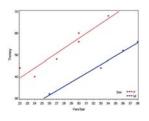
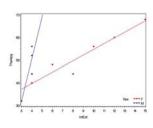


### Regression: Model assessment



Psychology 6140



# **Topics**

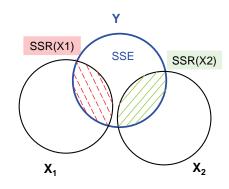
- How to assess the contributions of individual predictors to a regression model?
  - Type I (sequential) tests: added contribution of each new variable, in a given order
  - Type II (partial) tests: unique contribution of each variable, above all others
- More general methods: the general linear test: H<sub>0</sub>: Lβ=0
- The Marginality principle: always include low-order relatives
  - Testing hierarchical (ordered subset) models
  - Moderator variables: interaction effects

# **Uncorrelated predictors**

 When predictors are uncorrelated, their SSR are additive

$$SSR(X1 X2) = SSR(X1) + SSR(X2)$$

 This makes it easy to see & test the contributions of each predictor



This "Ballentine" diagram shows variance or SS by areas of circles

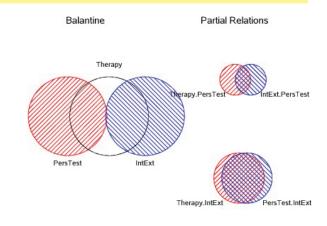
3

# Example: therapy data

For the therapy data, PersTest and IntExt turn out to be nearly uncorrelated.

Each have modest correlations with Therapy, but jointly account for 92%

In this example, IntExt acts as a suppressor variable for the test of PersTest, by removing effect of IntExt from error variance





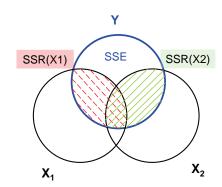
2

### Correlated predictors

- Typically, predictors are correlated
- So, the portions of variance of Y they account for overlap

$$SSR(X1 X2) < SSR(X1) + SSR(X2)$$

 How to assess contribution of each X?



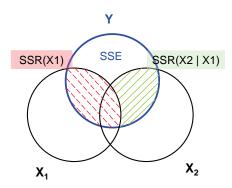
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## Sequential & partial SS

- Sequential SS (Type I)
  - 1st variable accts for all it can
  - Each next var: only what is left over
- ∴ contributions are additive

$$SSR(X1 X2) = SSR(X1) + SSR(X2 \mid X1)$$

- Only useful if there is a reason for ordering variables
  - e.g., polynomial models
  - e.g., hierarchical models



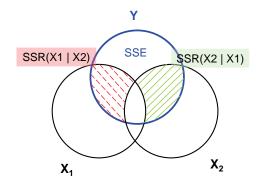
 $H_0$ :  $\beta_1 = 0$  (ignoring X2, X3)

 $H_0$ :  $\beta_2 = 0$  (adjusting for only X1)

 $H_0$ :  $\beta_3 = 0$  (adjusting for X1, X2)

# Sequential & partial SS

- Partial SS (Type II)
  - Each var accts for its unique contribution
  - Q: can we delete X<sub>i</sub> given that all others are included?
  - $t = b_i / s(b_i)$  is a partial test
- These are most generally useful, except where there is a hierarchical ordering of predictors
- In ANOVA designs there are also Type III (and IV) tests (take empty cells into account)



 $H_0$ :  $\beta_1 = 0$  (adjusting for X2, X3)

 $H_0$ :  $\beta_2 = 0$  (adjusting for X1, X3)

 $H_0$ :  $\beta_3 = 0$  (adjusting for X1, X2)

#### Multiple regression: therapy data

proc reg data=therapy;
 model therapy = perstest intext sx;
 sx=1 for female here
run;

