

# Gaussian distributions

## QQ plots

## t-tests

## Comparing two datasets



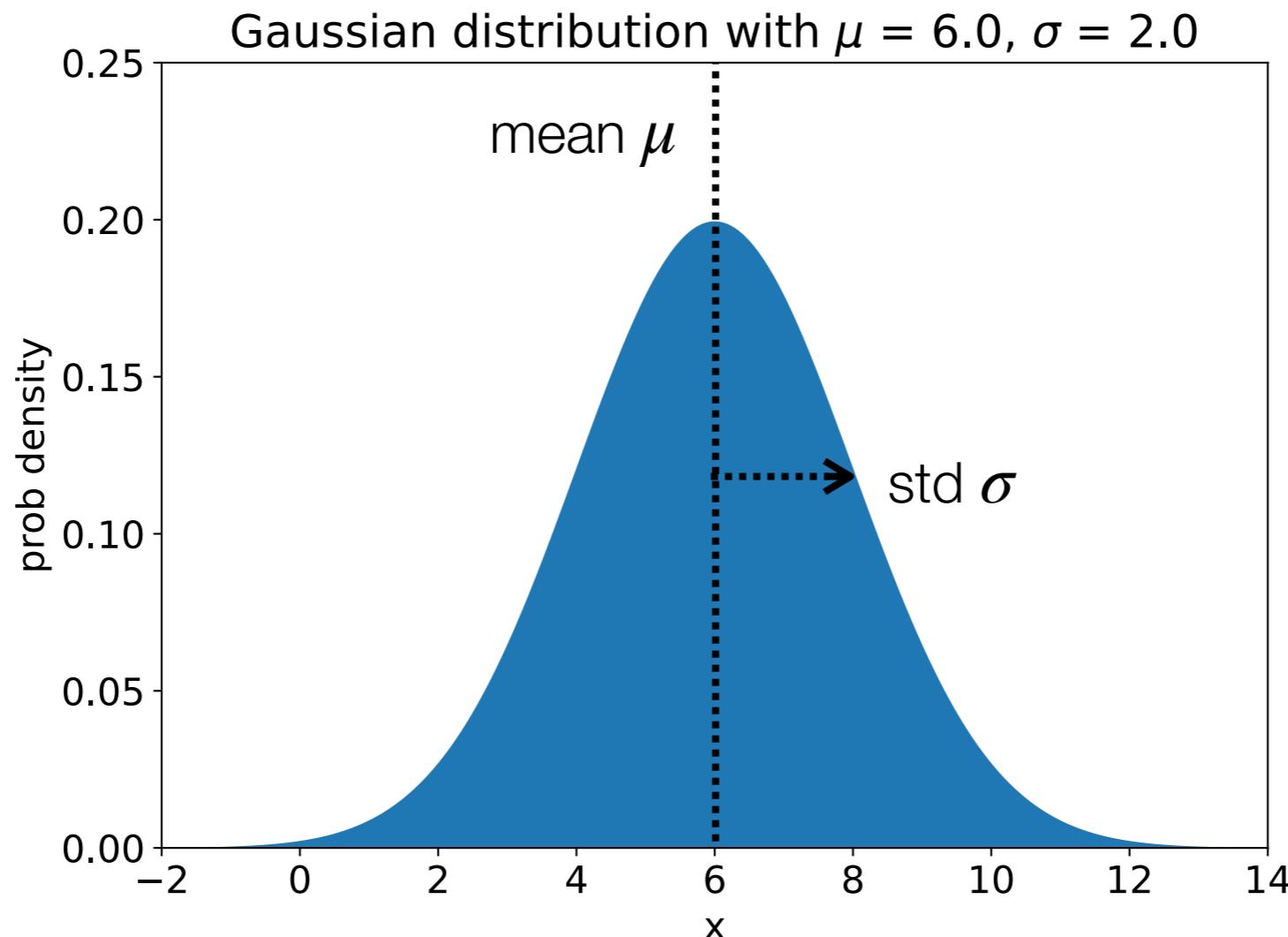
Biostatistics Course 2024  
Lecture 3  
Wednesday, 10 July 2024  
10:00am - 12:00pm

## Gaussian distributions

## The normal distribution is ubiquitous in statistics

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“Gaussian distribution” = “normal distribution”



$x \sim \text{Normal}(\mu, \sigma^2)$

drawn from      mean      variance

## Mean and variance

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Let  $X \sim N(\mu, \sigma^2)$

- Mean:  $E[X] = \mu$
- Variance:  $Var[X] = \sigma^2$
- Standard Deviation:  $SD_X = \sigma$

## Mean of standardized random variable

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Let

$$Z = (Y - \mu)/\sigma$$

$$\begin{aligned} E[Z] &= E\left[\frac{Y - \mu}{\sigma}\right] = \frac{1}{\sigma}E[Y - \mu] \\ &= \frac{1}{\sigma}(E[Y] - \mu) \\ &= \frac{1}{\sigma}(\mu - \mu) \\ &= 0 \end{aligned}$$

## Variance of standardized random variable

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$$\begin{aligned} \text{Var}[Z] &= \text{Var}\left[\frac{Y - \mu}{\sigma}\right] \\ &= \frac{1}{\sigma^2} \text{Var}[Y - \mu] \\ &= \frac{1}{\sigma^2} \text{Var}[Y] \\ &= \frac{1}{\sigma^2} \sigma^2 \\ &= 1 \end{aligned}$$

