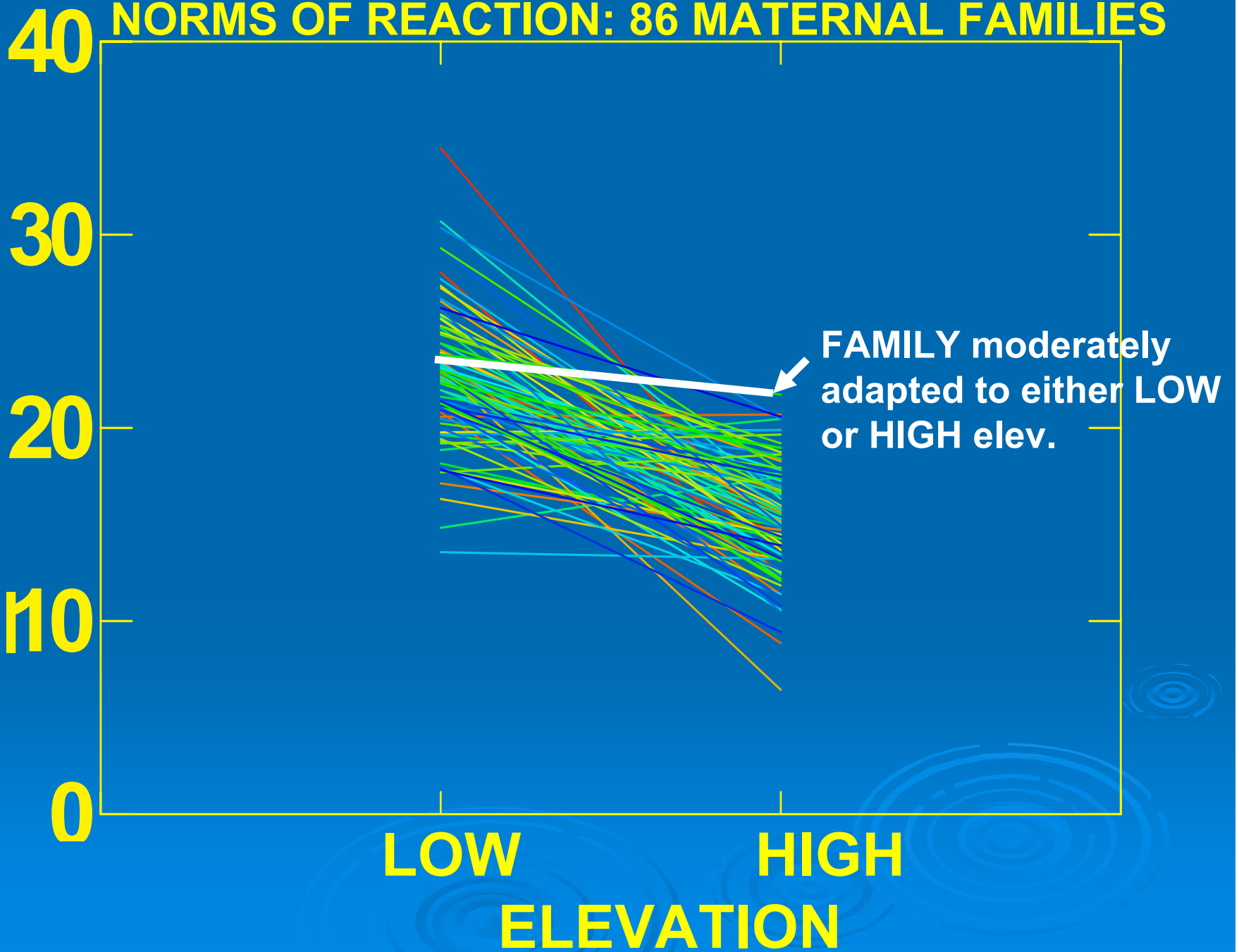
SQ RT(CANOPY AREA IN CM²)

NORMS OF REACTION: 86 MATERNAL FAMILIES



SQ RT(CANOPY AREA IN CM²)

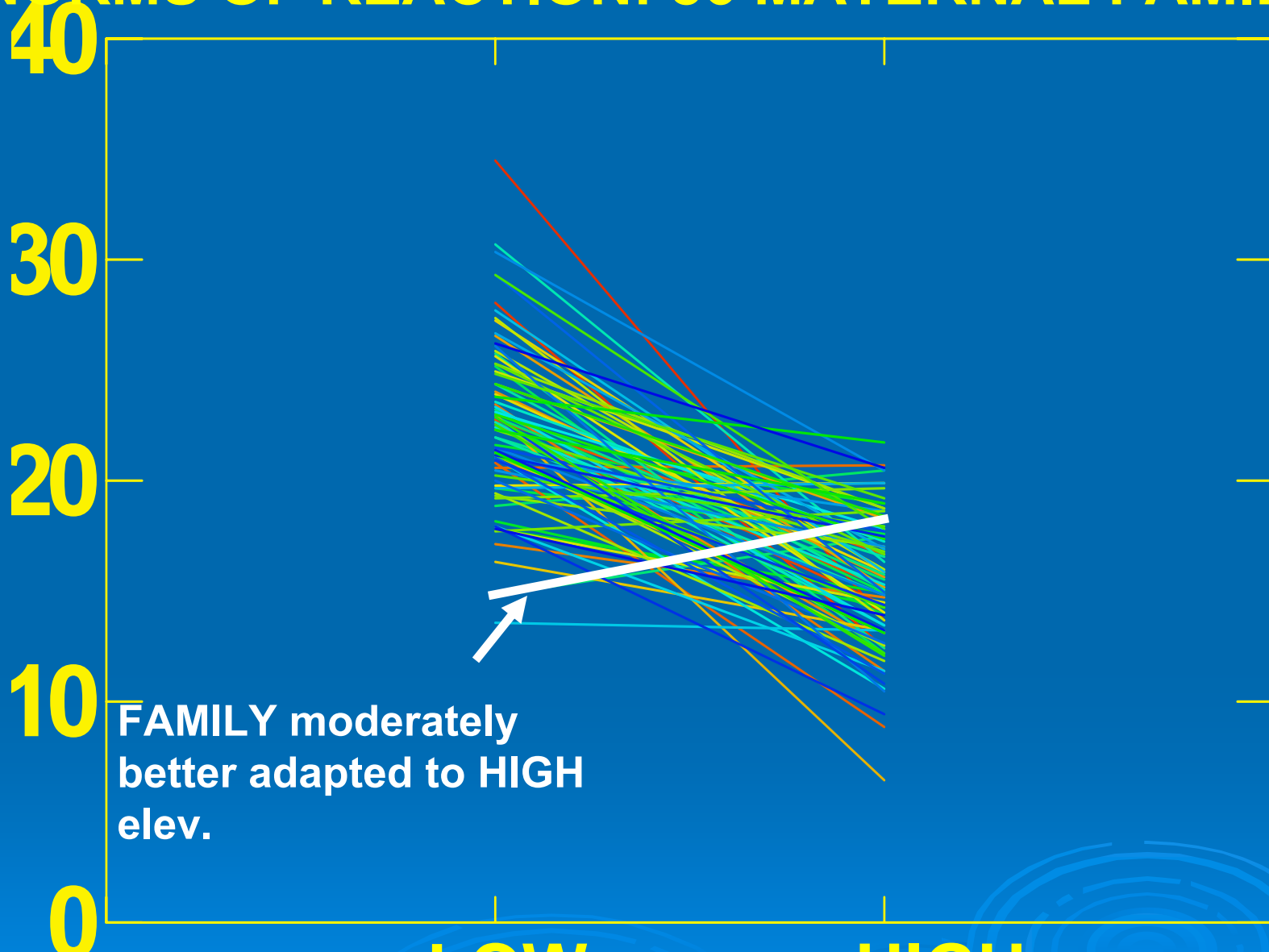
NORMS OF REACTION: 86 MATERNAL FAMILIES



FAMILY moderately adapted to either LOW or HIGH elev.

NORMS OF REACTION: 86 MATERNAL FAMILIES

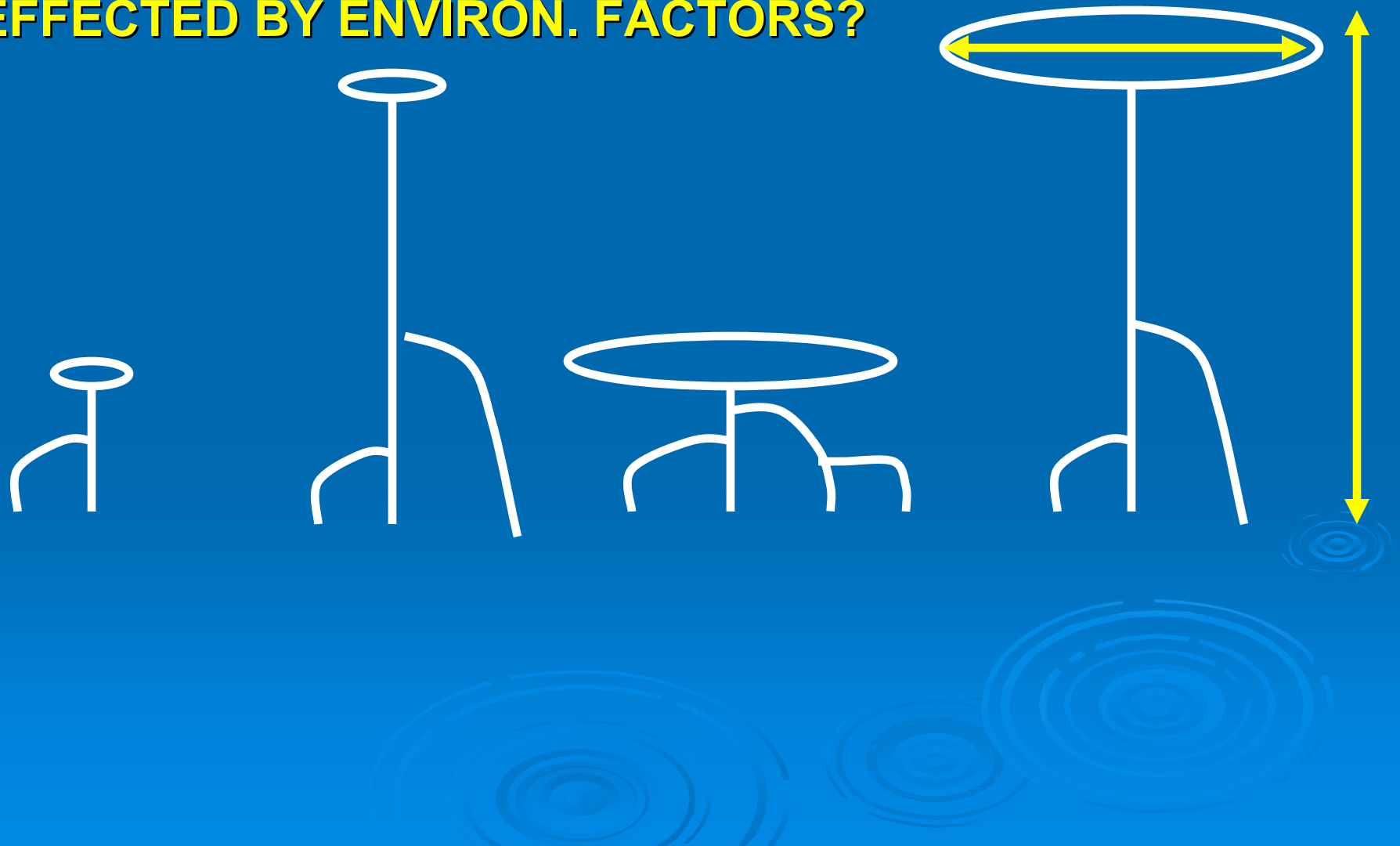
SQ RT(CANOPY AREA IN CM²)



LOW HIGH
ELEVATION

CANOPY AREA / HEIGHT RATIO

- ARE THERE DIFFERENT PLANT ARCHITECTURES?
- DOES IT VARY BY MATERNAL GENOTYPE?
- EFFECTED BY ENVIRON. FACTORS?



