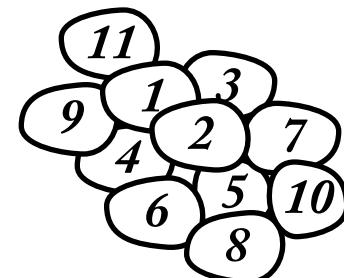


Statistical Rethinking

Winter 2019

Lecture 11 / Week 6

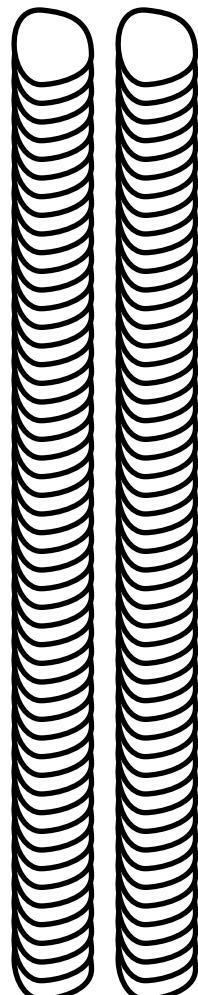
Big Entropy & The
Generalized Linear Model



100 pebbles



100



1



2



3

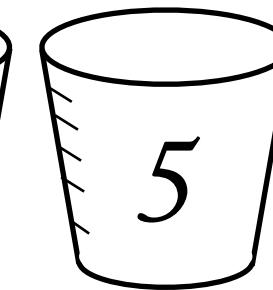
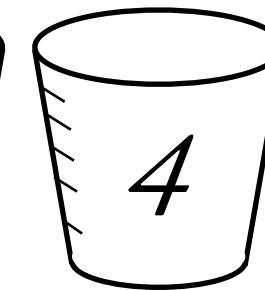
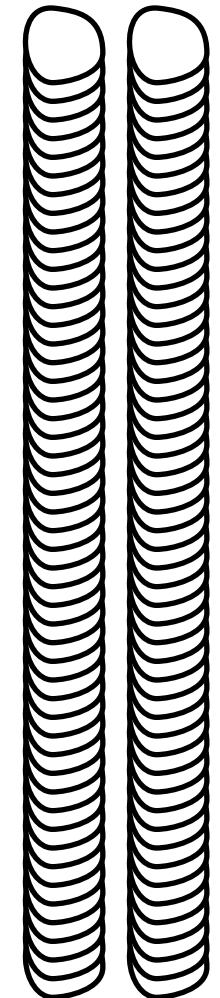


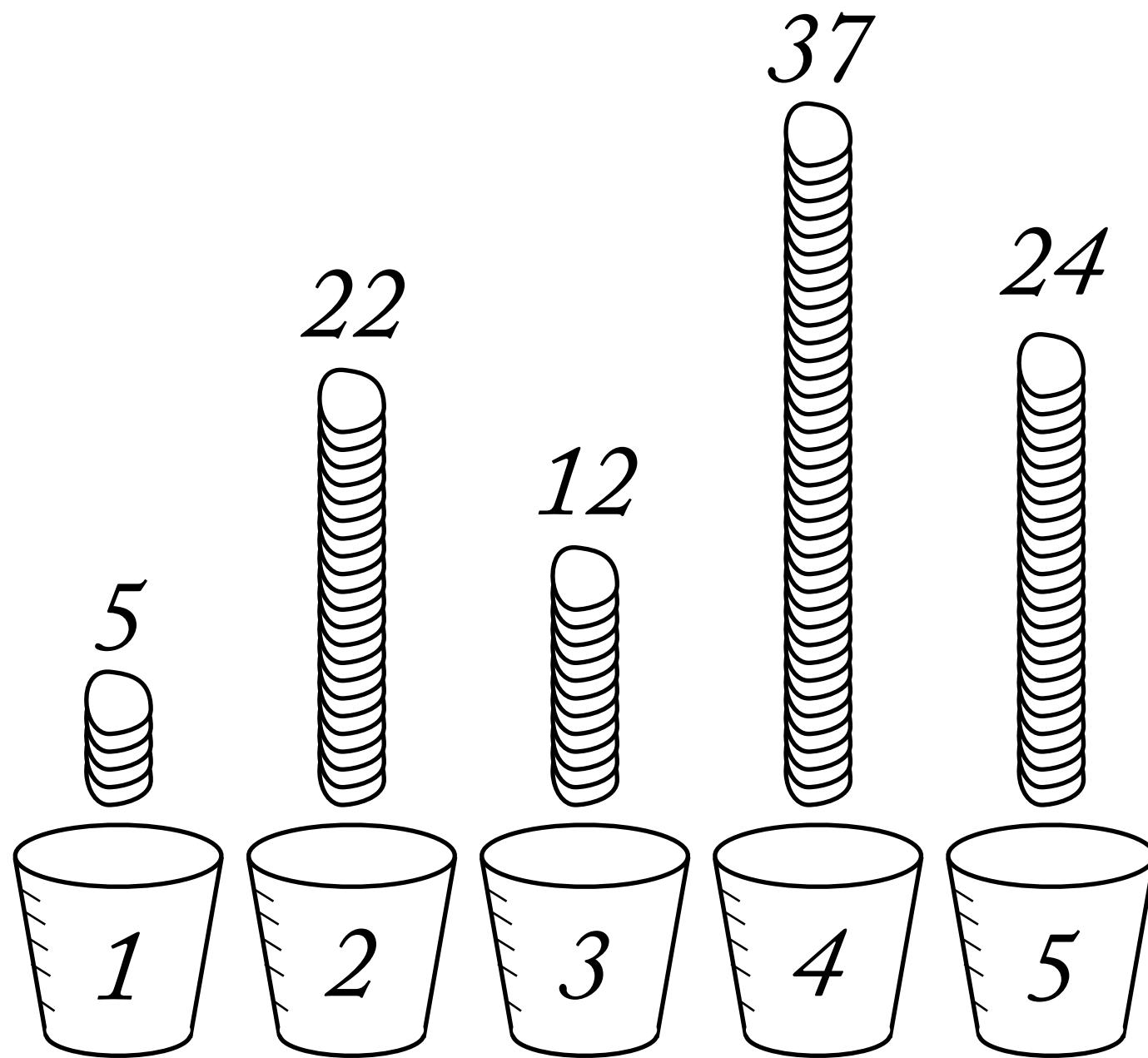
4



5

100





n_1 n_2 n_3 n_4 n_5 

Number of ways:

$$W = \frac{N!}{n_1! n_2! n_3! n_4! n_5!}$$

n_1

n_2

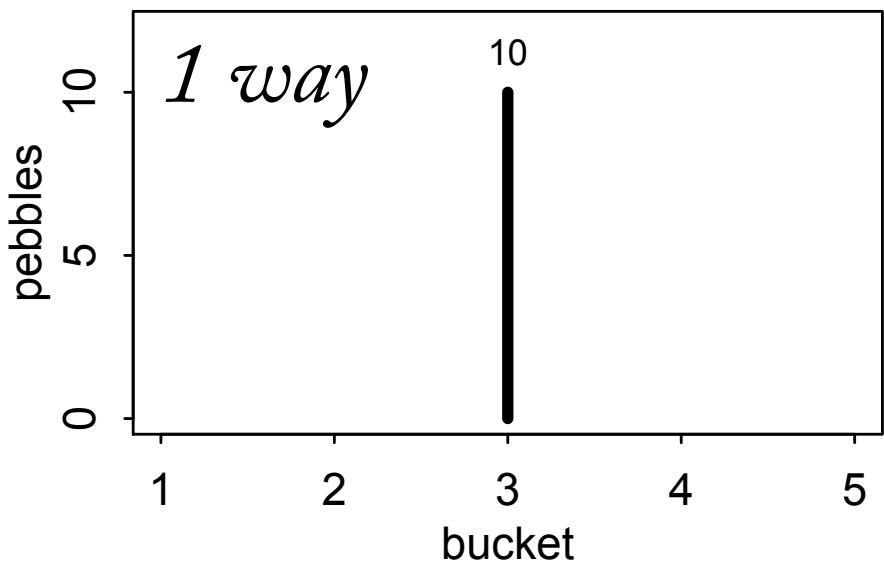
n_3

n_4

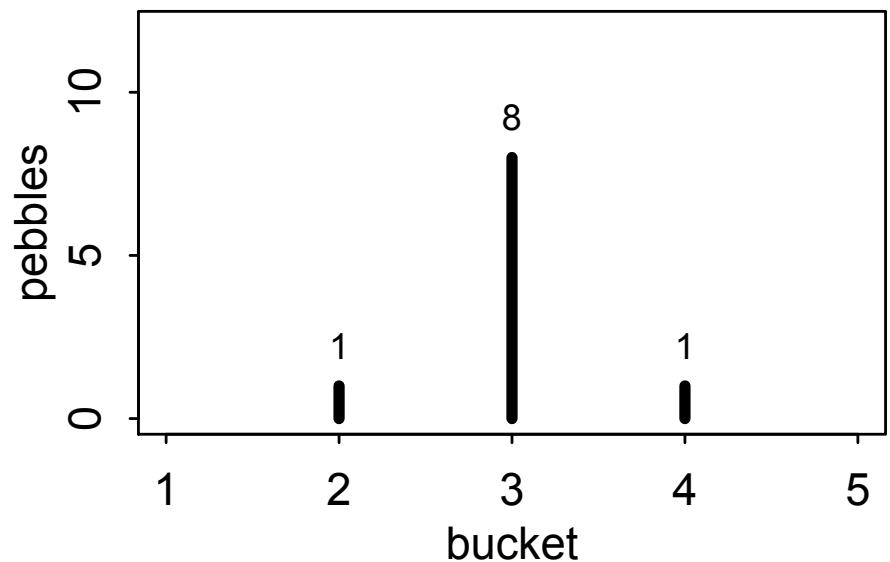
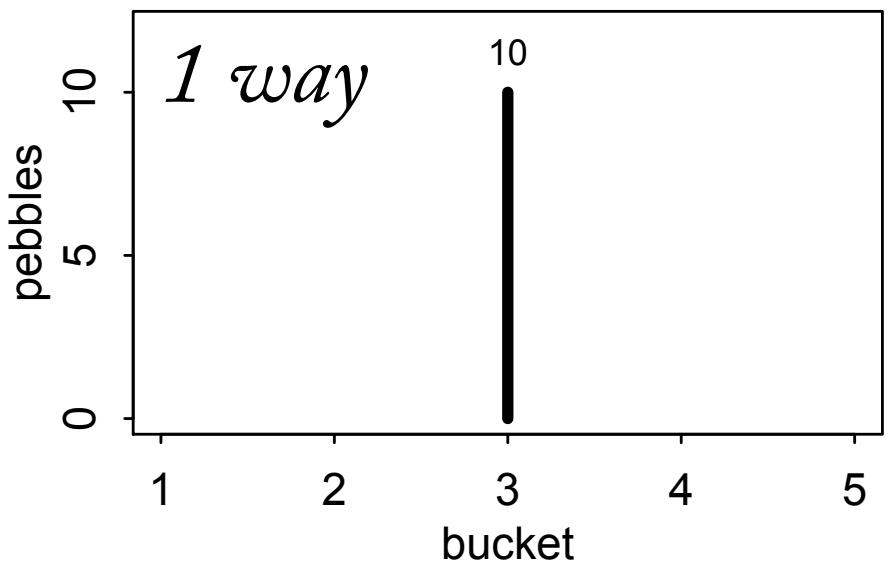
n_5



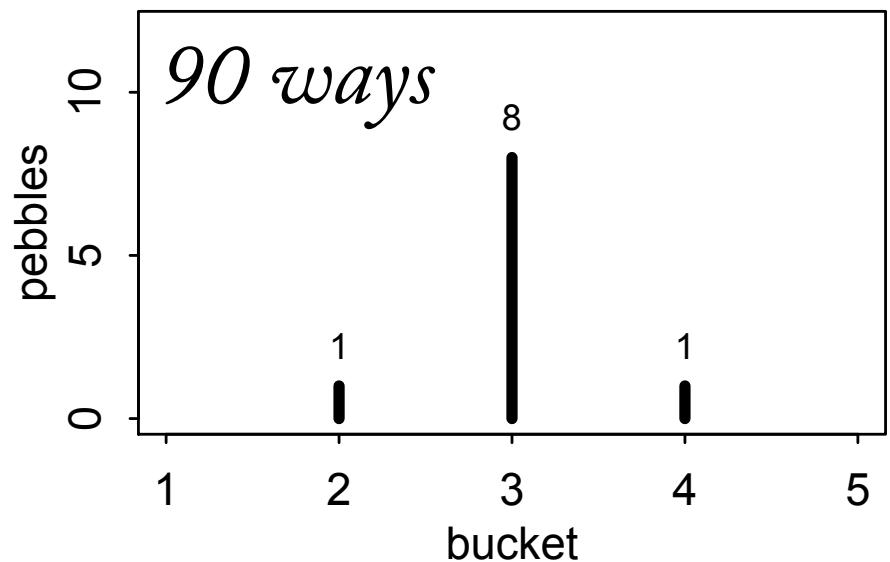
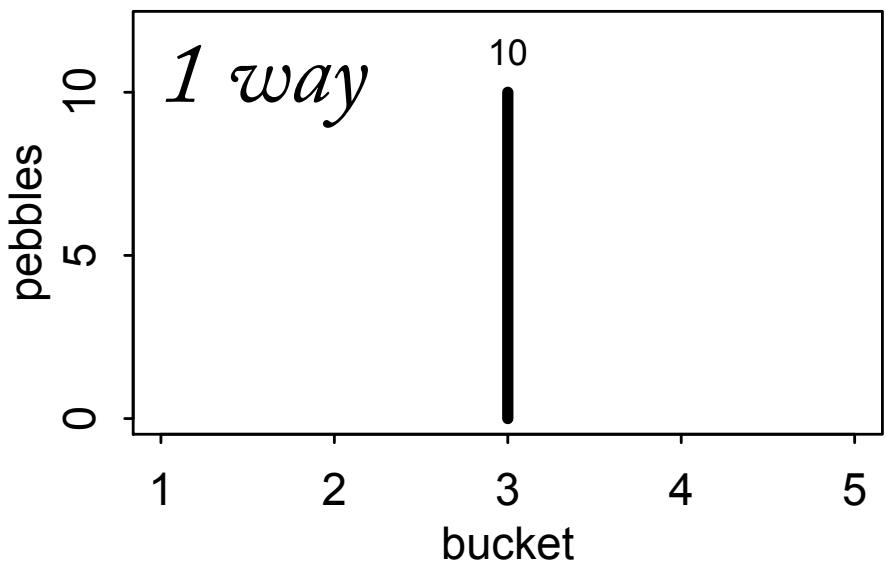
Suppose only 10 pebbles...



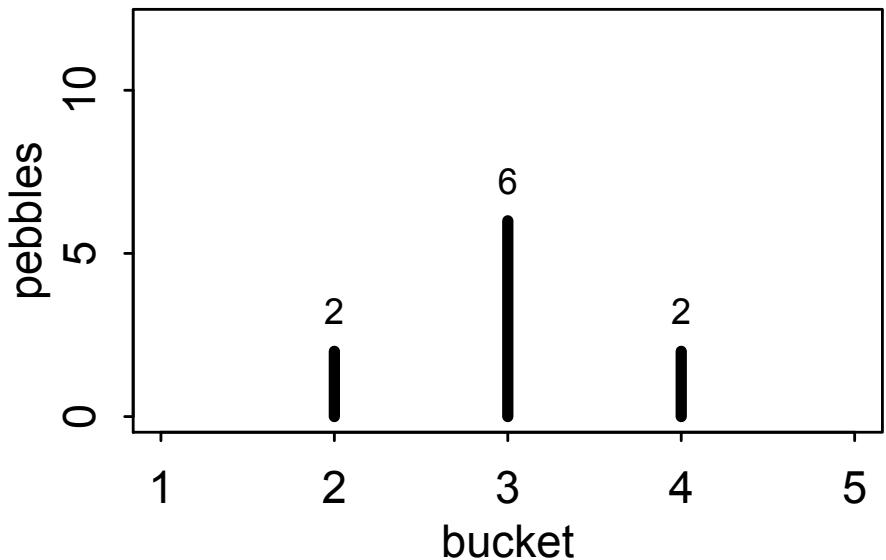
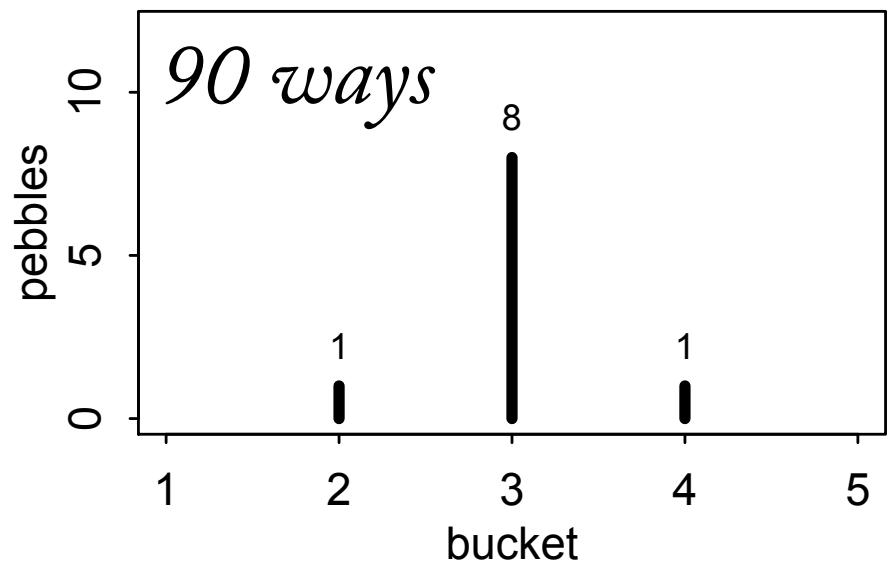
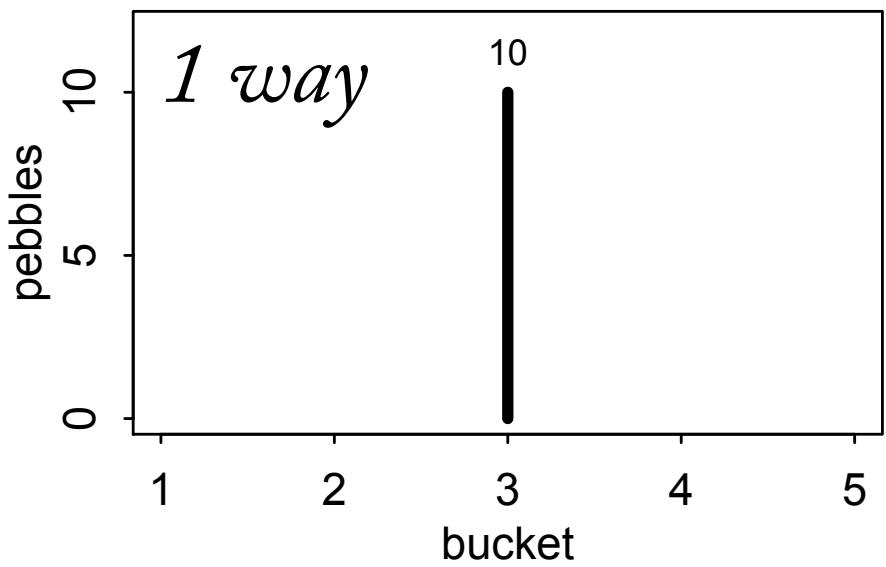
Suppose only 10 pebbles...