3					
Analysis of Va	riance				
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model Error Corrected Total	3 6 9	982.05152 17.94848 1000.00000	327.35051 2.99141	109.43	<.0001
Root MSE Dependent Mean Coeff Var	1.72957 50.00000 3.45914	R-Square Adj R-Sq	0.9821 0.9731		
	Paramet	ter Estimates			
Variable DF	Parameter Estimate	Standard Error		Pr >  t	Partial tests
Intercept 1 PERSTEST 1 INTEXT 1 SX 1	-14.79157 1.71897 0.96956 10.72600	5.22575 0.17268 0.25620 2.40251	-2.83 9.95 3.78 4.46	0.0299 <.0001 0.0091 0.0043	

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#### Sequential vs. partial tests

```
proc reg data=therapy;
  model therapy = perstest intext sx / SS1 SS2;
  run;
```

Parameter	Estimates					
Variable	Label	DF	Parameter Estimate	Standard Error		
Intercept PERSTEST INTEXT SX	Intercept Personality Test Score Internal External scale Sex	1 1 1 1	-14.79157 1.71897 0.96956 10.72600	5.22575 0.17268 0.25620 2.40251	9.95 3.78	
	Paramet	er Es	timates			
Variable	Label	DF	Pr >  t	Type I SS	Type II SS	
Intercept PERSTEST INTEXT SX	Intercept Personality Test Score Internal External scale Sex	1 1 1 1		25000 360.00000 562.42744 59.62408	23.96662 296.42129 42.84039 59.62408	
						' \

F = SSR / MSE gives the test statistic for each hypothesis

SSR(X1) SSR(X2 | X1) SSR(X3 | X1 X2) SSR(X1 | X2 X3) SSR(X2 | X1 X3) SSR(X3 | X1 X2) IVIO

# Model comparison

- All statistical tests resolve to comparisons between two models
  - E.g., simple linear regression:  $H_0$ :  $\beta_1 = 0$  vs  $H_a$ :  $\beta_1 \neq 0$ 
    - Full model:  $y_i = \beta_0 + \beta_1 x_i + \epsilon_i$
    - Reduced model:  $y_i = \beta_0 + \epsilon_i$
    - Test:

$$F^{\star} = \frac{(SSR_{full} - SSR_{reduced}) / (df_{full} - df_{reduced})}{MSE_{full}} = \frac{SS_{hyp} / df_{hyp}}{SS_{E} / df_{E}}$$

 More generally, we can compare any larger model to a subset model, using the extra sum of squares, e.g.,

$$H_0$$
:  $\beta_3 = \beta_4 = 0$   
 $SSR(X_3 X_4 | X_1 X_2) = \underbrace{SSR(X_1 X_2 X_3 X_4)}_{full} - \underbrace{SSR(X_1 X_2)}_{reduced}$ 

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### Testing composite hypotheses

proc reg;  
 model therapy = perstest intext sx;  
 test intext, sx;  
 run;
Test 
$$\beta_2 = \beta_3 = 0 \mid \beta_1$$

$$SSR(X_2 | X_3 | X_1) = SSR(X_1 | X_2 | X_3) - SSR(X_1) = 982.05 - 360 = 622.05$$

F\* = 
$$\frac{(SSR_{x_1,x_2,x_3} - SSR_{x_1})/(3-1)}{MSE_{x_1,x_2,x_3}} = \frac{622.05/2}{2.99} = 103.97$$

#### General Linear Hypothesis Tests

 Even more generally, any hypothesis test can be regarded as an example of a GLH of the form

$$H_0: \underset{q \times (p+1)}{\mathbf{L}} \quad \mathbf{\beta} = \mathbf{0}$$

where the hypothesis matrix,  $\mathbf{L}$ , contains specified constants and is of rank  $q = \mathrm{df}$  for hypothesis

• e.g.

$$\mathbf{L}\boldsymbol{\beta} = \begin{bmatrix} 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{pmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} = \begin{pmatrix} 0 \\ 0 \end{pmatrix} \qquad \qquad \mathbf{L}\boldsymbol{\beta} = \begin{bmatrix} 0 & 1 & -1 & 0 \end{bmatrix} \begin{pmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} = 0$$

SAS svntax

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test intext, sx;

test perstest - intext;

