

MR-MEETING UPDATES

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MRDoC Power Comparisons

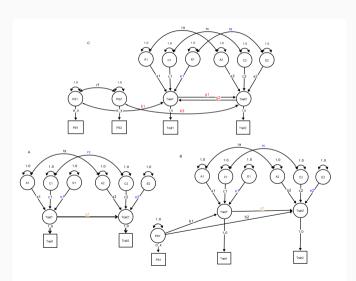
MRDoC Power Comparisons - Data generating model is MRDOC2

Longitudinal model

MRDoC Power Comparisons

SPECIFICATION





MR-DoC (A), DoC (B), and MR-DoC2 (C) model specifications for a single twin member. They include the effects of additive genetic (A), common environment (C) and specific environment (E) factors for both Trait 1 and Trait 2, and their effects may correlate (parameters ra, rc, and re). Path labels in red are important to the model's overall power, those susceptible to measurement error in blue, and in orange are those that are both susceptible to measurement error and are important to the model's overall power.

FACTORIAL DESIGN

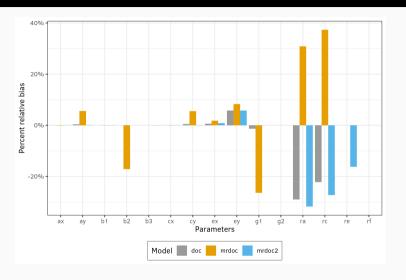


Table 1: Parameter levels on the three factorial designs, with respective total number of cells for each design simulation. The model specification can be seen in Figure 1.

theta	Design 1 (DoC)	Design 2 (MR-DoC)	Design 3 (MR-DoC2)
b1		sqrt(0.025, 0.05,0.75)	sqrt(0.025, 0.05,0.75)
b2		sqrt(0.025, 0.05,0.75)	
b3			sqrt(0.025, 0.05,0.75)
g1	sqrt(0.20, 0.40, 0.60)	sqrt(0.20, 0.40, 0.60)	sqrt(0.20, 0.40, 0.60)
g2			sqrt(0.20, 0.40, 0.60)
ra	.0,.25,.50	.0,.25,.50	.0,.25,.50
rc	.0,.25,.50	.0,.25,.50	.0,.25,.50
re			.0,.25,.50
rf			.0,.25,.50
ax	.0,.10,.25	.0,.10,.25	.0,.10,.25
ay	.0,.10,.25	.0,.10,.25	.0,.10,.25
СХ	.0,.10,.25	.0,.10,.25	.0,.10,.25
су	.0,.10,.25	.0,.10,.25	.0,.10,.25
Total cells	3^7=2187	3^9= 19683	3^12=531441

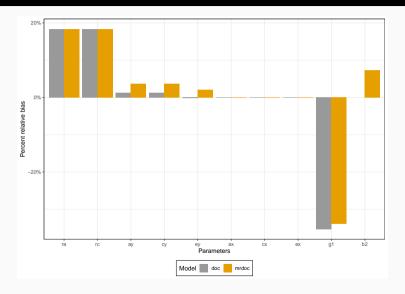
PERCENT BIAS - UNRELIABILITY FACTOR





PERCENT BIAS - RE UNMODELLED





MRDoC Power Comparisons - Data Generating model is MRDOC2

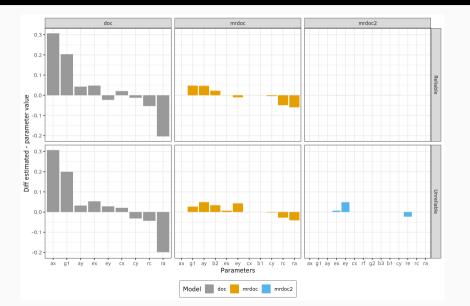
LEVELS



```
try(stopCluster(cl))
cl <- makeCluster(detectCores()-1, outfile = "")</pre>
registerDoParallel(cl)
local1 <- foreach(g1=c(sqrt(.020),sqrt(.040)), .combine =rbind,</pre>
                  .packages = c("umx", "MASS", "tidvr")) %:%
          foreach(axs=c(0,.10), .combine =rbind) %:%
          foreach(ays=c(0,.10), .combine = rbind) %:%
          foreach(g2=c(sqrt(.020),sqrt(.040)), .combine =rbind) %:%
          foreach(b3=c(sqrt(.025),sqrt(.05)), .combine =rbind) %:%
          foreach(re=c(0..10), .combine =rbind) %:%
          foreach(ra=c(0, .25), .combine =rbind) %:%
          foreach(reliability=c(FALSE, TRUE), .combine =rbind) %:%
          foreach(rc=c(0, .25), .combine =rbind) %dopar% {
  sim(est out = T. g1=g1.g2=g2. b3=b3. axs=axs. avs=avs. rc=rc. ra =ra.
      reliability = reliability )
stopCluster(cl)
```

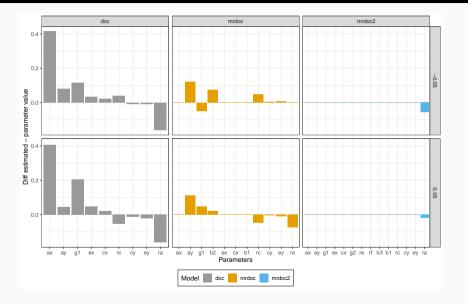
UNRELIABILITY FACTOR





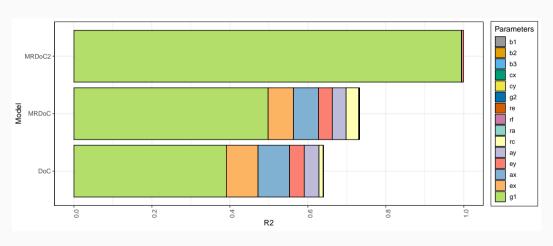
RE IN DATA, BUT NOT IN THE fIT





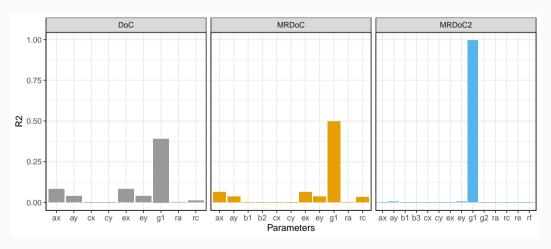
VARIANCE EXPLAINED IN STATISTICAL POWER - V1





