AP Statistics
Unit 08 – Day 01 Notes
Chi Square: Goodness of Fit Test

Name_	Key	
Period		

We use this type of Inference test when we want to examine the distribution of a single categorical variable in a population!

One-way table: a table used to display the distribution of a single categorical variable (ex: M&M color)

Goodness of Fit: a test that tests the null hypothesis that a categorical variable has a specified distribution in the population of interest

(ex: 20%, yellow, 25%, red, 15%, blue,

Observed Value: the actual # of individuals in the sample that fall in each cell of the one-way table (ex: 6 yellow M&Ms in)

Expected Value: the expected # of individuals in the

Sample that would fall in each cell of the one-way table if Ho The Chi-Square Distribution and P-Values Were true (ex: poin for each color of MAM

The sampling distribution of the chi-square statistic is **NOT** a **normal distribution**Yellow: 0, 2 - 24 = 4.8 yellow
MdNAC expects It is a **right-skewed** distribution that allows only **nonnegative values** ( $\chi^2$  = can't have a

negative number when you square) Large counts condition/

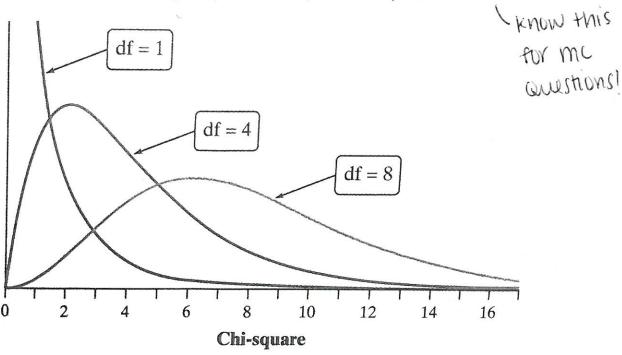
• When the expected counts are all at least 5, the sampling distribution of  $x^2$  statistic is modeled well by a chi-square distribution (you must list them all when checking conditions)

Chi-square distributions are a family of density curves (Kind of like t-distributions)

Chi-square distribution with degrees of freedom = the number of categories - 1 (not sample size)

A particular chi-square distribution is specified by giving its degrees of freedom (see picture below)

When df > 2, the mode (peak) of the chi-square density curve is df - 2



#### STATE:

 $H_o$  = the stated distribution of the <u>categorical variable</u> in the population of interest is correct.

 $H_A$  = the stated distribution of the categorical variable in the population of interest is incorrect.

#you can write these using #s but it takes foreverso words are easier :

### PLAN:

Random: The data must come from a random sample or randomized experiment.

10%: The sample must be less than 10% of the population if we are sampling without replacement

Large Counts: All counts must be 5 or more. YOU MUST WRITE THEM ALL OUT. TOO bad:

State Test: Because our conditions are met, we will perform a chi-square test for goodness of fit .

#### DO:

# Chi-square statistic

$$\chi^2 = \sum_{all \ cells} \frac{(observed - expected)^2}{expected}$$

- work shown for the above formula (at least 3 terms +...)
- degrees of freedom df =
- $\chi^2 =$
- P-value =

You have to show all of the stuff even if you do the test on your calculator. No need to put in what you actually enter into your calculator this time (and thank goodness because that would basically require writing out the lists you entered).

### CONCLUDE:

- Large values of  $\chi^2$  (the test statistic) are evidence against the  $H_o$  and in favor of the  $H_A$ .
- The P-value is the area to the right of the  $\chi^2$  under the chi-square distribution with degrees of freedom (df = number of categories 1). YOU (an find this using  $\chi^2$  (df ( , ) in the dismbutions menu (#8).

We either will reject the null or fail to reject the null and explain in context why. We can use the same sentence frames as before:

Because our p-value  $\_\_\_$  is (greater than/less than) our significance level =  $\_\_$ , we (reject/fail to reject) the null. There (is/is not) convincing evidence that (alternative hypothesis in context).