**NOTE:**  $\mu = 0$  and  $\sigma^2 = 1$  for **any** standardized random variable

## 68-95-99.7 Rule

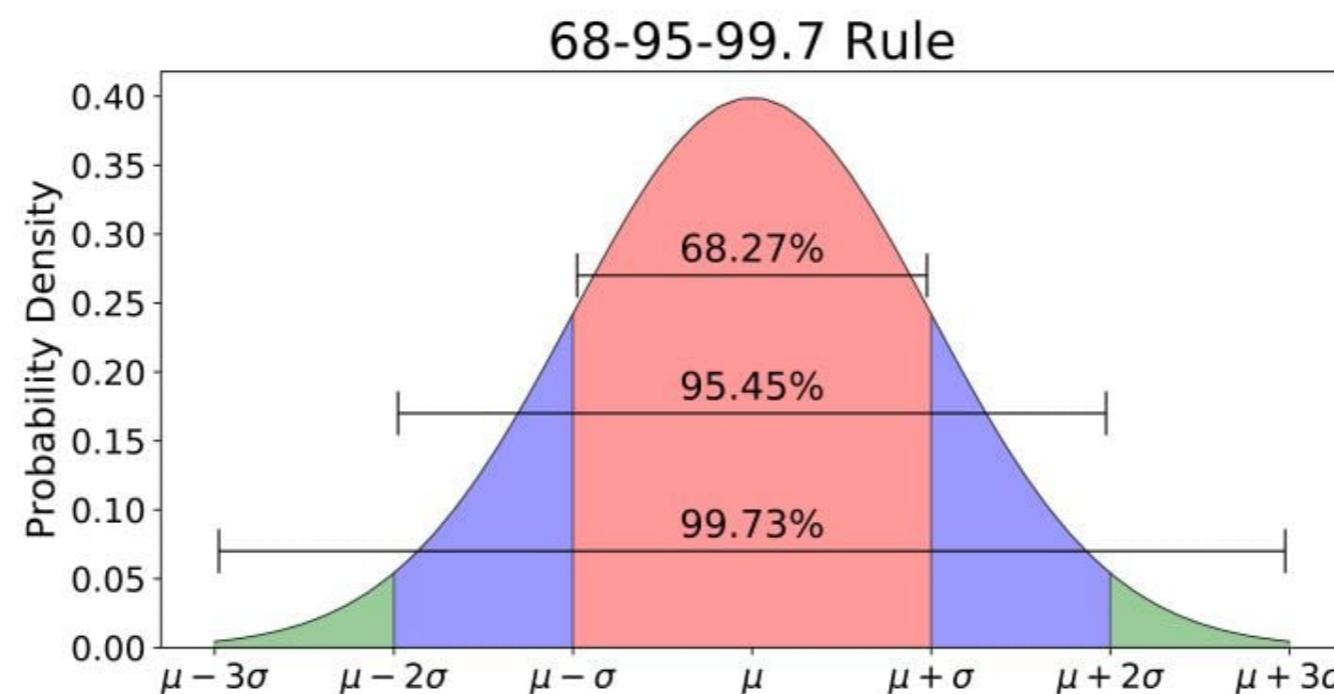
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Recall the 68-95-99.7 rule Note for a standard normal random variable,  
 $Z \sim N(0, 1)$

$$Pr(-1 < Z < 1) \approx 0.68$$

$$Pr(-2 < Z < 2) \approx 0.95$$

$$Pr(-3 < Z < 3) \approx 0.997$$



## The central limit theorem makes the normal distribution extremely relevant

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If a random variable  $X$  has population mean  $\mu$  and population variance  $\sigma^2$ , the sample mean  $\bar{X}$ -bar, based on  $n$  observations, is approximately normally distributed with mean  $\mu$  and variance  $\sigma^2/n$ , for sufficiently large  $n$ .

$$x_1 \sim p_1(x)$$

$$x_2 \sim p_2(x)$$

...

$$x_N \sim p_N(x)$$

$$\bar{x} = \frac{x_1 + x_2 + \cdots + x_N}{N} \rightarrow \bar{x} \sim \text{Normal}(\mu, \sigma^2)$$

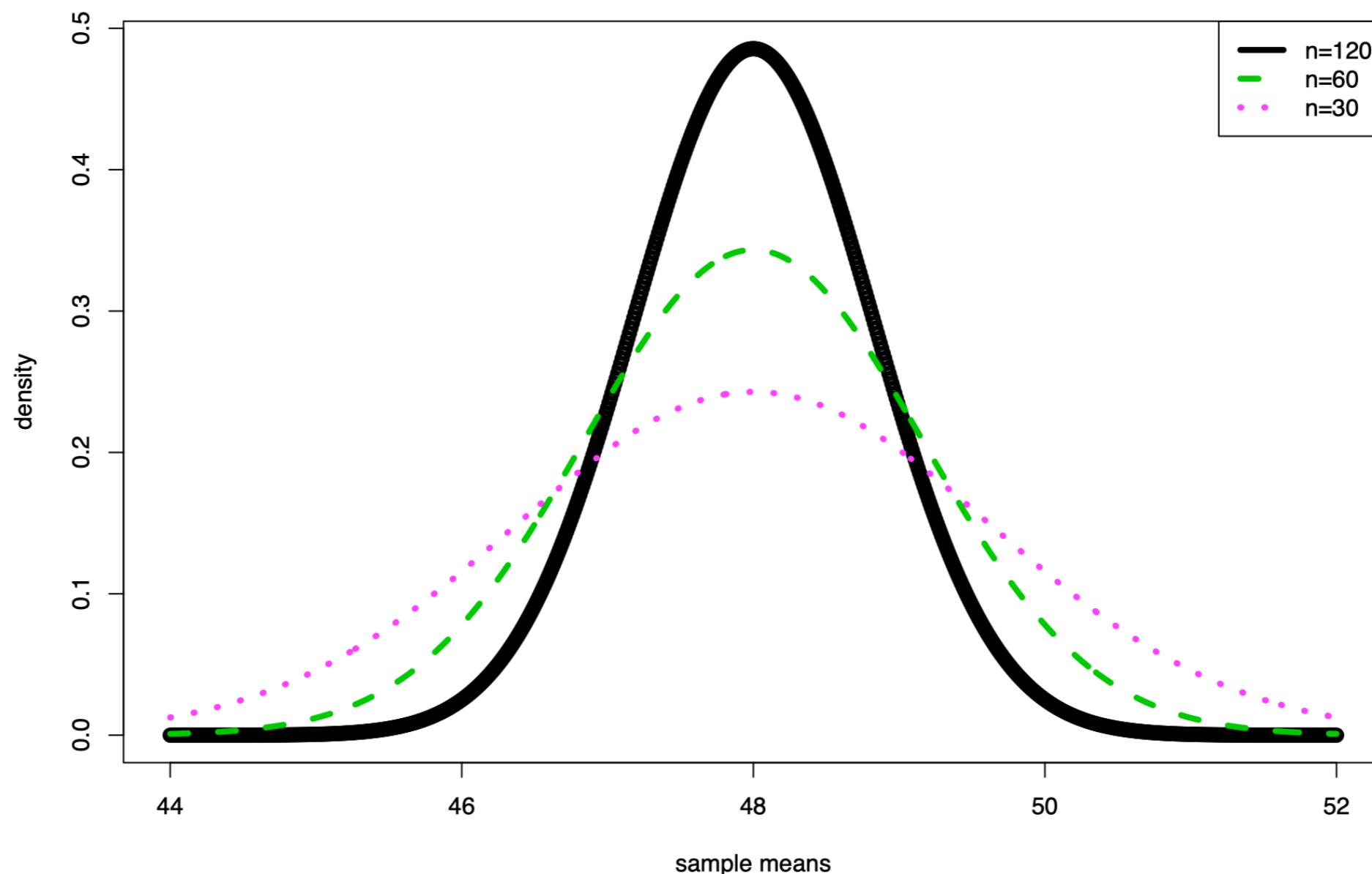
This means that, if many sources additively contribute to an experimental measurement, independent measurements will be approximately normally distributed.

