

Recall :-

## Hypothesis Test

z-test : Testing for sample mean when  $\sigma$  is known.

$$H_0: \mu = c$$

(Null)

$$H_A: \begin{cases} \mu < c \\ \mu > c \\ \mu \neq c \end{cases} \equiv \text{alternative}$$

Sample:  $x_1, x_2, \dots, x_n$  from population

Compute: 
$$\frac{\sqrt{n}(\bar{x} - c)}{\sigma}$$

$\bar{x} = \text{mean}$

Fix  $\alpha \in (0,1)$  and Find  $z_{\alpha/2}$  :  $TP(Z > z_{\alpha/2}) = \frac{\alpha}{2}$   
 $Z \sim \text{Normal}(0,1)$

check:  $\frac{\sqrt{n}(\bar{x} - c)}{\sigma} > z_{\alpha/2}$

if it happens then we would reject the null hypothesis.

$\Leftrightarrow$  Reject the null hypothesis if

$$P\left(Z \geq \frac{\sqrt{n}(\bar{x} - c)}{\sigma}\right) < \alpha$$

t-test:- Test sample mean when  $\sigma$  is not known.

$$H_0: \mu = c$$

(Null)

$$H_A: \begin{cases} \mu < c \\ \mu > c \\ \mu \neq c \end{cases} \equiv \text{alternative}$$

Sample:  $x_1, x_2, \dots, x_n$  from population

Compute:  $\frac{\sqrt{n} (\bar{x} - c)}{S}$

$S$  - sample variance  
 $\bar{x}$  - mean

Fix  $\alpha \in (0, 1)$

Reject null hypothesis if

$$P\left(T > \frac{\sqrt{n} (\bar{x} - c)}{S}\right) < \alpha \quad T \sim t_{n-1}$$

- Likelihood Ratio test statistic and derived the above test [last week notes]

# Hypothesis Testing– Proportions

Let  $n \geq 1$ ,  $X_1, X_2, \dots, X_n$  be i.i.d. Bernoulli ( $p$ ) random variables.

We want to test:

Null Hypothesis :  $p = 0.5$

Alternative Hypothesis:  $p \neq 0.5$

Use Binomial Central Limit Theorem that

$$\frac{\sqrt{n}(\bar{X} - p)}{\sqrt{p(1-p)}} \xrightarrow{d} Z,$$

where  $Z$  is standard Normal.

2-test :-

Assume convergence  
is "good"  
that

Normality assumption  
holds

Apply z-test

# Hypothesis Testing– Proportions

in built test

Use `prop.test`. Suppose  $n = 100$ ,  $\bar{X} = 0.43$ .

```
> prop.test(43,100)
```

```
1-sample proportions test with continuity correction
```

```
data: 43 out of 100, null probability 0.5
```

```
X-squared = 1.69, df = 1, p-value = 0.1936
```

```
alternative hypothesis: true p is not equal to 0.5
```

```
95 percent confidence interval:
```

```
0.3326536 0.5327873
```

```
sample estimates:
```

```
p
```

```
0.43
```

# Hypothesis Testing– Proportions

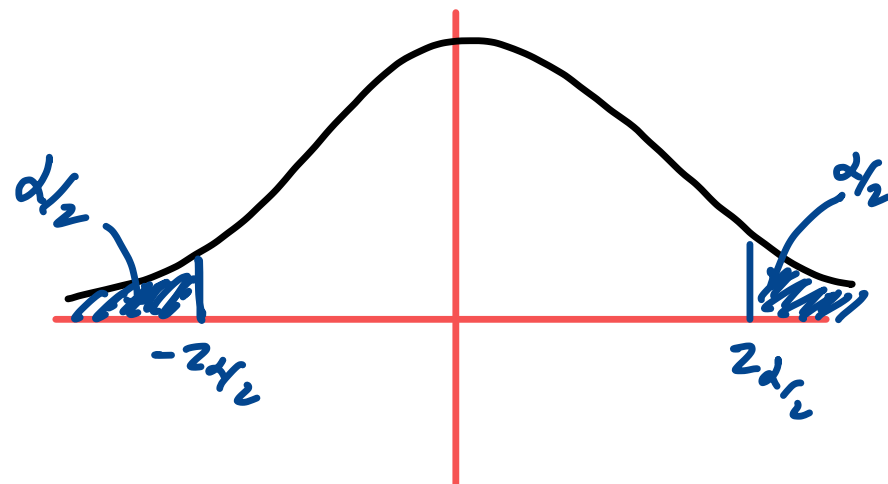
`prop.test` does the following:

