

Lecture 8

8.1. Inferences about Population Variances

Summary

In this lecture we presented statistical procedures that can be used to make inferences about population variances. In the process we introduced two new probability distributions: the chi-square distribution and the F distribution. The chi-square distribution can be used as the basis for interval estimation and hypothesis tests about the variance of a normal population.

We illustrated the use of the F distribution in hypothesis tests about the variances of the normal populations. In particular, we showed that with independent simple random samples of sizes n_1 and n_2 selected from two normal populations with equal variances $\sigma^2 = \sigma^2$, the sampling distribution of the ratio of the two sample variances s_1^2/s_2^2 has an F distribution with $n_1 - 1$ degrees of freedom for the numerator and $n_2 - 1$ degrees of freedom for the denominator.

Inferences about the difference between two population means were then discussed for the matched sample design. In the matched sample design each element provides a pair of data values, one from each population. The difference between the paired data values is then used in the statistical analysis. The matched sample design is generally preferred to the independent sample design because the matched-sample procedure often improves the precision of the estimate.

Finally, interval estimation and hypothesis testing about the difference between two population proportions were discussed. Statistical procedures for analyzing the difference between two population proportions are similar to the procedures for analyzing the difference between two population means.

Key formulas

Interval Estimate of a Population Variance

$$\frac{(n-1)s^2}{\chi_{\alpha/2}^2} \leq \sigma^2 \leq \frac{(n-1)s^2}{\chi_{(1-\alpha/2)}^2} \quad (8.1)$$

Test Statistic for Hypothesis Tests about a Population Variance

$$\chi^2 = \frac{(n-1)s^2}{\sigma_0^2} \quad (8.2)$$

Test Statistic for Hypothesis Tests about Population Variances with $\sigma_1^2 = \sigma_2^2$

$$F = \frac{s_1^2}{s_2^2} \quad (8.3)$$

Lecture supplementary material

FIGURE 11.1 EXAMPLES OF THE SAMPLING DISTRIBUTION OF $(n-1)s^2/\sigma^2$
(A CHI-SQUARE DISTRIBUTION)

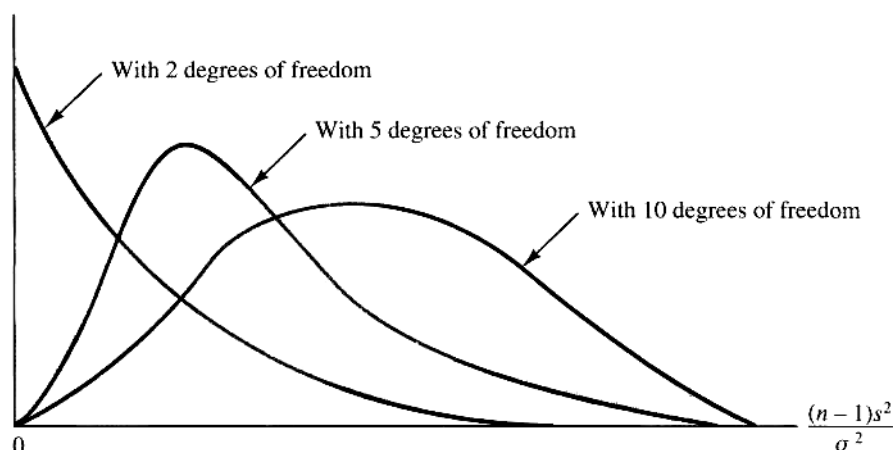
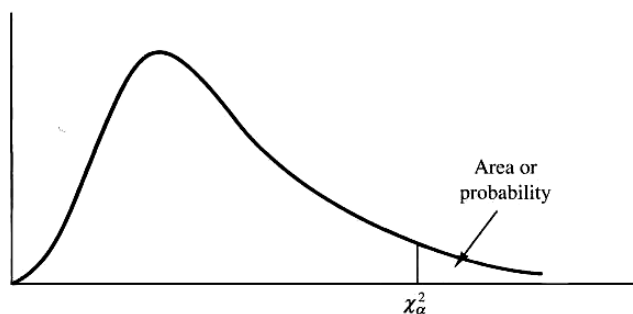


