From causal salads to causal inference

Francisco Rodríguez-Sánchez

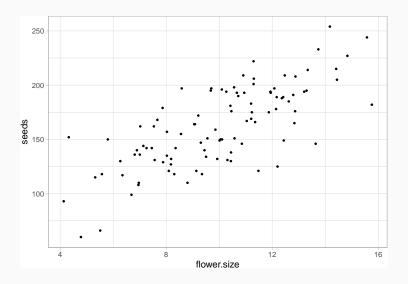
https://frodriguezsanchez.net



Self-learnt stuff ahead



Larger flowers produce more seeds



Larger flowers produce more seeds

lm(seeds ~ flower.size)

Variable	Beta	SE	p.value	
(Intercept)	57	10.1	<0.001	
flower.size	11	0.978	<0.001	

Does flower size

really cause

increased seed production?

Shall we select plants with large flowers to increase seed production?

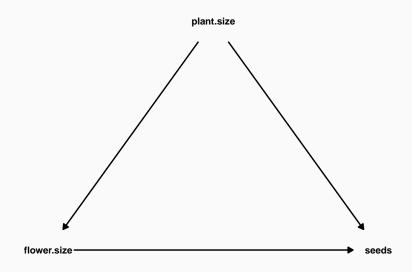
Shall we select plants with large flowers to increase seed production?

We tried but didn't get the expected benefits

Maybe large plants (e.g. growing on better soil) have large flowers AND produce more seeds?



Maybe plant size is a **CONFOUNDER**?



Adjusting for plant size (confounding)

lm(seeds ~ flower.size + plant.size)

Variable	Beta	SE	p.value	
(Intercept)	12	12.9	0.4	
flower.size	6.6	1.18	<0.001	
plant.size	0.82	0.168	<0.001	

Including pollinators (bees)

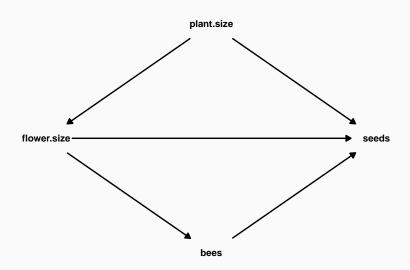


Including pollinators (bees)

lm(seeds ~ flower.size + plant.size + bees)

Variable	Beta	SE	p.value
(Intercept)	5.2	12.1	0.7
flower.size	2.1	1.56	0.2
plant.size	0.90	0.157	<0.001
bees	8.8	2.14	<0.001

Pollinators are a **MEDIATOR**



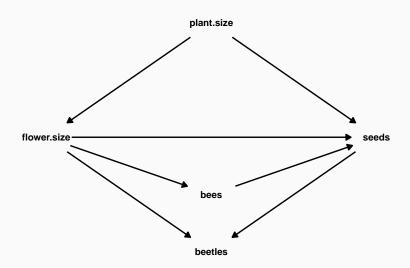
Including beetles (pollen & seed predators)

lm(seeds ~ flower.size + plant.size + bees +
beetles)

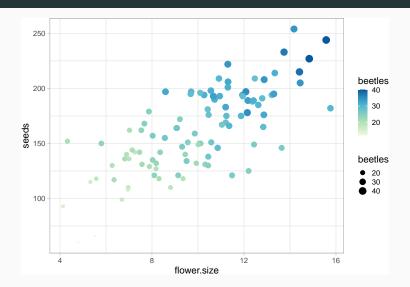
Variable	Beta	SE	p.value
(Intercept)	-11	8.67	0.2
flower.size	-3.8	1.25	0.003
plant.size	0.47	0.118	< 0.001
bees	4.8	1.56	0.003
beetles	5.2	0.529	<0.001

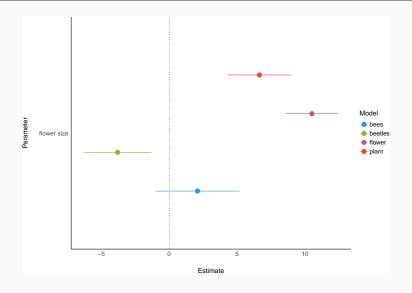
Now flower.size has negative coefficient!!

