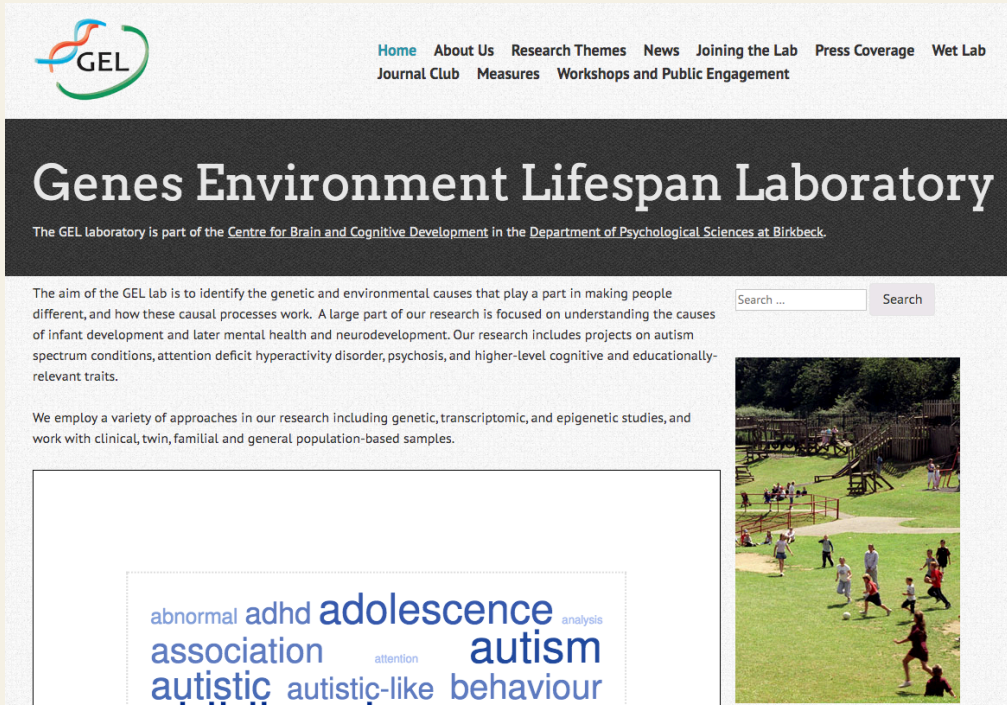


The first 10 years and beyond....

CBCD 20th Anniversary

Angelica Ronald

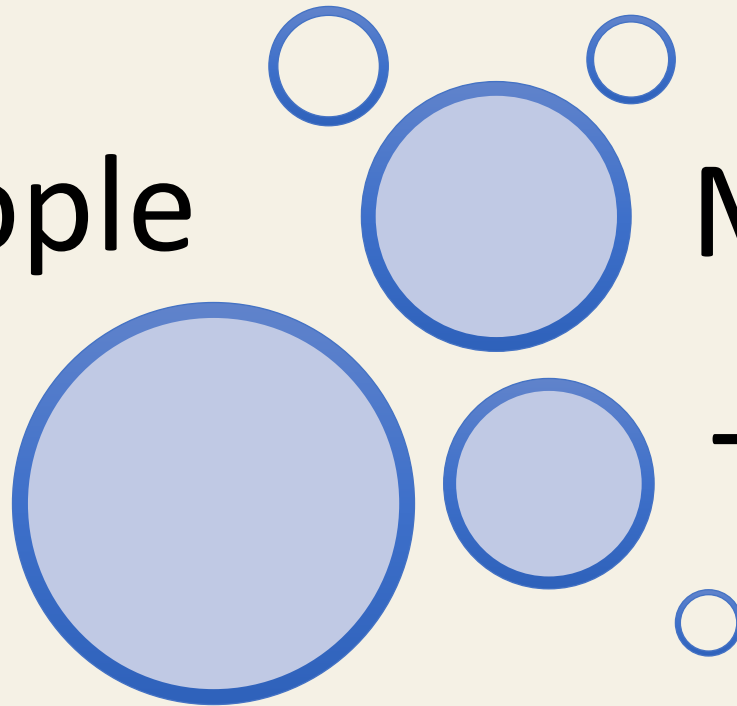
Joined CBCD in 2007



The screenshot shows the homepage of the Genes Environment Lifespan Laboratory (GEL). At the top left is the GEL logo, which consists of a stylized DNA double helix in blue and red, with the letters 'GEL' in green. To the right of the logo is a navigation menu with links: Home, About Us, Research Themes, News, Joining the Lab, Press Coverage, Wet Lab, Journal Club, Measures, and Workshops and Public Engagement. Below the navigation menu is a dark grey banner with the text 'Genes Environment Lifespan Laboratory' in white. Underneath the banner is a sub-header: 'The GEL laboratory is part of the Centre for Brain and Cognitive Development in the Department of Psychological Sciences at Birkbeck'. The main content area has a light grey background. On the left, there is a paragraph of text describing the lab's aim: 'The aim of the GEL lab is to identify the genetic and environmental causes that play a part in making people different, and how these causal processes work. A large part of our research is focused on understanding the causes of infant development and later mental health and neurodevelopment. Our research includes projects on autism spectrum conditions, attention deficit hyperactivity disorder, psychosis, and higher-level cognitive and educationally-relevant traits.' Below this paragraph is another paragraph: 'We employ a variety of approaches in our research including genetic, transcriptomic, and epigenetic studies, and work with clinical, twin, familial and general population-based samples.' To the right of the text is a search bar with the placeholder text 'Search ...' and a 'Search' button. Below the search bar is a photograph of a group of children playing soccer on a grassy field. At the bottom left of the screenshot is a word cloud containing terms such as 'abnormal', 'adhd', 'adolescence', 'analysis', 'association', 'attention', 'autism', 'autistic', 'autistic-like', and 'behaviour'.



People



Methods

Themes



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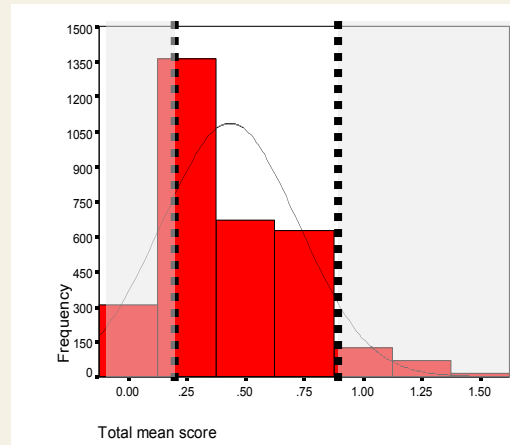
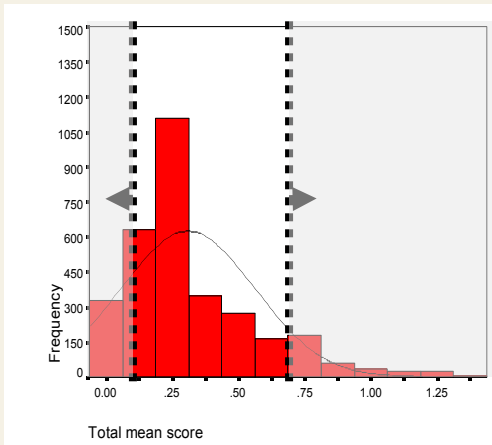
Back in 2007:

Behav Genet (2010) 40:31–45
DOI 10.1007/s10519-009-9308-6

ORIGINAL RESEARCH

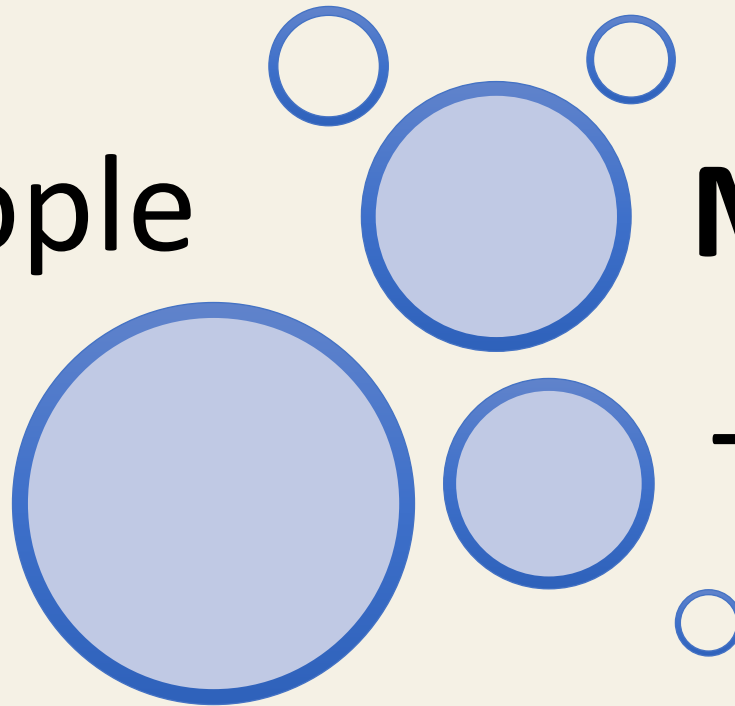
A Genome-Wide Association Study of Social and Non-Social Autistic-Like Traits in the General Population Using Pooled DNA, 500 K SNP Microarrays and Both Community and Diagnosed Autism Replication Samples

Angelica Ronald · Lee M. Butcher · Sophia Docherty ·
Oliver S. P. Davis · Leonard C. Schalkwyk ·
Ian W. Craig · Robert Plomin