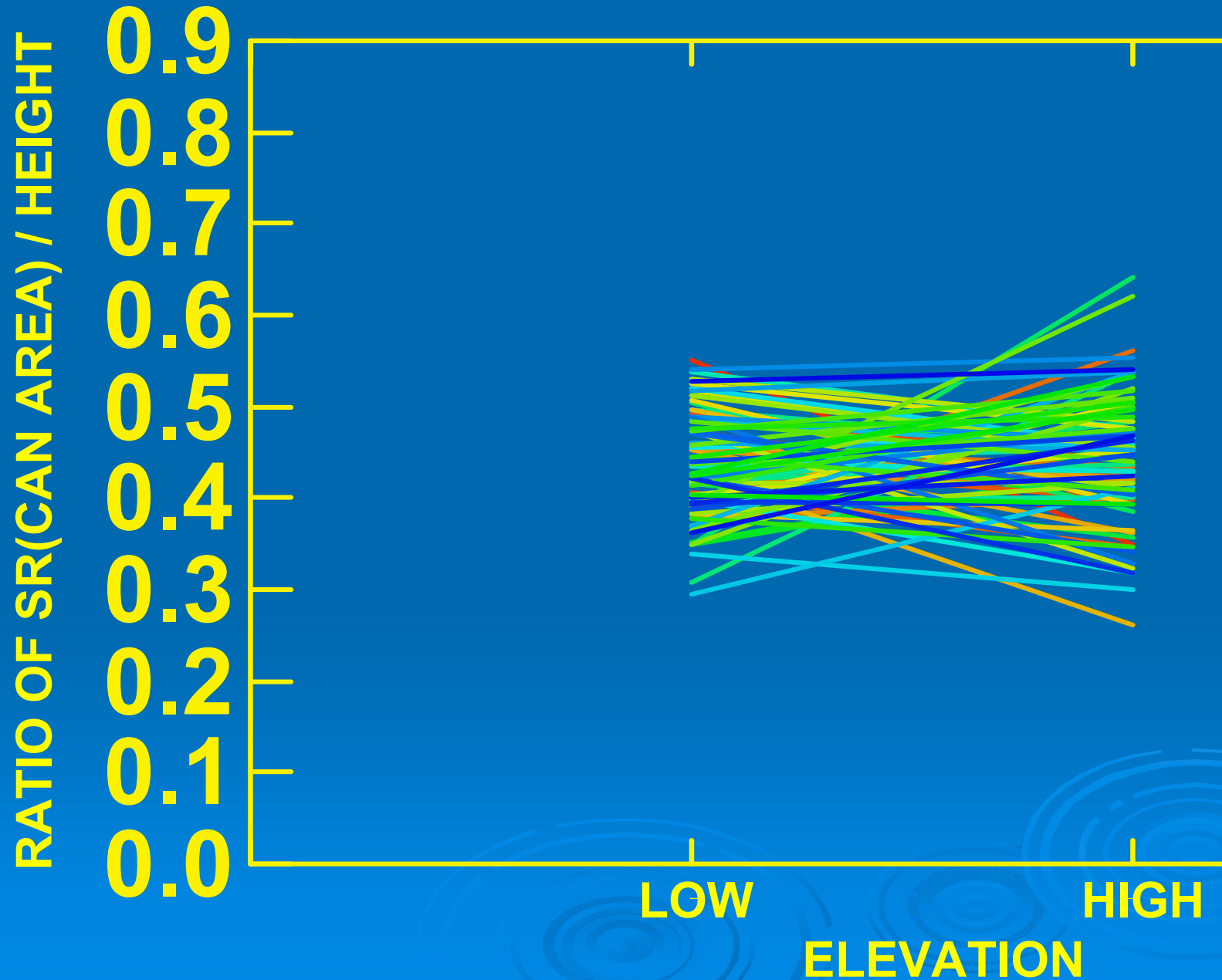
R. MANGLE PLANT ARCHITECTURE (3 YR OLD SEEDLINGS)

ANOVA SUMMARY: SR(CANOPY AREA) / HEIGHT: R²=0.34

| Source | DF | F Value | P |
|-------------------------|-----------|--------------|---------------|
| ELEVATION | 1 | 2.90 | 0.0890 |
| EMBAYMENT | 4 | 0.31 | 0.8687 |
| ISLAND PLANTED | 4 | 29.84 | 0.0001 |
| MAT. FAM(EMBAY.) | 81 | 1.51 | 0.0036 |
| ELV*FAM. (EMBAY.) | 85 | 1.06 | 0.3518 |
| ELEV.*ISLAND | 4 | 15.89 | 0.0001 |
| PROPLEN | 1 | 0.15 | 0.6976 |

- **EFFECT OF MATERNAL FAMILY**
- **EFFECT OF LOCAL ISLAND PLANTED ENVIRONMENT**
- **NO EFFECT OF ELEVATION**
- **NO COVARIANCE WITH PROPAGULE SIZE**

TREE ARCHITECTURE (RATIO OF CANOPY AREA/HEIGHT) 86 FAMILIES NORMS OF REACTIONS AT LOW & HIGH ELV.



PLANT ARCHITECTURE

- MATERNAL FAMILY HAS A LARGE, SIGNIFICANT EFFECT
- ELEVATION DOES NOT
- LOCAL SEEDLING ENVIRONMENT DOES



REPRODUCTION

- **BEGAN PRODUCING PROPAGULES**
 - **A FEW PLANTS AT YEAR 2**
 - **A NUMBER OF PLANTS AT YEAR 3**



2007 (3 yrs old): Reproduction



ISL 2 LOW
31

Reproduction: Propagules YR 3

Elevation

Low

High

% Plantings Producing Propagules

4.9%

4.6%



Reproduction: Propagules

**% of Maternal Families
whose seedlings Produced
Propagules in 2007**

40%

**% Families. Repro. at
LOW Elv. Only**

25.6%

**% Families. Repro. at
HIGH Elv. Only**

20.9%

**% Families. Repro. at
BOTH Elvs.**

7%



Potential for Local Adaptation Exists

- **May affect the rate at which *R. mangle* (relative to other species) moves with global change**
 - Interactions among species affected
- **Can influence restoration success**
 - Diversity of seedling parentage may be very important
- **Confirm with next experiment using F₁ generation of this Tampa Bay study**

Next Step 2: Red Mangrove Genetic Diversity

Florida *R. mangle* may have lost genetic diversity because of climate fluctuation bottlenecks

Does lower GD affect potential for further change with climate now? For interactions with other mangroves and salt marsh species?

Florida: Low gen. div. per AFLP
(Travis & Proffitt, unpubl.; Devlin, unpubl.)

Colombia: high Gen. Div. (microsatellite heterozygosity)
(Argelaez-Cortes et al. 2007)



Thanks

- Jordan Sanford, Katie Tiling, and Glenn Coldren for all the help in the field
- Brandt Henningsen for promoting use of the Schultz Family Park restoration site
- Randy Runnells for help in all phases of the Tampa Bay study
- USGS for funding of the first year and for logistic help

END



Field Experiment: SUMMARY

- **Seedling height, canopy area, and “architecture” (can. Area / ht) affected by**
 - **Maternal Sibling Family and probably genotype (NEXT F₁ experiment)**
 - **Seedling environment**
 - **Island scale**
 - **Several m within island (transect) scale**
 - **Propagule size (maternal effects + genotype)**
 - **NOT maternal tree environment (embayment)**

END



MANOVA OF GROWTH AT 2007



Correlations among the 3 “growth” response variables (2007): height, trunk diameter (dbh) and canopy area

| | dbh07 | SR_carea |
|-------|--------------------|--------------------|
| ht07 | 0.554507 <.0001 | 0.620549 <.0001 |
| dbh07 | | 0.490476 <.0001 |

MANOVA: response variables Height, Trunk Diameter, and Canopy Area all measured at 2007 (3 years)
Sig. = from Wilk's Lambda multivariate analysis

| <i>SOURCE</i> | <i>SIG.</i> | <i>Wilk's Lambda</i> |
|----------------------------------|---------------|----------------------|
| MATERNAL TREE EMBAYMENT | NS | |
| ELEVATION (LOW or HIGH) | 0.0001 | 0.427 |
| PLANTING ISLAND (1...5) | 0.0001 | 0.740 |
| MATERNAL TREE GENOTYPE | 0.0001 | 0.307 |
| PROPAGULE LENGTH (COVAR.) | 0.0001 | 0.955 |
| ELEV. X ISLAND | 0.0001 | 0.910 |
| ELEV. X GENOTYPE | 0.0002 | 0.577 |
| ISLAND x GENOTYPE | 0.0001 | 0.096 |
| ISL X GENO X ELEV | 0.9991 | 0.573 |

UNIVARIATE ANOVAS (P VALUES)

| <i>SOURCE</i> <i>R²</i> | <i>HT</i> <i>0.863</i> | <i>DBH</i> <i>0.822</i> | <i>CAN. AREA</i> <i>0.691</i> |
|---------------------------------------|---------------------------|----------------------------|----------------------------------|
| ELEVATION | 0.0001 | 0.0001 | 0.0001 |
| PLANTING ISLAND | 0.0001 | 0.0001 | 0.0001 |
| MATERNAL GENOTYPE | 0.0050 | 0.0580 | 0.0200 |
| PROPAGULE LENGTH | 0.0001 | 0.0002 | 0.1320 |
| ELEV. X ISLAND | 0.0001 | 0.0043 | 0.0001 |
| ELEV. X GENOTYPE | 0.0001 | 0.9954 | 0.9698 |
| ISLAND x GENOTYPE | 0.0001 | 0.0001 | 0.9989 |
| ISL X GENO X ELEV. | 0.2934 | 0.9973 | 0.9486 |

