

## Test statistic for Sample mean:

- Mean Weight of packet

↓  
450 gm

↓  
 $\mu$

- std. dev.

↓  
3 gm

↓  
 $\sigma$

} Population mean / std. dev.  
(from historical data)

⇒ test statistic: Z score

✓  $n$ : Sample size

↳  $\bar{x}$ : sample mean

$$\bar{x} \sim N\left(\mu, \frac{\sigma}{\sqrt{n}}\right)$$

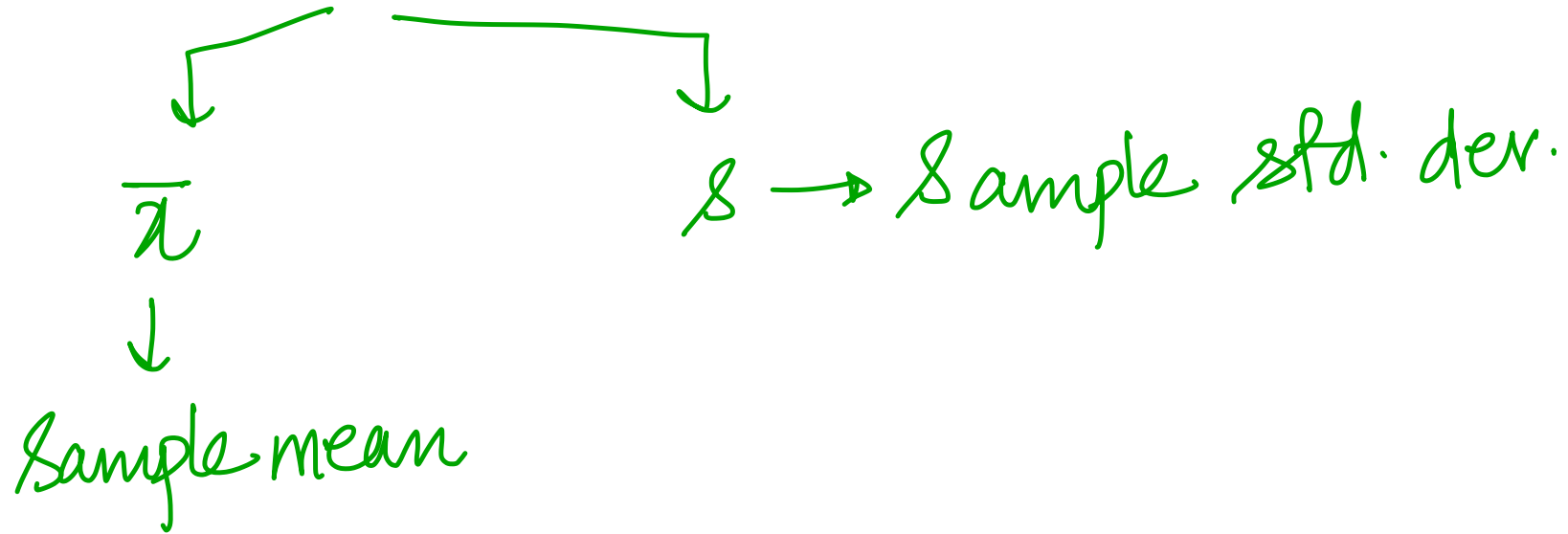
$$Z = \left( \frac{\bar{x} - \mu}{\sigma / \sqrt{n}} \right)$$

Normal distribution

- In hypothesis testing, we often don't have population standard deviation  $\rightarrow \sigma$ .

- what to do then?

• from 'n' samples



• If  $n$  is large enough; usually  $n > 30$   
we can safely assume

$$Z = \left( \frac{\bar{x} - \mu}{s/\sqrt{n}} \right)$$

$Z \sim$  Normal  
distribution  
=.

