## Homework3

Baxter Worthing

9/12/2018

 To begin our discussion of this paper, we should recall an important equation:

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$$V_P = V_G + V_E$$

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 Total phenotypic variation is the sum of the contributions of genetic variation and environmental variation

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Today we're going to focus on  $V_E$ 

•

$$V_E = V_{Eg} + V_{GxE} + V_{ES}$$

•

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•

$$V_{Eg}$$
 :

•

$$V_E = V_{Eg} + V_{G \times E} + V_{ES}$$

$$V_{Eg}$$
:

General environmental varience

•

$$V_E = V_{Eg} + V_{G \times E} + V_{ES}$$

•

$$V_{Eg}$$
:

General environmental varience

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$$V_{ES}$$
:

•

$$V_E = V_{Eg} + V_{G \times E} + V_{ES}$$

$$V_{Eg}$$
 :

General environmental varience

$$V_{ES}$$
:

• Specific environmental varience

$$V_E = V_{Eg} + V_{G \times E} + V_{ES}$$

$$V_{Eg}$$
 :

• General environmental varience

$$V_{ES}$$
:

• Specific environmental varience

$$V_{G\times E}$$
:

 $V_{5} - V_{5} + V_{6} = 1$ 

$$V_E = V_{Eg} + V_{G \times E} + V_{ES}$$

 $V_{Eg}$  :

• General environmental varience

 $V_{ES}$ :

• Specific environmental varience

 $V_{G\times E}$ :

 Genotype by environment interaction (what this paper was interested in)

# Characterizing GxE

Regulatory Divergence

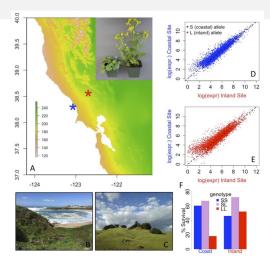
## Characterizing GxE

- Regulatory Divergence
- Allele-specific Expression (ASE)

### Main goals of this paper

- "determine the prevalence of GxE interactions affecting the expression of native and non-native alleles"
- "characterize the relative contributions of cis and/or trans regulatory variation in expression divergence"
- "establish the role of genome structure (e.g.inversions) in the evolution of potentially locally adaptive regulatory variation"
- "identify candidate genes underlying local adaptation in this system"

#### Figure 1



This is the first figure from this paper. A shows a map of the locations of their two transplant sites. B shows a picture of the blue location and C shows a picture of the red location. D compares the expression of coastal alleles at both sites, while E shows the same for inland allele. F shows the difference in survival between parental plants and F1 hybrids at each site.

# Figure 2