2004 STORM EFFECTS (BURIAL)

Buried

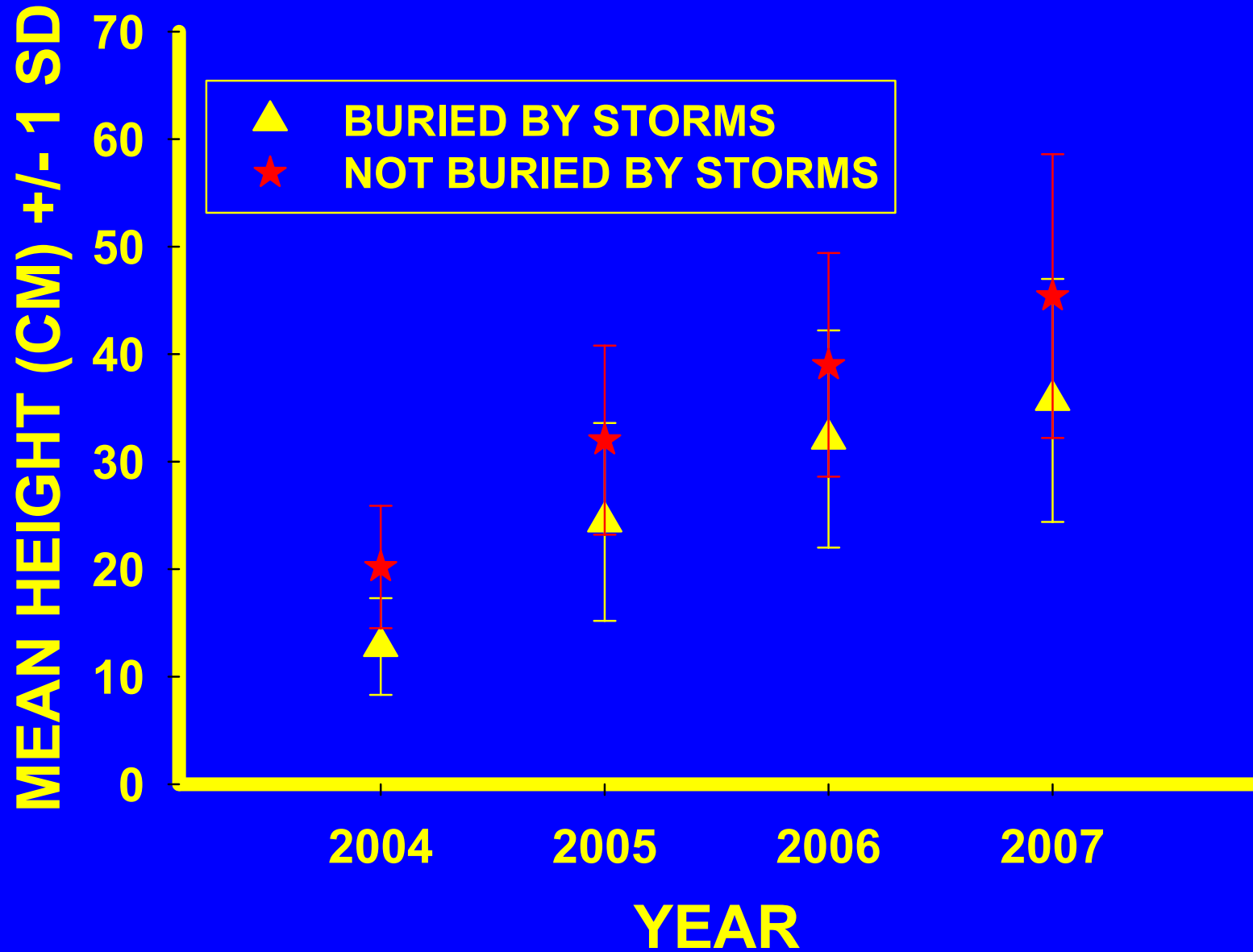


Seriously Buried



(PHOTOS 6 MONTHS POST 2004 STORMS)

GROWTH OF SEEDLINGS BURIED BY STORMS SHIFTING SAND AROUND



Spartina alterniflora (smooth
cordgrass)
and
Rhizophora mangle (red mangrove)

- Both are foundation species at diff. latitudes
- Interact with one another at transition latitudes (in part of Florida peninsula)

Spartina alterniflora salt marsh





Rhizophora mangle **(red mangrove)**

- **Foundation species in tropics & subtropics**
- **Low – mid intertidal**
- **Viviparous (seedling propagules live on maternal trees for 4-6 mo.)**
- **Evergreen – continuous growth & leaf production**

species comparisons

- *Spartina*
 - clonal grass
 - clonal genets may be very long-lived but individual ramets are not
 - sexual: near-obligate outcrosser
 - Not shade tolerant
- *Rhizophora*
 - tree (non-clonal although does make new trunks & canopy by iteration)
 - fairly long lived (maybe 70-100 years?)
 - sexual: highly selfing in many estuaries; but, this varies from 0-33% outcrossing
 - some degree of shade tolerance

N ↑

Spartina

Influenced strongly by
global change
Position
Size ?

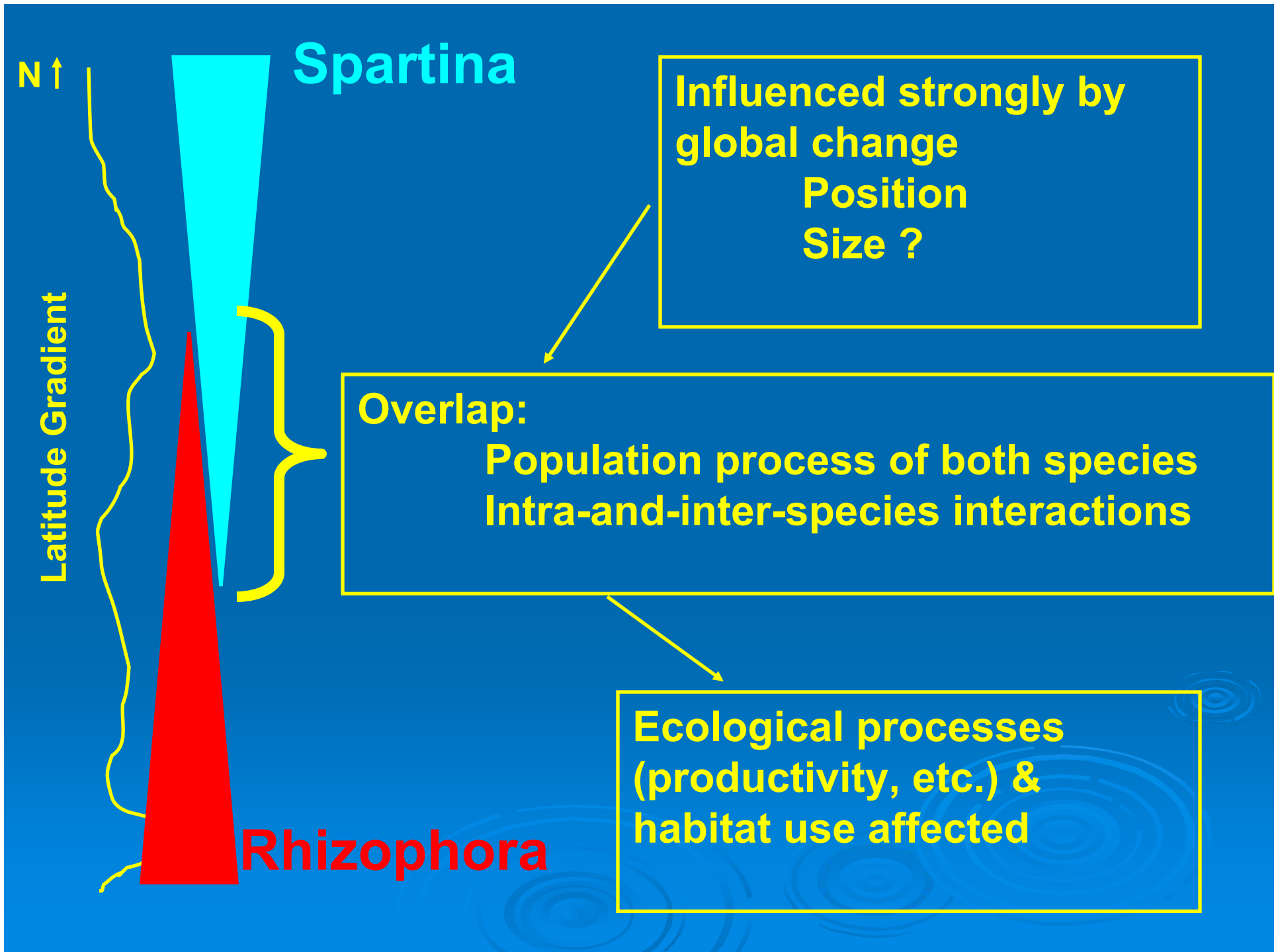
Latitude Gradient

Overlap:

Population process of both species
Intra-and-inter-species interactions

Ecological processes
(productivity, etc.) &
habitat use affected

Rhizophora



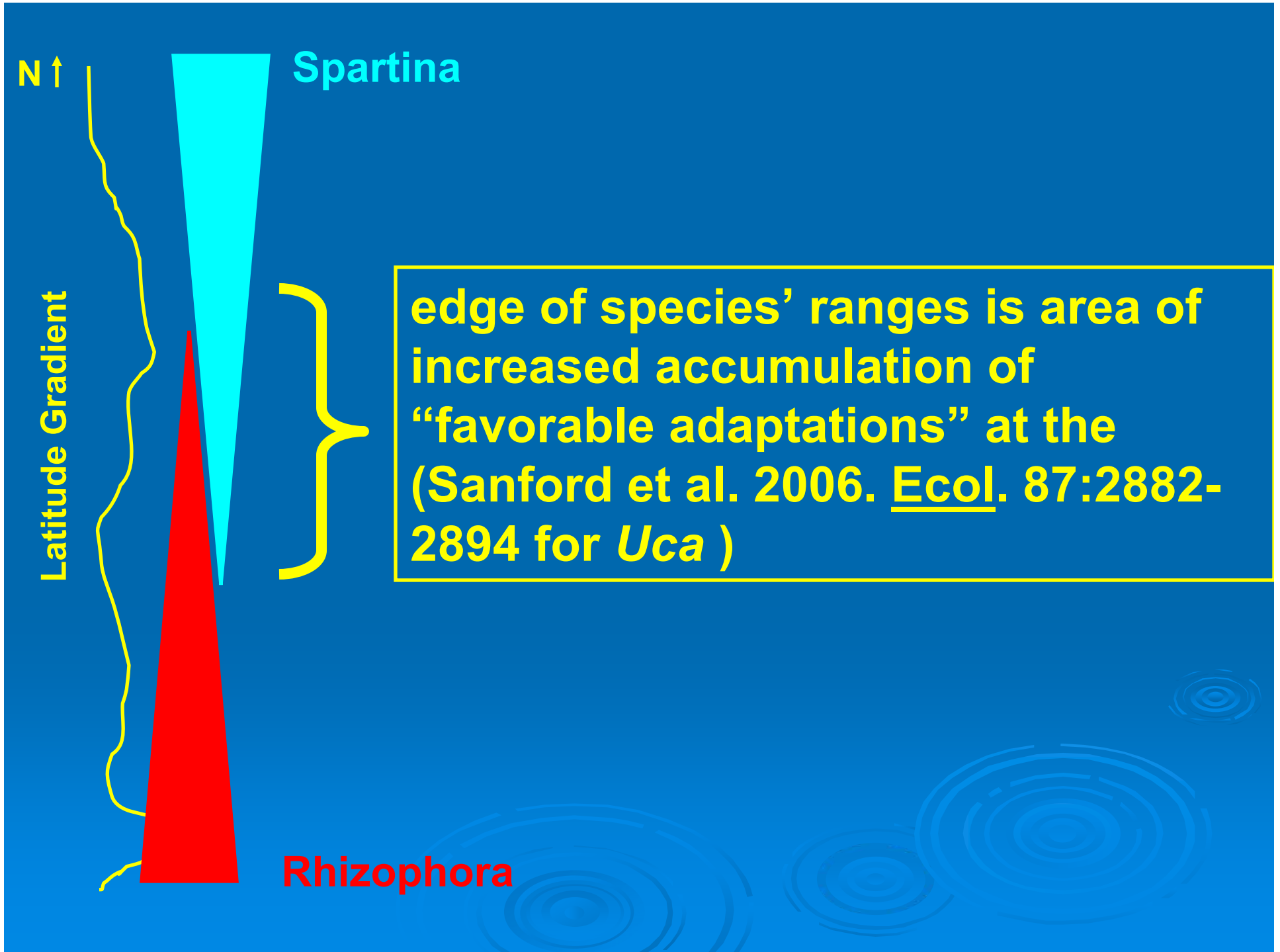
N ↑

Spartina

Latitude Gradient

edge of species' ranges is area of increased accumulation of "favorable adaptations" at the (Sanford et al. 2006. Ecol. 87:2882-2894 for *Uca*)

