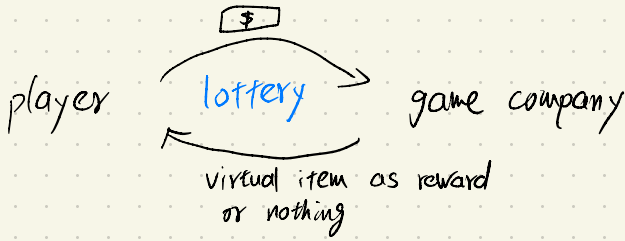


Appetizer: How unlucky is unlucky?



Rules

- \$1 per shot, non-refundable
- prob = $\frac{1}{10}$ to win.
- player may shoot as many times, may stop any time

Outcome

- 30 shots total
- 1 win, 29 lose \Rightarrow observed prob = $\frac{1}{30}$
- player sues the company

If you were the $\left. \begin{array}{l} \text{judge} \\ \text{player} \\ \text{company} \end{array} \right\}$, what would you do?

$$P(1 \text{ win } 29 \text{ lose} \mid P = \frac{1}{10}) = \binom{29}{1} \cdot \left(\frac{1}{10}\right)^1 \left(\frac{9}{10}\right)^{29} \approx 0.137$$

Is this enough to prove the company guilty?
or the player is having bad luck?

P-value

It's used to test the null hypothesis H_0 (the initial hypothesis)

If $p < .05$, then we reject H_0

Otherwise we don't reject H_0

Discrete case

p-value = the prob. of seeing sth that's equally rare

+ the prob of seeing sth rarer or more extreme

+ the prob of seeing the observed

↗
The prob calculated with H_0

eg.

H_0 = fair coin for 5 times

Is the coin fair?

result = 4 heads, 1 tail

$$P(4H, 1T) = \frac{5}{32} \quad P(5H) = P(5T) = \frac{1}{32}$$

$$P(4T, 1H) = \frac{5}{32} \quad \therefore p\text{-value} = \frac{5}{32} + \frac{5}{32} + \frac{1}{32} \times 2 = 0.375$$

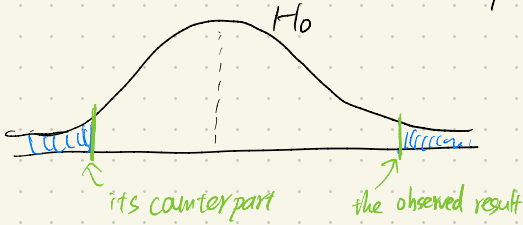
Fail to reject H_0

~~✗~~ accept H_0

More data is needed.

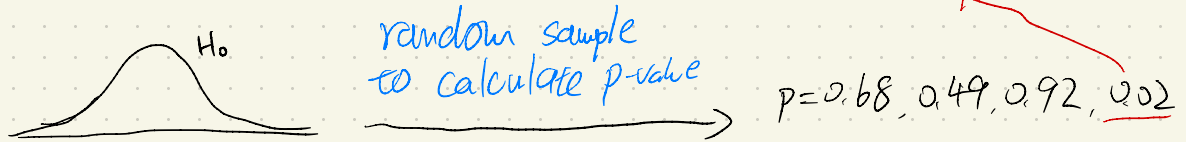
Continuous case

p-value = the blue area



If we reject H_0 , then it means there is a better distribution to fit the data
Otherwise H_0 is good enough

Approximate 5% of stats tests from the same distribution will be false positive



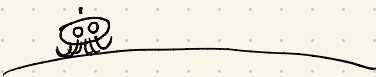
How tall is human?
I knew they're around 5.7 ft $= H_0$

What r u?

Unfortunately, the alien went to a kindergarten,
and recorded avg = 2.6 ft, SD = 0.4 ft
 $\Rightarrow p = 0.005 \Rightarrow$ reject H_0

\Rightarrow it's a false positive

Human are shorter than I thought



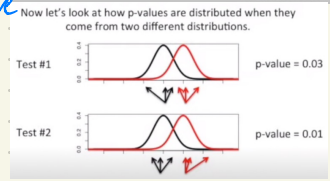
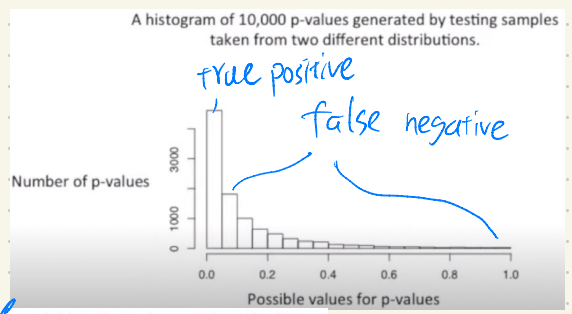
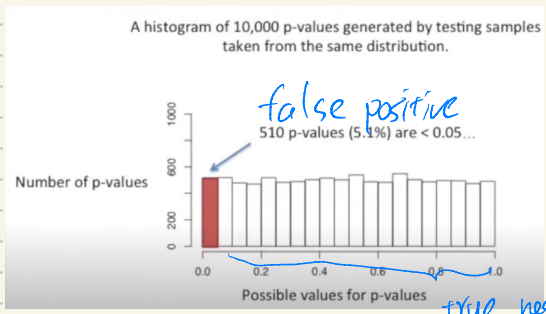
Confusion matrix

