

MAXIMUM LIKELIHOOD ESTIMATION
IN THE GENERAL LINEAR MODEL.

Suppose $f(\underline{w}|\underline{\theta})$ is the probability density function (pdf) or probability mass function (pmf) of a random vector \underline{w} , where $\underline{\theta}$ is a $k \times 1$ parameter.

Given a value of the parameter vector $\underline{\theta}$, $f(\underline{w}|\underline{\theta})$ is a real-valued function of \underline{w} .

The likelihood function $L(\underline{\theta}|\underline{w}) = f(\underline{w}|\underline{\theta})$ is a real-valued function of $\underline{\theta}$ for a given value of \underline{w} .

For any potential observed value \underline{w} ,
define $\hat{\underline{\theta}}(\underline{w})$ to be a parameter value
at which $L(\underline{\theta} | \underline{w})$ attains its maximum
value. A maximum likelihood estimator
(MLE) of $\underline{\theta}$ is $\hat{\underline{\theta}}(\underline{w}) = \hat{\underline{\theta}}$.

Invariance Property of MLEs:

The MLE of a function of $\underline{\theta}$, say $\underline{g}(\underline{\theta})$, is
the function evaluated at the MLE of $\underline{\theta}$:

$$\widehat{\underline{g}(\underline{\theta})} = \underline{g}(\hat{\underline{\theta}})$$

$$\text{Let } \ell(\underline{\theta} | \underline{w}) = \log L(\underline{\theta} | \underline{w}).$$

If $\ell(\underline{\theta} | \underline{w})$ is differentiable, candidates for the MLE of $\underline{\theta}$ can be found by

Equating the score function

$$\frac{\partial \ell(\underline{\theta} | \underline{w})}{\partial \underline{\theta}} = \begin{bmatrix} \frac{\partial \ell(\underline{\theta} | \underline{w})}{\partial \theta_1} \\ \vdots \\ \frac{\partial \ell(\underline{\theta} | \underline{w})}{\partial \theta_k} \end{bmatrix} \quad \text{to } \underline{0}$$

and solving for $\underline{\theta}$.

The likelihood equations are

$$\frac{\partial \ell(\underline{\theta} | \underline{w})}{\partial \underline{\theta}} = \underline{0} \iff \frac{\partial \ell(\underline{\theta} | \underline{w})}{\partial \theta_j} = 0 \quad \forall j=1, \dots, k.$$

One strategy for obtaining an MLE is to find values of $\underline{\theta}$ that satisfy the likelihood equations and verify that at least one such value maximizes $\ell(\underline{\theta} | \underline{w})$.

Example: Normal Theory Gauss-Markov Linear Model

$$\underset{n \times 1}{Y} = \underset{n \times p}{X} \underset{p \times 1}{\beta} + \underset{n \times 1}{\varepsilon}, \quad \underset{n \times 1}{\varepsilon} \sim N(\underline{0}, \sigma^2 I), \quad \underset{(p+1) \times 1}{\underline{\theta}} = \begin{bmatrix} \underline{\beta} \\ \sigma^2 \end{bmatrix}$$

$$f(Y|\underline{\theta}) = \frac{\exp \left\{ -\frac{1}{2} (Y - X\beta)' (\sigma^2 I)^{-1} (Y - X\beta) \right\}}{(2\pi)^{n/2} |\sigma^2 I|^{1/2}}$$

$$= \frac{1}{(2\pi\sigma^2)^{n/2}} \exp \left\{ -\frac{1}{2\sigma^2} (Y - X\beta)' (Y - X\beta) \right\}$$

$$\ell(\underline{\theta}|Y) = -\frac{n}{2} \log 2\pi\sigma^2 - \frac{1}{2\sigma^2} (Y - X\beta)' (Y - X\beta)$$

The score function is

$$\frac{\partial l(\underline{\theta} | \underline{y})}{\partial \underline{\theta}} = \begin{bmatrix} \frac{\partial l(\underline{\theta} | \underline{y})}{\partial \underline{\beta}} \\ \frac{\partial l(\underline{\theta} | \underline{y})}{\partial \sigma^2} \end{bmatrix} = \begin{bmatrix} \frac{1}{\sigma^2} (X' \underline{y} - X' X \underline{\beta}) \\ \frac{(\underline{y} - X \underline{\beta})' (\underline{y} - X \underline{\beta})}{2\sigma^4} - \frac{n}{2\sigma^2} \end{bmatrix}$$

The likelihood equations are

$$\frac{\partial l(\underline{\theta} | \underline{y})}{\partial \underline{\theta}} = \underline{0} \iff \begin{aligned} X' X \underline{\beta} &= X' \underline{y} \\ \sigma^2 &= (\underline{y} - X \underline{\beta})' (\underline{y} - X \underline{\beta}) / n \end{aligned}$$

Thus, a solution to the likelihood equations

is $\begin{bmatrix} \hat{\beta} \\ (Y - X\hat{\beta})'(Y - X\hat{\beta})/n \end{bmatrix}$, where $\hat{\beta}$ is any

solution to the normal equations.

