

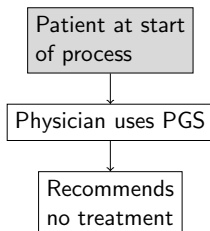
# Precision Medicine and Black-White Health Disparities

Rebecca Johnson<sup>1</sup>, Sam Trejo<sup>2</sup>, Jeremy Freese<sup>2</sup>, Ben Domingue<sup>2</sup>

1. Princeton University; 2. Stanford University

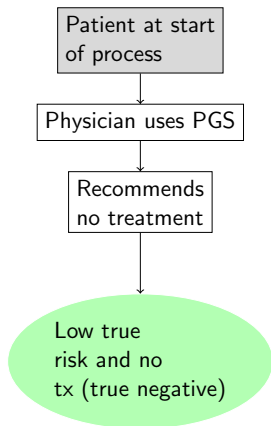
IGSS 2019

Motivation: past research focuses on how clinical use of polygenic scores (PGS) may widen Black-White disparities through mechanism of unequal accuracy



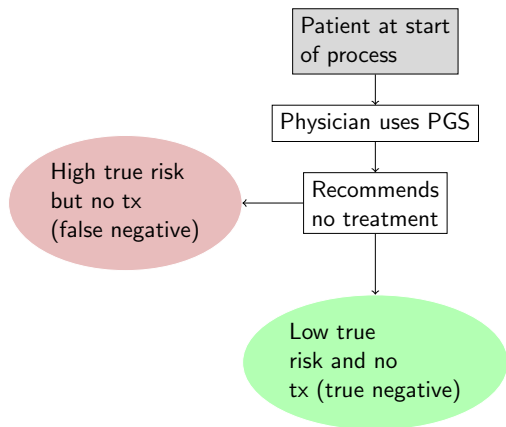
Duncan et al. (2019); Martin et al. (2019) Mills and Rahal (2019); Mostafavi et al. (2019)

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# Gaps and present study

1. **Unequal accuracy:**

high true risk

but not given high-risk  
label (false negative)

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but no scoring by PGS  
(false negative)

## 1. **Unequal accuracy:**

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## 2. **Unequal disease mgmt.:**

accurate high-risk label,  
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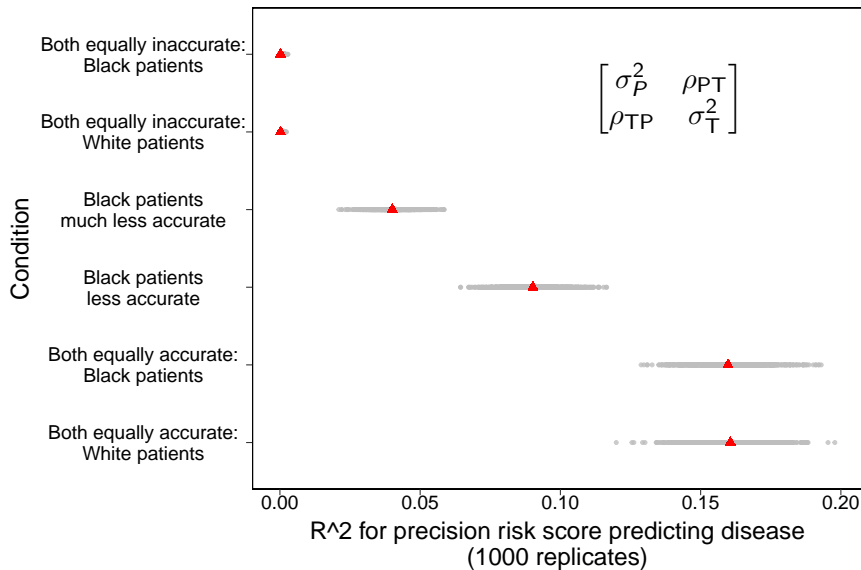
2. **Unequal disease mgmt.:**  
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## **Present study:**

