ACE/ADE Twin models

Sarah Medland and Hermine Maes 2024

Today - sarah/2024/monday

We pickup from the saturated model We will look at:

Extending the modelling to estimate heritability

Important structural stuff

- openMx has a very fluid and flexible structure
- Each code snippet is being saved as an object
- We tend to reuse the object names in our scripts
 - There are very few 'reserved' names
 - Naming a matrix "mean" does not make it a mean.

Important structural stuff

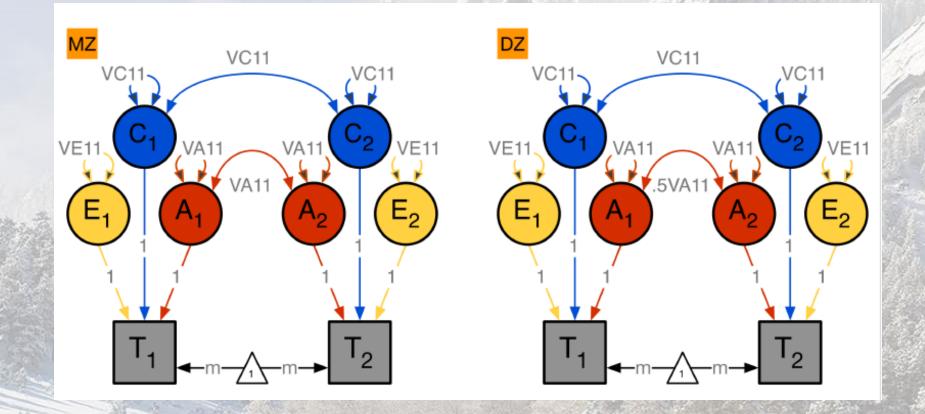
- No reserved names
 - We often use (and reuse) boring but meaningful names
 - make matrices VA VC VE
 - estimate VP = VA + VC + VE
 - But it would also work if we did this
 - make matrices Coffee Milk Ice
 - estimate Frappe = Coffee + Milk + Ice

Important structural stuff

- Because we tend to reuse the object names in our scripts you might need to remove these recycled objects from our work space to keep things tidy
 - rm(list=ls())
 - tends to freakout seasoned R uses sorry
- Remember the project also contains the data so these files can become very large.
- Storing projects that contain data requires careful thinking about data security and can be an IRB risk if the project 'leaves the building'

Today

• MZ and DZ pairs – estimating A, C and E



MZ	A+C+E	A+C	
	A+C	A+C+E	

covP <- mxAlgebra(expression= VA+VC+VE, name="V")
covMZ <- mxAlgebra(expression= VA+VC, name="CMZ")</pre>

V	cMZ
cMZ	V COLUMN

expCovMZ <- mxAlgebra(expression= rbind(cbind(V, cMZ),

cbind(t(cMZ), V)),
name="expCovMZ")

DZ

A+C+E	.5⊗A+C
.5⊗A+C	A+C+E

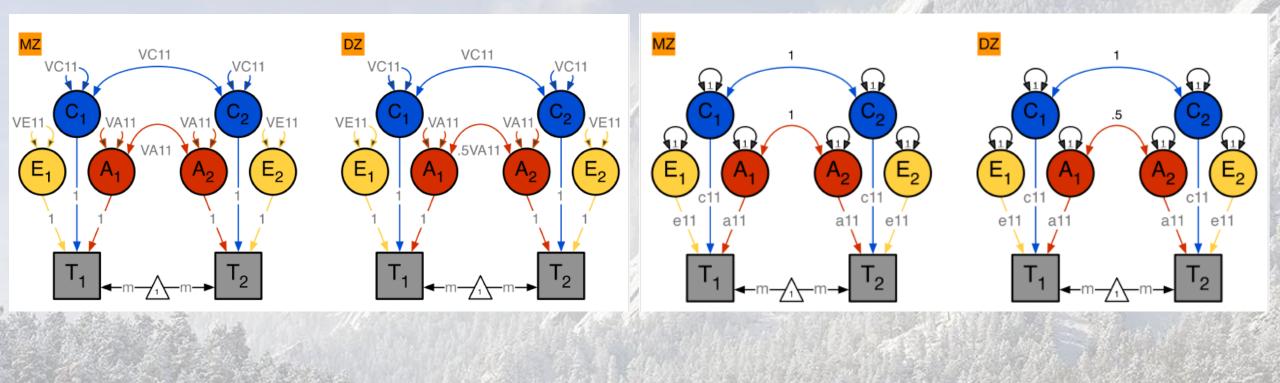
covP <- mxAlgebra(expression= VA+VC+VE, name="V")
covDZ <- mxAlgebra(expression= 0.5%x%VA+VC, name="cDZ")</pre>



expCovDZ <- mxAlgebra(expression= rbind(cbind(V, cDZ),

cbind(t(cDZ), V)),
name="expCovDZ")