- Computes  $P(|Z - 0.5| \geq | \frac{\sqrt{n}(\bar{X} - 0.5)}{0.5} - 0.5 |)$  towards  $p$ -value.

- Finds  $100(1 - \alpha)\%$ - Confidence Interval by finding the region of  $p$  where

$$\left| \frac{\sqrt{n}(\bar{X} - p)}{\sqrt{p(1 - p)}} \right| < z_{\frac{\alpha}{2}},$$

where  $P(Z > z_{\frac{\alpha}{2}}) = \frac{\alpha}{2}$ .



# Hypothesis Testing: z-test

Ex. Write a code for t-test

The below is code for z test and Confidence interval for a data x.

```
> ztestci = function(x, mu=0, sigma=1, alpha=0.95){  
+ z = qnorm( (1-alpha)/2, lower.tail=FALSE)  
+ sdx = sigma/sqrt(length(x))  
+ pvalue = pnorm(mean(x) - mu)/sdx,  
+ lower.tail=FALSE)  
+ c(mean(x) - z*sdx, mean(x) + z*sdx, pvalue)  
+ }  
> x=c(75,76,73,75,74,73,76,73,79) ; y = ztestci(x,76,1.5)  
> y  
[1] 73.9089069 75.8688709 0.9868659
```

$$P(|Z| \leq z) = \alpha$$

$$P(Z \geq \frac{\sigma_n(\bar{X} - \mu)}{s})$$

$$H_0: \mu = 76$$

$$H_A: \mu > 76$$

$$\bar{X} - z \cdot \frac{s}{\sqrt{n}} \quad \bar{X} + z \cdot \frac{s}{\sqrt{n}}$$



# Hypothesis Testing: $t$ -test

```
> t.test(x, mu=74)

    One Sample t-test

data:  x
t = 1.3571, df = 8, p-value = 0.2118
alternative hypothesis: true mean is not equal to 74
95 percent confidence interval:
 73.37848 76.39930
sample estimates:
mean of x
 74.88889
```

# Hypothesis Testing: $t$ -test

```
> wilcox.test(x,mu=74,alt="greater")
```

```
Wilcoxon signed rank test with continuity correction
```

```
data: x
```

```
V = 27, p-value = 0.1108
```

```
alternative hypothesis: true location is greater than 74
```

Applications: one needs to compare two population


Test for equality of means when variance is known

Assume:  $X \sim \text{Normal}(\mu_1, \sigma_1^2)$

$Y \sim \text{Normal}(\mu_2, \sigma_2^2)$

—  $\sigma_1$  is known and  $\sigma_2$  is known

$H_0: \mu_1 = \mu_2$  vs  $H_A: \mu_1 \neq \mu_2$

  
 $\mu_1 - \mu_2 = 0 \rightsquigarrow$  check:  $\bar{X} - \bar{Y} \sim \text{close to } 0$

Sample :-  $x_1, x_2, \dots, x_{n_1}$  from  $X$   
 $y_1, y_2, \dots, y_{n_2}$  from  $Y$

Under our assumptions :

$$\bar{X} - \bar{Y} \sim \text{Normal}(\mu_1 - \mu_2, \frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2})$$

Test :-  $Z \sim \text{Normal}(0,1)$

Fix  $\alpha \in (0,1)$

$$\text{It } \mathbb{P}\left(|Z| \geq \frac{|\bar{X} - \bar{Y}|}{\sqrt{\frac{\sigma_1^2}{n_1} + \frac{\sigma_2^2}{n_2}}}\right) < \alpha$$

Then reject null hypothesis.

