## **Power Calculation Practical**

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# Power Calculations Empirical

- Attempt to Grasp the NCP from Null
- Simulate Data under theorized model
- Calculate Statistics and Perform Test
- Given α, how many tests p < α</li>
- Power = (#hits)/(#tests)

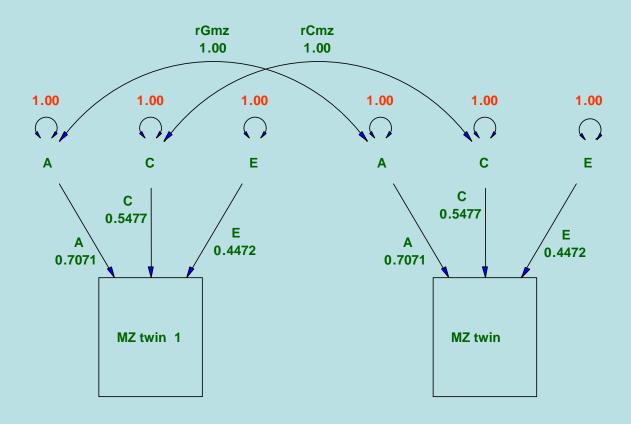
# Practical: Empirical Power 1

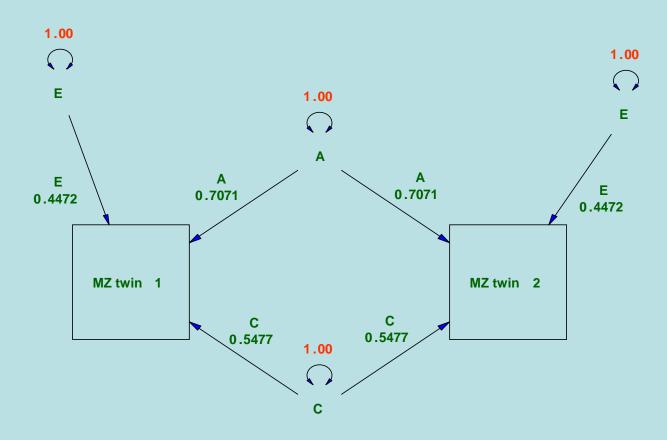
- We will Simulate Data under a model online
- We will run an ACE model, and test for C
- We will then submit our results and Jeff will collate the empirical values
- While that is being calculated, we'll talk about theoretical power calculations

# Practical: Empirical Power 2

- First get ace.mx and rprog.R from
- /faculty/ben/2006/power/practical/.
- We'll talk about what the R program does before we run it

## Simulation of the MZs: model

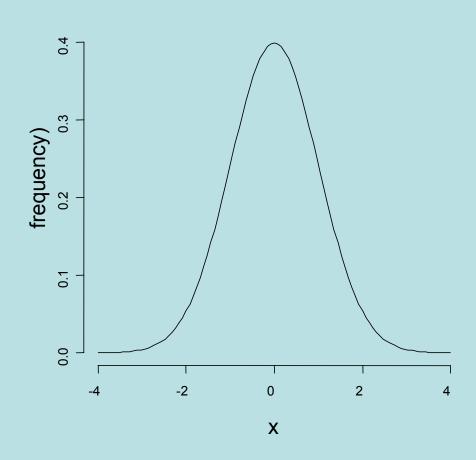


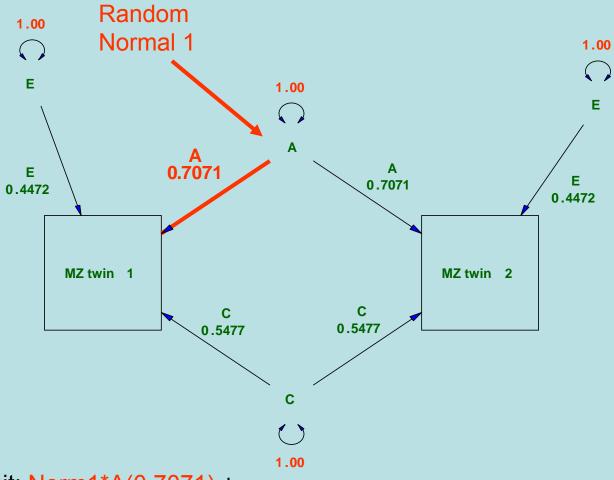


#### When we simulate

- From a path diagram, we can simulate trait values from simulating each latent trait
- These latent traits are assumed to be normal (µ=0,σ²=1 or □=0,☞²=1)
- The latent trait is then multiplied by the path coefficient