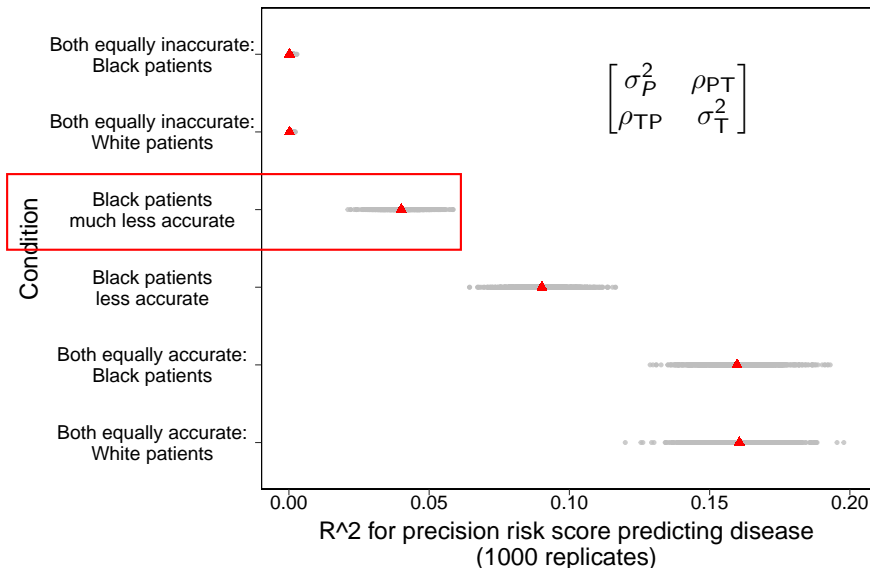
use agent-based modeling (ABM),  
informed by HRS data;  
start with false negatives due  
to unequal accuracy as baseline;  
investigate *relative magnitude*  
of unequal accuracy as mechanism for disparities  
vs. other mechanisms that persist  
after equalizing accuracy



Simulating unequal accuracy ( $T$ : true disease risk;  $P$ : measured PGS)



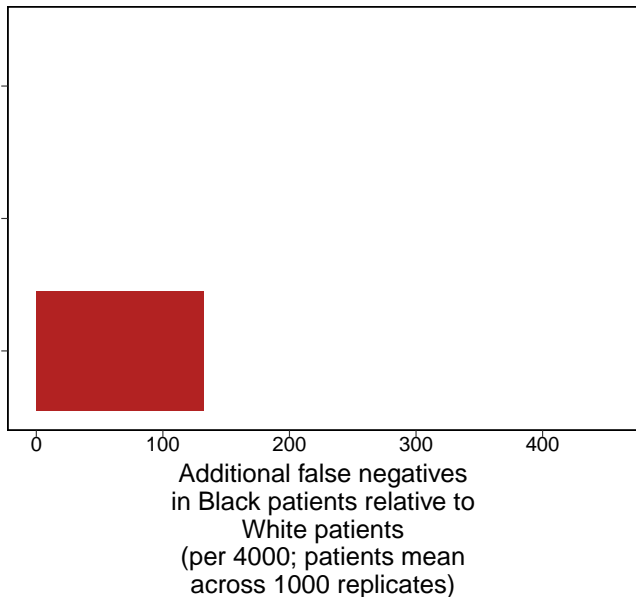
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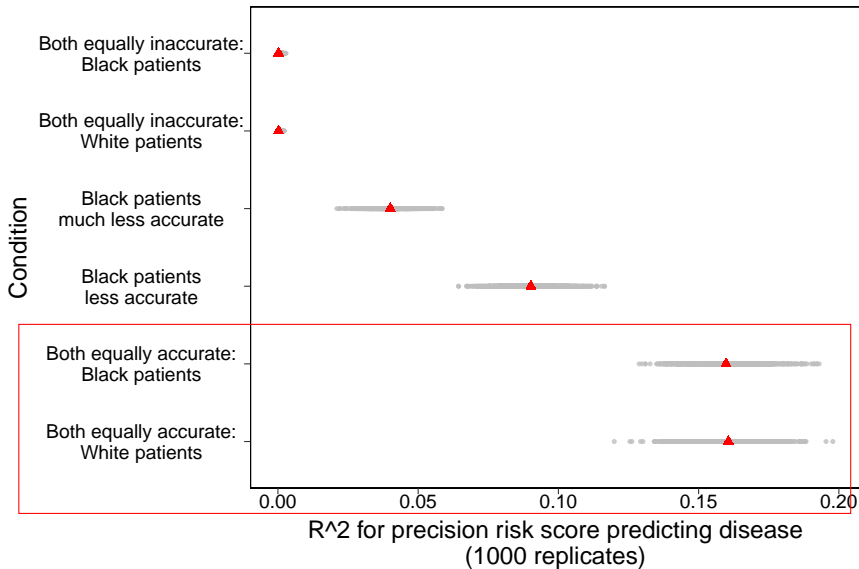
Equal accuracy and care,  
but unequal prevention

