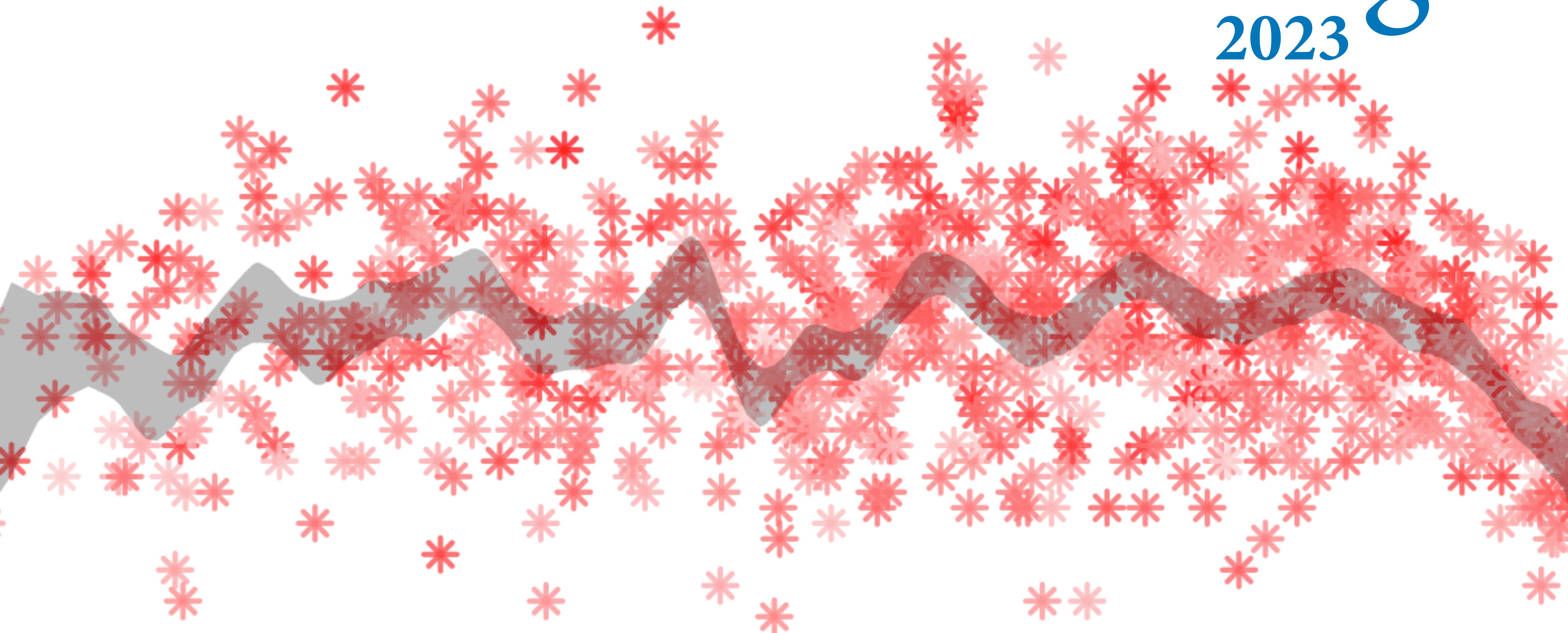
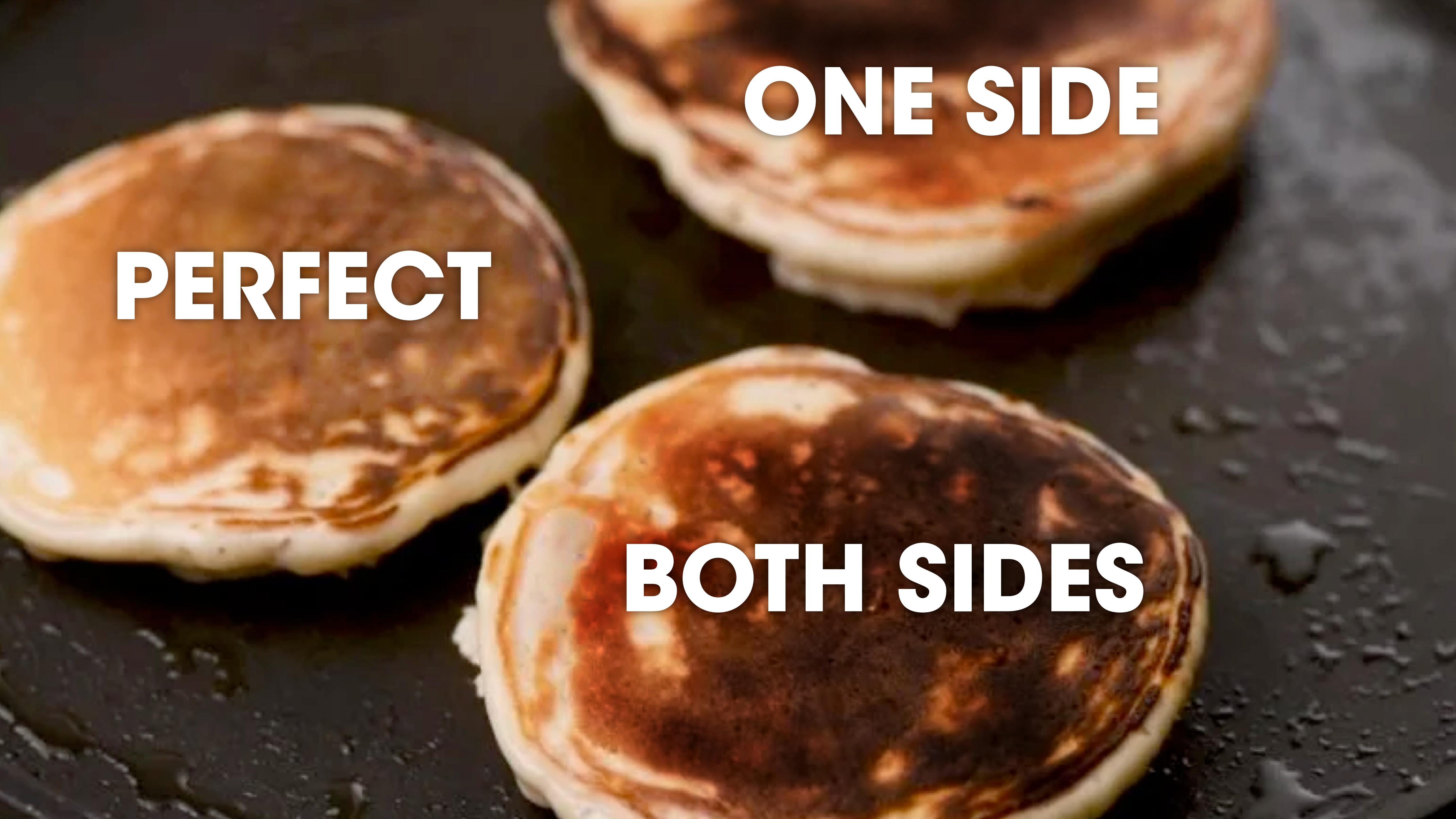


Statistical Rethinking

2023



17. Measurement & Misclassification

A close-up photograph of several round pancakes cooking in a dark, non-stick frying pan. The pancakes are stacked in a column, showing their fluffy, light-colored undersides and perfectly browned, slightly焦化的 tops. The lighting highlights the texture and color of the pancakes.

PERFECT

ONE SIDE

BOTH SIDES

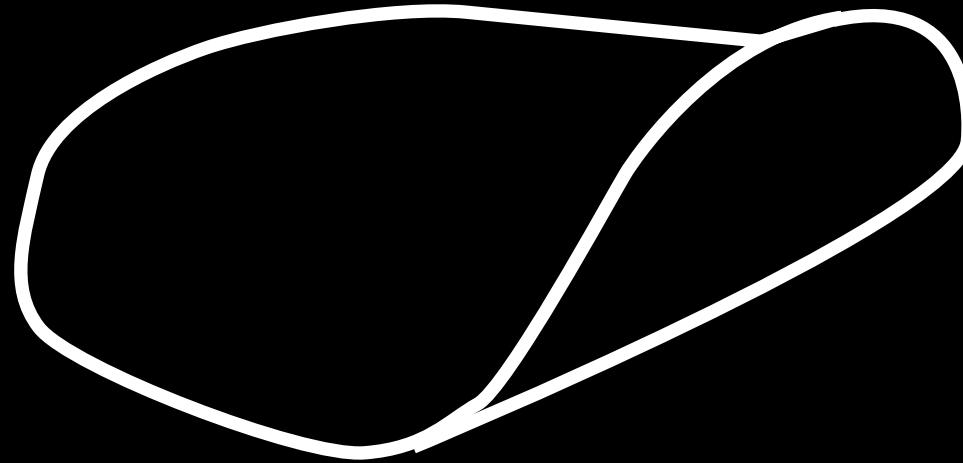
1



2



3



You are served:



What's the probability the other side is also burnt?

$$\Pr(\text{burnt down} \mid \text{burnt up}) = \frac{\Pr(\text{burnt up, burnt down})}{\Pr(\text{burnt up})}$$

$$\Pr(\text{burnt down} \mid \text{burnt up}) = \frac{\Pr(\text{burnt up, burnt down})}{\Pr(\text{burnt up})}$$



$$\Pr(\text{burnt up}) = (1/3)(1) + (1/3)(0.5) + (1/3)(0) = 1/2$$

$$\Pr(\text{burnt down} \mid \text{burnt up}) = \frac{\Pr(\text{burnt up, burnt down})}{\Pr(\text{burnt up})}$$



$$\Pr(\text{burnt up}) = (1/3)(1) + (1/3)(0.5) + (1/3)(0) = 1/2$$

$$\Pr(\text{burnt down} \mid \text{burnt up}) = \frac{1/3}{1/2} = \frac{2}{3}$$

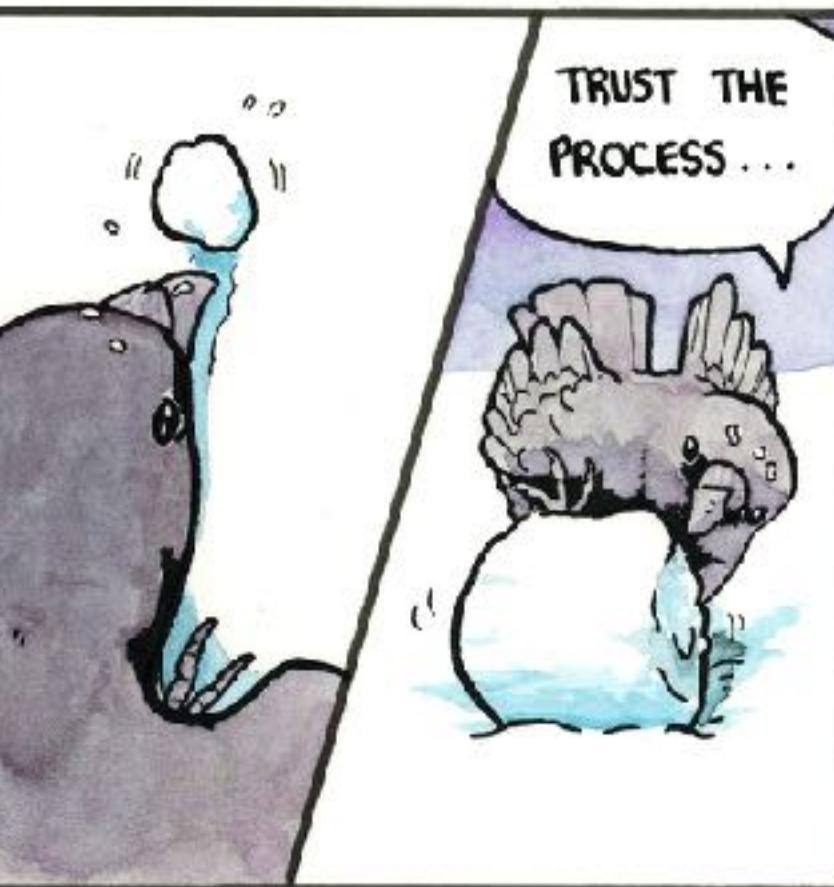
The Importance of Not Being Clever

Being clever is unreliable and opaque

Better to follow the axioms

Probability theory provides solutions
to challenging problems, if only we'll
follow the rules

Often nothing else to “understand”



Ye Olde Causal Alchemy

The Four Elemental Confounds

The Fork

$$X \leftarrow Z \rightarrow Y$$

The Pipe

$$X \rightarrow Z \rightarrow Y$$

The Collider

$$X \rightarrow Z \leftarrow Y$$

The Descendant

$$X \rightarrow Z \rightarrow Y$$

↓
A

Measurement Error

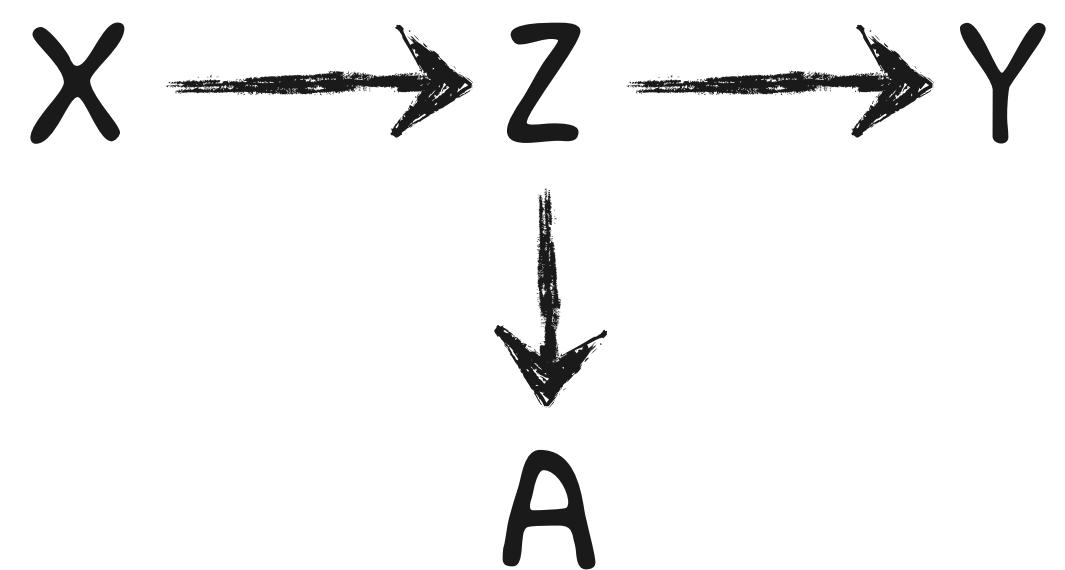
Many variables are proxies of the causes
of interest

Common to ignore measurement

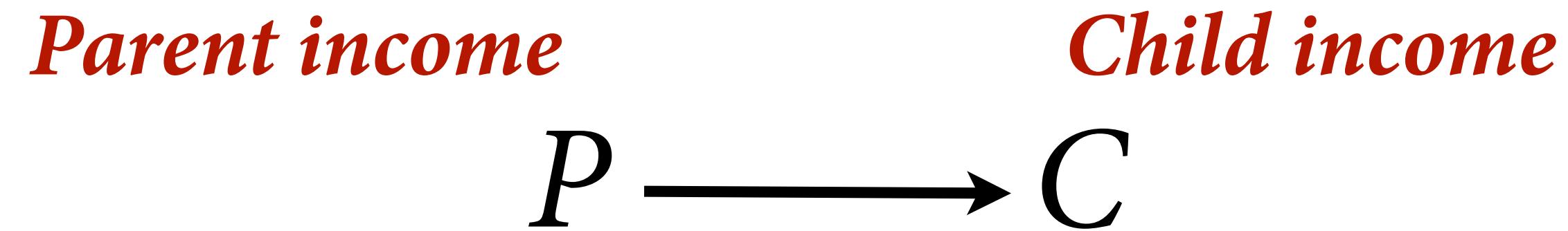
Many ad hoc procedures

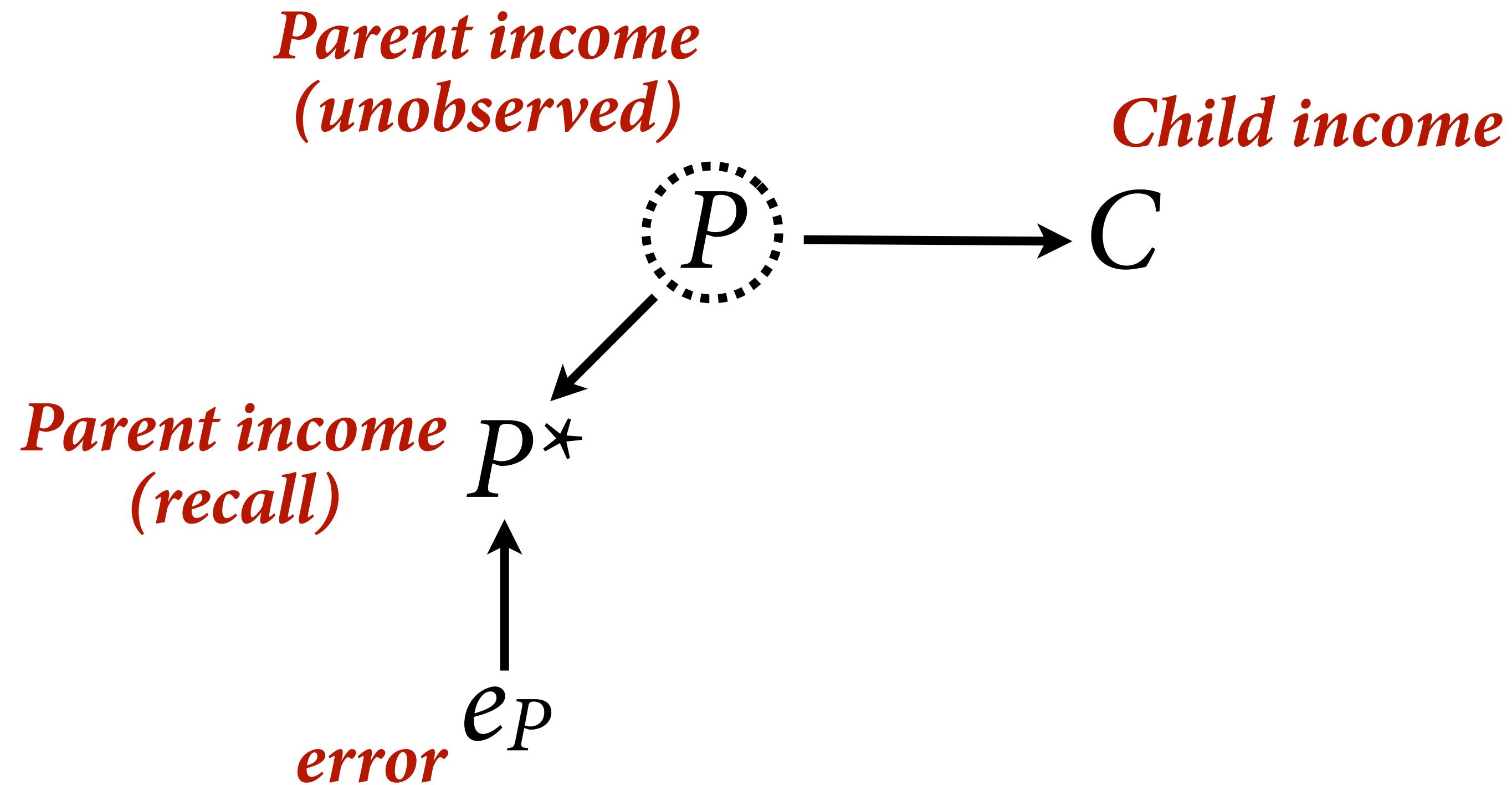
Think causally, lean on probability theory

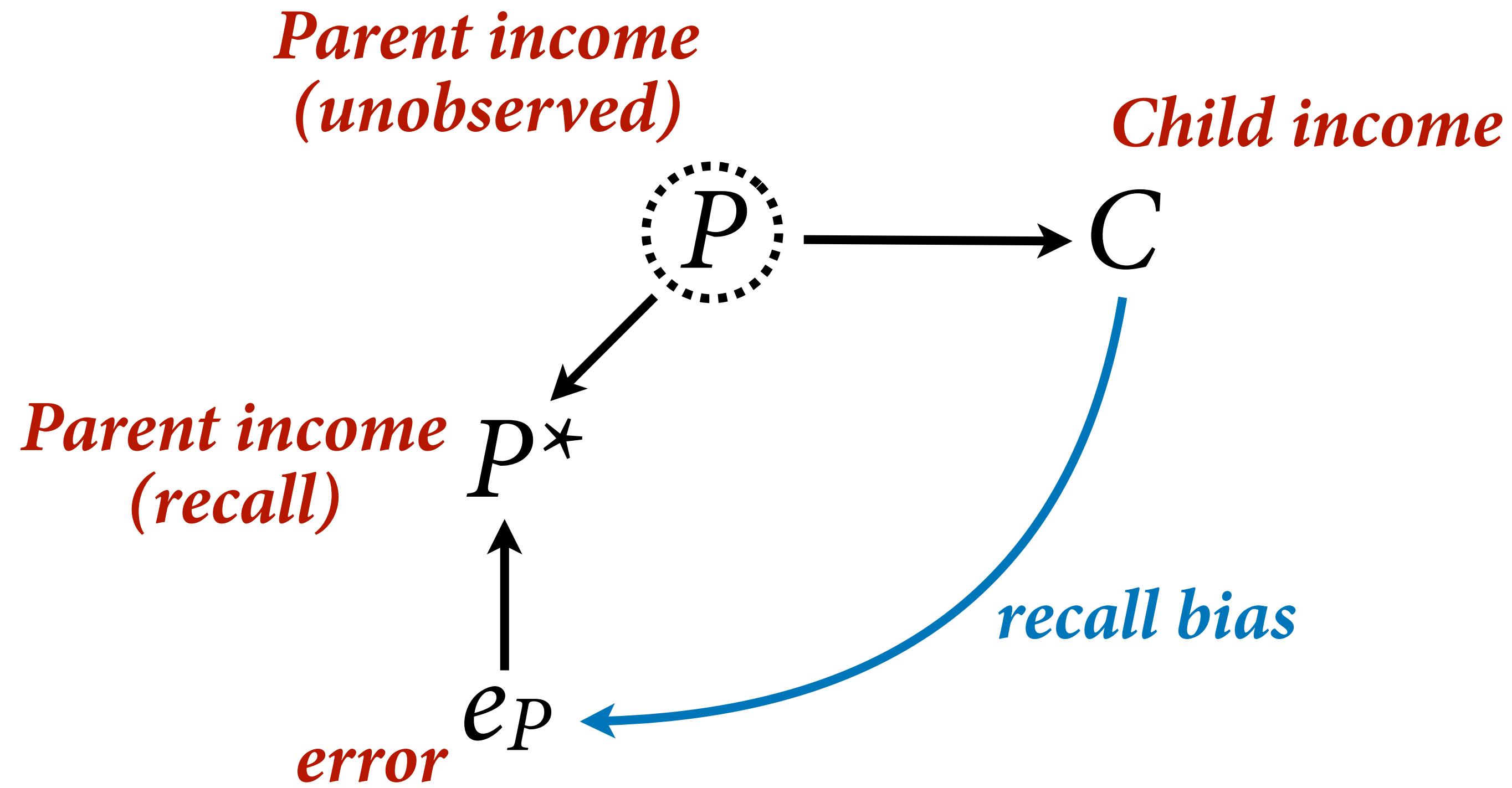
The Descendant



Myth: Measurement error only reduces effect estimates, never increases them







```

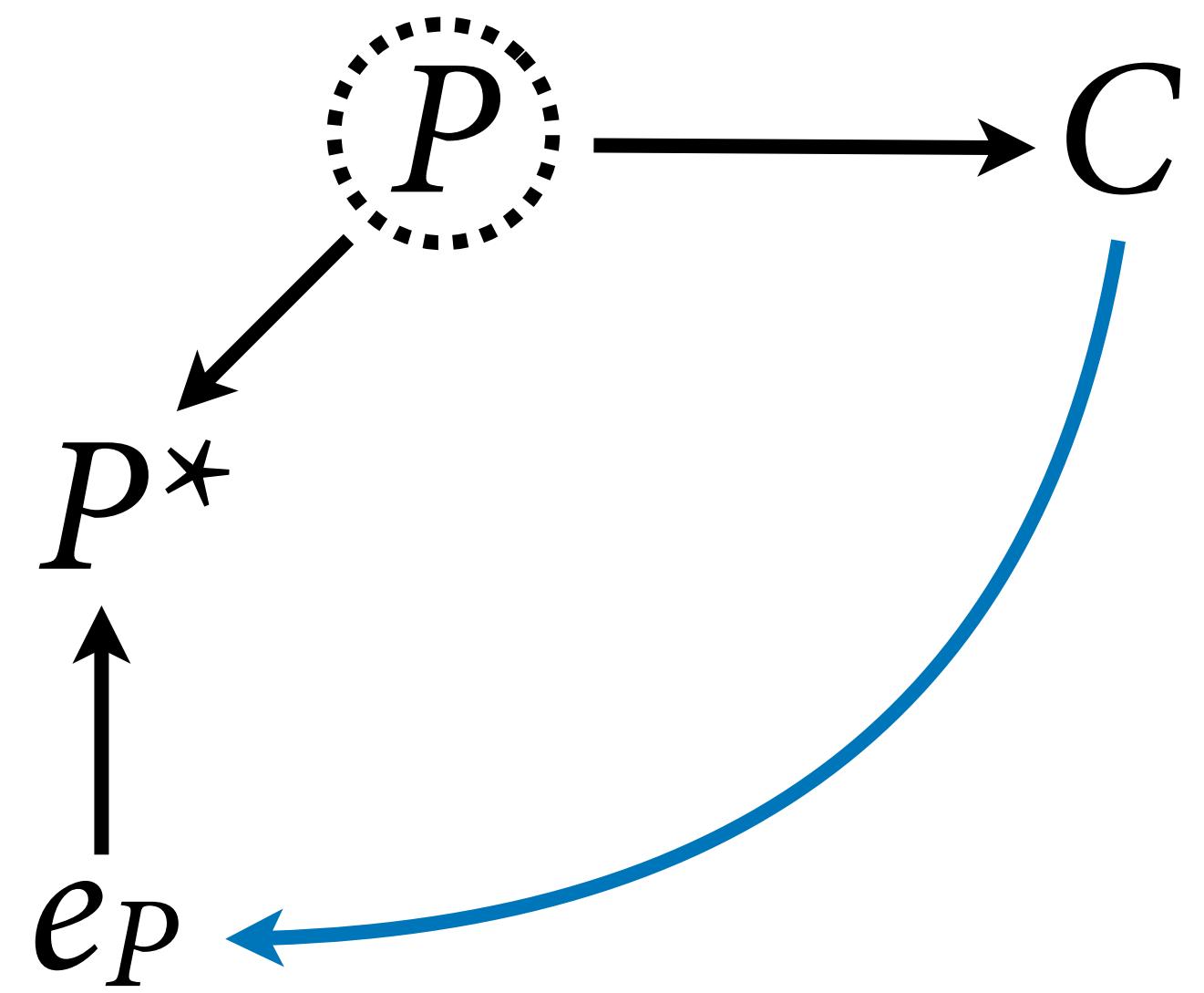
# simple parent-child income example
# recall bias on parental income

N <- 500
P <- rnorm(N)
C <- rnorm(N, 0*P)
Pstar <- rnorm(N, 0.8*P + 0.2*C )

mCP <- ulam(
  alist(
    C ~ normal(mu,sigma),
    mu <- a + b*P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

precis(mCP)

