# Stat 13 <br> http://www.stat.ucla.edu/~dinov/courses_students.html <br> Chapter 12 Problems/Solutions 

All Problems are from: Myra L. Samuels and Jeffrey A. Witmer, Statistics for the Life Sciences, 3rd edition, Prentice-Hall (2003)
12.5. SOCR (http://www.socr.ucla.edu) output is given below.
(a) cob-wt $=316-0.721$ plant-density
(b) scatterplot (not shown) shows a strong negative linear association between cobweight (gm grain/cob) and plant density (\# plants / pot).
(c) as plant density increases by 1 plant per plot, cob weight decreases by 0.72 gm of grain per cob, on average.
(d) $\mathrm{sY}=\operatorname{sqrt}(11831.8 / 19)=25 \mathrm{gm}$ and $\mathrm{sY} / \mathrm{X}=\operatorname{sqrt}(1337.3 / 18)=8.6 \mathrm{gm}$
(e) Predictions of cob weight based on the regression model tend to be off by 8.6 gm on average.

Equivalently, the data points deviate above or below the regression line by 8.6 gm on average.

Regression Analysis
The regression equation is
cob-wt = 316-0.721 plant-density

| Predictor | Coef | StDev | T | P |
| :--- | :---: | :---: | :---: | :---: |
| Constant | 316.376 | 8.000 | 39.55 | 0.000 |
| plant-de | -0.72063 | 0.06063 | -11.89 | 0.000 |
| $\mathrm{~S}=8.619$ | $\mathrm{R}-\mathrm{Sq}=88.7 \%$ | $\mathrm{R}-\mathrm{Sq}(\mathrm{adj})=88.1 \%$ |  |  |
| Analysis of Variance |  |  |  |  |


| Source | DF | SS | MS | F | P |  |
| :--- | :---: | :---: | :---: | :--- | :--- | :--- |
| Regression | 1 | 10495 | 10495 | 141.26 | 0.000 |  |

Residual Error $181337 \quad 74$
Total 1911832

## 12.6

(a) The slope and intercept of the regression line are
b_1 = -927.75/1303 = -.7120;
b_0 $=$ y-bar $-\mathrm{b} \_1 \mathrm{x}-\mathrm{bar}=23.64-(-.7120)(11.5)=31.83$
The fitted regression line is $\quad y$-hat $=31.83-.7120 \mathrm{X}$
(c) $\mathrm{s}_{-} \mathrm{Y} \mid \mathrm{X}=\operatorname{sqrt}[\mathrm{SS}($ resid $) / \mathrm{df}]=\operatorname{sqrt}[16.7812 /(12-2)]=1.3$
12.8a: b1 = 161.40/50667 = . $0003186 ; \mathrm{b} 0=.210-(.0003186)(433.3)=.0720$.

The fitted regression line is $\mathrm{Y}=.072-.0003186 \mathrm{X}$.
12.8c: As altitude of origin goes up by 1 m , respiration rate goes up by $.0003186 \mathrm{mul} / \mathrm{hr}-\mathrm{mg}$, on average.
12.8d: $s_{-}\{Y \mid X\}=\operatorname{sqrt}[.013986 / 10]=.0374$

### 12.12

The intercept of the regression line b0 is based on all 12 data points, not just on the two point for which $X=0$. If there is a linear relationship between $X$ and $Y$ (a scatter plot of the data strongly suggest that there is), then the best estimate of the average for Y at any given value of X is given by the regression line, taking into account all of the data. In contrast, the average $(33.3+31.0) / 2=32.15$ ignores most of the data.
12.14a: (See Exercise 12.5 for b0 and b1.
(i) plugging in plant-density $=100$ plants gives a predicted cob-wt of 316 $0.721(100)=244.34 \mathrm{gm}$
(ii) plugging in plant-density $=120$ plants gives a predicted cob-wt of 316 -
$0.721(120)=229.93 \mathrm{gm}$

### 12.14b:

(i) $(224.34)(100)=24434 \mathrm{gm}=24.43 \mathrm{~kg}$
(ii) $(229.928)(12)=27591 \mathrm{gm}=27.6 \mathrm{~kg}$

### 12.15

Using the fitted regression line found in Exercise 12.6 above, we substitute $\mathrm{X}=15$. This yields $\quad y$-hat $=31.83-(.7120)(15)=21.1$.

