

# Misinterpretations of admixture regression in racial hereditarian research

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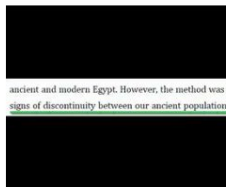
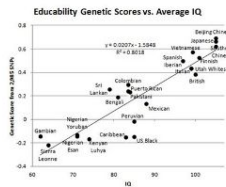
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**AABA 2024: Undermining the production of race science**

# Racist Science in the 21st century

The New York Times

## *Why White Supremacists Are Chugging Milk (and Why Geneticists Are Alarmed)*



## RESEARCH ARTICLE

## How White nationalists mobilize genetics: From genetic ancestry and human biodiversity to counterscience and metapolitics

Aaron Panofsky✉, Kushan Dasgupta, Nicole Iturriaga

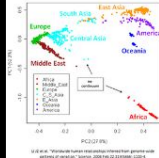
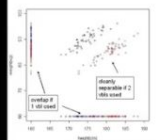
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## The Cultural Marxist War against Darwinism

**Creationists:** evolution is a social construct, not biologically real.  
**Liberal Creationists:** race is a social construct not biologically real.  
**Charles Darwin:** I'm not a creationist; I actually wrote: "There is, however, no doubt that various races, when carefully compared and measured, differ much from each other..."



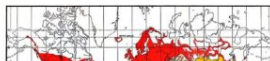
**Fact:** There are 100,000+ people of African descent in the Americas, but only 10% of the genetic research on the continent has included them.

When you sample and map them, patterns emerge across the continent.

This clustering is a result of evolution due to environmental factors.

Africans are central to the history of the Americas, but only 10% of the genetic research on the continent has included them.

**Lothrop Stoddard World Race Map (1920):**



**No Credible Scientists Believe in Race**

Some people feel that they do not have the needed expertise to judge the validity of race. So, they defer to the experts, and the experts tell them that race does not exist. The problem with this argument is, even though the most vocal anthropologists and biologists deny race, academic surveys show that there is no actual consensus on this topic.

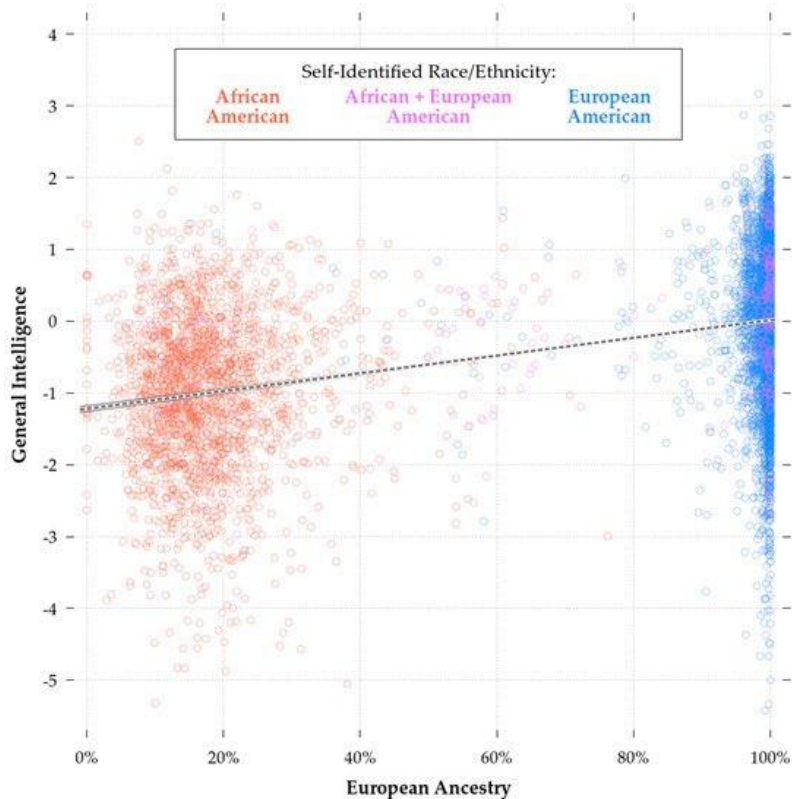
Year	Non-medical description of race	Medical description of race
1952-1962	90%	0%
1963-1983	64%	3%
1984-2002	34%	38%

**UC DAVIS**

# Racist Science in the 21st century

- **Reframing concepts and standards:** speculating about racial superiority and inferiority is not racism but exploring hypotheses about differences—or, data cannot be racist; repeating correlations is evidence of causation.
- **Challenging academic legitimacy and expertise:** attack academics as ideological to promulgate certainty about genetic causes of race and racial differences.
- **Building a cadre and a toolkit:** ideological foot soldiers armed with diverse race science materials from memes, to IQ classics, to cutting edge population genetics, from shitposting on Reddit, to commenting on New York Times articles, to publishing in scientific journals.