• If  $n < 30$ ;

test statistic  $\rightarrow \left( \frac{\bar{x} - \mu}{s/\sqrt{n}} \right)$  follows T-distribution

$\Rightarrow t = \left( \frac{\bar{x} - \mu}{s/\sqrt{n}} \right) \Rightarrow t \sim \text{T-distributions}$

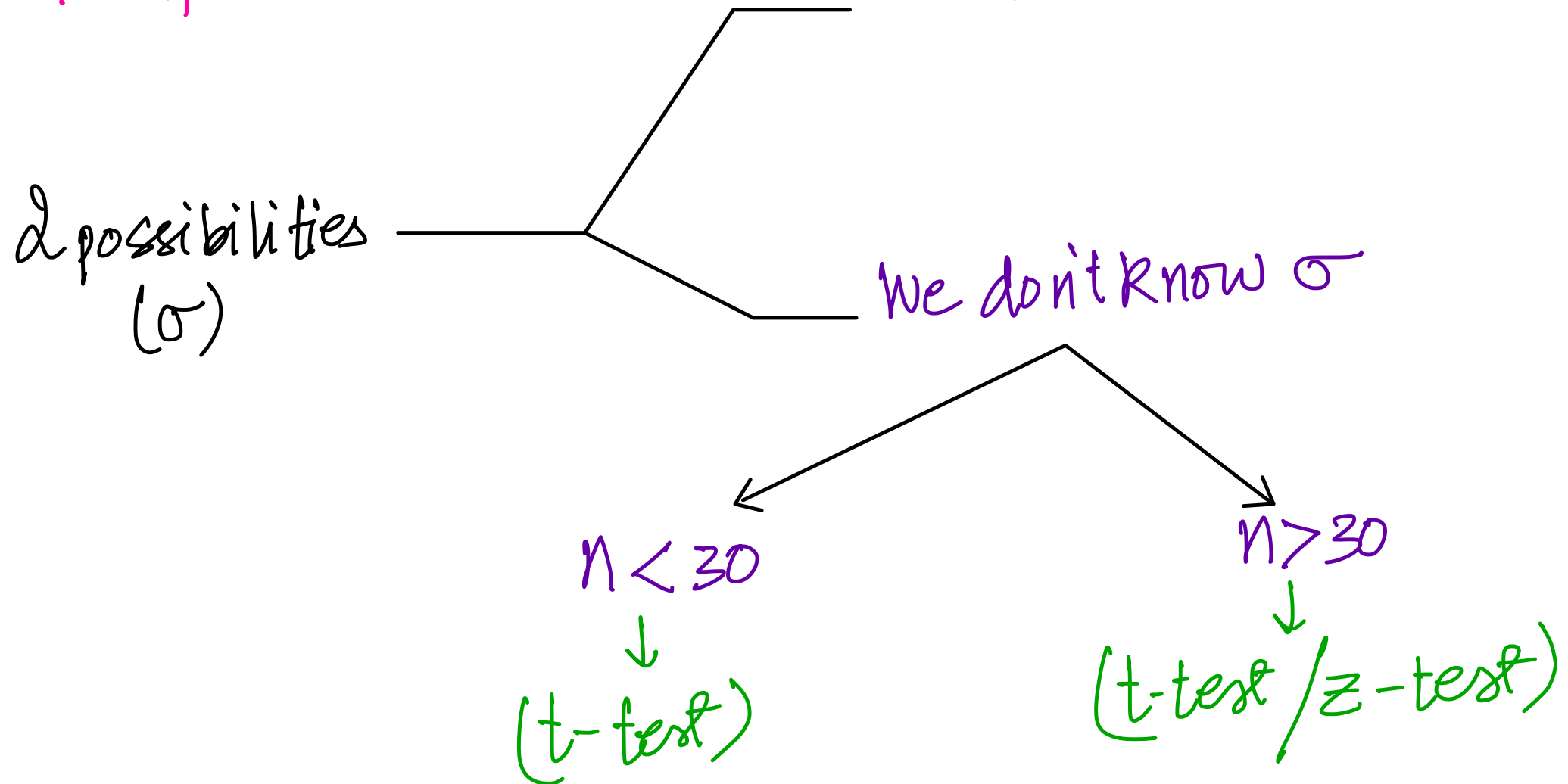
↓

we use  
T-distribution CDF to  
find out p-value.

Summary: When to use what.

$\sigma$ : population std. deviation

$s$ : sample std. dev.



