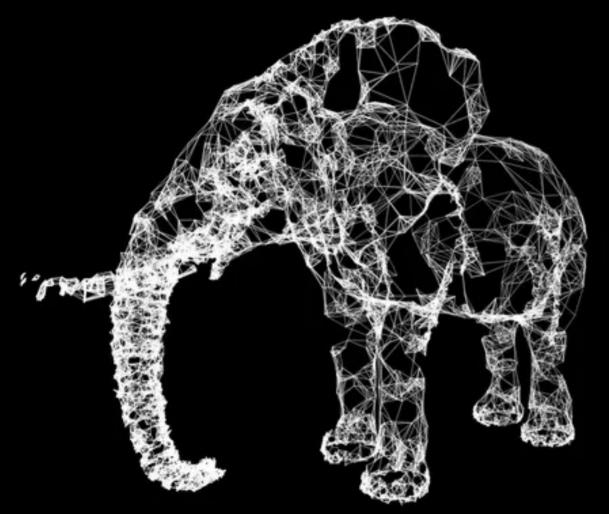
# The neglected importance of complexity in statistics and Metascience

Daniele Fanelli

### In this talk:

- 1) what's missing in the current paradigm
- 2) what a new paradigm might look like
- 3) evidence in support of this proposal

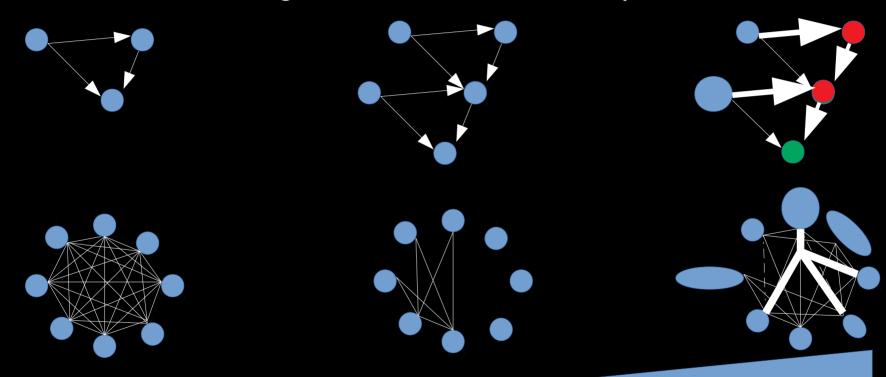


The elephant in the room of:

- 1) metascience
- e.g. reproducibility
- 2) statistics
- e.g. model "complexity"
- [see SW seminar 2021, Fanelli 2019, 2022]

# What is "complex"?

Many, diverse, interacting parts. Long to describe, difficult to predict.



Level of complexity

# example 1) complexity deflates the "reproducibility crisis".

year	project	discipline	N	result
2014	Many labs 1	psychology, misc.	13, 36 labs	77%
2016	COS	social+cognitive psychology	100	36-68%
2016	Camerer et al.	experimental economics	18	61-78%
2018	Many labs 2	social+cognitive psychology	28, 62 samples, 36 countries	54%
2018	Camerer et al.	social studies in Nature, Science	21	57-67%
2021	RPCB	cancer biology	188, 50 exper., 23 papers	3-82%

Lower reproducib:

- 1) complex phenomena
- 2) complex methods
- not just random noise
- structured, systematic diff.

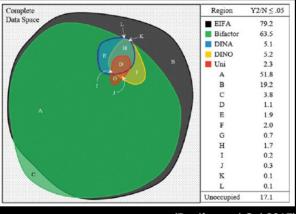
# example 2) complexity confuses statistical results

$$AIC:-2\log(L)+2k$$

### models vary by "fitting propensity"

("complexity" beyond n. of parameters)

universe of possible data



(Bonifay and Cai 2017)

### When is a theory <u>actually</u> supported?

### How might the elephant appear?

- 1) integrate complexity of phenomena & methods in measuring, forecasting, correcting reproducibility
- 2) penalize statistical models for complexity beyond number of parameters

integrating "complexity" ~ severity of testing

### Part 2: A candidate alternative



### K theory in a nutshell

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

### K = consilience

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

explain/predict/control more/diverse phenomena with fewer/simpler theories/methods