Ronald et al (2010) *Behavior Genetics*

People



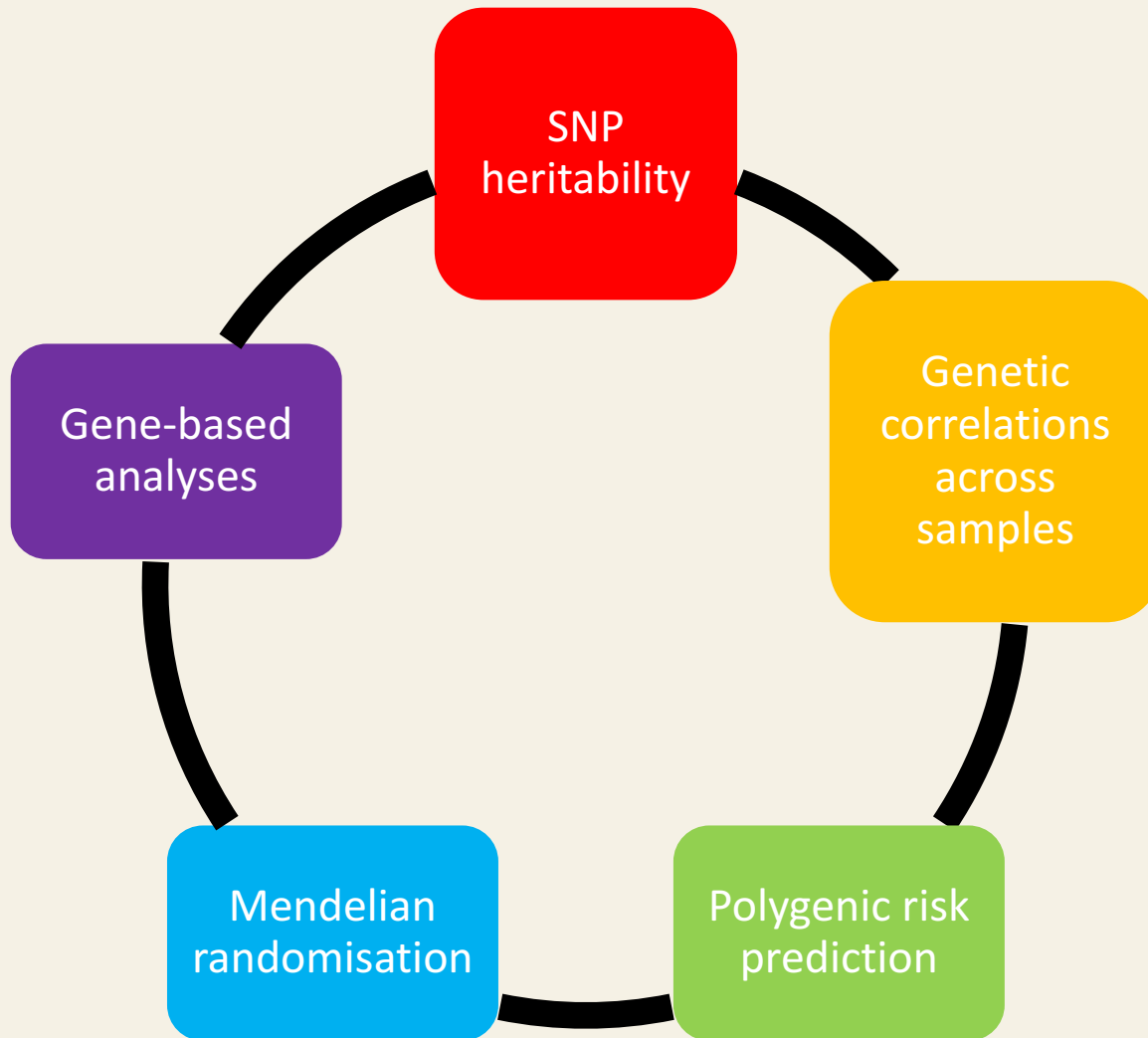
Methods

Themes



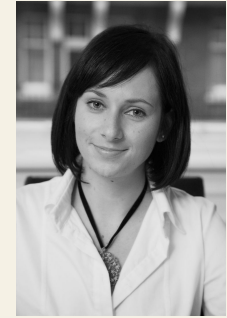
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Methods



SNP Heritability

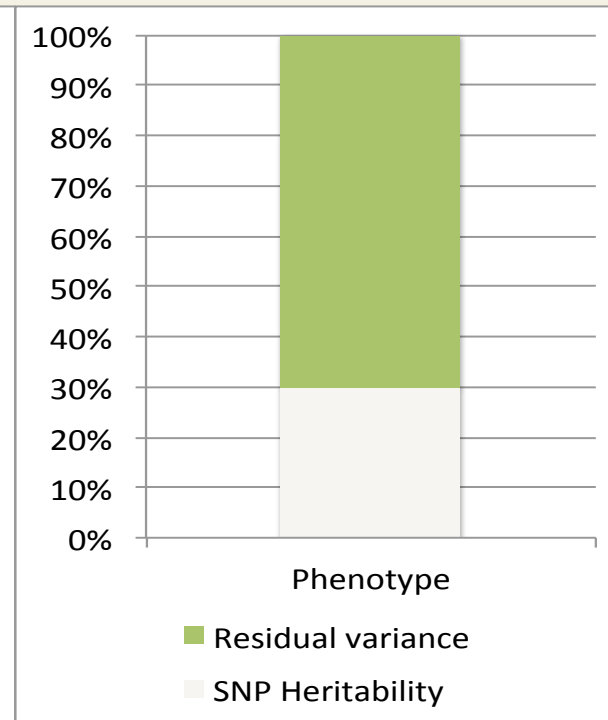
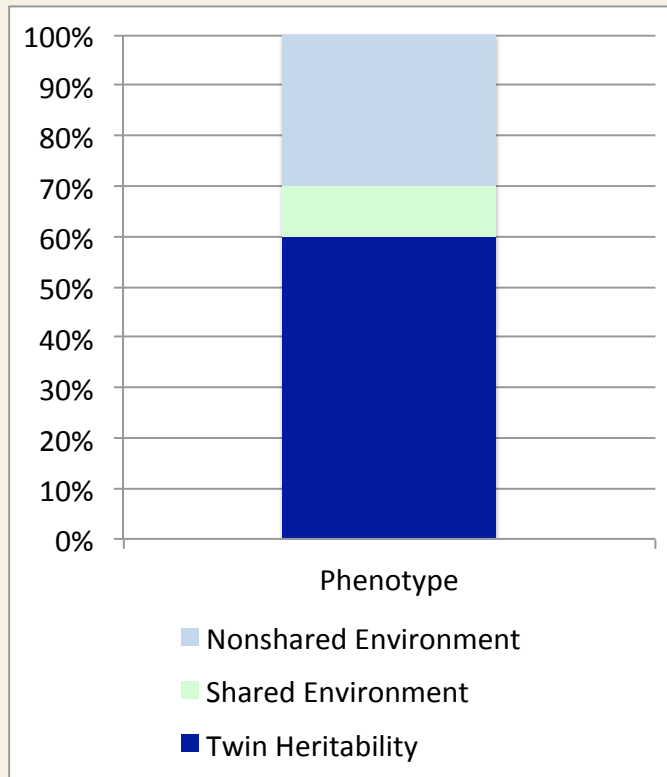
SNP; single nucleotide polymorphism



Dr Dominika Sieradzka



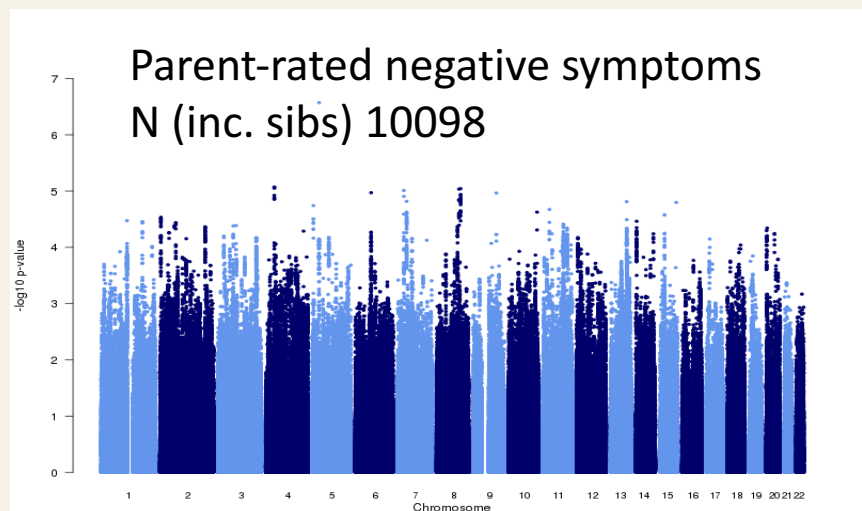
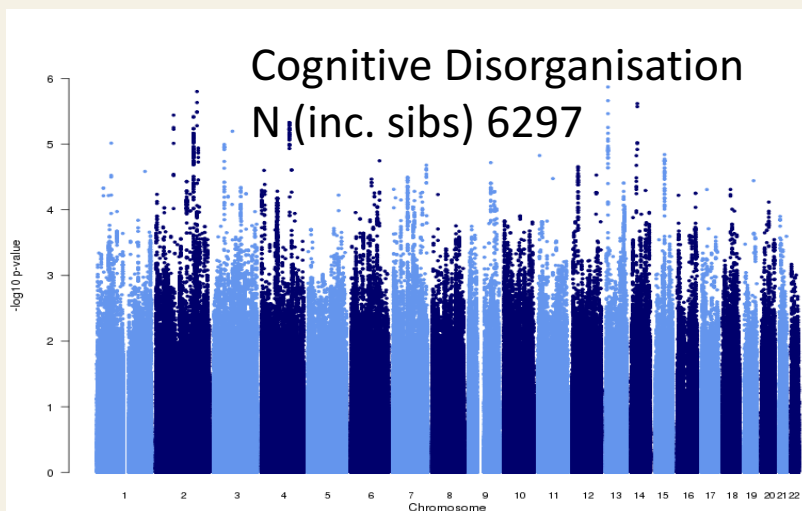
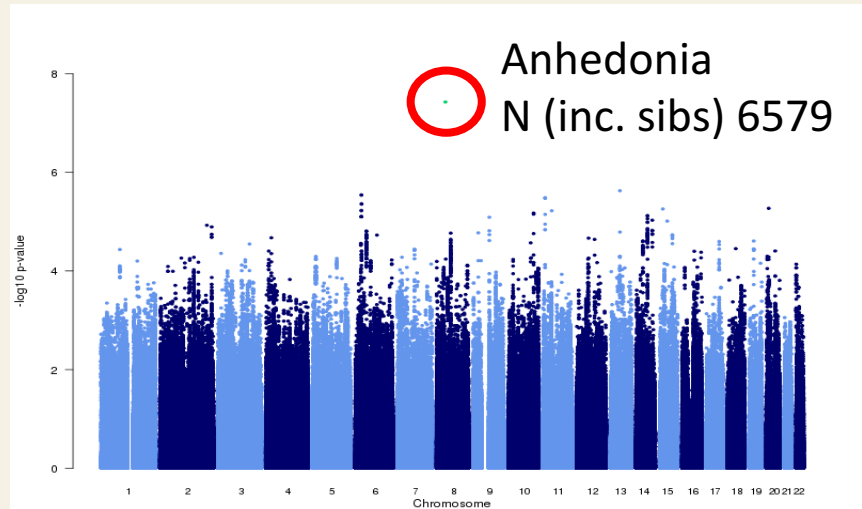
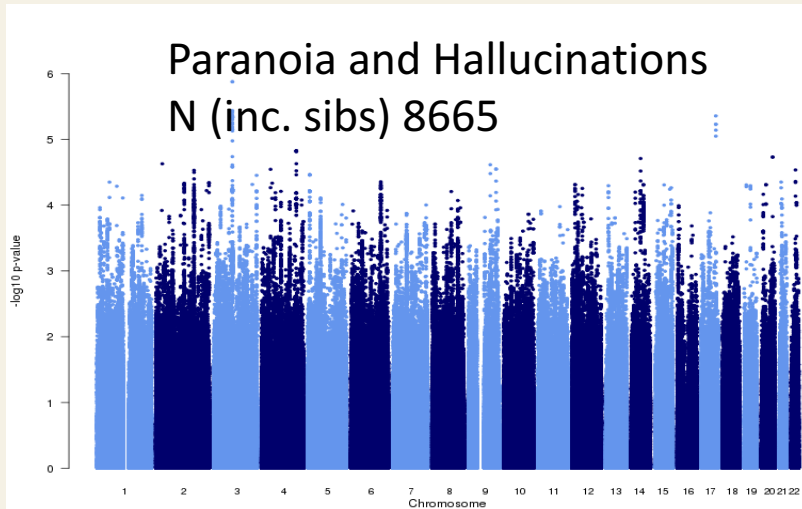
Dr Oliver Pain