Only one parameter  $\rightarrow$  t-distribution. degree of freedom (dof)

$\Rightarrow$  sample size  $= n$

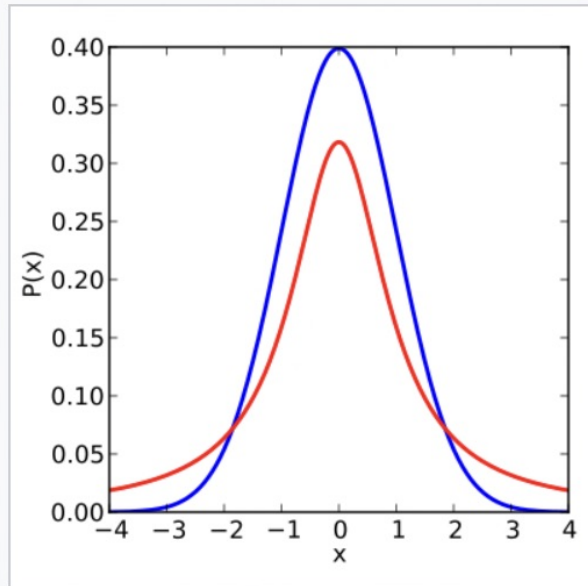
$\hookrightarrow$  test statistic

$$t \sim T[\text{dof} = n-1]$$

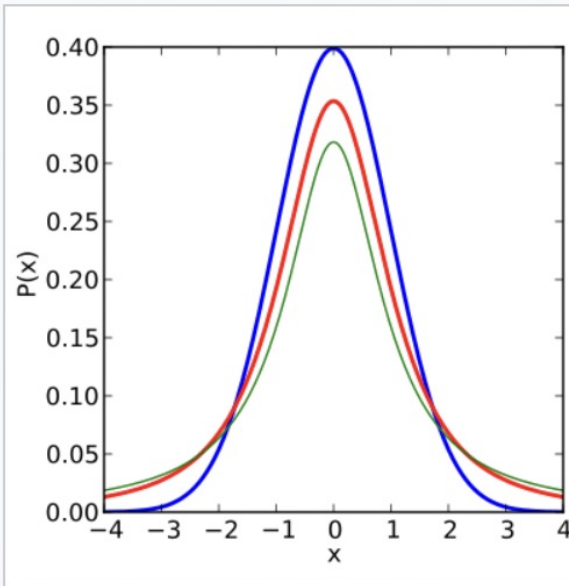
$\Rightarrow$  as  $n \uparrow$  T-distribution becomes more and more gaussian (Normal?)

**Density of the  $t$ -distribution (red) for 1, 2, 3, 5, 10, and 30 degrees of freedom compared to the standard normal distribution (blue).**