This is why statisticians so often assume that experimental measurements follow normal distributions.

## Impact of sample size on sampling distribution

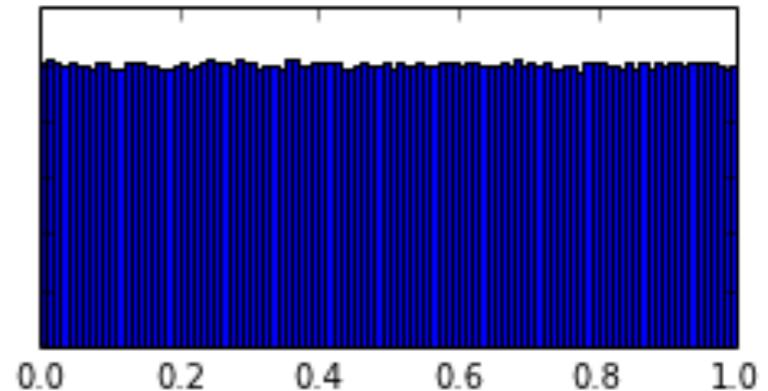
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Sample 1 (n=30); sample 2 (n=60); sample 3 (n=120)

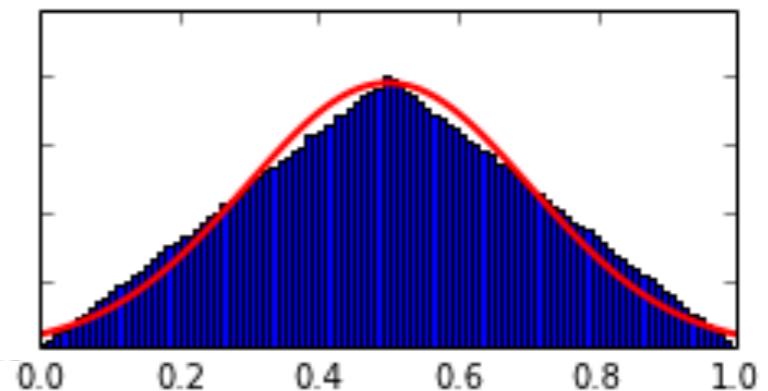


Suppose  $x_1, x_2, \dots, x_N$  are drawn from a uniform (i.e. flat) probability distribution that stands from 0 and 1

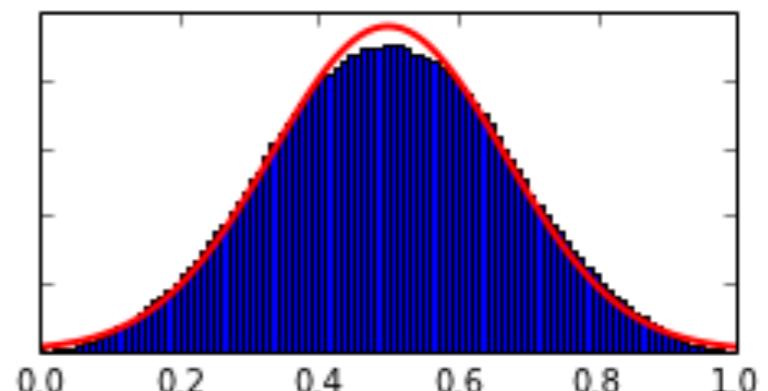
$x_1$



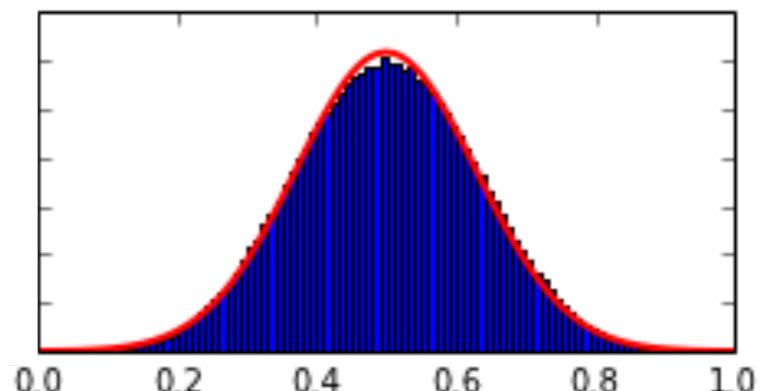
$$\frac{x_1 + x_2}{2}$$



$$\frac{x_1 + x_2 + x_3}{3}$$



$$\frac{x_1 + x_2 + x_3 + x_4 + x_5}{5}$$



## **Example 1: Human Sex Ratio**

## The human sex ratio at birth is slightly skewed towards boys rather than girls.

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	count
male	484382
female	453841
total	938223



**probability of male birth**

estimate: 51.63%

95% CI: [51.53%, 51.73%]

Arbuthnot J (1711). An Argument for Divine Providence, taken from the Constant Regularity observed in the Births of both Sexes.