Rhizophora



salt marsh & mangrove at latitudes where they dominate



**Mangrove “islands” within a salt marsh “matrix”
Merritt Island National Wildlife Refuge**



Merritt Island National Wildlife Refuge Banana River (Kennedy Space Center)



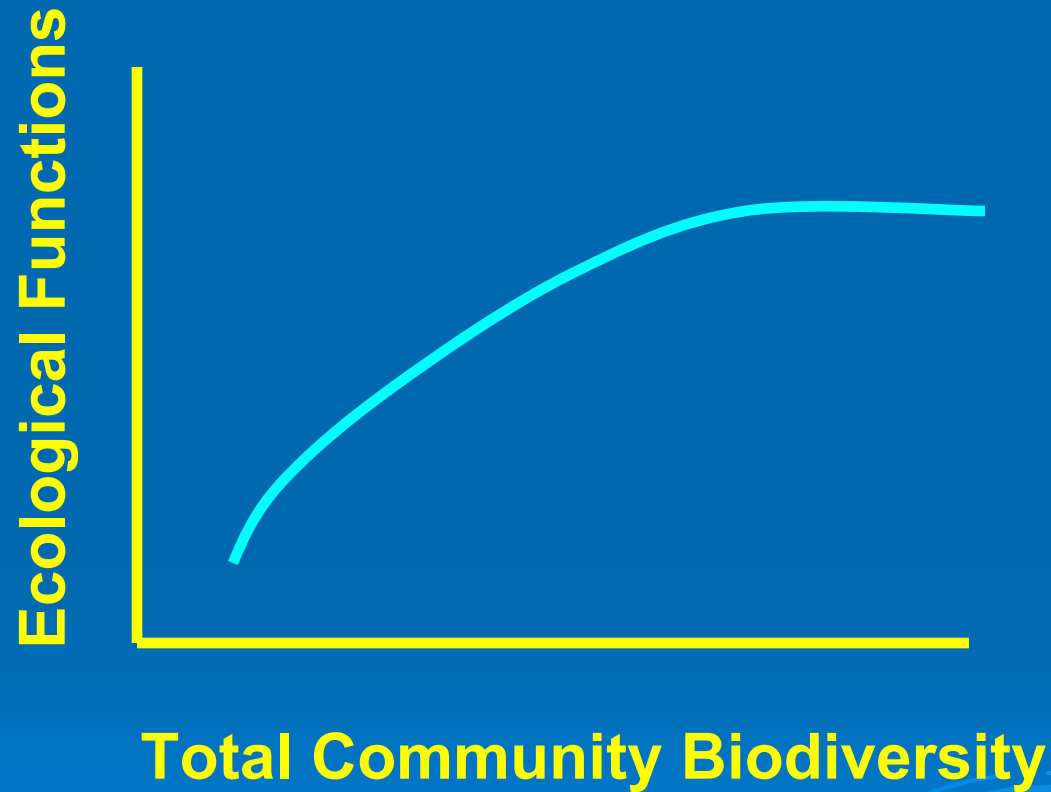
mangroves

Spartina alterniflora
displaced seaward

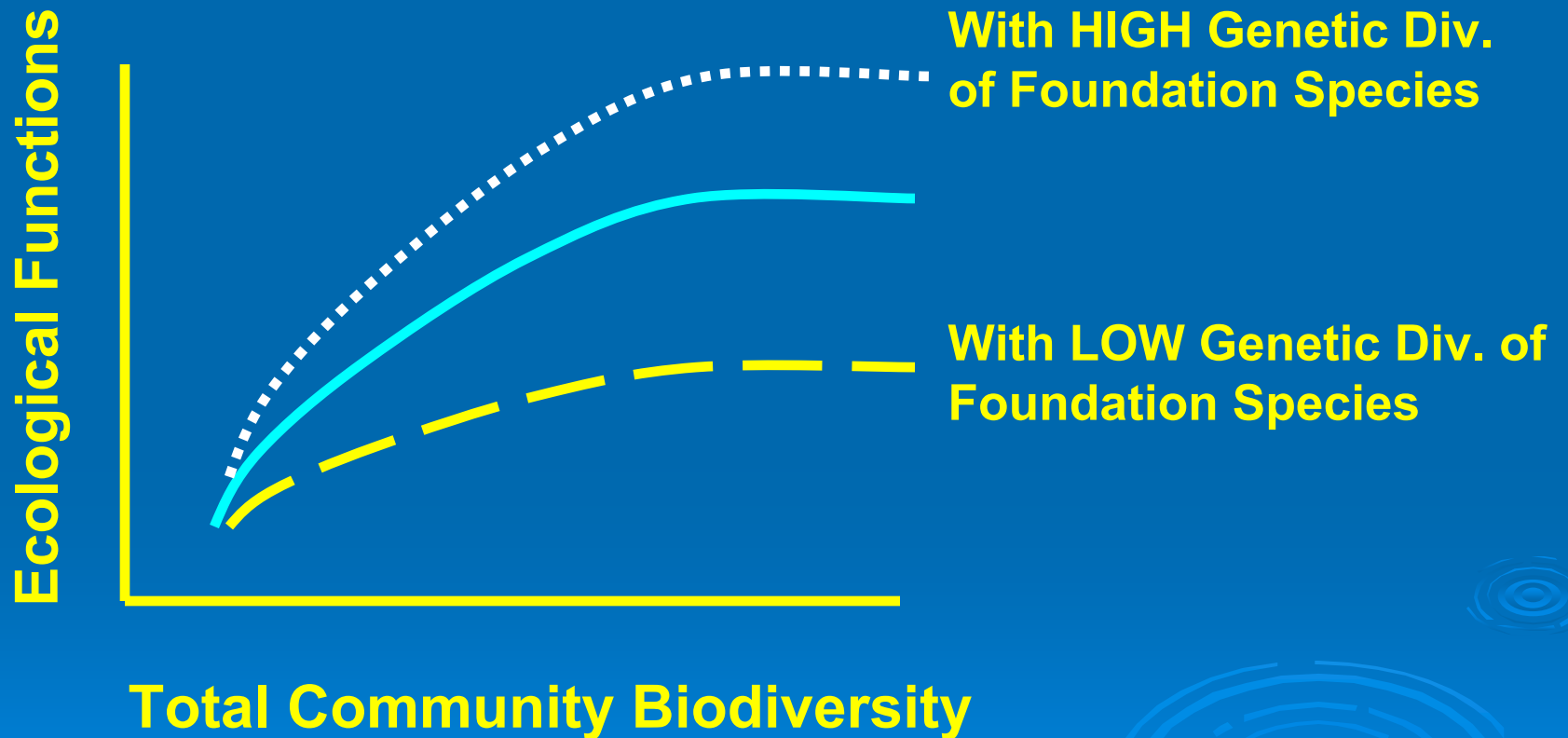
Population & Genetic Ecology

- **Essential to understanding change with climate and change due to human manipulations**
- **Critical to understand in Foundation Species because affects so many other species**

WHY IS DIVERSITY OF FOUNDATION SPECIES IMPORTANT?



WHY IS DIVERSITY OF FOUNDATION SPECIES IMPORTANT?



Knowledge Bases

- *Spartina alterniflora*: many studies
- *Rhizophora mangle*: few studies

Field Experiment
Spartina alterniflora
genetic ecology

***Spartina alterniflora* SALT MARSH
FOUNDATION SPECIES GENETIC
ECOLOGY: Genotypes vary:**

- In morphology and architecture (Proffitt et al. 2003)
- In ecological effects
 - On other species (competition and facilitation). Proffitt et al. 2005)
 - Intra-specific competition w/ other genets of *Spartina* (Proffitt & Travis in review)
- In genetic and genotypic structure among marshes and over latitudinal gradient (Travis et al. 2003).



Premises: With Climate Change

- **Mangroves will have to adapt to changing**
 - Physical environmental conditions
 - Biotic interactions
- **Overlap of temperate & tropic foundation species may be area of enhanced selection for traits needed with climate change**
- **Requires**
 - Phenotypic Plasticity
 - Genetic variability
 - Genetic component of plasticity x environment changes (evolution *for* plasticity)

Related Question

- **Restoration (i.e., the “Northern Estuaries” of Everglades Restoration)**
 - **Are there different requirements because of**
 - **Proximity of subtropical / temperate biogeographical limits**
 - **Changes in climate (and factors related to climate, like hurricane frequency, etc.)**