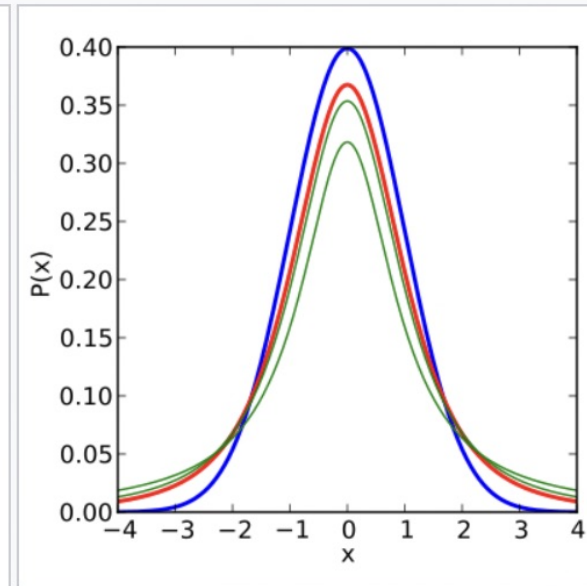
**Previous plots shown in green.**



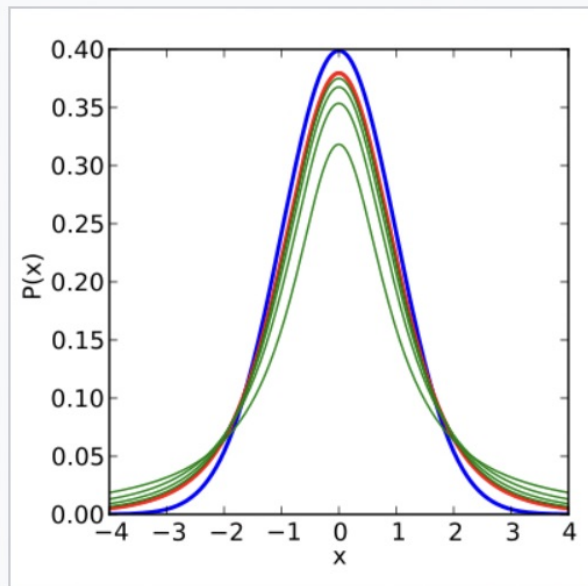
1 degree of freedom



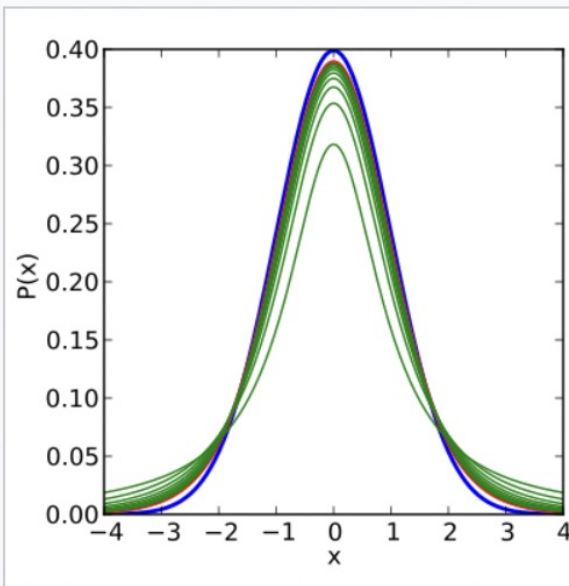
2 degrees of freedom



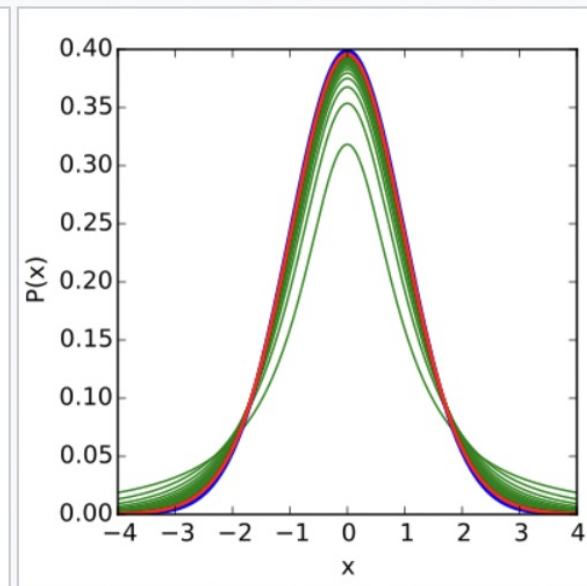
3 degrees of freedom



5 degrees of freedom



10 degrees of freedom



30 degrees of freedom

# One Sample t-test

A public health official claims that the mean home water use is 350 gallons a day.

To verify this claim, a study of 20 randomly selected homes was instigated with the result that the average daily water uses of these 20 homes were as follows:

usage = [340, 344, 362, 375, 356, 386, 354, 364, 332, 402, 340, 355, 362, 322, 372, 324, 318, 360, 338, 370]