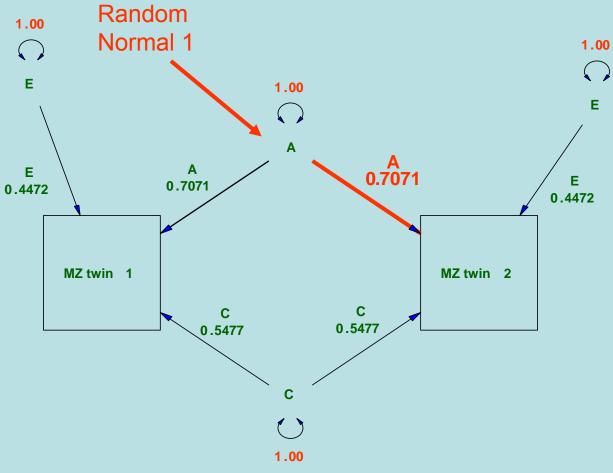
## What's a random normal





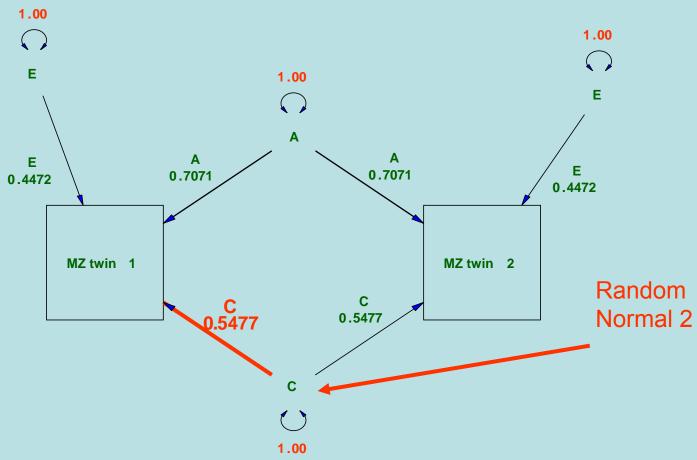
MZ twin 1 trait: Norm1\*A(0.7071) +

MZ twin 2 trait:



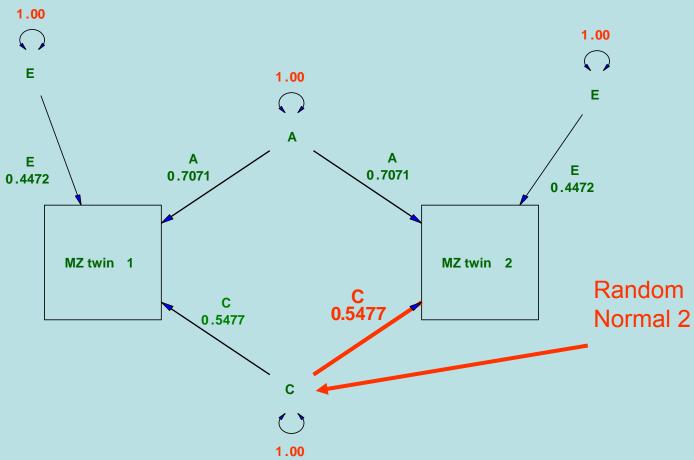
MZ twin 1 trait: Norm1\*A(0.7071) +

MZ twin 2 trait: Norm1\*A(0.7071) +



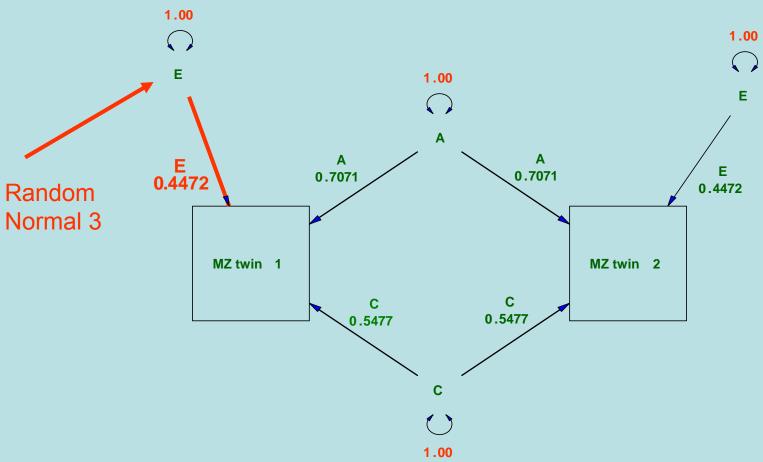
MZ twin 1 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477)

