

The Likelihood Ratio Test

The likelihood ratio test is a general purpose test designed evaluate nested statistical models in a way that is strictly analogous to the F-test for reduced models (RM) and full models (FM) commonly employed with linear models (see *Biostatistics Worksheet 402*). In both, failure to reject the null hypothesis results in model simplification. The likelihood ratio test works not only with linear models, shown here, but may be applied to a very wide array of problems involving Generalized Linear Models (GLM), where Maximum Likelihood (ML) or Restricted Maximum Likelihood (REML) methods are utilized to estimate model parameters. The latter methods/models include, among others, Logistic Regression (see *GLM 020*), Poisson Regression (see *GLM 040*), and Linear Mixed Models (see *LMM 060*) described in Worksheets on the *Biologist's Analytic Toolkit Website* under *Statistical Models*. For direct comparison of results, the data set analyzed here is the same as for the general F test (*Biostatistics Worksheet 402*). As can be seen, F test and likelihood ratio tests give similar but not exactly the same results. Helpful discussion of this approach appears in Kuter et al. (KNNL) *Applied Linear Statistical Models* 5th Edition, and numerous statistics websites.

Example in R:

```
#LOG LIKELIHOOD AND LIKELIHOOD RATIO TEST
setwd("c:/DATA/Models/")
K=read.table("KNNLCh9SurgicalUnit.txt")
K
attach(K)
```

> K

	X1	X2	X3	X4	X5	X6	Y
1	6.7	62	81	2.59	50	0	6.544
2	5.1	59	66	1.70	39	0	5.999
3	7.4	57	83	2.16	55	0	6.565
4	6.5	73	41	2.01	48	0	5.854
5	7.8	65	115	4.30	45	0	7.759
6	5.8	38	72	1.42	65	1	5.852
7	5.7	46	63	1.91	49	1	6.250
8	3.7	68	81	2.57	69	1	6.619
9	6.0	67	93	2.50	58	0	6.962
10	3.7	76	94	2.40	48	0	6.875
11	6.3	84	83	4.13	37	0	6.613
12	6.7	51	43	1.86	57	0	5.549
13	5.8	96	114	3.95	63	1	7.361
...							
44	6.5	56	77	2.85	41	0	6.288
45	3.4	77	93	1.48	69	0	6.178
46	6.5	40	84	3.00	54	1	6.416
47	4.5	73	106	3.05	47	1	6.867
48	4.8	86	101	4.10	35	1	7.170
49	5.1	67	77	2.86	66	1	6.365
50	3.9	82	103	4.55	50	0	6.983
51	6.6	77	46	1.95	50	0	6.005
52	6.4	85	40	1.21	58	0	6.361
53	6.4	59	85	2.33	63	0	6.310
54	8.8	78	72	3.20	56	0	6.478

Fitting Full and Reduced linear models:

Full Model:

```
#FITTING THE FULL LINEAR MODEL
FM=lm(Y~X1+X2+X3+X4+X5+factor(X6))
FMg=glm(Y~X1+X2+X3+X4+X5+factor(X6))
anova(FM)
anova(FMg)
```

Note: R's function `glm()` is also employed here since this function produces a data class for which the general wrapper `anova()` assumes `anova.glm()` which produces likelihood ratio results as default.

> anova(FM)

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
X1	1	0.7763	0.7763	12.5579	0.0009042 ***
X2	1	2.5888	2.5888	41.8803	5.187e-08 ***
X3	1	6.3341	6.3341	102.4704	2.157e-13 ***
X4	1	0.0246	0.0246	0.3976	0.5313820
X5	1	0.1265	0.1265	2.0460	0.1592180
factor(X6)	1	0.0522	0.0522	0.8448	0.3627348
Residuals	47	2.9053	0.0618		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

$$\sigma_{FM} = \sqrt{0.0618} = 0.2486$$

^ "standard error" (standard deviation of the residuals) for the full model

> anova(FMg)

Analysis of Deviance Table

Model: gaussian, link: identity

Response: Y

Terms added sequentially (first to last)

	Df	Deviance	Resid.	Df	Resid. Dev
NULL				53	12.8077
X1	1	0.7763		52	12.0315
X2	1	2.5888		51	9.4427
X3	1	6.3341		50	3.1085
X4	1	0.0246		49	3.0840
X5	1	0.1265		48	2.9575
factor(X6)	1	0.0522		47	2.9053

Reduced Model:**FITTING A REDUCED LINEAR MODEL****RM=lm(Y~X1+X2+X3+X5)****RMg=glm(Y~X1+X2+X3+X5)****anova(RM)****anova(RMg)****> anova(RM)**

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
X1	1	0.7763	0.7763	12.8495	0.000776 ***
X2	1	2.5888	2.5888	42.8528	3.349e-08 ***
X3	1	6.3341	6.3341	104.8499	9.118e-14 ***
X5	1	0.1484	0.1484	2.4561	0.123503
Residuals	49	2.9602	0.0604		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

$$\sigma_{RM} = \sqrt{0.0604} = 0.2458$$

^ "standard error" (standard deviation
of the residuals) for the reduced model

> anova(RMg)