### what the variables represent

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

explain/predict/control more/diverse phenomena with fewer/simpler theories/methods

### more information Y makes K GROW

$$H(Y) - H(Y|X, \tau)$$

$$H(Y) + n_X H(X) + D(\tau)$$

$$n_Y$$

### more information Y makes K GROW

$$= \frac{H(Y_1) + H(Y_2) + H(Y_3) - H(Y|X, \tau)}{H(Y_1) + H(Y_2) + H(Y_3) + n_X H(X) + D(\tau)}$$

$$n_Y$$

### K theory in a nutshell

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

### more info. Y|X, X, τ makes K small

$$\kappa = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + nH(X) + D(\tau)}$$

$$\frac{n_{Y}}{n_{Y}}$$

### more info. Y|X, X, τ makes K small

$$\kappa = \frac{H(Y) - H(Y_1|X, \tau) + H(Y_2|X, \tau) + H(Y_3|X, \tau)}{H(Y) + n_X + D(\tau_1) + D(\tau_2) + D(\tau_3)}$$

$$n_Y$$

### K theory in a nutshell

$$K = \frac{H(Y) - H(Y|X, T)}{H(Y) + n_X H(X) + D(T)} = 0, \text{ no knowledge}$$

$$= 0, \text{ no knowledge}$$

$$= 0, \text{ wrong}$$

### K theory in a nutshell

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

### K of a regression model

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + n_X H(X) + D(\tau)}$$

$$n_Y$$

$$Y = \alpha + \beta X + error$$

### this has been said before

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + \frac{n_x}{n_y} H(X) + D(\tau)}$$

$$N_y$$

$$Y = \alpha + \beta X + \text{error}$$

### key theoretical innovations

$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + D(\tau)}$$

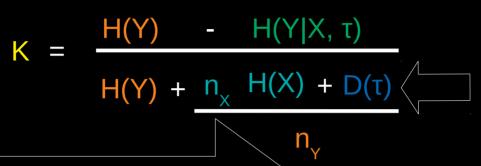
$$= \alpha + \beta X + error$$

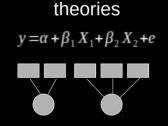
### key methodological innovations

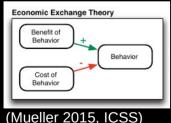
$$K = \frac{H(Y) - H(Y|X, \tau)}{H(Y) + \frac{n_x}{n_y} + D(\tau)}$$

$$Y = \alpha + \beta X + \text{error}$$

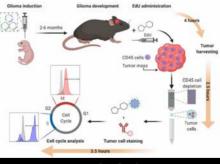
# graphs are everywhere in science





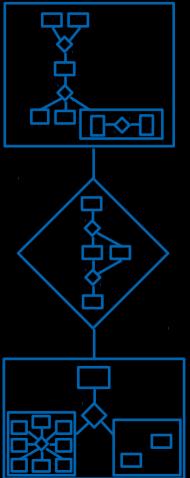


#### methodologies



WWW.protocols.io/view/an-optimized-protocol-for-in-vivo-analysis-of-tumo-3byl471m2lo5/v1

(Fanelli 2019, Fanelli 2022)