# Admixture regressions as a burgeoning method in race science



Open Access Article

## Global Ancestry and Cognitive Ability

by Jordan Lasker<sup>1</sup> , Bryan J. Pesta<sup>2,\*</sup> , John G. R. Fuerst<sup>3</sup> and Emil O. W. Kirkegaard<sup>4</sup>

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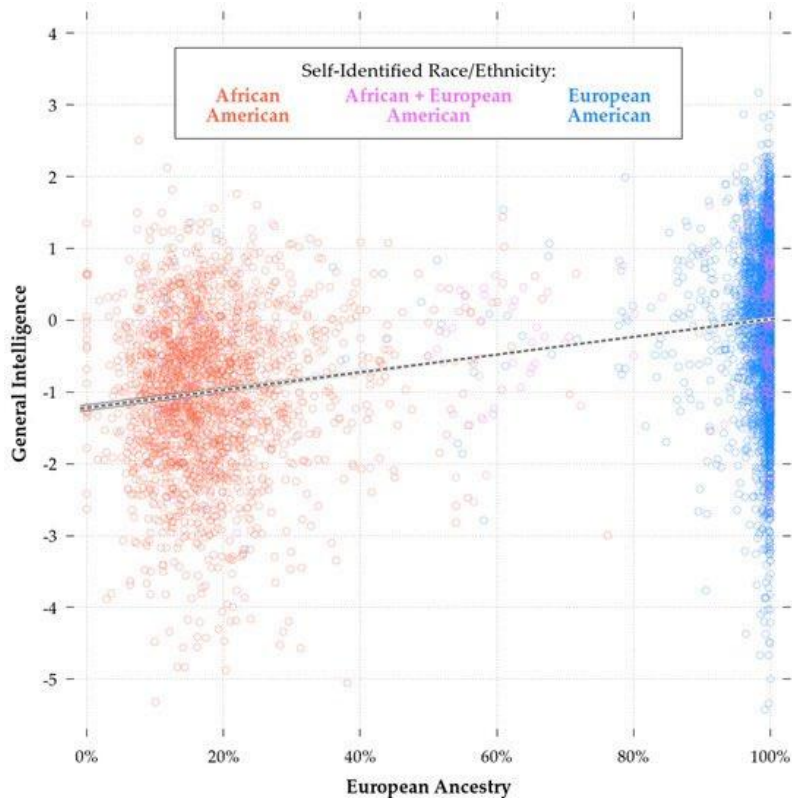
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# Admixture regressions as a burgeoning method in race science



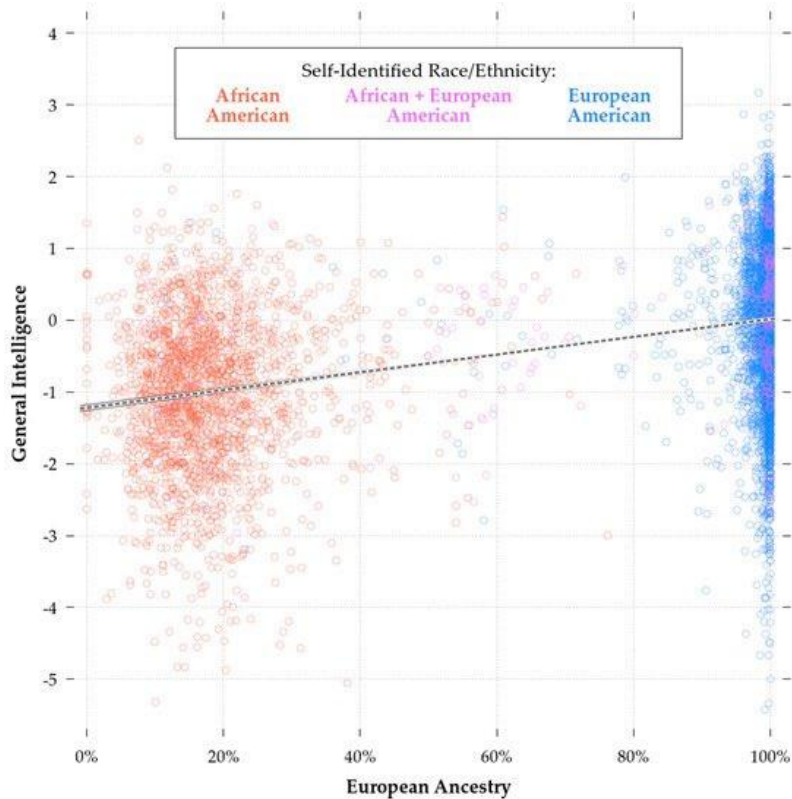
“landmark 2019 study... demonstrating a linkage between intelligence and racial admixture as measured by DNA.”  
- Steve Sailer, Unz Review Oct 13th, 2022