Change in approach



Two very useful papers in the ACE folder

Twin Research (1999) 2, 250–257 © 1999 Stockton Press All rights reserved 1369–0523/99 \$15.00

http://www.stockton-press.co.uk/tr

Genetic and environmental causes of variation in basal levels of blood cells

David M Evans¹, Ian H Frazer² and Nicholas G Martin¹

22



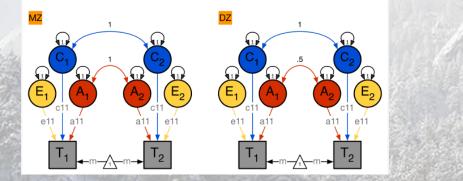
Biological Psychology 61 (2002) 33-51

BIOLOGICAL PSYCHOLOGY

www.elsevier.com/locate/biopsycho

Biometrical genetics

David M. Evans*, N.A. Gillespie, N.G. Martin



Practical time

• In R studio

system("cp -R /faculty/sarah/2024/ACE/* ./ ")

Qualtrics link is in ACE.txt

https://qimr.az1.qualtrics.com/jfe/form/SV_aXZTsTpDqACPQpM



Lets recap

- The % variation in a trait attributable to genetic effects in a population.
- The extent to which individual differences in genetics contribute to individual differences in observed behaviour in a large group of people
- Heritability should be thought about deterministically at the individual level
- These analyses don't tell us what the genetic or environmental factors are.

For example

- About 75% of the variation in ADHD can be explained by genetic effects
- Heritability of 75%

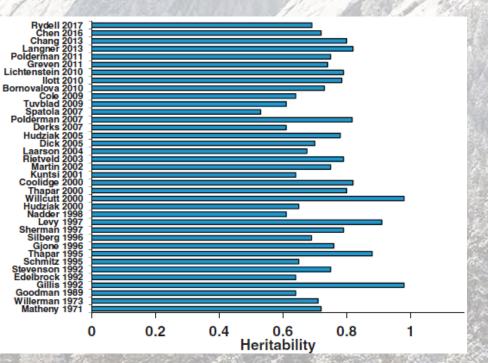
Molecular Psychiatry (2019) 24:562–575 https://doi.org/10.1038/s41380-018-0070-0

REVIEW ARTICLE

Genetics of attention deficit hyperactivity disorder

Stephen V. Faraone ¹ · Henrik Larsson^{2,3}

Received: 29 August 2017 / Revised: 31 January 2018 / Accepted: 19 February 2018 / Published online: 11 June 2018 \odot The Author(s) 2018. This article is published with open access



This does not mean...

- That 75% of an individual's behaviour is due to their genetics and the other 25% is due to their environment
 - Genetic control 3am-9pm. Environmental control 9pm-3am
- That 75% of people have ADHD because of a genetic reason and 25% have ADHD because of an environmental reason
- A child of someone with ADHD has a 75% chance of developing ADHD

Important to remember

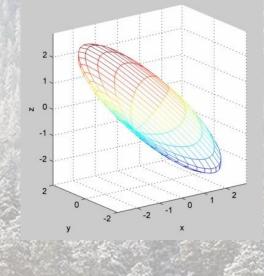
Our results are estimates based on the data we've analysed

- The won't necessarily generalize
 - Across time or space

Thinking out side the box...

Rather than thinking about estimates as fixed points I like to think about parameter space...

Imagine an ACE/ADE model as a solution space bounded by CIs



Important to remember

"Remember that all models are wrong; the practical question is how wrong do they have to be to not be useful"

George E P Box and Norman R Draper. 1986. Empirical Model-Building and Response Surface. John Wiley & Sons, Inc., New York, NY, USA.



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Questions?