		Reject null hypothesis	
		No $p > 0.05$	Yes $p < 0.05$
Null hypothesis	True	True negative	False positive <i>type 1 error</i>
	False	False negative <i>type 2 error</i>	True positive

eg.
 $H_0 \sim$ healthy, Test \sim virus test
 positive = reject H_0 = unhealthy
 = test positive

$$\text{specificity} = \frac{TN}{TN + FP}$$

$$\text{sensitivity} = \text{recall} = \frac{TP}{TP + FN}$$



p-values are uniform if draw from the same distr.

p-values are skewed or close to zero if draw from diff. distr.

Likelihood

$$L(\theta; x) = \text{Prob}(X=x | \theta = \theta) = f_x(x; \theta) \quad \leftarrow \text{PDF}$$

$$L(\theta) = L(\theta; x) = \prod_{i=1}^n f_i(x_i; \theta)$$

x = the data, θ = the parameters, n = # of samples

score $\frac{d}{d\theta} l(\theta)$

loglikelihood $l(\theta) = \log L(\theta)$

Score equation $\frac{d}{d\theta} l(\theta) = 0$ is to find the max $l(\theta)$

Observed information $I_o(\theta) = -\frac{d^2}{d\theta^2} l(\theta)$ quantifies the confidence in the MLE (max likelihood estimation)

Expected information $I(\theta) = \underline{E}[I_o(\theta; X)]$
 \leftarrow expected value

Variance $V(\hat{\theta}) \approx I(\theta)^{-1}$

If θ is multi-dim, $\vec{\theta}$, then score equ: $\frac{\partial}{\partial \theta_i} l(\vec{\theta}) = 0$

information: $I_{ij}(\vec{\theta}) = -\frac{\partial^2}{\partial \theta_i \partial \theta_j} l(\vec{\theta})$

Bayes Thm

$$P(\theta | D) = \frac{\overset{\text{likelihood}}{P(D|\theta)} \overset{\text{prior}}{P(\theta)}}{P(D) \leftarrow \text{(normalizing const)}}$$

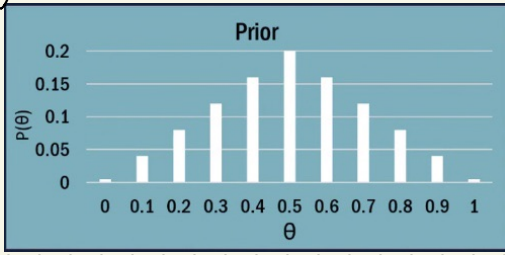
prior and posterior are distribution (pdf)

D = data
 θ = model or parameter

eg. flip a coin

proportion of heads

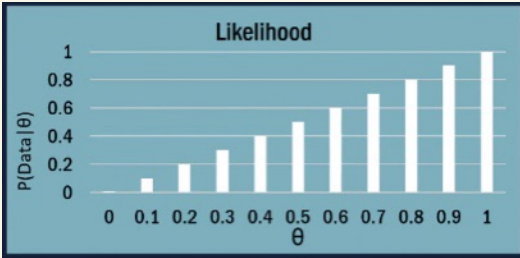
$$P(\theta)$$



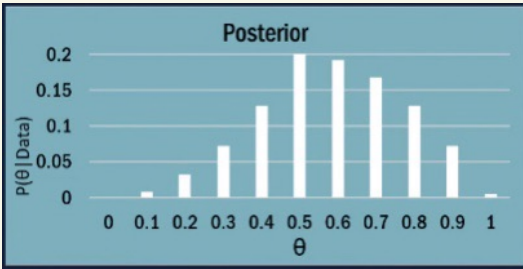
let $n=1$, head = 1

$$P(\text{Data} | \theta)$$

*(not a pdf)



$$P(\theta | \text{Data})$$



If posterior has the same parametric form as prior
Then the prior is called **conjugate prior**

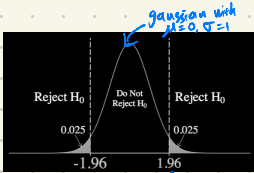
$$P(\theta | \text{Data}) = \frac{P(\text{Data} | \theta) P(\theta)}{\int P(\text{Data} | \theta) P(\theta) d\theta}$$

Z test (Suppose σ is known)

$$H_0: \mu = \mu_0$$

$$\text{Test stat: } Z = \frac{\bar{x} - \mu_0}{\sigma_{\bar{x}}}, \text{ where } \sigma_{\bar{x}} = \frac{\sigma}{\sqrt{n}}, \text{ where } \sigma \text{ is known}$$

If H_0 is true, then Z will be in normal distr.