```



```

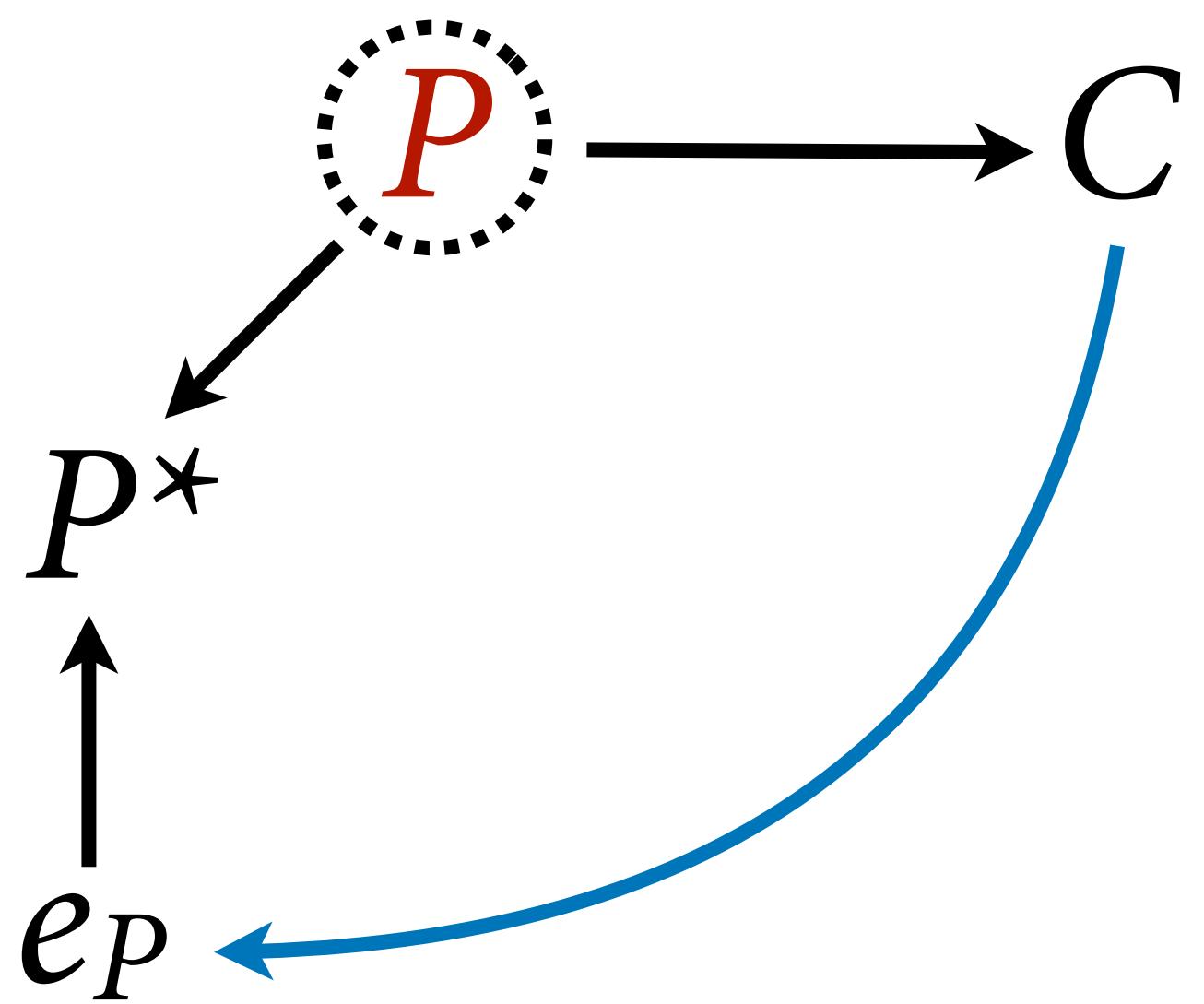
# simple parent-child income example
# recall bias on parental income

N <- 500
P <- rnorm(N)
C <- rnorm(N, 0×P)
Pstar <- rnorm(N, 0.8×P + 0.2×C)

mCP <- ulam(
  alist(
    C ~ normal(mu,sigma),
    mu <- a + b×P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

precis(mCP)

```



```

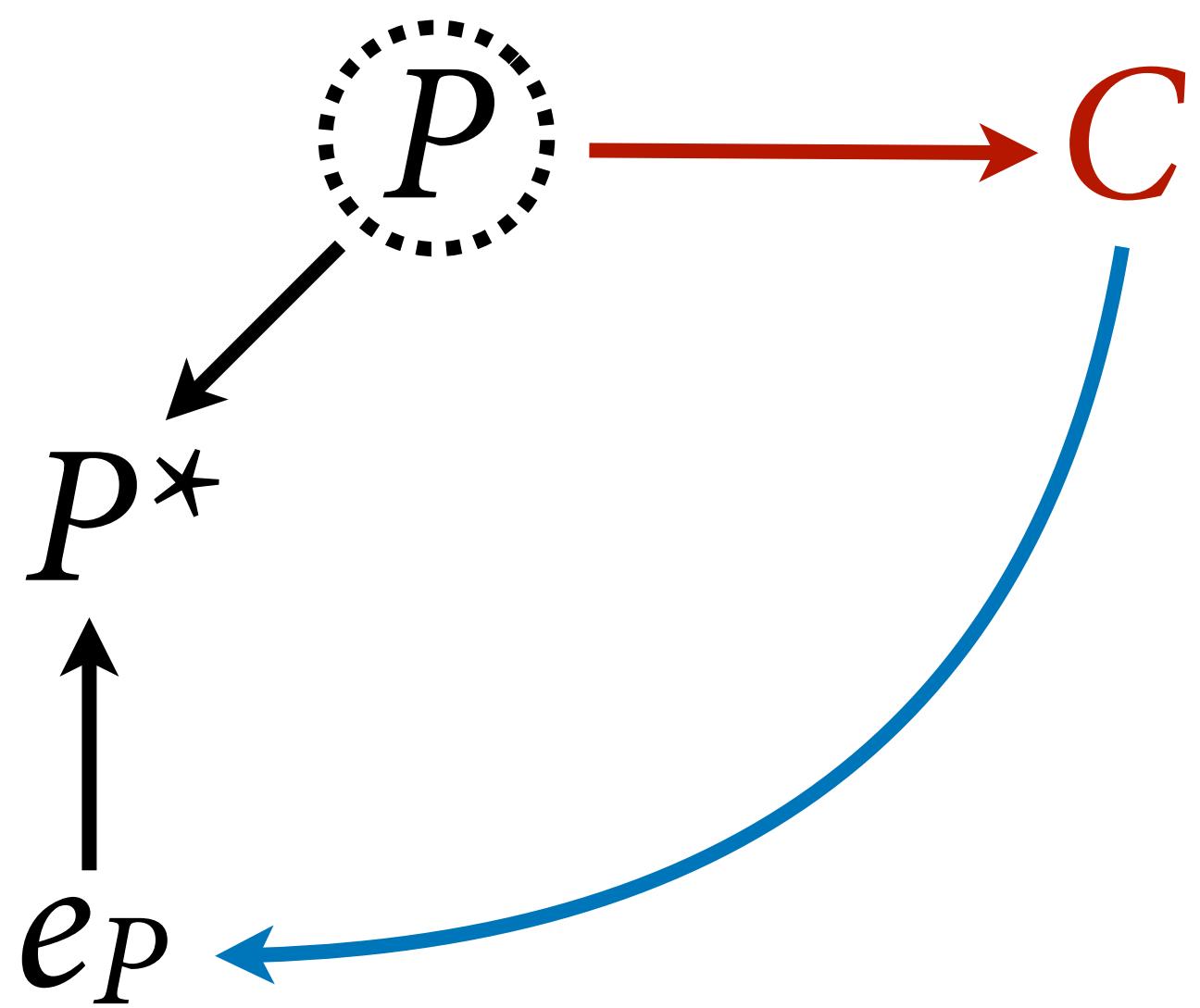
# simple parent-child income example
# recall bias on parental income

N <- 500
P <- rnorm(N)
C <- rnorm(N, 0*P)
Pstar <- rnorm(N, 0.8*xP + 0.2*xC)

mCP <- ulam(
  alist(
    C ~ normal(mu,sigma),
    mu <- a + b*P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

precis(mCP)

```



```

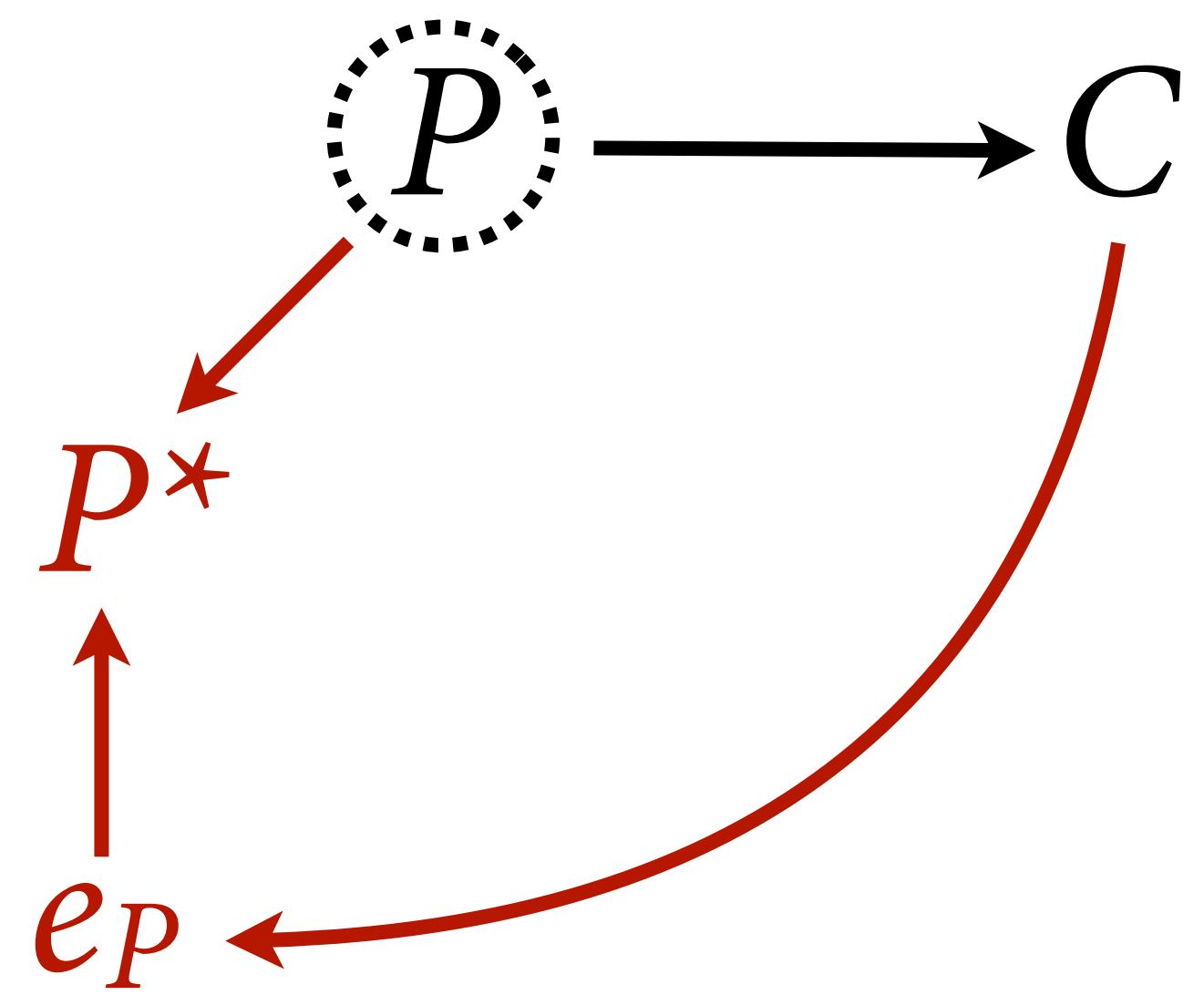
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# recall bias on parental income

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C <- rnorm(N, 0*P)
Pstar <- rnorm(N, 0.8*P + 0.2*C )

mCP <- ulam(
  alist(
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    mu <- a + b*P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

precis(mCP)

```



```

# simple parent-child income example
# recall bias on parental income

N <- 500
P <- rnorm(N)
C <- rnorm(N, 0*P)
Pstar <- rnorm(N, 0.8*P + 0.2*C )

mCP <- ulam(
  alist(
    C ~ normal(mu,sigma),
    mu <- a + b*P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

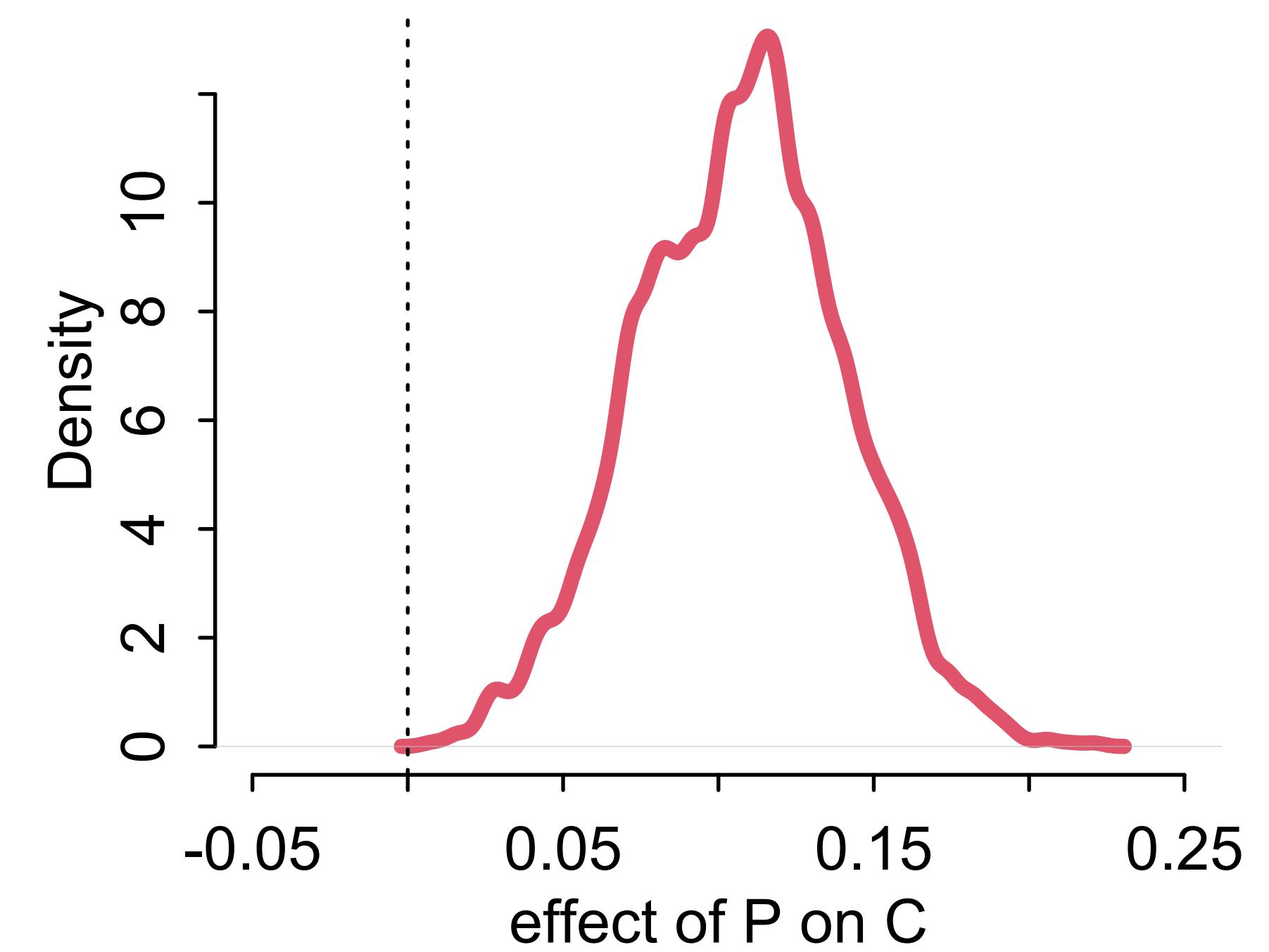
precis(mCP)

```

```

> precis(mCP)
      mean   sd  5.5% 94.5% n_eff Rhat4
a     0.01 0.04 -0.06  0.07  1764    1
b     0.11 0.03  0.05  0.16  1993    1
sigma 0.98 0.03  0.93  1.03  1929    1

```



```

# simple parent-child income example
# recall bias on parental income

N <- 500
P <- rnorm(N)
C <- rnorm(N, 0.75*P) (1)
Pstar <- rnorm(N, 0.8*P + 0.2*C )

```

```

mCP <- ulam(
  alist(
    C ~ normal(mu,sigma),
    mu <- a + b*P,
    a ~ normal(0,1),
    b ~ normal(0,1),
    sigma ~ exponential(1)
  ) , data=list(P=Pstar,C=C) , chains=4 )

```

```

precis(mCP)

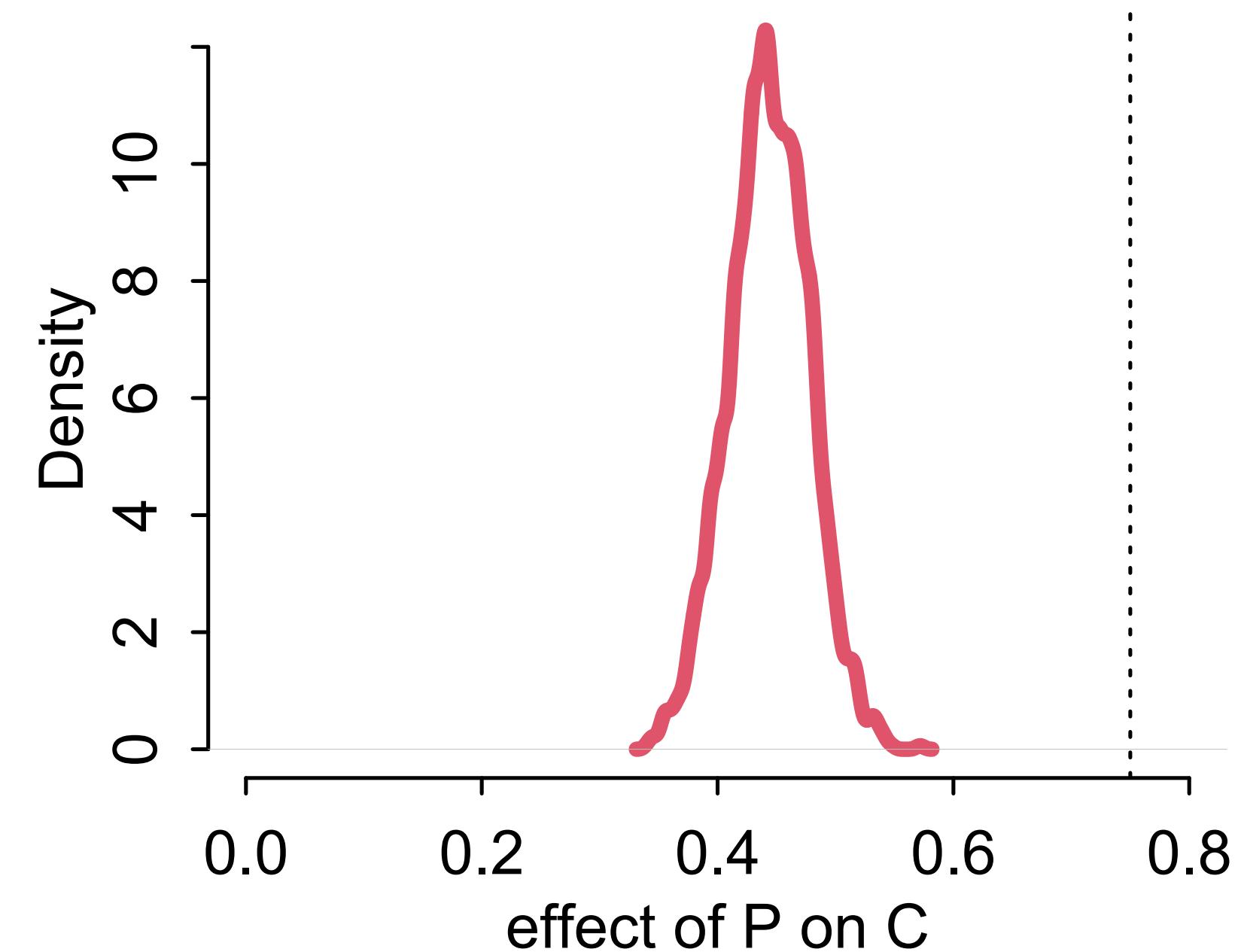
```

```

> precis(mCP)

```

	mean	sd	5.5%	94.5%	n_eff	Rhat4
a	-0.09	0.05	-0.17	-0.02	1782	1
b	0.44	0.03	0.39	0.50	1698	1
sigma	1.05	0.03	1.00	1.11	1947	1



