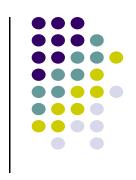


psychopathology, personality and genetics

Bob Krueger Hathaway Distinguished Professor University of Minnesota, USA

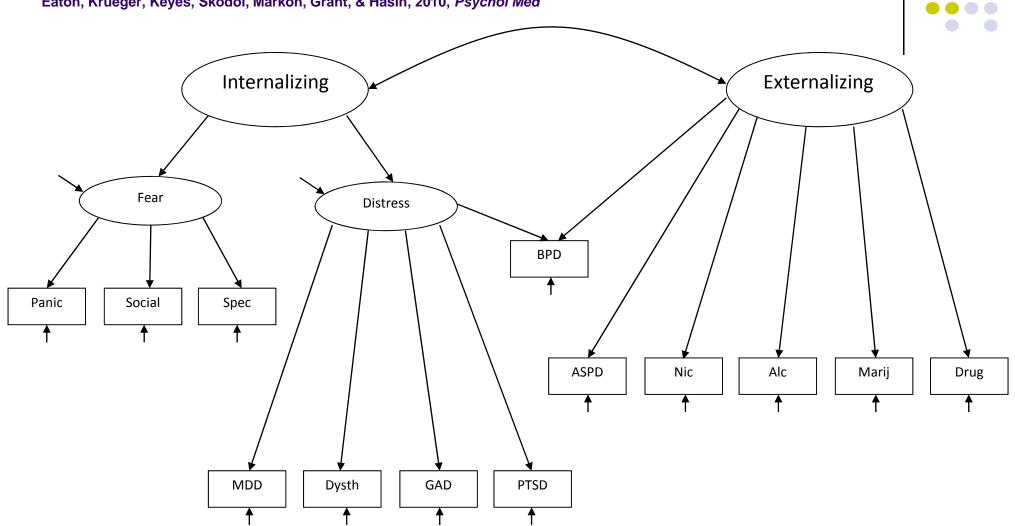
psychopathology



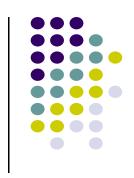
- has typically been thought of in terms of
- 297
- putatively discrete
- and separate categories
- that are polythetic
 - multiple combinations of criteria -> same diagnosis
- and each category is typically studied by different people publishing in different journals attending different scientific meetings funded by different parts of NIH (and so on)
- is this an accurate model of psychopathological variation?
 - and a good way to work on the problem?

a dimensional-spectrum model of common forms of psychopathology

Eaton, Krueger, Keyes, Skodol, Markon, Grant, & Hasin, 2010, Psychol Med

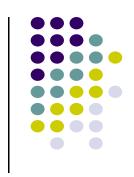


some things we've learned



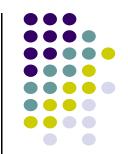
- i. the latent spectrum variables are continuous
 - as opposed to being discrete
 - via direct comparison of models with categorical vs. continuous latent variables
- ii. the phenotypic correlations among indicators are (primarily) genetically mediated
 - the latent variables appear more heritable than the indicator diagnoses
 - especially externalizing (> internalizing)
 - there are residual genetic effects on the indicators
 - alcohol is a good example
 - but the residuals are smaller effects

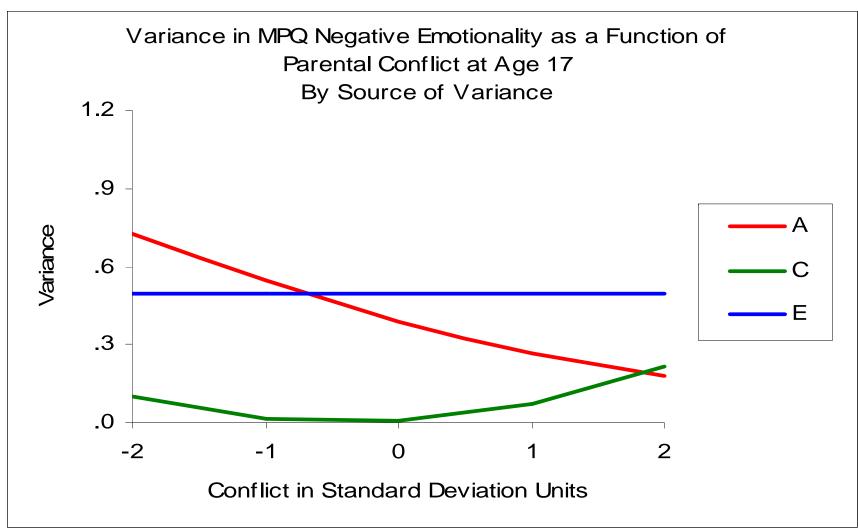
some things we've learned



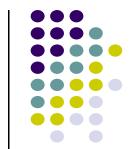
- iii. personality is at the core of the spectrums
 - dispositions function like diagnoses as indicators
 - genetically correlated with diagnoses
 - controlling for dispositions reduces "comorbidity"
- iv. this model is likely to frame key aspects of the DSM-5 meta-structure
- upshot: personality dispositions are key variables in behavioral public health
 - the social costs of psychopathology are unequivocal
 - negative emotionality (neuroticism) is at the core of internalizing
 - disinhibition (un-conscientious dis-agreeableness) is at the core of externalizing
 - understanding the etiology and neurobiology of these dispositions is important