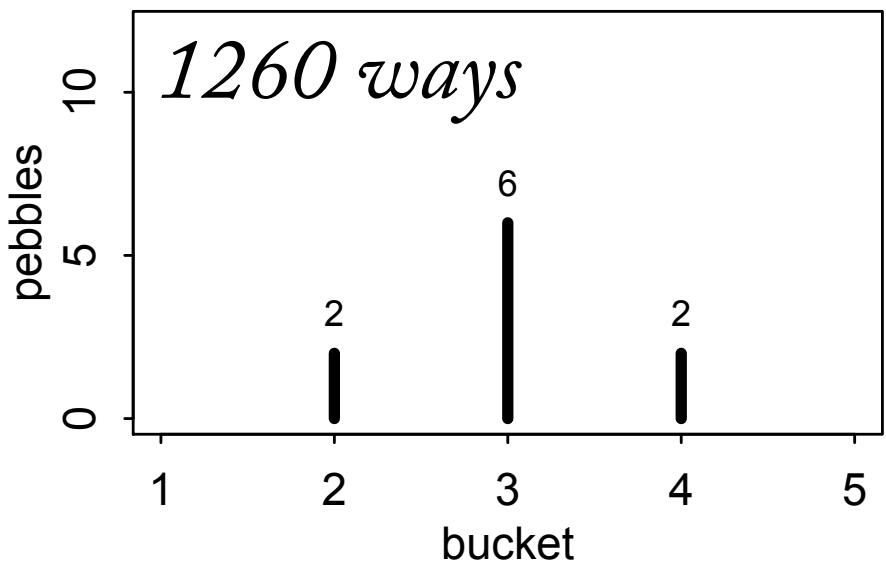
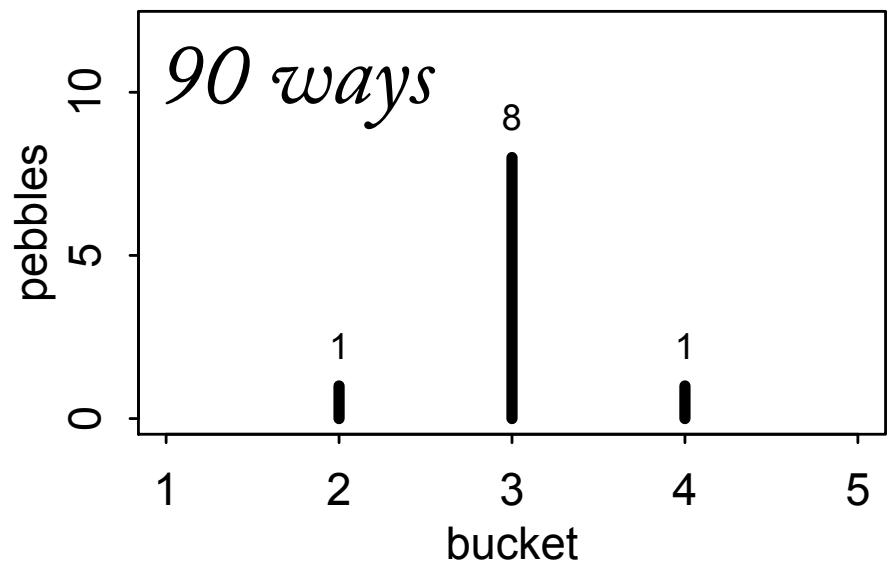
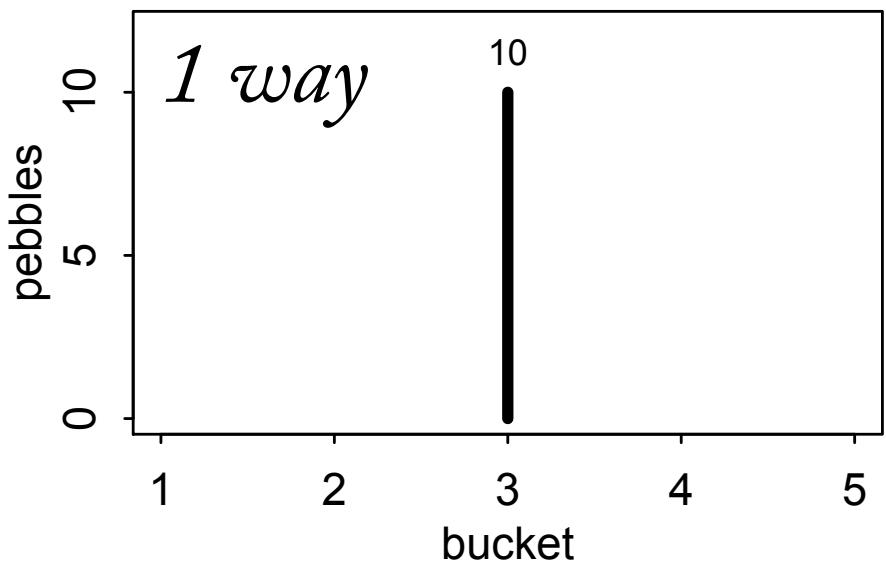
Suppose only 10 pebbles...



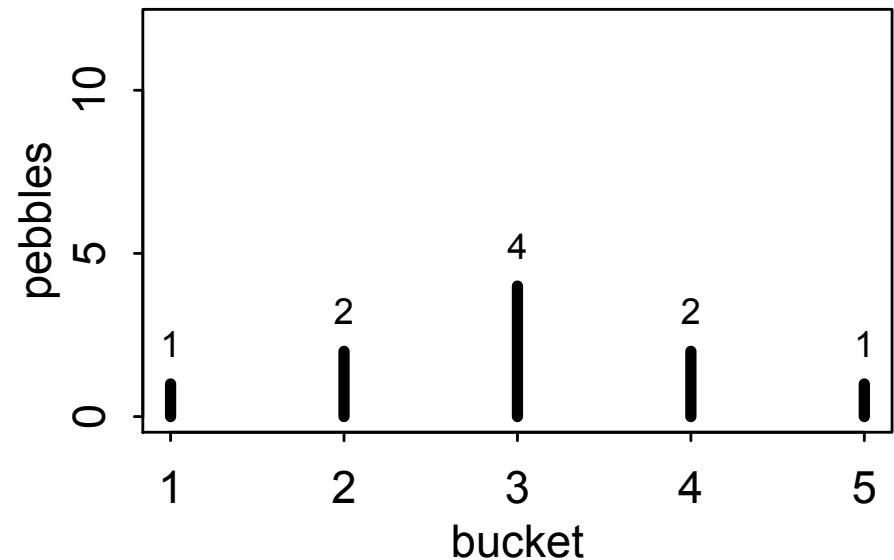
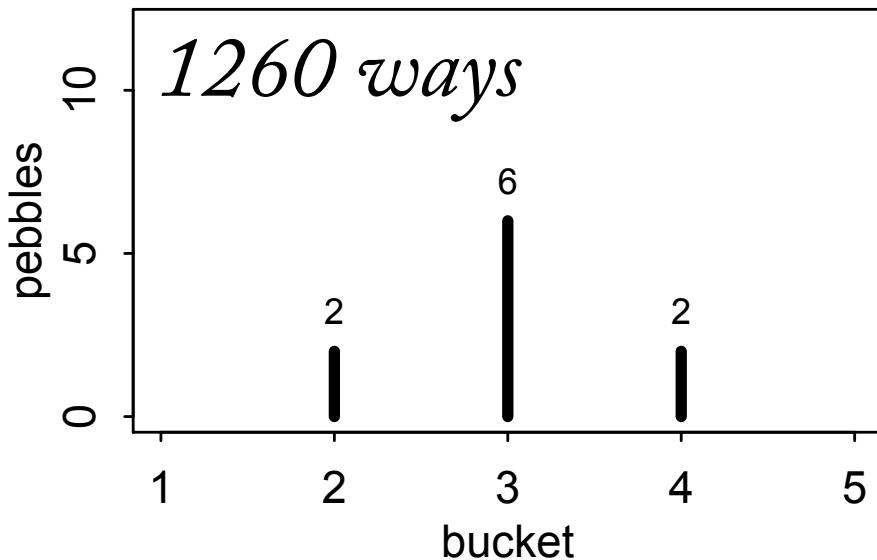
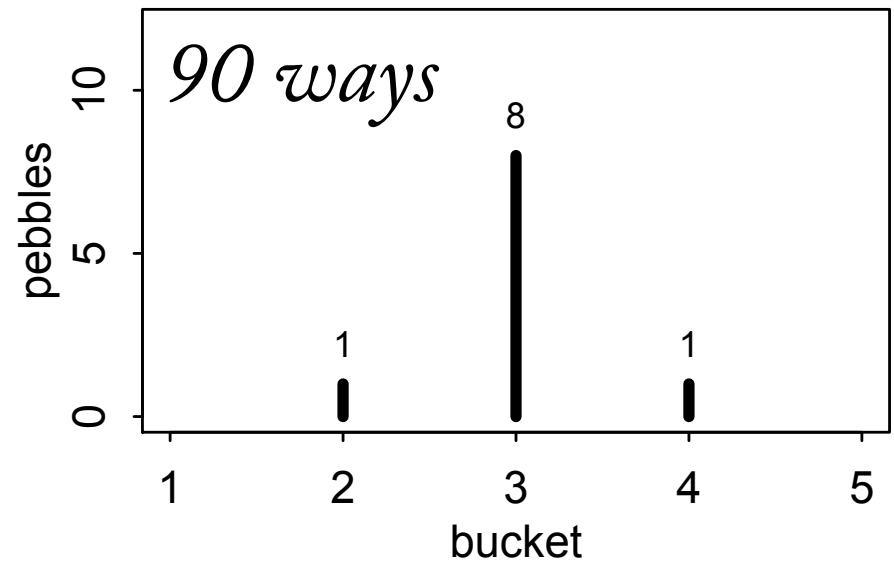
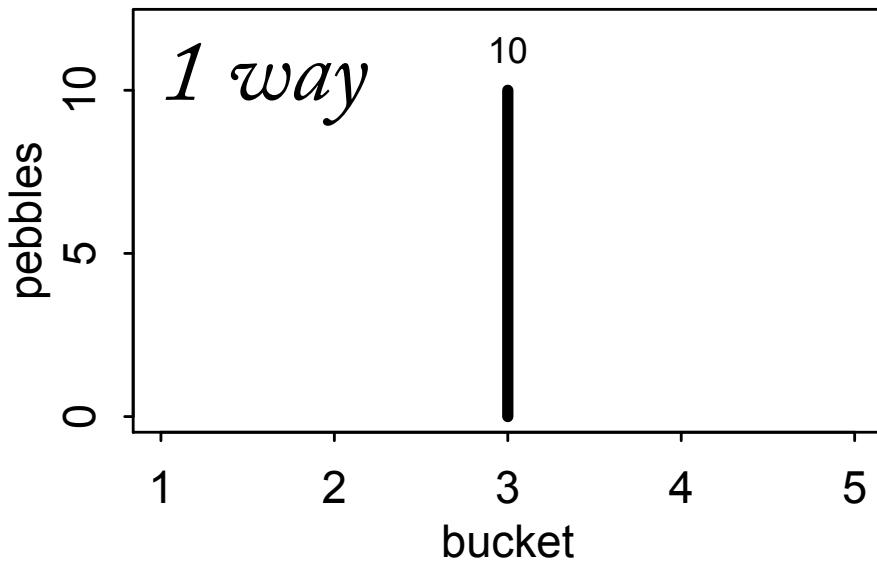
Suppose only 10 pebbles...



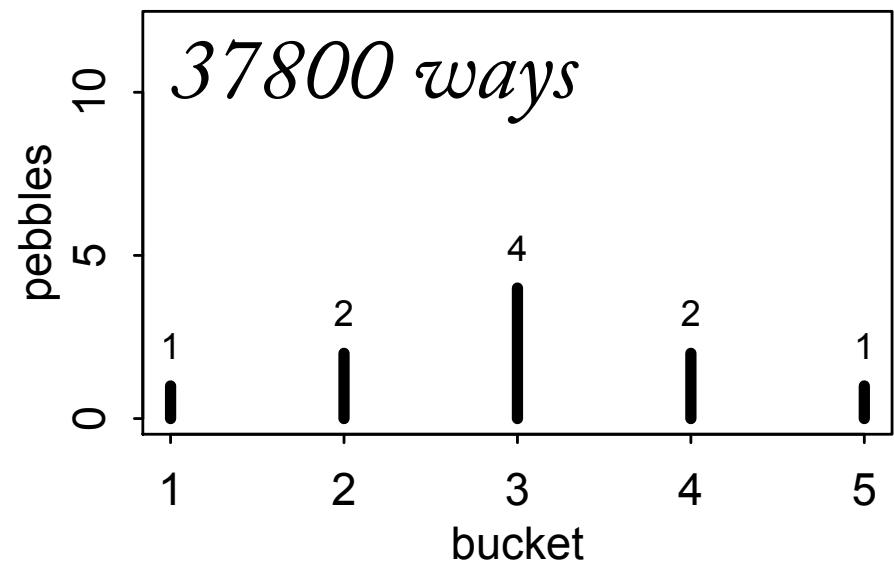
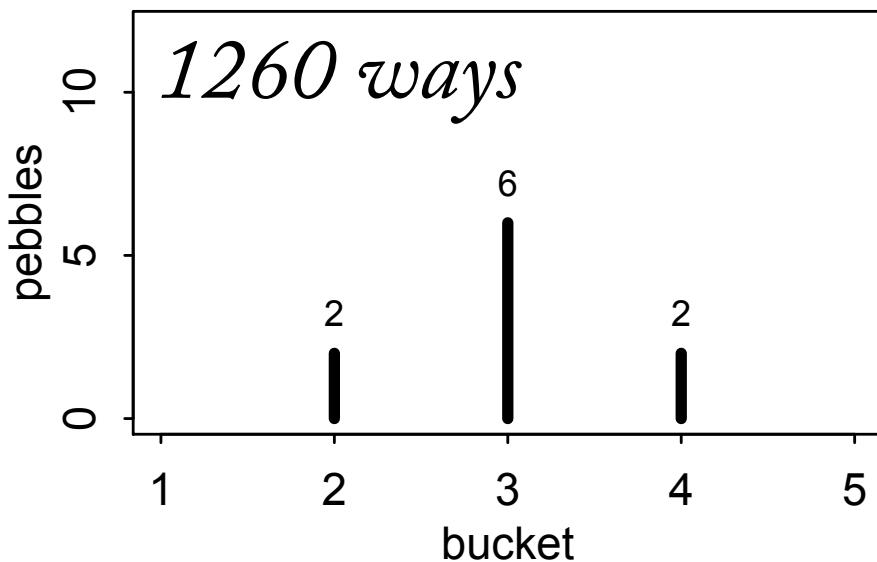
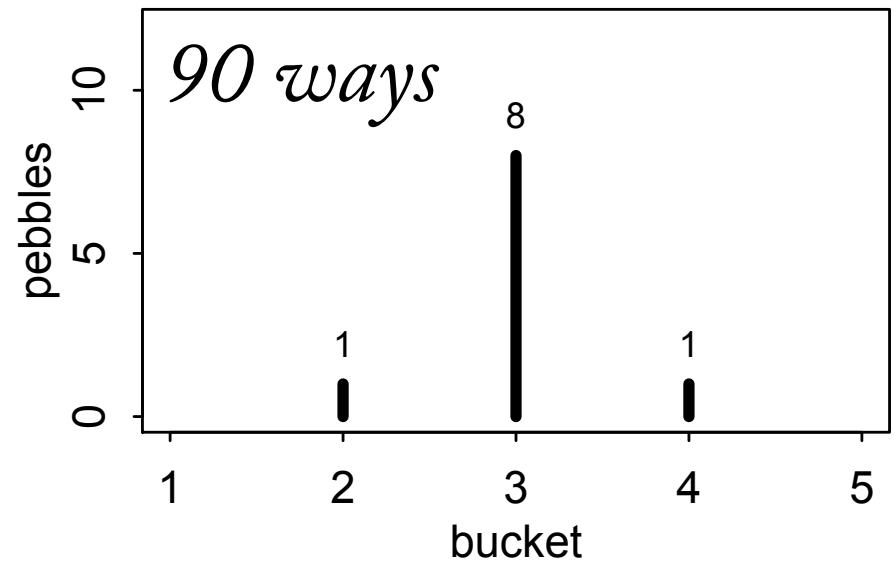
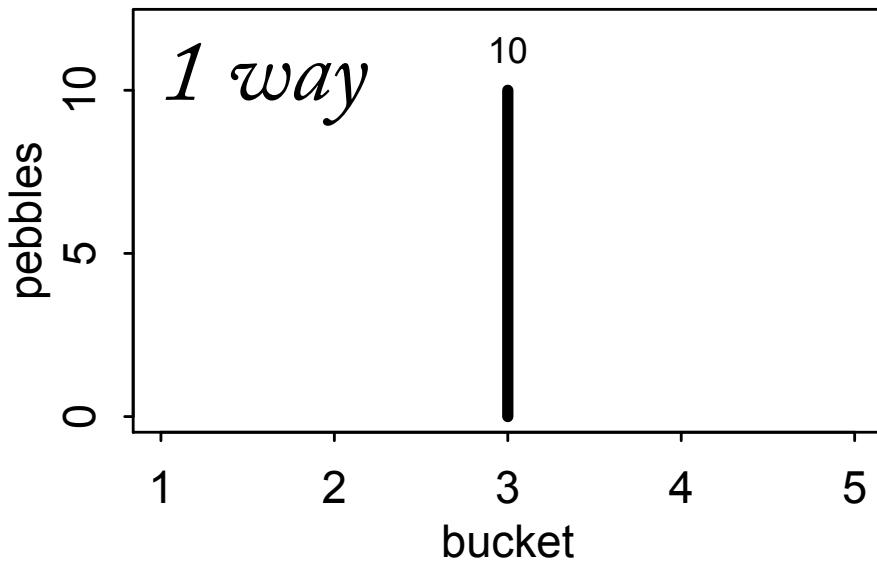
Suppose only 10 pebbles...