Modeling Measurement

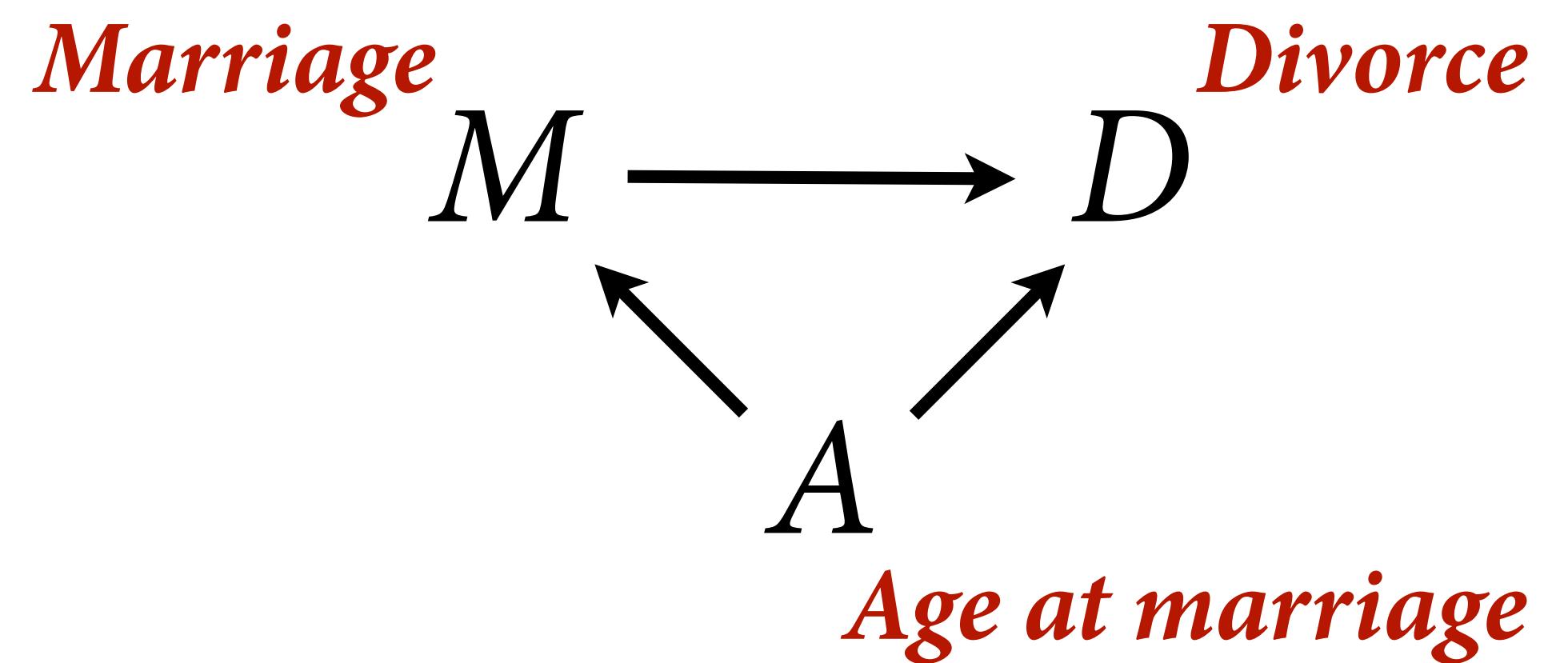
```
data(WaffleDivorce)
```

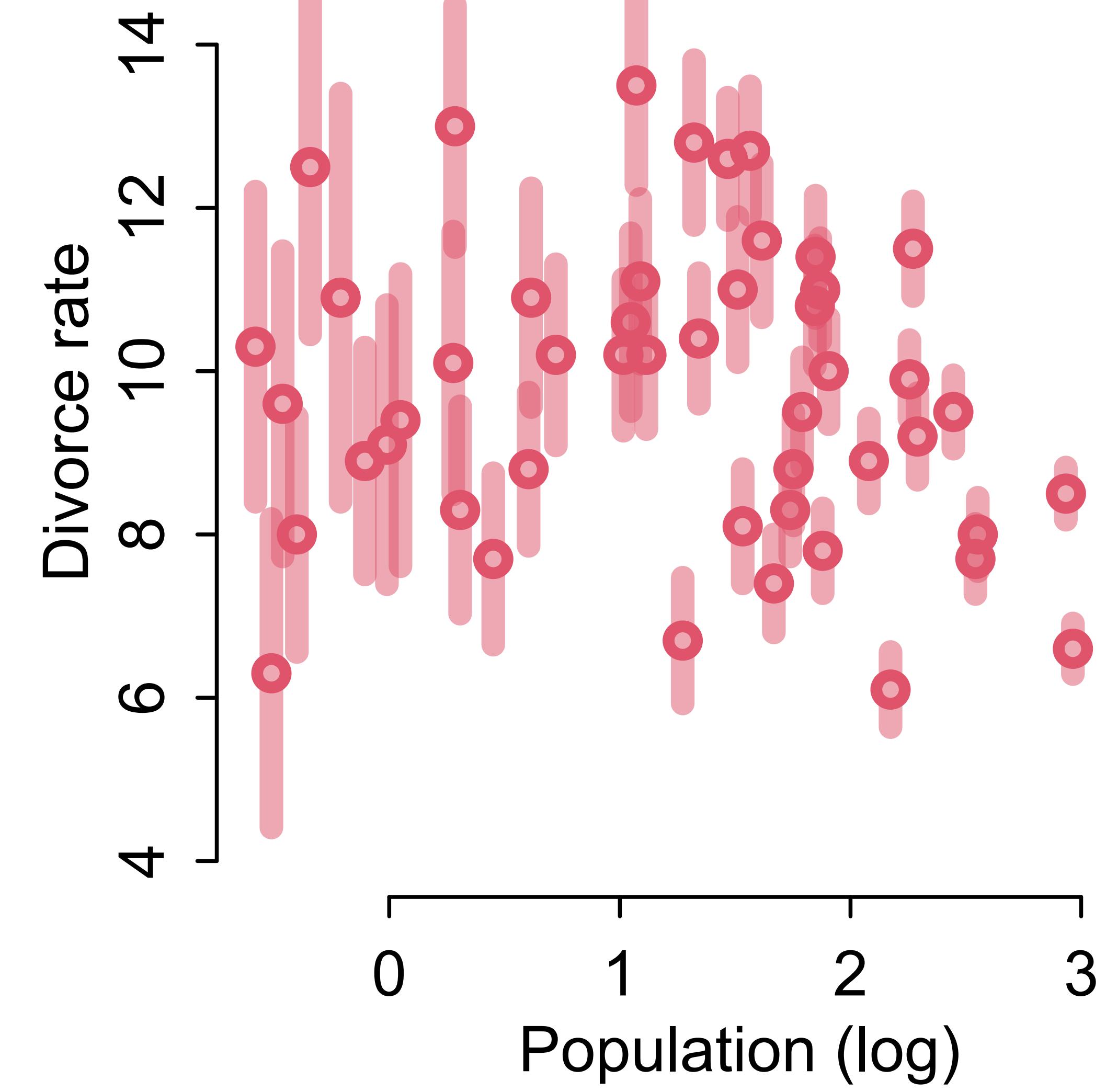
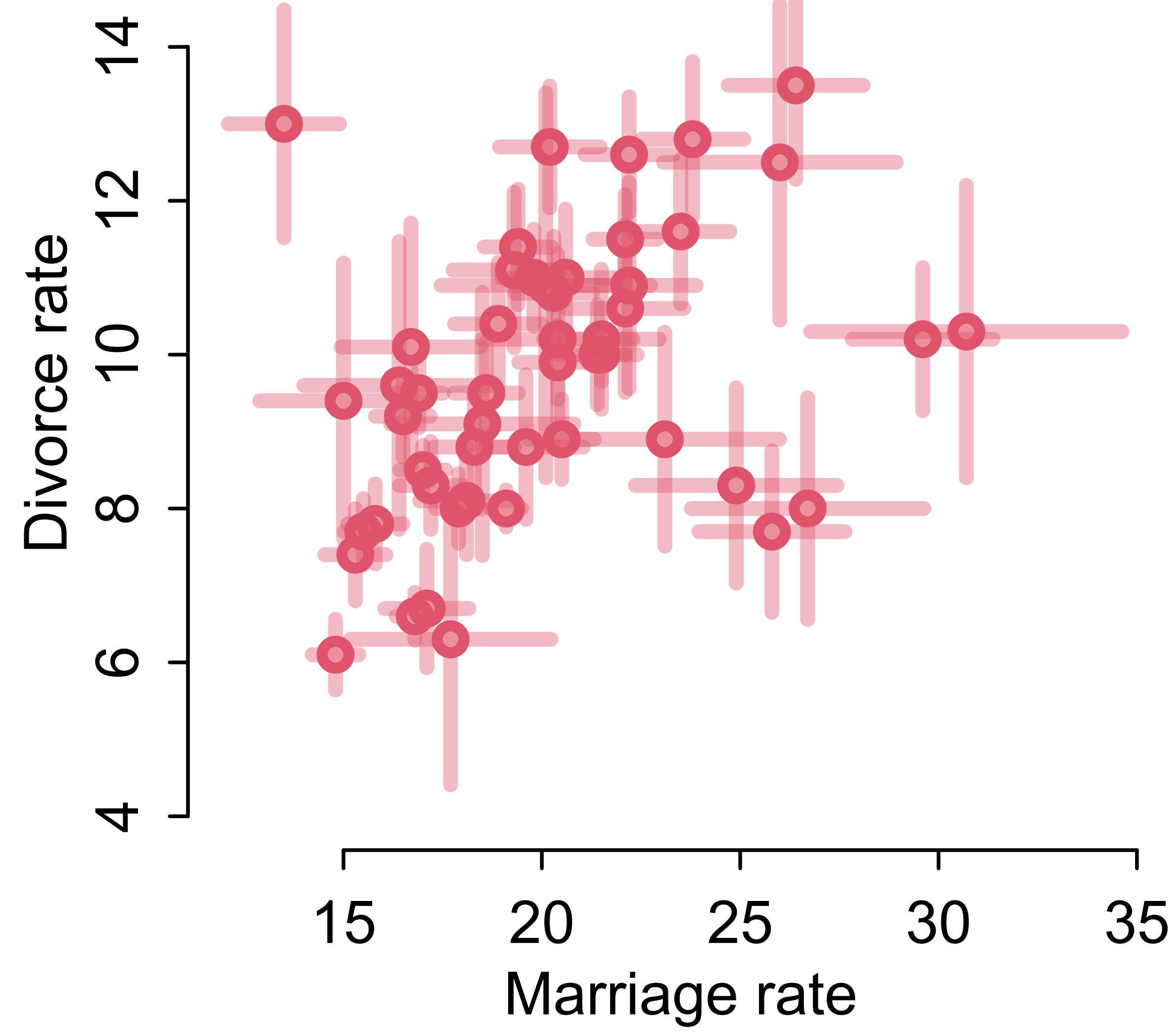
State estimates D, M, A measured
with error, error varies by State

Two problems:

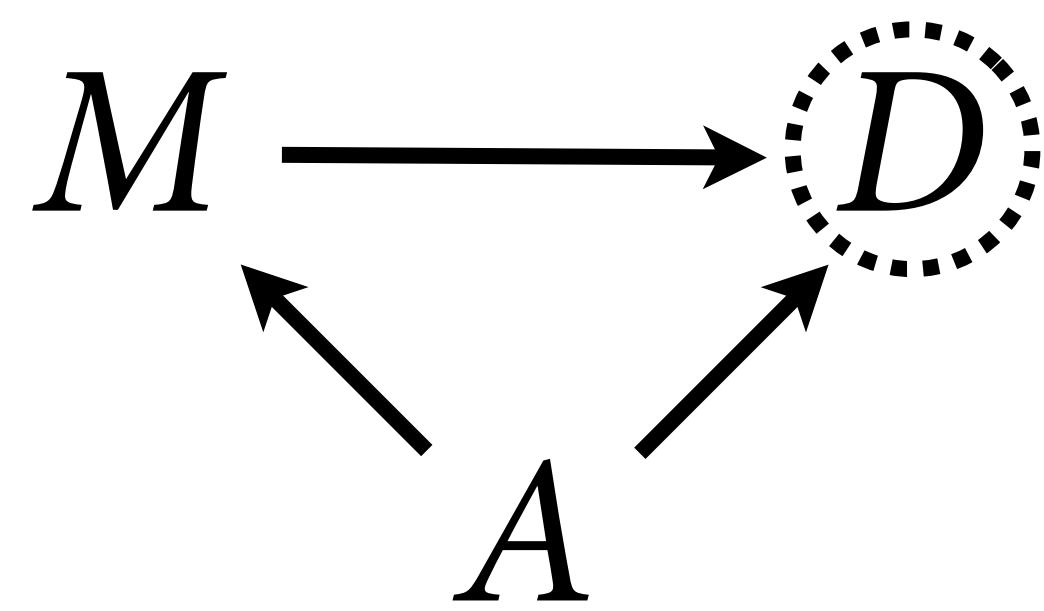
(1) Imbalance in evidence

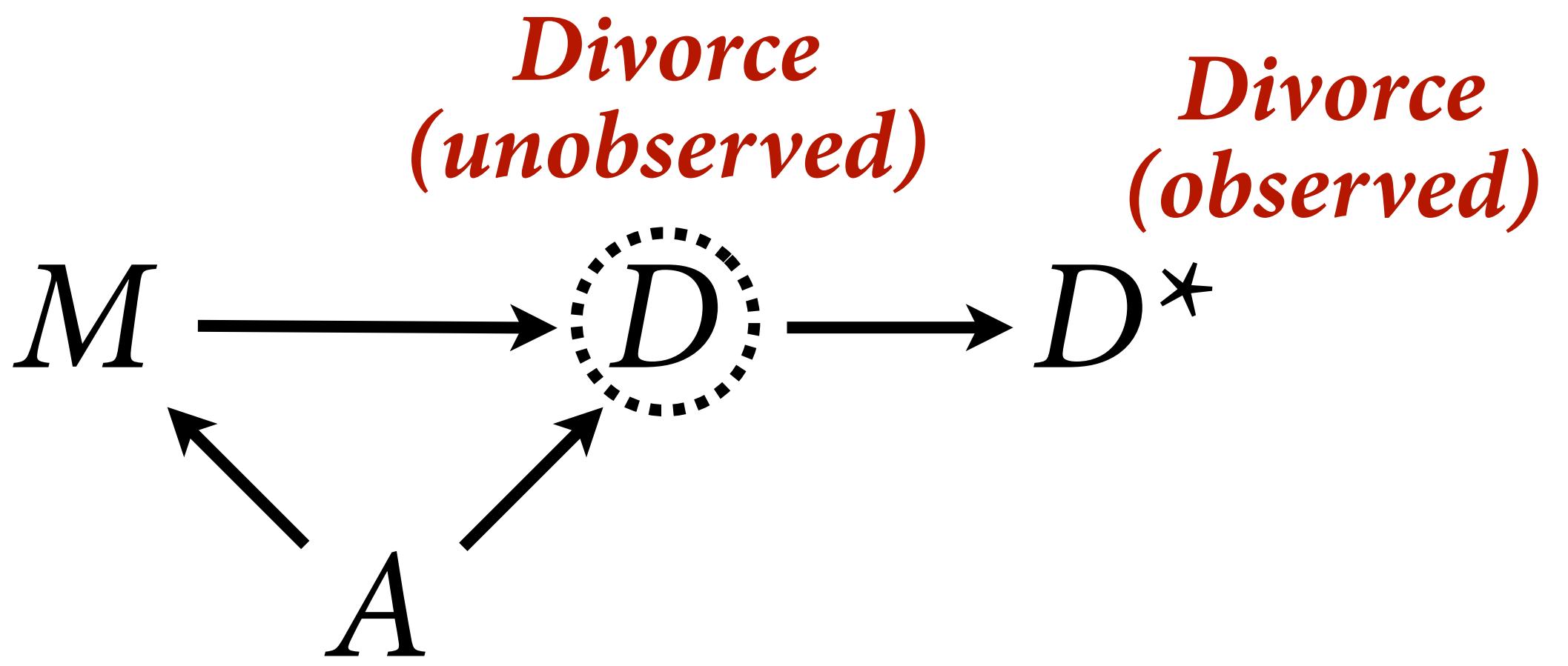
(2) Potential confounding

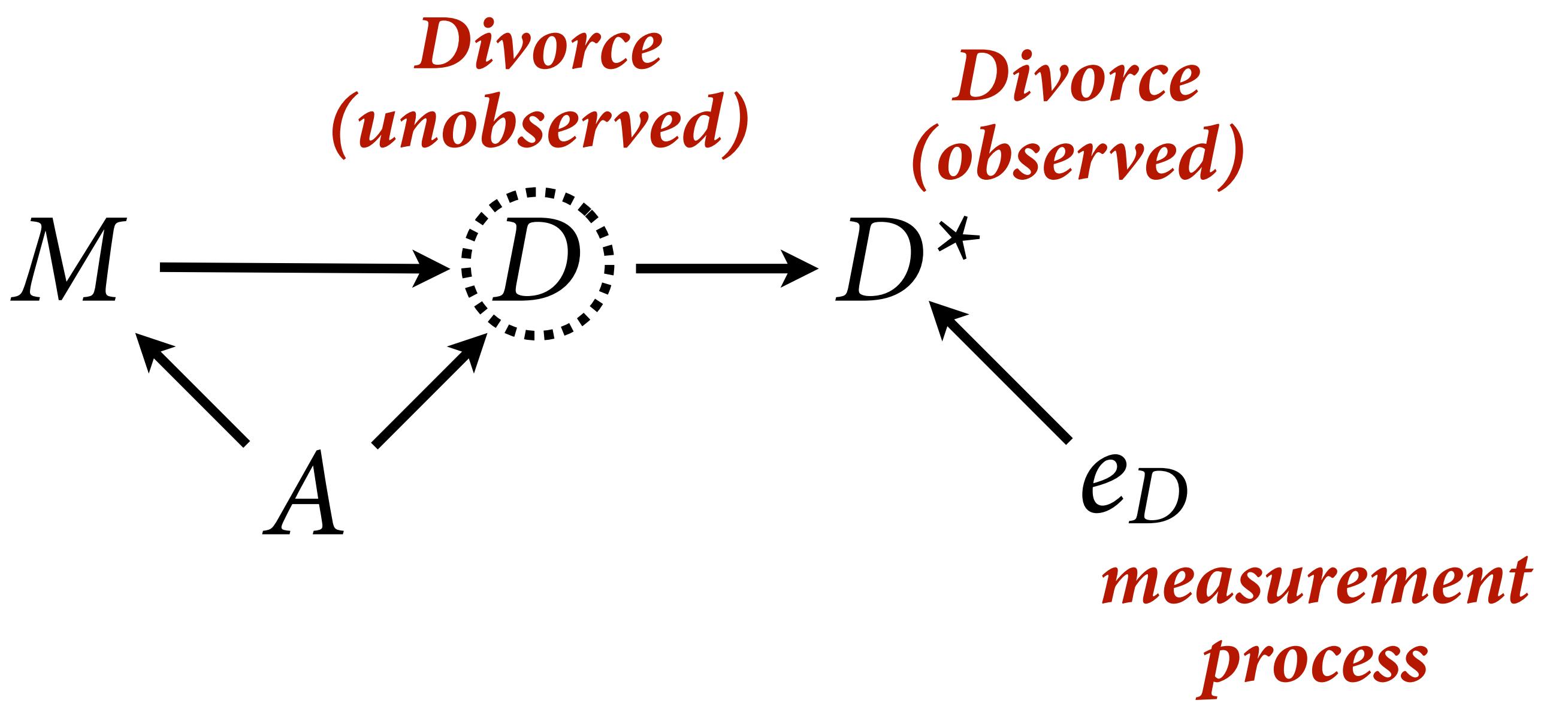


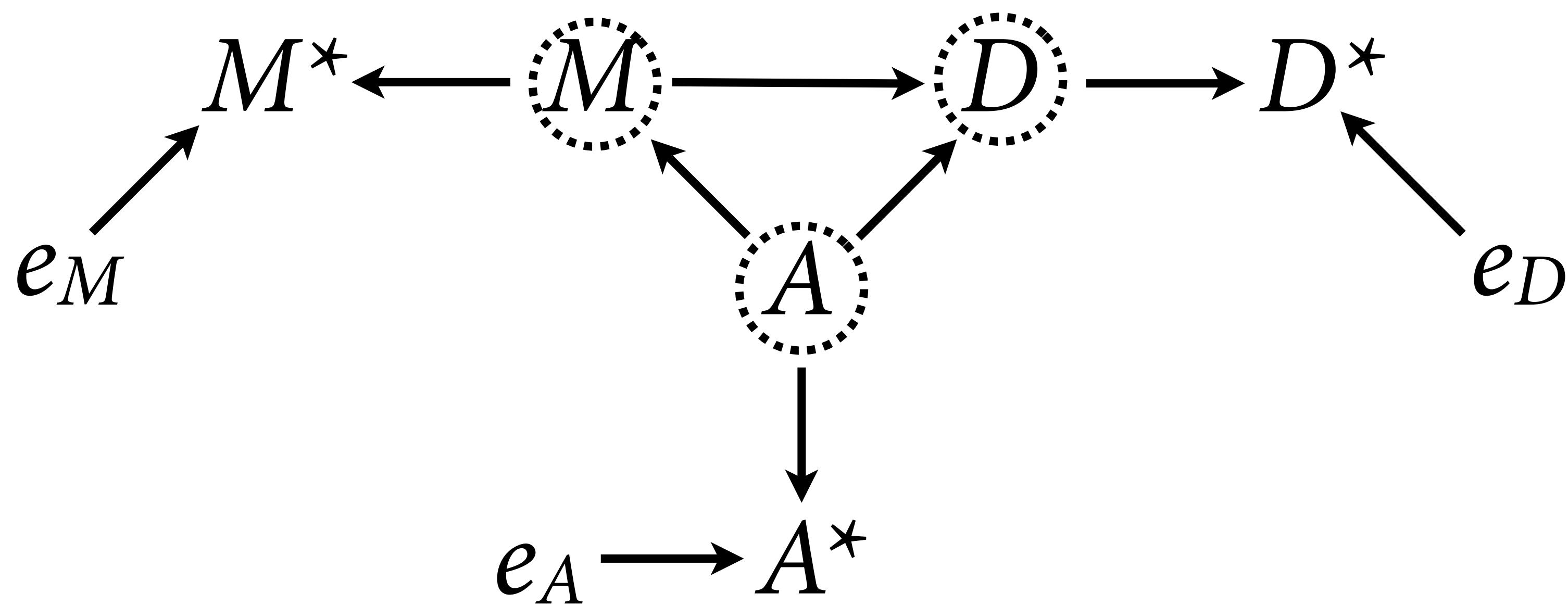


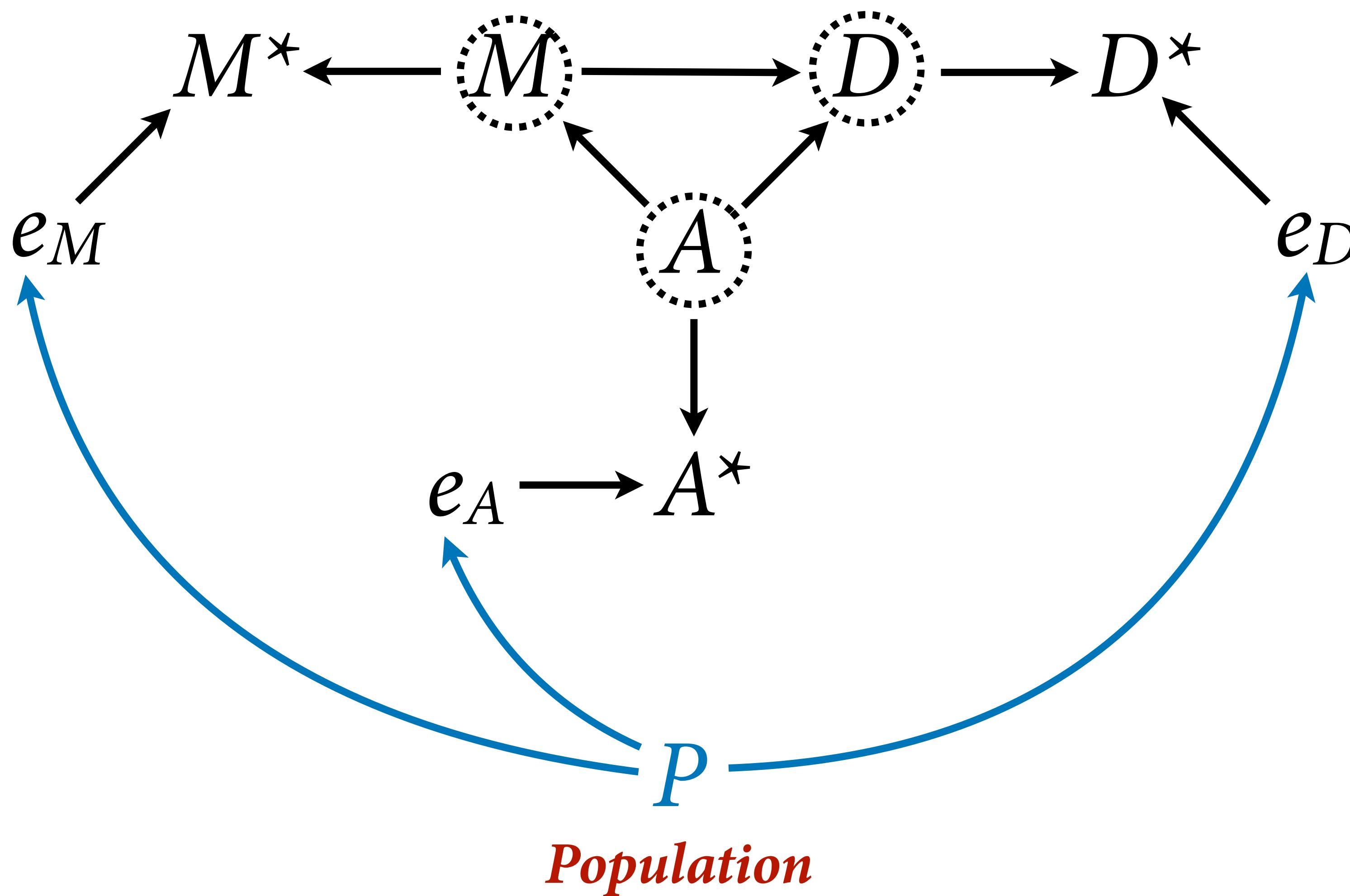
*Divorce
(unobserved)*

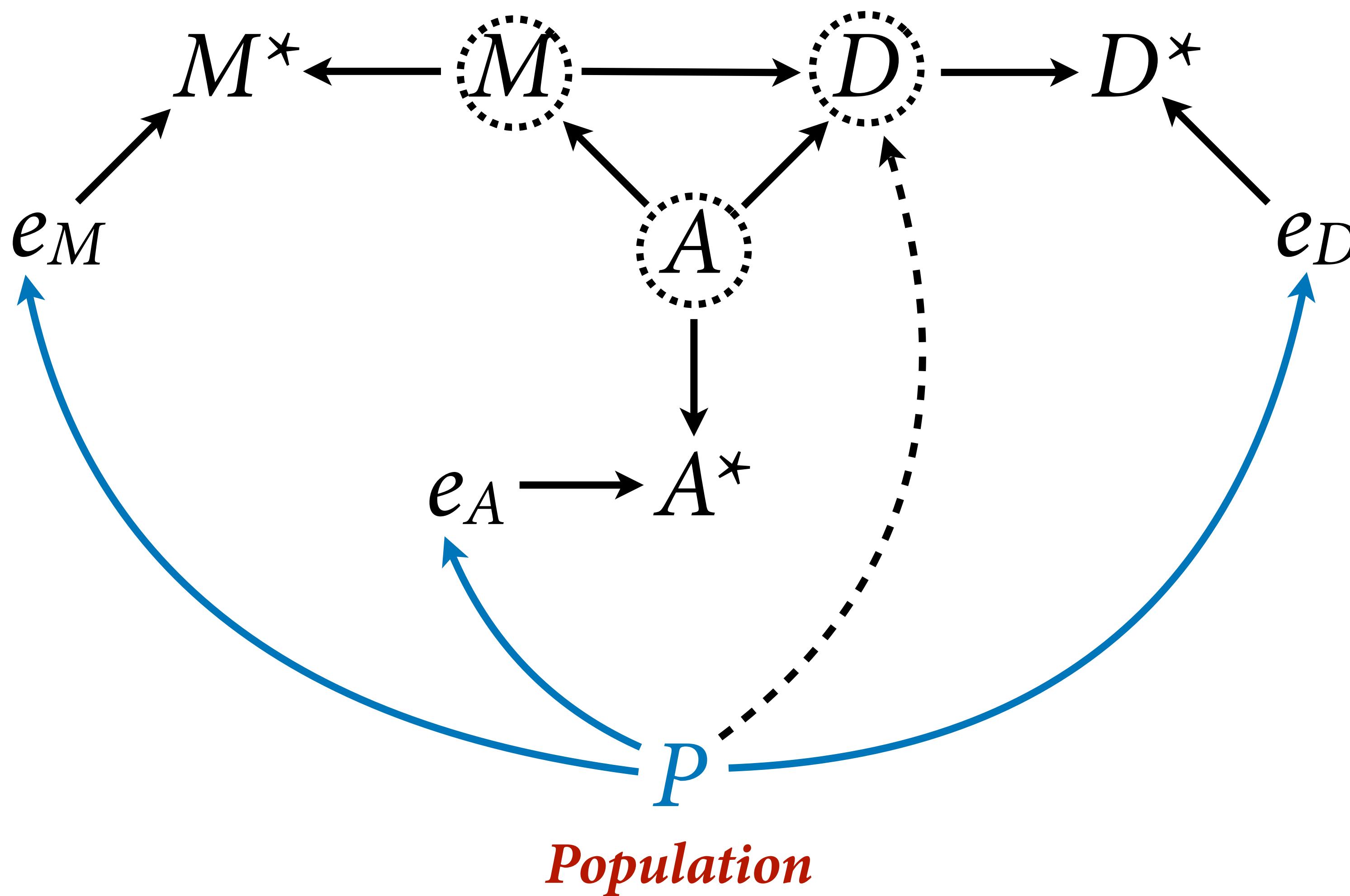


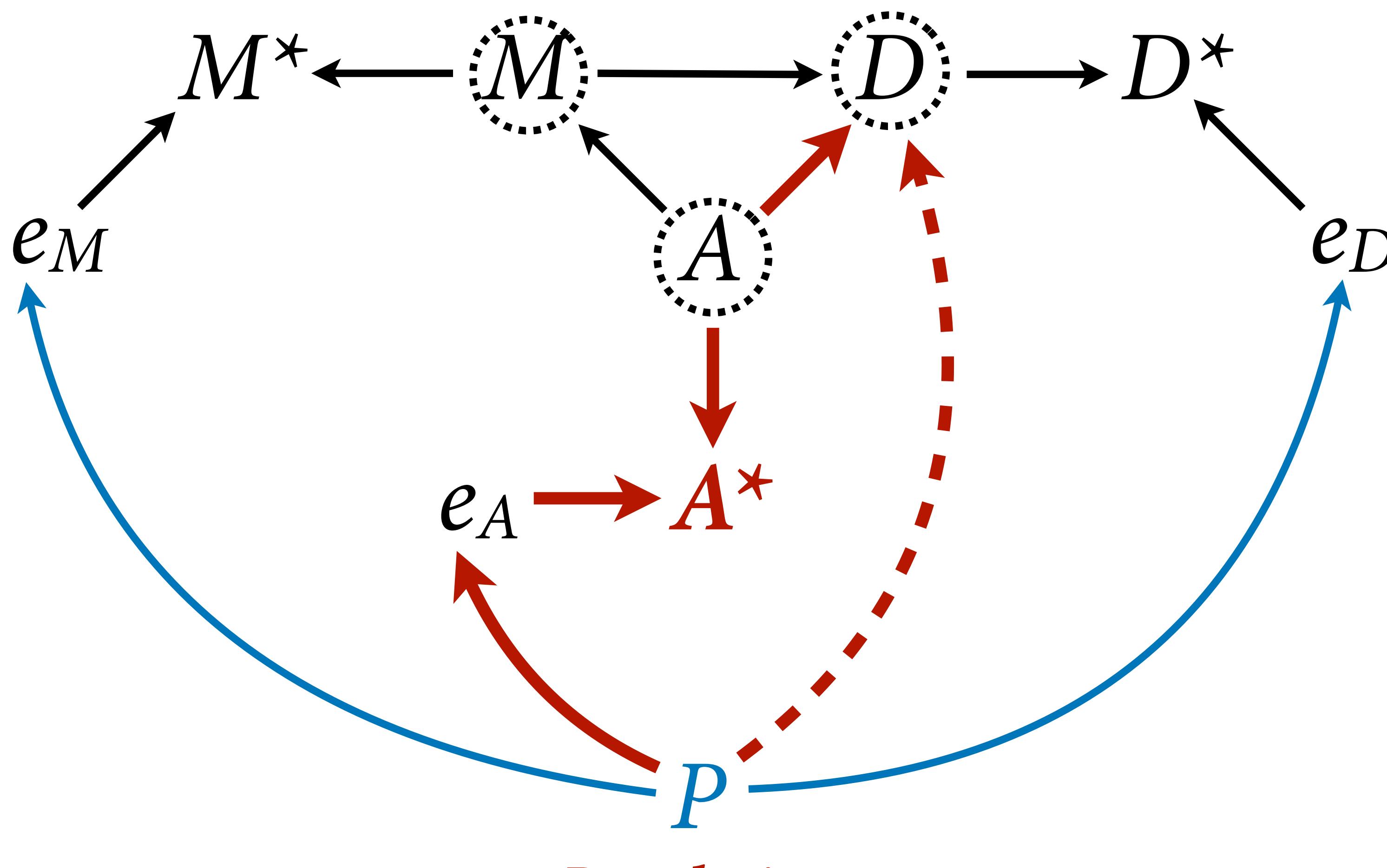












Thinking Like a Graph

Regressions (GLMMs) are special case machines

Thinking like a regression: Which predictor variables do I use?

Thinking like a graph: How do I model the network of causes?

this is
YOUR BRAIN

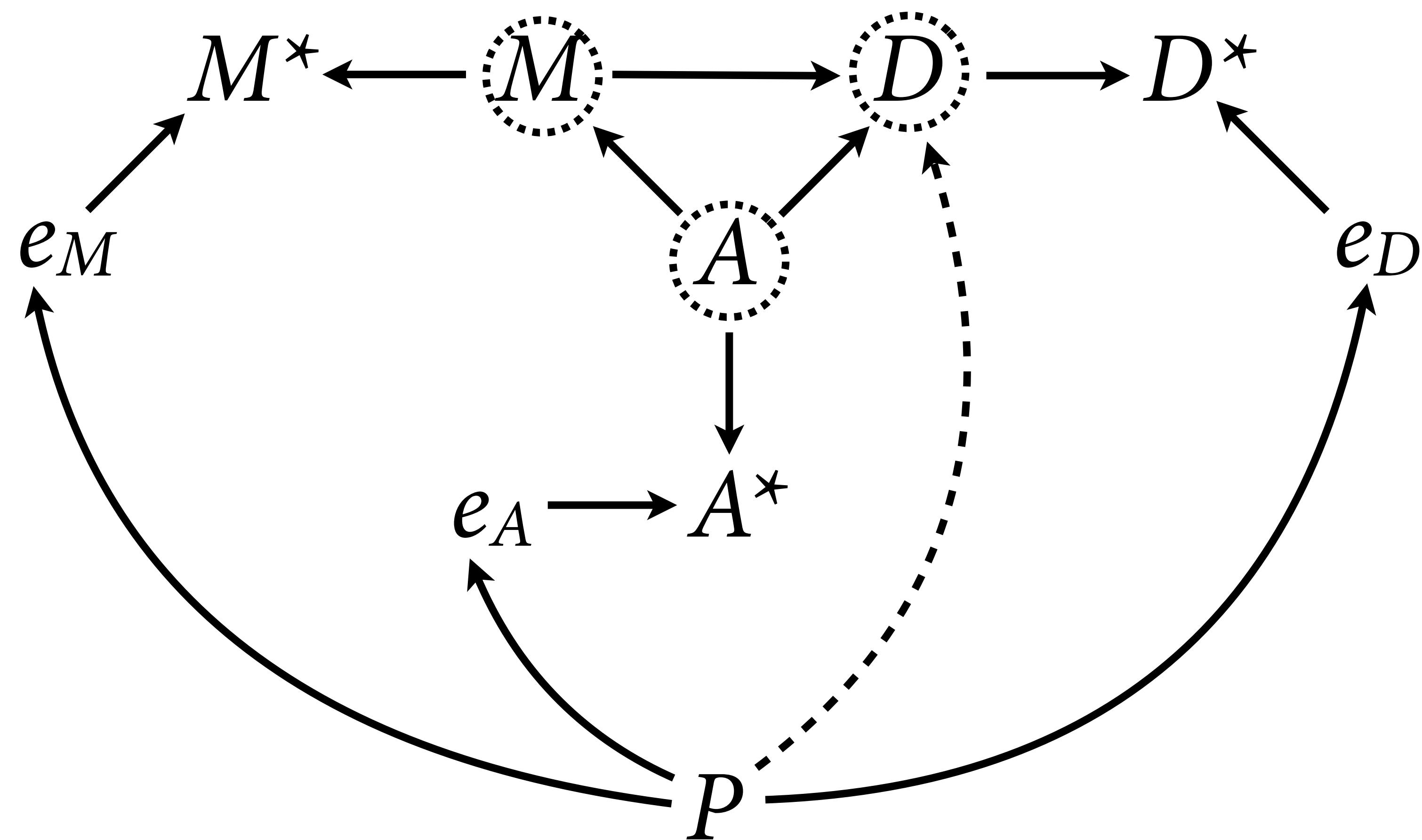
this is
REGRESSION

this is
YOUR BRAIN
on
REGRESSION

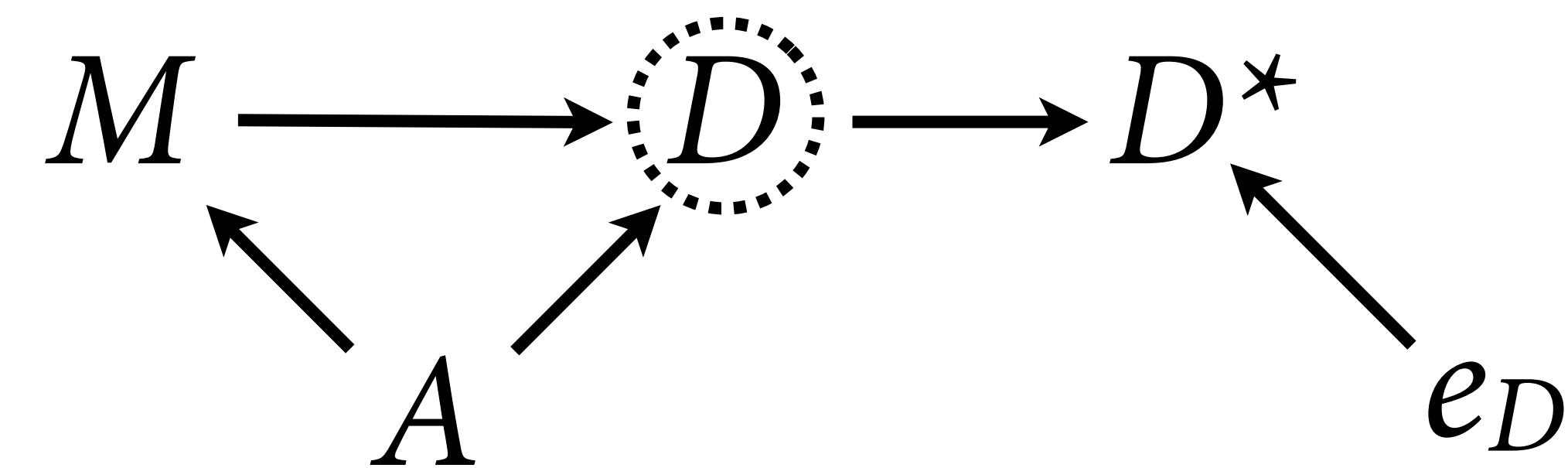
any questions?



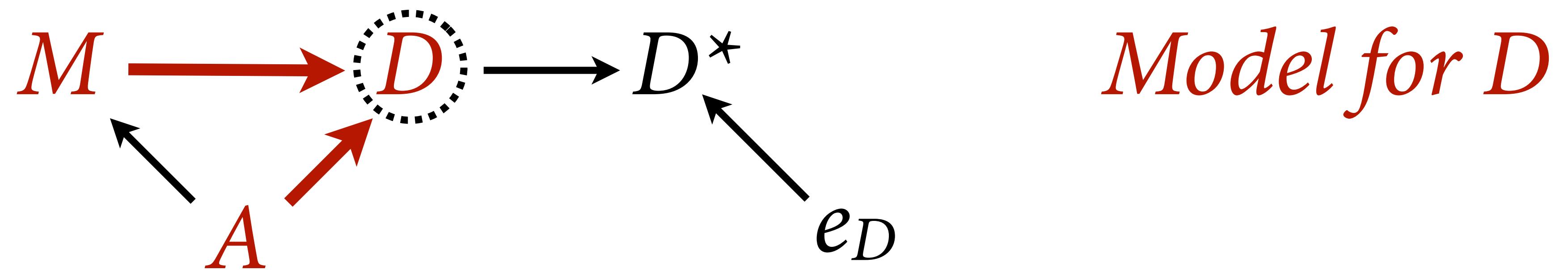
Thinking Like a Graph



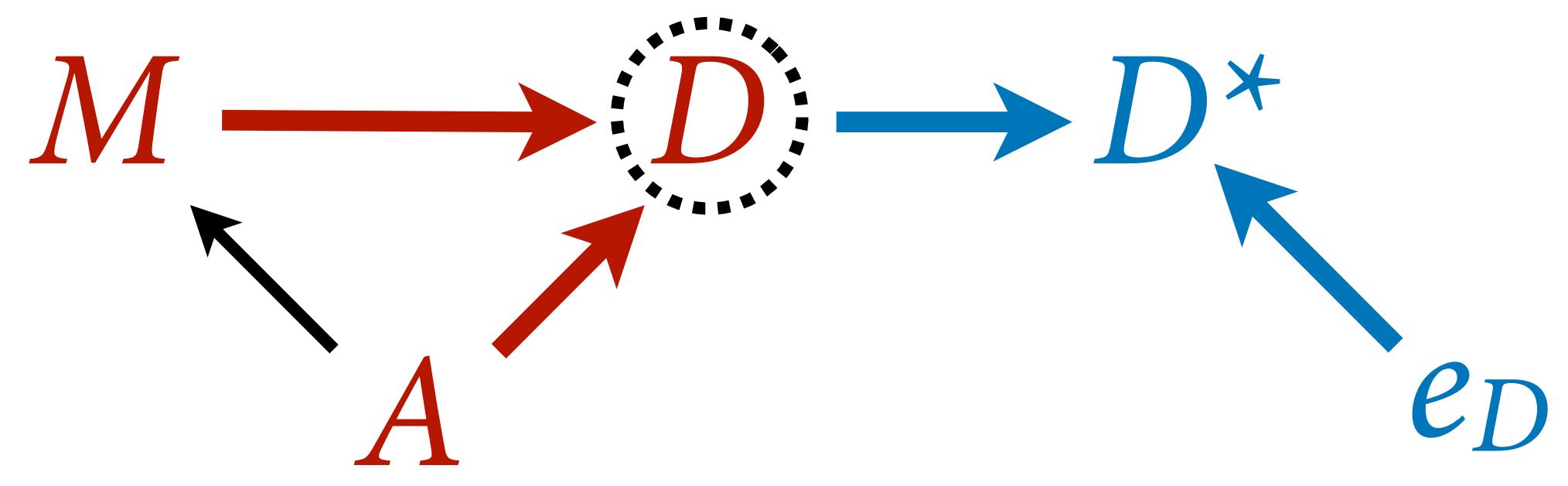
Thinking Like a Graph



Thinking Like a Graph



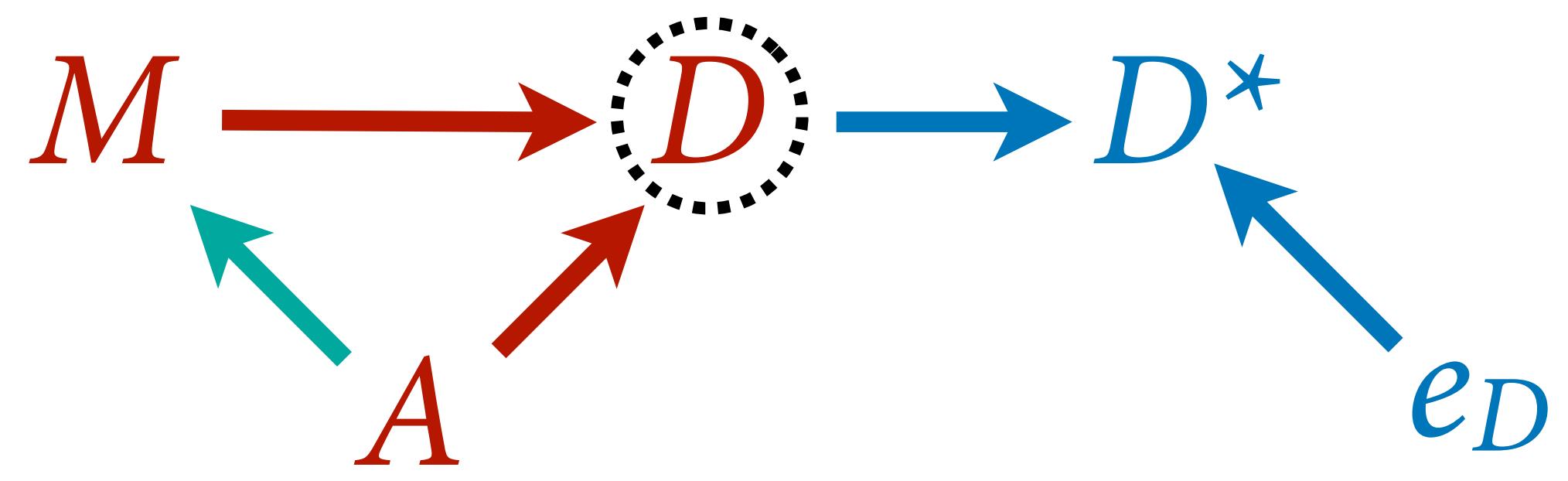
Thinking Like a Graph



Model for D

*Model for D^**

Thinking Like a Graph



Model for D

*Model for D**

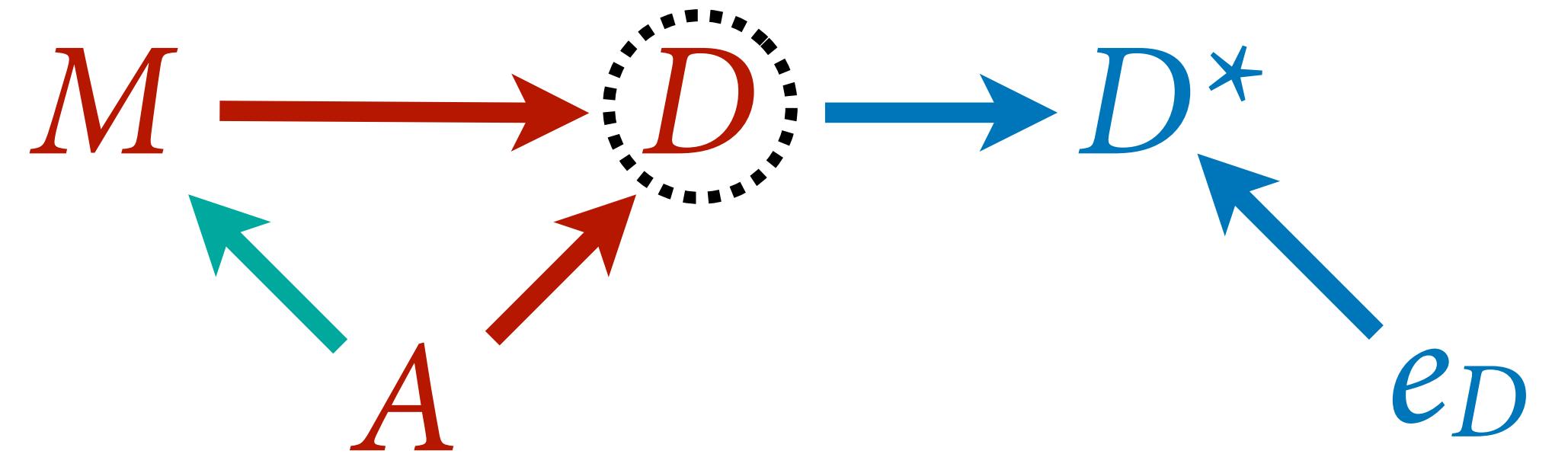
Model for M

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$D_i^\star = D_i + e_{D,i}$$

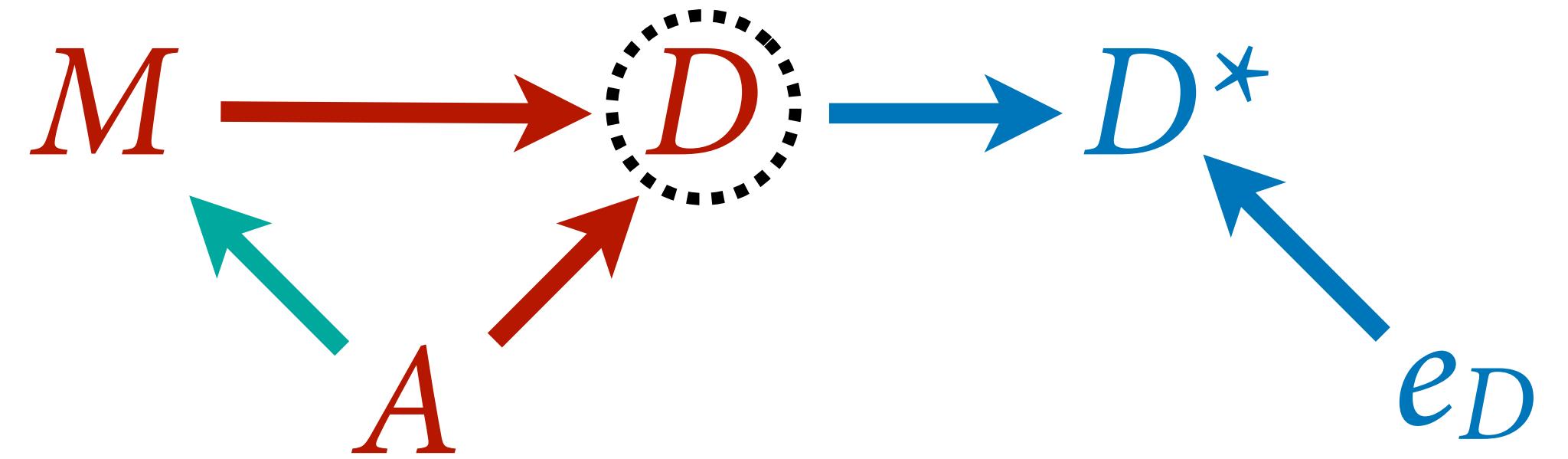
$$e_{D,i} \sim \text{Normal}(0, S_i)$$



$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$D_i^\star \sim \text{Normal}(D_i, S_i)$$

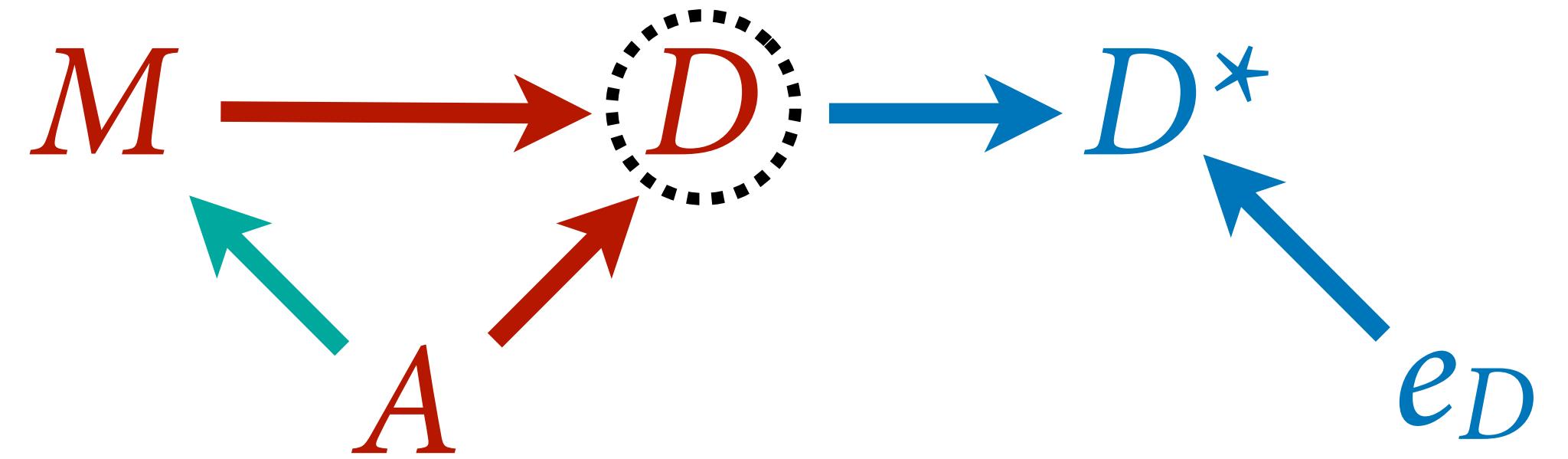


PAUSE

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$D_i^\star \sim \text{Normal}(D_i, S_i)$$



```

## R code 15.3
dlist <- list(
  D_obs = standardize( d$Divorce ),
  D_sd = d$Divorce.SE / sd( d$Divorce ),
  M = standardize( d$Marriage ),
  A = standardize( d$MedianAgeMarriage ),
  N = nrow(d)
)

m15.1 <- ulam(
  alist(
    # model for D* (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,

    # model for D (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp(1)
  ) , data=dlist , chains=4 , cores=4 )

precis( m15.1 , depth=2 )