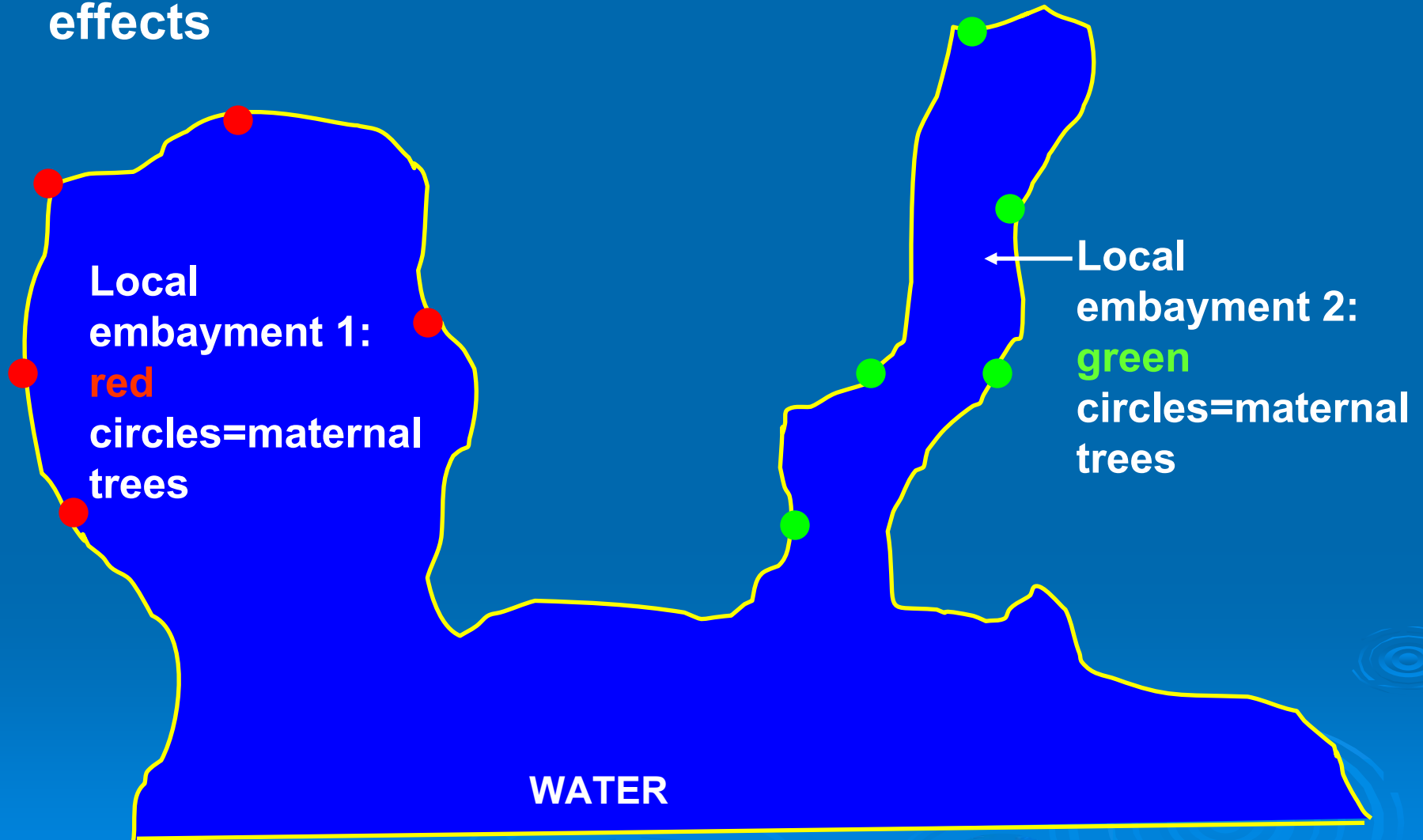
Needed Studies

- **Genetic variability and gene flow**
- **Outcrossing and inbreeding rates**
- **Experiments addressing:**
 - **What are the important physical and biotic stressors (and resources)?**
 - **Is there potential for local adaptation?**
 - **What affects colonization, dispersal, and recruitment of seedlings/saplings into canopy?**
 - **Interactions with other foundation species (e.g., salt marsh *Spartina alterniflora*)**

***R. mangle* genetic diversity ?**



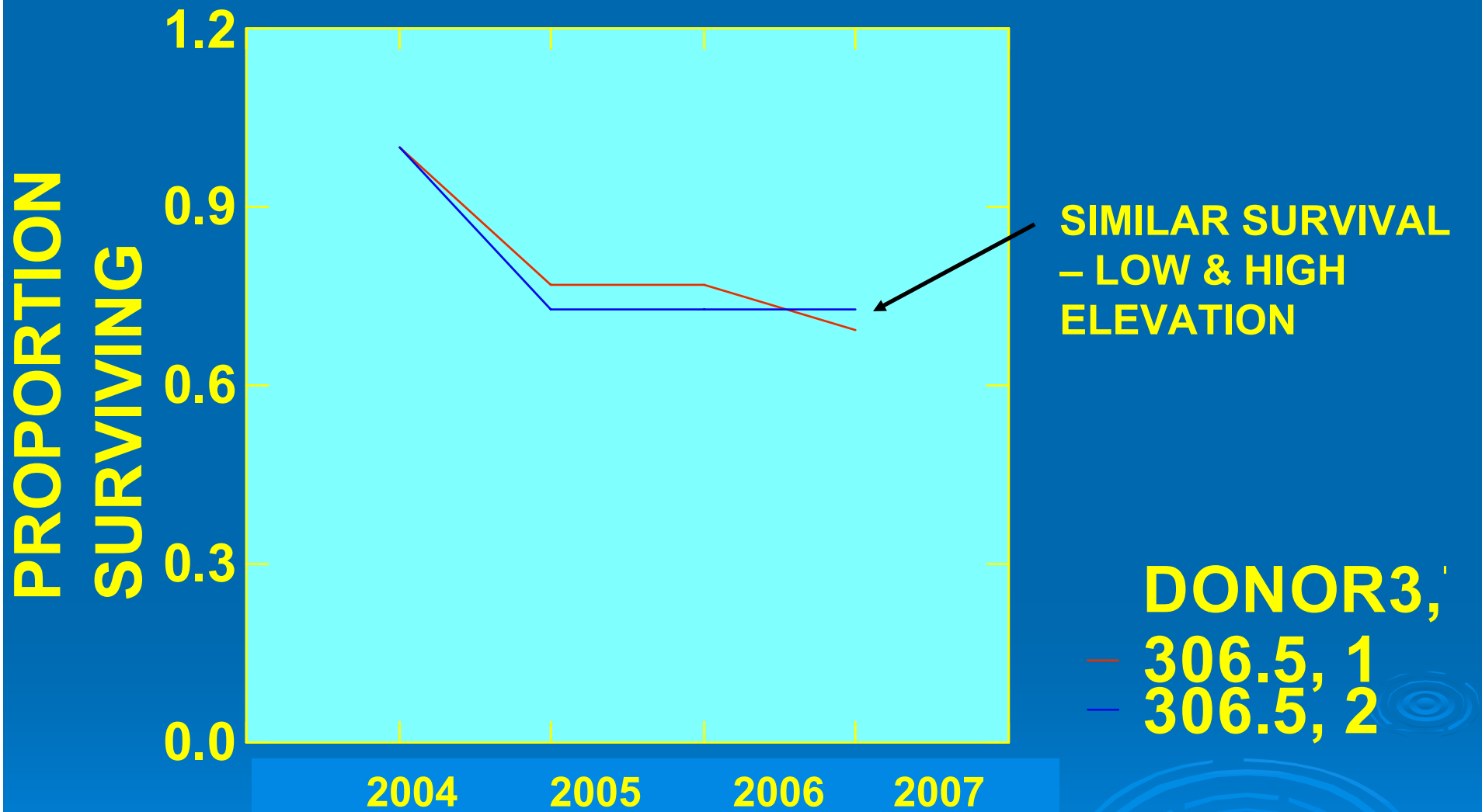
Account for **SITE** effects (local embayment environmental conditions) **DONOR TREE (maternal families)** **WITHIN SITE** effects