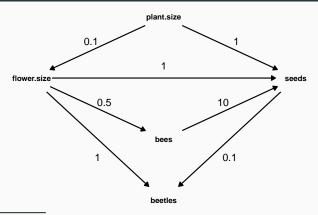
Beetles are a **COLLIDER**



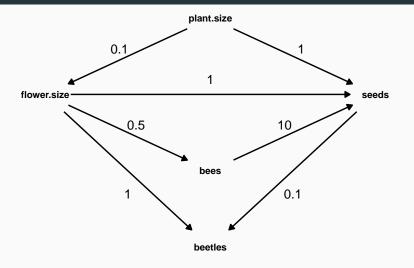
Colliders induce negative relation between treatment (flower.size) and outcome (seeds)



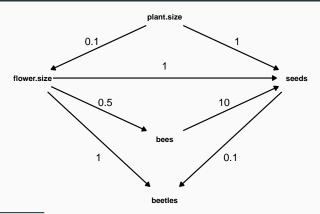




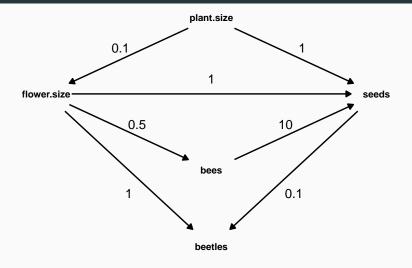
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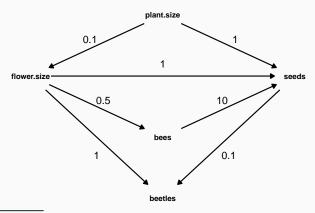
AVOID COLLIDERS!



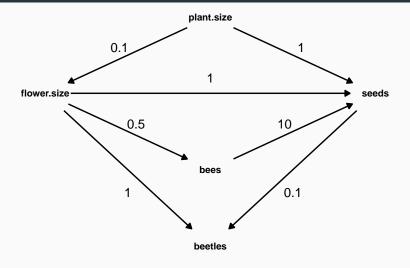
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MEDIATORS split total effect into direct and indirect effects



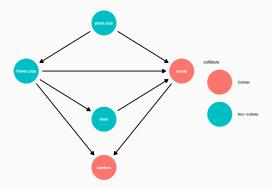
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Include CONFOUNDERS to avoid bias (backdoor criterion)

Tools to identify correct causal structure

```
dagify(
  seeds ~ plant.size + flower.size + bees,
  flower.size ~ plant.size,
  bees ~ flower.size,
  beetles ~ flower.size + seeds,
  coords = coords
) |>
  ggdag_collider(size = 2) + theme_dag_blank()
```



Causal salads

Causal salads

You put everything into a regression equation, toss with some creative story-telling, and hope the reviewers eat it

R. McElreath



Jerry Pank

Causal salads

Ian Vanhove

Throwing predictor variables into a statistical model hoping this will improve the analysis is a dreadful idea

Predictive criteria don't help for causal inference

Predictive criteria don't help to choose correct causal model

Making good predictions doesn't require accurate causal model

Model	AIC	R2
m.flower	933.3	0.5
m.flower.plant	913.2	0.6
m.flower.plant.bees	899.1	0.7
m.flower.plant.bees.beetles	829.9	0.8

[&]quot;Best model" (based on AIC or R2) not good for causal inference

Simpler (best) model provides biased causal estimates

Simulate response depending on two correlated variables (Hartig 2022)

```
x1 = runif(100)
x2 = 0.8*x1 + 0.2*runif(100)
y = x1 + x2 + rnorm(100)
```

У	x1	x2
-0.1	0.3	0.4
1.7	0.8	0.7
0.6	0.4	0.4
1.4	0.9	0.9
0.8	0.9	0.8
0.2	0.0	0.2

Simpler (best) model provides biased causal estimates

Simulate response depending on two correlated variables (Hartig 2022)

```
fullmodel = lm(y \sim x1 + x2)
```

Residual standard error: 0.9765 on 97 degrees of freedom Multiple R-squared: 0.237, Adjusted R-squared: 0.2212 F-statistic: 15.06 on 2 and 97 DF, p-value: 2.009e-06