```

$$D_i^{\star} \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

```

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  D_obs = standardize( d$Divorce ),
  D_sd = d$Divorce.SE / sd( d$Divorce ),
  M = standardize( d$Marriage ),
  A = standardize( d$MedianAgeMarriage ),
  N = nrow(d)
)

m15.1 <- ulam(
  alist(
    # model for D* (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,
    # model for D (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M,
    a ~ dnorm(0,0.2),
    bA ~ dnorm(0,0.5),
    bM ~ dnorm(0,0.5),
    sigma ~ dexp(1)
  ) , data=dlist , chains=4 , cores=4 )

precis( m15.1 , depth=2 )

```

$$D_i^* \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

```

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  M = standardize( d$Marriage ),
  A = standardize( d$MedianAgeMarriage ),
  N = nrow(d)
)

m15.1 <- ulam(
  alist(
    # model for D* (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,

    # model for D (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp(1)
  ) , data=dlist , chains=4 , cores=4 )

precis( m15.1 , depth=2 )

```



$$D_i^* \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

```

## R code 15.3
dlist <- list > precis( m15.1 , depth=2 )
  D_obs = s      mean   sd  5.5% 94.5% n_eff Rhat4
  D_sd = d$    1.17 0.37  0.59  1.78  2219    1
  M = stand    0.69 0.57 -0.23  1.59  2895    1
  A = stand    0.42 0.34 -0.13  0.97  2582    1
  N = nrow(    1.42 0.48  0.66  2.21  1938    1
)                                D_true[5] -0.90 0.13 -1.10 -0.70  2855    1
m15.1 <- ulam
  alist(
    # mod
    D_obs
    # mod
    vector
    mu ~
    a ~ d
    bA ~
    bM ~
    sigma
  ) , data=
precis( m15.1

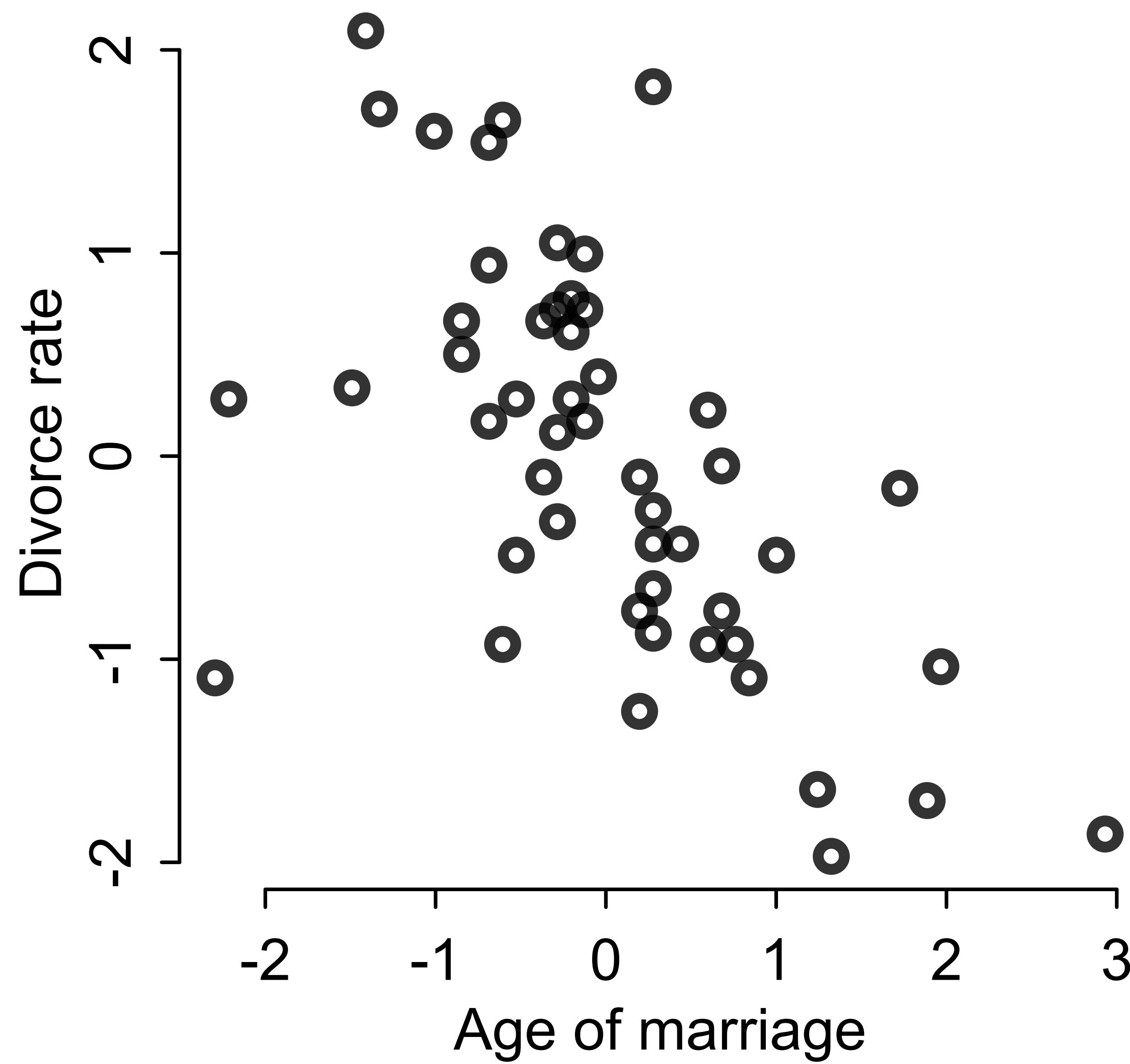
```

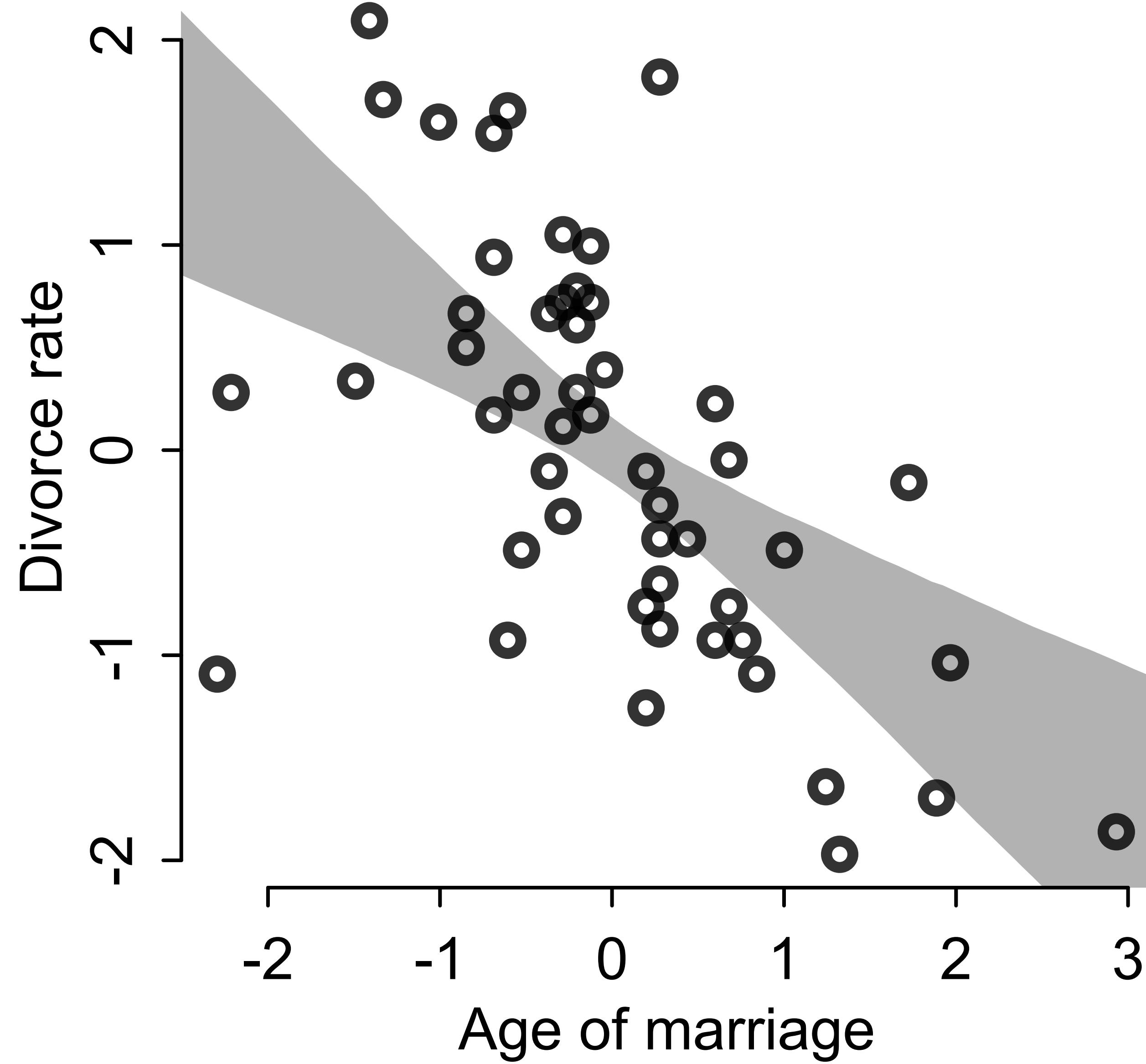


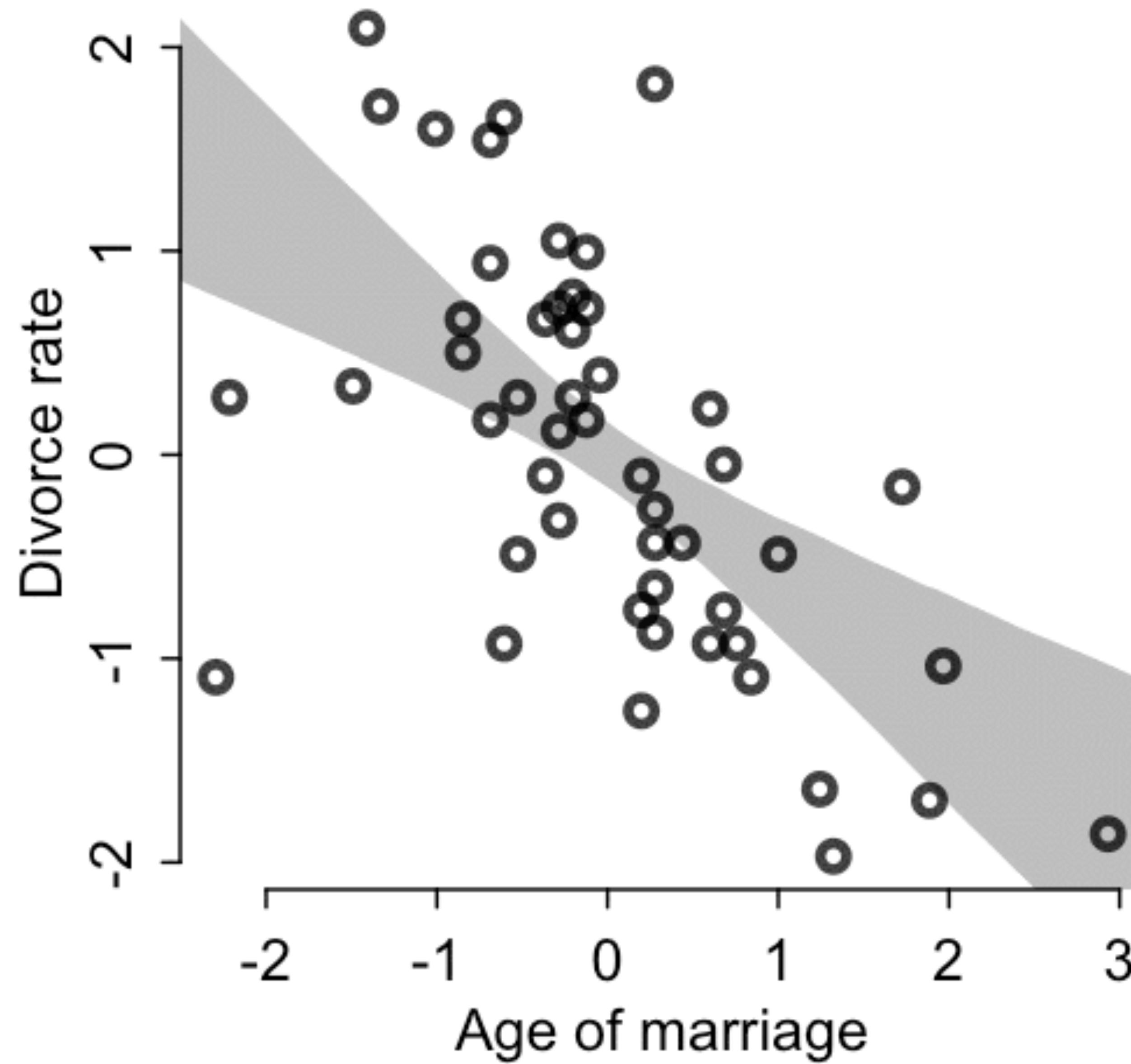
$$D_i^* \sim \text{Normal}(D_i, S_i)$$

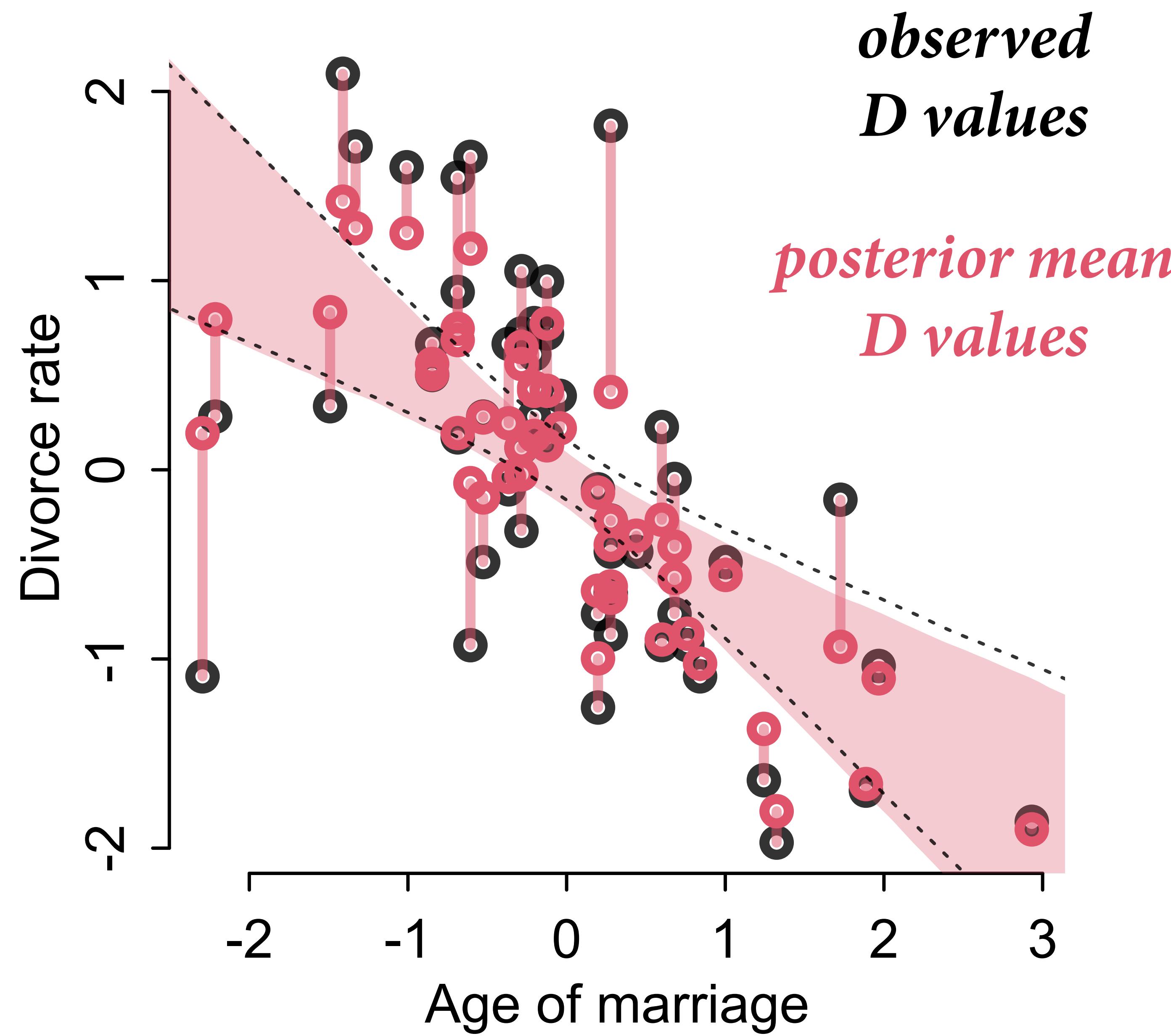
$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

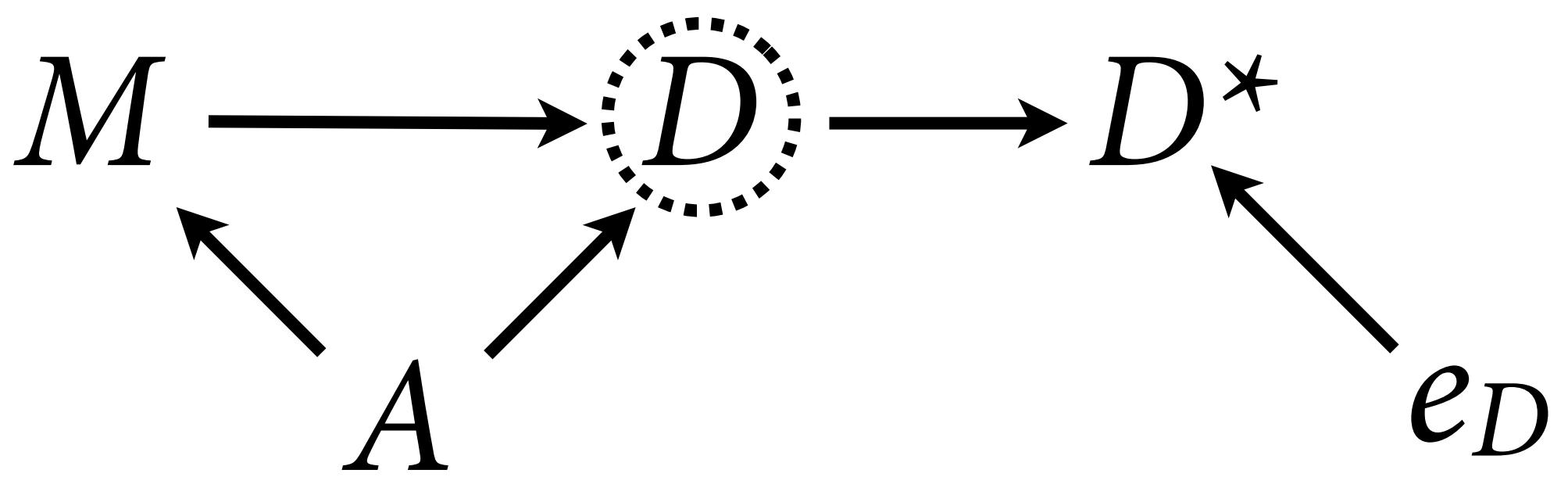
$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

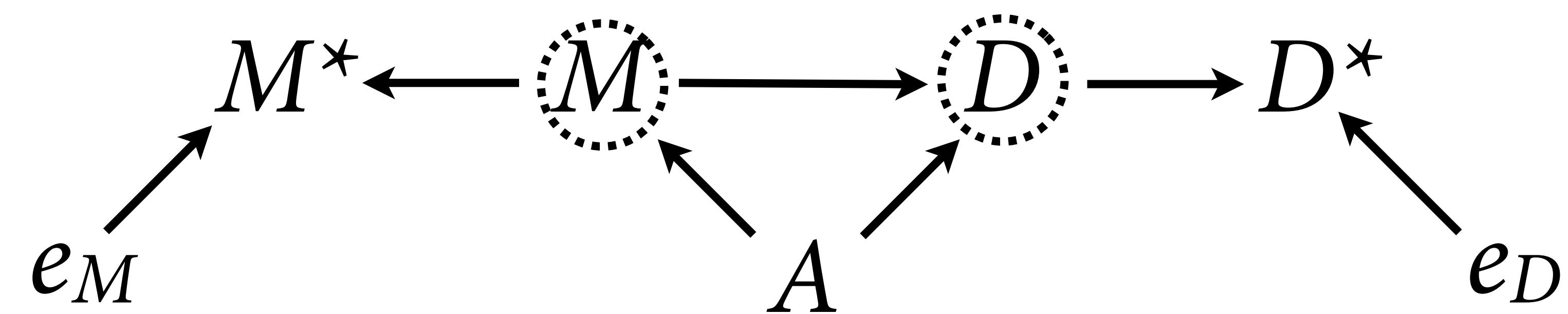






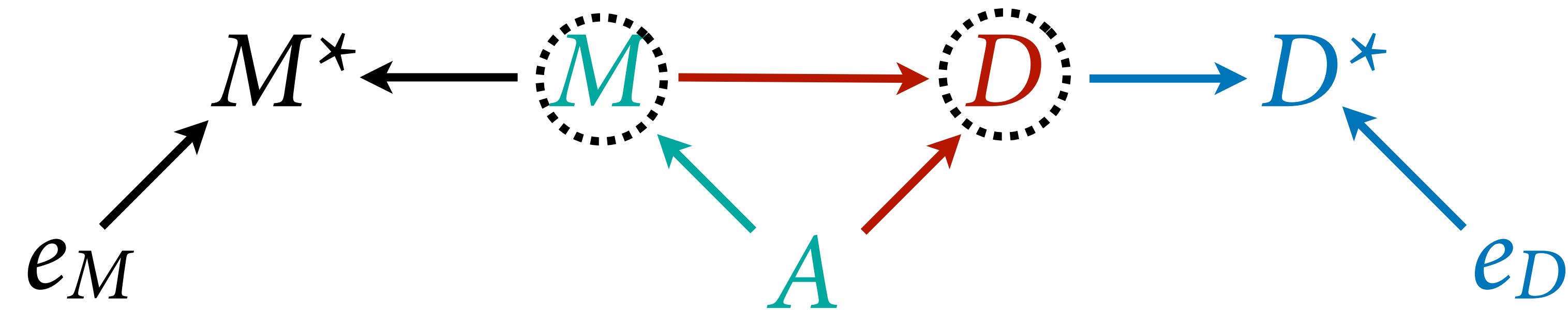






$$D_i \sim \text{Normal}(\mu_i, \sigma)$$
$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

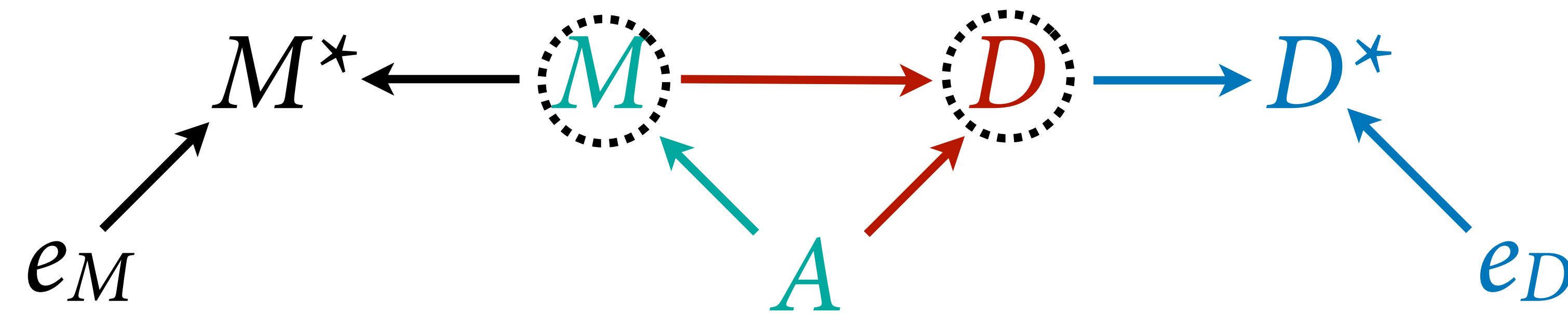
$$D_i^* \sim \text{Normal}(D_i, S_i)$$



$$D_i \sim \text{Normal}(\mu_i, \sigma)$$
$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$M_i^* \sim \text{Normal}(M_i, T_i)$$

$$D_i^* \sim \text{Normal}(D_i, S_i)$$



$$M_i \sim \text{Normal}(\nu_i, \tau)$$
$$\nu_i = \alpha_M + \beta_{AM} A_i$$

```

m15.2 <- ulam(
  alist(
    # D* model (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,

    # D model (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M_true[i] ,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp( 1 ) ,

    # M* model (observed)
    M_obs ~ dnorm( M_true , M_sd ) ,

    # M model (unobserved)
    vector[N]:M_true ~ dnorm( nu , tau ) ,
    nu <- aM + bAM*A ,
    aM ~ dnorm(0,0.2) ,
    bAM ~ dnorm(0,0.5) ,
    tau ~ dexp( 1 )

  ) , data=dlist2 , chains=4 , cores=4 )