# Admixture regressions as a burgeoning method in race science

Race science proponents have created their own whig history where everyone agreed admixture correlation was a dispositive method



# Admixture regressions as a burgeoning method in race science



Obvious issue with confounding!



# Problems with admixture regression: confounding

Schraiber and Edge (2024) recently formally derived a formula for expected genetic variation of a trait due to variation in global genetic ancestry

$$\mathbb{E}(Y_i | \theta_i) = \mu_1 + \theta_i \delta_G + h_E(\theta),$$

Expected trait value of individual with given ancestry proportion

mean genetic contribution to the trait in population 1

Ancestry fraction

Mean difference in genetic value between populations

Function of how environmental effect depends on an individual's global ancestry fraction



# Problems with admixture regression: confounding

If environmental and genetic effects of global ancestry are confounded we can't recover  $\delta_G$  and regressions slopes represent some combination of environmental & genetic effects

$$\mathbb{E}(Y_i | \theta_i) = \mu_1 + \theta_i \delta_G + h_E(\theta),$$

Expected trait value of individual with given ancestry proportion

mean genetic contribution to the trait in population 1

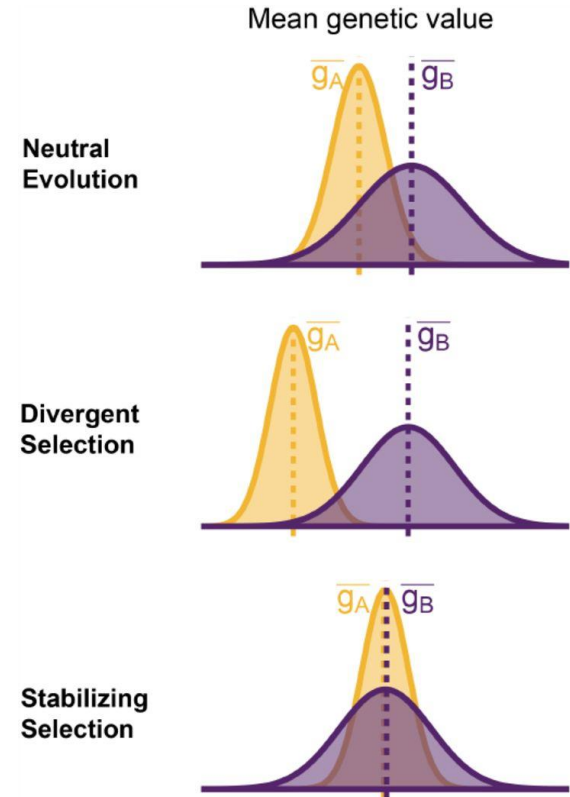
Ancestry fraction

Mean difference in genetic value between populations

Function of how environmental effect depends on an individual's global ancestry fraction

# Expectations of $\delta_G$ vary under different evolutionary scenarios

1. Neutral evolution: Differences in mean genetic value are expected to be small ( $\cong F_{st}$ ), can favor either population
2. Directional selection: Expect larger differences in mean genetic value compared to drift
3. Stabilizing selection: Expect smaller difference in mean genetic value compared to drift

