## **Lesson 9.1.3**

9-21. <u>Identify</u>: Chi-squared goodness of fit test.  $H_0$ : The outcomes on the die each occur with  $\frac{1}{6}$  probability.  $H_A$ : At least one outcome differs from the expected frequency. Sample evidence for  $H_A$ : observed frequencies do not match expected frequencies.  $\alpha = 0.05$ . <u>Check conditions</u>: It can be assumed that this data represents a SRS of all possible rolls. Independent trials and large counts, so the sampling distribution is  $\approx$  chi-squared. Each outcome is independent. Additionally, all expected counts are 100, well above five. <u>Calculate</u>:  $\chi^2 = \text{sum}(0.09, 0.49, 0.01, 0.16, 0.64, 0.81) = 2.2$  with a *p*-value of 0.8208 (df = 5).

<u>Conclude</u>: Because the p-value is large (much higher than 5%), students do not have sufficient evidence to reject  $H_0$ . It cannot be concluded that the die is unfair.

9-22. <u>Identify</u>: Chi-squared goodness of fit test.  $H_0$ : The traits occur in the expected 9:3:3:1 ratio.  $H_A$ : At least one trait differs from the expected frequency. Sample evidence for  $H_A$ : observed frequencies do not match expected frequencies.  $\alpha = 0.05$ . <u>Check conditions</u>: Random sample, to avoid bias. It can be assumed that this data represents a SRS of seeds. Independent trials and large counts, so the sampling distribution is  $\approx$  chi-squared. Each outcome is independent. All expected counts (144, 48, 48, 16) are above 5.

<u>Calculate</u>:  $\chi^2 = \text{sum}(0.6944, 0.1875, 1.6875, 1) = 3.5694$  with a *p*-value of 0.3119 (df = 3).

<u>Conclude</u>: Because the p-value is large (much higher than 5%), students do not have sufficient evidence to reject  $H_0$ . It cannot be concluded that the observed traits differ significantly from the expected frequencies predicted by Mendelian genetics.

9-23. a. <u>Identify</u>: Chi-squared goodness of fit test.  $H_0$ : The current city population remains in the same proportion as the historical census.  $H_A$ : At least one continent now differs in its representation in this city. Sample evidence for  $H_A$ : Observed frequencies do not match expected frequencies.  $\alpha = 0.05$ . <u>Check conditions</u>: Random sample, to avid bias. It is stated that this sample is a

SRS. Independent trials and large counts, so the sampling distribution is  $\approx$  chi-squared. The individuals can be considered independent because 500 is much less than 10% of the population of a "large city." All expected counts (65, 60, 140, 45, 175, 15) are above 5.

<u>Calculate</u>:  $\chi^2 = \text{sum}(12.0615, 0.15, 0.4571, 0.0222, 1.4629, 0.2667) = 14.4204$  with a *p*-value of 0.0131 (df = 5).

<u>Conclude</u>: Because the *p*-value is less than 5%, students have sufficient evidence to reject  $H_0$ . It be concluded that there <u>has</u> been a shift in the representation of at least one continent.

b. The continent of Asia had a contribution of 12.0615.

5

9-24. <u>Identify</u>: Chi-squared goodness of fit test.  $H_0$ : The digits in the decimal representation of  $\sqrt{2}$  each occur with approximately a  $\frac{1}{10}$  relative frequency.  $H_A$ : At least one digit differs from the expected frequency. Sample evidence for  $H_A$ : observed frequencies do not match expected frequencies.  $\alpha = 0.05$ .

<u>Check conditions</u>: Random sample, to avoid bias. Any bias found is evidence for  $H_A$ . Independent trials and large counts, so the sampling distribution is  $\approx$  chi-squared. It can be assumed that these digits are all independent of one another. All expected counts (1000 each) are well above 5.

Calculate:  $\chi^2 = \text{sum}(2.304, 0.025, 0.016, 0.4, 0.256, 0.001, 1.024, 1.296, 0.729, 0.361) = 6.4120$  with a *p*-value of 0.6981 (df = 9).

Conclude: Because the *p*-value is high (much higher than 5%) we, do not have sufficient evidence to reject  $H_0$ . It cannot be concluded that the digits of  $\sqrt{2}$  differ from the uniform distribution.

- 9-25. a. Expected value: \$3.802, standard deviation: \$5.425
  - b. \$599,173.60
  - c. <u>Identify</u>: Chi-squared goodness of fit test.  $H_0$ : The state is producing tickets at the advertised ratios.  $H_A$ : The state is not producing tickets at the advertised ratios. Sample evidence for  $H_A$ : observed frequencies do not match expected frequencies.  $\alpha = 0.05$ .

<u>Check conditions</u>: The sample was stated to be randomly gathered. Independent trials and large counts, so the sampling distribution is  $\approx$  chi-squared. It is assumed that there were more than 500(10) tickets available. The sample size of 500 was large enough that all expected counts but one were larger than 5.

Calculate:  $\chi^2 = 19.9975$ , df = 4, p-value = 0.0005.

<u>Conclude</u>: Because the p-value is so small (less than 5%), reject  $H_0$ . The evidence indicates that the state is not producing the tickets at the advertised ratios.

9-26. a. <u>Identify</u>: One-sample confidence interval of proportions.

<u>Check conditions</u>: Random, to avoid bias: SRS assumed, independent trials for an accurate  $\sigma$ : 297(10) < lake fish population assumed, large counts, so the sampling distribution is  $\approx$  normal: np = 14 and 14 > 10.

<u>Calculate</u>: SE = 0.0123, find  $z^* = 1.96$ , ME = 1.96(0.01213) = 0.0241, confidence interval =  $0.0471 \pm 0.0241$ ,  $\{0.0230, 0.0712\}$ .

<u>Conclude</u>: We are 95% confident that the interval from 0.0230 to 0.0712 contains the true population proportion of tagged fish in the lake.

- b.  $\frac{150}{0.0230} = 6512$ ,  $\frac{150}{0.0712} = 2106$ . The interval (2106 to 6512) seems too wide to be of much use.
- c. Increase the sample size of the fish and/or lower the confidence level.

9-27. See frequency table below. Multiply row by column to get numbers for events A, B, C, and D. Car alarms will sound in events A or D. That would be 99 + 8205 = 8304 alarms per year. Of those alarms only the 99 from event A are actual break-ins. If you hear a car alarm in Madelinton the probability the car is being broken into is just  $\frac{99}{8304} \approx 0.0119$ .

Alarm is correct 0.99 Alarm is *not* correct 0.01

100 are broken into	820,500 are <i>not</i> broken into
(A) 99	(B) 812,295
(C) 1	(D) 8205 false alarms