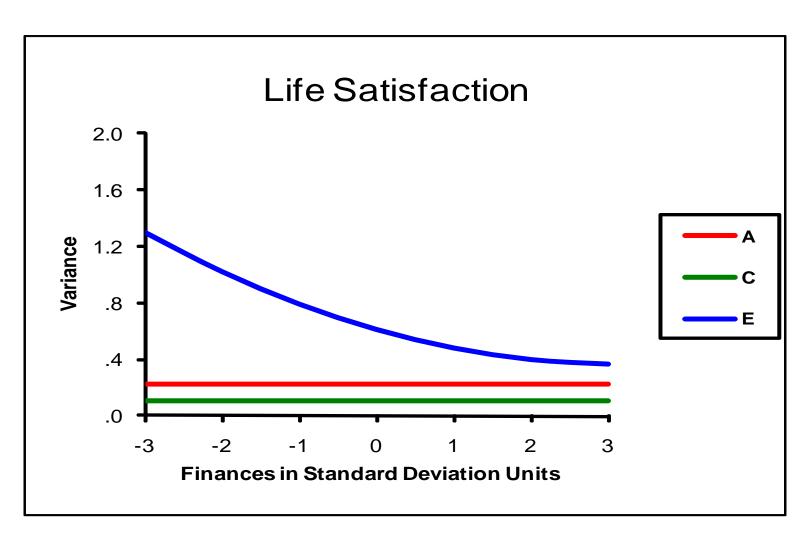
etiology of negative emotionality varies as a function of family conflict





finances moderate etiology of life satisfaction





Variance in Life Satisfaction as a Function of Finances, by source of variance.

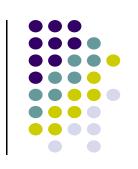
meta analytic GWAS of personality

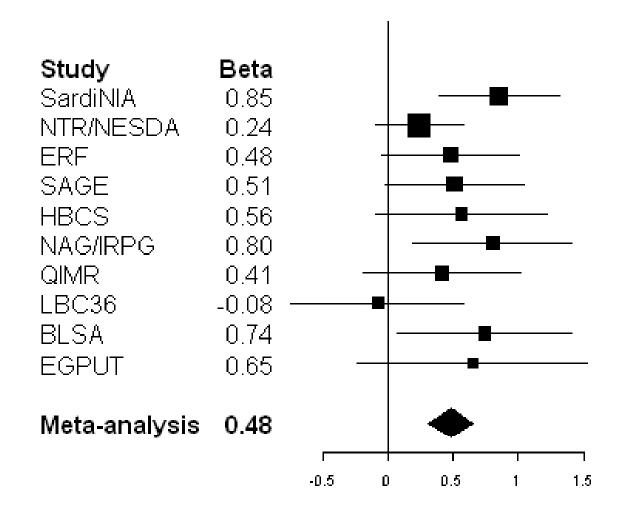
Marleen H.M. de Moor 1†, Paul T. Costa 2, Antonio Terracciano 2, Robert F. Krueger 3, Eco J.C. de Geus 1, Tanaka Toshiko 2, Brenda W.J.H. Penninx 4,5,6, Tõnu Esko 7,8,9, Pamela A F Madden 10, Jaime Derringer 3, Najaf Amin 11, Gonneke Willemsen 1, Jouke-Jan Hottenga 1, Marijn A. Distel 1, Manuela Uda 12, Serena Sanna 12, Philip Spinhoven 5, Catharina A. Hartman 4, Patrick Sullivan 13, Anu Realo 14, Jüri Allik 14, Andrew C Heath 10, Michele L Pergadia 10, Arpana Agrawal 10, Peng Lin 10, Richard Grucza 10, Teresa Nutile 15, Marina Ciullo 15, Dan Rujescu 16, Ina Giegling 16, Bettina Konte 16, Elisabeth Widen 17, Diana L Cousminer 17, Johan G. Eriksson 18,19,20, 21,22, Aarno Palotie 17,23,24, 31, Leena Peltonen 17,23,24, 31 **, Michelle Luciano 25, Albert Tenesa 26, Gail Davies 25, Lorna M. Lopez 25, Narelle K. Hansell 27, Sarah E. Medland 27, Luigi Ferrucci 2, David Schlessinger 2, Grant W. Montgomery 27, Margaret J. Wright 27, Yurii S. Aulchenko 11, A.Cecile J.W. Janssens 11, Ben A. Oostra 28, Andres Metspalu 7,8,9, Gonçalo R. Abecasis 29, Ian J. Deary 25, Katri Räikkönen 30, Laura J. Bierut 10, Nicholas G. Martin 27, Cornelia M. van Duijn 11*, and Dorret I. Boomsma, in press, *Mol Psychiatry*