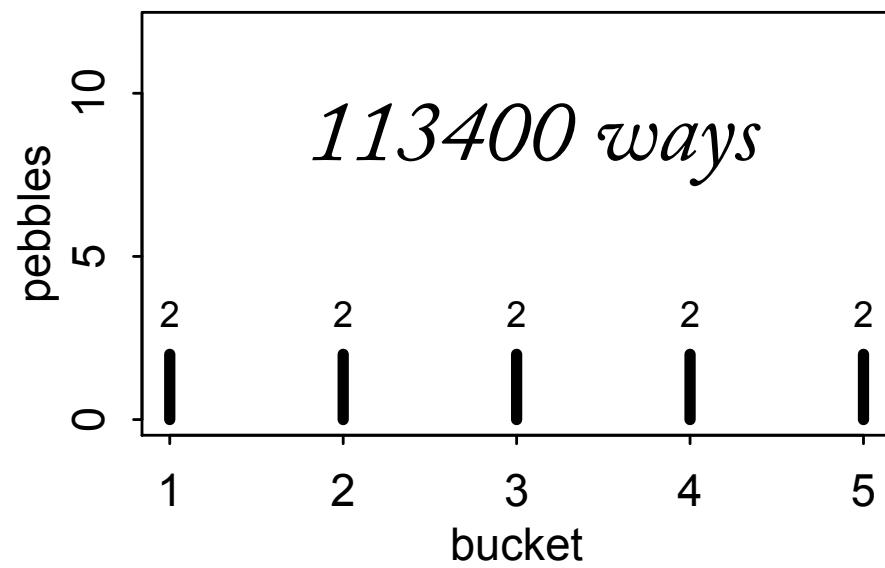
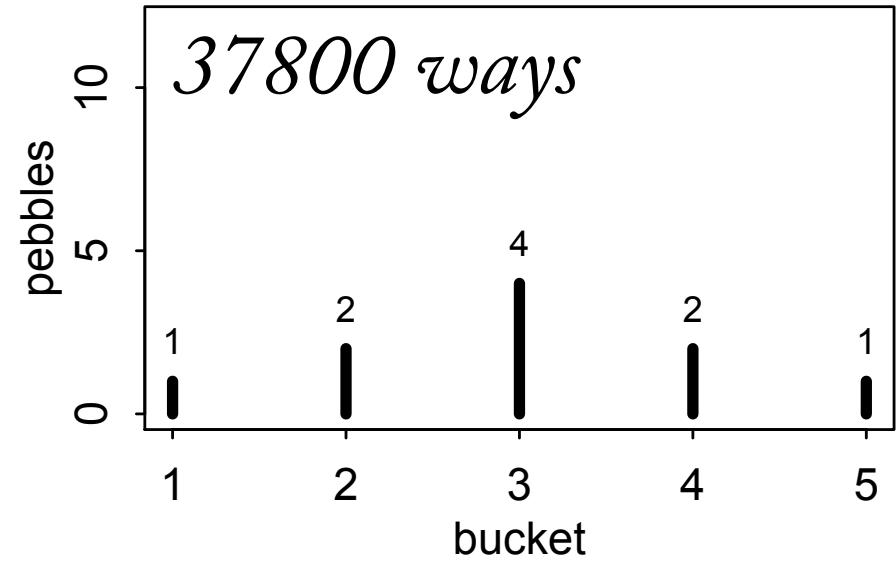
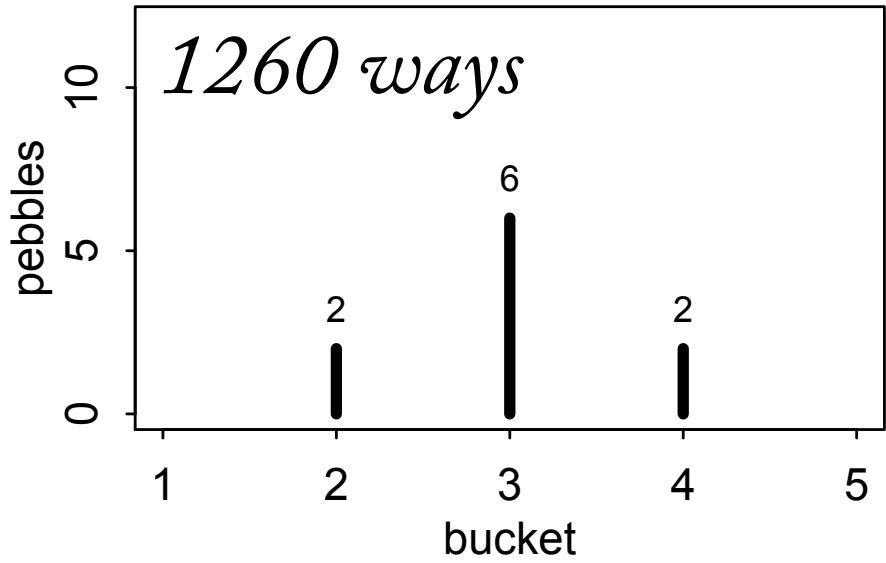


Suppose only 10 pebbles...



Suppose only 10 pebbles...





For large N :

$$\frac{1}{N} \log W \approx - \sum_i \frac{n_i}{N} \log \left(\frac{n_i}{N} \right)$$

n_1

n_2

n_3

n_4

n_5



For large N :

$$\frac{1}{N} \log W \approx - \sum_i \frac{n_i}{N} \log \left(\frac{n_i}{N} \right) = - \sum_i p_i \log p_i$$

n_1

n_2

n_3

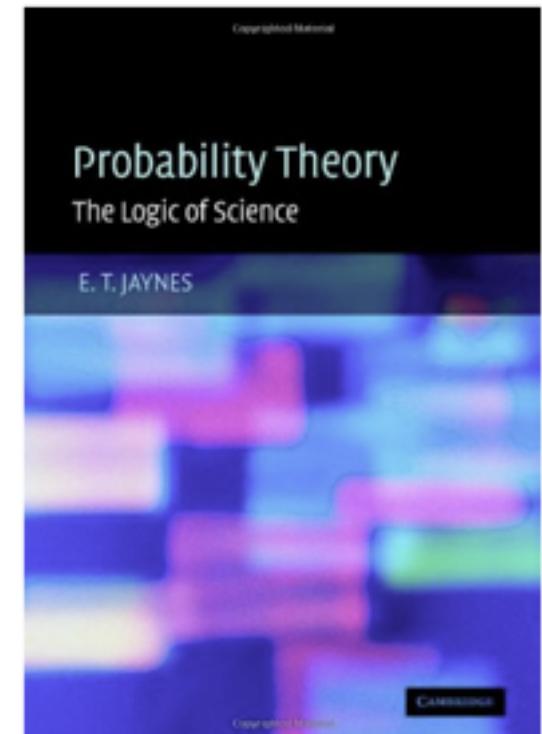
n_4

n_5



Maximum entropy

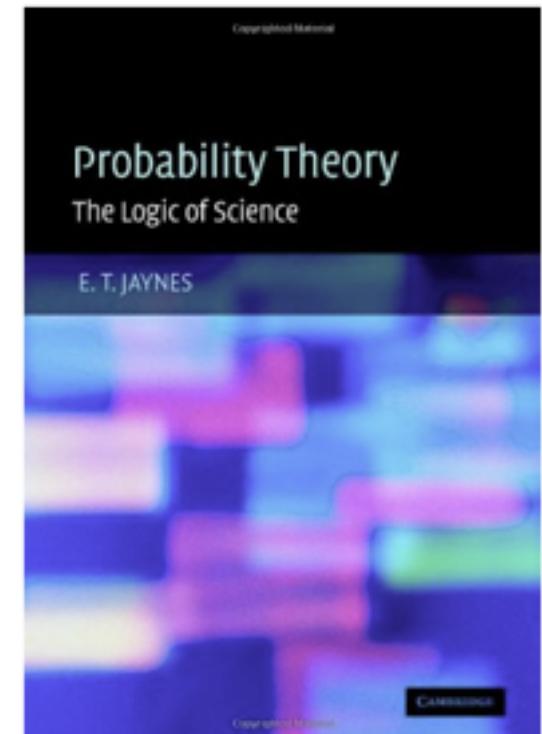
- Due to Edwin T. Jaynes (1922–1998)
- The maxent principle:
 - *Distribution with largest entropy is distribution most consistent with stated assumptions*
 - *Can happen the largest number of ways*



E. T. Jaynes (1922–1998)

Maximum entropy

- Due to Edwin T. Jaynes (1922–1998)
- The maxent principle:
 - *Distribution with largest entropy is distribution most consistent with stated assumptions*
 - *Can happen the largest number of ways*
- For parameters, provides way to understand priors
- For observations, way to understand likelihood
- Also reproduces Bayesian updating as special case (*minimum cross-entropy*)



E. T. Jaynes (1922–1998)

Maximum entropy

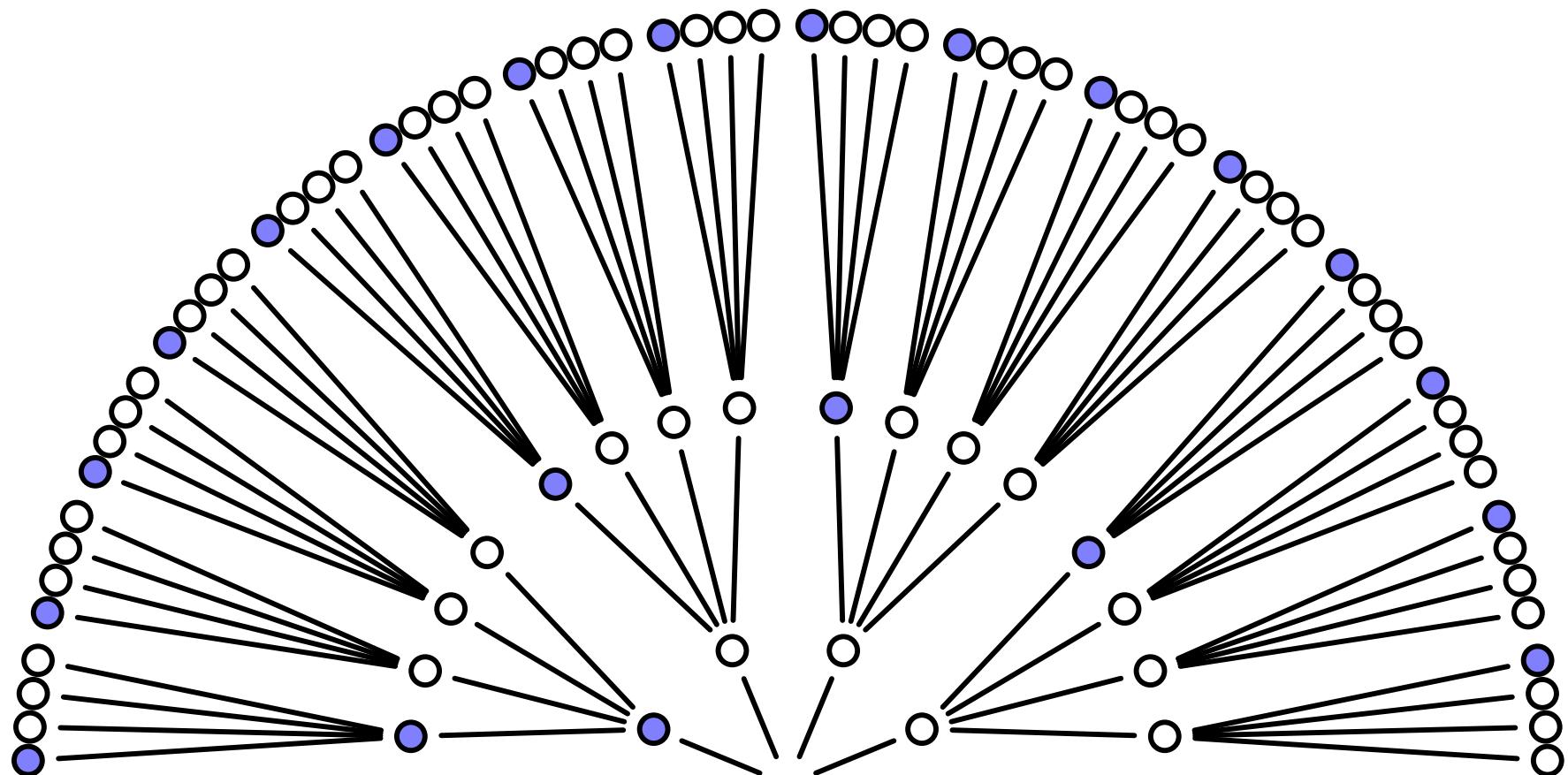
- Ye olde information entropy:

$$H(p) = - \sum_i p_i \log p_i$$

- Q: What kind of distribution maximizes this quantity?
- A: Flattest distribution still consistent with **constraints**. This is the distribution that can happen the most unique ways.
- Whatever does happen, bound to be one of those ways.

Conjecture: 

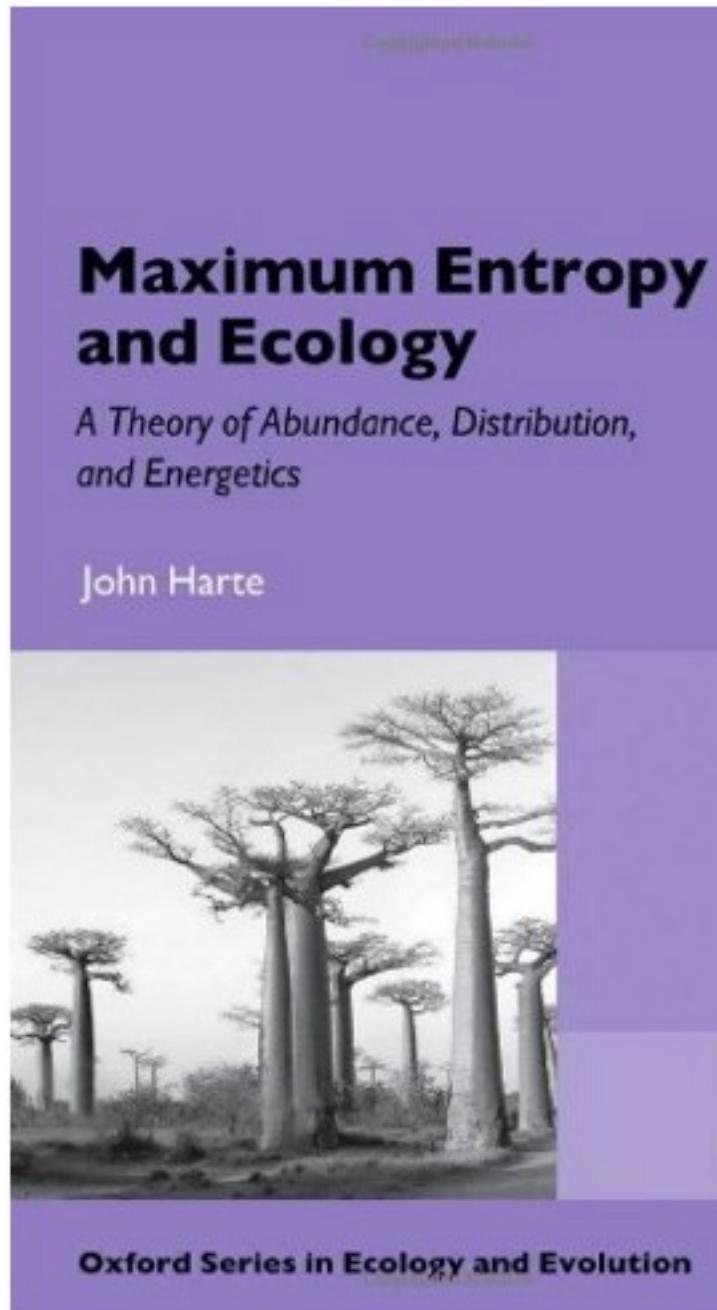
Data: 



*Counting paths produces flattest distribution
consistent with stated constraints.*

Maximum entropy

Constraints	Maxent distribution
Real value in interval	Uniform
Real value, finite variance	Gaussian
Binary events, fixed probability	Binomial
Non-negative real, has mean	Exponential



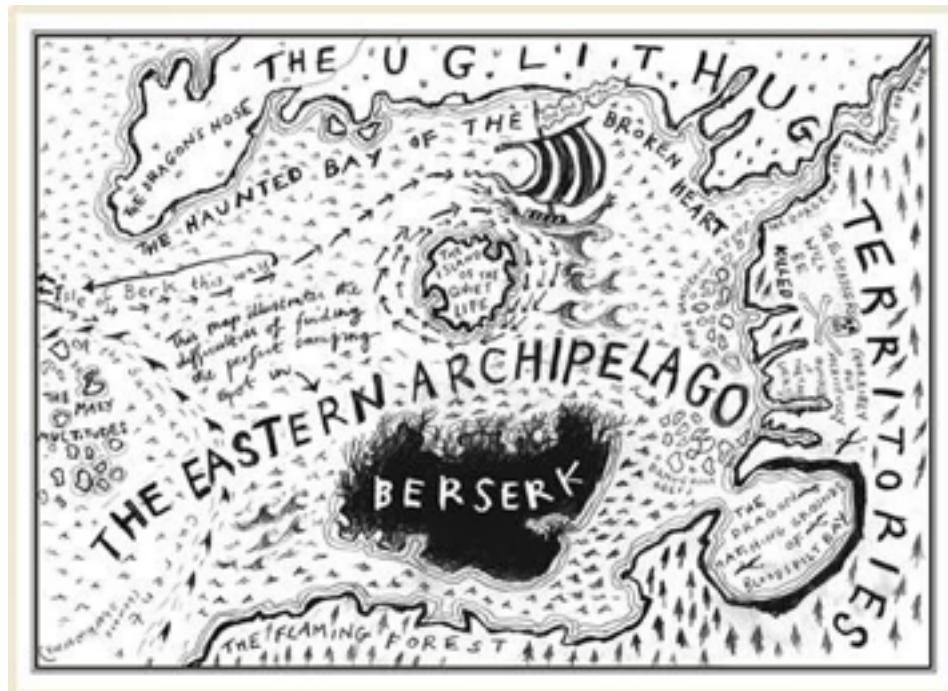
OSEE

Generalized Linear Models

- Goal: Connect linear model to outcome variable
- Still geocentric!
- Strategy:
 1. Pick an outcome distribution
 2. Model its parameters using links to linear models
 3. Compute posterior
- Can model multivariate relationships and non-linear responses
- Building blocks of multilevel models

Generalized Linear Models

- How to pick a data distribution
 - Mostly *exponential family*
 - Arise from natural processes
 - *Maximum entropy* interpretations
 - Select from first principles
 - Resist *histomancy*: Superstitious practice of picking likelihoods by gazing at a histogram



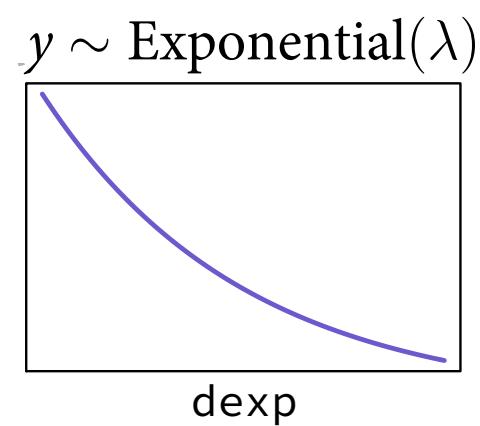
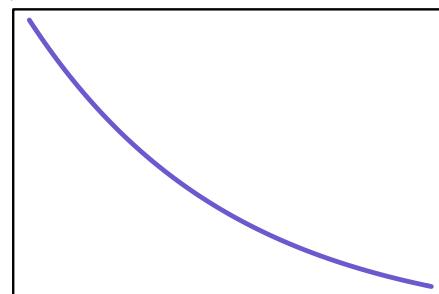
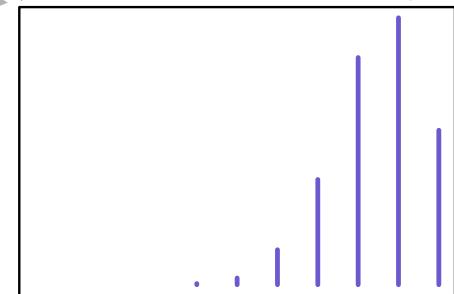