We already know that $(Y - X\beta)'(Y - X\beta)$ is minimized over $\beta \in \mathbb{R}^p$ by any solution to the normal equations.

Thus, $\forall \sigma^2 > 0$, $l\left(\begin{bmatrix} \hat{\beta} \\ \hat{\sigma}^2 \end{bmatrix} \mid Y\right) \geq l\left(\begin{bmatrix} \beta \\ \sigma^2 \end{bmatrix} \mid Y\right) \quad \forall \beta \in \mathbb{R}^p$

Examining the 2ND derivative of $l(\underline{\theta} \mid Y)$ with respect to σ^2 shows that

$\begin{bmatrix} \hat{\beta} \\ (\underline{Y} - X\hat{\beta})'(\underline{Y} - X\hat{\beta})/n \end{bmatrix}$ is an MLE of $\underline{\theta} = \begin{bmatrix} \beta \\ \sigma^2 \end{bmatrix}$.

Thus, if $C\beta$ is estimable, the MLE of $C\beta$ is $C\hat{\beta}$ (by the Invariance Property of MLEs), which is the BLUE of $C\beta$.

Note that the MLE of σ^2 is not the unbiased estimator we have been using.

$$\begin{aligned} E \left[\frac{(\mathbf{y} - \mathbf{X}\hat{\beta})'(\mathbf{y} - \mathbf{X}\hat{\beta})}{n} \right] &= E \left(\frac{SSE}{n} \right) \\ &= E \left[\frac{(n-r)MSE}{n} \right] = \frac{n-r}{n} \sigma^2 < \sigma^2 \end{aligned}$$

Thus, the MLE underestimates σ^2 on average.

Now consider the general linear model

$$\underline{y} = X\underline{\beta} + \underline{\varepsilon}, \quad \underline{\varepsilon} \sim N(\underline{0}, \Sigma), \quad \text{where}$$

Σ is a positive definite covariance matrix whose entries depend on unknown parameters in some vector $\underline{\gamma}$.

For example, $\Sigma = \begin{bmatrix} \sigma_e^2 + \sigma_u^2 & \sigma_u^2 & 0 \\ \sigma_u^2 & \sigma_e^2 + \sigma_u^2 & 0 \\ 0 & 0 & \sigma_e^2 + \sigma_u^2 \end{bmatrix}, \quad \underline{\gamma} = \begin{bmatrix} \sigma_e^2 \\ \sigma_u^2 \end{bmatrix}$

Our linear mixed effects model
is a special case where

$$\underline{\tilde{y}} = \underline{Z}\underline{u} + \underline{e} \quad \text{and} \quad \Sigma = \underline{Z}\underline{G}\underline{Z}' + \underline{R}.$$

In general, $\underline{\theta} = \begin{bmatrix} \underline{\beta} \\ \underline{\gamma} \end{bmatrix}$ and

$$f(\underline{y} | \underline{\theta}) = \frac{\exp\left\{-\frac{1}{2}(\underline{y} - \underline{X}\underline{\beta})' \Sigma^{-1}(\underline{y} - \underline{X}\underline{\beta})\right\}}{(2\pi)^{n/2} |\Sigma|^{1/2}}$$

$$\begin{aligned} \ell(\underline{\theta} | \underline{y}) = & -\frac{1}{2} \log |\Sigma| - \frac{1}{2} (\underline{y} - \underline{X}\underline{\beta})' \Sigma^{-1} (\underline{y} - \underline{X}\underline{\beta}) \\ & - \frac{n}{2} \log(2\pi) \end{aligned}$$

We know that for any positive definite covariance matrix Σ , $(y - X\beta)' \Sigma^{-1} (y - X\beta)$ is minimized over $\beta \in \mathbb{R}^p$ by

$$\hat{\beta}_{\Sigma} = (X' \Sigma^{-1} X)^{-1} X' \Sigma^{-1} y:$$

Thus, for any \underline{y} such that Σ is a positive definite covariance matrix,

$$\ell\left(\begin{bmatrix} \hat{\beta}_{\Sigma} \\ \underline{y} \end{bmatrix} \mid \underline{y}\right) \geq \ell\left(\begin{bmatrix} \beta \\ \underline{y} \end{bmatrix} \mid \underline{y}\right) \quad \forall \beta \in \mathbb{R}^p.$$

We define the profile log likelihood for $\underline{\gamma}$

to be $l^*(\underline{\gamma} | \underline{y}) = l\left(\begin{bmatrix} \hat{\beta}_{\underline{\gamma}} \\ \underline{\gamma} \end{bmatrix} | \underline{y}\right)$.