Sieradzka et al 2015 *Behaviour Genetics*; Pain et al (2018) *Neuropsychiatric Genetics*

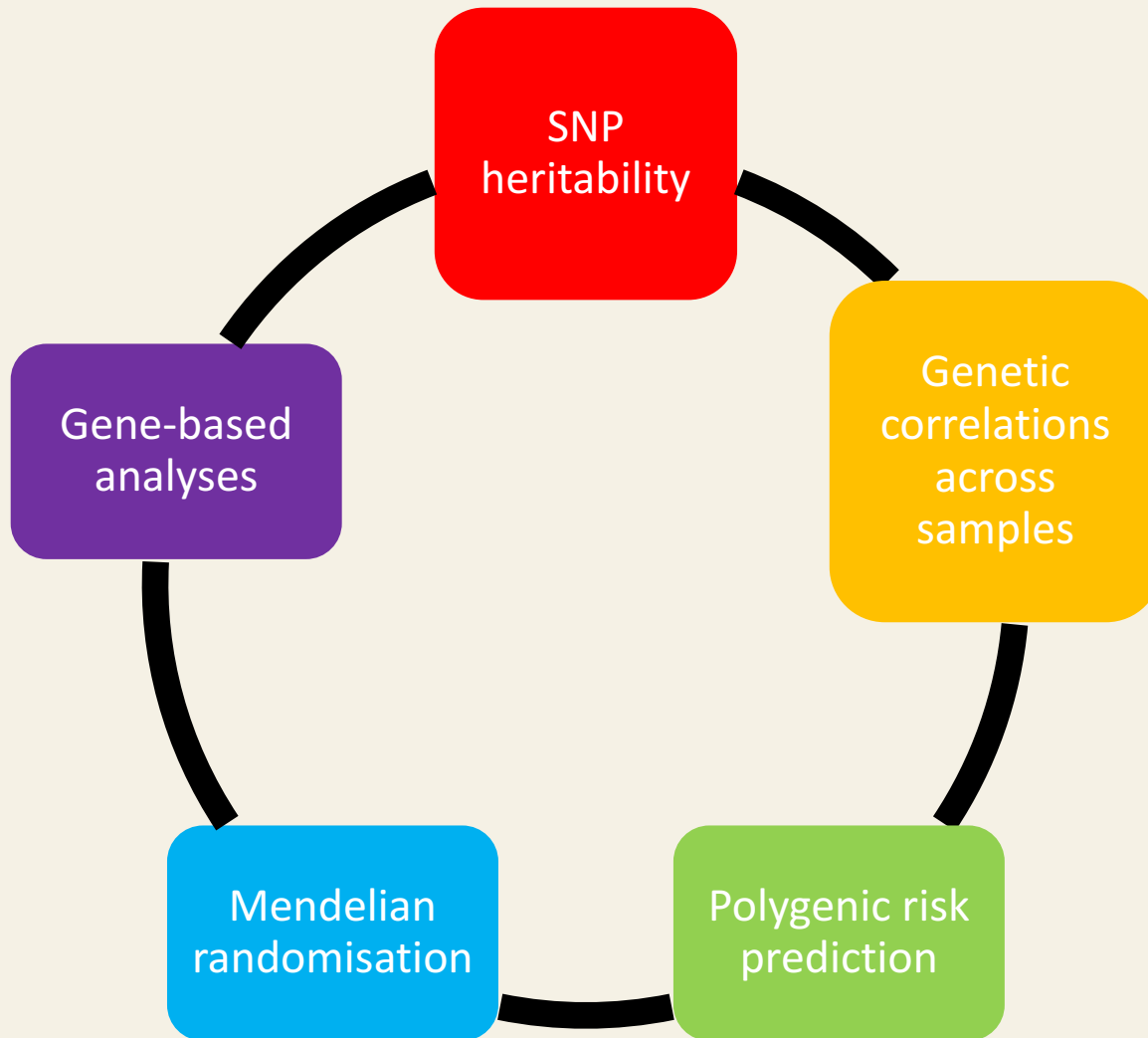
Method: Bulik-Sullivan et al (2015) *Nature Genetics*; Yang et al (2011) *Nature Genetics*

Gene discovery: Larger and cheaper



Pain, Dudbridge, Cardno, Freeman, Lu, Lundstrom, Lichtenstein, Ronald (2018) *Neuropsychiatric Genetics*

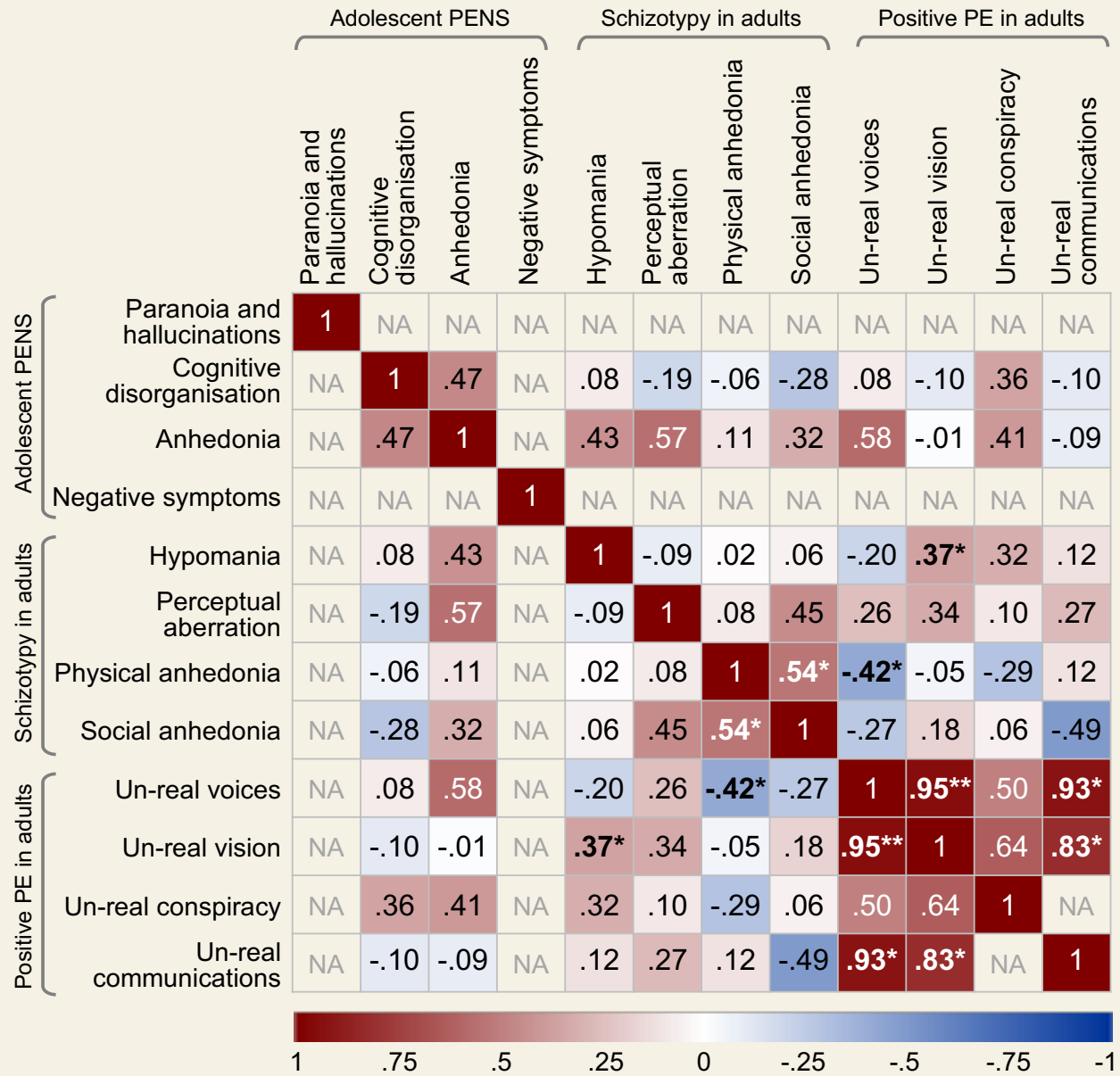
Methods





Wikus
Barkhuizen

Genetic correlations across samples!