Figure 9.5

$y \sim \text{Exponential}(\lambda)$ 

dexp

*count
events*

 $y \sim \text{Binomial}(n, p)$ 

dbinom

Figure 9.5

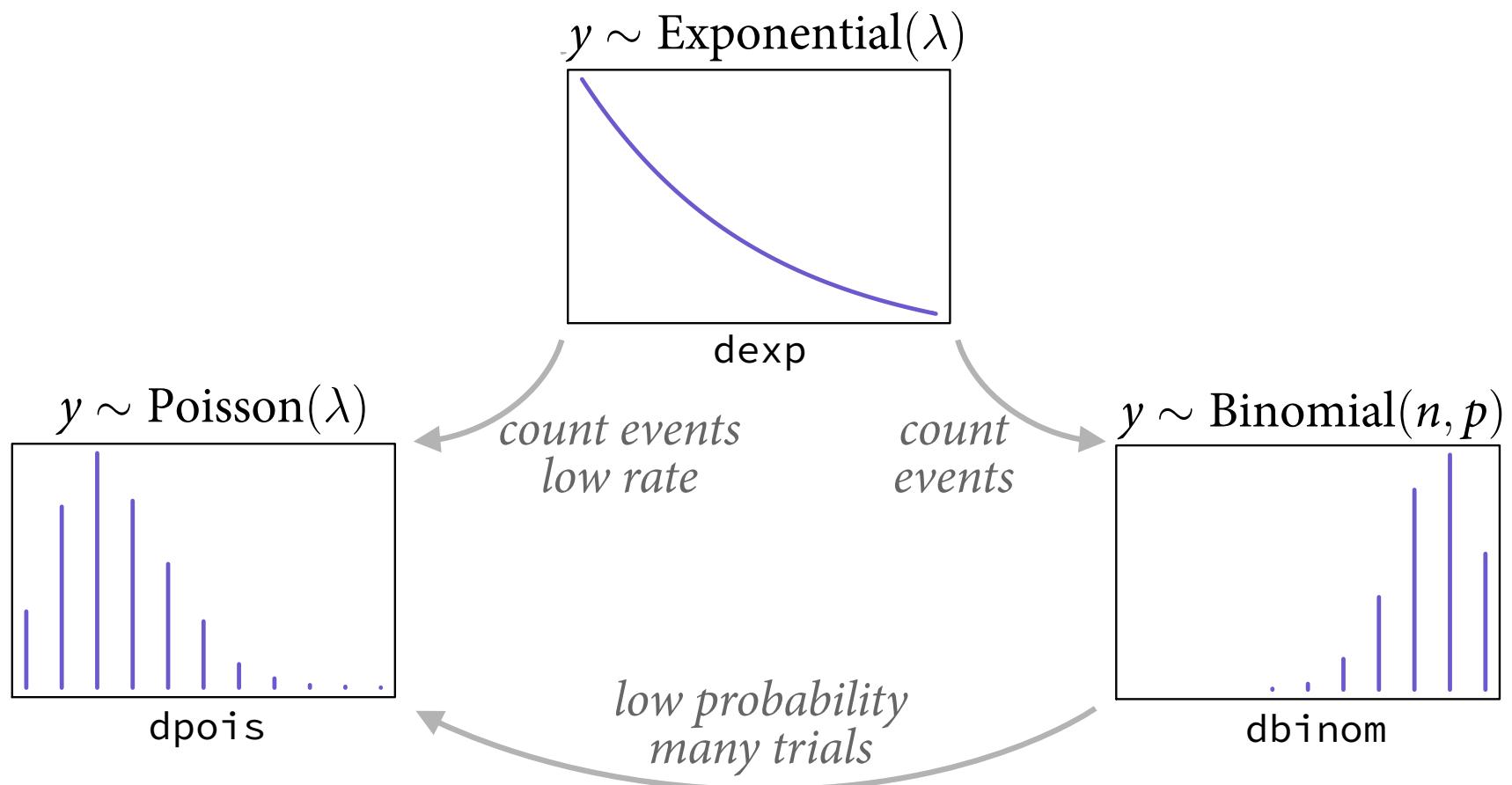
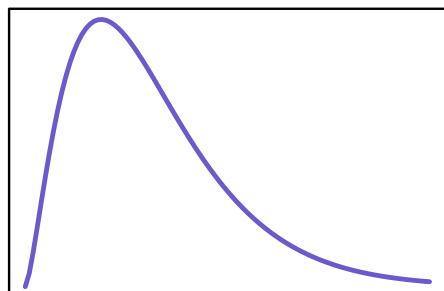


Figure 9.5

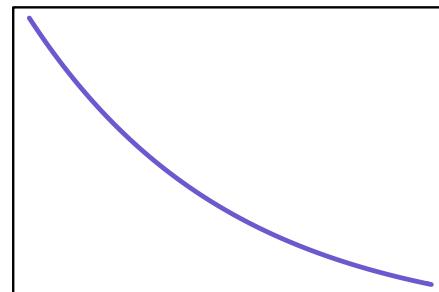
$y \sim \text{Gamma}(\lambda, k)$



dgamma

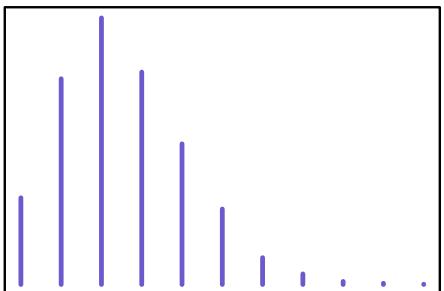
sum

$y \sim \text{Exponential}(\lambda)$



dexp

$y \sim \text{Poisson}(\lambda)$



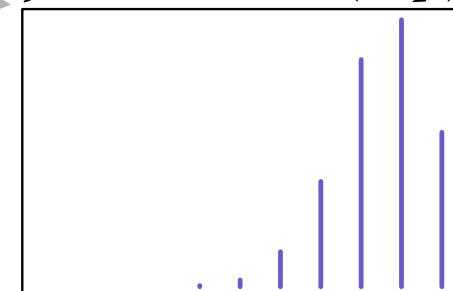
dpois

count events
low rate

count events

low probability
many trials

$y \sim \text{Binomial}(n, p)$



dbinom

Figure 9.5

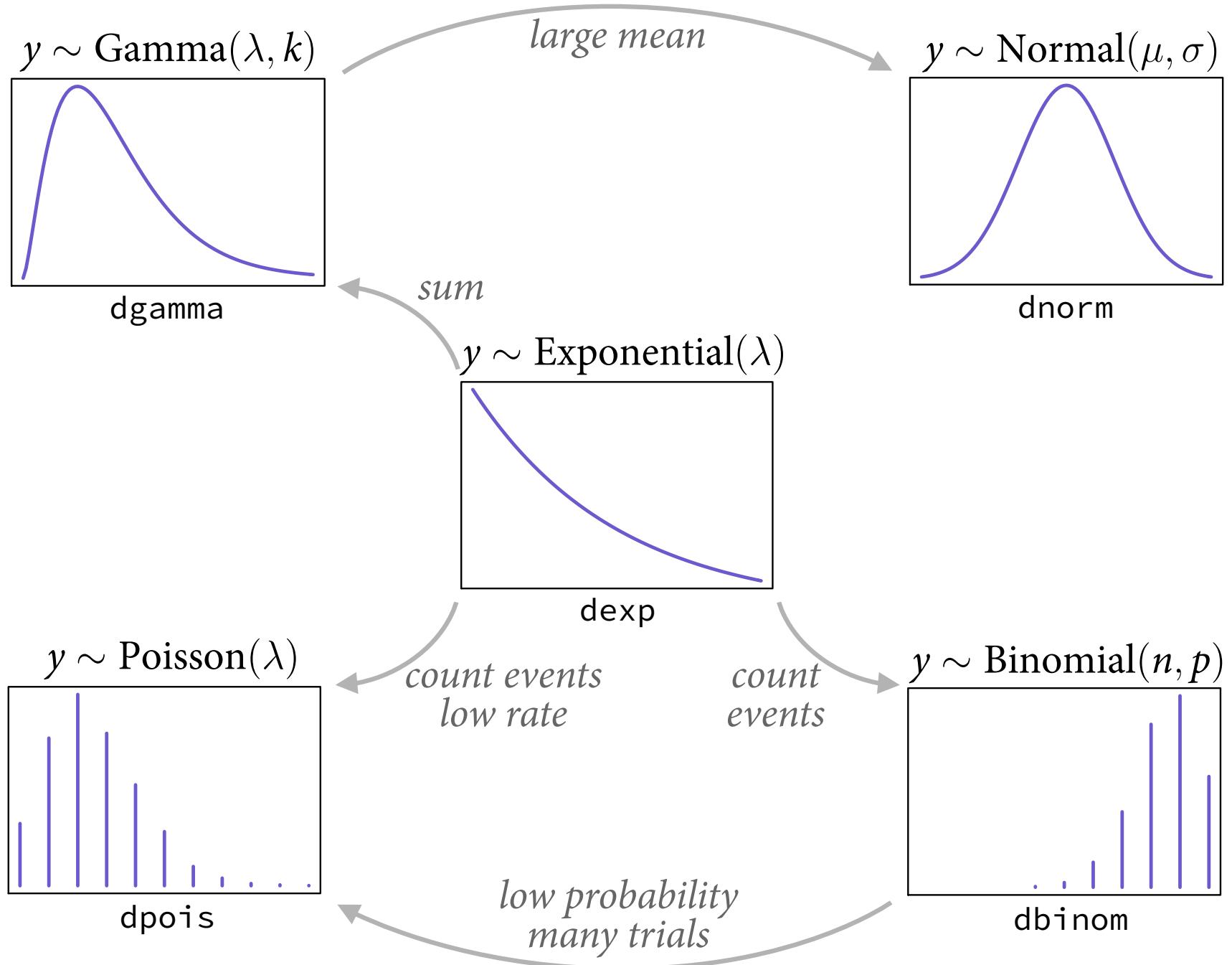
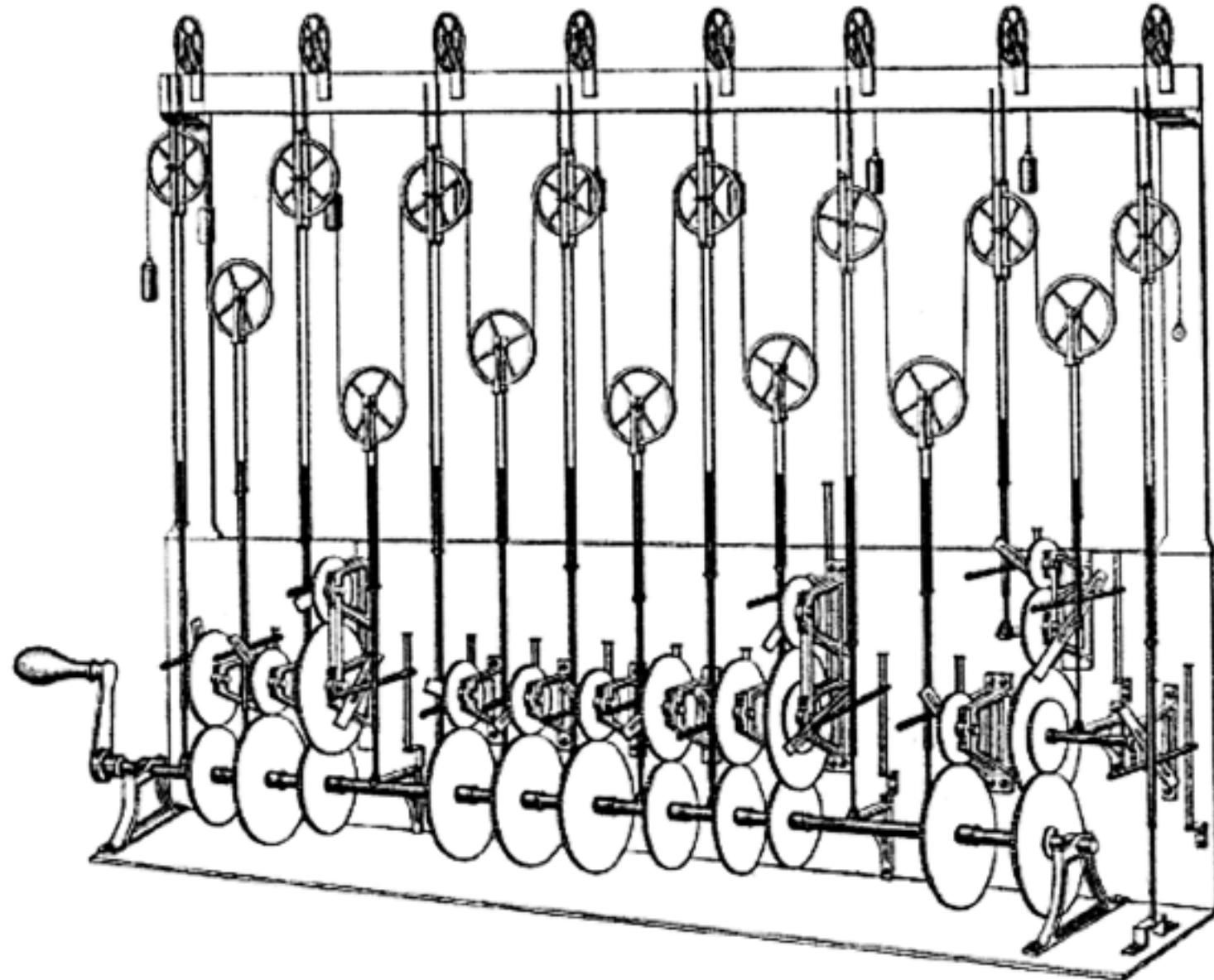


Figure 9.5



Tide prediction machine, 1879 design,
by William Thomson (Lord Kelvin, 1824–1907)

Generalized Linear Models

- (1) Pick an outcome distribution
 - Distances and durations: exponential, gamma (*survival* or *event history*)
 - Counts: Poisson, binomial, multinomial, geometric
 - Monsters: Ranks and ordered categories
 - Mixtures: Beta-binomial, gamma-Poisson, zero-inflated processes, occupancy models

Generalized Linear Models

- (2) Model parameters with a *link*

same units $\boxed{y_i \sim \text{Normal}(\mu_i, \sigma),}$
 $\boxed{\mu_i = \alpha + \beta x_i.}$

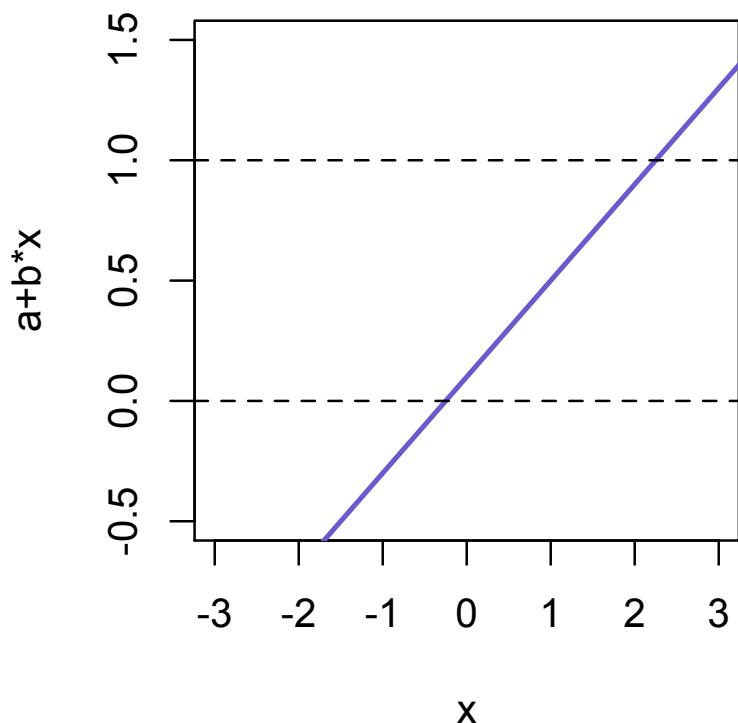
Generalized Linear Models

same units $y_i \sim \text{Normal}(\mu_i, \sigma),$

$$\mu_i = \alpha + \beta x_i.$$

count $y_i \sim \text{Binomial}(n_i, p_i)$

probability $p_i \stackrel{?}{=} \alpha + \beta x_i$



Generalized Linear Models

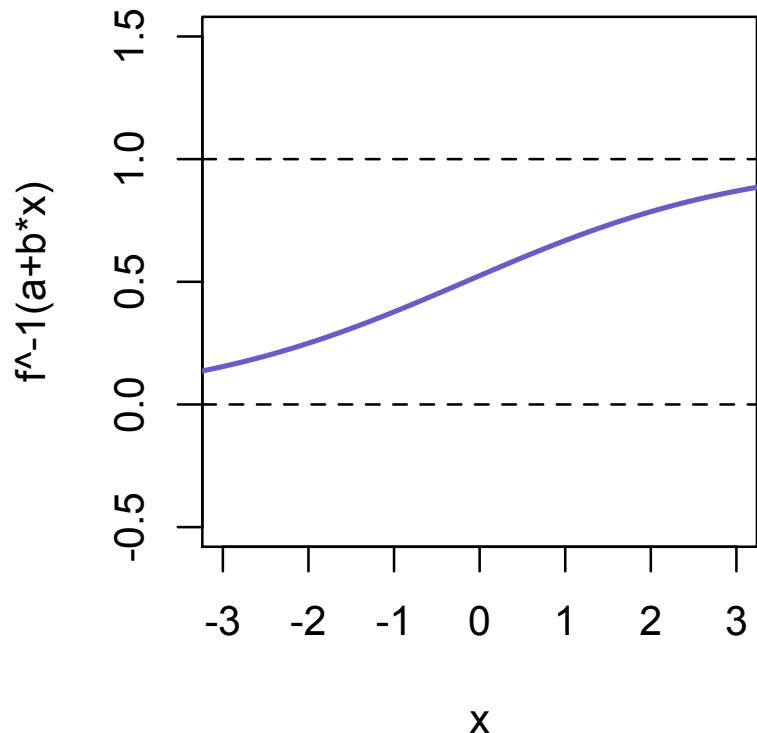
same units.....

$$\boxed{\begin{array}{l} y_i \sim \text{Normal}(\mu_i, \sigma), \\ \mu_i = \alpha + \beta x_i. \end{array}}$$

count.....