We assume the number of male babies (versus female babies) is drawn from a binomial distribution

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## data

$n = 484,382$ : number of male births

$N = 938,223$ : total number of births

## model

$$n \sim \text{Binom}(q, N)$$

$q$ : probity of a male birth

The assume probability distribution is called the sampling distribution

## goals

1. Compute a best estimate  $\hat{q}$  for  $q$
2. Compute a confidence interval for  $q$

## The standard estimate of probability is just the ratio of counts

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$n = 484,382$ : number of male births

$N = 938,223$ : total number of male births

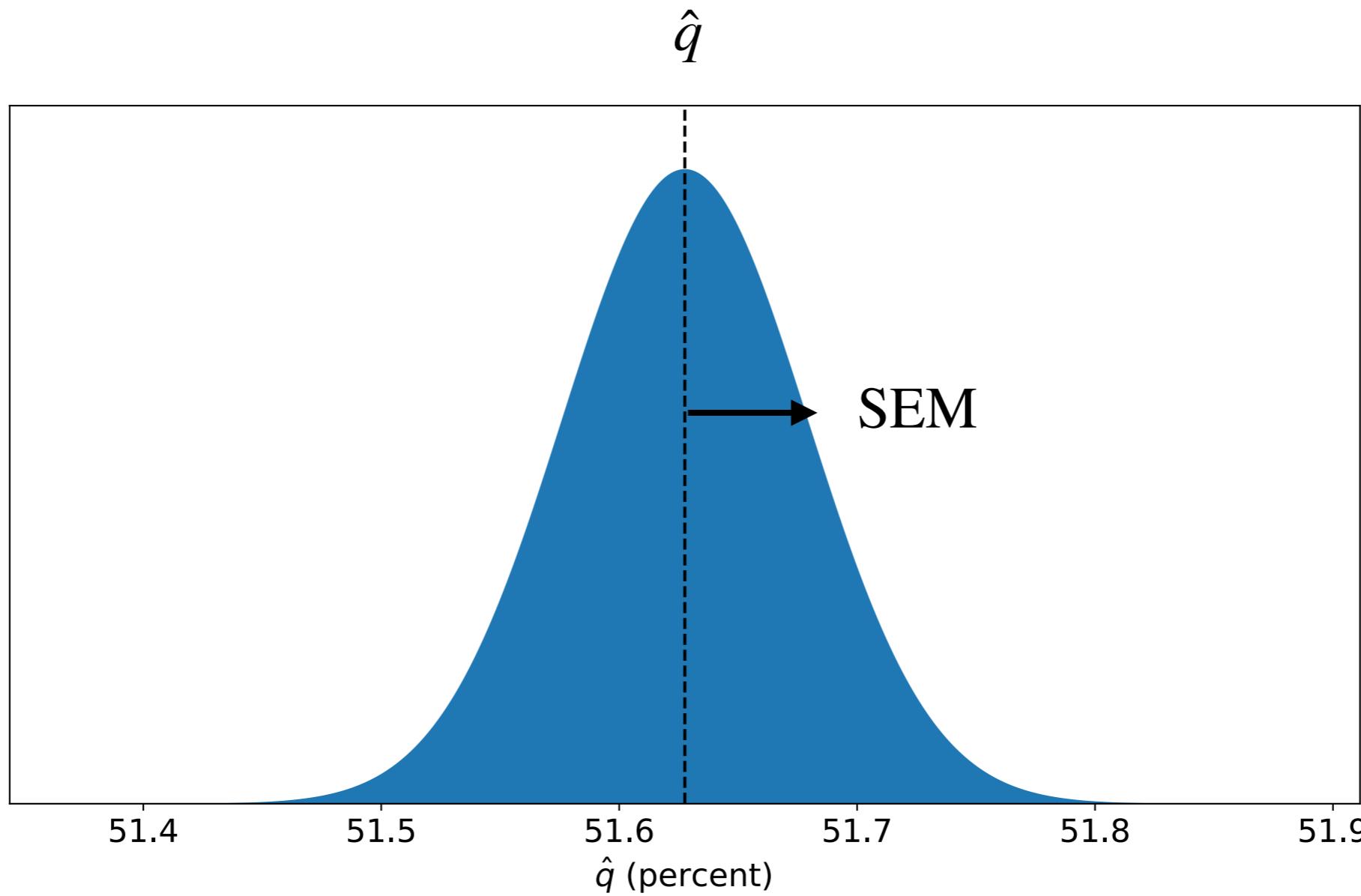
$\hat{q} = \frac{n}{N} = 51.63\%$  : estimated probability of a newborn being male

The lingering uncertainty in  $q$  is (verily nearly) described by a normal distribution centered on the estimate  $\hat{q}$ .