## Test for proportions when variance is not known

Assume: "Two coins" -  $p_1$  - Prob of heads of coin 1  
 $p_2$  - Prob of heads of coin 2

$$H_0: p_1 = p_2$$

Sample :-  $X_1^{(1)}, \dots, X_n^{(1)} \rightarrow \hat{p}_1 = \bar{X}^{(1)}$

$X_1^{(2)}, \dots, X_n^{(2)} \rightarrow \hat{p}_2 = \bar{X}^{(2)}$

Statistic

$$Z = \frac{\hat{p}_1 - \hat{p}_2}{\sqrt{\dots}}$$

"Pooled variance"

$$\hat{p} = \frac{\hat{p}_1 + \hat{p}_2}{2}$$

$$\frac{\hat{p}(1-\hat{p})}{n}$$

Use  $Z \sim \text{Normal}(0,1) \equiv \text{Requires proof}$

# Hypothesis Testing: Two Sample Proportion-test

$X_1^{(1)}, \dots, X_{n_1}^{(1)}$  ... from population 1  
 $X_1^{(2)}, \dots, X_{n_2}^{(2)}$  ... from population 2

- Want to test if proportion of success  $p_1 = p_2$  between two populations.
- Let  $\hat{p}_1 = \bar{X}^{(1)}$  and  $\hat{p}_2 = \bar{X}^{(2)}$
- The statistic is

$$Z = \frac{\hat{p}_1 - \hat{p}_2}{\sqrt{\hat{p}(1 - \hat{p})\left(\frac{1}{n_1} + \frac{1}{n_2}\right)}},$$
$$\hat{p} = \frac{n_1 \hat{p}_1 + n_2 \hat{p}_2}{n_1 + n_2}$$

Large  $n_1, n_2$  assume normality for  $Z$

└──────────> Requires proof

# Hypothesis Testing: Two Sample-test

Assume : value of the variance(s) is not known but they are equal

Let  $n, m \geq 1$ ,  $X_1, X_2, \dots, X_n$  be i.i.d.  $\text{Normal}(\mu_X, \sigma_1^2)$  and  $Y_1, Y_2, \dots, Y_m$  be i.i.d.  $\text{Normal}(\mu_Y, \sigma_2^2)$ .

Test Statistic:

$$T := \frac{\bar{X} - \bar{Y} - (\mu_X - \mu_Y)}{S_{pooled} \sqrt{\frac{1}{n} + \frac{1}{m}}} \sim t_{n+m-1}$$

Requires  
a  
Proof

- Equal Variance:  $\sigma_1 = \sigma_2$

$$S_{pooled}^2 = \frac{(n-1)S_X^2 + (m-1)S_Y^2}{n+m-2}$$

$$H_0: \mu_X = \mu_Y$$

vs

$$H_A \equiv \begin{cases} \mu_X > \mu_Y \\ \mu_X < \mu_Y \\ \mu_X \neq \mu_Y \end{cases}$$



## Hypothesis Testing: Two Sample-test

Assume : value of the variances is not known but they are unequal

Let  $n, m \geq 1$ ,  $X_1, X_2, \dots, X_n$  be i.i.d.  $\text{Normal}(\mu_X, \sigma_1^2)$  and  $Y_1, Y_2, \dots, Y_m$  be i.i.d.  $\text{Normal}(\mu_Y, \sigma_2^2)$ .

Test Statistic:

$$\tilde{T} := \frac{\bar{X} - \bar{Y} - (\mu_X - \mu_Y)}{S_{pooled}} \quad \tilde{T} \sim t_d$$

- Equal Variance:  $\sigma_1 \neq \sigma_2$

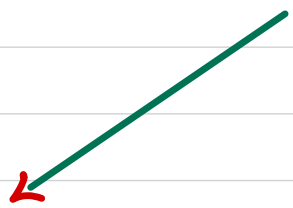
$$S_{pooled}^2 = \frac{S_X^2}{n-1} + \frac{S_Y^2}{m-1}$$


$d = \dots -$  complicated expression

## $\chi^2$ - Goodness of fit test :-

G. Mendel :- - seed shape      a and A governed by allele  
(pea)                      - cross breed to produce  
aa, aA, AA

Estimate :-       $P(aa) = ..$     $P(aA) = ..$     $P(AA) = ..$

  
Genetic laws  
Hypothesis

  
Verity  
??