$$y_i \sim \text{Binomial}(n_i, p_i)$$
$$f(p_i) = \alpha + \beta x_i$$

link function

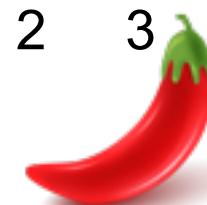
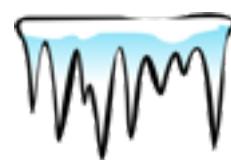
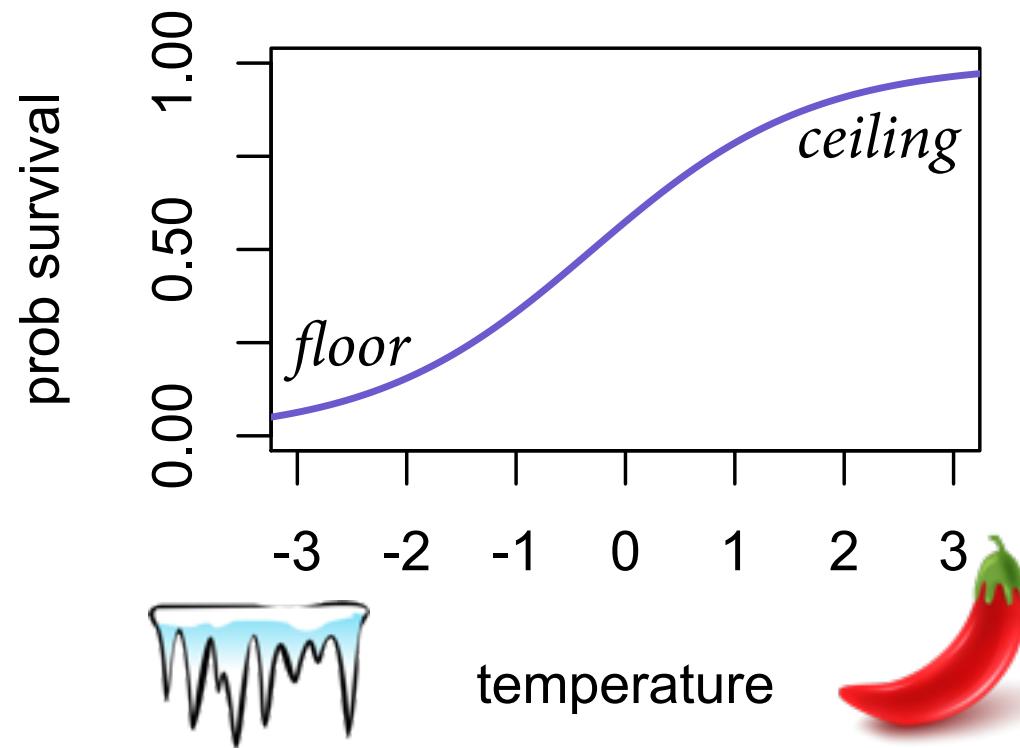


Generalized Linear Models

- (3) Compute posterior
 - Search is harder
 - Interpretation is harder
 - Links matter
 - Quadratic approximation often works, but not always
 - Safer to rely on MCMC

Everything interacts

- There are *floor* and *ceiling* effects



Everything interacts

- Linear regression:

$$\mu = \alpha + \beta x \quad \partial\mu/\partial x = \beta$$

- Logistic regression:

$$p = \frac{\exp(\alpha + \beta x)}{1 + \exp(\alpha + \beta x)}$$

$$\frac{\partial p}{\partial x} = \frac{\beta}{2(1 + \cosh(\alpha + \beta x))}$$

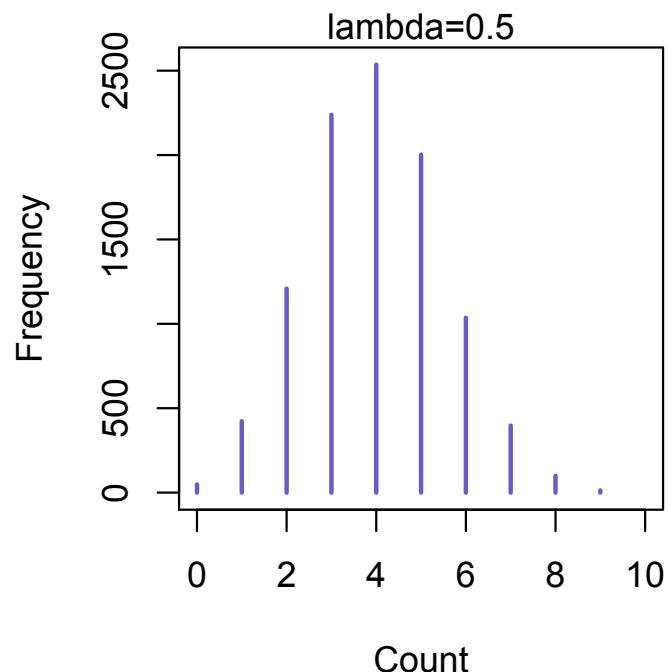
Binomial distribution

- Counts of a specific event out of n possibilities
- Constant expected value
- Maxent: Binomial

$$y \sim \text{Binomial}(n, p)$$

count *number of trials* *probability of success*

“successes”



Binomial distribution

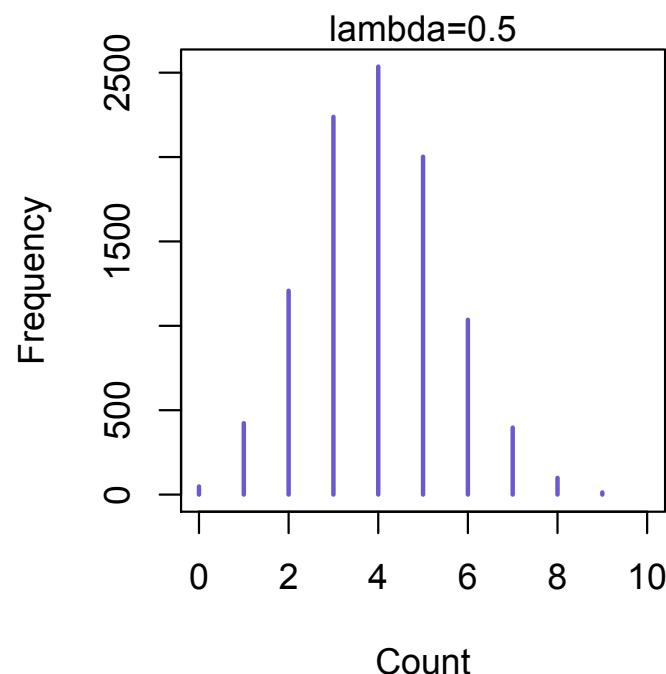
- Counts of a specific event out of n possibilities
- Constant expected value
- Maxent: Binomial

$$y \sim \text{Binomial}(n, p)$$

$$\text{E}(y) = np$$

$$\text{var}(y) = np(1 - p)$$

*Mean and variance
not independent*



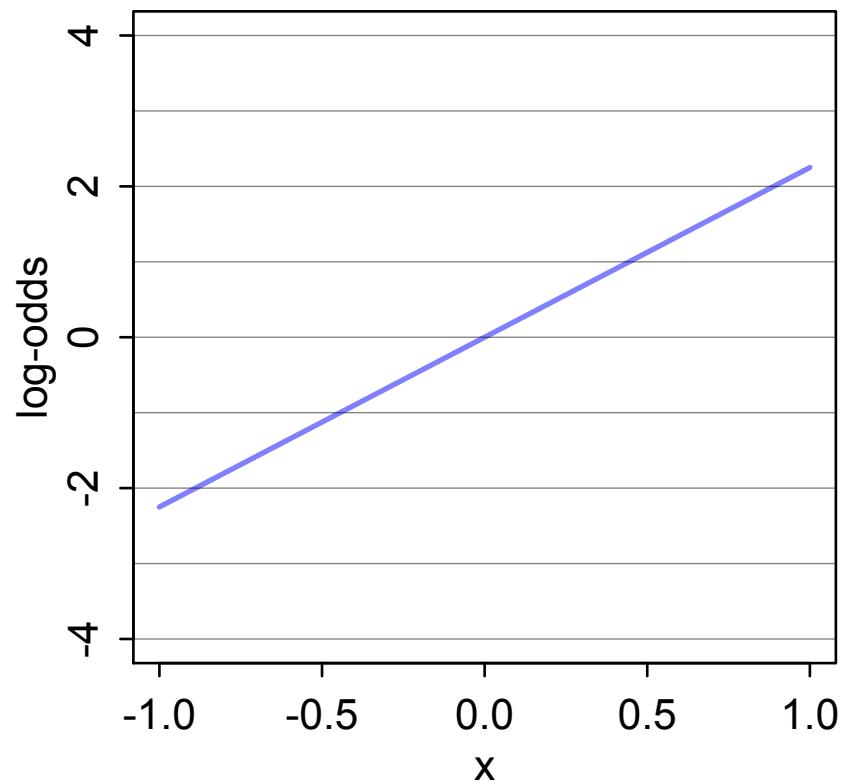
Need a link

$$y \sim \text{Binomial}(n, p)$$

- y and p on different scales
- y : count
- p : probability
- Want to model p as function of predictor variables
- Must bound it to $[0,1]$ interval