```

$$D_i^{\star} \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$M_i^{\star} \sim \text{Normal}(M_i, T_i)$$

$$M_i \sim \text{Normal}(\nu_i, \tau)$$

$$\nu_i = \alpha_M + \beta_{AM} A_i$$

```

m15.2 <- ulam(
  alist(
    # D* model (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,
    # D model (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M_true[i] ,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp( 1 ) ,
    # M* model (observed)
    M_obs ~ dnorm( M_true , M_sd ) ,
    # M model (unobserved)
    vector[N]:M_true ~ dnorm( nu , tau ) ,
    nu <- aM + bAM*A ,
    aM ~ dnorm(0,0.2) ,
    bAM ~ dnorm(0,0.5) ,
    tau ~ dexp( 1 )
  ) , data=dlist2 , chains=4 , cores=4 )

```

$$D_i^* \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$M_i^* \sim \text{Normal}(M_i, T_i)$$

$$M_i \sim \text{Normal}(\nu_i, \tau)$$

$$\nu_i = \alpha_M + \beta_{AM} A_i$$

```

m15.2 <- ulam(
  alist(
    # D* model (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,

    # D model (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M_true[i] ,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp( 1 ) ,

    # M* model (observed)
    M_obs ~ dnorm( M_true , M_sd ) ,

    # M model (unobserved)
    vector[N]:M_true ~ dnorm( nu , tau ) ,
    nu <- aM + bAM*A ,
    aM ~ dnorm(0,0.2) ,
    bAM ~ dnorm(0,0.5) ,
    tau ~ dexp( 1 )

  ) , data=dlist2 , chains=4 , cores=4 )

```

$$D_i^* \sim \text{Normal}(D_i, S_i)$$

$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$M_i^* \sim \text{Normal}(M_i, T_i)$$

$$M_i \sim \text{Normal}(\nu_i, \tau)$$

$$\nu_i = \alpha_M + \beta_{AM} A_i$$

```

m15.2 <- ulam(
  alist(
    # D* model (observed)
    D_obs ~ dnorm( D_true , D_sd ) ,

    # D model (unobserved)
    vector[N]:D_true ~ dnorm( mu , sigma ) ,
    mu <- a + bA*A + bM*M_true[i] ,
    a ~ dnorm(0,0.2) ,
    bA ~ dnorm(0,0.5) ,
    bM ~ dnorm(0,0.5) ,
    sigma ~ dexp( 1 ) ,

    # M* model (observed)
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    vector[N]:M_true ~ dnorm( nu , tau ) ,
    nu <- aM + bAM*A ,
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    bAM ~ dnorm(0,0.5) ,
    tau ~ dexp( 1 )

  ) , data=dlist2 , chains=4 , cores=4 )

```

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    nu <- aM + bAM*A ,
    aM ~ dnorm(0,0.2) ,
    bAM ~ dnorm(0,0.5) ,
    tau ~ dexp( 1 )