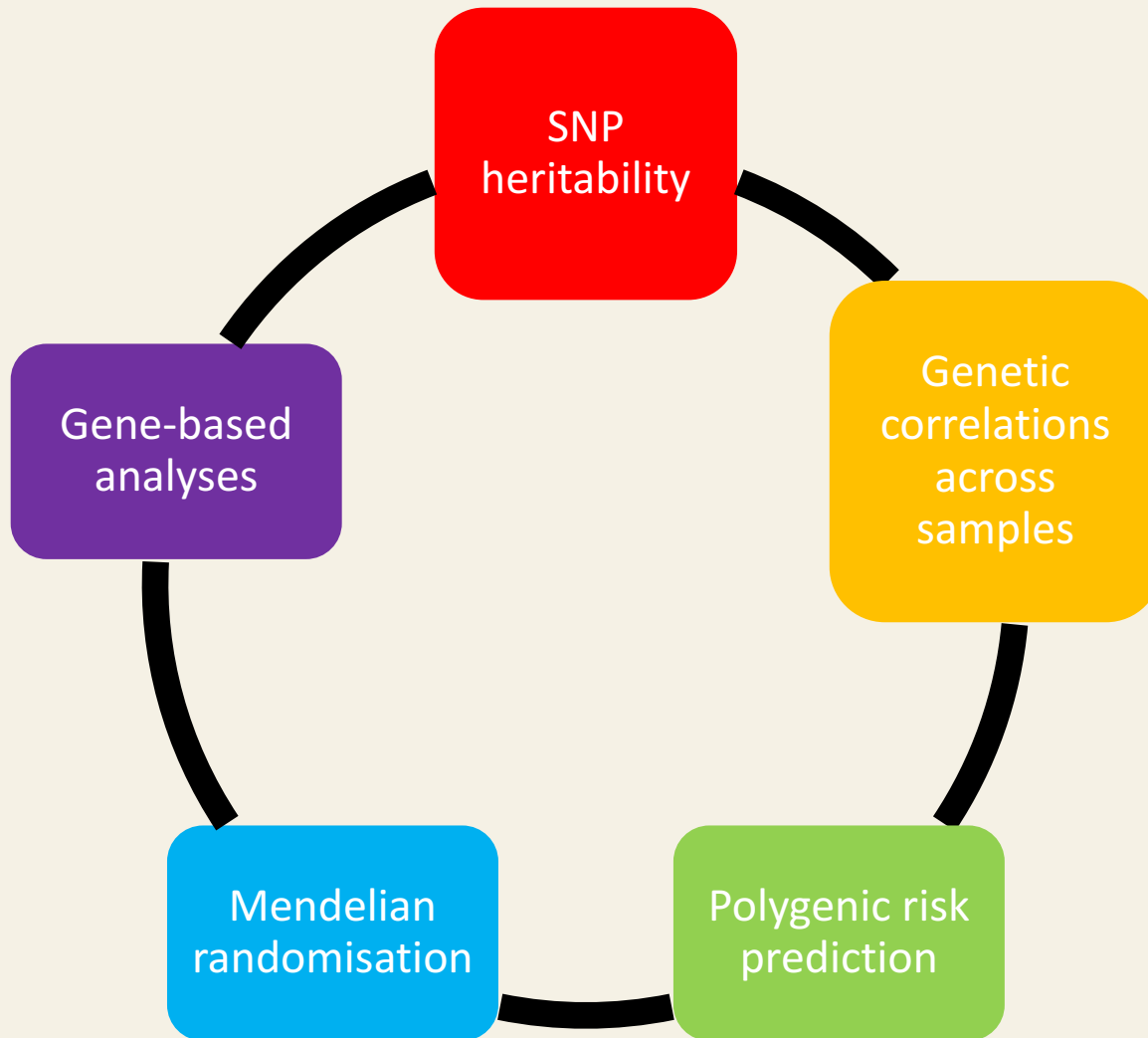
Barkhuizen, Pain, Dudbridge & Ronald (2019) BioRxiv Preprint

<https://www.biorxiv.org/content/10.1101/718015v1>

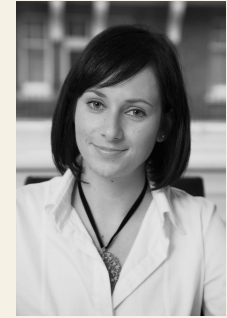
Method: Bulik-Sullivan et al (2015) *Nature Genetics*



Methods



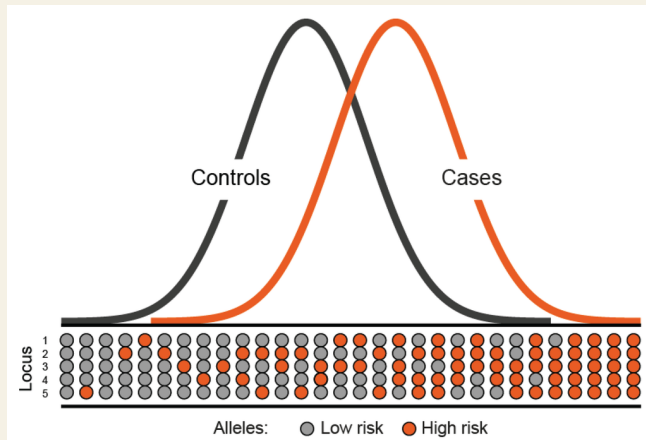
Polygenic Risk Score prediction



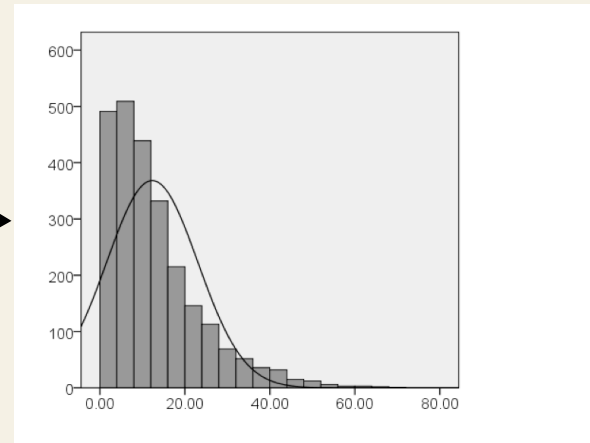
Dr Dominika Sieradzka



Dr Oliver Pain



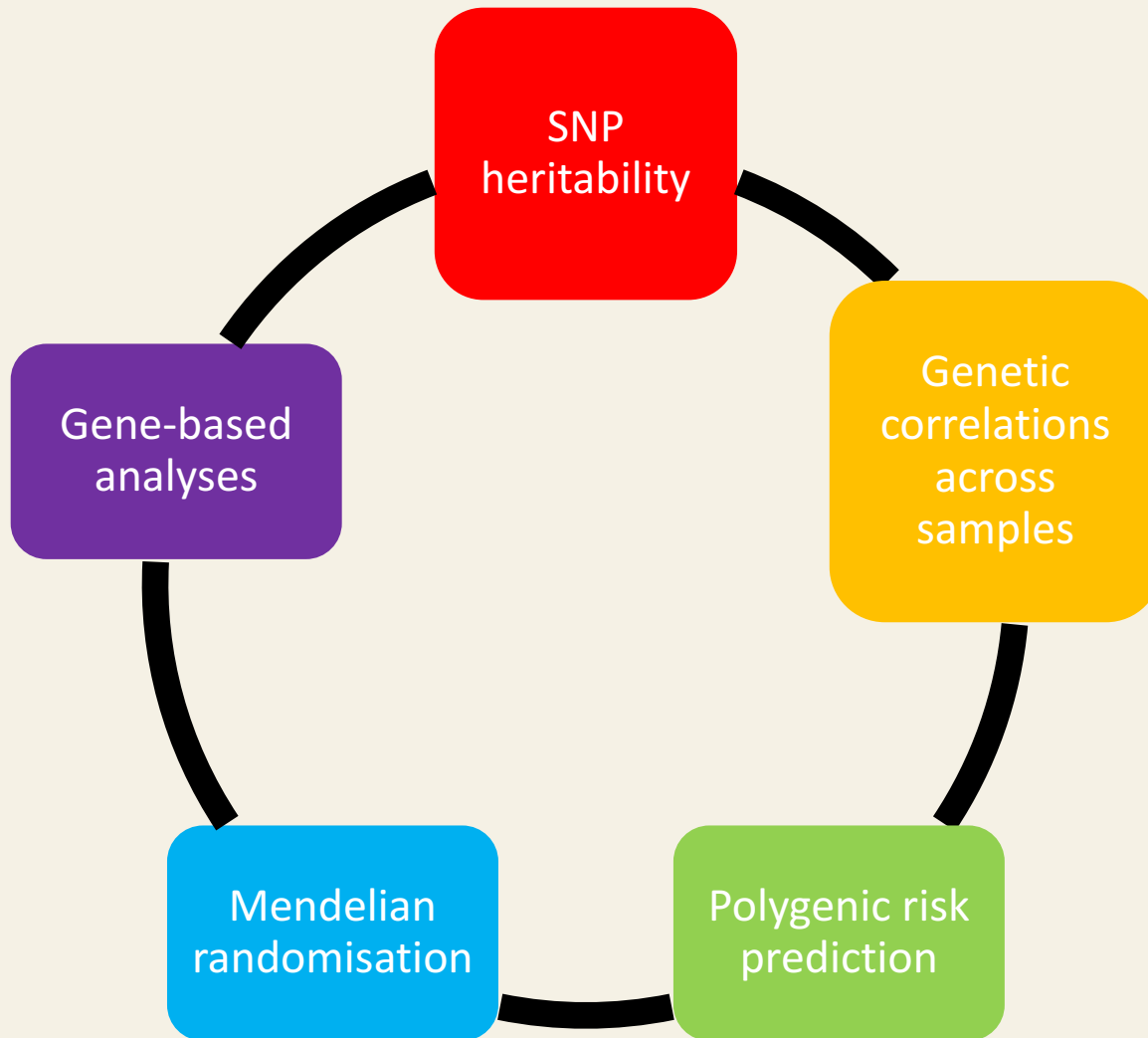
Polygenic risk score for each individual