MZ twin 2 trait: Norm1\*A(0.7071) +



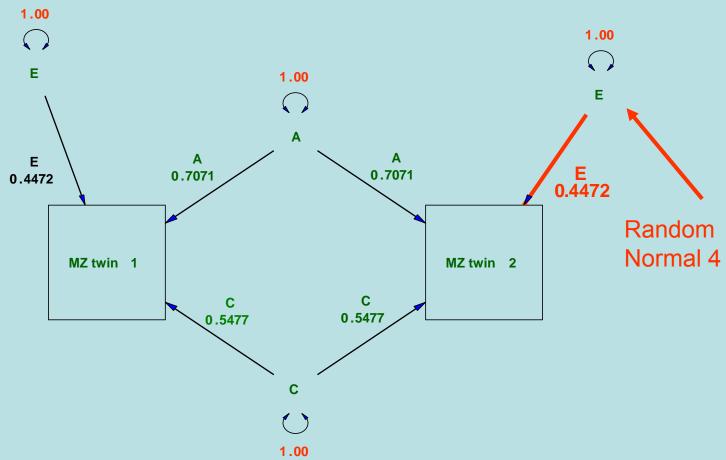
MZ twin 1 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477)

MZ twin 2 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477)



MZ twin 1 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477) + Norm3\*E(0.4472)

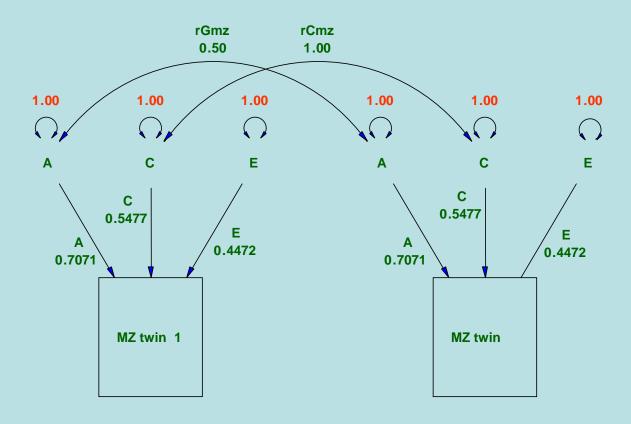
MZ twin 2 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477) +

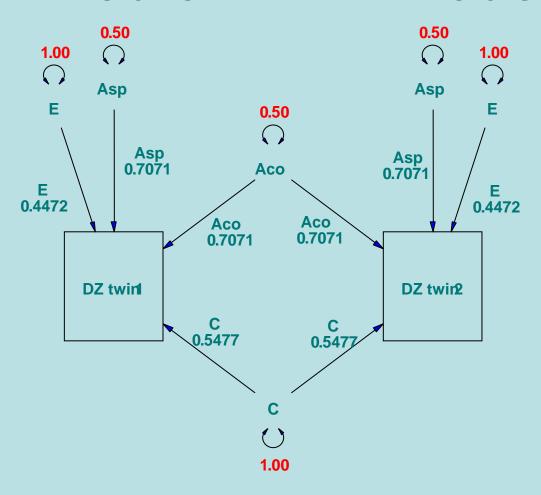


MZ twin 1 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477) + Norm3\*E(0.4472)

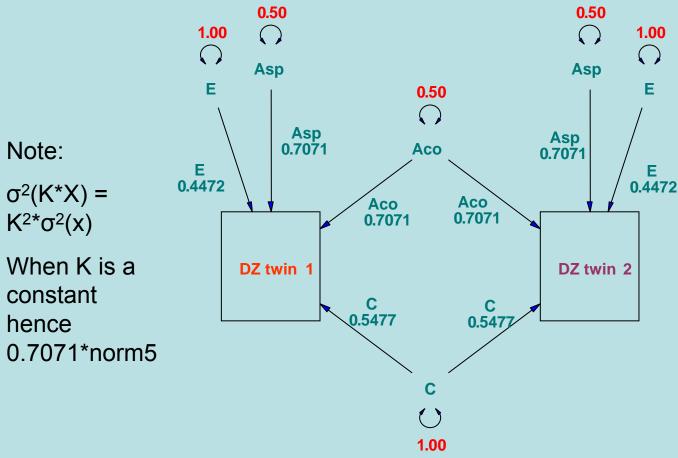
MZ twin 2 trait: Norm1\*A(0.7071) + Norm2\*C(0.5477) + Norm4\*E(0.4472)

## Simulation of the DZs: model





How many random normals will we need to supply a trait value for both DZ twins?