Equal accuracy  
but unequal care

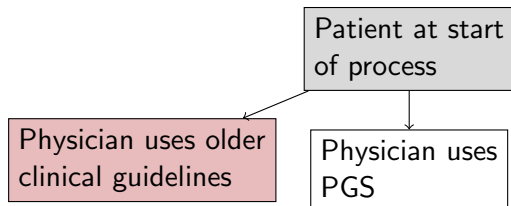
Unequal accuracy  
(1/4th as accurate)



# Now suppose we equalize accuracy...



## Two additional mechanisms

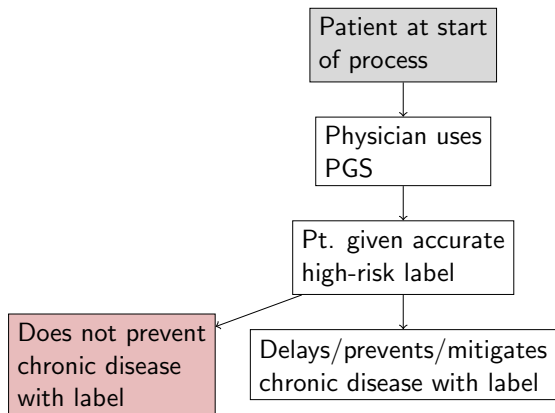


Parameters:

1. Redundancy between PGS and existing risk assess. methods (less redundant; more it matters)
2. How race affects Doctors using new tech./evidence-based care

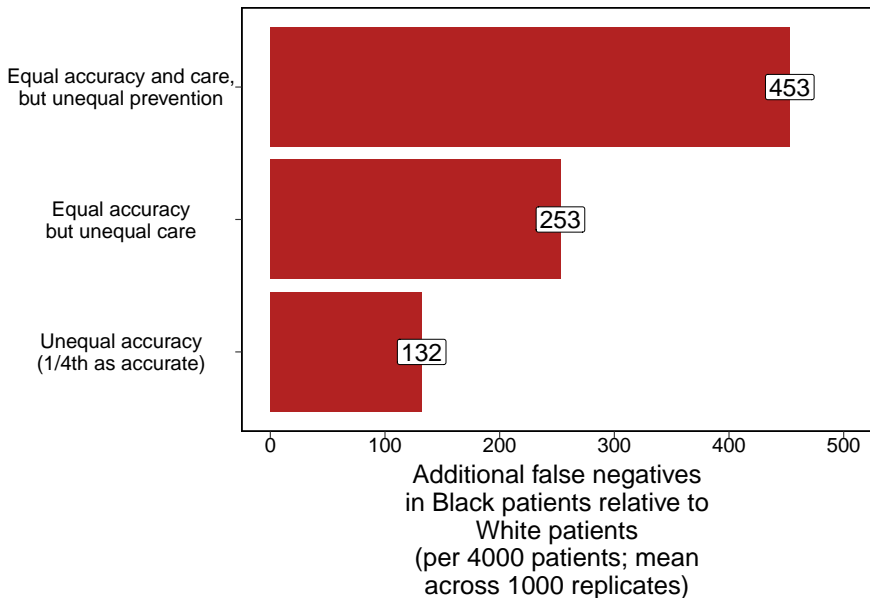
Torkamani et al. 2018; Khera et al. 2018

## Two additional mechanisms



Parameters:

1. Correlation between race and SES
2. Disparities in chronic disease management among those at high risk



Future directions: improve simulation + supplement work on improving accuracy disparities with work investigating other mechanisms

<b>Mechanism</b>	<b>Example research</b>
Whose care features newer risk assessment methods?	$Pr(\text{Dr. offers})$ ; $Pr(\text{Pt. accepts} \text{Dr. offers})$ (how much does the selection into genotyping in pop. cohort studies—White, healthier, higher-SES—generalize to clinical contexts?)
How does disclosure of polygenic risk affect health behaviors?	RCTs of candidate gene disclosure (e.g., Green et al. ApoE4 REVEAL study)...similar RCTs for PGS? How does interpretability of risk info. influence its impact?



Thanks!

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[https://github.com/rebeccajohnson88/precisionmed\\_disparities](https://github.com/rebeccajohnson88/precisionmed_disparities)