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The standard deviation of this distribution is called the standard error of the mean (SEM).

$$\text{SEM} = \sqrt{\hat{q}(1 - \hat{q})/N}$$

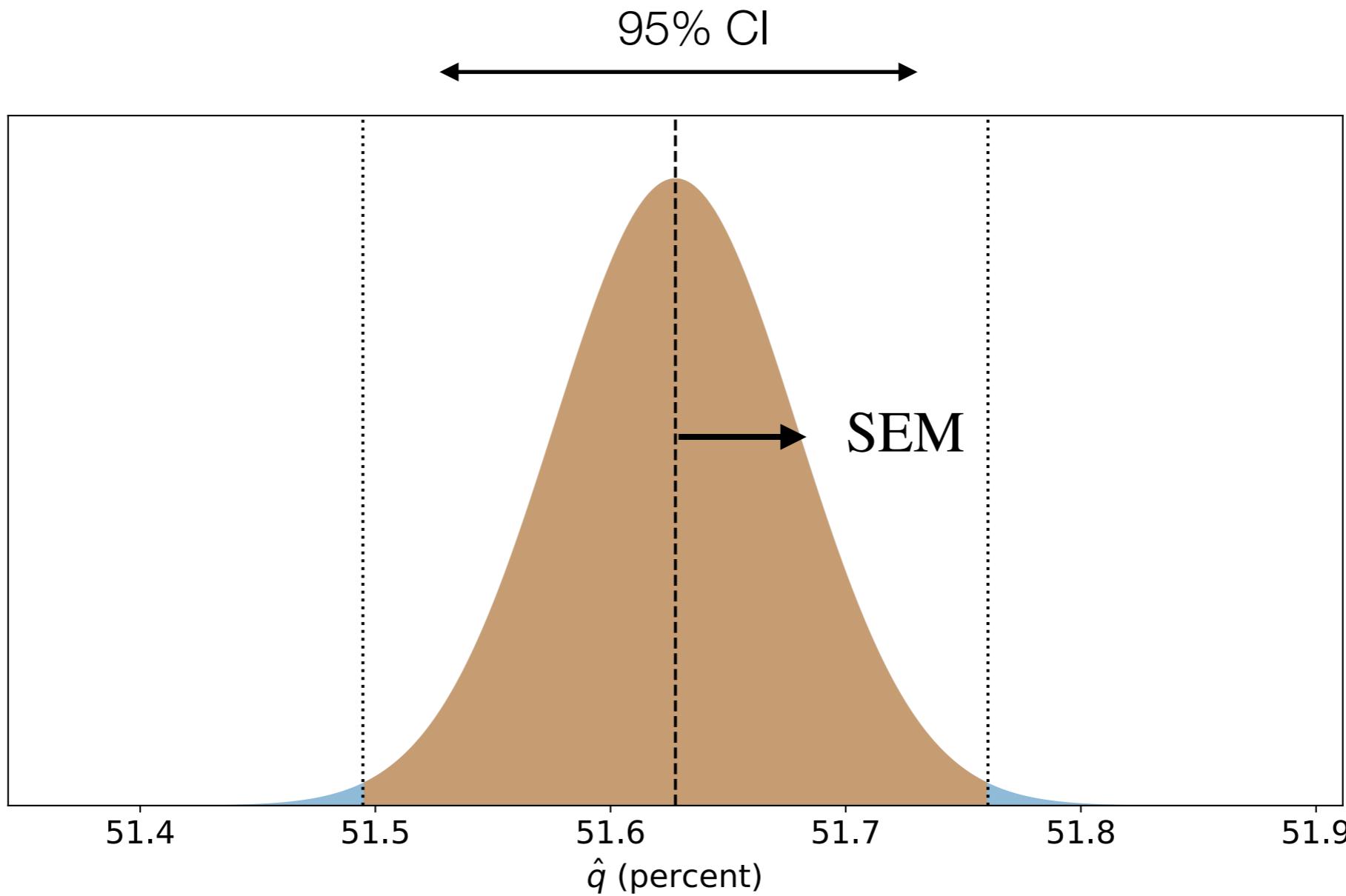


The 95% confidence interval, describing plausible values of  $q$ , is computed using both  $\hat{q}$  and SEM.

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The corresponding 95% confidence interval (CI) is

$$[\hat{q} - W, \hat{q} + W] \text{ where } W = 1.96 \times \text{SEM}$$



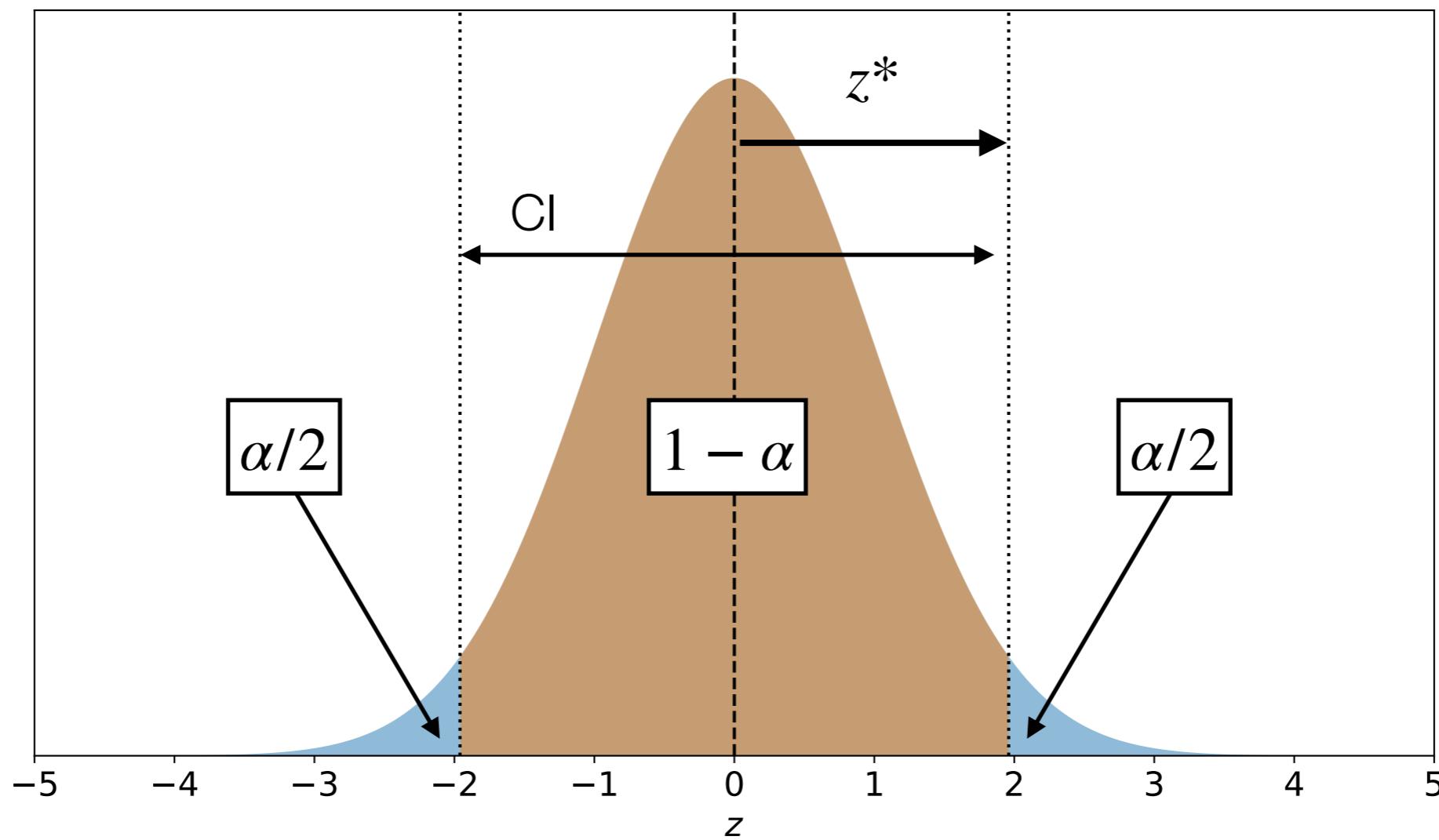
## Uncertainty in $q$ is summarized by a $z$ -statistic