## ON THE CALCULATOR:

- 1. Enter your data into lists:
  - a. STAT → Enter
    - i. Observed Values in L1
    - ii. Expected Values in L2
- 2. Choose the appropriate test:
  - a. STAT → TESTS
  - b. Option D:  $\chi^2$ GOF-Test
    - i. df: # categories 1
    - ii. Calculate

## AND/OR

- 1. Calculate the test statistic by hand and then use the calculator to find the p-value:
  - a. 2<sup>ND</sup> VARS
  - b. Option 8:  $\chi^2$  cdf(
    - i. lower: your  $\chi^2$  value
    - ii. upper: 10,000 (a very large number since it is skewed to the right)
    - iii. df: # categories 1

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#### **EXAMPLE 1:**

Baseball is a remarkable sport, in part because so much data are available. We have the birth dates of every one if the 16,804 players who ever played in a major league game. Since the effect we're suspecting may be due to relatively recent policies (and to keep the same size moderate), we'll consider the birth months of the 1478 major league players born since 1975 to 2006. We can also look up the national demographic statistic to find what percentage of people who were born in each month. Let's test whether the observed distribution of ball players birth months shows just random fluctuations or whether it represents a real deviation from the national pattern.

Month	Ballplayer Count	National Birth %	Month	Ballplayer Count	National Birth %
1	137	8%	7	102	9%
2	121	7%	8	165	9%
3	116	8%	9	134	9%
4	121	8%	10	115	9%
5	126	8%	11	105	8%
6	114	8%	12	122	9%
0			TOTAL	1478	100%

STATE:

H<sub>o</sub>: The distribution of birthdays of all MLB players is the same as the gen. pop.  $H_A$ : The distribution of birthdays of all MLB players is not the (national patern) same of the general population.

PLAN:

Random: all players between 1975 and 2006

10% condition: n=1478 14780<16804 players e verin MLB

Large Counts: (this part sucks)

L) Find the expected values for each category (month)

Month	Expected	Month	Expected	Month	Expected
1	1478 - 0.08 = 118.24	5	118.24	9	133,02
2	103.46	6	118.24	10	133,02
3	110 24	7	133, 02	11	118.24
4	118 14	8	133.07	12	133.02

all expected count are >5.

Because our conditions are met... We will perform a  $\chi^2$  HST for goodness of fit.

DO: 
$$\chi^2 = \frac{(137 - 118.24)^2}{118.24} + \frac{(121 - 103.46)^2}{103.46} + \frac{(116 - 118.24)^2}{118.24} + \dots = 26.48$$

$$df = 12 - 1 = 11$$

$$x^2 = 26.48$$

P-value = 0.00549

conclusion: Because our p-value = 0.00549 is less than our significance this level d=0.05, we reject the null. There is convincing evidence that the distribution of MLB players' birndays deviates from the national pattern.

#### **EXAMPLE 2:**

We have counts of 256 randomly selected executives in 12 zodiac sign categories. The natural null hypothesis is 'hat birth dates of executives are divided equally among all the zodiac signs. The test statistic looks at how closely the observed data math this idealized situation. Are zodiac signs of CEO's distributed uniformly?

Births		Signs
Observed Values	Expected Values	
23	256. Y12 = 21.3	Aries
20	21.3	Taurus
18	21.3	Gemini
23	21.3	Cancer
20	21.3	Leo
19	21.3	Virgo
18	21.3	Libra
21	21.3	Scorpio
19	21.3	Sagittarius
22	21.3	Capricorn
24	21.3	Aquarius
29	21.3	Pisces
TOTAL = 256		

State: Ho: Birthdates of executives are evenly distributed across oul zodiac signs.

HA: Birtholates of executives are not evenly distributed across our zodiac signs.

d= 0.05.

Plan: random: 256 randomly selected executives ver 10% condition: n=256 2560 < all executives ever Large counts: if birth dates are evenly distributed across all 12 zodiacs in the year, then the expected count for each zodiac is 256. Y12 = 21.3 executives > 5. Decause our conditions are met, we will perform a 12 test for goodness of fit.

$$\underline{D0:} \quad \chi^2 = \frac{(23 - 21.3)^2}{21.3} + \frac{(20 - 21.3)^2}{21.3} + \frac{(18 - 21.3)^2}{21.3} + \dots 5.09$$

df= 11 p-value = 0.9265

conclude: Because our p-value = 0.9265 is greater than our Significance level d=0.05, we fail to reject the null. There is not convincing evidence that the birthdates of executives are not evenly distributed among all 12 zodiac signs.

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