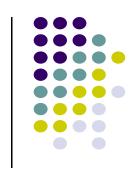
- 17,375 unrelated individuals of European ancestry from Europe, the United States and Australia
- 10 contributing studies
- genotyping platforms rendered commensurate via imputation
 - ~2.5M common SNPs included in HapMap, using the HapMap phase II CEU data as the reference sample
 - ~2,500,000 data points per person
- phenotypes are NEO-FFI (Five Factor Model; FFM) scales
 - Openness ,Conscientiousness, Extraversion, Agreeableness, Neuroticism
 - OCEAN

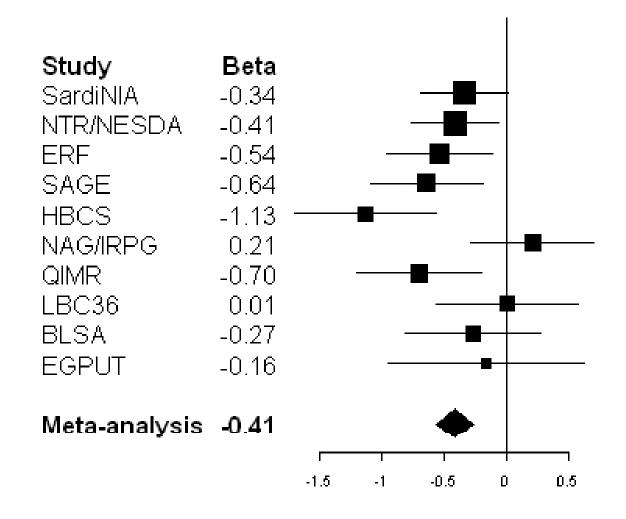
FFM meta analysis: snp on chromosome 5 for openness





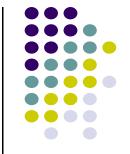
FFM meta analysis: snp on chromosome 18 for conscientiousness

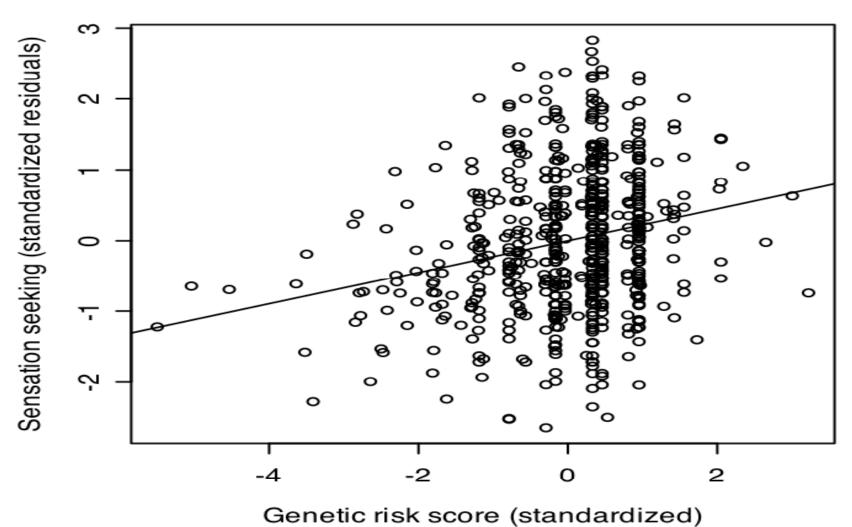




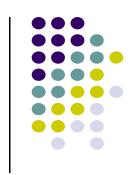
aggregate of 12 snps in the dopamine (DA) system predicts sensation seeking

Derringer J, Krueger RF, Dick DM, Saccone S, Grucza RA, Agrawal A, ... Bierut LJ, 2010, Psychol Sci





genetics of personality and psychopathology: ongoing steps



- whole genome approaches will (and should) continue
 - much of what is found in "medical" molecular genetics would not have been predicted from a candidate approach
- candidate sets of variants for association can be pursued simultaneously
 - neurobiological variables are key intermediate phenotypes
- twin research continues to offer valuable insights
 - can study G-E processes at an aggregate level

full moderator model (cholesky with moderation effects; 17 parameter estimates)