The  $z$ -statistic is defined by:  $z = \frac{q - \hat{q}}{\text{SEM}}$

Because of the central limit theorem,  $z \sim \text{Normal}(0, 1)$ .

The user chooses a value for  $\alpha$ , the probability that  $q$  is not within the confidence interval.

Choosing  $\alpha$  fixes the value of  $z^*$ . Using  $\alpha = 5\%$  gives  $z^* = 1.96$ .

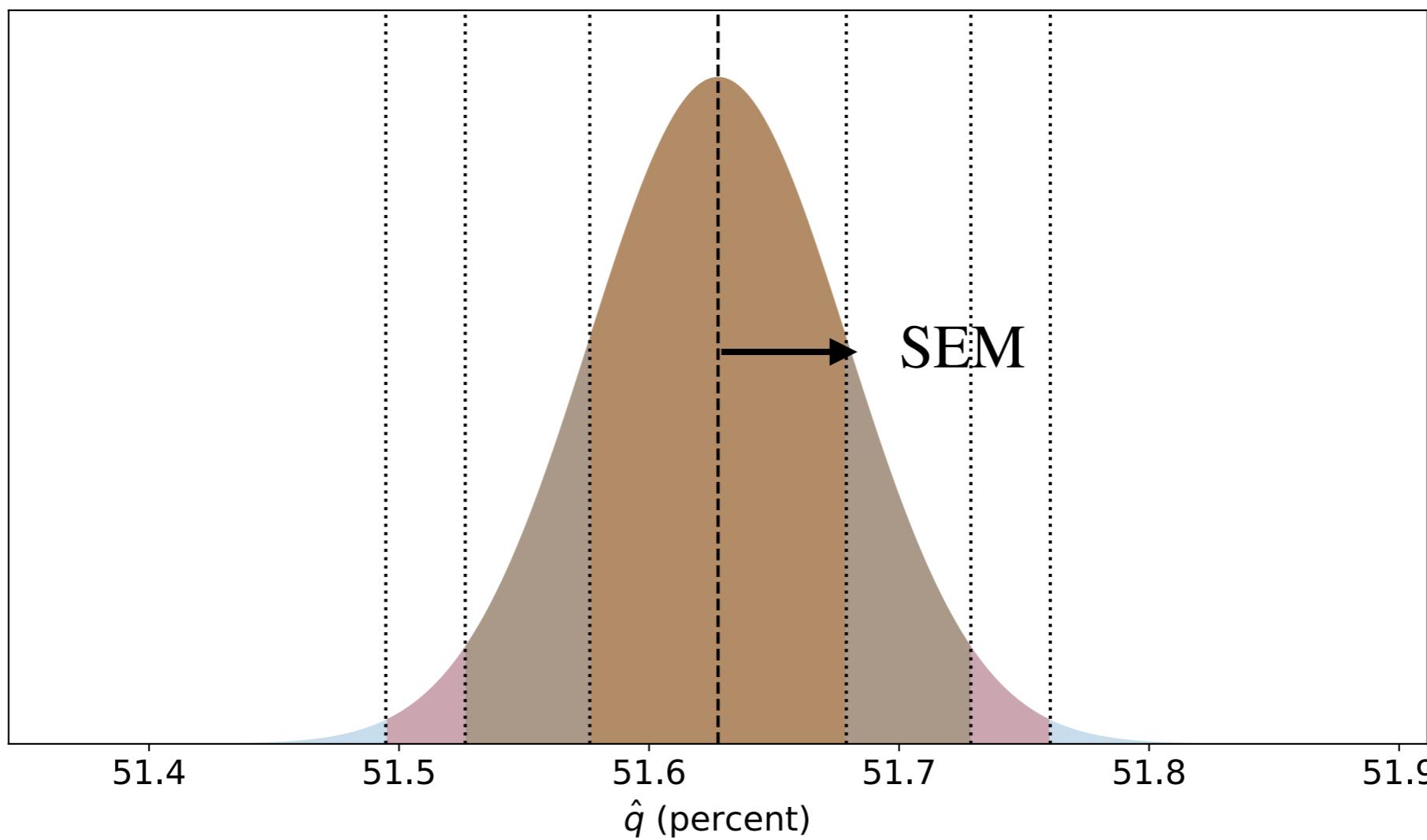


## Confidence intervals of different stringency can be computed using different z-statistic thresholds

Other confidence intervals are given by  $[\hat{q} - W, \hat{q} + W]$  where

$$\text{margin of error: } W = z^* \times \text{SEM}$$

- $\xleftarrow{\hspace{2cm}}$  99% CI:  $z^* = 2.58$
- $\xleftarrow{\hspace{1.5cm}}$  95% CI:  $z^* = 1.96$
- $\xleftarrow{\hspace{0.8cm}}$  68% CI:  $z^* = 0.99$