### General Linear Hypothesis Tests

In all cases, the sums of squares for the hypothesis, H<sub>0</sub>:
 L β = 0 has the same form,

$$SS_{hyp} = (\mathbf{Lb})^T \mathbf{L} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{L}^T \mathbf{Lb}$$

This measures the squared distance of L  $\beta$  from 0

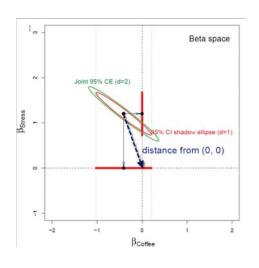
- GLH tests extend in a natural way to
  - MANOVA, MMReg: Y = X B + E

$$LB = 0$$

Repeated measures designs

$$LBM=0$$

#### Example: Heart disease, coffee and stress



In the model Im(Heart ~ Coffee + Stress, data=coffee)

Test: 
$$H_0$$
:  $\beta_{\text{Coffee}} = \beta_{\text{Stress}} = 0$   
(X'X)<sup>-1</sup> is covariance matrix of  $\beta = \begin{pmatrix} \beta_0 \\ \beta_1 \\ \beta_1 \end{pmatrix}$ 

$$\mathbf{L} = \begin{bmatrix} 0 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \quad \text{selects } \beta_1, \, \beta_2$$

$$(\mathbf{Lb})^{\mathsf{T}} \ \mathbf{L}(\mathbf{X}^{\mathsf{T}}\mathbf{X})^{-1}\mathbf{L}^{\mathsf{T}} \ (\mathbf{Lb})$$

is squared distance of **b** from (0,0)

ance of **b** from (0,0)

# Marginality principle

- Any model including a high-order term should normally include all low-order relatives
  - Interactions: Perstest \* Sex → Perstest + Sex ("main effects")
  - Polynomial models: X<sup>3</sup> → X + X<sup>2</sup>
- We can neither test nor interpret main effects of variables that interact
  - $X_1 * X_2 \rightarrow \text{effect } (\beta_1) \text{ of } X_1 \text{ varies with } X_2$
- Similarly, if X<sup>3</sup> is important, X and X<sup>2</sup> must remain in the model (even if NS!)

# Hierarchical testing

- Variables in regression are sometimes ordered in terms of research questions & hypotheses
  - Include necessary control variables (age, IQ)
  - Test for effects of new predictor(s) beyond old ones
- In such cases, do hierarchical (blockwise) tests

proc reg data=mydata;

var y age IQ reading math depression anxiety;

block1: model y = age IQ;

block2: model y = age IQ reading math;

test reading, math;

block3: model y = age IQ reading math depression anxiety;

test depression, anxiety;

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#### Moderator variables

- A moderator effect occurs when the effect of one variable, x<sub>1</sub>, depends on, or varies with variable, x<sub>2</sub>.
  - i.e., interaction of x₁ and x₂.
  - i.e., slope (b₁), for x₁ varies with x₂.
- In regression, this is modeled by including the product, x<sub>1</sub> \* x<sub>2</sub> in the model

$$y = b_0 + b_1 x_1 + b_2 x_2 + b_3 (x_1 \times x_2) + \epsilon$$
  
=  $b_0 + (b_1 + b_3 x_2) x_1 + b_2 x_2 + \epsilon$ 

- In SAS proc reg & SPSS must calculate x<sub>1</sub> \* x<sub>2</sub> explicitly
- (often useful for interpretation to center x<sub>1</sub>, x<sub>2</sub>)
- Must include x<sub>1</sub> and x<sub>2</sub> (marginality)
- Must test x<sub>1</sub>\*x<sub>2</sub> by partial test
- Conclude no moderator effect if b<sub>3</sub> is non-significant

run;							
Variable	Label	DF	Parameter Estimate	Standard Error	t Value	Pr >  t	
Intercept PERSTEST SX	Intercept PersTest Sex	1 1 1	-20.70091 2.00604 9.94015	11.41755 0.34022 14.44245	-1.81 5.90 0.69	0.1198 0.0011 0.5170	