Observations  
from  
cross - breeding

R. A. Fisher :- "Controversy" data was not repeatable. — 1936 Annals of statistics

"Data was too good a fit for the distribution".

Based on test :- identify if the data comes from a distribution

# $\chi^2$ - goodness of fit test

Some questions:

were the dice really fair?

Q1 • Are the dice we roll in our experiments in class really fair?

Q2 • Are two populations  $X$  and  $Y$  actually independent?

Rephrase:

- How well the distribution of the data fit the model?
- Does one variable affect the distribution of the other?

## Specific Question:

- To understand how "close" are the observed values to those which would be expected under the fitted model ?

## Towards Answer:

- In this case we seek to determine whether the distribution of results in a sample could plausibly have come from a distribution specified by a null hypothesis.
- The test statistic is calculated by comparing the observed count of data points within specified categories relative to the expected number of results in those categories (under Null).

$x_1, \dots, x_n$  — sample from  $X$

$H_0: X \sim \begin{cases} \text{Normal} \\ \vdots \\ \text{Bernoulli}(p) \end{cases}$

Test ?

# $\chi^2$ - goodness of fit test

- Let  $T$  be a random variable with finite range  $\{c_1, c_2, \dots, c_k\}$  for which

Null Hypothesis  $P(T = c_j) = p_j > 0$  for  $1 \leq j \leq k$ .

Suppose there are  $k$  possible outcomes and each occurring with a specified Probability.

- Let  $X_1, X_2, \dots, X_n$  be the sample from the distribution  $T$  and let

Typo:  $Y_j = |\{k : X_k = c_j\}|$   $Y_j = |\{j : X_j = c_j\}|$  for  $1 \leq j \leq k$ .

“Counting the number of sample points in each bin”

$Y_j$  is the number of sample points whose outcome was  $c_j$

- Then the statistic

$$\chi^2 := \sum_{j=1}^k \frac{(Y_j - np_j)^2}{np_j} \equiv \sum_{j=1}^k \frac{(\text{Observed} - \text{Expected})^2}{\text{Expected}}$$

Pearson's Chi-square Test Statistic

# $\chi^2$ - goodness of fit test

$$\mathbf{X}^2 := \sum_{j=1}^k \frac{(Y_j - np_j)^2}{np_j} \equiv \sum_{j=1}^k \frac{(\text{Observed} - \text{Expected})^2}{\text{Expected}}$$

- $\mathbf{X}^2$ - has  $\chi_{k-1}^2$  degrees of freedom, asymptotically as  $n \rightarrow \infty$ . Requires a proof which we will omit for this course
- **Null Hypothesis:** Distribution comes from Multinomial with parameters  $p_1, p_2, \dots, p_k$
- **Alternate Hypothesis:** Distribution comes from Multinomial with parameters with at least one parameter different from  $p_1, p_2, \dots, p_k$

Fix level of significance “alpha”

And use the distribution fact about  $\mathbf{X}^2$  — as chi-square to compute the p-value

# $\chi^2$ - goodness of fit test

Example has three outcomes: NDA, UPA, Third-Front

Probability of each outcome: 0.38, 0.32, 0.3

Observed: 35, 40, 25

Sample Size  $n = 100$

## Example:

We divide the political parties in India into 3 large alliances: NDA, UPA, and Third-Front. In the previous election the support had been 38%, 32% and 30% support respectively. Super-Nation TV channel takes a sample of 100 people and finds that there are 35 for NDA, 40 for UPA and 25 for Third-Front. It concludes that the vote share has not changed. Is this hypothesis correct ?

Expected ::= (38, 32, 30)

Observed - Expected : === (35-38, 40-32, 25-30)



# Contingency Tables

- Bivariate Data is often presented as a two-way table.