Analysis of Deviance Table

Model: gaussian, link: identity

Response: Y

Terms added sequentially (first to last)

	Df	Deviance	Resid. Df	Resid. Dev
NULL			53	12.8077
X1	1	0.7763	52	12.0315
X2	1	2.5888	51	9.4427
X3	1	6.3341	50	3.1085
X5	1	0.1484	49	2.9602

Estimating Standard Error using Maximum Likelihood:

difference in number of parameters >
between models FM & RM

n := 54

k := 7

r := 2

 $\sigma_{FM} := 0.2486247$ $\sigma_{RM} := 0.2457874$

$$ML\sigma_{FM} := \sqrt{\frac{n-k}{n}} \cdot \sigma_{FM}$$

ML $\sigma_{FM} = 0.231951$

$$ML\sigma_{RM} := \sqrt{\frac{n-k+r}{n}} \cdot \sigma_{RM}$$

ML $\sigma_{RM} = 0.234132$ **> #CALCULATING MAXIMUM LIKELIHOOD
STANDARD DEVIATION****> n=length(K[,1])****> n #NUMBER OF CASES IN DATASET K**

[1] 54

> k=length(K)**> k #NUMBER OF VARIABLES IN FM**

[1] 7

**> r=2 #DIFFERENCE IN NUMBER OF
VARIABLES FM VS RM****> r**

[1] 2

> #EXTRACTING STANDARD ERRORS:**> #FOR FM:****> FMsigma = summary(FM)\$sigma****> FMsigma**

[1] 0.2486247

> #FOR RM:**> RMsigma = summary(RM)\$sigma****> RMsigma**

[1] 0.2457874

> #MAXIMUM LIKELIHOOD STANDARD DEVIATIONS**> #FOR FM:****> FMsigma.ML = FMsigma*sqrt((n-k)/n)****> FMsigma.ML**

[1] 0.2319511

> #FOR RM:**> RMsigma.ML = RMsigma*sqrt((n-k+r)/n)****> RMsigma.ML**

[1] 0.234132

Calculating Log Likelihoods for Each Model:

```

> # LOG LIKELIHOOD OF MODELS
> # FM:
> sum(log(dnorm(x = Y, mean = predict(FM), sd = FMsigma.ML)))
[1] 2.28368
> logLik(FM)
'log Lik.' 2.28368 (df=8)
> # RM:
> sum(log(dnorm(x = Y, mean = predict(RM), sd = RMsigma.ML)))
[1] 1.778311
> logLik(RM)
'log Lik.' 1.778311 (df=6)

In likelihood value for FM >
In likelihood value for RM >
^ ln ="natural logs" in base e

```

Note: log likelihoods for each model are calculated here using maximum likelihood estimates of standard error for *each model separately*. This contrasts with the use of standard error using only the FM in the test below.

Likelihood Ratio Test:

Assumptions:

- Standard Linear Regression depends on specifying in advance which variable is to be considered 'dependent' and which 'independent'. This decision matters as changing roles for Y & X usually produces a different result.\

- $Y_1, Y_2, Y_3, \dots, Y_n$ (dependent variable) is a random sample.

Note: Although a Normal distribution is assumed here for Y in a linear model, in other instances of the likelihood ratio test, this assumption doesn't apply.

- $X_1, X_2, X_3, \dots, X_n$ (independent variable) with each value of X_i matched to Y_i

Within this setup, two models for the relationship between X and Y variables are explicitly compared:

Full Model: where: Y_i and $[X_1, X_2, \dots, X_i]$ are matched dependent and independent variables, and

$$Y_i = \beta_0 + \sum \beta_j X_i + \varepsilon_i$$

β_0 is the y **intercept** of the regression line (translation)

Reduced Model:

β_j are slope coefficients for the *full set* of independent variables X_1, X_2, \dots, X_j

β_k are slope coefficients for a smaller *set* of independent variables *within* X_i

$$Y_i = \beta_0 + \sum \beta_k X_i + \varepsilon_i$$

ε_i is the error factor in prediction of Y_i and a random variable $\sim N(0, \sigma^2)$.

Hypotheses:

H_0 : coefficients in j but NOT INCLUDED in k = 0.

Note: this is always the more parsimonious (i.e., smaller) model

H_1 : at least some of these coefficients not 0

Degrees of Freedom:

n = 54 < n = number of matched observations in dataset
k = 7 < k = number of variables in FM
r = 2 < r = differences in number of variables between FM & RM

Sum of Squares and Standard Error for FM:**> anova(FM)**

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
X1	1	0.7763	0.7763	12.5579	0.0009042
X2	1	2.5888	2.5888	41.8803	5.187e-08
X3	1	6.3341	6.3341	102.4704	2.157e-13
X4	1	0.0246	0.0246	0.3976	0.5313820
X5	1	0.1265	0.1265	2.0460	0.1592180
factor(X6)	1	0.0522	0.0522	0.8448	0.3627348
Residuals	47	2.9053	0.0618		

> anova(RM)

Analysis of Variance Table

Response: Y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
X1	1	0.7763	0.7763	12.8495	0.000776 ***
X2	1	2.5888	2.5888	42.8528	3.349e-08 ***
X3	1	6.3341	6.3341	104.8499	9.118e-14 ***
X5	1	0.1484	0.1484	2.4561	0.123503
Residuals	49	2.9602	0.0604		

s := 0.2486247

SSE_{FM} := 2.90527SSE_{RM} := 2.960161

^ Standard errors are the square root of MSE, see above.