## **Example 2: Healthy Human Body Temperature**

## Example 2: Human body temperature

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<b>Body Temp</b>	<b>Sex</b>	<b>Heart Rate</b>
96.3	2	70
96.7	2	71
96.9	2	74
97.0	2	80
97.1	2	73
97.1	2	75
97.1	2	82
97.2	2	64
97.3	2	69
97.4	2	70

⋮

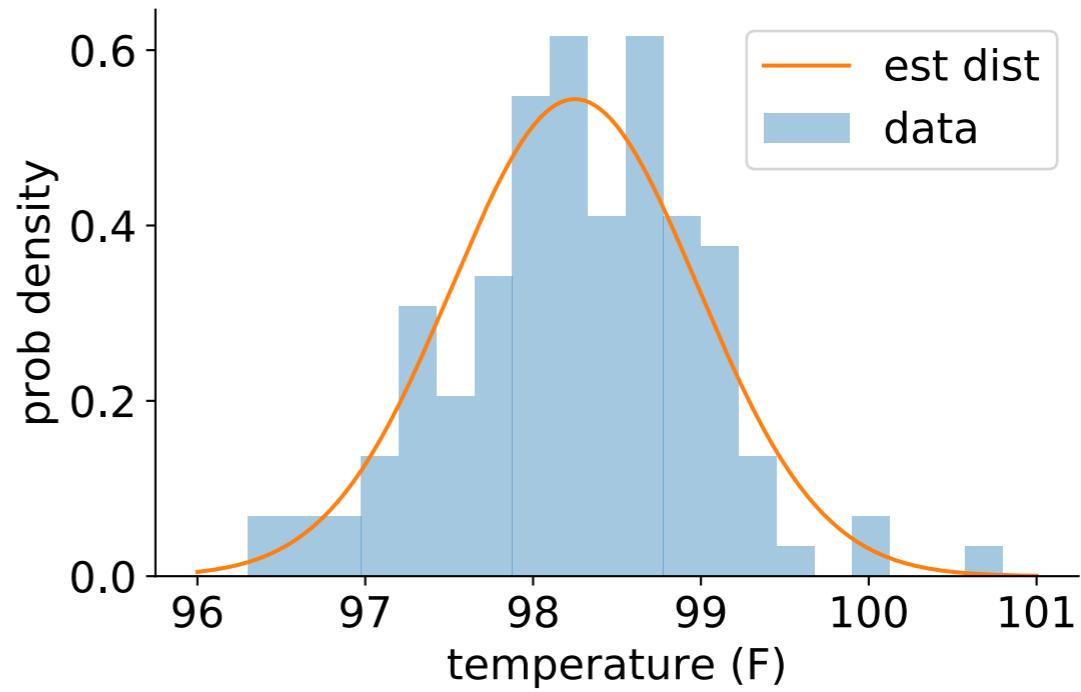
Mackowiak PA, Wasserman SS, Levine MM. (1992) A Critical Appraisal of 98.6°F, the Upper Limit of the Normal Body Temperature, and Other Legacies of Carl Reinhold August Wunderlich. *JAMA*. 268(12):1578–1580.

(Sex: 1 = female, 2 = male)

## Example 2: Human Body Temperature

We model temperature using a normal distribution

Body Temp
96.3
96.7
96.9
97.0
97.1
97.1
97.1
97.2
97.3
97.4



**temperature mean  $\mu$**

estimate: 98.25 F

95% CI: [98.12 F, 98.38 F]