  ) , data=dlist2 , chains=4 , cores=4 )

```

$$D_i^* \sim \text{Normal}(D_i, S_i)$$

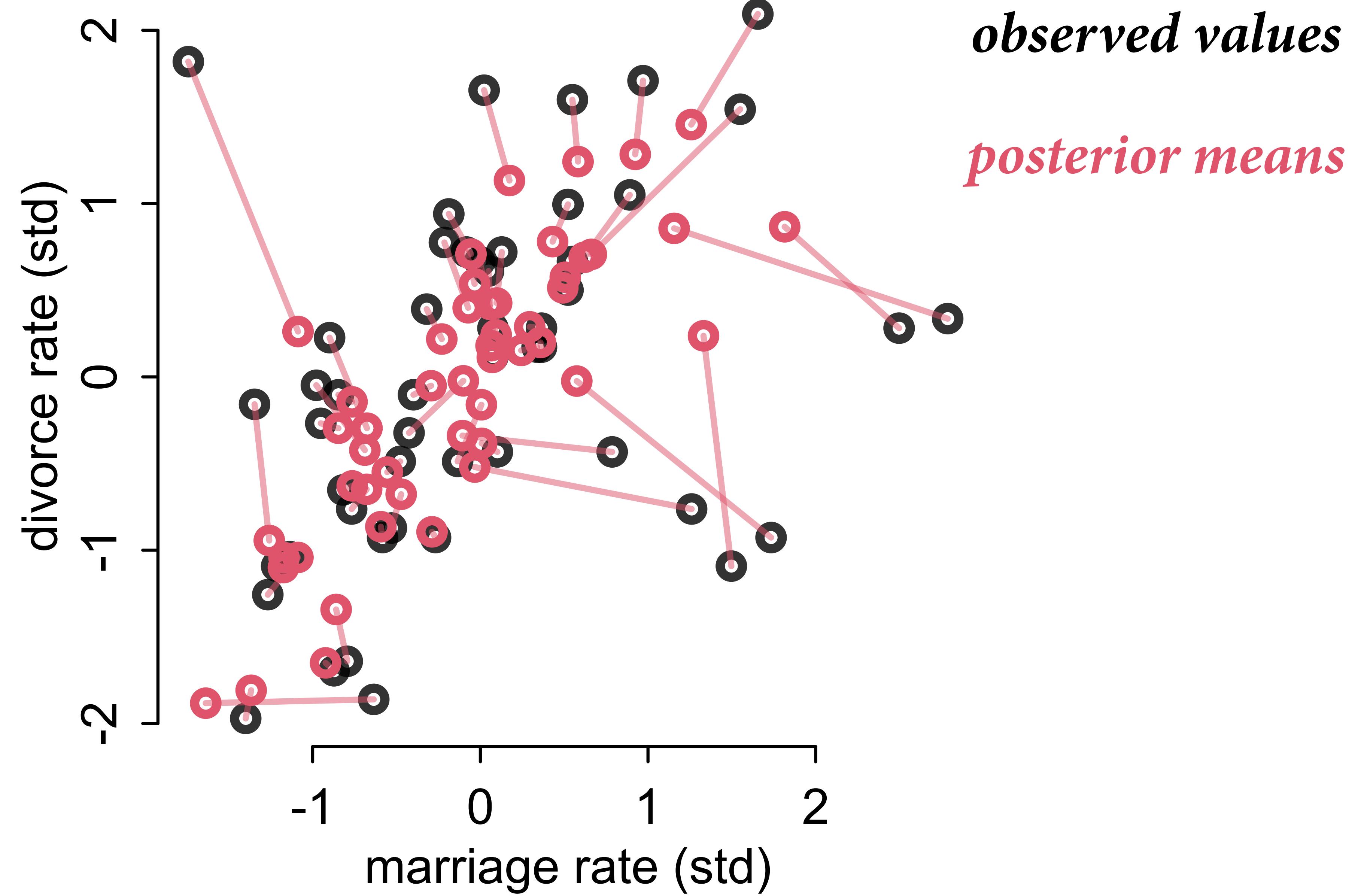
$$D_i \sim \text{Normal}(\mu_i, \sigma)$$

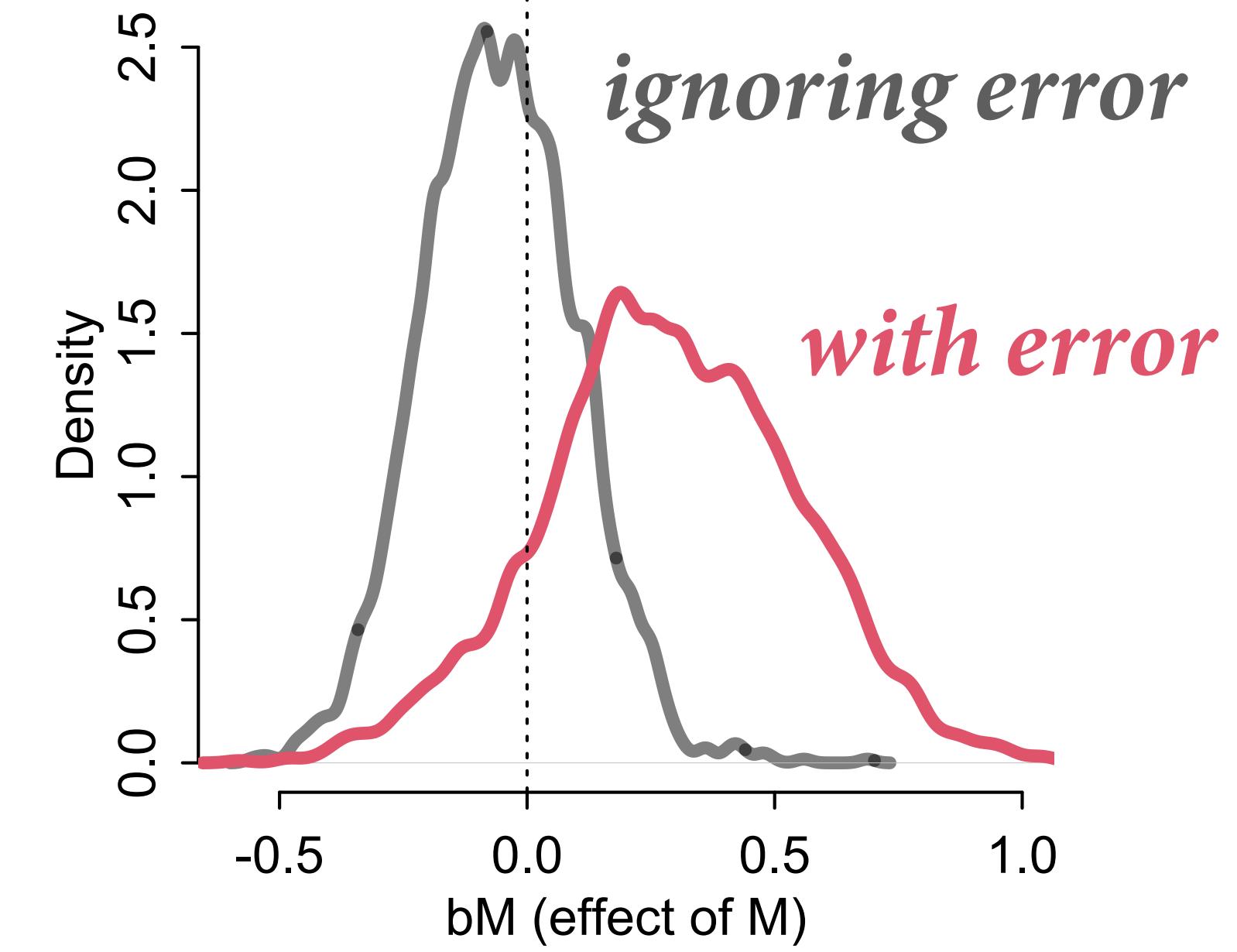
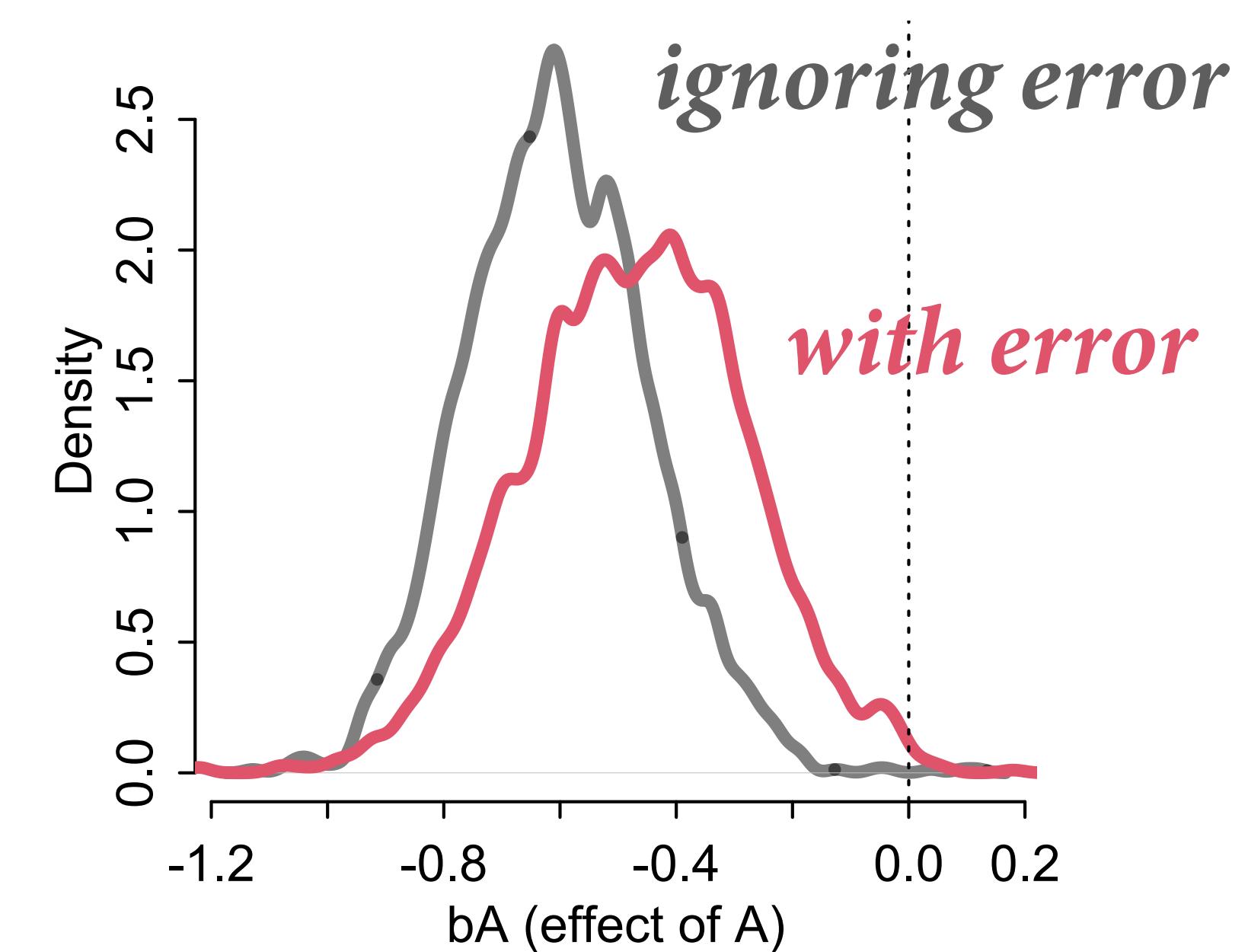
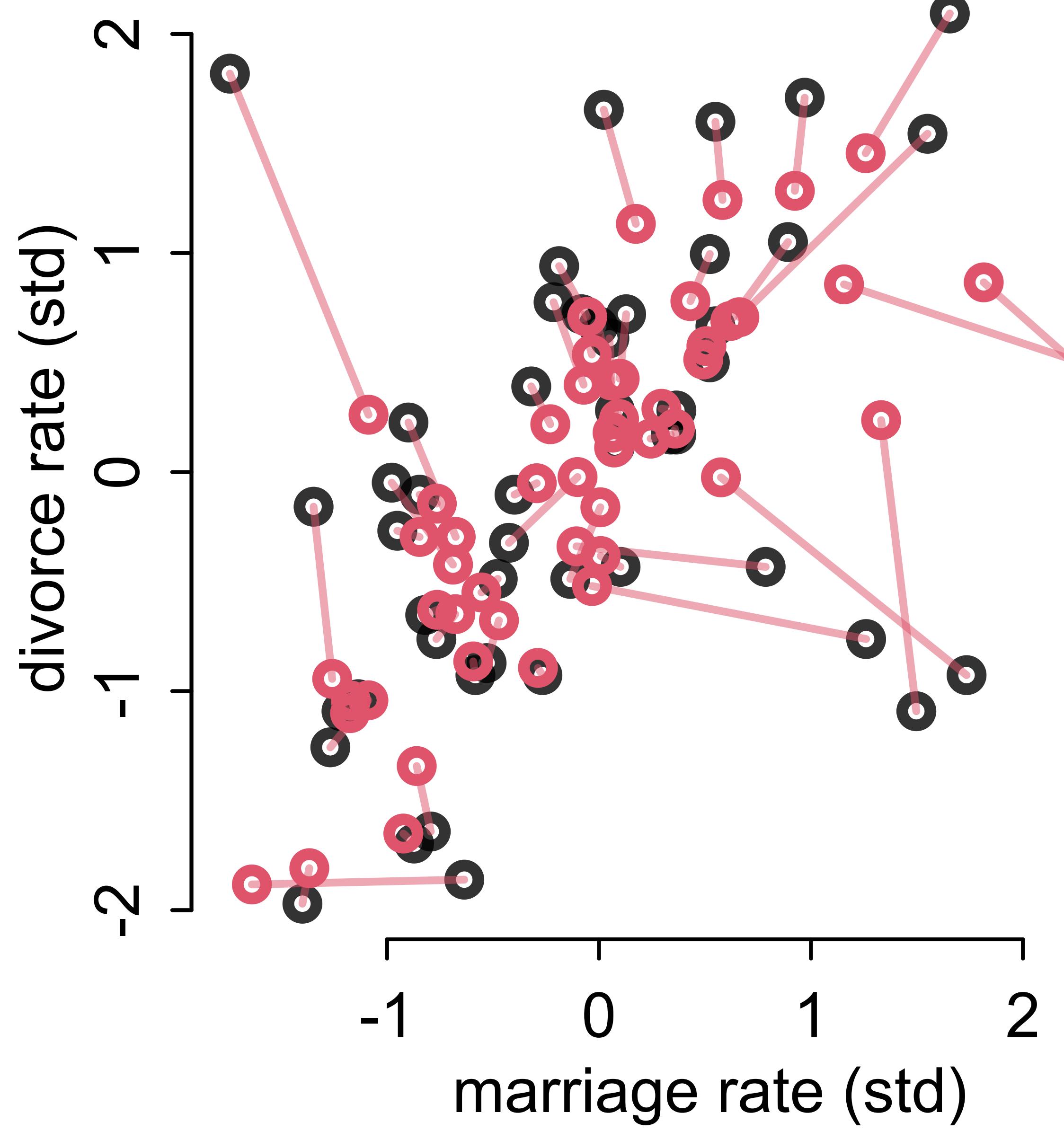
$$\mu_i = \alpha + \beta_A A_i + \beta_M M_i$$

$$M_i^* \sim \text{Normal}(M_i, T_i)$$

$$M_i \sim \text{Normal}(\nu_i, \tau)$$

$$\nu_i = \alpha_M + \beta_{AM} A_i$$





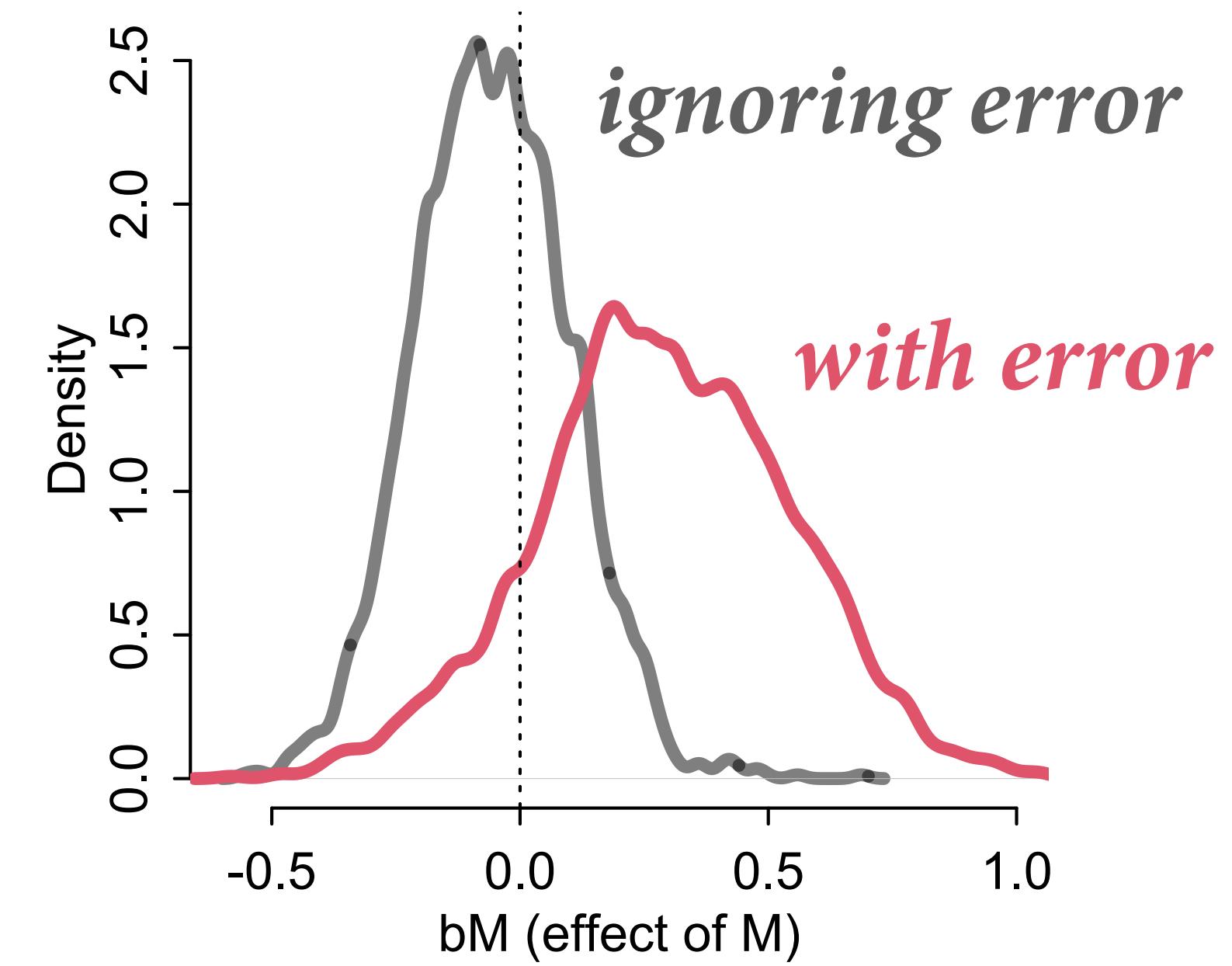
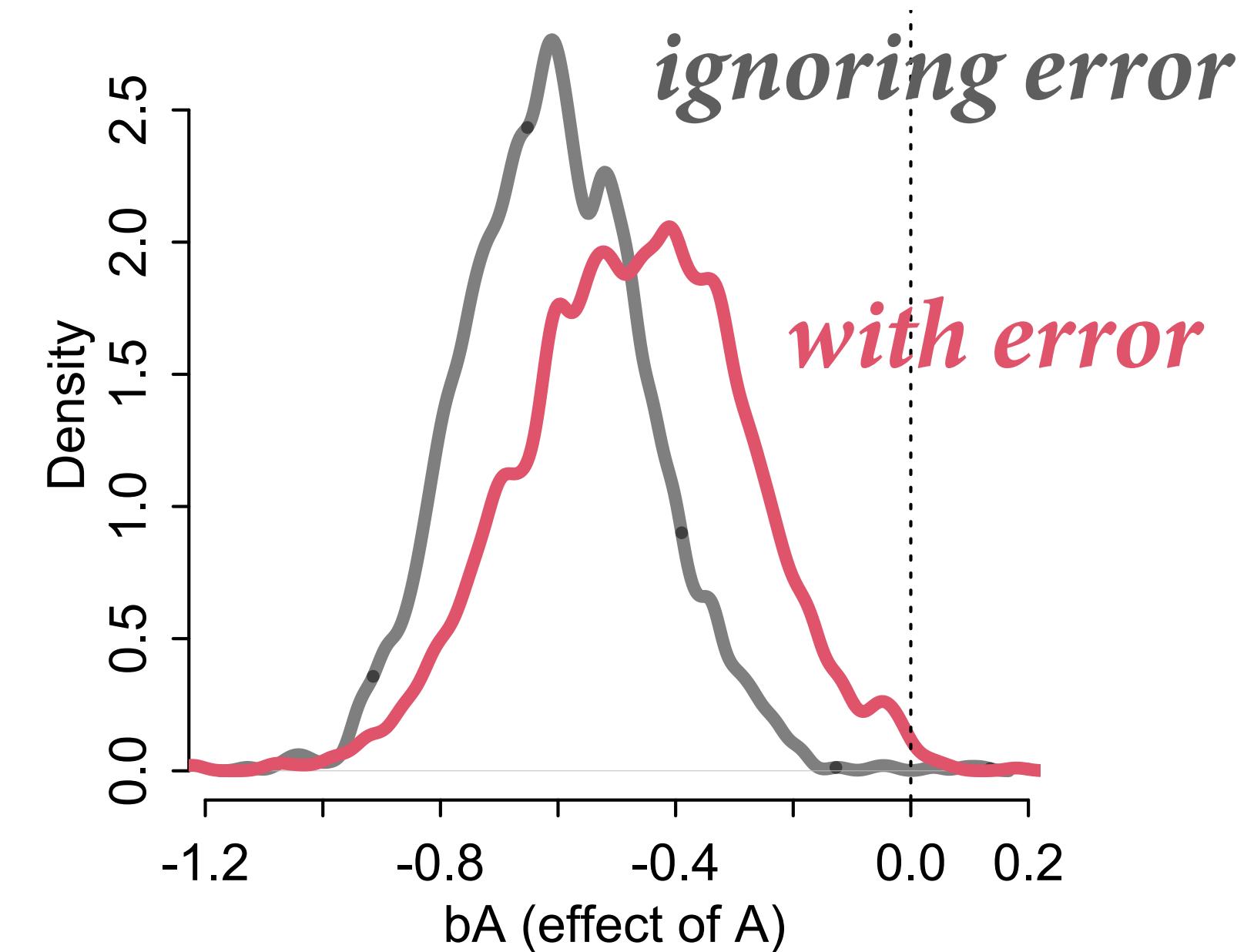
Unpredictable errors

Including error on M increases evidence for an effect of M on D

Why?

Down-weighting of unreliable estimates

Errors can hurt or help, but only honest option is to attend to them



PAUSE

Bantu pastoralists

50k people, 50k km²

Bilineal descent

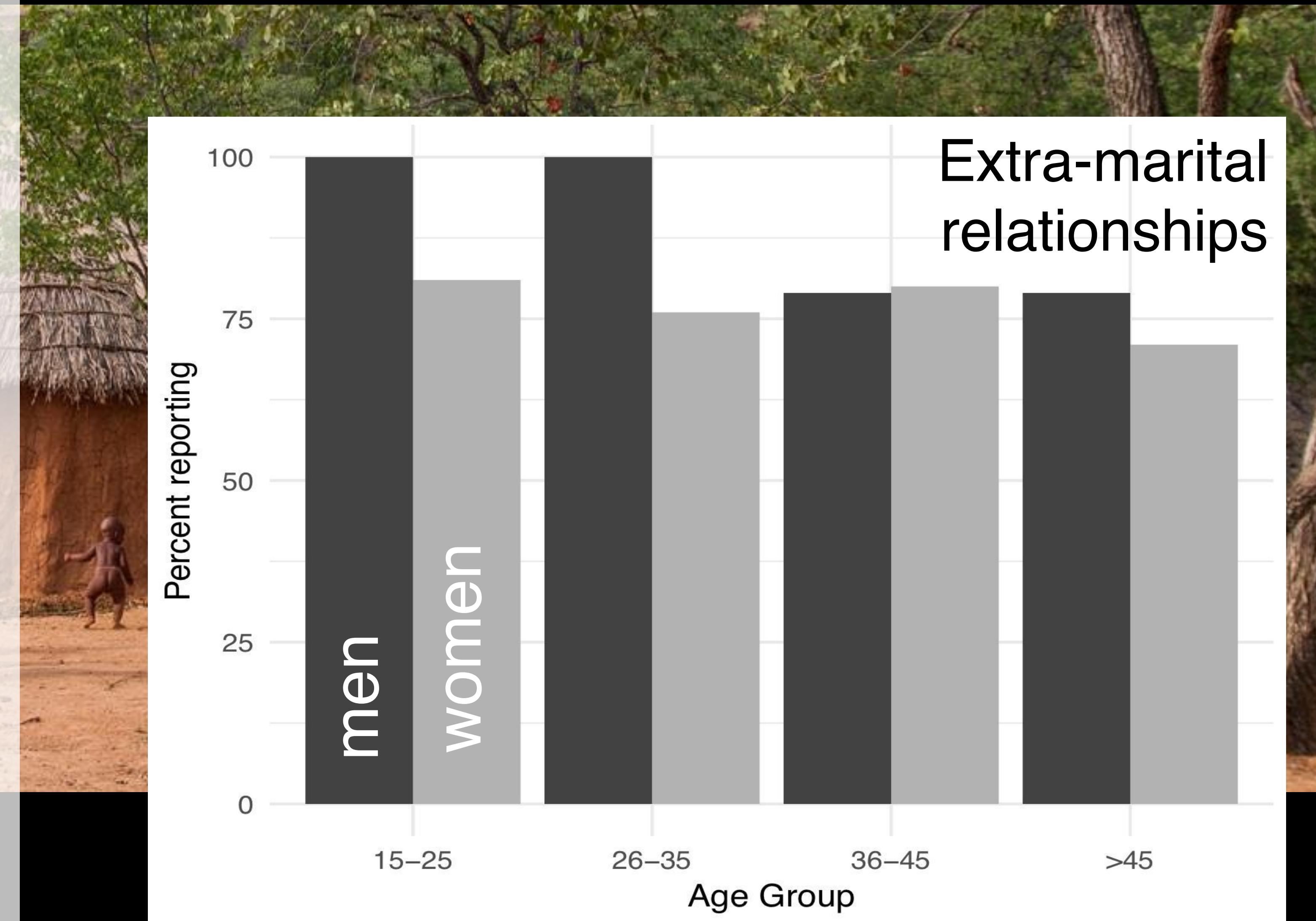
Matrilineal inheritance

Patrilineal status

Patrilocality

Polygyny common

“Open” marriages



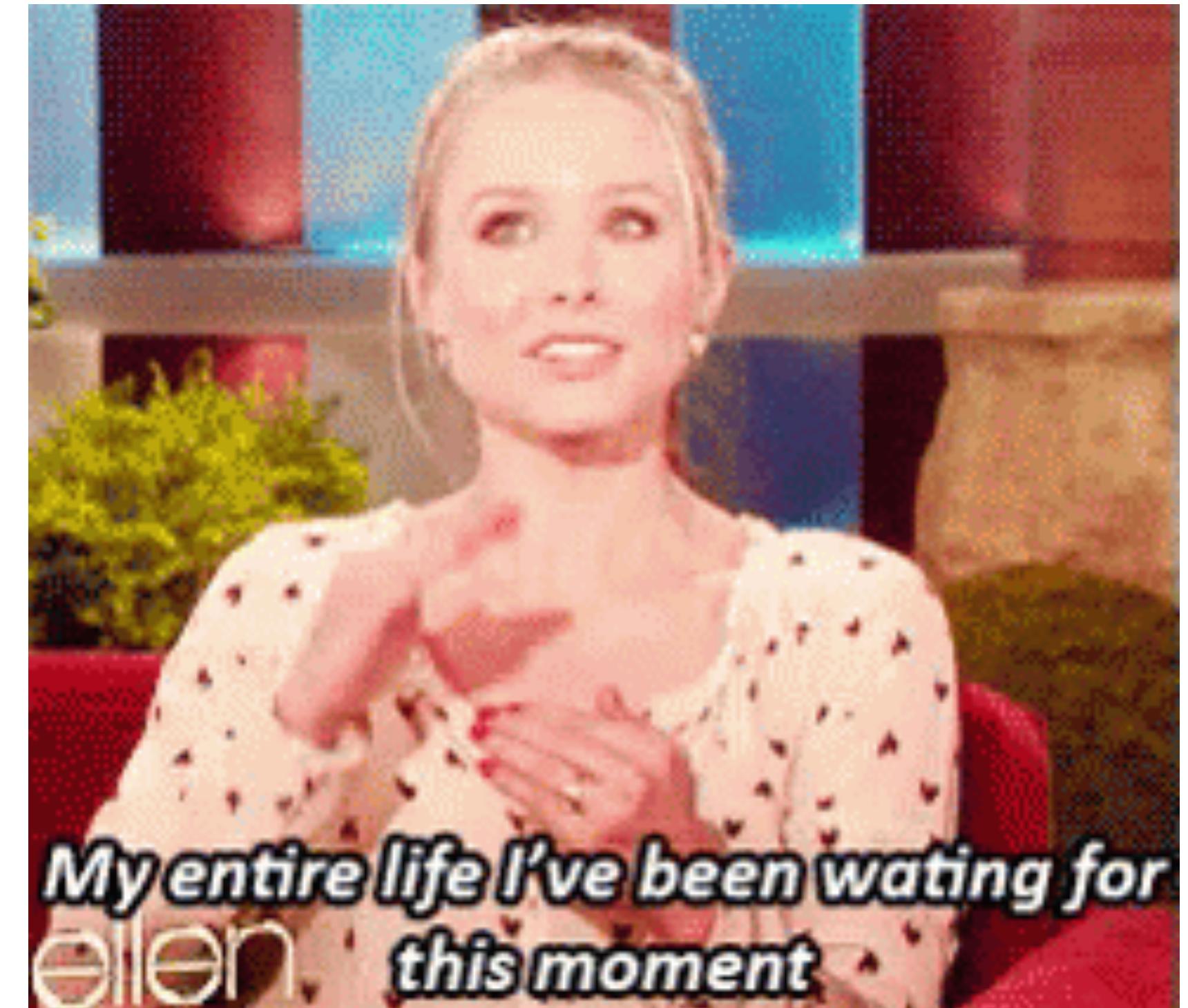
Misclassification

Estimand: Proportion of children fathered by extra-pair men

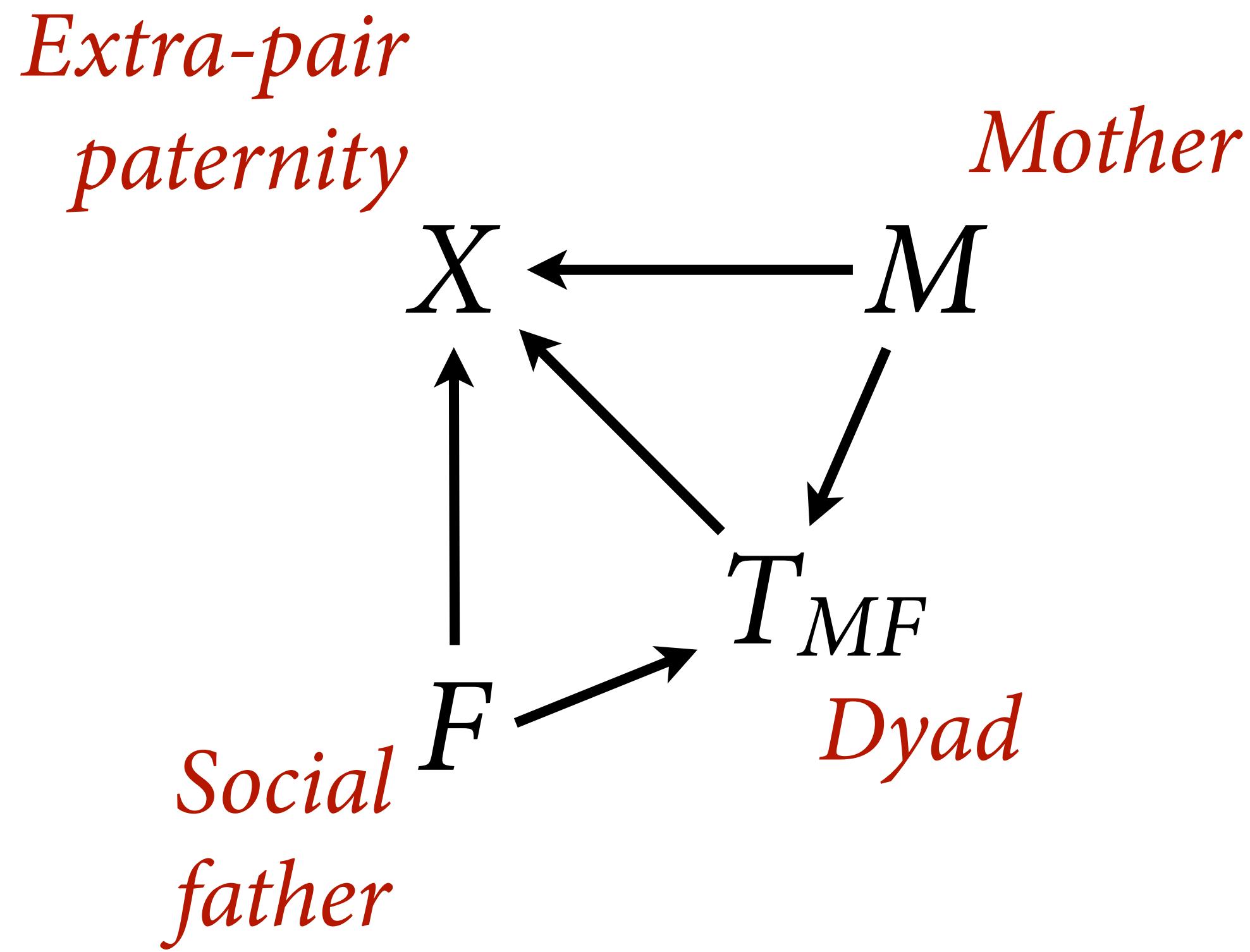
Problem: 5% false-positive rate

Misclassification: Categorical version of measurement error

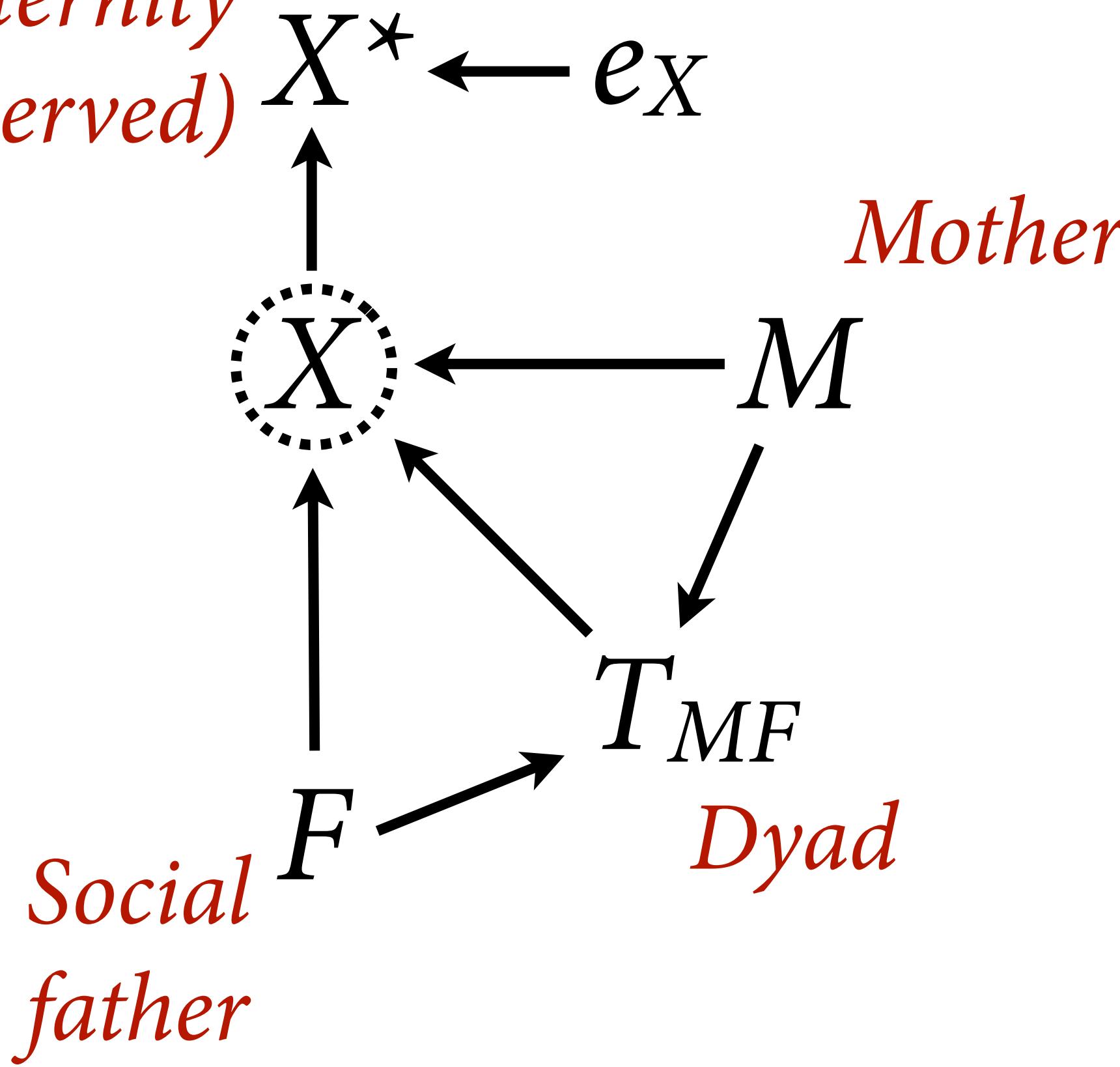
How do we include misclassification?



When my colleagues ask me for help with misclassification



*Extra-pair
paternity
(observed)*



*Social
father*

Mother

Dyad

*Generative
model*

$$X_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha + \mu_{M[i]} + \delta_{D[i]}$$

Mom Dyad

*Generative
model*

$$X_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha + \mu_{M[i]} + \delta_{D[i]}$$

Mom *Dyad*

*Measurement
model*

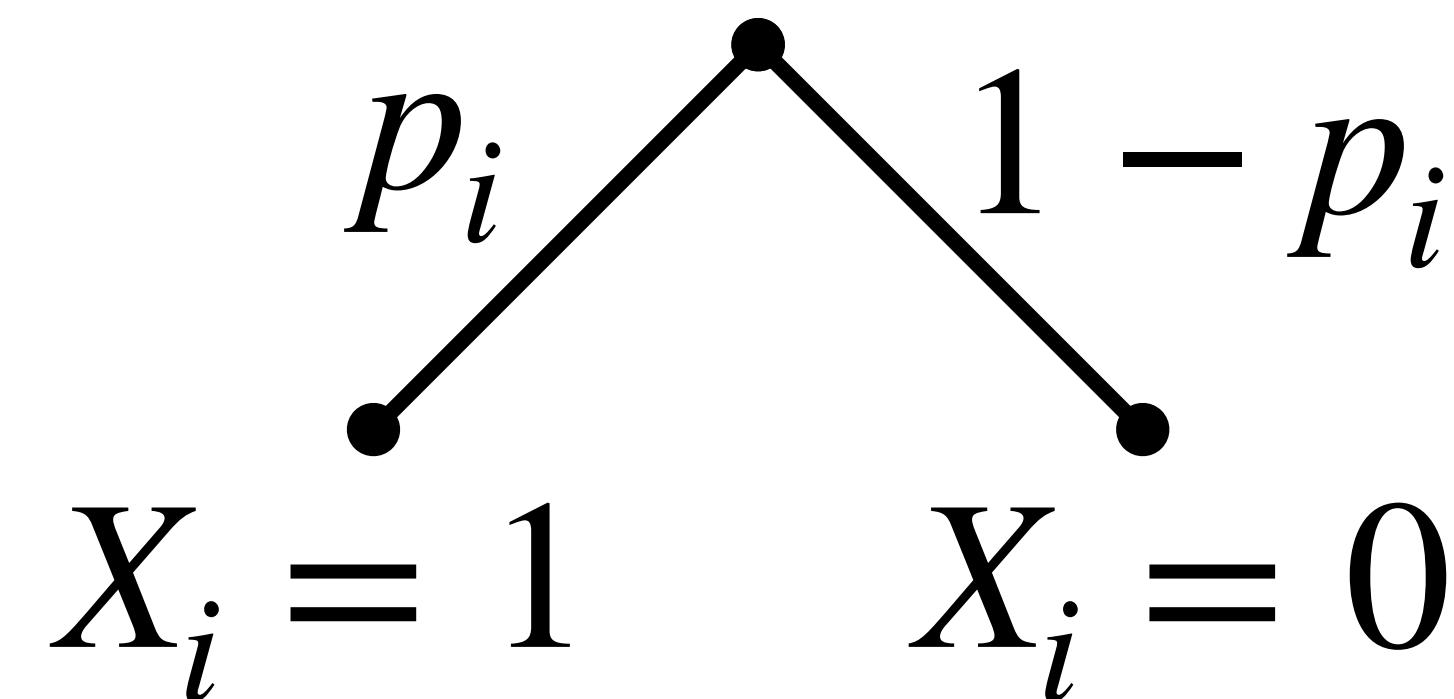
$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

*Measurement
model*

$$\Pr(X_i^{\star} = 1 | p_i) = p_i + (1 - p_i)f$$

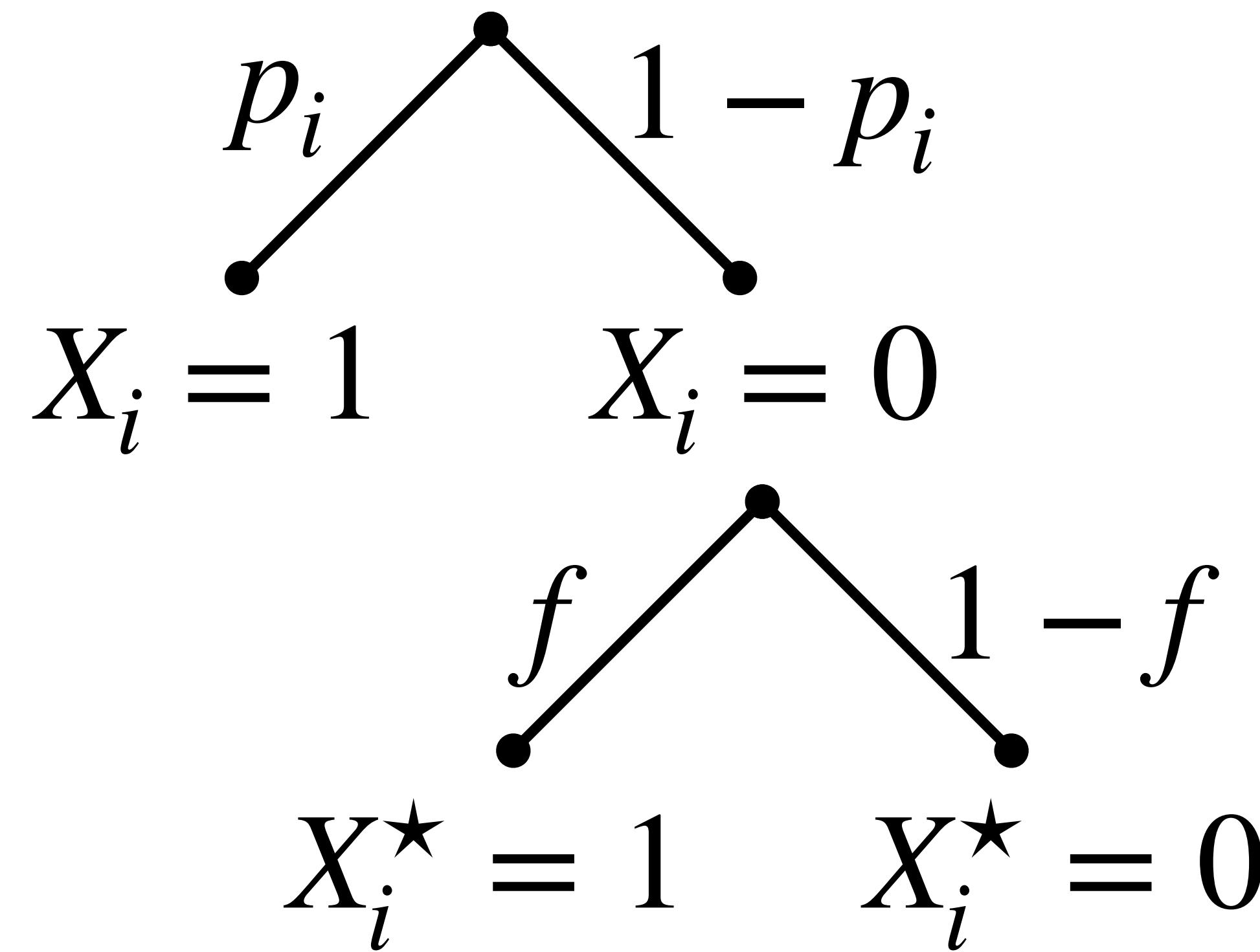
$$\Pr(X_i^{\star} = 0 | p_i) = (1 - p_i)(1 - f)$$



*Measurement
model*

$$\Pr(X_i^{\star} = 1 | p_i) = p_i + (1 - p_i)f$$

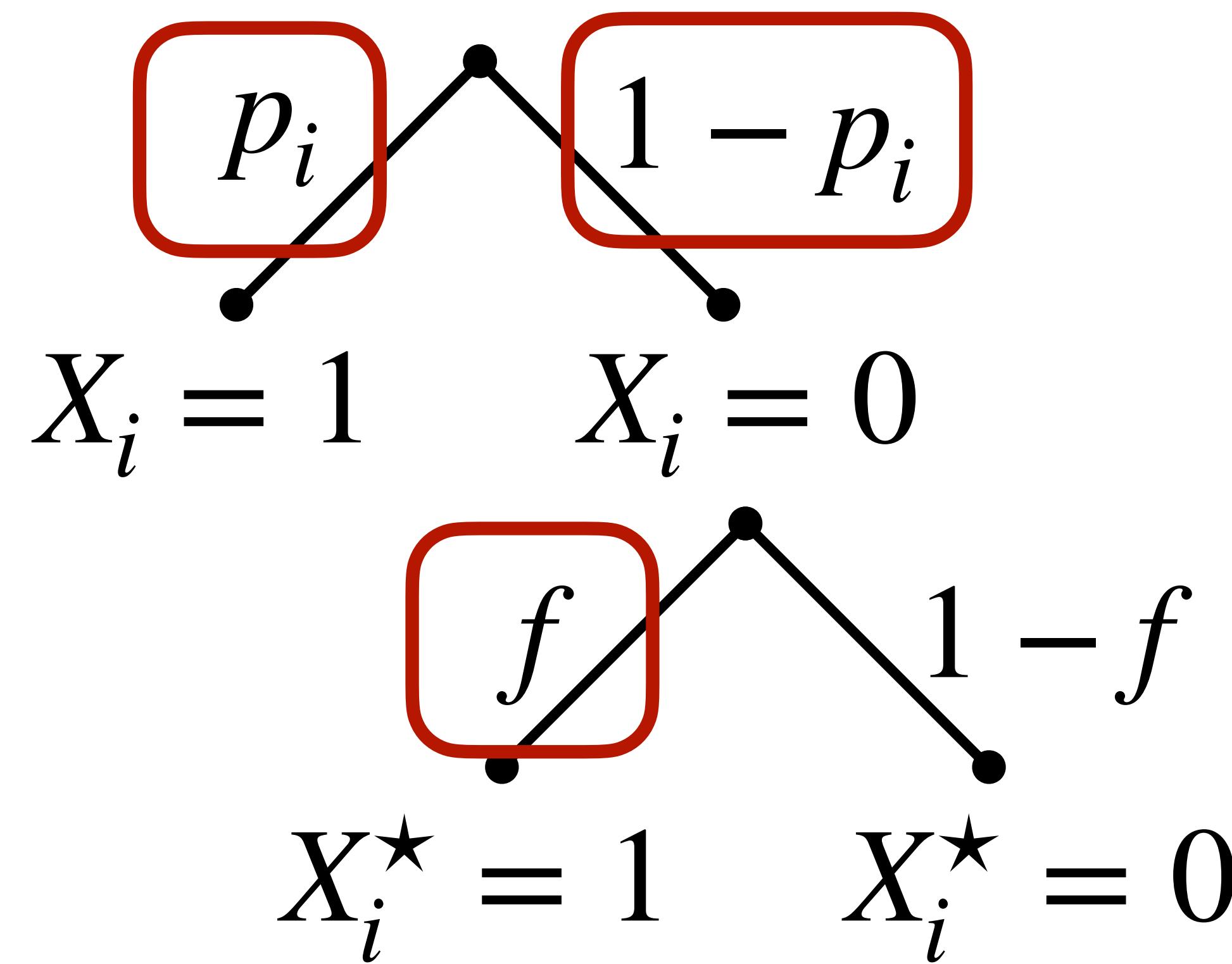
$$\Pr(X_i^{\star} = 0 | p_i) = (1 - p_i)(1 - f)$$



*Measurement
model*

$$\Pr(X_i^* = 1 | p_i) \neq p_i + (1 - p_i)f$$

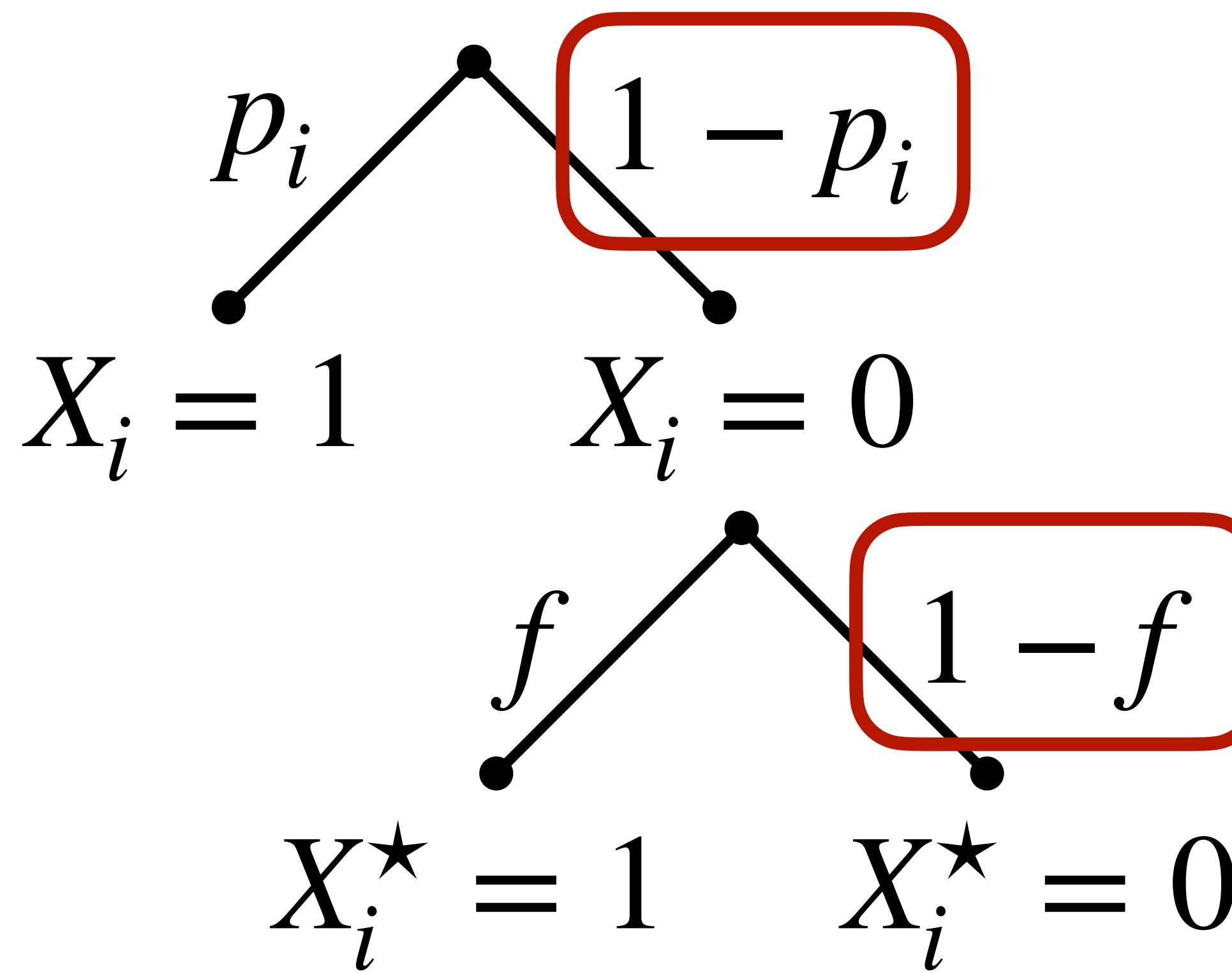
$$\Pr(X_i^* = 0 | p_i) = (1 - p_i)(1 - f)$$



*Measurement
model*

$$\Pr(X_i^* = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^* = 0 | p_i) = \boxed{(1 - p_i)(1 - f)}$$