Does the score predict a second phenotype

Ronald & Pain (2018) *Human Molecular Genetics*
Sieradzka, Power, Freeman, ... & Ronald (2014) *PLoS ONE*
Pain et al (2018) *Neuropsych Genetics*
Method: Purcell et al (2009)

Methods



Mendelian Randomisation

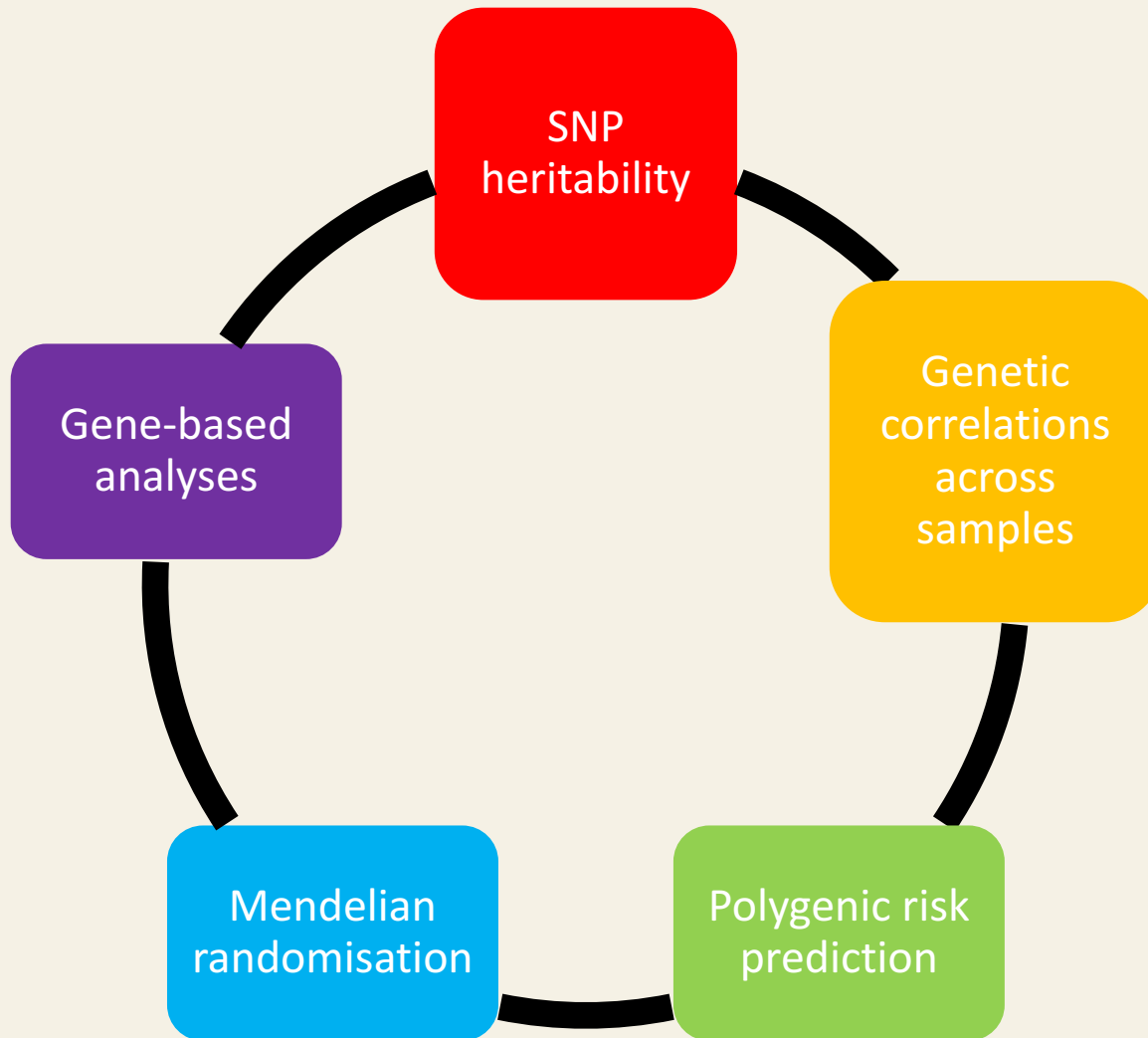


Wikus
Barkhuizen

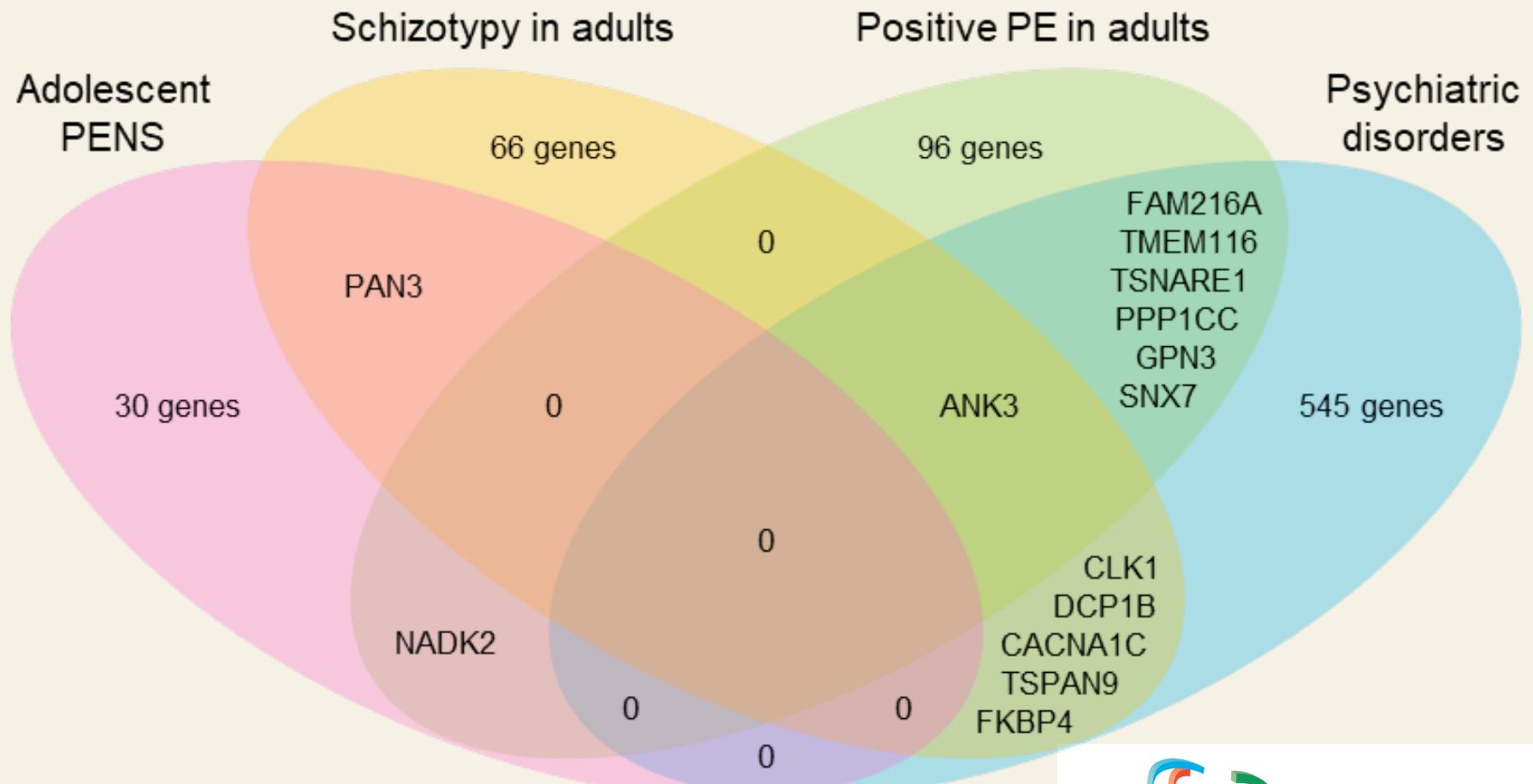


Barkhuizen, Dudbridge & Ronald (in preparation)

Methods



Gene-based analyses: Overlapping genes across age and with psychiatric disorders



Barkhuizen, Pain, Dudbridge & Ronald (2019) BioRxiv Preprint

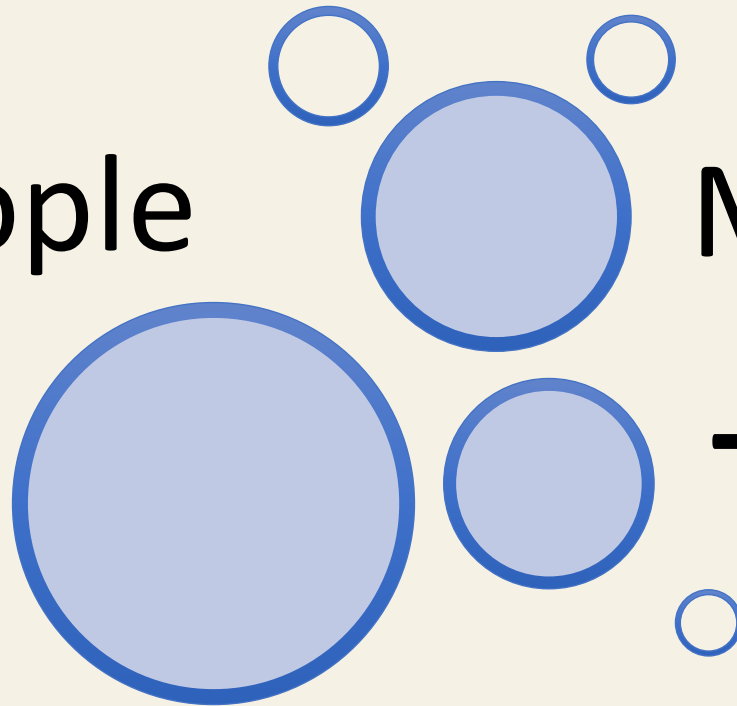
<https://www.biorxiv.org/content/10.1101/718015v1>