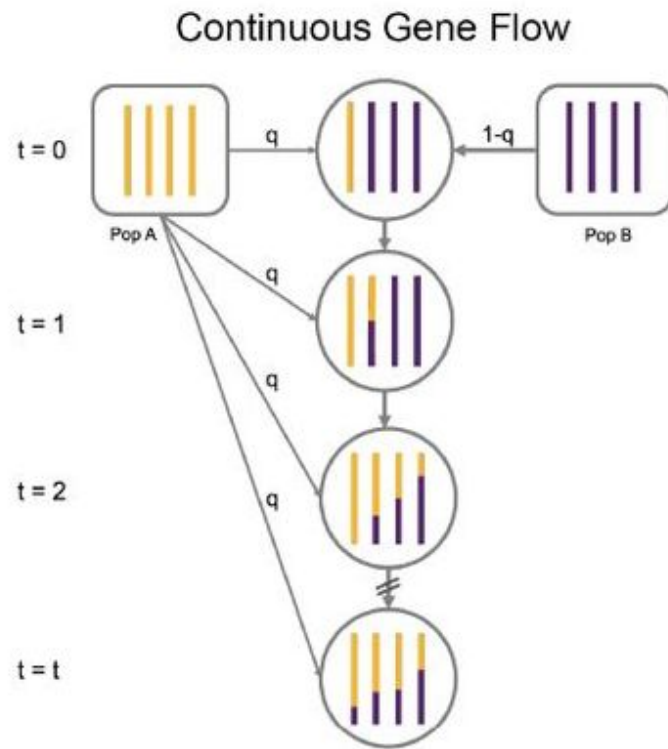


# Explore limitations and behavior of trait-ancestry correlations

- What is ancestry-trait correlation under different evolutionary scenarios?
- How much can ancestry-trait correlations be affected by environmental confounding?
- How effective is using a proxy variable as a statistical control?

# Methods:

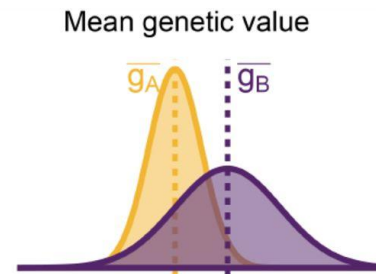
- Simulate an admixed population with variable ancestry fractions
  - 10,000 individuals, across 20 generations. Replicated 10 times



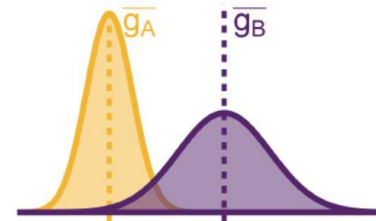
# Methods:

- Simulate an admixed population with variable ancestry fractions
  - 10,000 individuals, across 20 generations. Replicated 10 times
- Simulate genetic values (polygenic scores) for individuals under different evolutionary scenarios

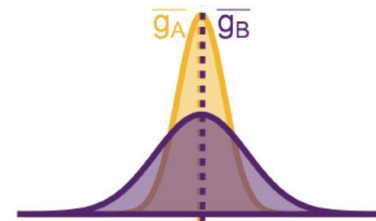
Neutral Evolution



Divergent Selection



Stabilizing Selection



# Methods:

Simulate trait values ( $y$ ) based on :

- $g_i$  - Individual's genetic value
- $\theta_i$  - Individual's ancestry fraction
- $a$  - slope between environment and ancestry
- $e$  - random noise