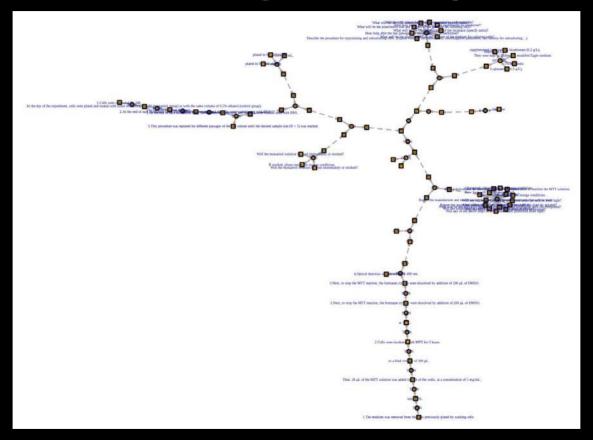


# Part 3: supporting evidence



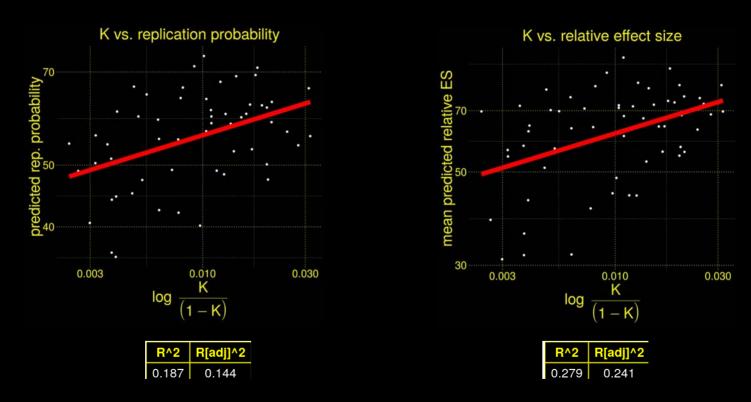
# 1) K predicts perceived and actual reproducibility

### "tau" of biological experiments



(Fanelli, Tan, Amaral & Neves, 2022, MetaArxiv)

### K vs. perceived reproducibility



(Fanelli, Tan, Amaral & Neves, 2022, MetaArxiv)

### K vs. actual reproducibility

$$K_r = K_o 2^{-\lambda \cdot d}$$

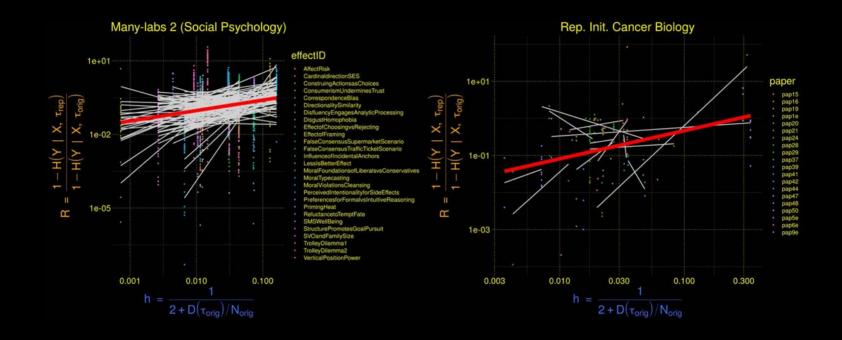
$$k_r h_r = k_o h_o 2^{-\lambda \cdot d}$$

$$\log \frac{k_r}{k_o} = \log \frac{h_o}{h_r} - \lambda \cdot d$$

$$R \equiv \log \frac{H(Y) - H(Y|X, \tau_r)}{H(Y) - H(Y|X, \tau_o)} = \alpha + \beta \log \frac{1}{D(\tau_r)/N}$$

## 1) K predicts <u>actual</u> reproducibility

(part of collaboration with Brazilian Reproducibility Initiative)



### independent new predictor

#### multiple regression, Y=reproducibility

	social psychology	cancer biol.
median (ir)reproducibility	-4.76 [-7.01; -2.5] ***	-2.65 [-3.21; -2.1] ***
- $log(2 + D(\tau_{orig}))$	0.47 [0.25; 0.69] ***	1.27 [0.68; 1.86] ***
$\log(P_{\text{orig}})$	-0.13 [-0.15; -0.1] ***	0 [0; 0.01]
$\sqrt{N_{orig}}$	0.01 [-0.11; 0.13]	0.19 [0.08; 0.3] ***

### very easy to measure, automatize

#### multiple regression, Y=reproducibility

	social psychology	cancer biol.
median (ir)reproducibility	-4.76 [-7.01; -2.5] ***	-2.65 [-3.21; -2.1] ***
- $\log(2 + D(\tau_{orig}))$	0.47 [0.25; 0.69] ***	1.27 [0.68; 1.86] ***
$\log(P_{orig})$	-0.13 [-0.15; -0.1] ***	0 [0; 0.01]
$\sqrt{N_{orig}}$	0.01 [-0.11; 0.13]	0.19 [0.08; 0.3] ***

(here  $D(\tau)$  based on sentences in replication protocol!)