```

# with false positive rate
dat$f <- 0.05
mX <- ulam(
  alist(
    X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    logit(p) <- a + z[mom_id]*sigma + x[dyad_id]*tau,
    a ~ normal(0,1.5),
    z[mom_id] ~ normal(0,1),
    sigma ~ normal(0,1),
    x[dyad_id] ~ normal(0,1),
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )

```

$$X_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha + \mu_{M[i]} + \delta_{D[i]}$$

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```

# with false positive rate
dat$f <- 0.05
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  alist(
    X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    logit(p) ~ a + z[mom_id]*sigma + x[dyad_id]*tau,
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    z[mom_id] ~ normal(0,1),
    sigma ~ normal(0,1),
    x[dyad_id] ~ normal(0,1),
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$$X_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha + \mu_{M[i]} + \delta_{D[i]}$$

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

Floating Point Monsters

Probability calculations tend to underflow (round to zero) and overflow (round to one)

Solution: Calculate on log scale

Ancient weapons:

`log_sum_exp`, `log1m`, `log1m_exp`



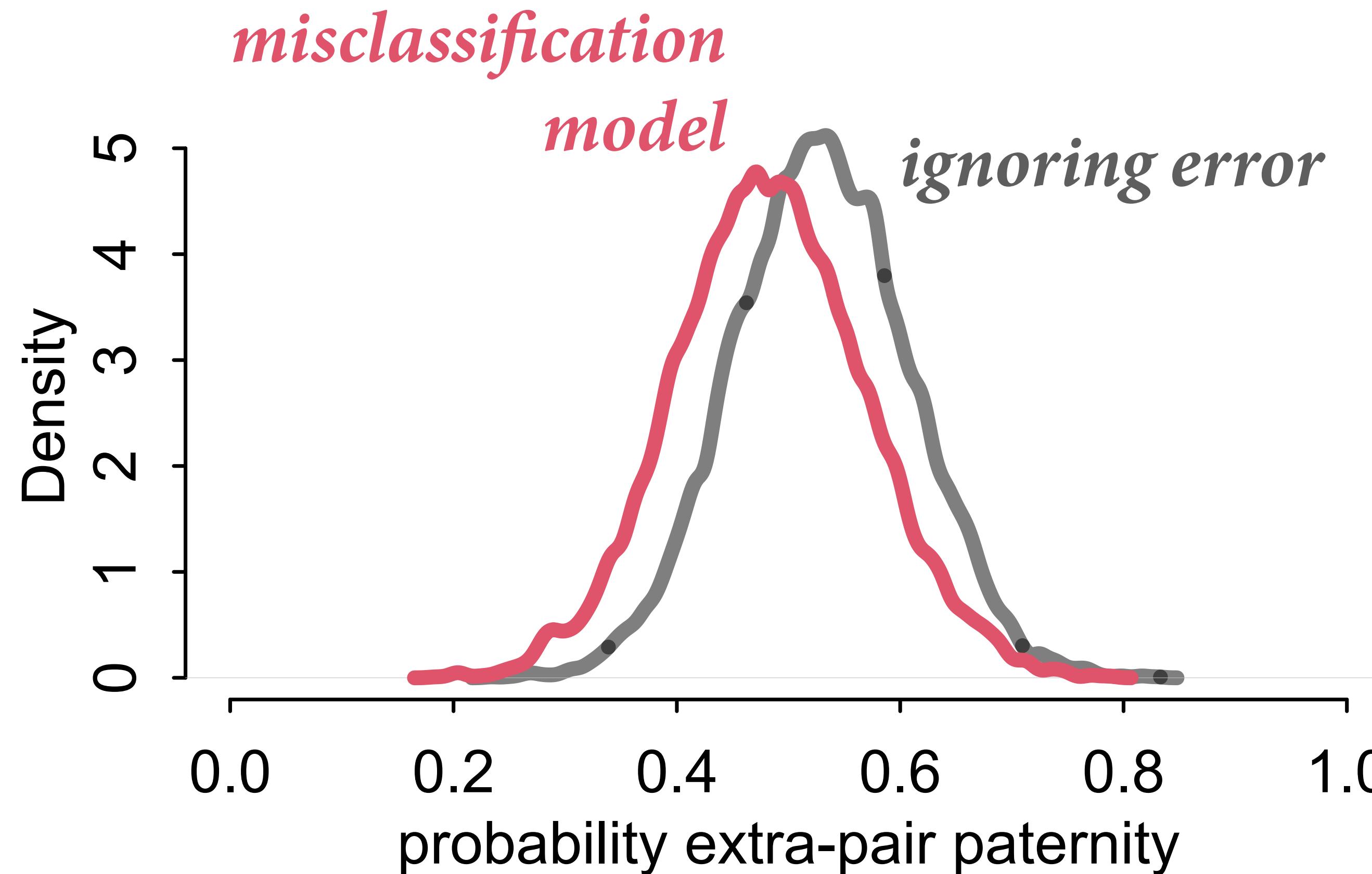
$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# numerically stable version of model - see BONUS for explanation
mX3 <- ulam(
  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==1 ~ custom( log_sum_exp( log_p , log1m_exp(log_p)+log(f) ) ) ,

    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m_exp(log_p) + log1m(f) ) ,

    log_p <- log_inv_logit( a + z[mom_id]*sigma + x[dyad_id]*tau ) ,
    a ~ normal(0,1.5),
    z[mom_id] ~ normal(0,1),
    sigma ~ normal(0,1),
    x[dyad_id] ~ normal(0,1),
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```



$$X_i \sim \text{Bernoulli}(p_i)$$

$$\text{logit}(p_i) = \alpha + \mu_{M[i]} + \delta_{D[i]}$$

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

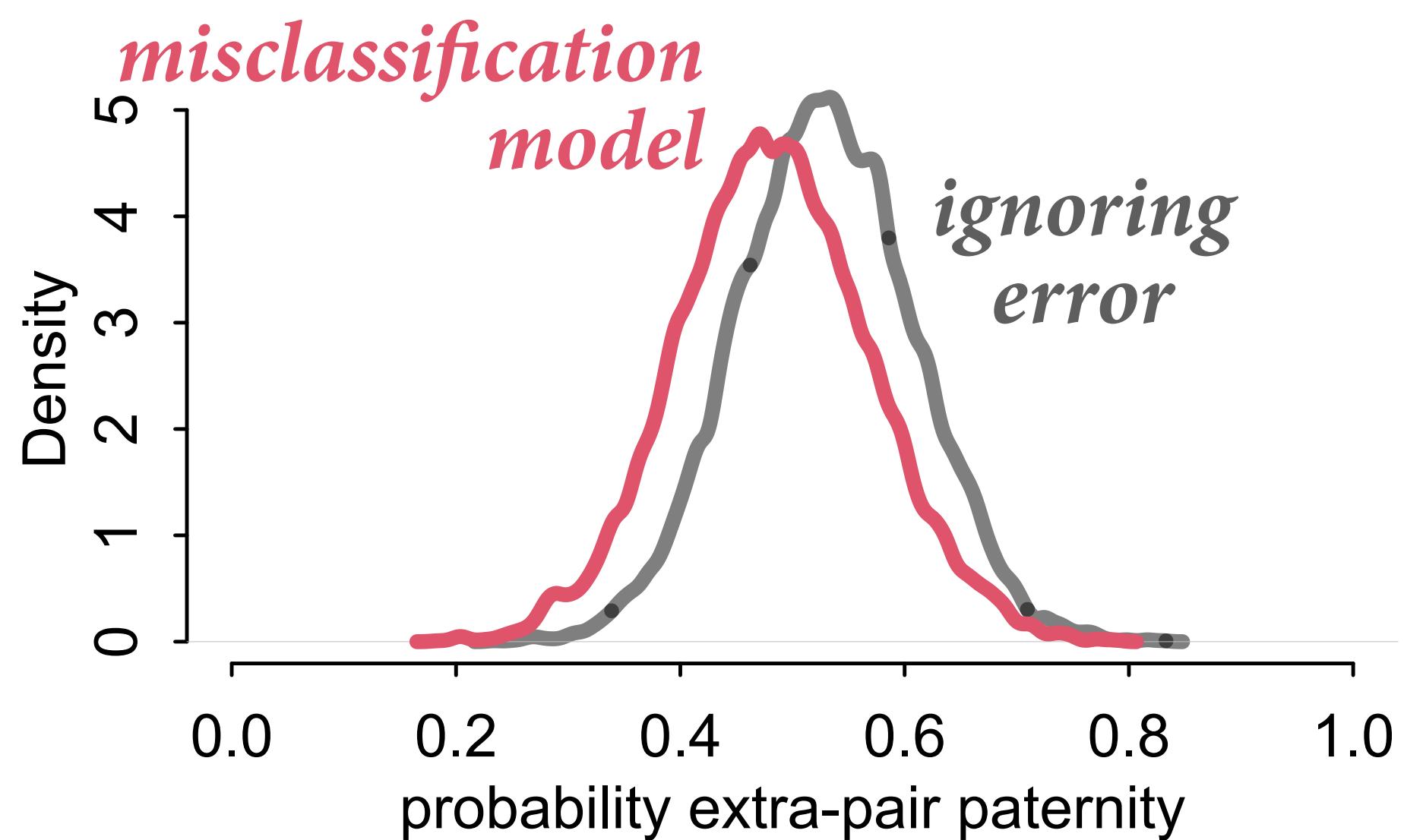
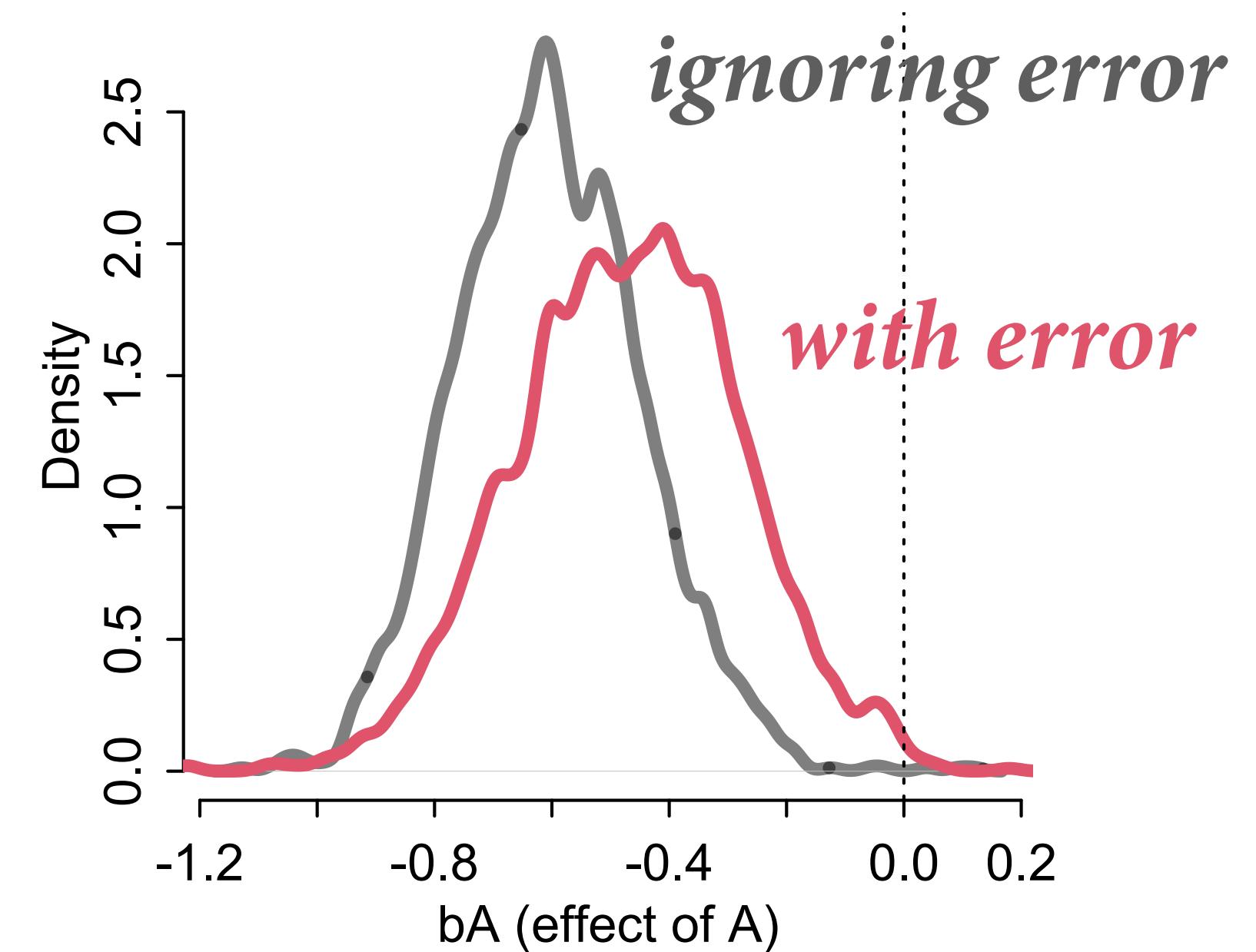
Measurement Horizons

Plenty of related problems and solutions

Rating and assessment: Judges and tests
are noisy — item response theory and
factor analysis

Hurdle models: Thresholds for detection

Occupancy models: Not detecting
something doesn't mean it isn't there



Course Schedule

Week 1	Bayesian inference	Chapters 1, 2, 3
Week 2	Linear models & Causal Inference	Chapter 4
Week 3	Causes, Confounds & Colliders	Chapters 5 & 6
Week 4	Overfitting / MCMC	Chapters 7, 8, 9
Week 5	Generalized Linear Models	Chapters 10, 11
Week 6	Ordered categories & Multilevel models	Chapters 12 & 13
Week 7	More Multilevel models	Chapters 13 & 14
Week 8	Social Networks & Gaussian Processes	Chapter 14
Week 9	Measurement & Missingness	Chapter 15
Week 10	Generalized Linear Madness	Chapter 16

https://github.com/rmcelreath/stat_rethinking_2023

BONUS

Floating Point Monsters

Probability calculations tend to underflow (round to zero) and overflow (round to one)

Solution: Calculate on log scale

Ancient weapons:

`log_sum_exp`, `log1m`, `log1m_exp`



Logarithms make sense

$$\Pr(X_i^\star = 0 \mid p_i) = (1 - p_i)(1 - f)$$

Logarithms make sense

$$\Pr(X_i^\star = 0 \mid p_i) = (1 - p_i)(1 - f)$$

$$\log \Pr(X_i^\star = 0 \mid p_i) = \log[(1 - p_i)(1 - f)]$$

Logarithms make sense

$$\Pr(X_i^\star = 0 \mid p_i) = (1 - p_i)(1 - f)$$

$$\begin{aligned}\log \Pr(X_i^\star = 0 \mid p_i) &= \log[(1 - p_i)(1 - f)] \\ &= \log(1 - p_i) + \log(1 - f)\end{aligned}$$

Devil in the details

$$\log \Pr(X_i^\star = 0 | p_i) = \log(1 - p_i) + \log(1 - f)$$

If p_i is close to zero, $\log(1-p_i)$ could evaluate to zero

Devil in the details

$$\log \Pr(X_i^\star = 0 | p_i) = \log(1 - p_i) + \log(1 - f)$$

If p_i is close to zero, $\log(1-p_i)$ could evaluate to zero

```
> log( 1 - 0.01 )
[1] -0.01005034
> log( 1 - 1e-10 )
[1] -1e-10
> log( 1 - 1e-90 )
[1] 0
```

Devil in the details

$$\log \Pr(X_i^\star = 0 | p_i) = \log(1 - p_i) + \log(1 - f)$$

If p_i is close to zero, $\log(1-p_i)$ could evaluate to zero

```
> log( 1 - 0.01 )
[1] -0.01005034
> log( 1 - 1e-10 )
[1] -1e-10
> log( 1 - 1e-90 )
[1] 0
```

```
> log1p( -1e-90 )
[1] -1e-90
```

$$\log1p(-x) = \log1m(x)$$

Devil in the details

Okay, but what is $\log1p(x)$? And how do we program it?

When $x > 1e-4$, no reason to worry

When $x < 1e-4$, use a Taylor series approximation:

$$\log(1 + x) = x - \frac{x^2}{2} + \frac{x^3}{3} + \dots$$

second-order approximation, for x near zero

log_sum_exp

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\log \Pr(X_i^\star = 1 | p_i) = \log[p_i + (1 - p_i)f]$$

log_sum_exp

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\log \Pr(X_i^\star = 1 | p_i) = \log[p_i + (1 - p_i)f]$$

$$= \text{log_sum_exp}(\log p_i, \log(1 - p_i) + \log f)$$

log_sum_exp

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\log \Pr(X_i^\star = 1 | p_i) = \log[p_i + (1 - p_i)f]$$

$$= \text{log_sum_exp}(\log p_i, \log(1 - p_i) + \log f)$$

$$= \text{log_sum_exp}(\log p_i, \text{log1m}(p_i) + \log f)$$

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# numerically kosher version
mX2 <- ulam(
  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==1 ~ custom( log_sum_exp( log(p) , log1m(p)+log(f) ) ) ,

    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m(p) + log1m(f) ) ,

    logit(p) <- a + z[mom_id]*sigma + x[dyad_id]*tau,
    a ~ normal(0,1.5),
    z[mom_id] ~ normal(0,1),
    sigma ~ normal(0,1),
    x[dyad_id] ~ normal(0,1),
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^{\star} = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^{\star} = 0 | p_i) = (1 - p_i)(1 - f)$$

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    X|X==0 ~ custom( log1m(p) + log1m(f) ) ,
    logit(p) <- a + z[mom_id]*sigma + x[dyad_id]*tau ,
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    z[mom_id] ~ normal(0,1) ,
    sigma ~ normal(0,1) ,
    x[dyad_id] ~ normal(0,1) ,
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

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    x[dyad_id] ~ normal(0,1) ,
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
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```

$$\Pr(X_i^\star = 1 | p_i) = p_i - \boxed{(1 - p_i)f}$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# numerically kosher version
mX2 <- ulam(
  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
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    x[dyad_id] ~ normal(0,1) ,
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# numerically kosher version
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  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==1 ~ custom( log_sum_exp( log(p) , log1m(p)+log(f) ) ) ,
    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m(p) + log1m(f) ) ,

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    a ~ normal(0,1.5),
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    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

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  alist(
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    X|X==1 ~ custom( log_sum_exp( log(p) , log1m(p)+log(f) ) ) ,

    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m(p) + log1m(f) ) ,
```

log1m(p) + log1m(f)

```
    logit(p) <- a + z[mom_id]*sigma + x[dyad_id]*tau ,
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    x[dyad_id] ~ normal(0,1) ,
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# double kosher
mX3 <- ulam(
  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==1 ~ custom( log_sum_exp( log_p , log1m_exp(log_p)+log(f) ) ) ,

    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m_exp(log_p) + log1m(f) ) ,

    log_p <- log_inv_logit( a + z[mom_id]*sigma + x[dyad_id]*tau ) ,
    a ~ normal(0,1.5),
    z[mom_id] ~ normal(0,1),
    sigma ~ normal(0,1),
    x[dyad_id] ~ normal(0,1),
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
constraints=list(sigma="lower=0",tau="lower=0") )
```

$$\Pr(X_i^\star = 1 | p_i) = p_i + (1 - p_i)f$$

$$\Pr(X_i^\star = 0 | p_i) = (1 - p_i)(1 - f)$$

```
# double kosher
mX3 <- ulam(
  alist(
    #X|X==1 ~ custom( log( p + (1-p)*f ) ) ,
    X|X==1 ~ custom( log_sum_exp( log_p , log1m_exp(log_p)+log(f) ) ) ,
    #X|X==0 ~ custom( log( (1-p)*(1-f) ) ) ,
    X|X==0 ~ custom( log1m_exp(log_p) + log1m(f) ) ,
    log_p <- log_inv_logit( a + z[mom_id]*sigma + x[dyad_id]*tau ) ,
    a ~ normal(0,1.5) ,
    z[mom_id] ~ normal(0,1) ,
    sigma ~ normal(0,1) ,
    x[dyad_id] ~ normal(0,1) ,
    tau ~ normal(0,1)
  ) , data=dat , chains=4 , cores=4 , iter=4000 ,
  constraints=list(sigma="lower=0",tau="lower=0") )
```

Floating Point Monsters

Ancient weapons:

`log_sum_exp`, `log1m`, `log1m_exp`

Built into most math libraries

Often unnecessary, but when you
need them, no better defense

**FLOATING
POINT ARITHMETIC**

log_sum_exp