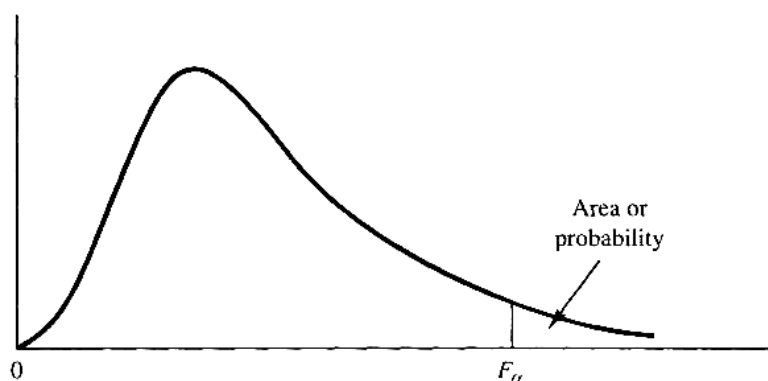
TABLE 11.1 SELECTED VALUES FROM THE CHI-SQUARE DISTRIBUTION TABLE*

| Degrees of Freedom | Area in Upper Tail | | | | | | | |
|-----------------------|--------------------|--------|--------|--------|---------|---------|---------|---------|
| | .99 | .975 | .95 | .90 | .10 | .05 | .025 | .01 |
| 1 | .000 | .001 | .004 | .016 | 2.706 | 3.841 | 5.024 | 6.635 |
| 2 | .020 | .051 | .103 | .211 | 4.605 | 5.991 | 7.378 | 9.210 |
| 3 | .115 | .216 | .352 | .584 | 6.251 | 7.815 | 9.348 | 11.345 |
| 4 | .297 | .484 | .711 | 1.064 | 7.779 | 9.488 | 11.143 | 13.277 |
| 5 | .554 | .831 | 1.145 | 1.610 | 9.236 | 11.070 | 12.832 | 15.086 |
| 6 | .872 | 1.237 | 1.635 | 2.204 | 10.645 | 12.592 | 14.449 | 16.812 |
| 7 | 1.239 | 1.690 | 2.167 | 2.833 | 12.017 | 14.067 | 16.013 | 18.475 |
| 8 | 1.647 | 2.180 | 2.733 | 3.490 | 13.362 | 15.507 | 17.535 | 20.090 |
| 9 | 2.088 | 2.700 | 3.325 | 4.168 | 14.684 | 16.919 | 19.023 | 21.666 |
| 10 | 2.558 | 3.247 | 3.940 | 4.865 | 15.987 | 18.307 | 20.483 | 23.209 |
| 11 | 3.053 | 3.816 | 4.575 | 5.578 | 17.275 | 19.675 | 21.920 | 24.725 |
| 12 | 3.571 | 4.404 | 5.226 | 6.304 | 18.549 | 21.026 | 23.337 | 26.217 |
| 13 | 4.107 | 5.009 | 5.892 | 7.041 | 19.812 | 22.362 | 24.736 | 27.688 |
| 14 | 4.660 | 5.629 | 6.571 | 7.790 | 21.064 | 23.685 | 26.119 | 29.141 |
| 15 | 5.229 | 6.262 | 7.261 | 8.547 | 22.307 | 24.996 | 27.488 | 30.578 |
| 16 | 5.812 | 6.908 | 7.962 | 9.312 | 23.542 | 26.296 | 28.845 | 32.000 |
| 17 | 6.408 | 7.564 | 8.672 | 10.085 | 24.769 | 27.587 | 30.191 | 33.409 |
| 18 | 7.015 | 8.231 | 9.390 | 10.865 | 25.989 | 28.869 | 31.526 | 34.805 |
| 19 | 7.633 | 8.907 | 10.117 | 11.651 | 27.204 | 30.144 | 32.852 | 36.191 |
| 20 | 8.260 | 9.591 | 10.851 | 12.443 | 28.412 | 31.410 | 34.170 | 37.566 |
| 21 | 8.897 | 10.283 | 11.591 | 13.240 | 29.615 | 32.671 | 35.479 | 38.932 |
| 22 | 9.542 | 10.982 | 12.338 | 14.041 | 30.813 | 33.924 | 36.781 | 40.289 |
| 23 | 10.196 | 11.689 | 13.091 | 14.848 | 32.007 | 35.172 | 38.076 | 41.638 |
| 24 | 10.856 | 12.401 | 13.848 | 15.659 | 33.196 | 36.415 | 39.364 | 42.980 |
| 25 | 11.524 | 13.120 | 14.611 | 16.473 | 34.382 | 37.652 | 40.646 | 44.314 |
| 26 | 12.198 | 13.844 | 15.379 | 17.292 | 35.563 | 38.885 | 41.923 | 45.642 |
| 27 | 12.878 | 14.573 | 16.151 | 18.114 | 36.741 | 40.113 | 43.195 | 46.963 |
| 28 | 13.565 | 15.308 | 16.928 | 18.939 | 37.916 | 41.337 | 44.461 | 48.278 |
| 29 | 14.256 | 16.047 | 17.708 | 19.768 | 39.087 | 42.557 | 45.722 | 49.588 |
| 30 | 14.953 | 16.791 | 18.493 | 20.599 | 40.256 | 43.773 | 46.979 | 50.892 |
| 40 | 22.164 | 24.433 | 26.509 | 29.051 | 51.805 | 55.758 | 59.342 | 63.691 |
| 60 | 37.485 | 40.482 | 43.188 | 46.459 | 74.397 | 79.082 | 83.298 | 88.379 |
| 80 | 53.540 | 57.153 | 60.391 | 64.278 | 96.578 | 101.879 | 106.629 | 112.329 |
| 100 | 70.065 | 74.222 | 77.929 | 82.358 | 118.498 | 124.342 | 129.561 | 135.807 |

