## Analyzing Categorical Data using the Pearson Statistic and R

Consider a multinomial experiment with k = 4 possible outcomes. After n = 100 trials are performed, we observe the following counts of each outcome.

Suppose that we wanted to determine whether the probabilities of each outcome differ; this analysis will echo that of Example 14.1. More precisely, we want to test the null hypothesis  $H_0: p_1 = p_2 = p_3 = p_4 = 0.25$ . To do so, we use the Pearson test statistic  $X^2$  with approximate  $\chi^2$  distribution; in this case,  $X^2$  has k-1 degrees of freedom since the only assumption used about the outcome probabilities is that  $p_1 + p_2 + p_3 + p_4 = 1$ .

We can calculate  $X^2$  using the following code.

```
# Set outcome probabilities under HO.
p <- rep(0.25, 4)

# Define vector of observed counts.
y <- c(20, 17, 31, 32)
n <- sum(y)

# Calculate X^2.
Xsq = sum((y - p*n)^2/(p*n))</pre>
```

This gives  $X^2 = 6.96$ .

We have an  $\alpha$ -level test if we reject  $H_0$  when  $X^2 > \chi^2_{\alpha,k-1}$ . For  $\alpha = 0.05$  and k-1=3, we have the rejection region

$$X^2 > 7.815$$
.

In this case, we cannot reject  $H_0$  since the observed value of  $X^2$  is outside the rejection region.

## Using the chisq.test function

We could have also performed the test using the chisq.test function, as in the following code.

```
chisq.test(y)
```

```
##
## Chi-squared test for given probabilities
##
## data: y
## X-squared = 6.96, df = 3, p-value = 0.07318
```

When called, chisq.test(y) calculates the value of  $X^2$  for the observed counts stored in the vector y, as well as the observed attained significance level. Here, we have p = 0.07318 which indicates that we cannot

Outcome	Observed Count
1	20
2	17
3	31
4	32

reject  $H_0$  (using  $\alpha = 0.05$ ).