Thus, we estimate that the mean fungus growth would be 21.1 mm at a laetisaric acid concentration of $15 \mathrm{microg} / \mathrm{ml}$.

According to the linear model, the standard deviation of fungus growth does not depend on X . Our estimate of this standard deviation from the regression line is the Residual Standard Deviation sigma_\{Y|X\} = sqrt[SS(resid)/(n-2)] = sqrt[16.7812/10] = 1.3 mm .

Thus we estimate that the standard deviation of fungus growth would be 1.3 mm at a laetisaric acid concentration of $15 \mathrm{microg} / \mathrm{ml}$.

For $\mathrm{X}=15$, we have y -hat $=21.1+/-1.3 \mathrm{~mm}$.
12.19a: $\mathrm{b} 1=81.90 / 2800=0.02925 \mathrm{ng} / \mathrm{min}$ (the rate of incorporation)
b0 $=0.83-(0.02925)(30)=-0.05$

$$
\text { s_ }\{y \mid x\}=\text { sqrt[SS(resid)/(n-2)] }=\operatorname{sqrt}[0.035225 / 5]=0.0839
$$

To construct a $95 \%$ confident interval, we consult the z-table (Table 4) with $\mathrm{df}=\mathrm{n}-2=7$ $2=5$; the multiplier is $\mathrm{t} \_\{4,0.025\}=2.571$. The confidence interval is

$$
\begin{aligned}
& \mathrm{b} 1+/-\mathrm{t} \_\{4,0.025\} \text { SEb1 }=0.02925+/-(2.571)(0.00159) \\
& \text { or } 0.0252<\text { beta } 1<0.033 \mathrm{ng} / \mathrm{min}
\end{aligned}
$$

12.19ba: We are 95\% confident that the rate at which leucine is incorporated into protein in the population of all Xenopus oocytes is between $0.0252 \mathrm{ng} / \mathrm{min}$ and $0.0333 \mathrm{ng} / \mathrm{min}$
12.21a: SEb1 $=8.6 / \operatorname{sqrt}(20209)=0.0605$, so $95 \%$ CI for b1 is

$$
-0.7206+/-(2.101)(0.0605) \text { or }-0.7206+/-0.1271 \text { or }(-0.848,-0.593)
$$

12.21b: We are $95 \%$ confident that as plant density increases by 1 plant per plot, average cob weight decreases by between 0.848 gm and 0.593 gm of grain per cob.
12.22a: From Exercise 12.6, s_Y|X = sqrt[SS(resid)/df] = sqrt[16.7812/(12-2)] = 1.3 The standard error of the slope is
SE_b1 = s_Y|X / sqrt[sum(x - x-bar)^2] = 1.3/sqrt[1303] $=0.36$
12.22b: H0: Leatisaric acid has no effect on fungus growth (beta_1 = 0)