*Note: A more extensive table is provided as Table 3 of Appendix B.

Table . Summary of hypothesis test about a population variance

| | Lower Tail Test | Upper Tail Test | Two-Tailed Test |
|--|---|---|--|
| Hypotheses | $H_0 : \sigma^2 \geq \sigma_0^2$ $H_a : \sigma^2 < \sigma_0^2$ | $H_0 : \sigma^2 \leq \sigma_0^2$ $H_a : \sigma^2 > \sigma_0^2$ | $H_0 : \sigma^2 = \sigma_0^2$ $H_a : \sigma^2 \neq \sigma_0^2$ |
| Test Statistic | $\chi^2 = \frac{(n-1)s^2}{\sigma_0^2}$ | $\chi^2 = \frac{(n-1)s^2}{\sigma_0^2}$ | $\chi^2 = \frac{(n-1)s^2}{\sigma_0^2}$ |
| Rejection Rule: p-Value Approach | Reject H_0 if $p\text{-value} \leq \alpha$ | Reject H_0 if $p\text{-value} \leq \alpha$ | Reject H_0 if $p\text{-value} \leq \alpha$ |
| Rejection Rule: Critical Value Approach | Reject H_0 if $\chi^2 \leq \chi_{(1-\alpha)}^2$ | Reject H_0 if $\chi^2 \geq \chi_{\alpha}^2$ | Reject H_0 if $\chi^2 \leq \chi_{(1-\alpha/2)}^2$ or if $\chi^2 \geq \chi_{\alpha/2}^2$ |