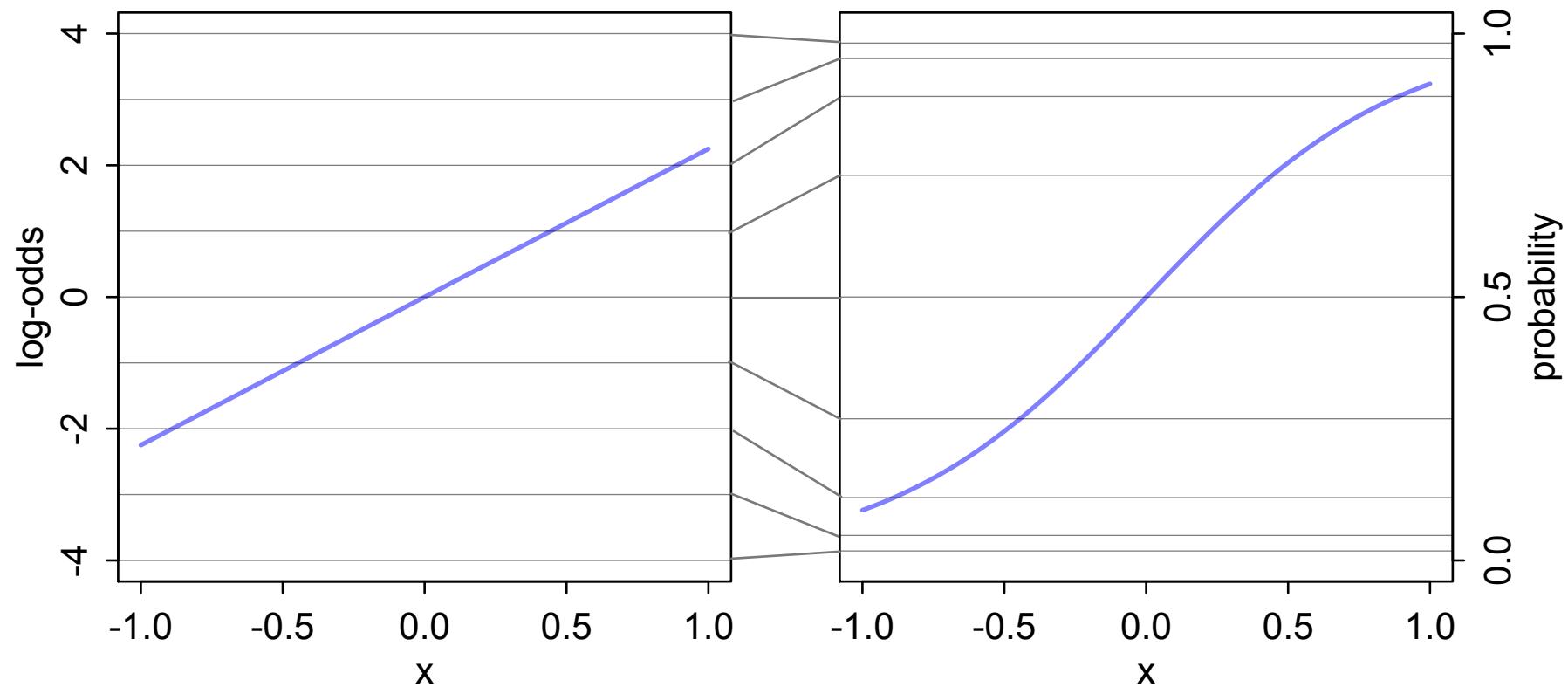
Logit link

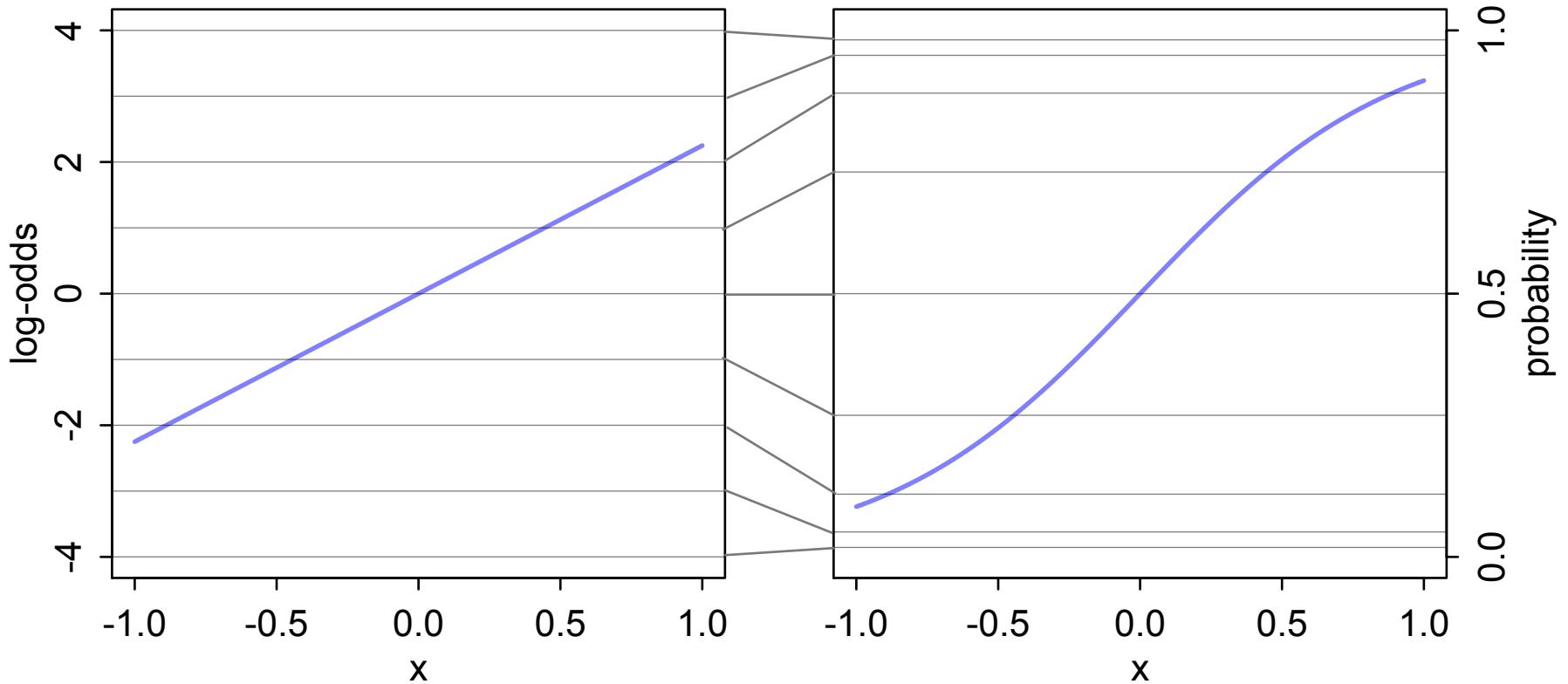
- Goal: map linear model to $[0,1]$



Logit link

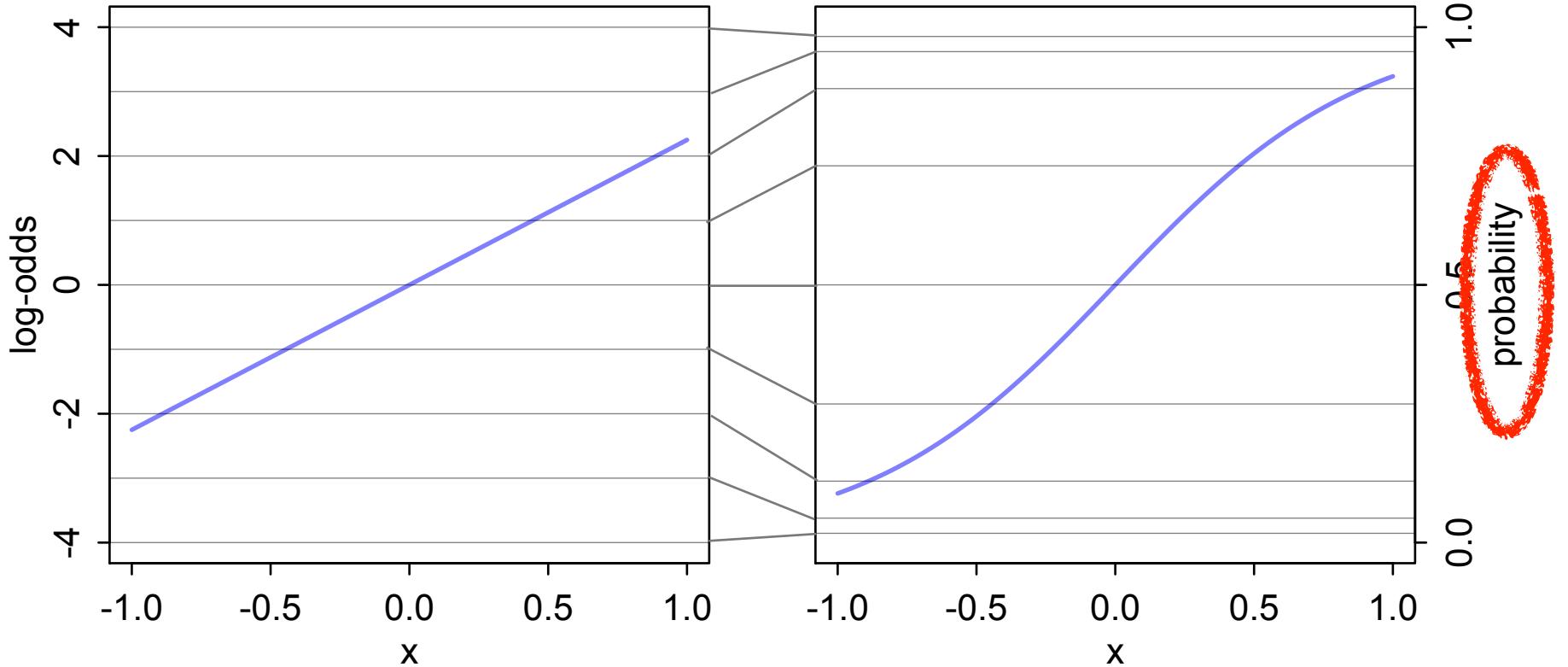
- Goal: map linear model to $[0,1]$



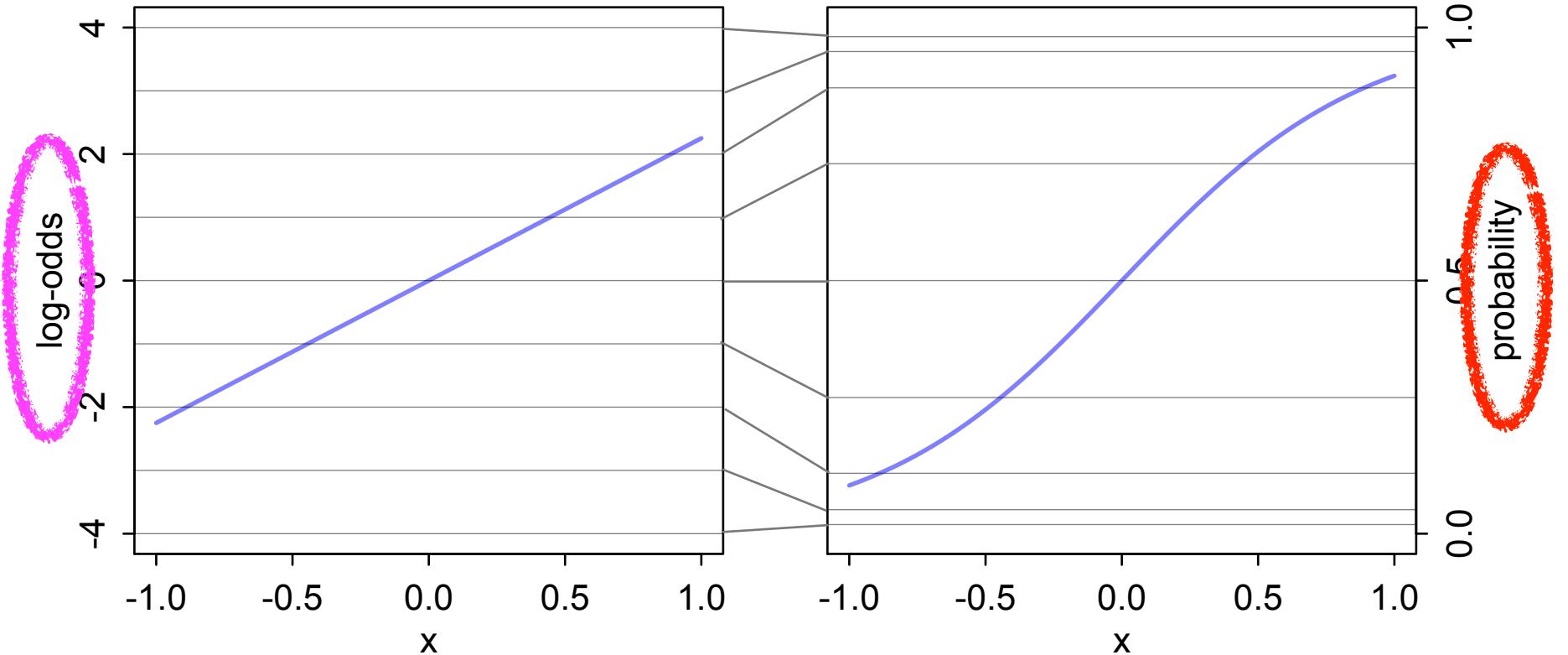


$$y_i \sim \text{Binomial}(n, p_i)$$

$$\text{logit}(p_i) = \alpha + \beta x_i$$



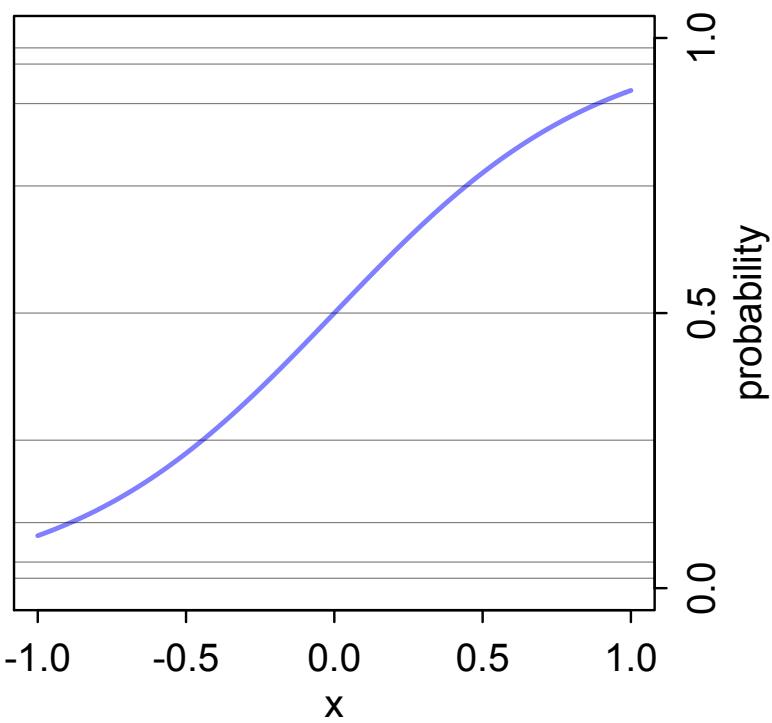
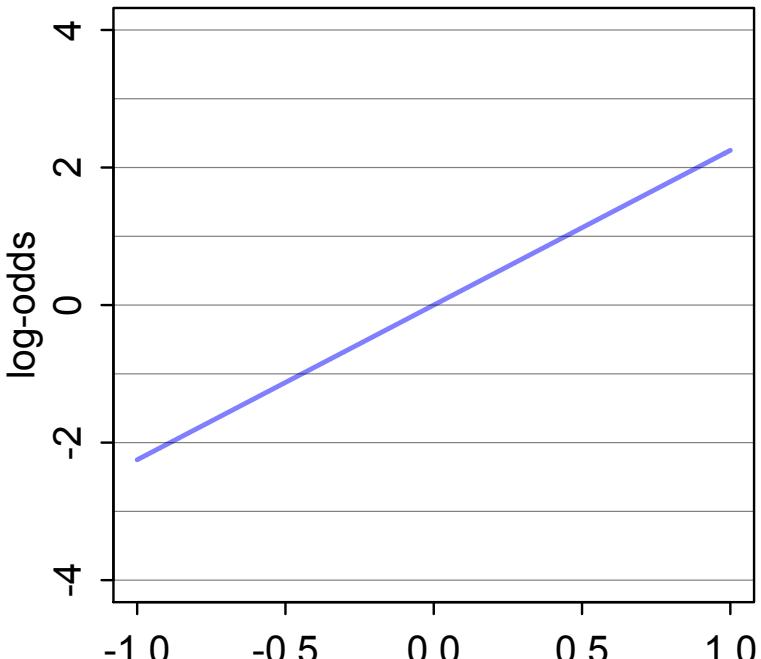
$$y_i \sim \text{Binomial}(n, p_i)$$
$$\text{logit } p_i = \alpha + \beta x_i$$



$$y_i \sim \text{Binomial}(n, p_i)$$
$$\text{logit } p_i = \alpha + \beta x_i$$

$$y_i \sim \text{Binomial}(n, p_i)$$

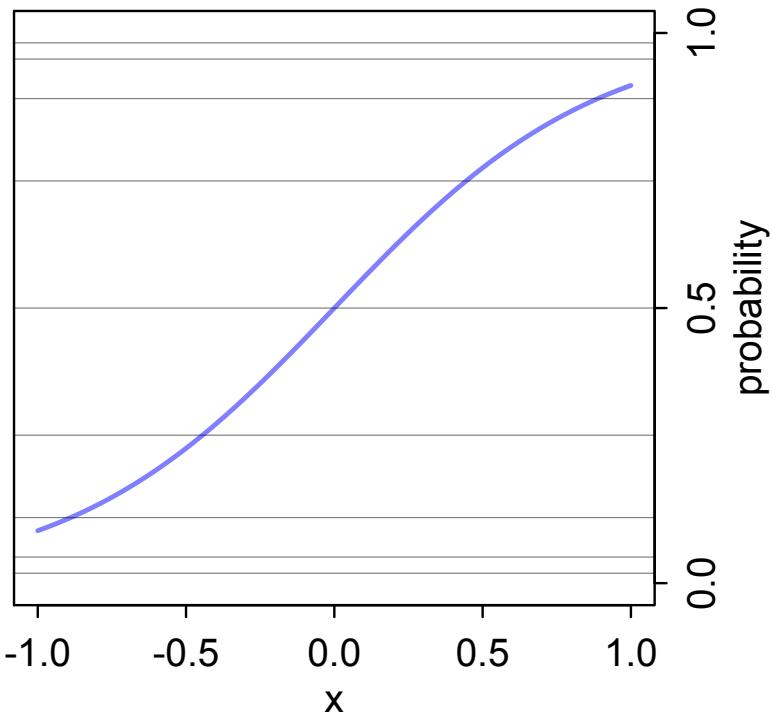
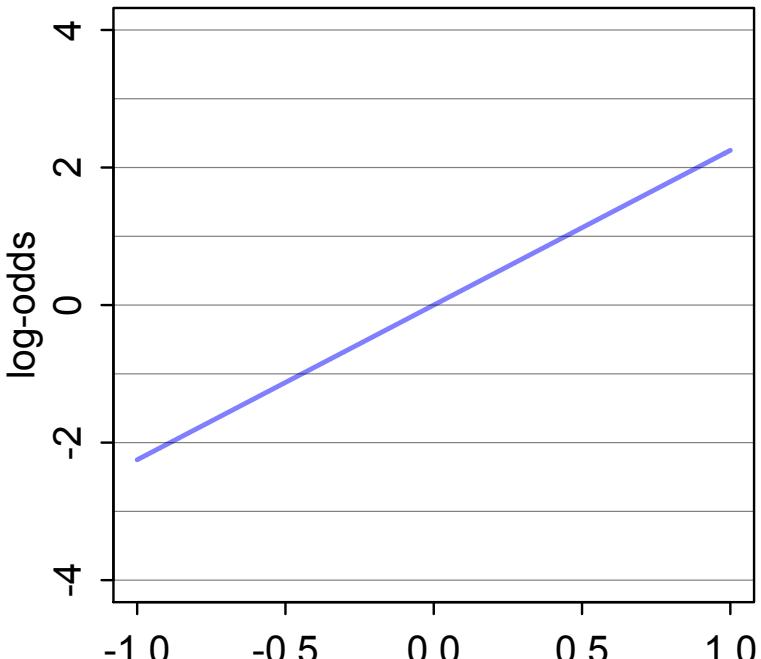
$$\text{logit}(p_i) = \alpha + \beta x_i$$



$$y_i \sim \text{Binomial}(n, p_i)$$

$$\text{logit}(p_i) = \alpha + \beta x_i$$

$$\text{logit}(p_i) = \log \frac{p_i}{1 - p_i} = \alpha + \beta x_i$$



$$y_i \sim \text{Binomial}(n, p_i)$$

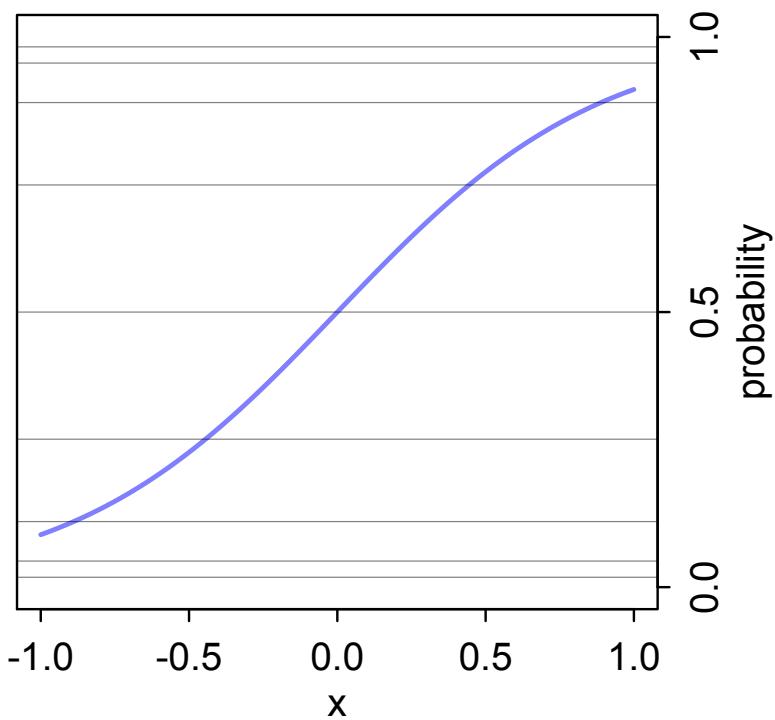
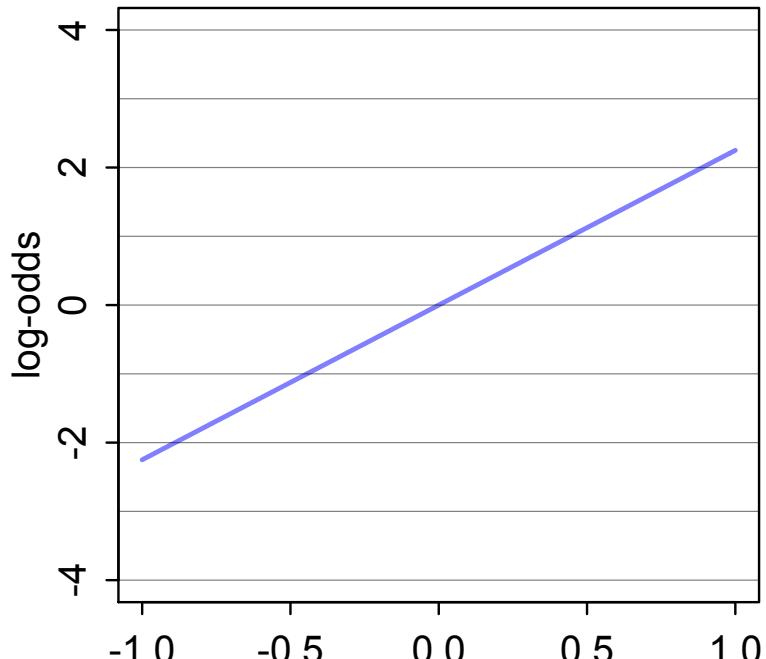
$$\text{logit}(p_i) = \alpha + \beta x_i$$

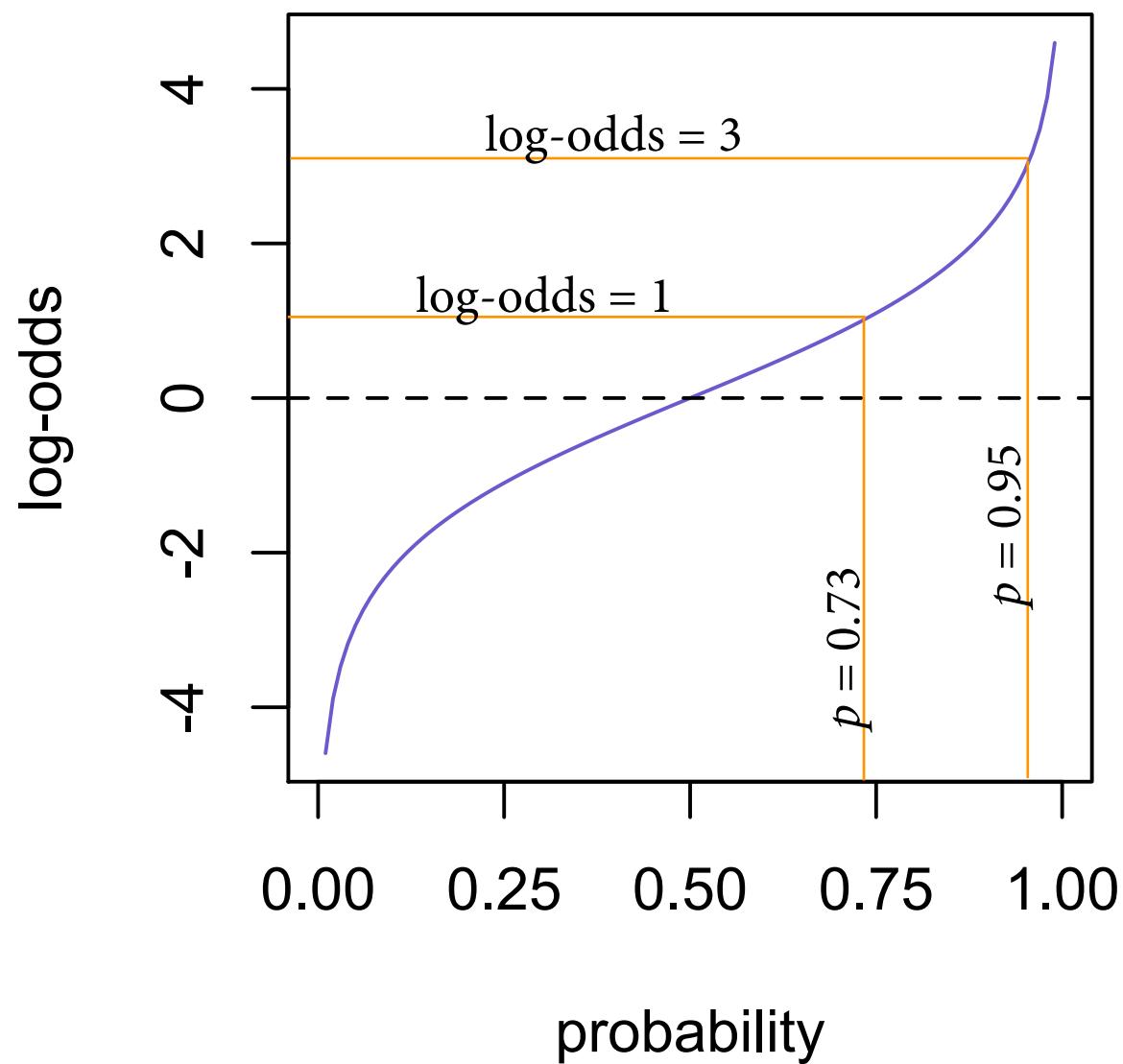
$$\text{logit}(p_i) = \log \frac{p_i}{1 - p_i} = \alpha + \beta x_i$$