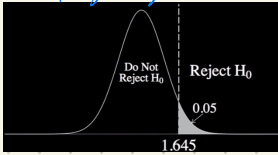


If $|Z| \geq 1.96$, then reject H_0 given $\alpha = 0.05$

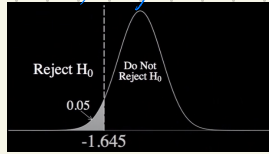
Z value threshold use ppf to find

We don't calculate p-value
We convert α to the threshold

$$H_1: \mu > \mu_0$$



$$H_1: \mu < \mu_0$$



T test (Suppose σ is unknown)

$$H_0: \mu = \mu_0$$

$$\text{Test stat: } t = \frac{\bar{x} - \mu_0}{SE}, \text{ where } SE = \frac{s}{\sqrt{n}}, \text{ } s = \text{SD of } x \text{ (observed)}$$

If H_0 is true, then t will be in t -distr. with $n-1$ df

Chi-squared test

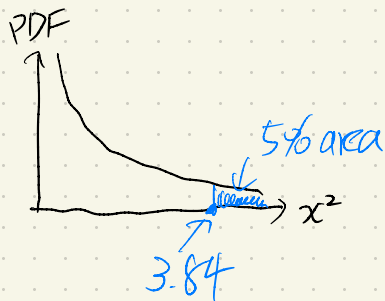
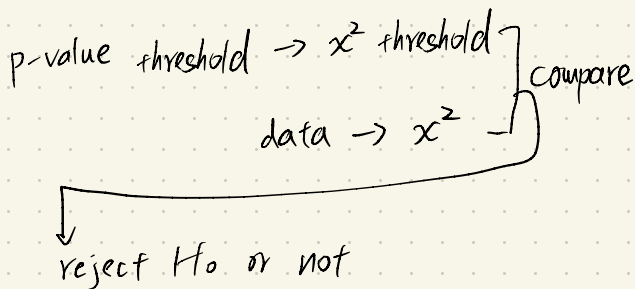
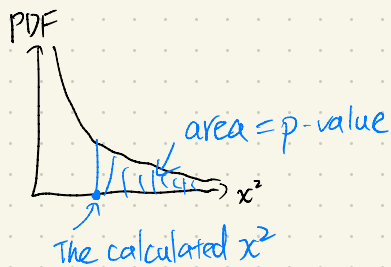
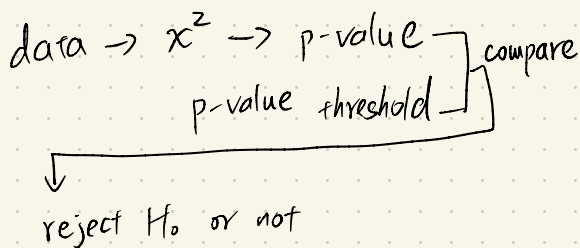
- goodness of fit
- test for indep

degree of freedom $df = (r-1) \cdot (c-1)$, $r = \#$ of rows, $c = \#$ of columns

$$\chi^2 = \sum_i \frac{(o_i - e_i)^2}{e_i}, \quad o_i = \text{observed}, \quad e_i = \text{expected (calculated by } H_0)$$

Reject H_0 if $\chi^2 > \chi^2_{ppf}(0.95, df) = 3.84$

or if $p = 1 - \chi^2_{cdf}(\chi^2, df) < 0.05$



Inference for one variance

Let σ be SD of the population
S be SD of the samples

$$H_0: \sigma = \sigma_0$$
$$\text{test stat: } \chi^2 = \frac{(n-1)s^2}{\sigma_0^2}$$

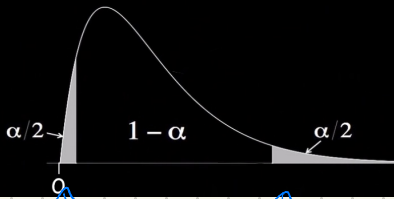
s^2 is unbiased estimator of σ^2 , i.e. $E(s^2) = \sigma^2$

$\frac{(n-1)s^2}{\sigma^2}$ has χ^2 -distrib with $df = n-1$ (if the population is gaussian)

eg. $\alpha = 0.05$

A $(1-\alpha)100\%$ confidence interval for σ^2 is given by:

$$\left(\frac{(n-1)s^2}{\chi_{\alpha/2}^2}, \frac{(n-1)s^2}{\chi_{1-\alpha/2}^2} \right)$$



$\chi_{1-\alpha/2}^2$

$\chi_{\alpha/2}^2$ use pft to calculate

Calculate the confidence interval

$$P\left(\chi_{1-\alpha/2}^2 < \frac{(n-1)s^2}{\sigma^2} < \chi_{\alpha/2}^2\right) = 1-\alpha$$

$$\Rightarrow P\left(\frac{(n-1)s^2}{\chi_{\alpha/2}^2} < \sigma^2 < \frac{(n-1)s^2}{\chi_{1-\alpha/2}^2}\right) = 1-\alpha$$

lower bound upper bound