0.46331

There is no evidence that the slope for PersTest varies with Sex

data therapy; set therapy;

PTxSX = perstest \* sx;

\*-- test moderator of Sex on Perstest;

model therapy = perstest sx PTxSX;

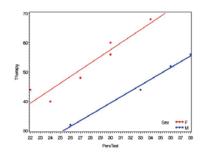
IExSX = intext \* sx;

proc reg data=therapy;

PTxSX

The slope for females is only 0.27 less than that for males

Interpreting such models is easiest if you plot the fitted relationships



0.59

0.5775

calculate products

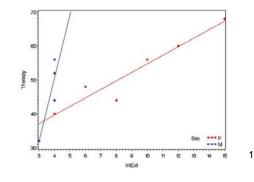
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```
*-- test moderator of Sex on IntExt;
proc reg data=therapy;
  model therapy = intext sx IExSX;
  run;
```

			Parameter	Standard			
Variable	Label	DF	Estimate	Error	t Value	Pr >  t	
Intercept	Intercept	1	-24.00000	19.53597	-1.23	0.2653	
INTEXT	IntExt	1	18.66667	5.17520	3.61	0.0113	
SX	Sex	1	53.60825	20.14653	2.66	0.0375	
IExSX		1	-16.15120	5.19916	-3.11	0.0209	

There is strong evidence that the slope for IntExt varies with Sex

The slope for females is 16.15 less than for males



Arguably, it might be better to test the full model, with both interactions:

```
*-- test both moderators with sex;
proc reg data=therapy;
  model therapy = perstest intext sx PTxSX IExSX;
  test PTxSX, IExSX;
  run;
```

Parameter   Standard   Pr   Ferror   Value   Pr   Pr   Pr   Pr   Pr   Pr   Pr   P	Variable         Label         DF         Estimate         Error         t Value         Pr >  t            Intercept         Intercept         1         -16.73684         3.48499         -4.80         0.0086           PERSTEST         PersTest         1         2.42105         0.22056         10.98         0.0004           INTEXT         IntExt         1         -4.73684         2.31673         -2.04         0.1104           SX         Sex         1         22.97128         4.53565         5.06         0.0072							
PERSTEST         PersTest         1         2.42105         0.22056         10.98         0.0004           INTEXT         IntExt         1         -4.73684         2.31673         -2.04         0.1104           SX         Sex         1         22.97128         4.53565         5.06         0.0072           PTxSX         1         -1.22461         0.26226         -4.67         0.0095	PERSTEST         PersTest         1         2.42105         0.22056         10.98         0.0004           INTEXT         IntExt         1         -4.73684         2.31673         -2.04         0.1104           SX         Sex         1         22.97128         4.53565         5.06         0.0072           PTxSX         1         -1.22461         0.26226         -4.67         0.0095	Variable	Label	DF			t Value	Pr >  t
	1 0.10934 2.32193 2.00 0.0000	PERSTEST INTEXT SX PTxSX	PersTest IntExt	1 1 1 1	2.42105 -4.73684 22.97128 -1.22461	0.22056 2.31673 4.53565 0.26226	10.98 -2.04 5.06 -4.67	0.0004 0.1104 0.0072 0.0095

Joint test for both interactions: Do I need any interactions with sex?

Source	DF	Mean Square	F Value	Pr > F
Numerator Denominator	2 4	7.74189 0.61618	12.56	0.0189

#### Testing and comparing models in R

- In R, fit a model using mod <-lm(y ~ x1+x2+ ...)</li>
- Test terms in that model using summary (mod)
- Type II F-tests with car::Anova(mod)
- Compare models using anova (mod1, mod2, ...)
- Linear hypotheses: car::linearHypothesis()

```
mod1 <- lm(therapy ~ perstest, data= therapy)</pre>
mod2 <- lm(therapy ~ perstest + intext, data=therapy)</pre>
mod3 <- lm(therapy ~ perstest + intext + sex, data=therapy)</pre>
summary(mod3)
Anova(mod3) # F tests
# test interactions
mod4 <- lm(therapy ~ perstest*sex + intext*sex, data=therapy)</pre>
# compare models
anova(mod1, mod2, mod3, mod4)
```