Method: Watanabe et al (2017) *Nature Communication*



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People



Methods

Themes



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Themes

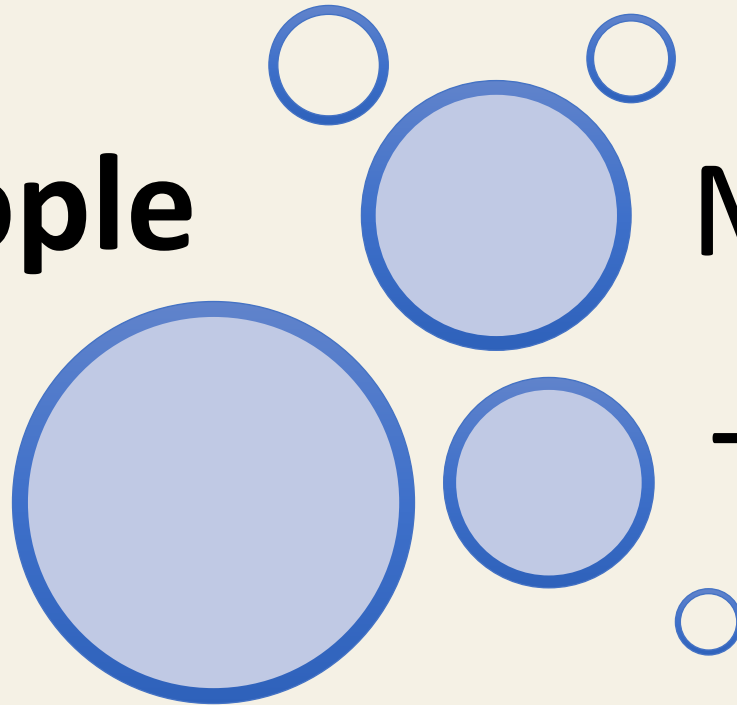


Dr Emma Meaburn, GEL lab
co-director



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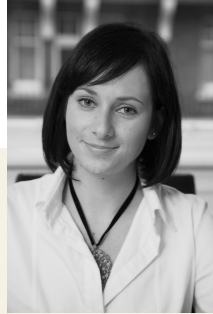
Genes Environment Lifespan
laboratory

★ Past & present postdocs and PhD students ★

Dr Sania Shakoor



Dr Kostas Papageorgiou



Dr Dominika Zavos Sieradzka



Dr Helena



Dr Charlotte Willfors



Dr Elise Robinson



Dr Oliver Pain

Dr Mark Taylor



Aislinn Bowler



Chloe Austerberry



Wikus Barkhuizen



Monica Siqueiros

Dr Aline Scherff



Dr Karla Holmboe



Dr Victoria Hallett



Laura Havers



Genes Environment Lifespan laboratory

Looking forward



Multi-method
study of
evocative parent-
child effects

Chloe Austerberry*

Longitudinal
infant twin
study in
Stockholm



Monica Siqueiros*



Stability and
change in
psychotic
experiences

Laura Havers*

iCASE PhD to
develop app for
cohort research



Aislinn Bowler

*See their posters today!



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Thank you

www.gel.bbk.ac.uk

Twitter @gelironald



Acknowledgements

Research Team:

Wikus Barkhuizen

Professor Frank Dudbridge

Dr Oliver Pain

Funding

Camara-Rivers David grant fund

UK Medical Research Council

Wellcome Trust ISSF fund

Summary statistics: Samples and Resources

ALSPAC, CATSS and TEDS samples

Neale Lab <http://www.nealelab.is/uk-biobank>

UK Biobank

North Finland Birth Cohort

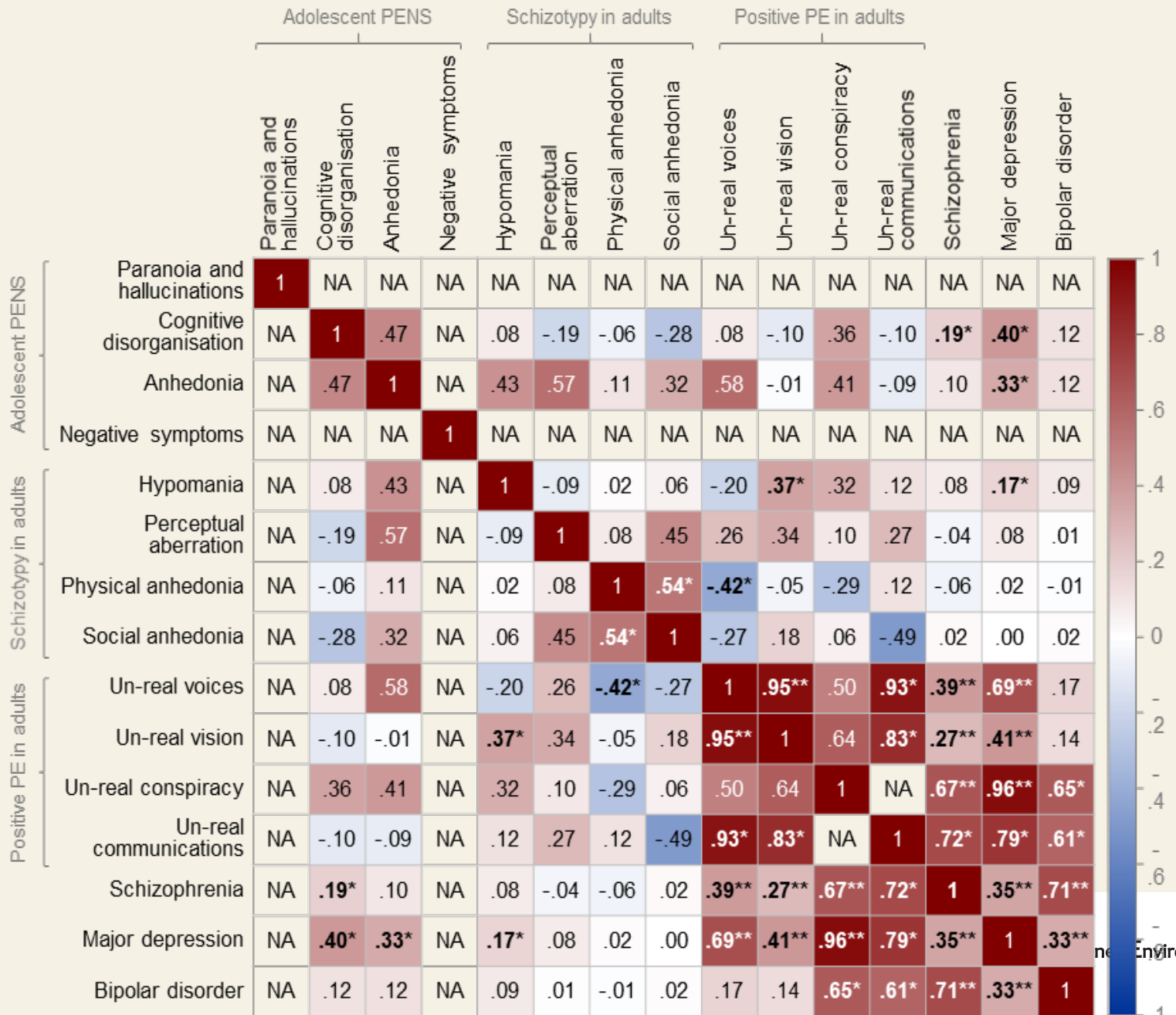
William Hennah and Alfreda Ortega-Alonso

Psychiatric Genomics Consortium
www.med.unc.edu/pgc/results-and-downloads/

All the cohort teams and participants

- Notes on WCPG Guidelines: -
- Contact Information on final slide
- Use common fonts: Times New Roman, Arial, and Courier.
- Save versions in both 4:3 and 16:9!
- BEFORE YOU GO TO YOUR TALK DO THE FOLLOWING:
- After you put together your presentation save it. Then:
 - a) Go to Design on the top Tab
 - b) Go to Slide Size and click on it
 - c) You will see the choice of 4:3 or 16:9
 - d) Whatever size it currently isn't on, change to that size
 - e) Save the presentation with another name or the name plus 4:3 or 16:9
 - f) Go through the new presentation and correct the sizing and placement of your photos
 - g) When you arrive at your talk determine whether you need a 4:3 or a 16:9 presentation and use the correct one.
- Regardless of whether you take your computer or a thumb drive include both your 4:3 and 16:9 presentations.