$$y_i = g_i + a * \theta_i + e$$

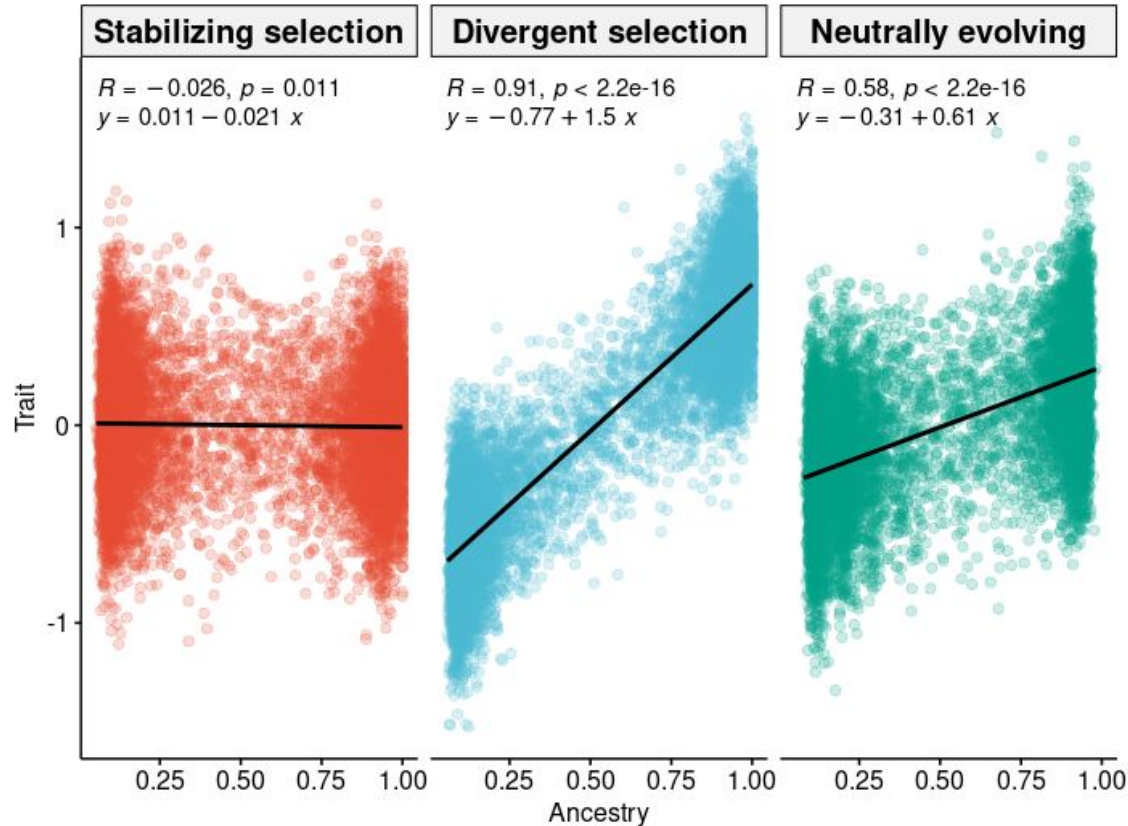


# Questions:

- **What is ancestry-trait correlation under different evolutionary scenarios?**
- How much can ancestry-trait correlations be affected by environmental confounding?
- How effective is using a proxy variable as a statistical control?



# Selection regime affects ancestry-trait correlations



\*All results presented are for generation 20

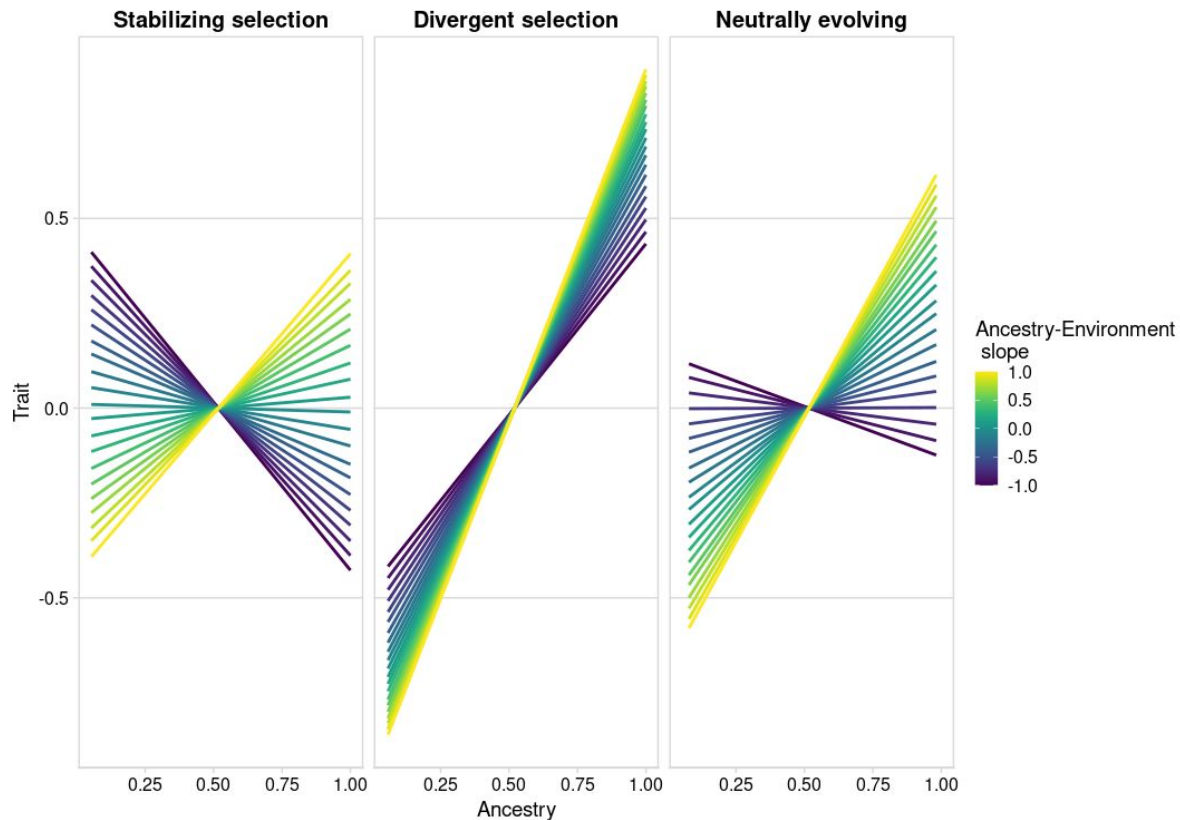
# Questions:

- What is ancestry-trait correlation under different evolutionary scenarios?
- **How much can ancestry-trait correlations be affected by environmental confounding?**
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# Effect of environmental confounding?