TABLE 11.3 SELECTED VALUES FROM THE F DISTRIBUTION TABLE*

| Denominator Degrees of Freedom | Area in Upper Tail | Numerator Degrees of Freedom | | | | |
|--------------------------------------|--------------------------|------------------------------|------|------|------|------|
| | | 10 | 15 | 20 | 25 | 30 |
| 10 | .10 | 2.32 | 2.24 | 2.20 | 2.17 | 2.16 |
| | .05 | 2.98 | 2.85 | 2.77 | 2.73 | 2.70 |
| | .025 | 3.72 | 3.52 | 3.42 | 3.35 | 3.31 |
| | .01 | 4.85 | 4.56 | 4.41 | 4.31 | 4.25 |
| 15 | .10 | 2.06 | 1.97 | 1.92 | 1.89 | 1.87 |
| | .05 | 2.54 | 2.40 | 2.33 | 2.28 | 2.25 |
| | .025 | 3.06 | 2.86 | 2.76 | 2.69 | 2.64 |
| | .01 | 3.80 | 3.52 | 3.37 | 3.28 | 3.21 |
| 20 | .10 | 1.94 | 1.84 | 1.79 | 1.76 | 1.74 |
| | .05 | 2.35 | 2.20 | 2.12 | 2.07 | 2.04 |
| | .025 | 2.77 | 2.57 | 2.46 | 2.40 | 2.35 |
| | .01 | 3.37 | 3.09 | 2.94 | 2.84 | 2.78 |
| 25 | .10 | 1.87 | 1.77 | 1.72 | 1.68 | 1.66 |
| | .05 | 2.24 | 2.09 | 2.01 | 1.96 | 1.92 |
| | .025 | 2.61 | 2.41 | 2.30 | 2.23 | 2.18 |
| | .01 | 3.13 | 2.85 | 2.70 | 2.60 | 2.54 |
| 30 | .10 | 1.82 | 1.72 | 1.67 | 1.63 | 1.61 |
| | .05 | 2.16 | 2.01 | 1.93 | 1.88 | 1.84 |
| | .025 | 2.51 | 2.31 | 2.20 | 2.12 | 2.07 |
| | .01 | 2.98 | 2.70 | 2.55 | 2.45 | 2.39 |

Note: A more extensive table is provided as Table 4 of Appendix B.

Table . Summary of hypothesis tests about two population variances

| | Upper Tail Test | Two-Tailed Test |
|--|---|--|
| Hypotheses | $H_0 : \sigma_1^2 \leq \sigma_2^2$ $H_a : \sigma_1^2 > \sigma_2^2$ | $H_0 : \sigma_1^2 = \sigma_2^2$ $H_a : \sigma_1^2 \neq \sigma_2^2$ Note: Population I has the larger sample variance |
| Test Statistic | $F = \frac{s_1^2}{s_2^2}$ | $F = \frac{s_1^2}{s_2^2}$ |
| Rejection Rule: p-Value Approach | Reject H_0 if $p\text{-value} \leq \alpha$ | Reject H_0 if $p\text{-value} \leq \alpha$ |
| Rejection Rule: Critical Value Approach | Reject H_0 if $F \geq F_\alpha$ | Reject H_0 if $F \geq F_\alpha$ |

8.2. Tests of Goodness of Fit and Independence

Summary

In this lecture we introduced the goodness of fit test and the test of independence, both of which are based on the use of the chi-square distribution. The purpose of the goodness of fit test is to determine whether a hypothesized probability distribution can be used as a model for a particular population of interest. The computations for conducting the goodness of fit test involve comparing observed frequencies from a sample with expected frequencies when the hypothesized probability distribution is assumed true. A chi-square distribution is used to determine whether the differences between observed and expected frequencies are large enough to reject the hypothesized probability distribution. We illustrated the goodness of fit test for multinomial, Poisson, and normal distributions.

A test of independence for two variables is an extension of the methodology employed in the goodness of fit test for a multinomial population. A contingency table is used to determine the observed and expected frequencies. Then a chi-square value is computed. Large chi-square values, caused by large differences between observed and expected frequencies, lead to the rejection of the null hypothesis of independence.

Glossary

Multinomial population A population in which each element is assigned to one and only one of several categories. The multinomial distribution extends the binomial distribution from two to three or more outcomes.

Goodness of fit test A statistical test conducted to determine whether to reject a hypothesized probability distribution for a population.

Contingency table A table used to summarize observed and expected frequencies for a test of independence.