•  $H_0: \mu = 350$

$H_A: \mu \neq 350$

Two tailed

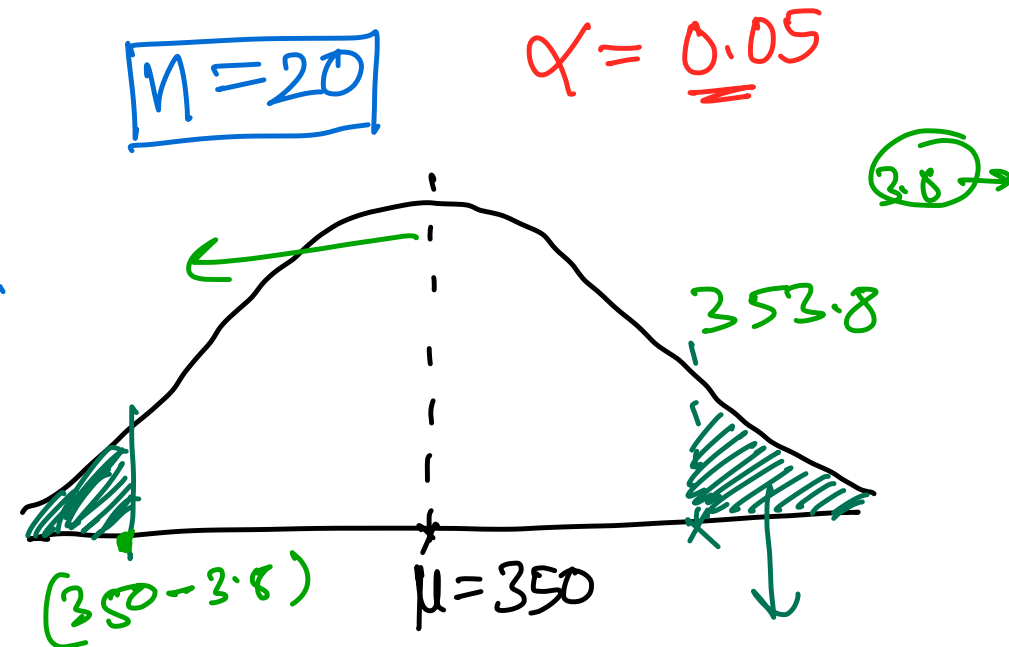
$\bar{m} \rightarrow$  Sample mean ;  $s \rightarrow$  sample std. dev.

$\bar{m} = 353.8$

• test statistic

$$t = \left( \frac{\bar{m} - \mu}{s / \sqrt{n}} \right)$$

$\Rightarrow \boxed{t = 0.798}$



Since 2 tailed

from scipy.stats import t

p\_value = 2\*(1 - t.cdf(t\_stat, df = n-1))

p\_value

0.798  
 $\rightarrow df = 19$

p\_value = 0.4347

$\because p\_value > \alpha \Rightarrow$  failed to reject  $H_0$

# Two Samples

## Drug Recovery

Suppose two companies develop a drug for a disease.

Drug 1 was tested on 100 people, and the recovery days look like this  
[8, 5, 9, 10, ....., 16]

The mean recovery time was 7.1 days

Drug 2 was tested on 120 people, and the recovery days look like this  
[12, 4, 7, 13, ....., 8]

The mean recovery time was 8.07 days

Can we say one drug was better than the other?

Or was this small difference a coincidence?

For such cases we use the **two-sample Z-test** or **two-sample T-test**

\* Comparing 2 versions (A & B)  $\Rightarrow$  **A/B TESTING** ✓



Let  $\mu_1$  be the average recovery for drug 1, and  $\mu_2$  be the average for drug 2

$$H_0: \mu_1 = \mu_2$$

$$H_a: \mu_1 \neq \mu_2$$

• from data of drug<sub>1</sub> and drug<sub>2</sub>

$$\downarrow$$
$$n_1 = 100$$

$$m_1 = 7.1$$

$$s_1$$

$$\downarrow$$
$$n_2 = 120$$

$$m_2 = 8.07$$

$$s_2$$

— number of samples  
— sample mean  
— sample std. dev<sup>n</sup>.