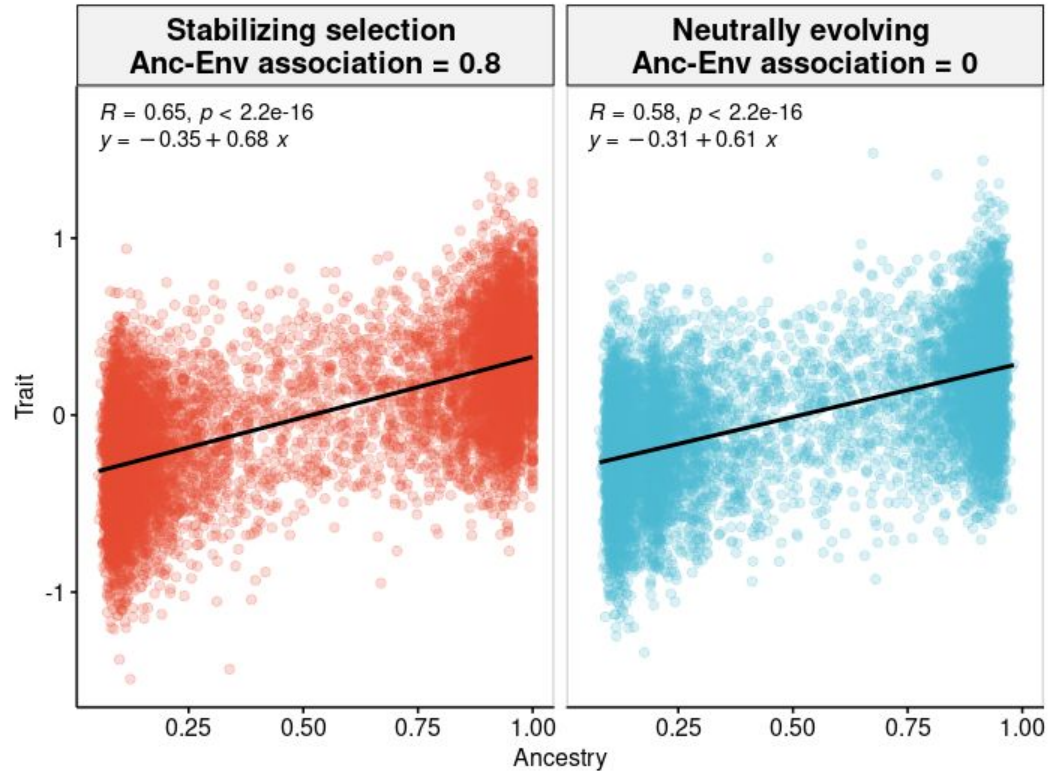
Environmental effects can change magnitude and direction of ancestry-trait correlation