```
> coef(mod3)
                                                               Linear hypotheses
(Intercept)
             perstest
                          intext
 -4.065574 1.718970 0.969555 -10.725995
                                                               for 1 or more
> linearHypothesis(mod3, c("intext", "sexM"))
                                                               coefficients in mod3
Linear hypothesis test
Hypothesis:
intext = 0
sexM = 0
Model 1: restricted model
Model 2: therapy ~ perstest + intext + sex
 Res.Df RSS Df Sum of Sq
                            F Pr(>F)
     8 640.00
      6 17.95 2 622.05 103.97 2.206e-05 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
> tests <- matchCoefs(mod4, ":")</pre>
> linearHypothesis(mod4, tests)
                                                              Test all interactions
Linear hypothesis test
                                                              (":" in name) in mod4
Hypothesis:
perstest:sexM = 0
sexM:intext = 0
Model 1: restricted model
Model 2: therapy ~ perstest * sex + intext * sex
 Res.Df RSS Df Sum of Sq
                                  F Pr(>F)
     6 17.9485
      4 2.4647 2 15.484 12.564 0.01886 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

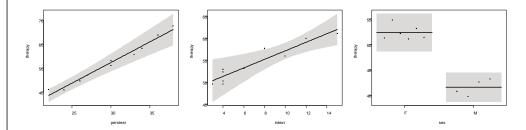
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```
> summary(mod3)
Call:
lm(formula = therapy ~ perstest + intext + sex, data = therapy)
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
                                                          Partial t-tests for
(Intercept) -4.0656 3.7572 -1.082 0.32078
                                                          coefs in mod3
perstest
            1.7190 0.1727 9.954 5.94e-05 ***
           0.9696 0.2562 3.784 0.00913 **
           -10.7260 2.4025 -4.464 0.00426 **
sexM
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.73 on 6 degrees of freedom
Multiple R-squared: 0.9821, Adjusted R-squared: 0.9731
F-statistic: 109.4 on 3 and 6 DF, p-value: 1.256e-05
> anova(mod1, mod2, mod3, mod4)
                                                        Hierarchical tests of
Analysis of Variance Table
                                                        mod, vs. mod, 1
Model 1: therapy ~ perstest
Model 2: therapy ~ perstest + intext
Model 3: therapy ~ perstest + intext + sex
Model 4: therapy ~ perstest * sex + intext * sex
                                                         These assume nested
 Res.Df RSS Df Sum of Sq F Pr(>F)
                                                         models
    8 640.00
      7 77.57 1
                    562.43 912.770 7.149e-06 ***
      6 17.95 1 59.62 96.765 0.0005989 ***
      4 2.46 2 15.48 12.564 0.0188571 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Visualizing model effects

• In R, the effects and visreg packages make it easy to visualize the effects of terms in models

```
> library(visreg)
> visreg(mod3)
```

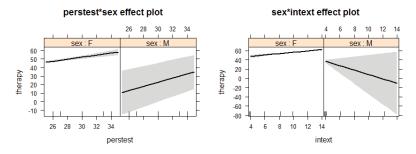


Conditional plots for each predictor, setting all others to their median value

### Visualizing model effects

 The effects package is most useful for plotting models with interactions

```
> library(effects)
> plot(allEffects(mod4))
```



Conditional plots for each high-order term, setting others to mean value

#### Summary

- Sequential (Type I) and Partial (Type II) SS provide different ways of assessing the contribution of a given predictor
  - Type I: added contribution of each new variable, in order
  - Type II: added contribution of each variable above all others
- Each of these essentially give a test comparing a "full" model against a "reduced" model
- This idea extends to the General Linear Test, H<sub>0</sub>: Lβ=0

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# Summary

- Tests of complex models must respect the Marginality Principle—include low-order relatives
- Testing hierarchical models and moderator variables are examples of these ideas.
- Always important to plot model terms for interpretation
- We will consider model selection problems more generally later