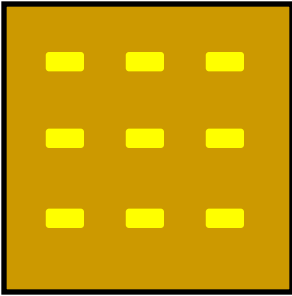
The MLE of $\underline{\theta}$ is $\hat{\underline{\theta}} = \begin{bmatrix} \hat{\beta}_{\hat{\underline{\gamma}}} \\ \hat{\underline{\gamma}} \end{bmatrix}$,

where $\hat{\underline{\gamma}}$ is a maximizer of $l^*(\underline{\gamma} | \underline{y})$

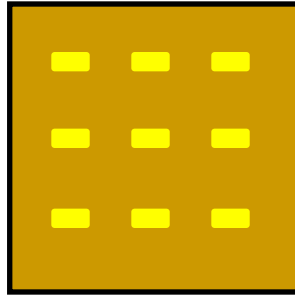
and $\hat{\Sigma}$ is Σ with $\hat{\underline{\gamma}}$ in place of $\underline{\gamma}$.

In general, numerical methods are required to find $\hat{\beta}$, a maximizer of $l^*(\beta | y)$.

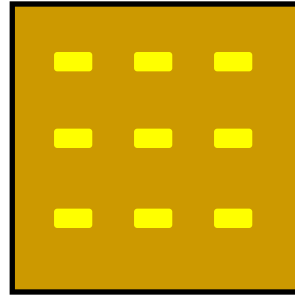
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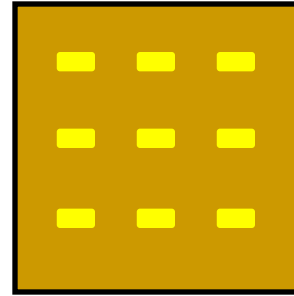
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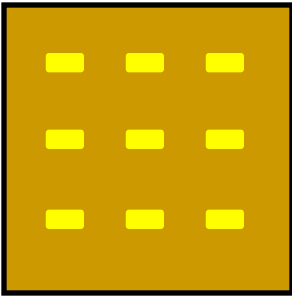
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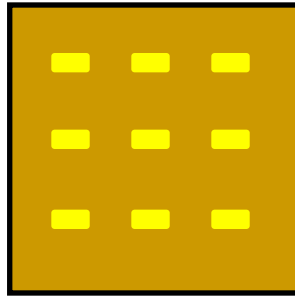
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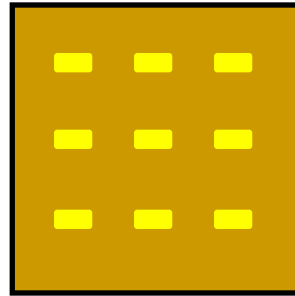