### better than just P-values and N

### multiple regression, Y=reproducibility R

	social psychology	cancer biol.
median (ir)reproducibility	-4.76 [-7.01; -2.5] ***	-2.65 [-3.21; -2.1] ***
- $\log(2 + D(\tau_{\text{orig}}))$	0.47 [0.25; 0.69] ***	1.27 [0.68; 1.86] ***
$\log(P_{orig})$	-0.13 [-0.15; -0.1] ***	0 [0; 0.01]
$\sqrt{N_{orig}}$	0.01 [-0.11; 0.13]	0.19 [0.08; 0.3] ***

## leads to progress in metascience

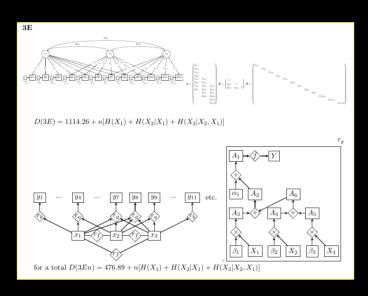
#### multiple regression, Y=reproducibility

	social psychology	cancer biol.
median (ir)reproducibility	-4.76 [-7.01; -2.5] ***	-2.65 [-3.21; -2.1] ***
- $\log(2 + D(\tau_{orig}))$	0.47 [0.25; 0.69] ***	1.27 [0.68; 1.86] ***
$\log(P_{\text{orig}})$	-0.13 [-0.15; -0.1] ***	0 [0; 0.01]
$\sqrt{N_{orig}}$	0.01 [-0.11; 0.13]	0.19 [0.08; 0.3] ***

# 2) D(τ) might predict fitting propensity

# preregistered test

### preregistered test

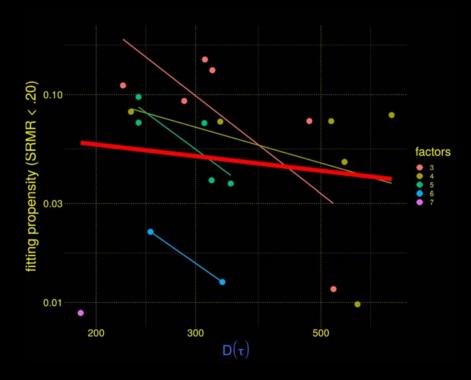


- 1) generated N=20 models, all with 36 parameters
- 2) derived D(t), predicted their **fitting propensity**
- 3) tested them on 20,000 random covariance matrices

(Fanelli & Bonifay, in prep)

### D(τ) uniquely reflects model complexity

(pre-registered test)



(Fanelli & Bonifay, in progress)

### NO alternative theory explains this!

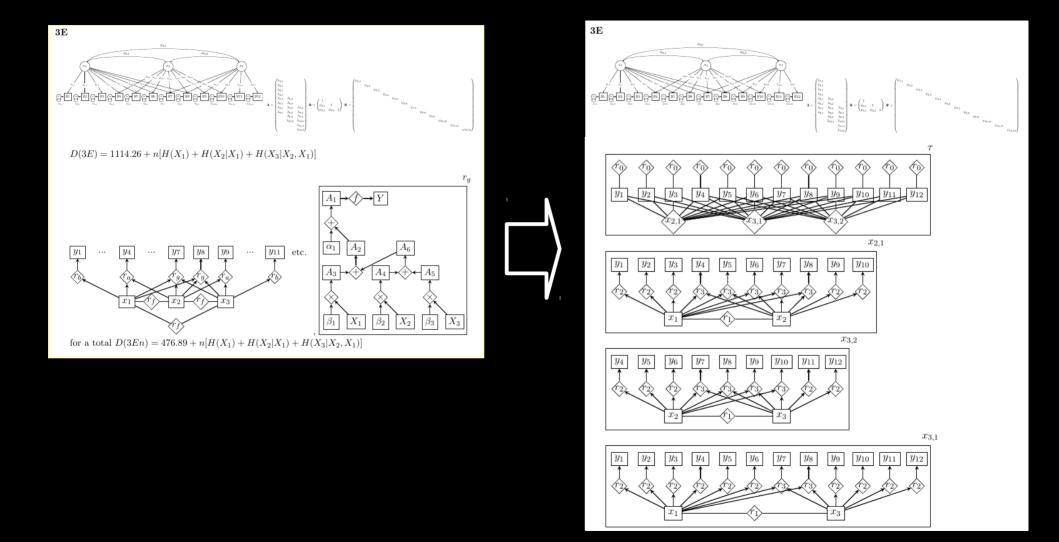
(pre-registered test)