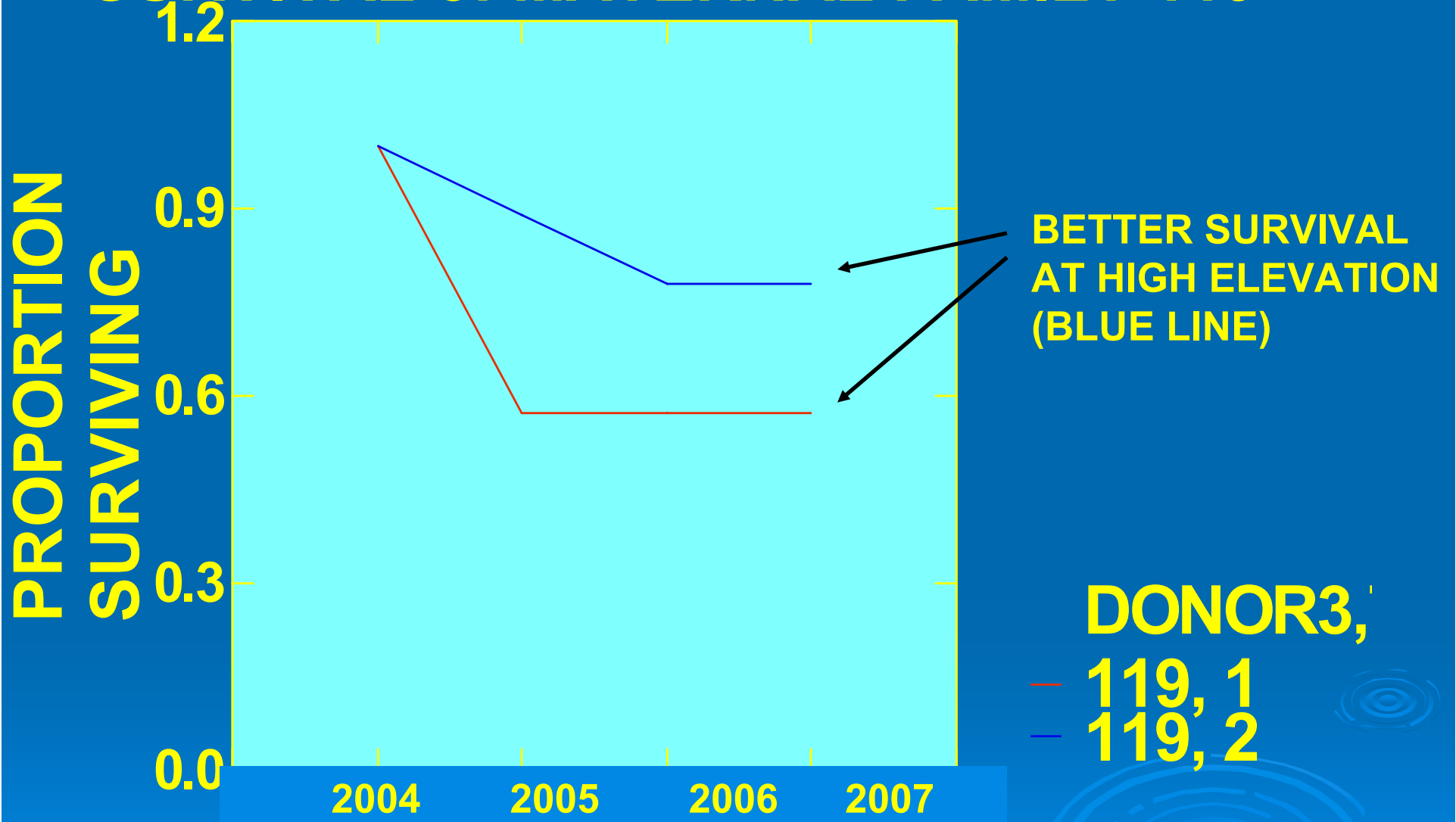
SURVIVAL of MATERNAL FAMILY 201.5



SURVIVAL of MATERNAL FAMILY 306.5



SURVIVAL of MATERNAL FAMILY 119

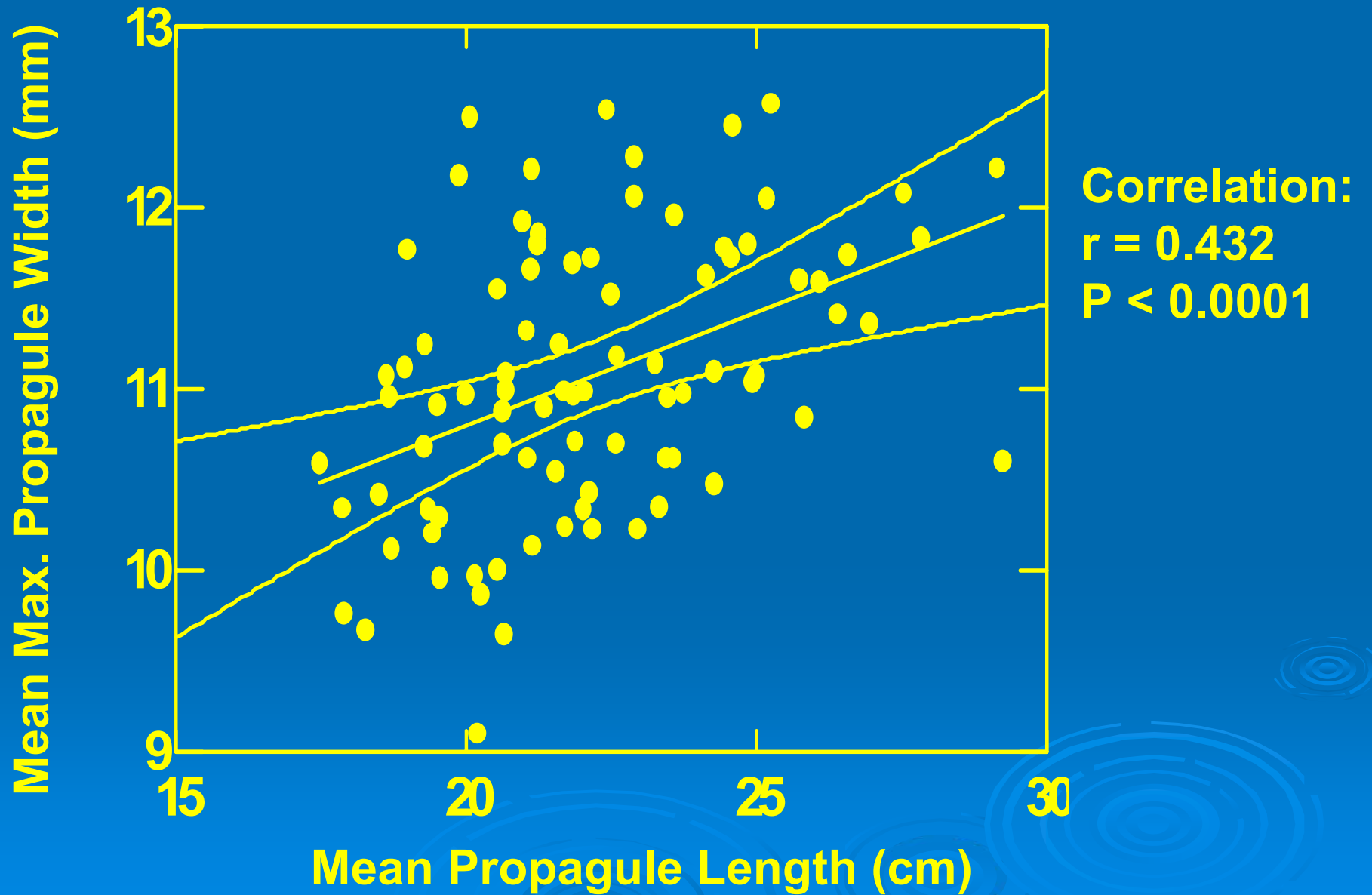


Propagule Length vs Max. Width

- Are propagule size measurements correlated?



Propagule Length vs Width (Family Means)



Does Propagule Size Differ

- Among Embayments (sets of maternal trees from same general location)
- Among maternal trees within embayment

ANOVA Response variable: **Propagule Length**

$R^2 = 0.46$

Source

p

Embayment

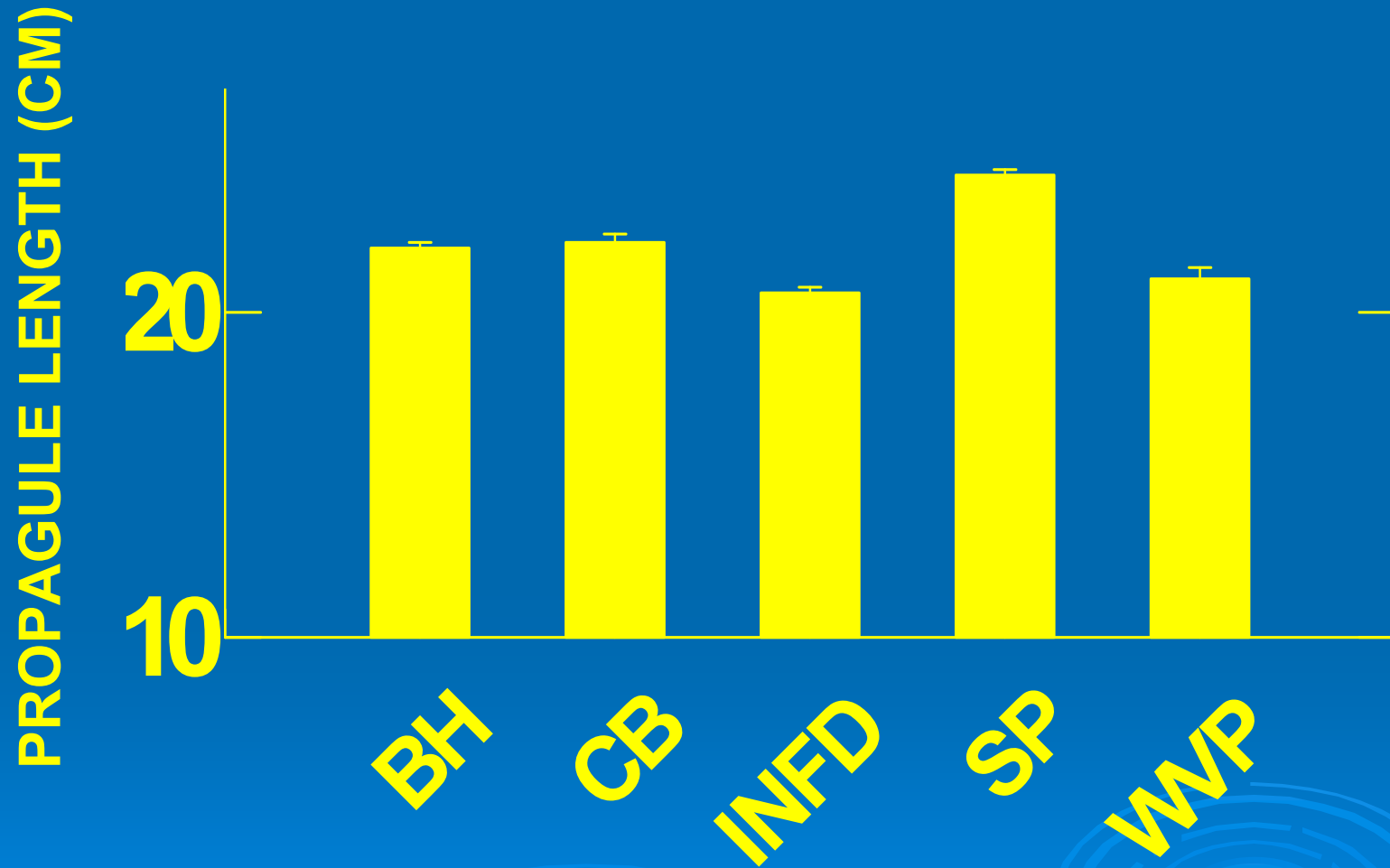
0.0001

Maternal Family(Embay.)

0.0001

PROPAGULE LENGTH by Embayment

ANOVA $F_{4,1596} = 85.411$; $p < 0.0001$



Embayment within Tampa Bay

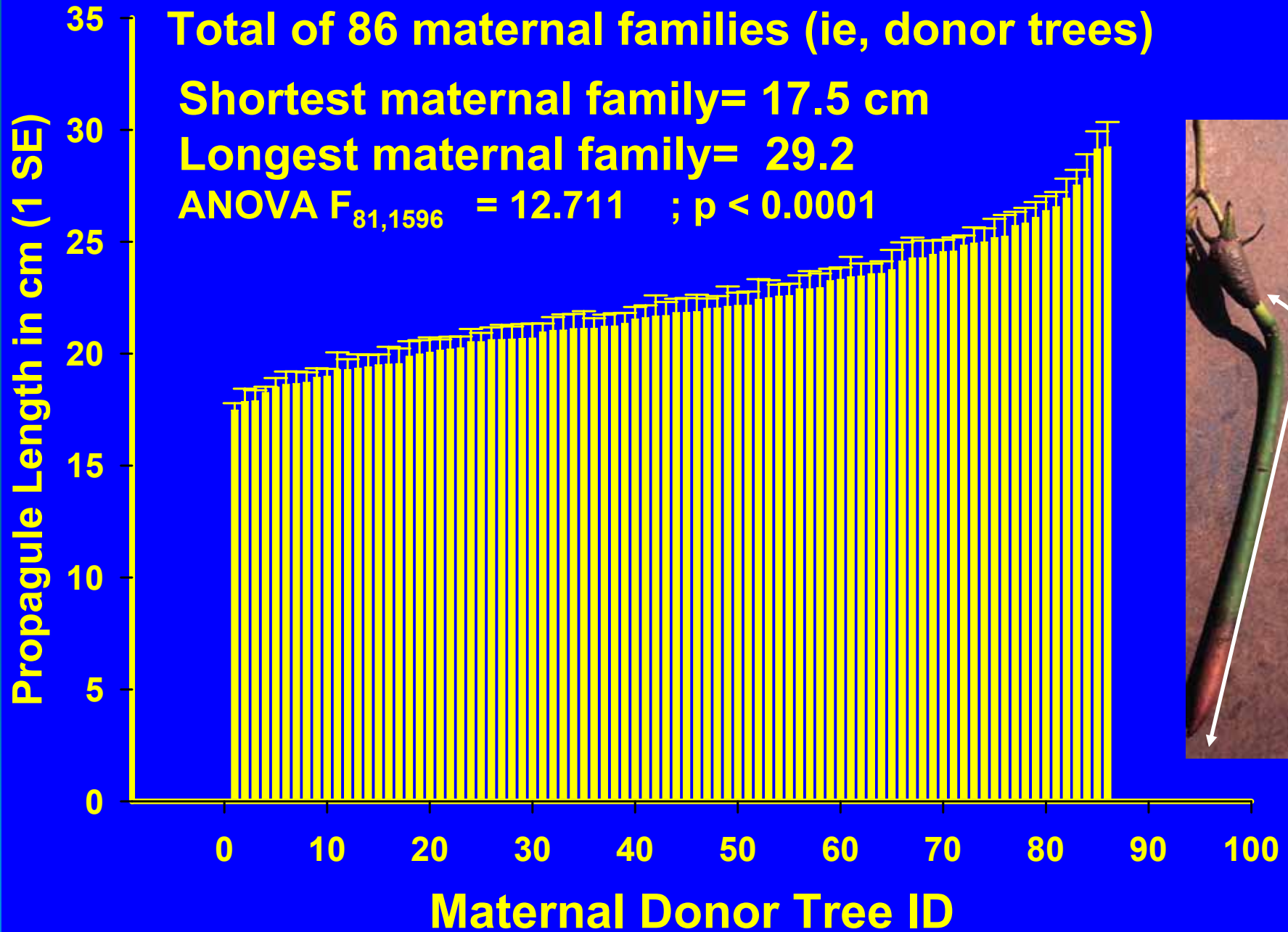
Rhizophora propagule length (mean & SE by donor tree)

Total of 86 maternal families (ie, donor trees)