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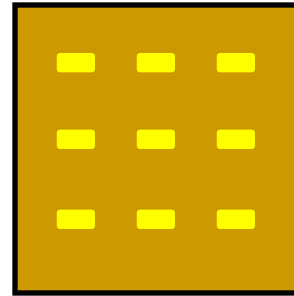
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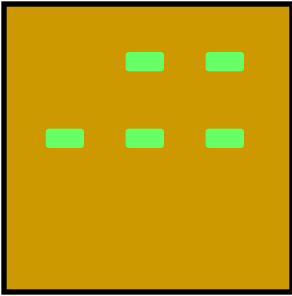
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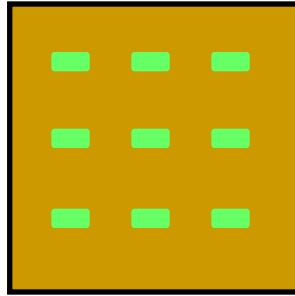
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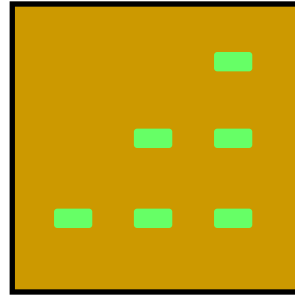
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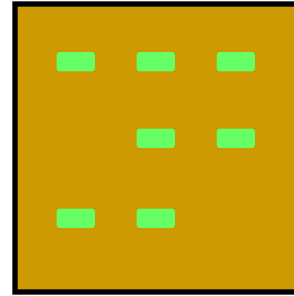
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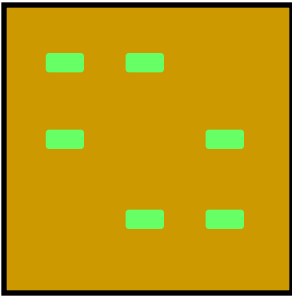
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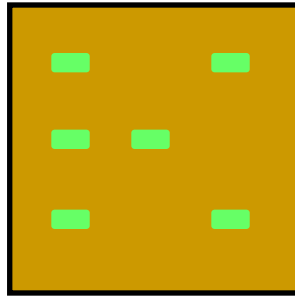
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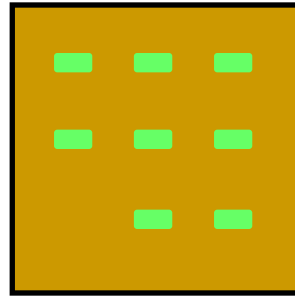
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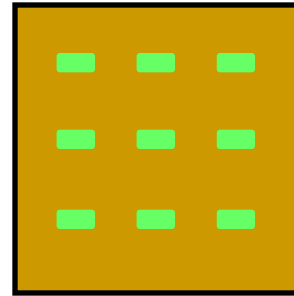
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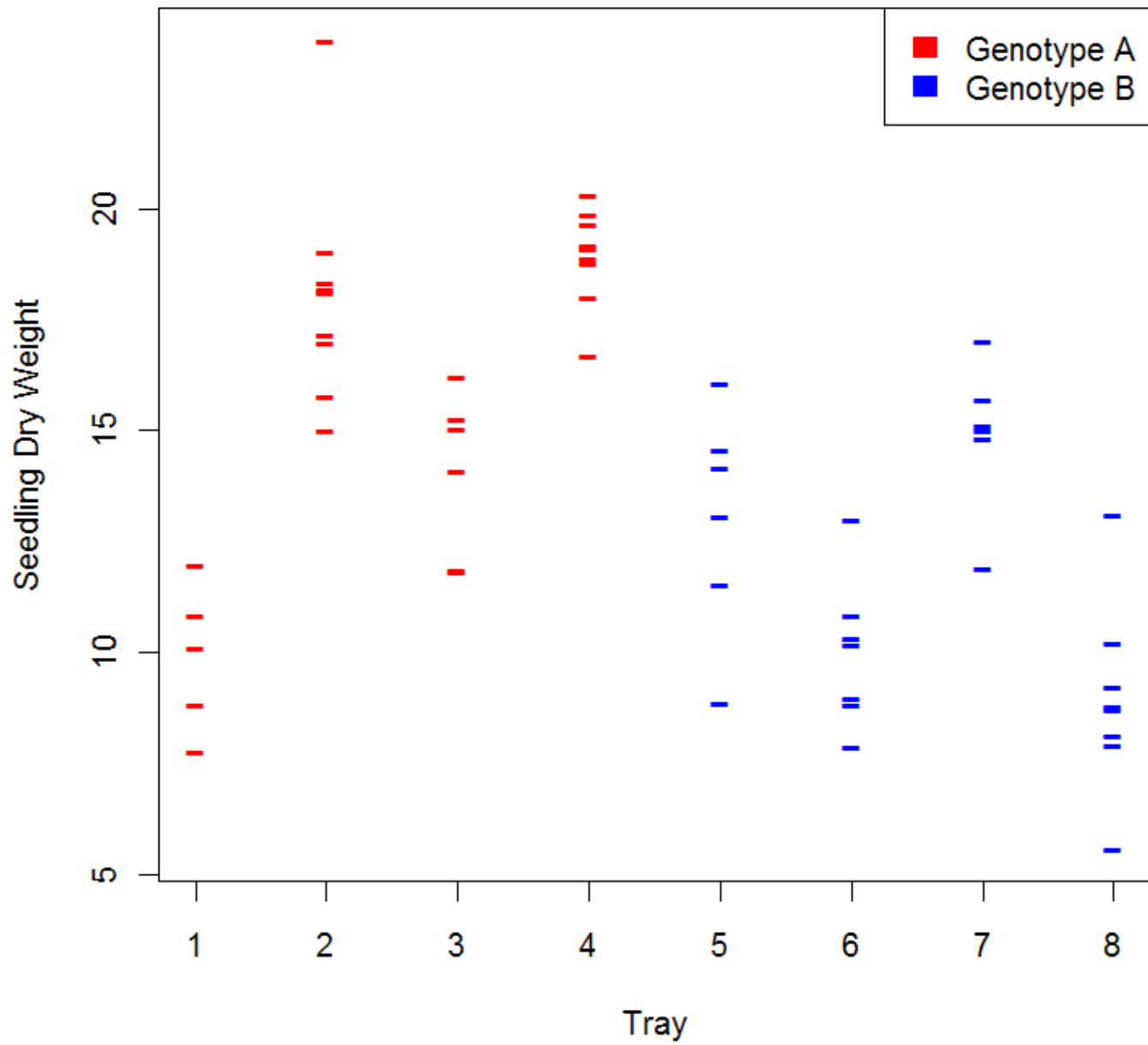
```
d=read.delim(  
"http://www.public.iastate.edu/~dnett/S511/SeedlingDryWeight2.txt"  
)  
d
```