VARIANCE EXPLAINED IN STATISTICAL POWER - V2





VARIANCE EXPLAINED IN STATISTICAL POWER BY MODEL PARAMETERS - OLD



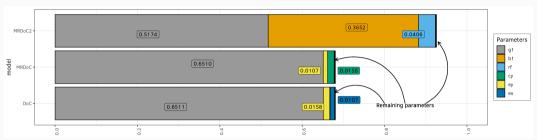
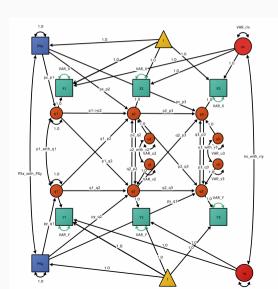


Figure 1: Linear regression of NCP on the parameter values used for 19683 and 4097 exact data simulation power analyses in the ACE model for DoC, MR-DoC and MR-DoC2, respectively. The hypotheses are g1=0 for all. These are stacked bar plots with semipartial correlation squared for each of the parameters. The total R2 for both DoC and MR-DoC models (including all parameters in the regression) was 0.945, and 0.93 for MR-DoC2. One can interpret these as g1 having the largest effect on DoC and MR-DoC overall power, and g1, b1 and rf having the largest effects on the MR-DoC2 overall power.

LONGITUDINAL MODEL

SPECIFICATION





The model is identified as depicted (3 waves). It comprises 5 main elements: (A) the observed variables in green; (B) the between-person variances in orange; (C) the polygenic scores in blue; (D) the means in yellow; and (E) the random intercepts in red. Free paths are in black, and named. Fixed paths are marked with 1. The specific variances for the observed variables are equal across study waves, marked in green.

STATIONARITY



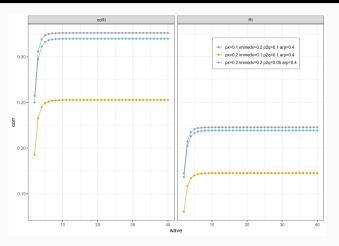
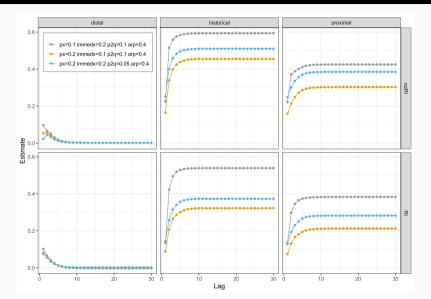


Figure 2: Correlation between Xt and Yt, where t is the study wave up to 40. With the extension of the model, the correlations stabilize. This model reaches stationarity at wave 10.

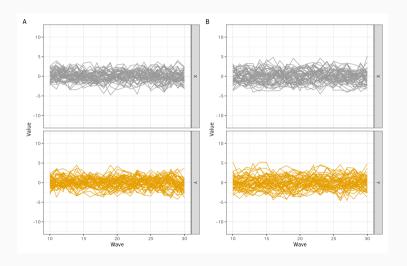
ESTIMATES WITH INCREASING LAG





WIREFRAME OF THE MEANS OF THE MEASURED VARIABLES ACROSS WAVES





The means of the observed variables are plotted individually for both Xt and Yt, when t is the study wave, starting in t=20. The panel B depict the means when the model with random intercepts. Panel A depict the model without random intercepts. Panel B shows more spread than panel A. Theta (px =py = 0.1; pt_qt+1=qt_pt+1 = 0.05; pt_qt+1 = qt_pt+1 = 0.1; pt_qt+1 = qt_qt+1 = 0.4)

HEATMAP FOR THE EXPECTED COV



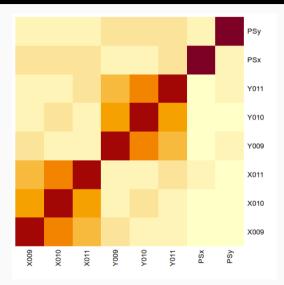
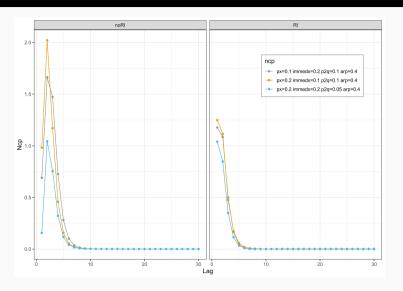


Figure 4: The expected covariance matrix for the observed variables.

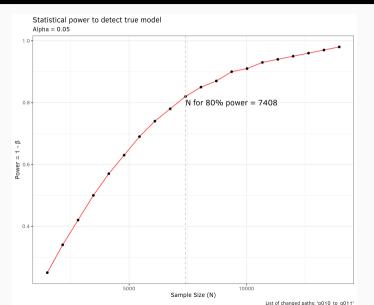
POWER WITH INCREASING LAG





POWER AT THE FIRST STATIONARY WAVE





ACKNOWLEDGEMENTS



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Contact



• THANK YOU