- For example in Dengue Data from Manipal Hospital

```
> y = read.table("dengueb.csv", header=TRUE)
> head(y)          > tail(y)
```

	DIAGNO	BICARB1		DIAGNO	BICARB1
1	DSS	16.2	45	D	22.0
2	DSS	22.0	46	D	16.6
3	DSS	16.0	47	D	18.3
4	DSS	21.3	48	D	23.0
5	DSS	19.0	49	D	24.0
6	DSS	18.7	50	D	21.0

# Contingency Tables

- Bivariate Data is often presented as a two-way table.
- For example in Dengue Data from Manipal Hospital

Diagnosis		
Cat.Marker	D	DSS
0	0	6
1	17	15
2	8	4

where we have grouped values of Marker to be 0, 1, 2 depending on the values being less than or equal to 16, between 16 and 21, and greater than 21.

# $\chi^2$ - test of independence

## Specific question:

- Does one variable affect the distribution of the other ?

## Notation:

- Let  $n_r$  be the number of rows in the table.
- Let  $n_c$  be the number of columns in the table.
- Let  $n = n_r n_c$  be the total number of observations.

If marker does not work then the diagnosis should be independent of the marker.

## Model:

- Let  $T \equiv (p_{ij})$  with  $1 \leq i \leq n_r, 1 \leq j \leq n_c$  be a probability distribution on  $\{(i, j) : 1 \leq i \leq n_r \text{ and } 1 \leq j \leq n_c\}$
- Let  $p_i^R = \sum_{j=1}^{n_c} p_{ij}$  and  $p_j^C = \sum_{i=1}^{n_r} p_{ij}$

# $\chi^2$ - test of independence

- Null Hypothesis: Variables are independent i.e

$$p_{ij} = p_i^R p_j^C \text{ for all } 1 \leq i \leq n_r \text{ and } 1 \leq j \leq n_c$$

- Alternate Hypothesis: Variables are not independent

# $\chi^2$ - test of independence

- Let  $y_{ij}$  record the frequency in the  $(i, j)$  cell.
- Let

$$\hat{p}_i^R = \frac{\sum_{j=1}^{n_c} y_{ij}}{\sum_{i=1}^{n_r} \sum_{j=1}^{n_c} y_{ij}} \text{ and } \hat{p}_j^C = \frac{\sum_{i=1}^{n_r} y_{ij}}{\sum_{i=1}^{n_r} \sum_{j=1}^{n_c} y_{ij}}$$

Individual Probabilities

Let

$$\hat{p}_{ij} = \hat{p}_i^R \hat{p}_j^C \quad \text{Under Independence}$$

and

$$\mathbf{x}^2 := \sum_{i=1}^{n_r} \sum_{j=1}^{n_c} \frac{(y_{ij} - n\hat{p}_{ij})^2}{n\hat{p}_{ij}}$$

# $\chi^2$ - test of independence

- Test Statistic:

$$\mathbf{X}^2 := \sum_{i=1}^{n_r} \sum_{j=1}^{n_c} \frac{(y_{ij} - n\hat{p}_{ij})^2}{n\hat{p}_{ij}}$$

Omit Proof for this class

is  $\chi_q^2$  distributed asymptotically as  $n \rightarrow \infty$  with  $q = (n_r - 1)(n_c - 1)$  degrees of freedom.

- Decide on level of significance:  $\alpha$

- Compute  $p$ -value:

$$\mathbb{P}(\chi_q^2 \geq \mathbf{X}^2)$$

- Reject Null Hypothesis:

if  $p$ -value is less than  $\alpha$

# $\chi^2$ - test of independence

For example in Dengue Data from Manipal Hospital:

```
> T = table(Cat.Marker, Diagnosis)
```

```
> T
```

	Diagnosis		
Cat.Marker	D	DSS	
0	0	6	
1	17	15	
2	8	4	

Can we test if the Marker value is independent of the characterisation of Dengue as normal or severe ?

Doctor's needs:

A patient arrives with Dengue

Based on Marker doctor needs to decide on Treatment

Statistical test performed:

We collected data of patients : Marker and final diagnosis

We test if Marker is independent of Diagnosis

# $\chi^2$ - test of independence

For example in Dengue Data from Manipal Hospital:

```
> chisq.test(T)
```

```
Pearson's Chi-squared test
```

```
data:  T
```

```
X-squared = 7.4583, df = 2, p-value = 0.02401
```