Solve for p_i :

$$p_i = \frac{\exp(\alpha + \beta x_i)}{1 + \exp(\alpha + \beta x_i)}$$

inverse-link is logistic





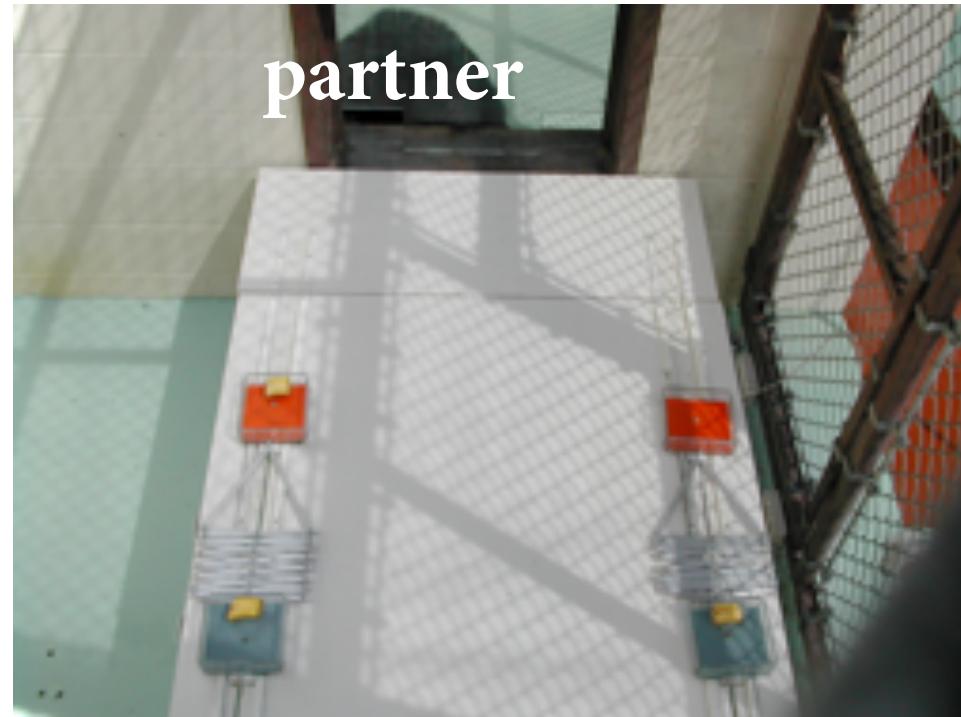
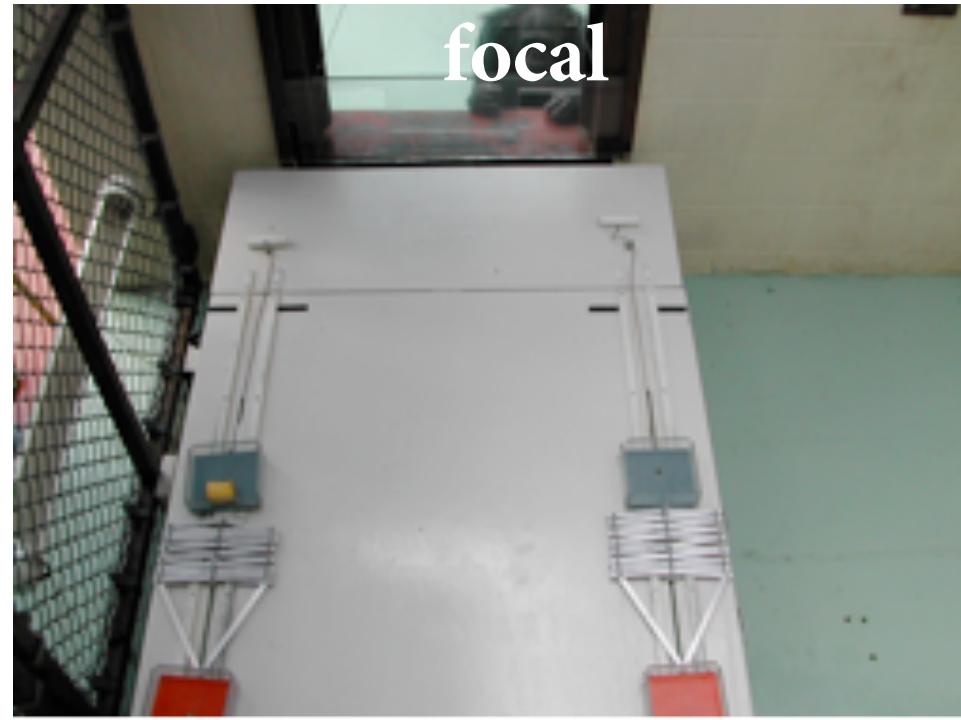
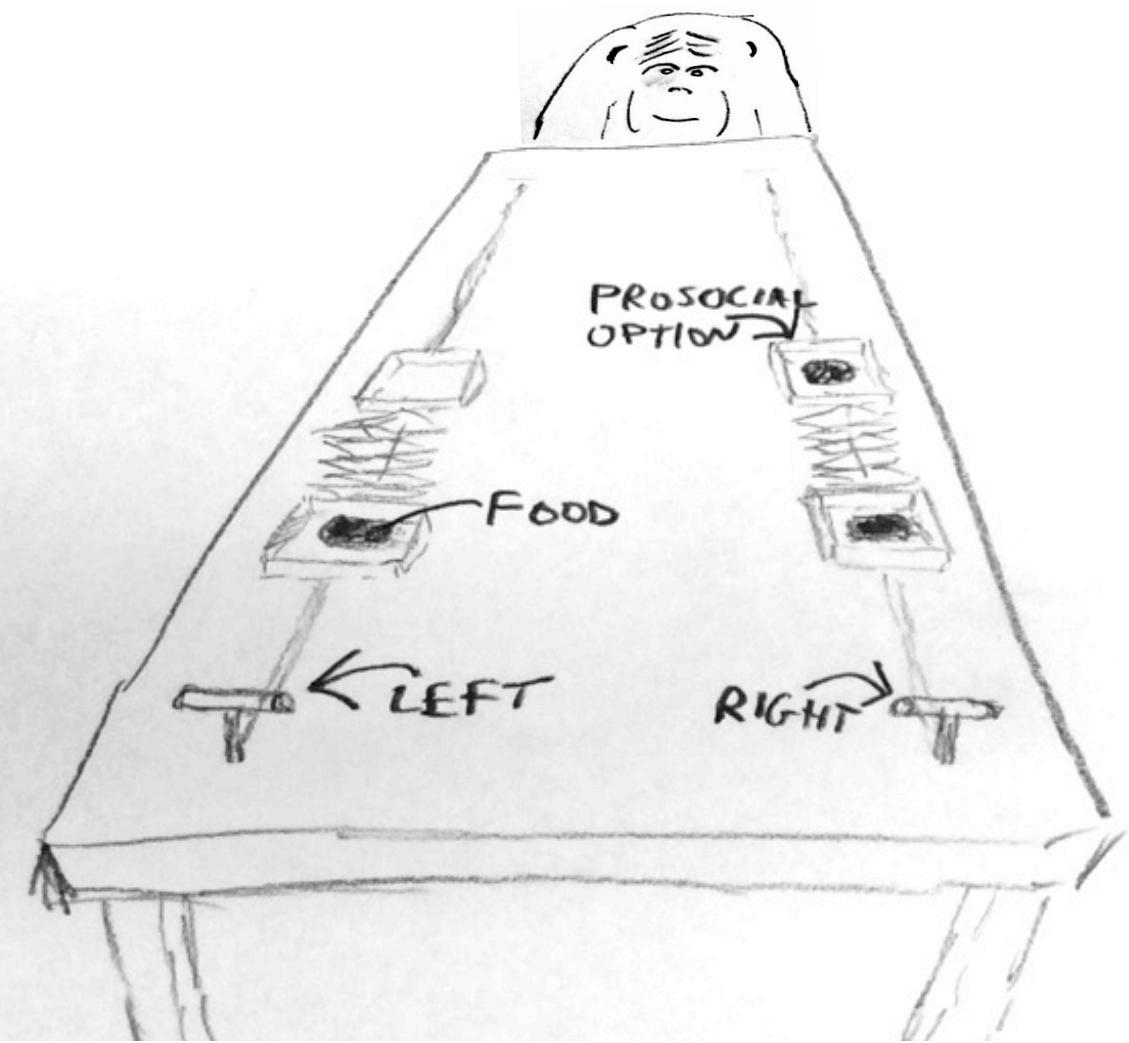
Logit link

$$y_i \sim \text{Binomial}(n, p_i)$$

$$\text{logit}(p_i) = \alpha + \beta x_i$$

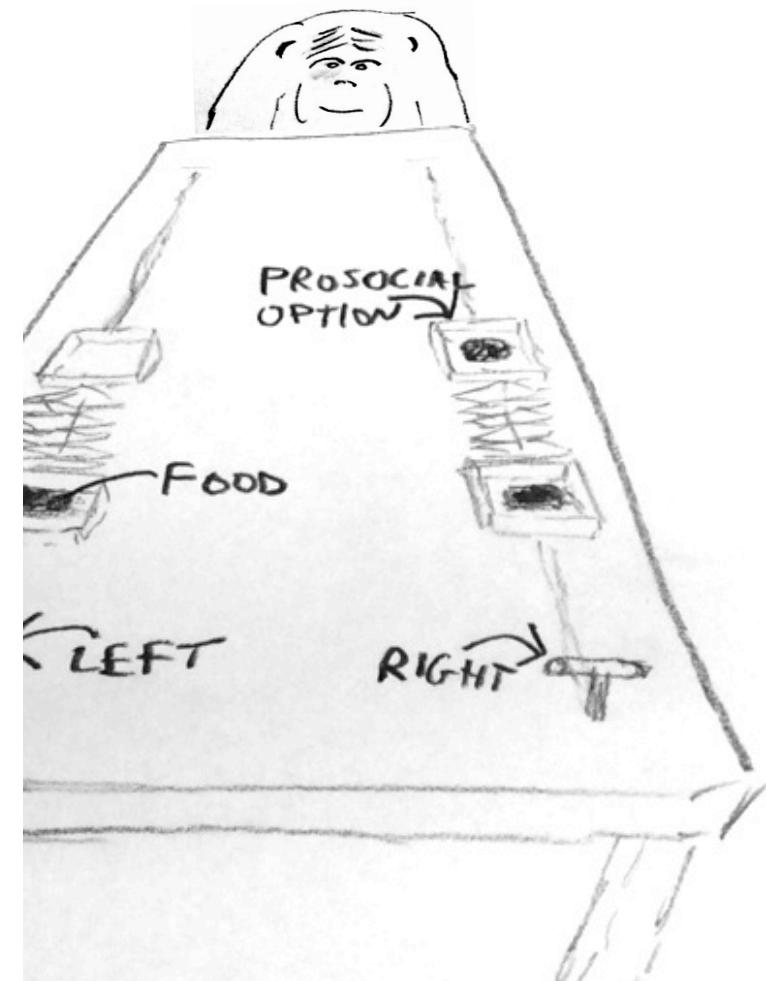
- Where does this thing come from?
- Several good answers:
 - “Natural” link inside probability formula
 - log-odds is fundamental parameter
 - See Overthinking box, pages 313–314
- Other links sometimes justified
 - Probit (common in economics)
 - Complementary-log-log (cloglog)
- If you have a real scientific model, link is automatic

Prosocial chimpanzees



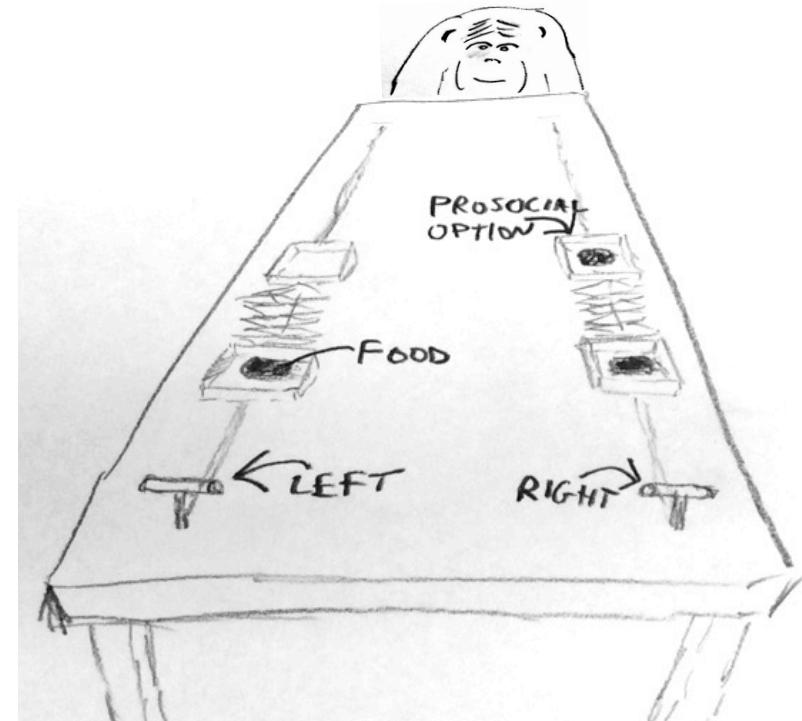
Prosocial chimpanzees

- Two *conditions*: (1) partner, (2) alone
- Two *options*: (1) prosocial, (2) asocial
- Two *outcomes*: (1) left lever, (2) right lever
- Want to predict *outcome* as function of *condition* and which side *option* is on
 - Do chimps prefer *left* lever when *partner* present and *prosocial* on *left*? => interaction!



Prosocial chimpanzees

- Coding treatments:
 - (1) right/no-partner
 - (2) left/no-partner
 - (3) right/partner
 - (4) left/partner



$$L_i \sim \text{Binomial}(1, p_i)$$

$$\text{logit}(p_i) = \alpha_{\text{ACTOR}[i]} + \beta_{\text{TREATMENT}[i]}$$

$\alpha_j \sim$ to be determined

$\beta_k \sim$ to be determined

$\text{Binomial}(1, p)$ often called
logistic regression

Same as $\text{Bernoulli}(p)$

Logit link priors

- Prior on logit scale not same shape as prior on probability scale
- Use prior simulation to understand

$$L_i \sim \text{Binomial}(1, p_i)$$

$$\text{logit}(p_i) = \alpha$$

$$\alpha \sim \text{Normal}(0, \omega)$$

Logit link priors

$$L_i \sim \text{Binomial}(1, p_i)$$

$$\text{logit}(p_i) = \alpha$$

$$\alpha \sim \text{Normal}(0, \omega)$$

```
m11.1 <- quap(  
  alist(  
    pulled_left ~ dbinom( 1 , p ) ,  
    logit(p) <- a ,  
    a ~ dnorm( 0 , 10 )  
  ) , data=d )
```

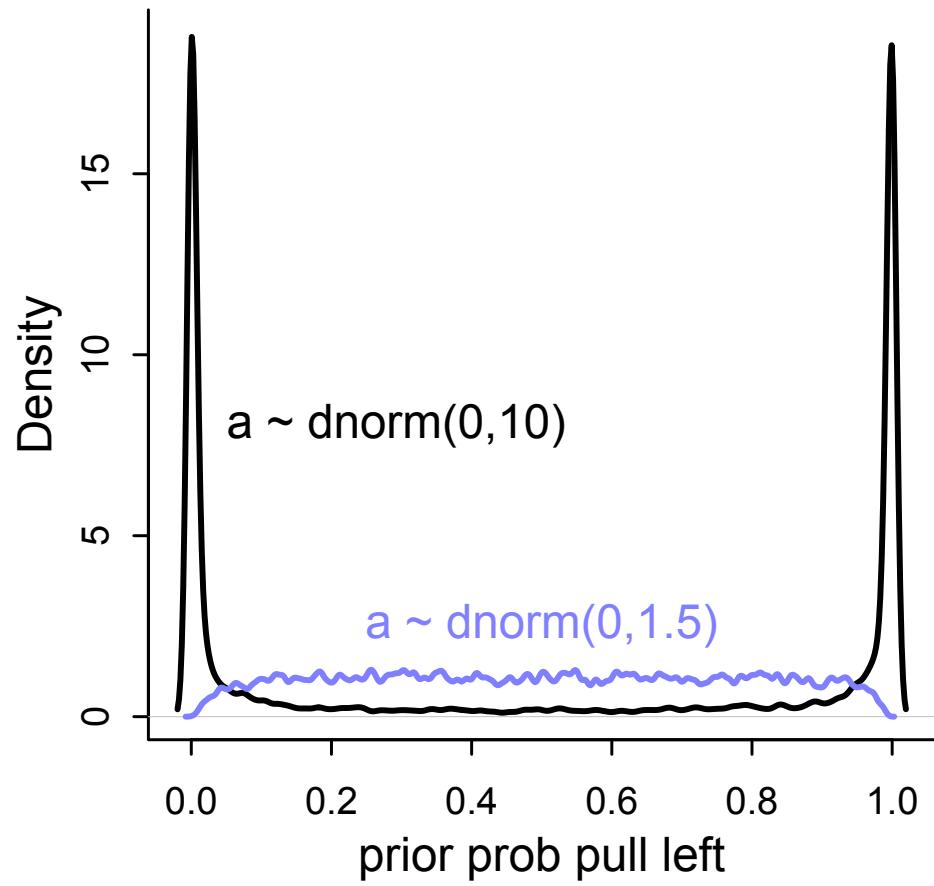


Figure 11.3

Logit link priors

- What about slopes?

R code
11.7

```
m11.2 <- quap(  
  alist(  
    pulled_left ~ dbinom( 1 , p ) ,  
    logit(p) <- a + b[treatment] ,  
    a ~ dnorm( 0 , 1.5 ) ,  
    b[treatment] ~ dnorm( 0 , 10 )  
  ) , data=d )
```