Weaker effect under divergent selection



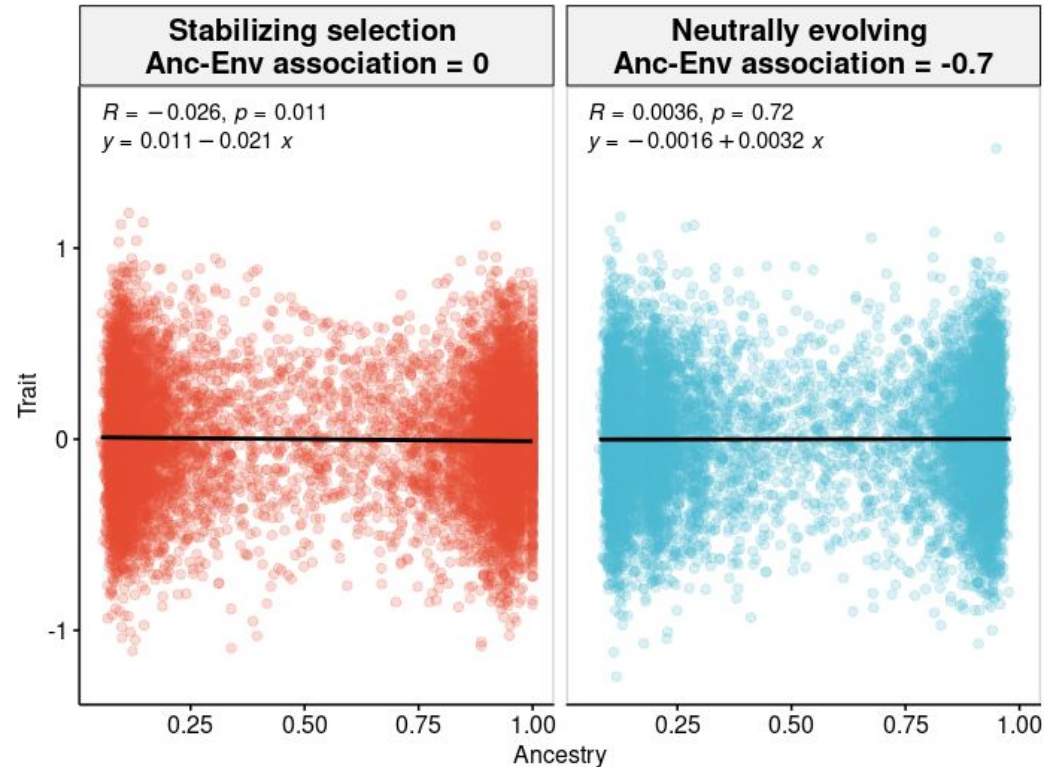
# Multiple scenarios produce identical ancestry-trait correlations

Ancestry-trait correlation that is entirely driven by environmental effects (left) is indistinguishable from an ancestry-trait correlation driven by neutral genetic differences and no environmental effects (right)



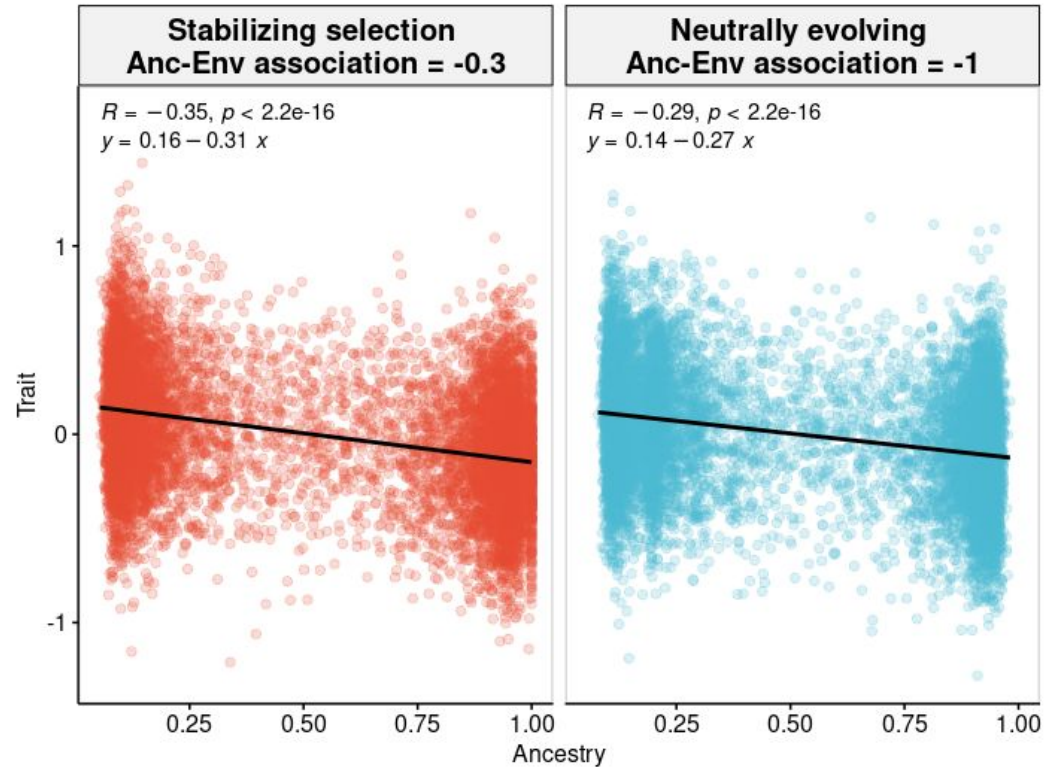
# Multiple scenarios produce identical ancestry-trait correlations