Background/ 2 – traits and disorders

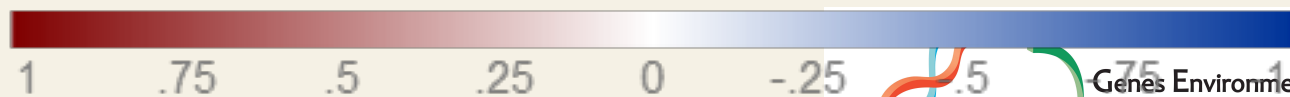
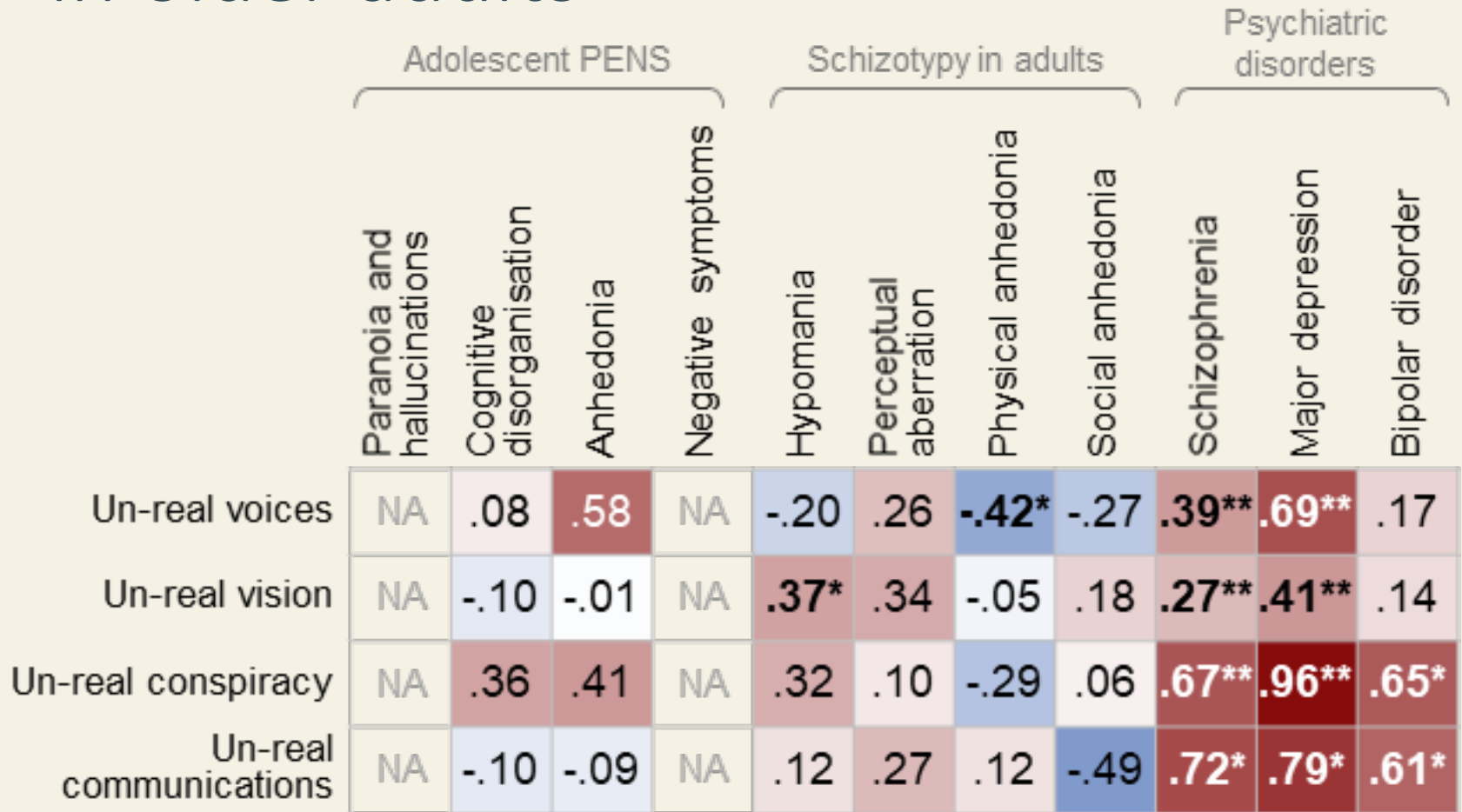
Psychopathology trait measures show genetic associations with disorders

Table 1. Effect Sizes of Genome-wide Polygenic Score Predictions for Four Psychiatric Disorders and Their Related Dimensional Traits

Disorder GPS	Related Trait	Variance Predicted in Disorder Itself by GPS (A)	Variance Predicted in Related Trait by Disorder GPS (B)	B/A
Autism	Autistic traits	1.13% (6)	0.1% (6)	9%
Schizophrenia	Negative symptom traits	7% (3)	0.7% (7)	10%
Depression	Depressive traits	0.72% (8)	0.11% (9)	15%
ADHD	ADHD traits	3.71% (10)	0.8% (10)	22%

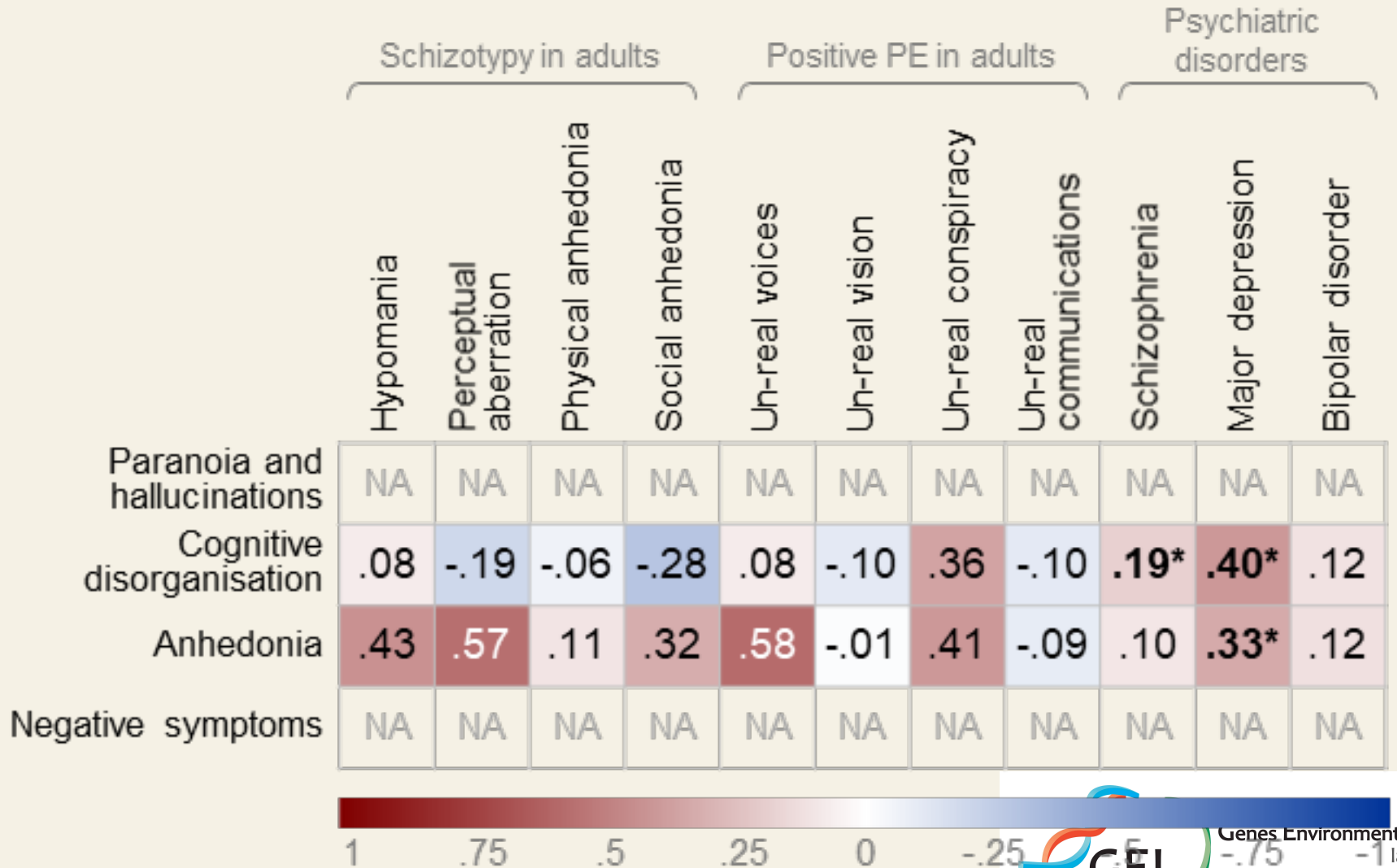
ADHD, attention-deficit/hyperactivity disorder; GPS, genome-wide polygenic score.

LD score regression with positive PE in older adults



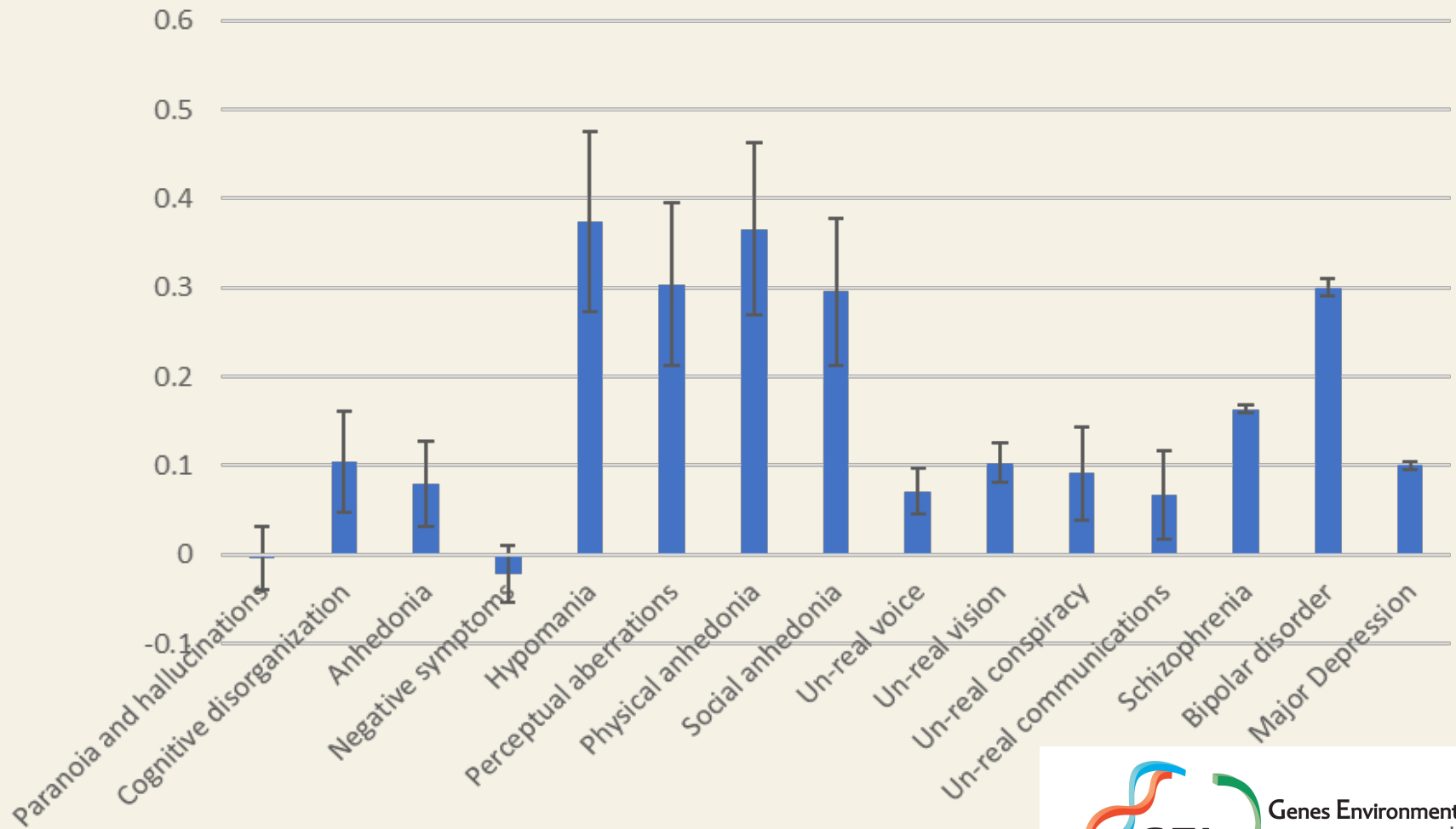
**Survived correction for multiple testing *Nominally significant