> #LIKELIHOOD RATIO TEST:**> #SUM OF SQUARES ERROR FOR MODELS:****> SSE.FM = sum((Y-predict(FM))^2) #SSE for FM****> SSE.FM**

[1] 2.90527

> SSE.RM = sum((Y-predict(RM))^2) #SSE for RM**> SSE.RM**

[1] 2.960161

> #STANDARD ERROR FOR FM:**> s=summary(FM)\$sigma****> s**

[1] 0.2486247

> s=sqrt(summary(FM)\$dispersion)**> s**

[1] 0.2486247

Relative Likelihoods:

see eq 1.26 in KNNL

$$\text{LFM} := e^{-\frac{1}{2} \frac{\text{SSE}_{\text{FM}}}{s^2}}$$

$$\text{LFM} = 6.2241 \times 10^{-11}$$

$$\text{LRM} := e^{-\frac{1}{2} \frac{\text{SSE}_{\text{RM}}}{s^2}}$$

$$\text{LRM} = 3.9926 \times 10^{-11}$$

$$C := \frac{1}{\left(2 \cdot \pi \cdot s^2\right)^{\frac{n}{2}}}$$

$$C = 1.2296 \times 10^{11}$$

> #RELATIVE LIKELIHOODS FOR THE MODELS:**> LFM = exp(-(1/2)*(SSE.FM/s^2)) #TIMES CONSTANT C****> LFM**

[1] 6.224145e-11

> LRM = exp(-(1/2)*(SSE.RM/s^2)) #TIMES CONSTANT C**> LRM**

[1] 3.992573e-11

> #CONSTANT C:**> n #NUMBER OF CASES IN DATASET K**

[1] 54

> C=1/((2*pi*s^2)^(n/2)) #CONSTANT IN EQ 1.26 IN KNNL**> C**

[1] 122956414826

Likelihoods:

$$\Lambda_{\text{FM}} := C \cdot \text{LFM}$$

$$\Lambda_{\text{FM}} = 7.653$$

$$\Lambda_{\text{RM}} := C \cdot \text{LRM}$$

$$\Lambda_{\text{RM}} = 4.9091$$

> LCFM=C*LFM**> LCFM**

[1] 7.652985

> LCRM=C*LRM**> LCRM**

[1] 4.909125

Likelihood Ratio Test Statistic:

$SSE_{FM} := 2.9053$

$SSE_{RM} := 2.9602$

$$LRT := \frac{(SSE_{RM} - SSE_{FM})}{s^2}$$

$LRT = 0.8881$ < difference here due to rounding...

```
> #LOG LIKELIHOOD RATIO STATISTIC:
> LRT=(SSE.RM - SSE.FM)/s^2
> LRT #LOG LIKELIHOOD RATIO STATISTIC
[1] 0.8880001
```

Critical Value of the Test:

$\alpha := 0.05$ < Probability of Type I error must be explicitly set

$CV := qchisq(1 - \alpha, r)$

$CV = 5.9915$

< note degrees of freedom reflect difference between the models

Decision Rule:

IF $F > CV$, THEN REJECT H_0 OTHERWISE ACCEPT H_0

$LRT = 0.8881$

$CV = 5.9915$

Probability Value:

$P := 1 - pchisq(LRT, r)$

$P = 0.6414$

```
> #PROBABILITY OF NULL HYPOTHESIS RM
> P=1-pchisq(LRT,2)
> P #PROBABILITY
[1] 0.6414654
```

IMPORTANT NOTE:

FALURE to reject H_0 in this test means that the MORE PARSIMONIOUS model RM is PREFERRED!

Prototype in R:

```
#LIKELIHOOD RATIO TEST:
anova(RM,FM,test="LRT")
anova(RMg,FMg,test="LRT")
```

```
> #LIKELIHOOD RATIO TEST:
> anova(RM,FM,test="LRT")
Analysis of Variance Table
```

```
Model 1: Y ~ X1 + X2 + X3 + X5
Model 2: Y ~ X1 + X2 + X3 + X4 + X5 + factor(X6)
  Res.Df  RSS Df Sum of Sq Pr(>Chi)
1     49 2.9602
2     47 2.9053  2  0.054891  0.6415
```

```
> anova(RMg,FMg,test="LRT")
Analysis of Deviance Table
```

```
Model 1: Y ~ X1 + X2 + X3 + X5
Model 2: Y ~ X1 + X2 + X3 + X4 + X5 + factor(X6)
  Resid. Df Resid. Dev Df Deviance Pr(>Chi)
1     49     2.9602
2     47     2.9053  2  0.054891  0.6415
```