A true Ancestry-trait correlation of zero (left) is indistinguishable from a case where a true positive ancestry-trait correlation exists but is counteracted by a negative environmental effect (right)



# Multiple scenarios produce identical ancestry-trait correlations

Environmental effects of different strength can create nearly identical negative slopes in cases where no correlation exists (left) or when the correlation should be positive (right)



# Questions:

- What is ancestry-trait correlation under different evolutionary scenarios?
- How much can ancestry-trait correlations be affected by environmental confounding?
- **How effective is using a proxy variable as a statistical control?**



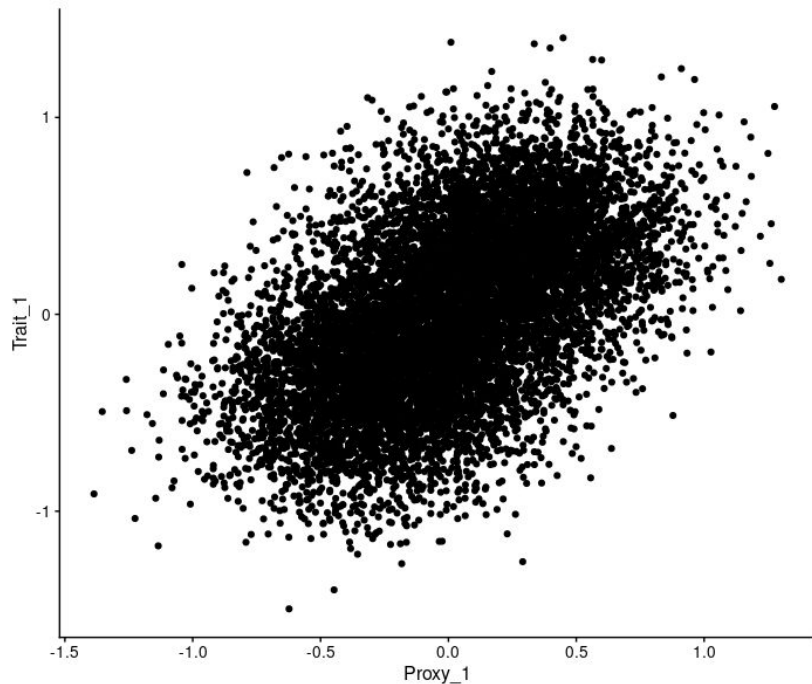
# How effective is using a proxy variable as a statistical control?

- Racial hereditarian researchers often include variables related to socioeconomic status (parental education, self-reported household income, etc)
- Claim the failure of such controls to fully attenuate ancestry-trait associations is evidence there is a true genetic effect

# How effective is using a proxy variable as a statistical control?

Simulate a noisy proxy trait that is partially correlated ( $r=0.5$ ) with actual environmental effects.

Include proxy trait as covariate in scenario where a strong ancestry-trait correlation is completely spurious.



# Statistical controls only partially attenuate spurious correlations

## Regression model without control

```
lm(formula = Trait_1 ~ Ancestry, data = .)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.16284	-0.20275	-0.00116	0.20038	1.10251

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-0.436678	0.005032	-86.78	<2e-16 ***
Ancestry	0.843741	0.007792	108.28	<2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.3009 on 9998 degrees of freedom

Multiple R-squared: 0.5398, Adjusted R-squared: 0.5397

F-statistic: 1.173e+04 on 1 and 9998 DF, p-value: < 2.2e-16

# Statistical controls only partially attenuate spurious correlations

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## Regression model with control

```
lm(formula = Trait_1 ~ Ancestry + Proxy_1, data = .)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.11126	-0.20082	-0.00159	0.20089	1.07666

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-0.391343	0.005917	-66.14	<2e-16 ***
Ancestry	0.756145	0.009876	76.56	<2e-16 ***
Proxy_1	0.138260	0.009733	14.21	<2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.298 on 9997 degrees of freedom  
Multiple R-squared: 0.5489, Adjusted R-squared: 0.5488  
F-statistic: 6081 on 2 and 9997 DF, p-value: < 2.2e-16

# Conclusions

- Ancestry-environment correlations are uninterpretable without additional knowledge of evolutionary dynamics and environmental effects.
- Environmental variables that only partially capture true environmental effects will be unable to fully attenuate even completely spurious ancestry-trait correlations
  - Showing that ancestry-trait associations survive controls for SES can't serve as evidence that a true genetic effect exists.