DZ twin 1 trait: 0.7071\*Norm5\*Aco(0.7071) + 0.7071\*Norm6\*Asp(0.7071) + Norm7\*C(0.5477) + Norm8\*E(0.4472)

DZ twin 2 trait: 0.7071\*Norm5\*Aco(0.7071) + 0.7071\*Norm9\*Asp(0.7071) + Norm7\*C(0.5477) + Norm10\*E(0.4472)

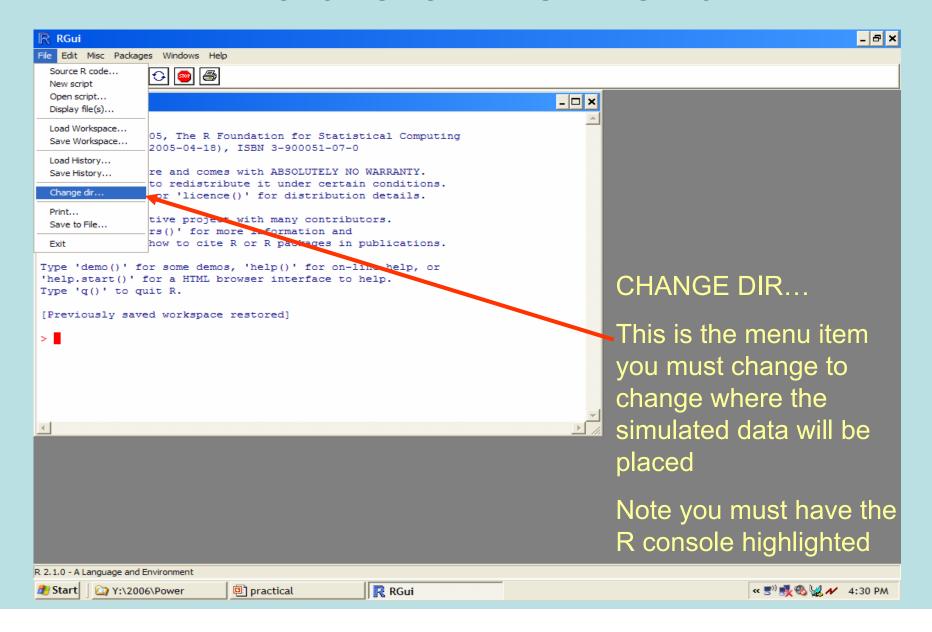
#### Simulation conditions

- 50% additive genetic variance
- 30% common environment variance
- 20% specific environment variance

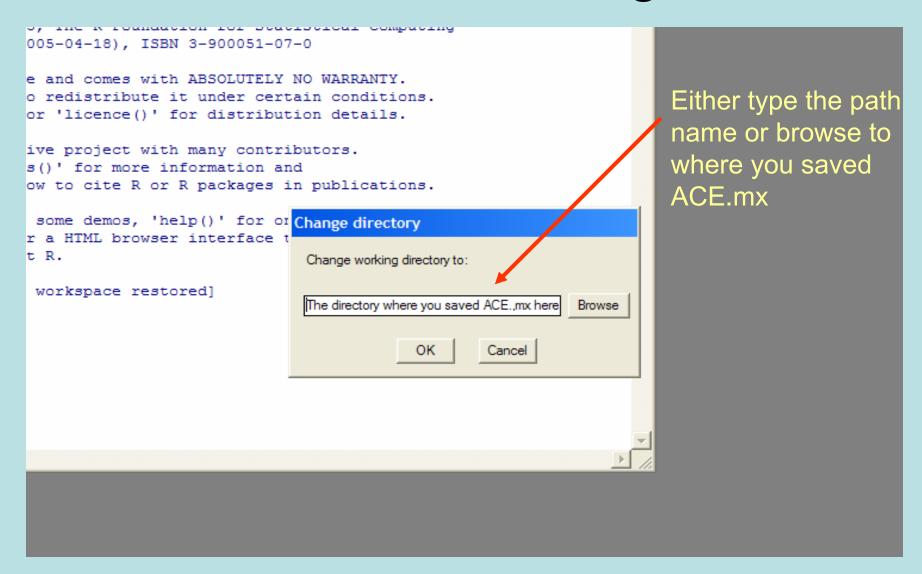
## Notes on the R program

- When you run the R program it is essential that you change your working directory to where you saved the Mx script.
- File menu then Change dir...
- After changing directory, load the R program.
- A visual guide to this follows this slide