HA: Laetisaric acid inhibits fungus growth (beta_1 < 0)
$\mathrm{t} \_\mathrm{s}=-.7120 / 0.36=-19.8$. With $\mathrm{df}=10$, the t -table (Table 4) gives $\mathrm{t} \_.0005=4.587$.
Thus the P-value $<.0005$, so we reject H0. There is sufficient evidence (P-value $<.0005$ ) to conclude that laetisaric acid inhibits fungus growth.
12.27a: $\quad \mathrm{r}=82.8977 / \mathrm{sqrt}[(28465.7)(.363708)]=.8147$
12.27b: $\quad$ s_Y $=\operatorname{sqrt}[(.363708 /(13-1)]=.1741 \mathrm{gm}$
s_Y|X = sqrt[SS(resid)/df] = sqrt[.1223/(13-2)] = . 1054 gm
$.1054 / .1741=.605 ; \operatorname{sqrt}[1-.8147 \wedge 2]=.580$
12.27c: b_1 = 82.8977/28465.7 = .002912;
b_0 $=2.174-(.002912)(443.8)=.882$
The fitted regession line is $\quad \mathrm{y}$-hat $=.882-.002912 \mathrm{X}$
12.28a: $r=-14563.1 /$ sqrt[ (20209)*(11831.8) ] $=-0.942$
12.28b: from Exercise 12.5 (d), sY $=25 \mathrm{gm}$ and $\mathrm{sY} / \mathrm{X}=8.6 \mathrm{gm}$, so sY/X / sY $=0.344$ further, sqrt( $1-\mathrm{r} 2$ ) $=0.3356$, which is nearly equal to 0.344 , so the approximate relationship is indeed verified.
12.28c: b1 = -14563.1/20209.0 = -.7206;
b0 $=224.1-(-.7206)(128.05)=316.4$
The fitted regression line is $\mathrm{Y}=316.4$ - . 7206X.
12.30: Let $X=$ age and let $Y=$ blood pressure. The Residual Standard Deviation is s_\{Y|X\} = sqrt[1-r^2](s_Y)sqrt[(n-1)/(n-2)] = sqrt[1-.43^2](19.5)sqrt[2668/2667] = 17.6 mm Hg .
s_ $\{\mathrm{Y} \mid \mathrm{X}\}=\operatorname{sqrt}\left[(\mathrm{y}-\mathrm{y} \text {-hat })^{\wedge} 2 /(\mathrm{n}-2)\right]$ is a measure of the variability about the regression line $y$-hat $=\mathrm{b} 1 \mathrm{x}+\mathrm{b} 0$.

But s_Y $=\operatorname{sqrt}\left[(y-y-b a r)^{\wedge} 2 /(n-1)\right]$ is a measure of the variability about the mean $y$-bar.
12.41a: with (iii),
12.41b: with (ii), and
12.41 c : with (i).
12.45a: The slope and intercept of the regression line are
b1 $=-.342 / .1512=-2.262$
b0 $=1.117-(-2.262)(.12)=1.39$
The fitted regression line is $\mathrm{Y}=1.39-2.262 \mathrm{X}$.
12.45c: s_\{Y|X\} = sqrt[SS(resid)/(n-2)] = sqrt[.2955/10] = . 1719 kg .
12.46a: If $\mathrm{x}=.24$, then predicted $\mathrm{y}=1.39-2.262(.24)=.84512$. But the variability of this prediction is given by s_\{Y|X\}=.17.

If x is unknown, then the best prediction is y -bar = 1.117, and the precision of this prediction is $+/-s_{-} \mathrm{Y}=.31175$. We write $\mathrm{y}=1.117+/-.31175 \mathrm{~kg}$.

However, if $\mathrm{x}=.24$ is known, then the best prediction for y is given by the regression line $\mathrm{y}=$ $1.39-2.262(.24)=.84512$, but the precision of this prediction is $+/-\mathrm{s} \_\{\mathrm{Y} \mid \mathrm{X}\}=+/-.17$. We write $\mathrm{y}=.84512+/-.17 \mathrm{~kg}$.
12.46b: The condition that sigma_ $\{\mathrm{Y} \mid \mathrm{X}\}$ does not depend on X appears to be doubtful. Rather, the scatterplot shows that there is more variability in Y when X is small than when X is large.

| X | SD |
| :--- | :--- |
| .00 | .21 |
| .06 | .28 |
| .12 | .11 |
| .30 | .06 |

12.47: The hypotheses are

H0: sulfur dioxide has no effect on yield (beta1 $=0$ ) and
HA: Increasing sulfur dioxide tends to decrease yield (beta1 < 0).
The sample slope is b1 $=-.342 / .1512=-2.262$
We note that b1 $<0$, so the data do deviate from H 0 in the direction specified by HA.

The residual standard deviation is s_\{Y|X\} = sqrt[SS(resid)/(n-2)] = sqrt[.2955/10] = .1719 kg .

