Q1 a) Let B; be the slope in the linear relationship between 'Alue' and 'NT3'. We are testing

Ho: B; = 0 against H; B; = 0

The test statistic is

t = 0.639 - 0 = 3.54

We compare the test statistic with the to-distribution.

P(T6 > 3143) = 0.01 and P(T6 > 3.707) = 0.005

P-value = 2 × min {P(T6 > 3.54), P(T6 (3.54)}

which is between 0.01 and 0.02. This is moderate evidence against the null hypothesis, suggesting a linear association between 'Alive' and 'NT3'.

b) The sum of the residuals is zero, since

$$\hat{e}_i = q_i - \hat{\beta}_o - \hat{\beta}_i x_i$$

and

$$\hat{Z} \hat{e}_i = \hat{Z} (y - \hat{\beta}_0 - \hat{\beta}_i x_i) = 0$$

(This is one of the equations we solve to get the least squares estimates $\hat{\beta}_0$, and $\hat{\beta}_1$, see page 130 of notes.)

c) We need an estimate of the mean percentage of alive cells for NT3 concentration 25g/ml and the Standard error.

estimated mean at 25g/ml of NT3 = 55.04 + 0.639 x 25 = 71.015.

The standard error of this mean is computed by taking the square root of

~ 7.3520

s.e. (estimate) = 17-3520 = 2.7115

The 95% confidence interval is formed by

estimate ± (critical-value) × s.e. (estimate)

= 71.015 ± t_{6;0.975} × 2.7115

= 71-015 = 2.447 x 2.7115

= 71-015 ± 6.6349 (percent)

We are 95% confident that the true mean percentage of Alive cells at NT3 concentration of Dryml is between 64.38% and 77.65%

- d) The assumptions of the linear regression model are
 - linearity of mean in explanatory variable
 - constant variance
 - normality
 - independence.

Departures in the first three assumptions can be detected with these plots. There is no obvious trend or change in the spread of residuals in the residuals a fitted values plot, so first two assumptions are ok. The normal probability plot appears straight which is consistent with the normality assumption.

(92 a) Let p be the correlation between CAA concentration and initial bigilm thathers.

Test Ho: p=0 against H; = p +0.

Test statistic $r-c_0 = \frac{-0.3508}{\sqrt{(1-r^2)/(n-2)}} = \frac{-1.5893}{\sqrt{1-(-0.3508)^2}}$

Under the null hypothesis, the test stedistic has a tn-2 distributions

p-value = 2 × min (P(T15 > -1-5893), P(T15 5-1-5893)}

P(T18 5-1-5893) = P(T18 > 1-5893) by symmetry

P(T18 > 1-330) = 0.1 and P(T18 > 1.734) = 0.05

So the p-value is between 0.1 and 0.2. This is inconclusive evidence against the null, suggesting no correlation between CAA concentrations and initial brightn thickness.

- b) Units are um/(ug/ml)
- c) estimated mean reduction in thickness is

5.002 + 0.02956 × 75 = 7-2190 um

d) Let B, be the true slope in the linear relationship between biofilm thickness reduction and CAA consentration. We test

Ho: β,=0 against H,; β, \$0

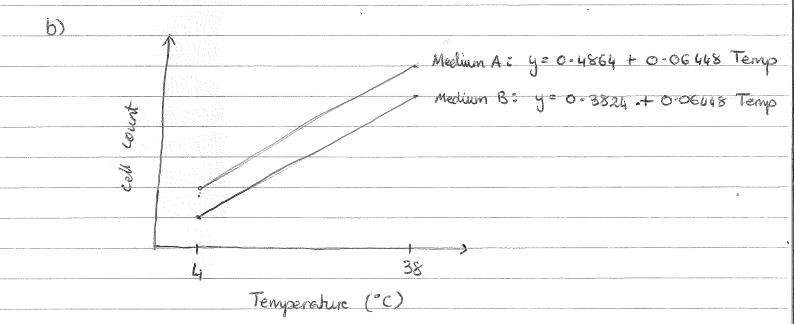
Test Stahishic

t = 0.02956 = 1.5476

p-value = 2 x min {P(T, > 1-5476), P(T, \$1-5476)}

As $P(T_{18} > 1.336) = 0.1)$ and $P(T_{18} > 1.734) = 0.05$, the p-value is between 0.1 and 0.2. There is inconclusive evidence of a linear association between CAA concentrations and the reduction in biofilm thickness.