# Test statistic :

$$t = \left( \frac{(m_1 - m_2)}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}} \right)$$

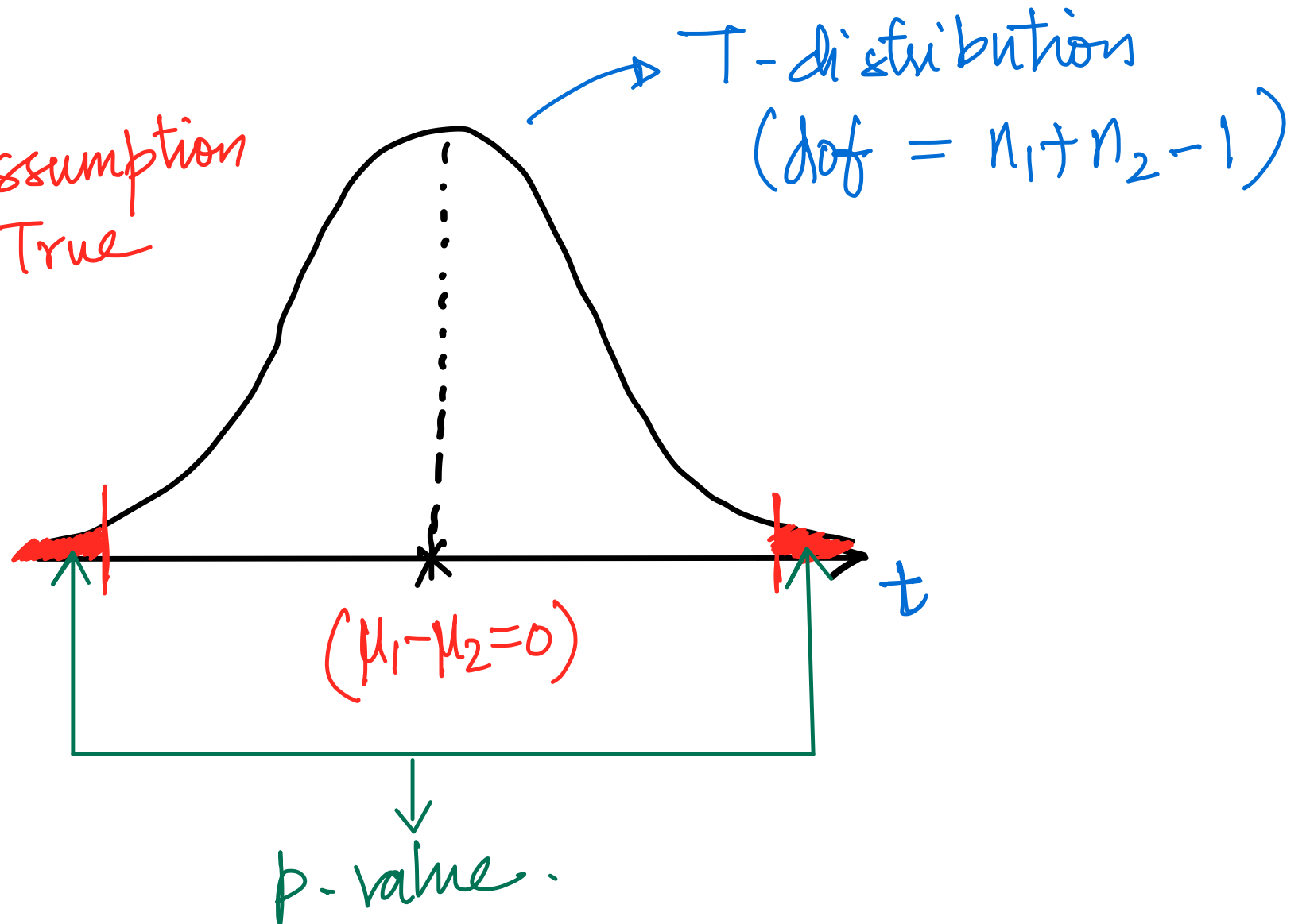
→ if  $n_1, n_2 < 30$   
Use 2 sample t-test

→ if  $n_1, n_2 > 30$   
2 sample t test / 2 sample z-test

# Two Tailed

# p-value

• under the assumption  
that  $H_0$  is True  
( $\mu_1 = \mu_2$ )



# If  $p\text{-value} < \alpha$  (let's say  $\alpha = 0.05$ ).  
we reject  $H_0$  (No statistically significant  
diff. b/w mean recovery  
time of  $D_1$  &  $D_2$ ).

And conclude that. there is  
statistically significant.

diff. b/w mean recovery  
time of  $D_1$  &  $D_2$

Else; failed to Reject  $H_0$ .