Shortest maternal family= 17.5 cm

Longest maternal family= 29.2

ANOVA $F_{81,1596} = 12.711$; $p < 0.0001$

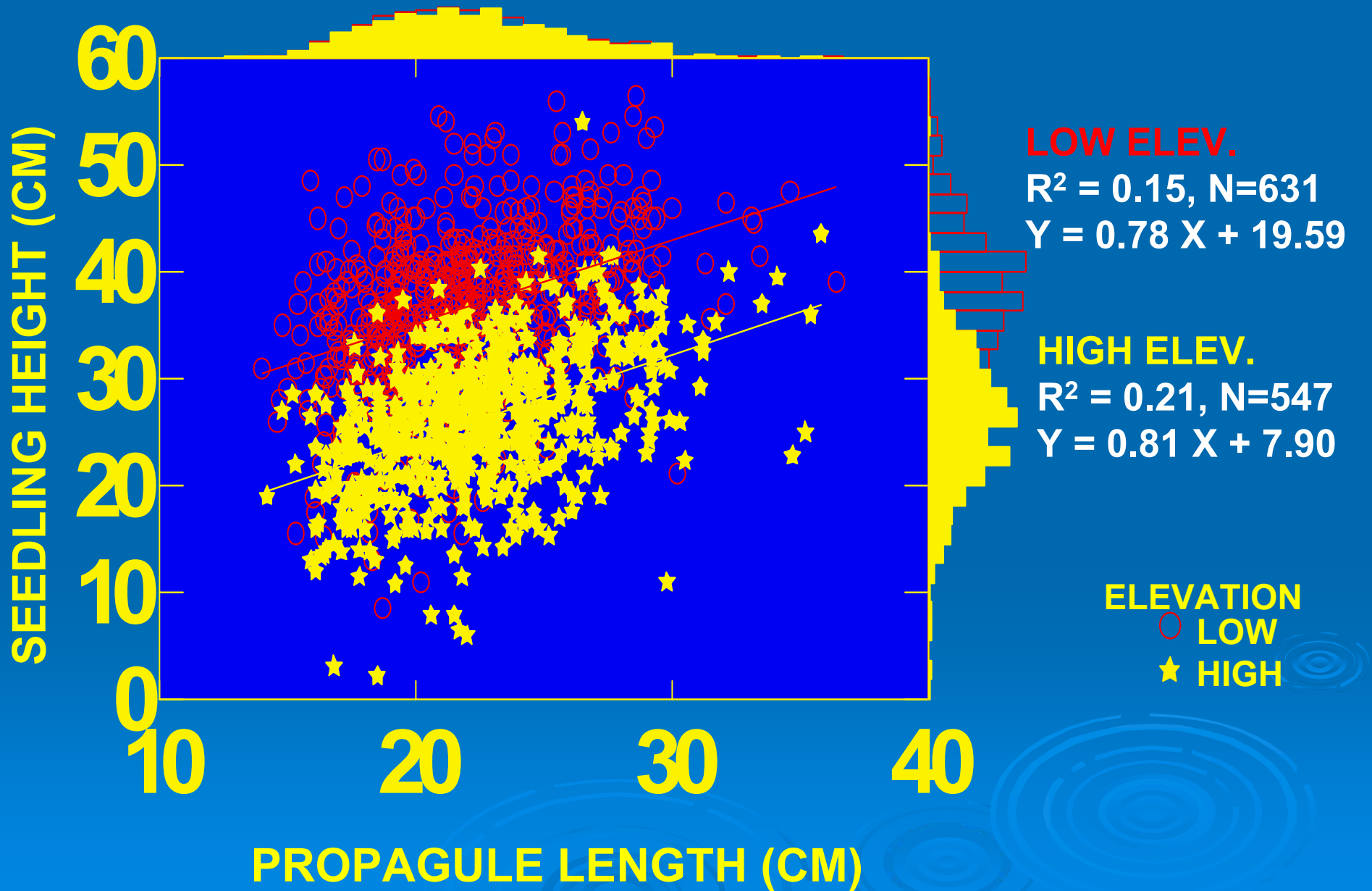


IS SEEDLING HEIGHT AFFECTED BY PROPAGULE SIZE?

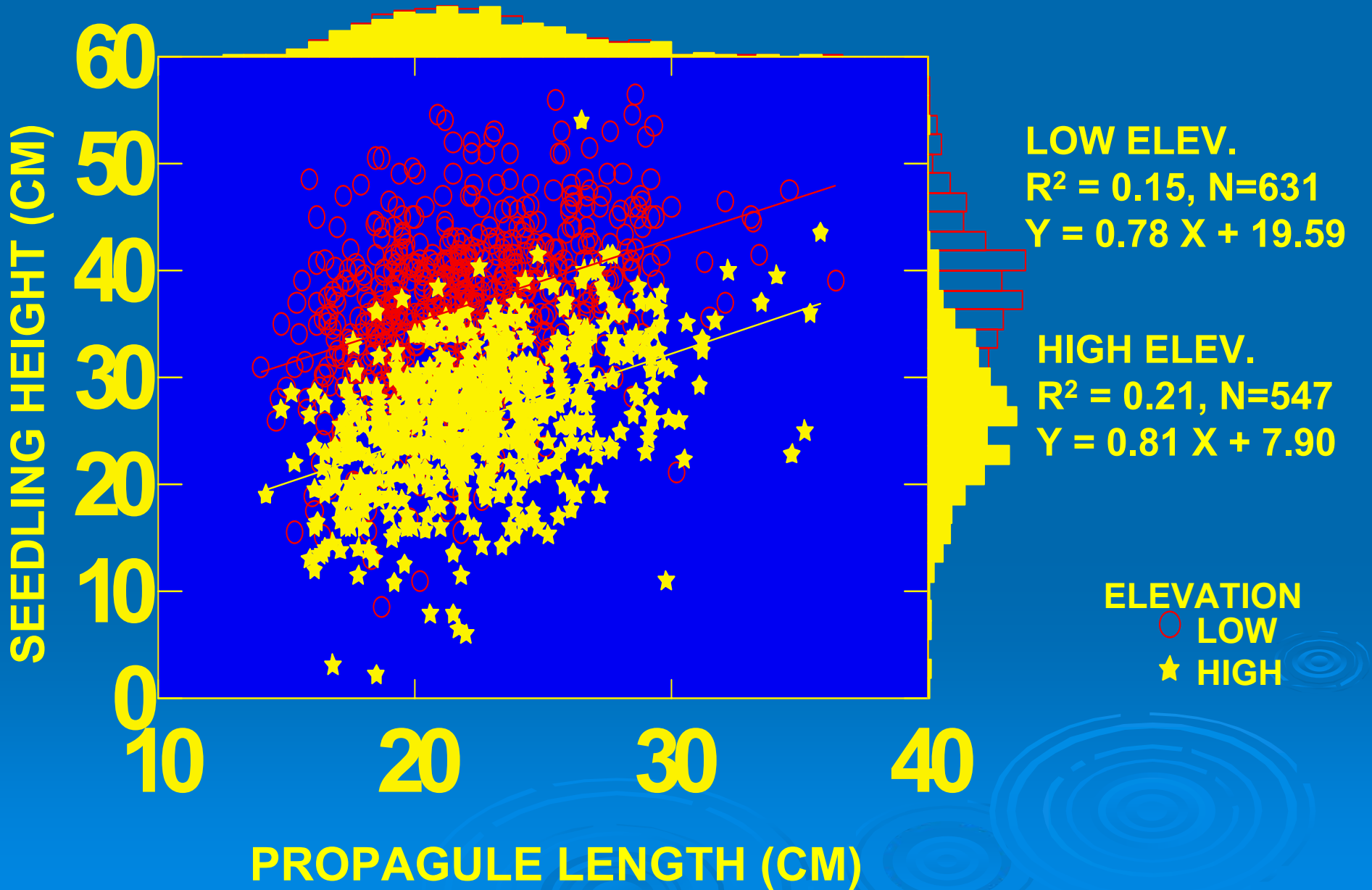
Summay of Linear Regression Results:

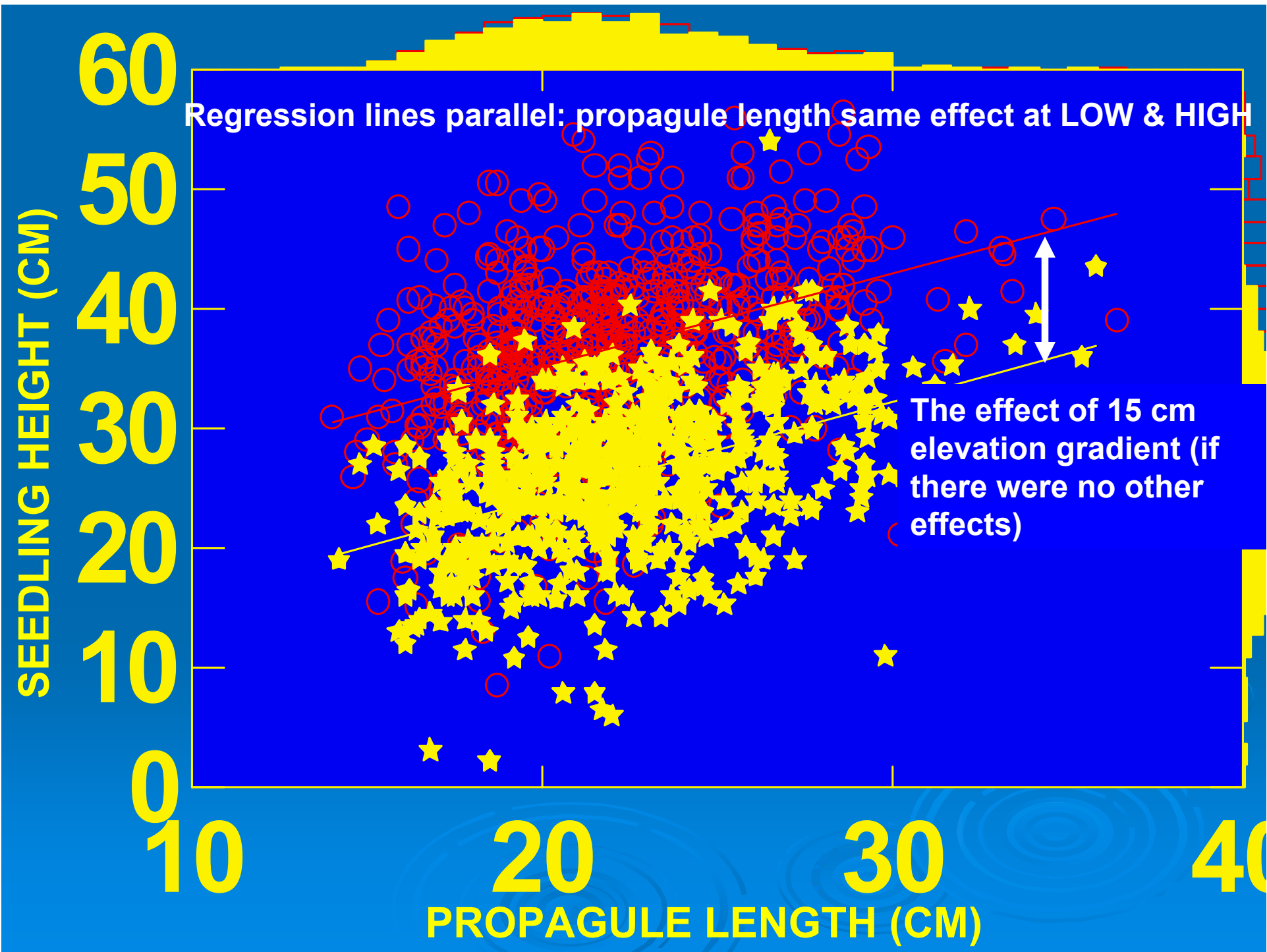
- **Weak, but significant + relationship**
- **Occurs in first year**