	Genotype	Tray	Seedling	SeedlingWeight
1	A	1	1	8
2	A	1	2	9
3	A	1	3	11
4	A	1	4	12
5	A	1	5	10
6	A	2	1	17
7	A	2	2	17
8	A	2	3	16
9	A	2	4	15
10	A	2	5	19
11	A	2	6	18
12	A	2	7	18
13	A	2	8	18
14	A	2	9	24
15	A	3	1	12
16	A	3	2	12
17	A	3	3	16

18	A	3	4	15
19	A	3	5	15
20	A	3	6	14
21	A	4	1	17
22	A	4	2	20
23	A	4	3	20
24	A	4	4	19
25	A	4	5	19
26	A	4	6	18
27	A	4	7	20
28	A	4	8	19
29	A	4	9	19
30	B	5	1	9
31	B	5	2	12
32	B	5	3	13
33	B	5	4	16
34	B	5	5	14
35	B	5	6	14
36	B	6	1	10
37	B	6	2	10
38	B	6	3	9

39	B	6	4	8
40	B	6	5	13
41	B	6	6	9
42	B	6	7	11
43	B	7	1	12
44	B	7	2	16
45	B	7	3	17
46	B	7	4	15
47	B	7	5	15
48	B	7	6	15
49	B	8	1	9
50	B	8	2	6
51	B	8	3	8
52	B	8	4	8
53	B	8	5	13
54	B	8	6	9
55	B	8	7	9
56	B	8	8	10

```
plot(d[,2],d[,4]+rnorm(56,0,.2),  
     xlab="Tray",ylab="Seedling Dry Weight",  
     col=2*(1+(d[,1]=="B")),pch="-",cex=2)  
  
legend("topright",c("Genotype A","Genotype B"),  
      fill=c(2,4),border=c(2,4))
```



```
library(nlme)
```

```
lme(SeedlingWeight~Genotype,random=~1|Tray,  
method="ML",data=d)
```

```
Linear mixed-effects model fit by maximum likelihood
```

```
Data: d
```

```
Log-likelihood: -126.3709
```

```
Fixed: SeedlingWeight ~ Genotype
```

```
(Intercept)      GenotypeB
```

```
15.301832      -3.567017
```

```
Random effects:
```

```
Formula: ~1 | Tray
```

```
(Intercept) Residual
```

```
StdDev:      2.932294 1.882470
```

```
Number of Observations: 56
```

```
Number of Groups: 8
```

```
library(lme4)
```

```
lmer(SeedlingWeight~Genotype+(1|Tray),REML=F,data=d)
```

```
Linear mixed model fit by maximum likelihood
```

```
Formula: SeedlingWeight ~ Genotype + (1 | Tray)
```

```
Data: d
```

AIC	BIC	logLik	deviance	REMLdev
260.7	268.8	-126.4	252.7	247.4

```
Random effects:
```

Groups	Name	Variance	Std.Dev.
Tray	(Intercept)	8.5984	2.9323
Residual		3.5437	1.8825

```
Number of obs: 56, groups: Tray, 8
```


Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	15.302	1.510	10.14
GenotypeB	-3.567	2.136	-1.67

Correlation of Fixed Effects:

(Intr)	
GenotypeB	-0.707