

# Problem Set 11

## (Vinay Kumar Ranganath Babu)

### # Problem1

```
observed = c(29,19,18,25,17,10,15,11)
expected = rep(sum(observed)/length(observed),8)
X2 = sum((observed-expected)^2/expected)
p.value = 1 - pchisq(X2,df=7)
```

| Values   |                                   |
|----------|-----------------------------------|
| expected | num [1:8] 18 18 18 18 18 18 18 18 |
| observed | num [1:8] 29 19 18 25 17 10 15 11 |
| p.value  | 0.0222394774623906                |
| X2       | 16.3333333333333                  |

Since our p-value is on the lower side of 0.02, we can reject the null hypothesis. Hence horse's starting position does not affect its chance of winning.

### # Problem2

```
observed = c(121,84,118,226,226,123)
prapos = c(0.13,0.14,0.13,0.24,0.20,0.16)
expected = prapos * sum(observed)
X2 = sum((observed-expected)^2/expected)
p.value = 1 - pchisq(X2,df=length(observed)-1)
```

| Values   |  |
|----------|--|
| expected | num [1:6] 117 126 117 216 180 ...      |
| observed | num [1:6] 121 84 118 226 226 123       |
| p.value  | 1.860202696502e-05                     |
| prapos   | num [1:6] 0.13 0.14 0.13 0.24 0.2 0.16 |
| X2       | 29.4874007766547                       |

Since our p-value is too small, we can reject the null hypothesis. Hence the claimed proportions are not credible in light of the provided data.

### # Problem3

n=1000

observed=c(30, 93, 159, 184, 195, 171, 92, 45, 31)

expected=c(sum(dbinom(0:1, 16, .29)\*n),dbinom(2, 16, .29)\*n,dbinom(3, 16, .29)\*n,dbinom(4, 16, .29)\*n,dbinom(5, 16, .29)\*n,dbinom(6, 16, .29)\*n, dbinom(7, 16, .29)\*n,  
dbinom(8, 16, .29)\*n, sum(dbinom(9:16, 16, .29)\*n))

X2=sum((observed-expected)^2/expected)

G2=2 \* sum(observed \* log(observed/expected))

p.valueX = 1-pchisq(X2, 1)

p.valueG = 1-pchisq(G2, 1)

| Values   |   |
|----------|---|
| expected | num [1:9] 31.4 83.5 159.1 211.2 207.1 ... |
| G2       | 11.9693183070806                          |
| n        | 1000                                      |
| observed | num [1:9] 30 93 159 184 195 171 92 45 31  |
| p.valueG | 0.000540837238129144                      |
| p.valueX | 0.00038265779331359                       |
| X2       | 12.6150280615715                          |

The Test static for LR is 11.96932 and P-value 0.000540

The Test static for Pearson's is 12.6150 and P-value 0.00038

Hence we can reject null hypothesis that the researchers' coloring algorithm was not performed as intended.

### # Problem4

a)  $\log(x+1) - \log(x)$

$\log(2) - \log(1) + \log(3) - \log(2) + \dots + \log(10) - \log(9) - \log(1) + \log(10)$

= 1

Hence verified.

b) observed = c(107,55,39,22,13,18,13,23,15)

n = sum(observed)

p = log10(1+1/(1:9))

expected = n\*p

X2 = sum((observed-expected)^2/expected)

p.valueX = 1-pchisq(X2 , 8)

# using likelihood ratio

G2 = 2 \* sum(observed \* log(observed/expected))

p.valueY = 1-pchisq(G2 , 8)

| Values   |   |
|----------|---|
| expected | num [1:9] 91.8 53.7 38.1 29.6 24.2 ...          |
| G2       | 15.5558864933756                                |
| n        | 305   |
| observed | num [1:9] 107 55 39 22 13 18 13 23 15           |
| p        | num [1:9] 0.301 0.1761 0.1249 0.0969 0.0792 ... |
| p.valueX | 0.0639909378017312                              |
| p.valueY | 0.0491962228990812                              |
| X2       | 14.7596477007757                                |

We get values of 0.049 approximately 0.05.

This is not very clear, however we can say leading digits hold Bedford's law and doesn't say how the law can't be too sure with this one.

### **# Problem5**

observed = c(173,125,150,73)

dumm = sum(observed)

expected = c((323\*298/dumm),(198\*298/dumm),(323\*223/dumm),(198\*223/dumm))

```
X2 = sum((observed-expected)^2/expected)
```

```
p.valueX = 1 - pchisq(X2,df=1)
```

```
G2 = 2* sum(observed * log(observed/expected))
```

```
p.valueG = 1 - pchisq(G2,df=1)
```

| Values   |                                  |
|----------|----------------------------------|
| dumm     | 521                              |
| expected | num [1:4] 184.7 113.3 138.3 84.7 |
| G2       | 4.62306318005088                 |
| observed | num [1:4] 173 125 150 73         |
| p.valueG | 0.0315448566517489               |
| p.valueX | 0.0321033323560087               |
| X2       | 4.59297035792607                 |

P-value is less than 0.05 hence we reject the null hypothesis, the panama sandflies do not vary with height above ground.

### **# Problem6**

```
positive = c(74,68,154,18)
```

```
partial = c(18,16,54,10)
```

```
none = c(12,12,58,44)
```

```
n.positive = sum(positive)
```

```
n.partial = sum(partial)
```

```
n.none = sum(none)
```

```
n = n.positive + n.partial + n.none
```

```
disease = (positive + partial + none) / n
```

```
positive.expected = disease * n.positive
```

```
partial.expected = disease * n.partial
```

```
none.expected = disease * n.none
```

```
X2 = sum((positive-positive.expected)^2/positive.expected,(partial-
partial.expected)^2/partial.expected,(none-none.expected)^2/none.expected)
```

```
p.value = 1 - pchisq(X2,df=3*2)
```

| Values            |                                   |
|-------------------|-----------------------------------|
| disease           | num [1:4] 0.193 0.178 0.494 0.134 |
| n                 | 538                               |
| n.none            | 126                               |
| n.partial         | 98                                |
| n.positive        | 314                               |
| none              | num [1:4] 12 12 58 44             |
| none.expected     | num [1:4] 24.4 22.5 62.3 16.9     |
| p.value           | 2.52020626589911e-14              |
| partial           | num [1:4] 18 16 54 10             |
| partial.expected  | num [1:4] 18.9 17.5 48.5 13.1     |
| positive          | num [1:4] 74 68 154 18            |
| positive.expected | num [1:4] 60.7 56 155.2 42        |
| X2                | 75.8901487931342                  |

Test static is 75.9 and P-value 2.5e-14. Hence we reject null hypothesis- patient's response to treatment for Hodgkin's disease does not vary by histological.