Q3 a) The error degrees of freedom is 23 = n - p. The number of terms in the linear regression is 3 (intercept, temp, medium B). So the number of observations/experimental trials used in the analysis is 23 + 3 = 26.



e) We are testing whother the coefficient for temperature in the linear relationship is zero. Denote this by \$1. We test

Ho: β, = 0 against H, = β, ±0

Test stanstic t = 0.064485 = 39.978 0.601613

We compare this to the t23-distribution

P-value = 2 × min { P(T23 > 39.978), P(T23 6 39-978)}

P(T23 >4.415) =0-0001

So the p-value is less than 0.0002. This is very strong evidence against the null hypothesis, suggesting a linear association between cell count and temperature.

d) Confidence interval: estimate ± (critical value) × 5.e. (estimate)

-0.104045 ± t23;0.95 × 0.040080

t23;0.95 = 1.714 (from tables). So

 $-0.104045 \pm 1.714 \times 0.040080$ $= -0.104045 \pm 0.0687$

We are 90% confidence the true coefficient for Medium B is between -0.1727 and -0.0353.

e) (Estimated mean cell count) = 0.4864 + 0.664485 × 30

(at 30°C in Medium A

= 2-4209 (x106 cells).

f) We have the estimated means for medium A at 30°C from part (e). We now need the Standard error. The solimate for σ is 0.1267 (Root Mean Square Error in output), we need to compute $x(X^TX)^{-1}x^T$ where x=(1300].

[1 30 0] 0.1223 2.143e-05 5.334e-04 [1]
2.143e-05 1.6207e-04 1.186e-05 30
5-334e-04 1.186e-05 0.1001 0

 $= \begin{bmatrix} 0.1229 & 0.0049 & 0.0009 \end{bmatrix} \begin{bmatrix} 1 \\ 36 \\ 0 \end{bmatrix}$

The standard error for the estimated mean for medium A at 30°C is 0-1267 × Jo-2694' = 0-0658.

The 95% confidence interval is

2.4209 ± ±23;0.975 x 0.0658 = 2.4209 ± 2.069 x 0.0658 = 2.4209 ± 0.1361 (x10° cells)

Q4 a) We assume permax ~ N(βo+β; weight, 6²) and that
the individuals are independent. Note that this includes

- assumption of lihearity of mean in explanatory variable

- assumption of constant variance

- assumption of Normality.

b) (A) = Estimate = 63.5456 = 12.7015 [From rearranging] +Stat = 5.003 [tstat = Estimate] SE

- (C) Error degrees of freddom = N-2 = 25-2=23
 - (B) p-value = 2 × min {P(T23 > 3.944), P(T23 < 3.944)}

As $P(T_{23} > 3.768) = 0.0005$ and $P(T_{23} > 4.415) = 0.0001$, the p-value is between 0.0002 and 0.001

- c) estimated mean persex = 63.5456 + 1.1867 × 50 = 122.8806 (units)
- d) $R^2 = r^2 = 0.635^2 = 0.4032$.
- e) Let β, denote the coefficient of weight in the linear relationship between permax and weight.

Test Ho: β, = 0 egainst H, = β, +0

The test statistic is 3.944 and p-value between 0.0002 and 0.001 (from Part b). This is strong evidence against the null hypothesis, suggesting a linear association between weight and persons.

f) The normal probability plot is relatively straight indicating normal distrabution for residuals. The residual is fitted value shows some fenning/increasing variability as fitted values suggesting the assumption of constant variance is violated. There is no clear trend in the residual plot so linearity assumption maybelse is ok.

- Q5 a) Error degrees of freedom = 11-2 = 15 so there were 17 soil samples
 - b) Pp = Bo + B, Pi + U, where U~ N(0,02).
 - c) The estimates of intercept and slepe are 62.57 and 1.23, respectively. The estimated mean availability of phosphorous at 17 ppm of inorganic phosphorous is

62.57 +1=23×17 = 83.48 (ppm)

d) tStat = Estimate = 1-2291 = 4.019294 (A) SE = 0.3058

p-value = 2x min {P(T,5 > 4.01924), P(T,5 < 4-01924)} = (B).

As $P(T_{15} > 3.733) = 0.001$ and $P(T_{15} > 4.073) = 0.0005$, The p-value is between 0.001 and 0.002.

e) Let β; be the true slope in the linear relationship between inorganic phosphorous and available phosphorous.

Test Ho: B, = 1 against H, = B, \$1

Test statistic t = estimate - hypothesised 5.e (estimate)

 $= \frac{1.9291 - 1}{0.3058} = 0.74918$

p-value = 2x min {P(T,5 > 0.74918), P(T,5 50.74918)}

As $P(T_p > 0.691) = 0.25$ and $P(T_{15} > 1.341) = 0.1$, the p-value is between 0.2 and 0.5.

There is no evidence against the null hypothesis, suggesting β , is equal to one.

- f) residual = observation estimated mean = $51 - (62.5694 + 1.2291 \times 12.6)$ = 51 - 178.0561 = -27.0561.
 - g) Normal probability plot to check the assumption of normality. Plot residuals egainst fitted values to check constraint variance and linearity (residuals should diaplay no hend.
 - h) The matrix is symmetric so (E) = 0.2763 $(D) = (SE of intercept)^2 = 4.4519^2 = 19.81941$ $(F) = (SE of slope)^2 = 0.3058^2 = 0.093514$