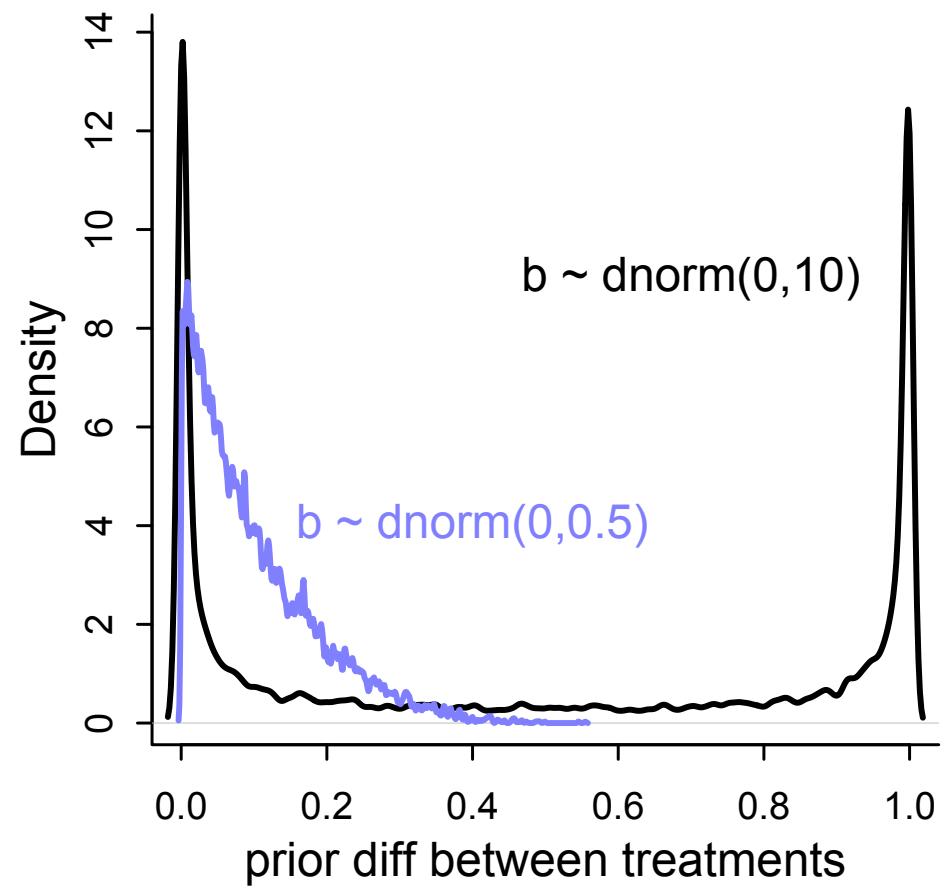
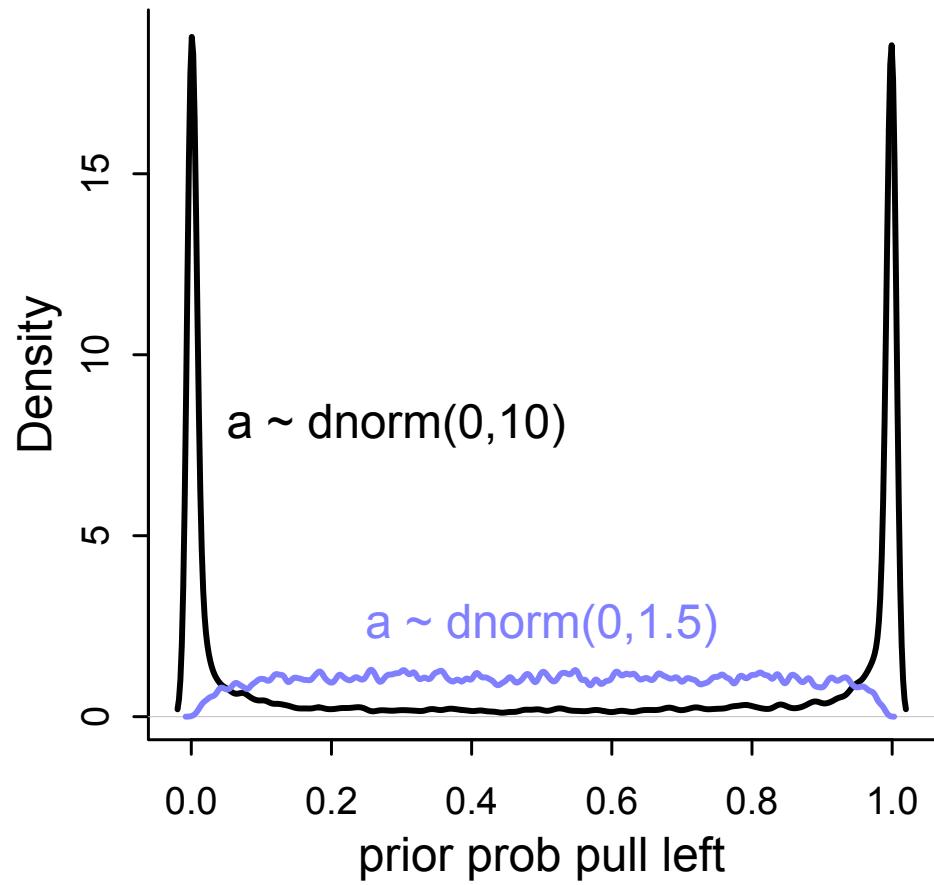


Figure 11.3

```

# particles in 11-dimensional space
m11.4 <- ulam(
  alist(
    pulled_left ~ dbinom( 1 , p ) ,
    logit(p) <- a[actor] + b[treatment] ,
    a[actor] ~ dnorm( 0 , 1.5 ) ,
    b[treatment] ~ dnorm( 0 , 0.5 )
  ) ,
  data=dat_list , chains=4 )
precis( m11.4 , depth=2 )

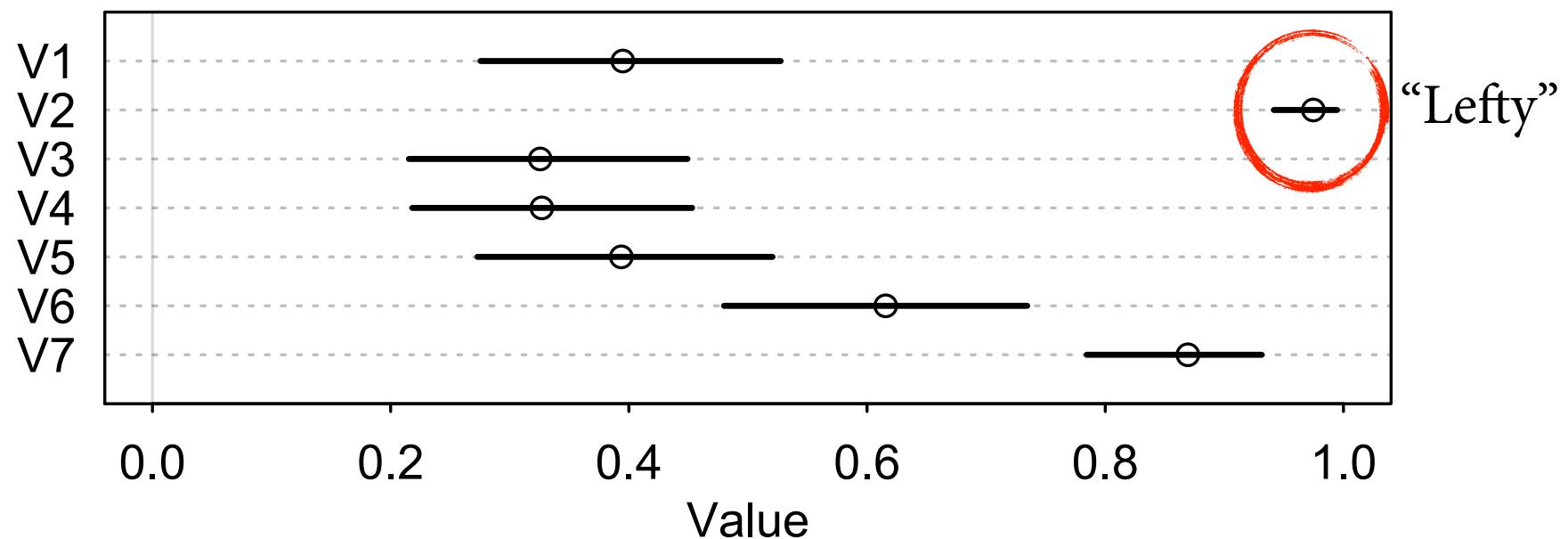
```

		mean	sd	5.5%	94.5%	n_eff	Rhat	
Chimpanzees	a[1]	-0.44	0.34	-0.97	0.11	736	1	
	a[2]	3.90	0.77	2.78	5.22	921	1	
	a[3]	-0.75	0.34	-1.29	-0.20	886	1	
	a[4]	-0.74	0.34	-1.28	-0.19	770	1	
	a[5]	-0.44	0.34	-0.98	0.08	832	1	
	a[6]	0.48	0.34	-0.08	1.02	854	1	
	a[7]	1.96	0.41	1.29	2.61	847	1	
Treatments	RN	b[1]	-0.05	0.29	-0.51	0.42	781	1
	LN	b[2]	0.48	0.29	0.03	0.94	657	1
	RP	b[3]	-0.39	0.28	-0.83	0.07	669	1
	LP	b[4]	0.36	0.29	-0.11	0.81	732	1

Individual differences

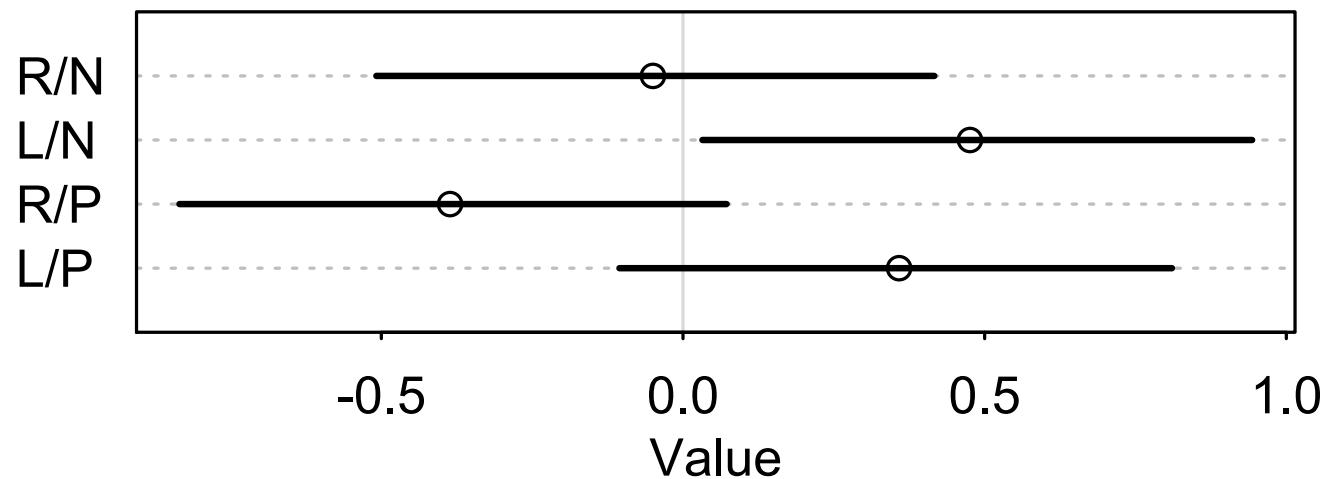
R code
11.11

```
post <- extract.samples(m11.4)
p_left <- inv_logit( post$a )
plot( precis( as.data.frame(p_left) ) , xlim=c(0,1) )
```



Treatments

```
labs <- c("R/N","L/N","R/P","L/P")
plot( precis( m11.4 , depth=2 , pars="b" ) , labels=labs )
```



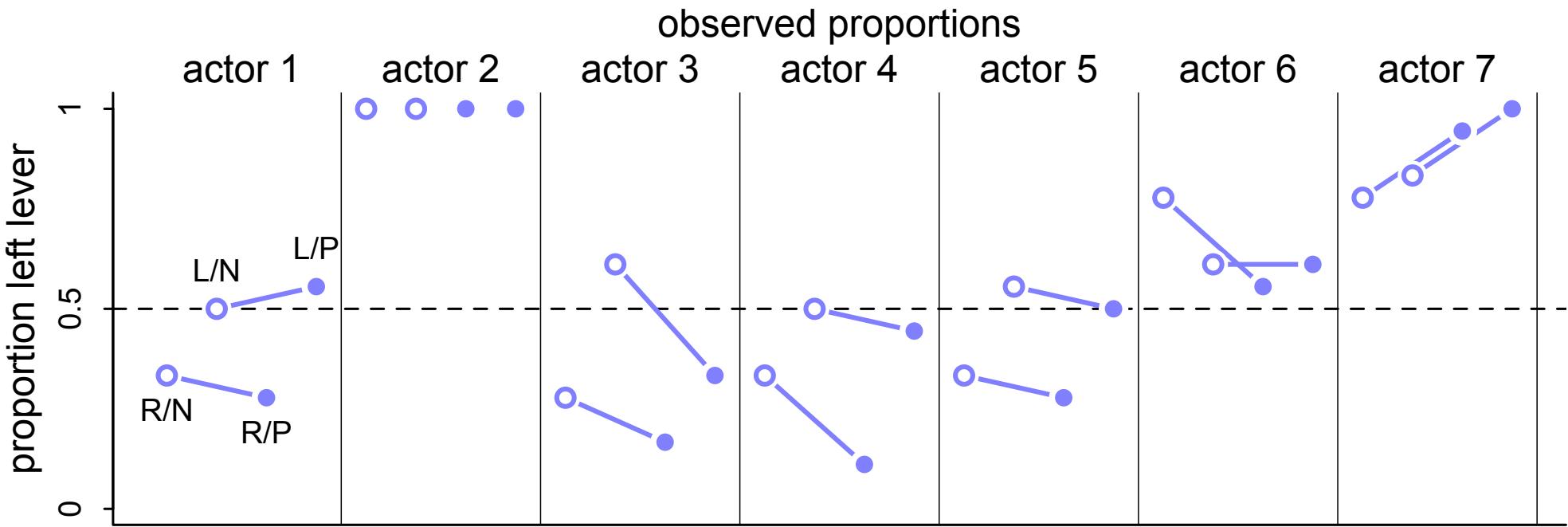


Figure 11.4

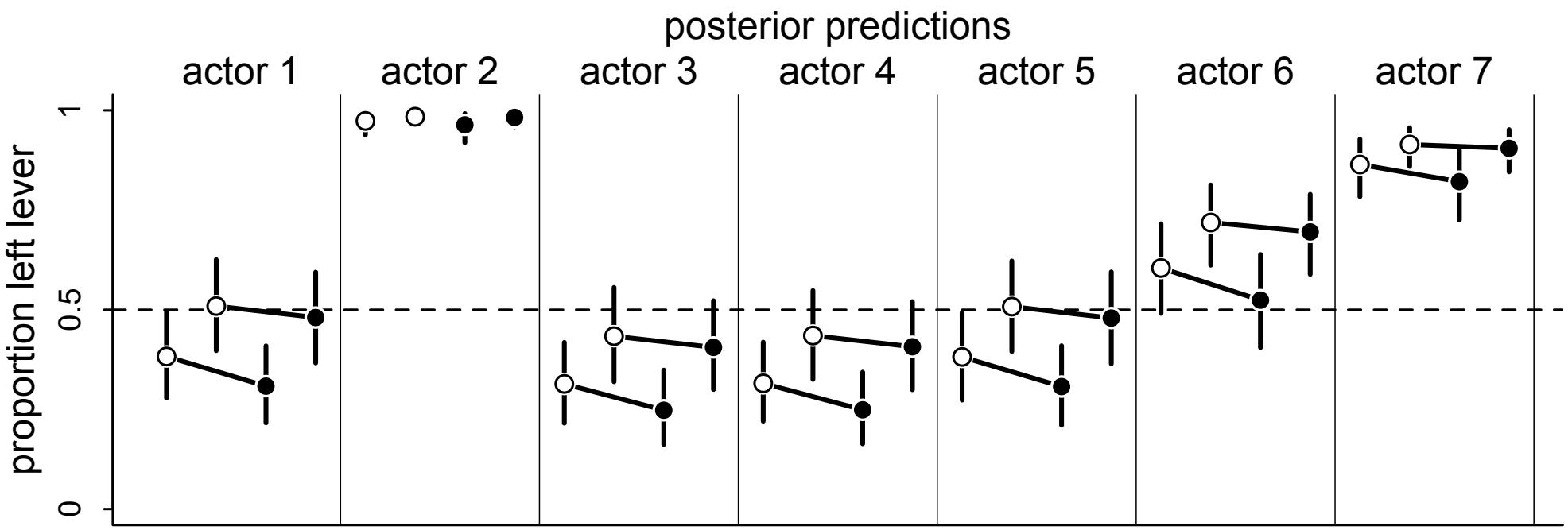
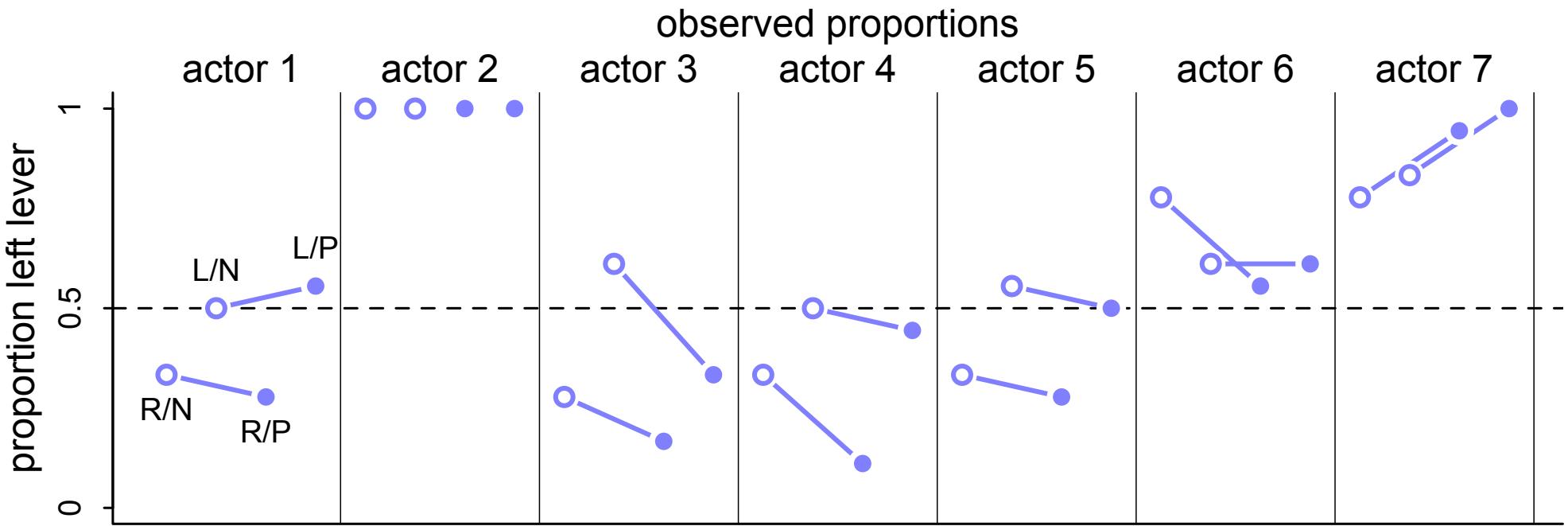


Figure 11.4

Comparing no-interaction

```
m11.5 <- ulam(  
  alist(  
    pulled_left ~ dbinom( 1 , p ) ,  
    logit(p) <- a[actor] + bs[side] + bc[cond] ,  
    a[actor] ~ dnorm( 0 , 1.5 ) ,  
    bs[side] ~ dnorm( 0 , 0.5 ) ,  
    bc[cond] ~ dnorm( 0 , 0.5 )  
  ) ,  
  data=dat_list2 , chains=4 , log_lik=TRUE )
```

R code
11.19

```
compare( m11.5 , m11.4 , func=L00 )
```

	L00	pL00	dL00	weight	SE	dSE
m11.5	531.2	7.9	0.0	0.66	19.17	NA
m11.4	532.6	8.7	1.4	0.34	19.01	1.28

Relative and absolute effects

- Parameters on *relative* effect scale
- Predictions on *absolute* effect scale
- Proportional odds: Relative effect measure

R code
11.22

```
post <- extract.samples(m11.4)
mean( exp(post$b[,4]-post$b[,2]) )
```

[1] 0.9206479

Relative and absolute effects

- Parameters on *relative* effect scale
- Predictions on *absolute* effect scale
- Using relative effects may exaggerate importance of predictor
 - Good for scaring people, getting published
 - Not so good for public health, scientific progress
 - But needed for causal inference



relative shark



absolute penguin



Deer kill
130 people
annually



Cows kill
22 people
annually



Jellyfish kill
40 people
annually



Sharks kill
5 people
annually



Ants kill
30 people
annually



Hippos kill
2,900 people
annually



Horses kill
20 people
annually

Risk communication

- Many people mistake relative risk for absolute risk
- Example:
 - 1/1000 women develop blood clots
 - 3/1000 women on birth control develop blood clots
 - => 200% increase in blood clots!
 - Change in probability is only 0.002
 - Pregnancy much more dangerous than blood clots



Deadly risk of pill used by 1m GP in Britain told to warn about popular contraceptive

- Bestselling brands of birth control tablets linked to
- They are believed to double the risk compared to older ones
- 'Third-generation' contraceptives caused 14 deaths
- UK doctors have been ordered to alert women to the