REGRESSION: HEIGHT AT 1 YEAR ON PROPAGULE LENGTH



REGRESSION: HEIGHT AT 1 YEAR ON PROPAGULE LENGTH





Reproduction: Propagules

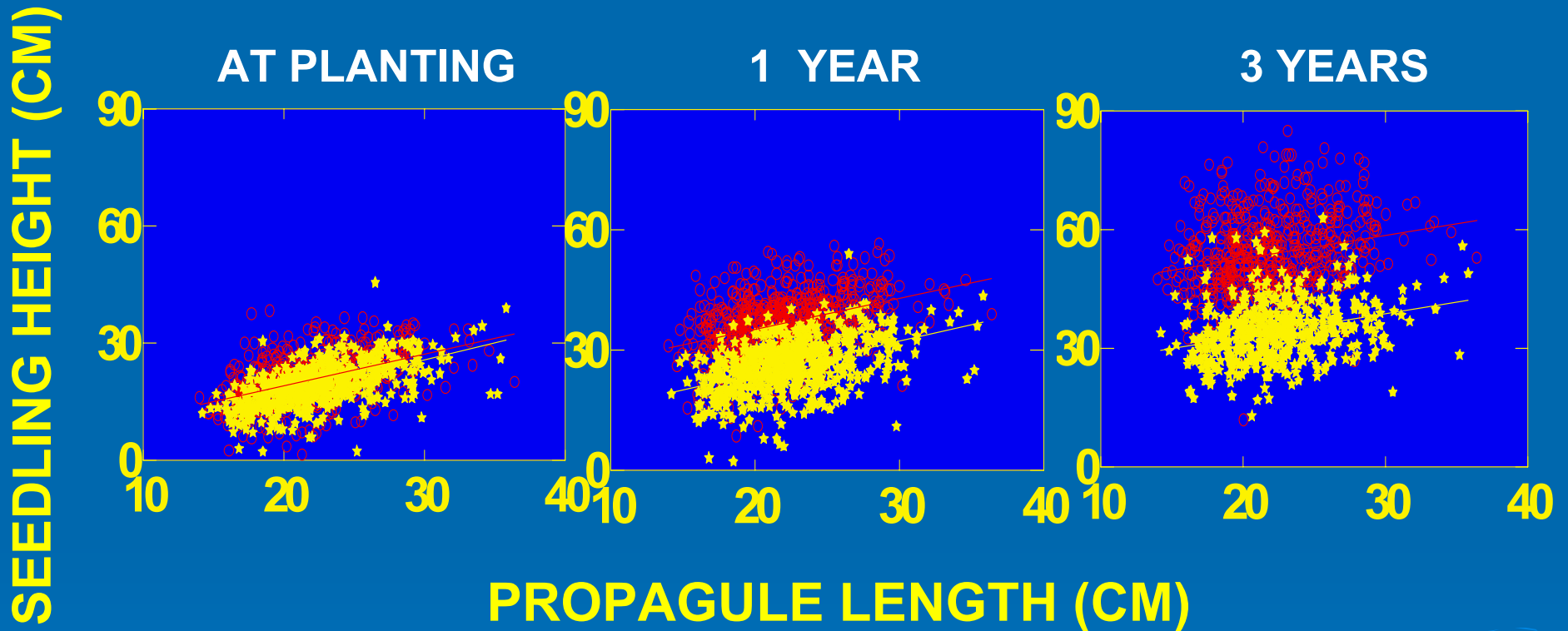
For Logistic Regression: 0 = Produced 1 propagule
1 = Produced >1 propagule

Logistic Regression Results:

Significant variation in propagule production (0 or 1) with CANOPY AREA *BUT, not a big biological diff.* (odds ratio: 1.1), variance explained: McFadden's Rho-Sq=10.6%)

No diff. at either ELEVATION or COMBINED ELEV's:

PLANT HEIGHT VS PROPAGULE LENGTH: **LOW** & **HIGH** ELEVATIONS



- PROPAGULE SIZE EFFECT IN YR 1 (SAME AT LOW & HIGH)
- SCATTER INCR. IN LATER YRS AS OTHER FACTORS BEC. INCREASINGLY MORE IMPORTANT

ISLAND 4

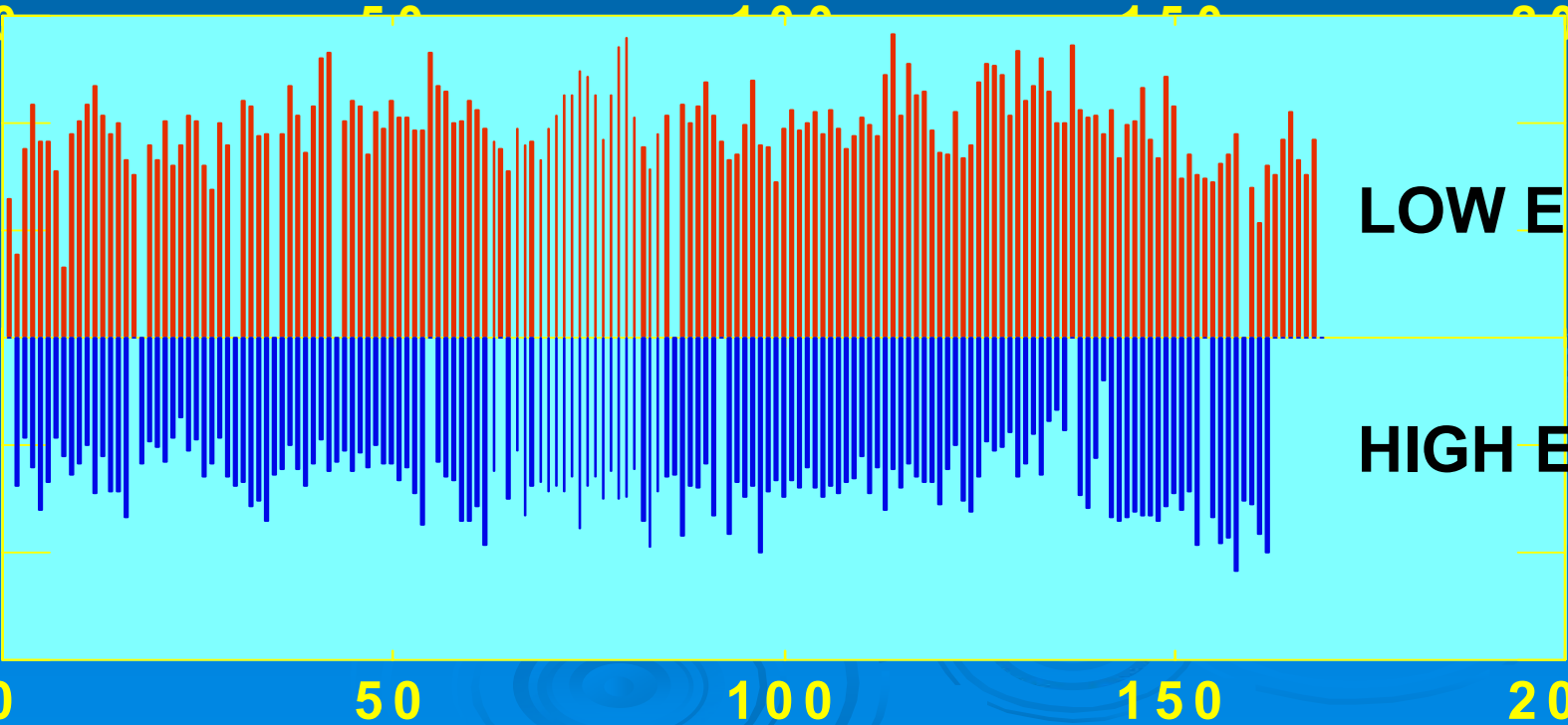
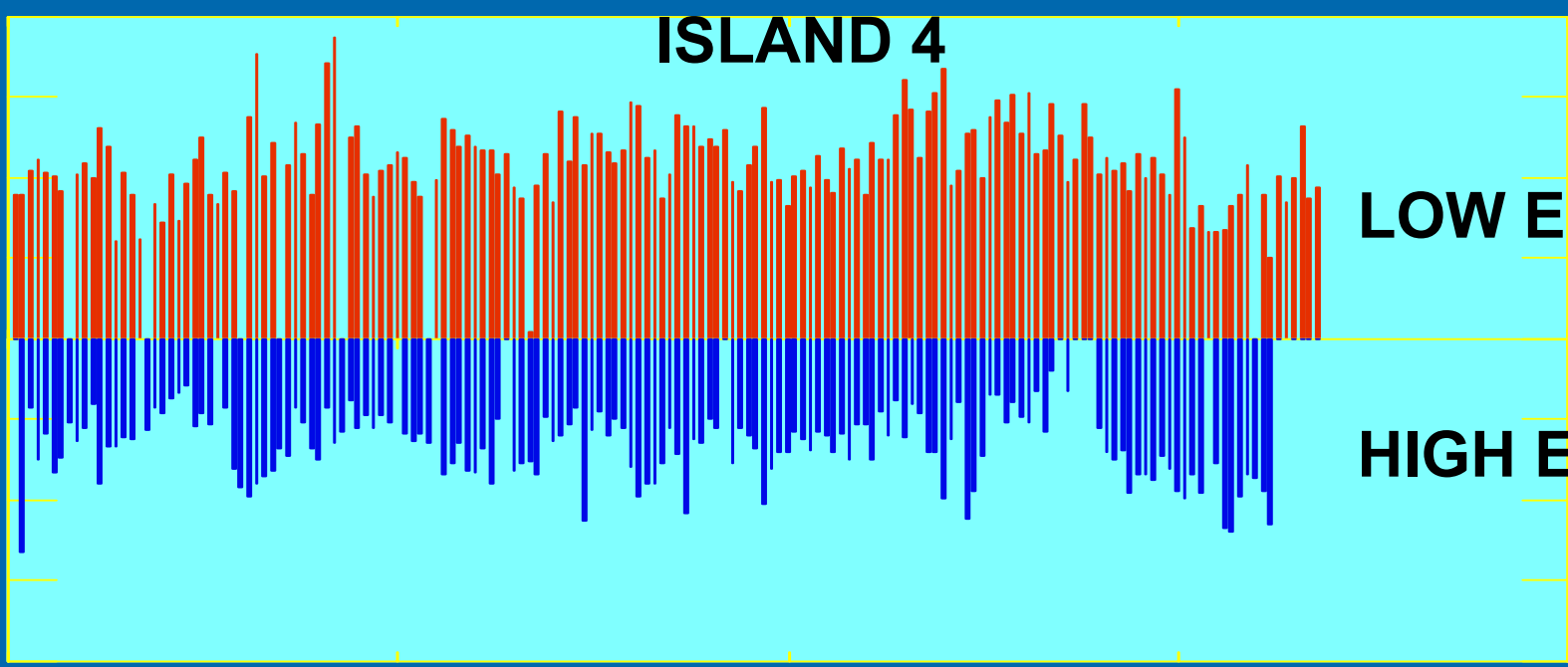
HT07

HT07

HT805

HT805

90
70
50
30
10
0
-10
-30
-50
-70
-90
60
40
20
0
-20
-40
-60



LOW Elv.

HIGH Elv.

LOW Elv.

HIGH Elv.

°C

R-M ANOVA for Height (summary)

- Height varied with 15 cm elevation gradient
- Height varied with island planted (ie, spatial scale of seedlings over 10's of meters)
- Height varied with maternal tree family (genotype?) BUT NOT with maternal tree location (ie., the embayment)
- Effects varied over the 3 years
 - Island Planted
 - Elevation
- Effects stable over 3 years
 - Maternal family (maternal genotype)