Simpler (best) model provides biased causal estimates

```
simplemodel = MASS::stepAIC(fullmodel, trace = 0)
Call:
lm(formula = v \sim x1)
Residuals:
   Min 10 Median 30 Max
-1.9047 -0.6292 -0.1019 0.6077 3.3394
Coefficients:
          Estimate Std. Error t value Pr(>|t|)
x1 1.88350 0.34295 5.492 3.13e-07 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.9725 on 98 degrees of freedom
Multiple R-squared: 0.2353, Adjusted R-squared: 0.2275
F-statistic: 30.16 on 1 and 98 DF, p-value: 3.134e-07
```

Automated model selection (dredge)

Simulating data with 10 random predictors

У	x.1	x.2	x.3	X.4	x.5	x.6	x.7	x.8	x.9	x.10
-0.1	0.6	0.6	0.3	0.8	0.2	0.0	0.4	0.4	0.3	0.2
0.8	0.4	0.4	0.9	0.2	0.5	0.1	0.6	0.2	0.0	0.0
-0.5	0.0	0.3	0.4	0.3	0.1	0.1	0.9	0.9	0.5	0.8
-0.6	0.7	0.7	0.4	0.5	0.2	0.7	0.7	0.8	0.5	0.3
0.7	0.0	0.6	0.9	0.1	0.2	0.4	0.8	0.6	0.6	0.1
-0.1	0.4	0.2	0.9	0.4	0.6	0.5	0.9	0.1	0.8	0.8

Automated model selection

Running MuMIn::dredge with 10 random predictors

```
full.model <- lm(y ~ ., data = dat)

dd <- MuMIn::dredge(full.model)</pre>
```

Best model:

Parameter	Coefficient	SE	р
(Intercept)	-1.50	0.36	0.00
x.2	0.78	0.36	0.03
x.5	0.59	0.32	0.07
x.6	0.61	0.35	0.09
x.9	0.87	0.34	0.01

Extract from dredge help

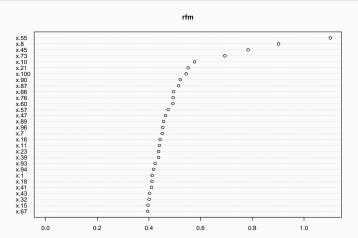
"Let the computer find out" is a poor strategy and usually reflects the fact that the researcher did not bother to think clearly about the problem of interest and its scientific setting

Burnham and Anderson 2002

Variable importance in machine learning

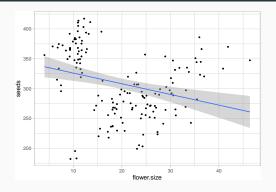
Random forest on 100 random predictors

```
dat <- data.frame(x = matrix(runif(50000), ncol = 100), y = runif(500))
rfm <- randomForest::randomForest(y ~ ., data = dat)
varImpPlot(rfm)</pre>
```



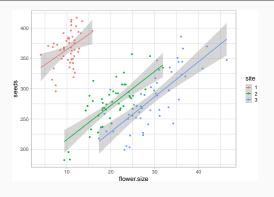
Simpson's paradox as a causal problem

Simpson's paradox



Variable	Beta	SE	p.value
(Intercept)	344	10.7	<0.001
flower.size	-1.8	0.486	<0.001

Simpson's paradox

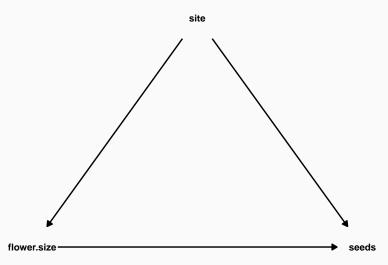


Variable	Beta	SE	p.value
(Intercept)	308	6.50	<0.001
flower.size	5.7	0.500	<0.001
site			
1			

40

Simpson's paradox

Site is a confounder!



Causal interpretation requires

external knowledge

To estimate causal effects accurately we require more information than can be gleaned from statistical tools alone

D'Agostino et al

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D'Agostino et al

No amount of data reliably turns salad into sense

R. McElreath

· Draw generative model (causal graph) beforehand

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- · Predictive criteria not fit for causal inference

To learn more

Suchinta Arif's papers

McElreath's workshop on causal inference

https://www.r-causal.org

https://theeffectbook.net