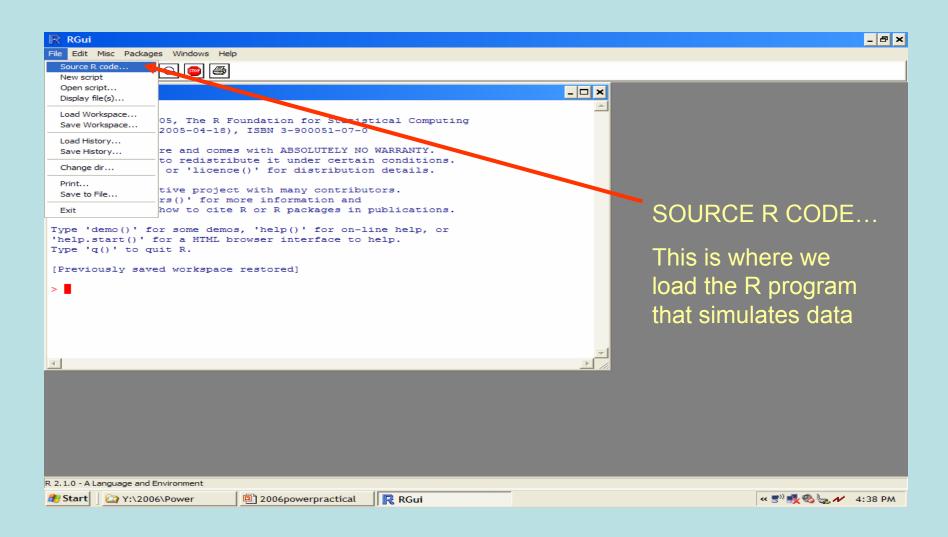
## Picture of the menu



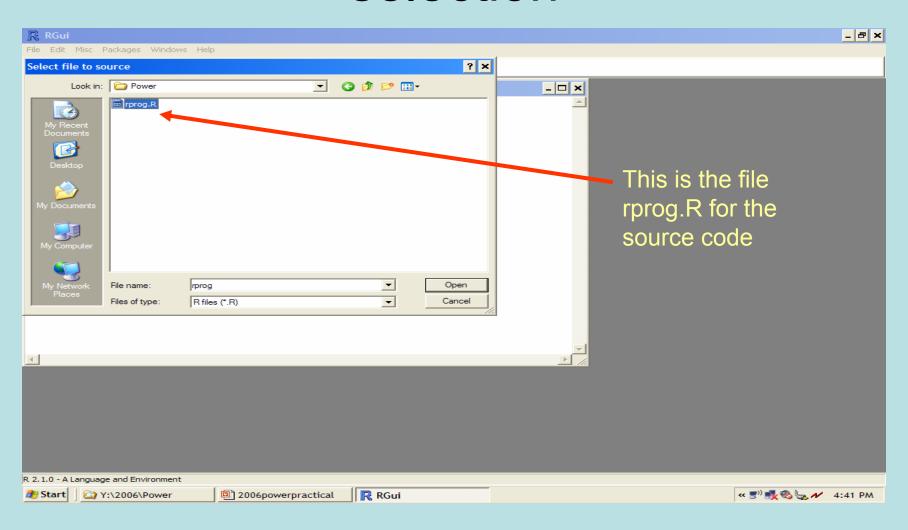
## Picture of the dialog box



# Running the R script



# Screenshot of source code selection



#### How do I know if it has worked?

- If you have run the R program correctly, then the file sim.fun ought to be in the directory where your rprog.R and ACE.mx is.
- If not, try again or raise your hand.

## When you have finished

- Note your likelihoods and your parameter estimates and complete the survey at:
- https://ibgwww.colorado.edu/phpsurveyor/i ndex.php?sid=4

# Theoretical power calculations

- Either derive the power solutions by hand (though this requires lots of time and more IQ points than I have)
- Use Mx to setup the variance covariance structure and use option power to generate power levels

# Quick note on the power calculations for Mx

- Total sample size is reported at the end of the script
- The sample size proportions for your groups are maintained.
- For example if we say 50 MZ pairs and 100 DZ pairs, then Mx will assume 1/3 of your sample is MZ and 2/3 is DZ

# Time to look at a script

- Open power.mx, and we'll chat about it.
- Quick overview of what the script does:
  - Generates the variance covariance structure under the full model (1st half)
  - Intentionally fits the wrong model (by dropping the parameter of interest for power calculations) (2<sup>nd</sup> half)
  - Based on the number of observations that you supply generates power estimates.

# Theoretical script

- Following chatting, depending on time, here are some suggestions:
  - Change ratio of MZ and DZ keeping same total sample size
  - Drop A rather than C
  - Change effect sizes for A, C, or E