Genetic Heterogeneity in Vulnerability to Depression During Changing Economic Conditions

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# Abstract

# Introduction

This study tests the genetic vulnerability or resistance to depressive events in the Health and Retirement Study (HRS). We approach this in the framework of the diathesis-stress model, which suggests excess genes confer excess vulnerability to negative outcomes when exposed to stress. We also test an opposing theory, that genees confer excess resilience ot negative outcomes when exposed to stress. Combining data from the Health and Retirement Study (HRS) with information on regional unemployment rates from the Beureau of Labor Statistics (BLS), we test sensitivities to changing economic conditions between 2007 and 2012 as changes in depressive symptoms. Then, using a polygenic score from a large published Genome Wide Association Study (GWAS), we test whether individuals’ reaction to the changing economic conditions vary with polygenic risk scores (PGS). Using white non-hispanic adults with genetic data in the HRS (n=9,946), we find (1) individuals with higher subjective well-being PGS have lower levels of depressive symptoms, but (2) individuals with higher subjective well-being PGS are more sensitive to changing economic conditions, as measured by the unemployment rate.

# Background

The stress process model has been used to explain the development of all sorts of adverse health events–including continuous measures of psychological distress and depressive symptoms which do not reach clinically diagnosable levels (Gayman and Barragan 2013). In its most basic formulation, the stress process model suggests that organisms respond negatively to negative stimuli from the environment (Pearlin et al. 1981). Heterogeneity in negative responses depend on mechanisms that can either buffer or exacerbate the stress response. Studies have found that economic conditions related to “job churn” generate poor health outcomes (Strully 2009). Importantly, poor economic conditions associated with poor economic conditions, like rising unemplyment rates, are associated with large-scale changes in mental health and psychological well-being (Lam, Fan, and Moen 2014).

This study asks whether there is variation in the response to these generalized stressful conditions is associated with differnces in genetic risk scores. The theory of genetic hetergoeneity in vulnerability to stress response is known as the diathesis-stress model (Domingue et al. 2016). While controversial, studies have found evidence of genetic vulnerability to stress, such as .

Hypothesis 1:

Hypothesis 2:

Hypothesis 3: survival: if these models pan out

# Data

## Polygenic Risk Scores

We follow the procedure outlined in Dominigue, et al. (2016) to calculate our polygenic risk scores. First, we match SNPs in the HRS to SNPS in the GWAS conducted by Okbay et al. (2016) available through the Social Science Genetic Association Consortium. Second, we sum the weighted estimates of well-being associated alleles from the GWAS to calculate a polygenic score. Third, we adjust the residulized PGS using 10 principal components and PLINK (command: pca), and normalize the PGS residual distribution, which we use as the measure of genetic predisposition to well-being.

# Analytic Strategy

Treating changes in unemployment rates as exogenous, we employ a standard fixed effects design to test our environemnetal models (Conley 2009). We then interact environmental conditions with the PGS. We do not include a direct effect of the PGS in our models because it is not identified.

Environment Model:

Interaction Model:

Where indicates the outcome, indexes individuals, and indexes time, is various time-varying covariates as described above, and is the PGS calculated as described above.

# Results

see

# Discussion and Conclusion

# References

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