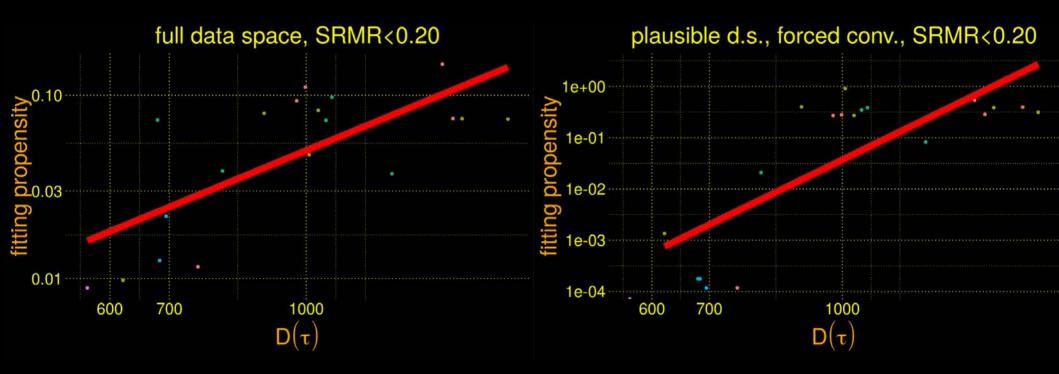
multiple regression: Y=fitting propensity

```
intercept 0.23 [0.16; 0.31] ***

D(τ) -1.46 [-2.43; -0.48] **

n. factors -2.71 [-3.86; -1.55] ***
```





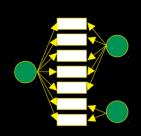
Spearman's  $\rho = 0.64$ , P<0.002

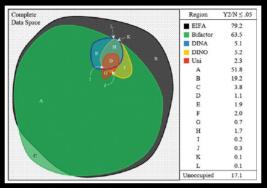
Spearman's  $\rho = 0.73$ , P<0.001

### how K improves theory testing

example of application

Bifactor confirmatory (theoretical) 20 parameters



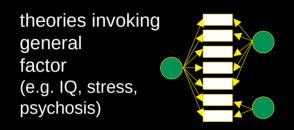


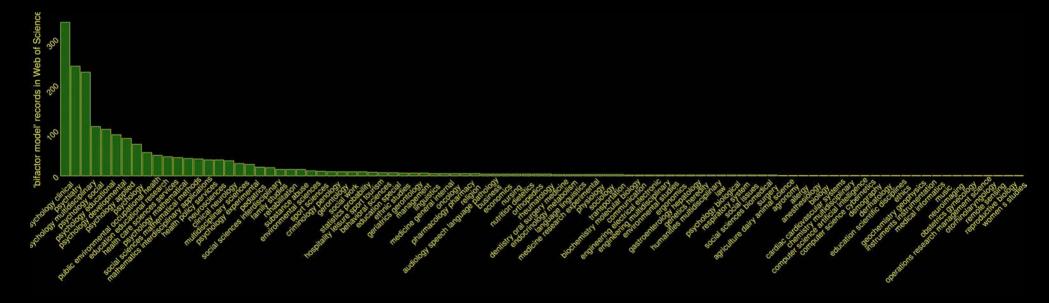




EIFA: exploratory (a-theoretical) 20 parameters

### Bifactor widely used to "test" theories

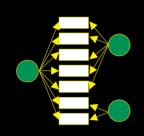


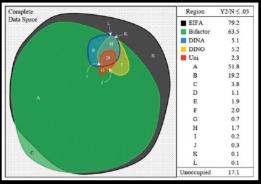


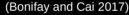
### when is a theory *actually* supported?

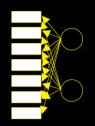
example of application

Bifactor confirmatory (theoretical) 20 parameters









EIFA: exploratory (a-theoretical) 20 parameters





"The [bifactor-encoded] theory is specifically supported by the data"

(Fanelli & Bonifay, in progress)

### Summary of this talk:

- 1) what's missing in the current paradigm?
  - we pretend **complexity** is irrelevant
- 2) what might a new paradigm look like?
  - measuring D(τ), integrating/penalizing with K
- 3) evidence in support of this proposal?
  - increasingly promising
  - any alternative, better suggestions?