Key formulas

Test Statistic for Goodness of Fit

$$\chi^2 = \sum_{i=1}^k \frac{(f_i - e_i)^2}{e_i} \quad (8.4)$$

Expected Frequencies for Contingency Tables Under the Assumption of Independence

$$e_{ij} = \frac{(\text{Row } i \text{ Total})(\text{Column } j \text{ Total})}{\text{Sample Size}} \quad (8.5)$$

Test Statistic for Independence

$$\chi^2 = \sum_i \sum_j \frac{(f_{ij} - e_{ij})^2}{e_{ij}} \quad (8.6)$$

Lecture supplementary material

MULTINOMIAL DISTRIBUTION GOODNESS OF FIT TEST: A SUMMARY

1. State the null and alternative hypotheses.

H_o : The population follows a multinomial distribution with specified probabilities for each of the k categories

H_a : The population does not follow a multinomial distribution with the specified probabilities for each of the k categories

2. Select a random sample and record the observed frequencies f_i for each category.

3. Assume the null hypothesis is true and determine the expected frequency e_i in each category by multiplying the category probability by the sample size.

4. Compute the value of the test statistic.

$$\chi^2 = \sum_{i=1}^k \frac{(f_i - e_i)^2}{e_i}$$

5. Rejection rule:

p -value approach: Reject H_o if $p\text{-value} < \alpha$

Critical value approach: Reject H_o if $\chi^2 > \chi^2_{\alpha}$

where α is the level of significance for the test and there are $k - 1$ degrees of freedom.

TEST OF INDEPENDENCE: A SUMMARY

1. State the null and alternative hypotheses

H_0 : The column variable is independent of the row variable

H_a : The column variable is not independent of the row variable

2. Select a random sample and record the observed frequencies for each cell of the contingency table.

3. Use equation $e_{ij} = \frac{(\text{Row } i \text{ Total})(\text{Column } j \text{ Total})}{\text{Sample Size}}$ (8.5) to compute the expected frequency for each cell.

4. Use equation $\chi^2 = \sum_i \sum_j \frac{(f_{ij} - e_{ij})^2}{e_{ij}}$ (8.6) to compute the value of the test statistic.

5. Rejection rule:

p-value approach: Reject H_0 if $p\text{-value} \leq \alpha$

Critical Value Approach: Reject H_0 if $\chi^2 \geq \chi^2_\alpha$

where α is the level of significance, with n rows and m columns providing $(n-1)(m-1)$ degrees of freedom.

POISSON DISTRIBUTION GOODNESS OF FIT TEST: A SUMMARY

1. State the null and alternative hypotheses.

H_0 : The population has a Poisson distribution

H_a : The population doesn't have Poisson distr.

2. Select a random sample and

a. Record the observed frequency f_i for each value of the Poisson random variable.

b. Compute the mean number of occurrences μ .

3. Compute the expected frequency of occurrences e_i for each value of the Poisson random variable. Multiply the sample size by the Poisson probability of occurrence for each value of the Poisson random variable. If there are fewer than five expected occurrences for some values, combine adjacent values and reduce the number of categories as necessary.

4. Compute the value of the test statistic.

$$\chi^2 = \sum_{i=1}^k \frac{(f_i - e_i)^2}{e_i}$$

5. Rejection rule:

p-value approach: Reject H_0 if $p\text{-value} < \alpha$

Critical value approach: Reject H_0 if $\chi^2 > \chi^2_\alpha$

where α is the level of significance and there are $k - 2$ degrees of freedom.

NORMAL DISTRIBUTION GOODNESS OF FIT TEST: A SUMMARY

1. State the null and alternative hypotheses.

H_0 : The population has a normal distribution

H_a : The population does not have a normal distribution

2. Select a random sample and

a. Compute the sample mean and sample standard deviation.

b. Define intervals of values so that the expected frequency is at least five for each interval. Using equal probability intervals is a good approach.

c. Record the observed frequency of data values f_i in each interval defined.

3. Compute the expected number of occurrences e_i for each interval of values defined in step 2(b). Multiply the sample size by the probability of a normal random variable being in the interval.

4. Compute the value of the test statistic.

$$\chi^2 = \sum_{i=1}^k \frac{(f_i - e_i)^2}{e_i}$$

5. Rejection rule:

p-value approach: Reject H_0 if $p\text{-value} < \alpha$

Critical value approach: Reject H_0 if $\chi^2 > \chi^2_\alpha$

where α is the level of significance and there are $k - 3$ degrees of freedom.