LD score regression with mid-adolescent PEs



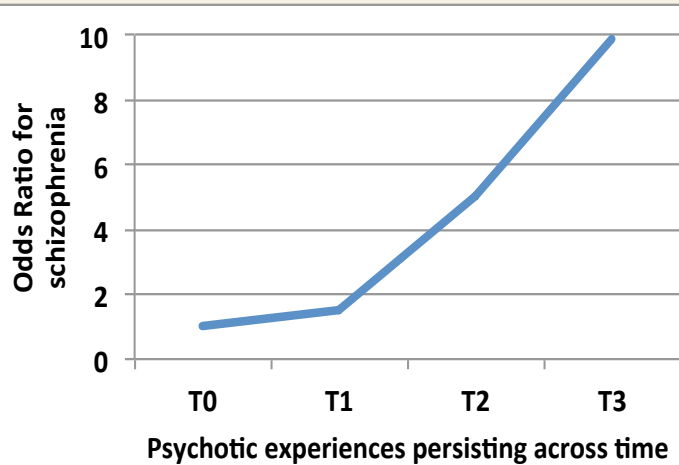
**Survived correction for multiple testing *Nominally significant

LD score regression: SNP h^2 estimates

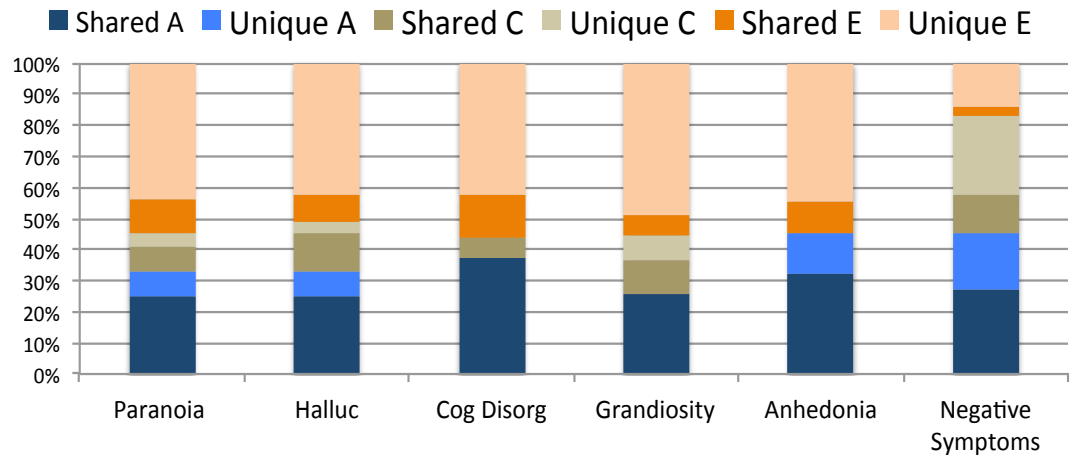


Pain et al., 2018 *Neuropsych Gen*; Ortega-Alonso et al., 2017 *Schz Bull*; Wray et al 2018 *Nat Gen*;
www.nealelab.is/uk-biobank; www.med.unc.edu/pgc;

Stability and Change across Age



Dominguez et al (2011) *Schz Bull*

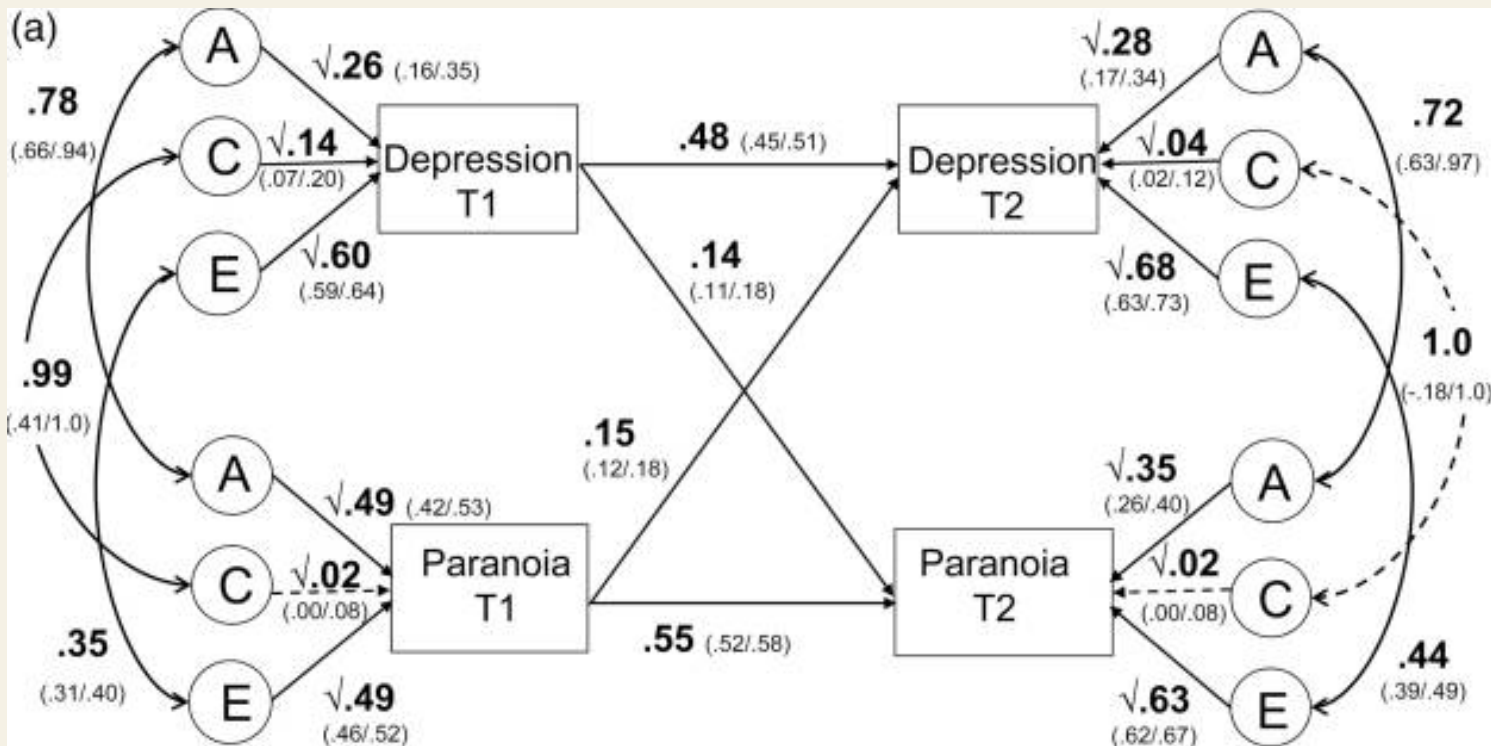


Variance in each subscale at time 2 explained by genetic and environmental influences unique and shared with time 1

Havers et al (2019) *Journal of Child Psychology and Psychiatry*

Causal Effects

Evidence of longitudinal causal effects between psychotic experiences and depression in adolescents



Zavos et al (2016) *Schz Bull*



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Aims

1 To evaluate genetic overlap between PE traits and clinical disorders

2 To explore stability of common genetic variation on PEs across adolescence → adulthood

3 Overlapping genes?

If genetic overlap:

3 Causal associations?

Analyses



LD score regression

→ To assess degree of genome-wide genetic overlap

FUMA

→ **Gene mapping**

- Positional
- Functional annotations

Mendelian randomization

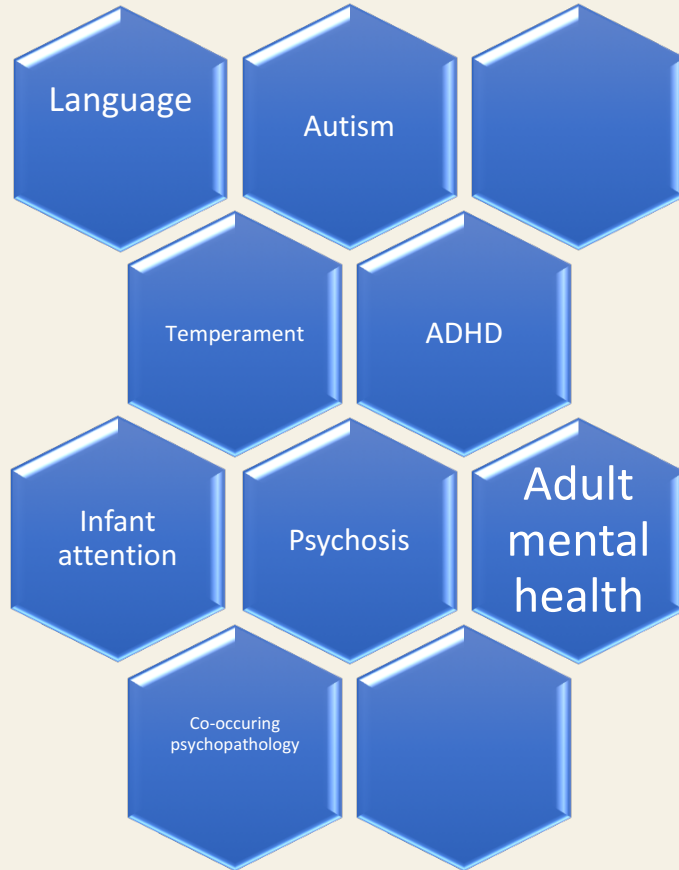
→ **Generalized summary based MR**

- Sensitivity analyses using MR methods that make different assumptions
- Heidi-outliers removed from main and sensitivity MR



Themes

adolescence analysis
attention autism
autistic-like behaviour
cognitive community-
development differences disorder
environmental etiological etiology evidence
genetic genome-wide
individual infancy influences
molecular nature neuropsychiatric
population-based problems
psychology
relationship research schizophrenia
traits twins



Development



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Genomic Structural Equation Modelling