The standard error of the slope is SE_\{b1\} = .1719/sqrt[.1512] = . 4421 .
The test statistic is ts $=(\mathrm{b} 1-0) /$ SE_ $\{\mathrm{b} 1\}=-2.261 / .4421=-5.12$.
Consulting Table 4 with $\mathrm{df}=\mathrm{n}-2=10$, we find that P -value $<.0005$, so we reject H 0 .
There is strong evidence (P-value $<.0005$ ) to conclude that increasing sulfur dioxide tends to decrease yield.
12.54: SE_\{b1\} = s_\{Y|X\}/sqrt[n-1]s_X = .0374/sqrt[506667] $=.0000525$

So $95 \% \mathrm{CI}$ is $.0003186+/-(2.228)(.0000525) \quad(\mathrm{df}=10)$ or $(.00020, .00044)$ or $.00020<$ beta $1<.00044$.

### 12.59

## Regression Analysis

| Predictor | Coef | StDev | T | P |
| :---: | :---: | :---: | :---: | :---: |
| Constant | 156.95 | 11.87 | 13.22 | 0.000 |
| dose | -23.580 | 7.358 | -3.20 | 0.009 |
| $S=26.01$ | $\mathrm{R}-\mathrm{Sq}=$ |  | adj) = | 7\% |

Analysis of Variance

| Source | DF | SS | MS | F | P |
| :--- | ---: | ---: | ---: | ---: | ---: |
| Regression | 1 | 6950.2 | 6950.2 | 10.27 | 0.009 |
| Residual Error | 10 | 6766.7 | 676.7 |  |  |
| Total | 11 | 13716.9 |  |  |  |

12.59a: b0 = 156.95; b1 = -23.580 (from SOCR, http://www.socr.ucla.edu printout)

The fitted regression line is $y$-hat $=156.95-23.580 \mathrm{x}$
12.59b:

12.59d: H0: Amphetamine dose has no affect on water consumption (beta1 $=0$ )

HA: Increasing amphetamine dose tends to reduce water consumption (beta1 < 0)
ts $=-3.20$ (SOCR, http://www.socr.ucla.edu printout) and the P -value $=0.009 / 2=0.0045$. Thus we reject H0.

There is strong evidence $(\mathrm{P}$-value $=0.0045)$ to conclude that increasing amphetamine dose tends to reduce water consumption.

### 12.59e: One-way Analysis of Variance

| Analysis of Variance for water-co |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Source | DF | SS | MS | $\begin{array}{r} F \\ 4.65 \end{array}$ | P |  |  |
| dose | 2 | 6972 | 3486 |  | 0.041 |  |  |
| Error | 9 | 6745 | 749 |  |  |  |  |
| Total | 11 | 13717 |  |  |  |  |  |
|  |  |  |  | Individual 95\% CIs For Mean |  |  |  |
| Level | N | Mean | StDev | --+---- | - -+ |  | -+--- |
| - |  |  |  |  |  |  |  |
| 0.00 | 4 | 156.00 | 25.32 |  |  |  | --) |
| 1.25 | 4 | 129.38 | 27.85 |  | (- | - |  |
| 2.50 | 4 | 97.05 | 28.84 | (---- | *- |  |  |
| - |  |  |  |  |  |  |  |
| Pooled | StDev = | 27.38 |  | 70 | 105 | 140 | 175 |

H0: The three doses produce the same mean water consumption level (mu1 = mu2 = mu3)
HA: The mean water consumption levels are not all equal (the mu's are not all equal)

Fs $=4.65$ (SOCR, http://www.socr.ucla.edu printout) and the P-value $=0.041$. Note HA cannot be directional because there are three doses). Thus we reject H0.

The conclusion here is similar to that in part (d), in that we reject H0. However, the analysis from (d) gave a smaller P-value, as it made use of the fact that the means are not only different, but they decrease as dose increases.
12.59f: The analysis in part (d) requires linearity; that is, the mean water consumption levels must have a linear relationship to dose for the regression model to make sense. The ANOVA in part (e) does not require this condition.
12.59g: s_\{pooled $\}=27.38$ (from ANOVA printout), which is similar to s_\{Y|X\}=26.01 (from regression printout)