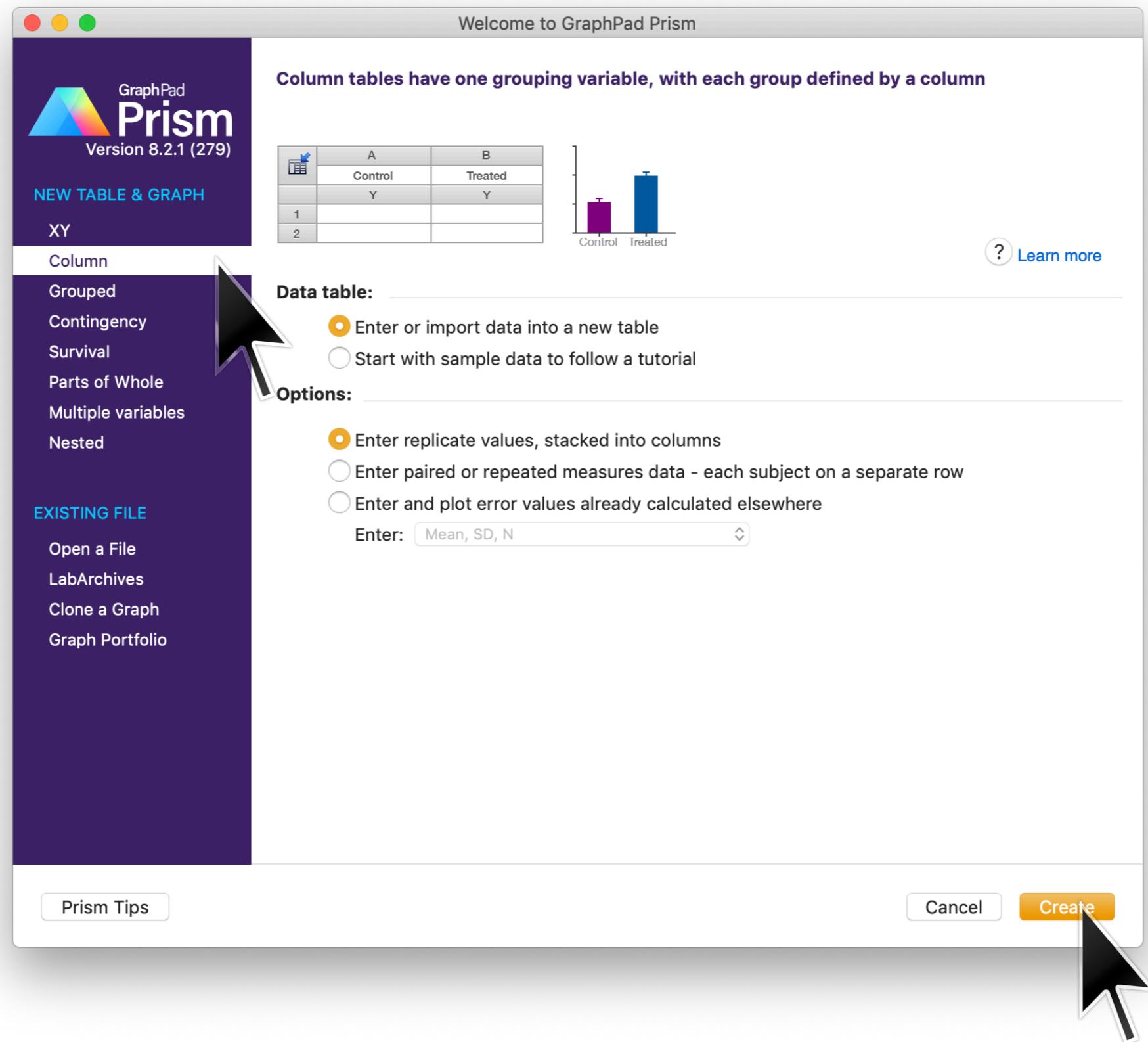
⋮

**temperature standard deviation  $\sigma$**

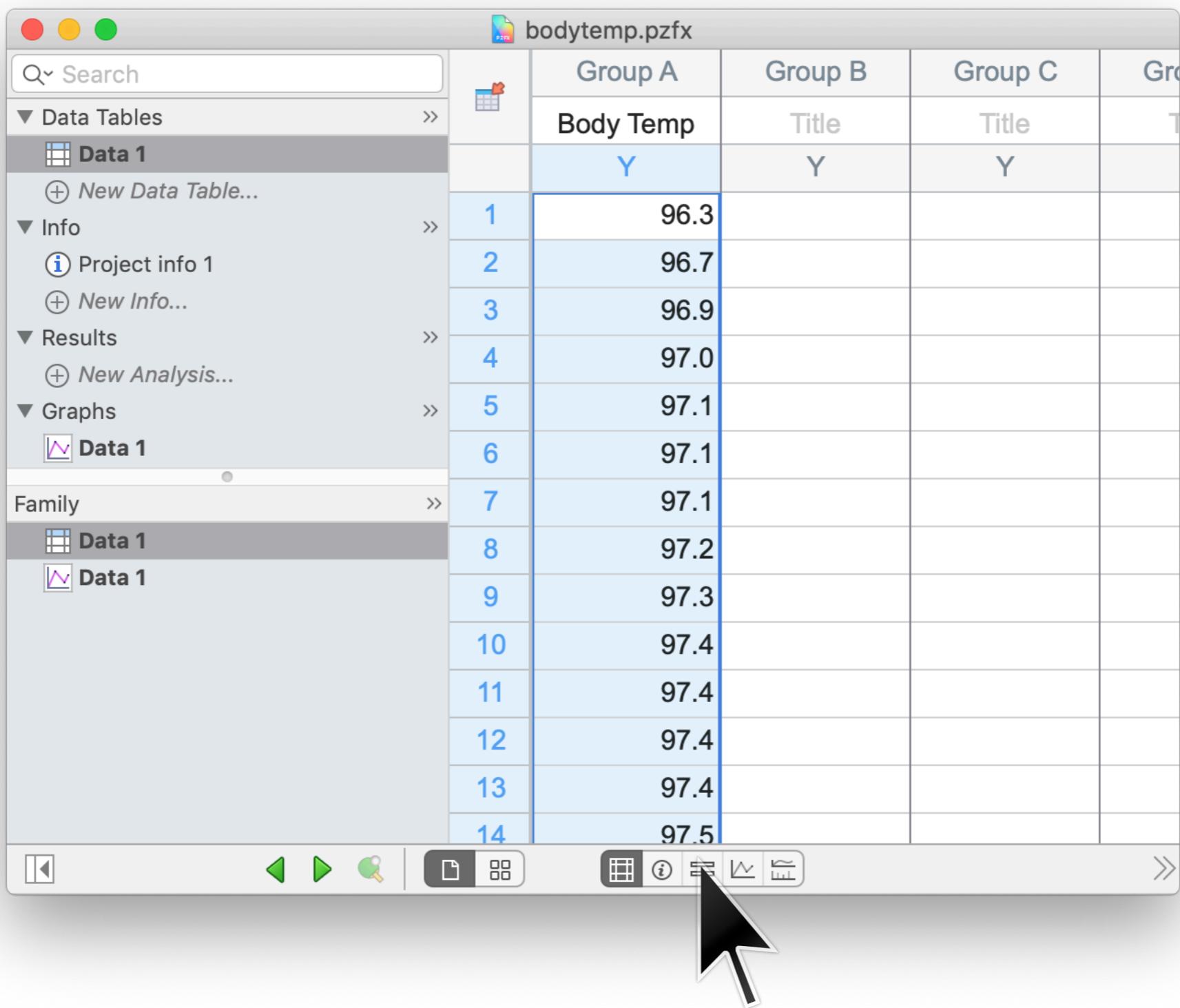
estimate: 0.73 F

95% CI: [0.65 F, 0.83 F]

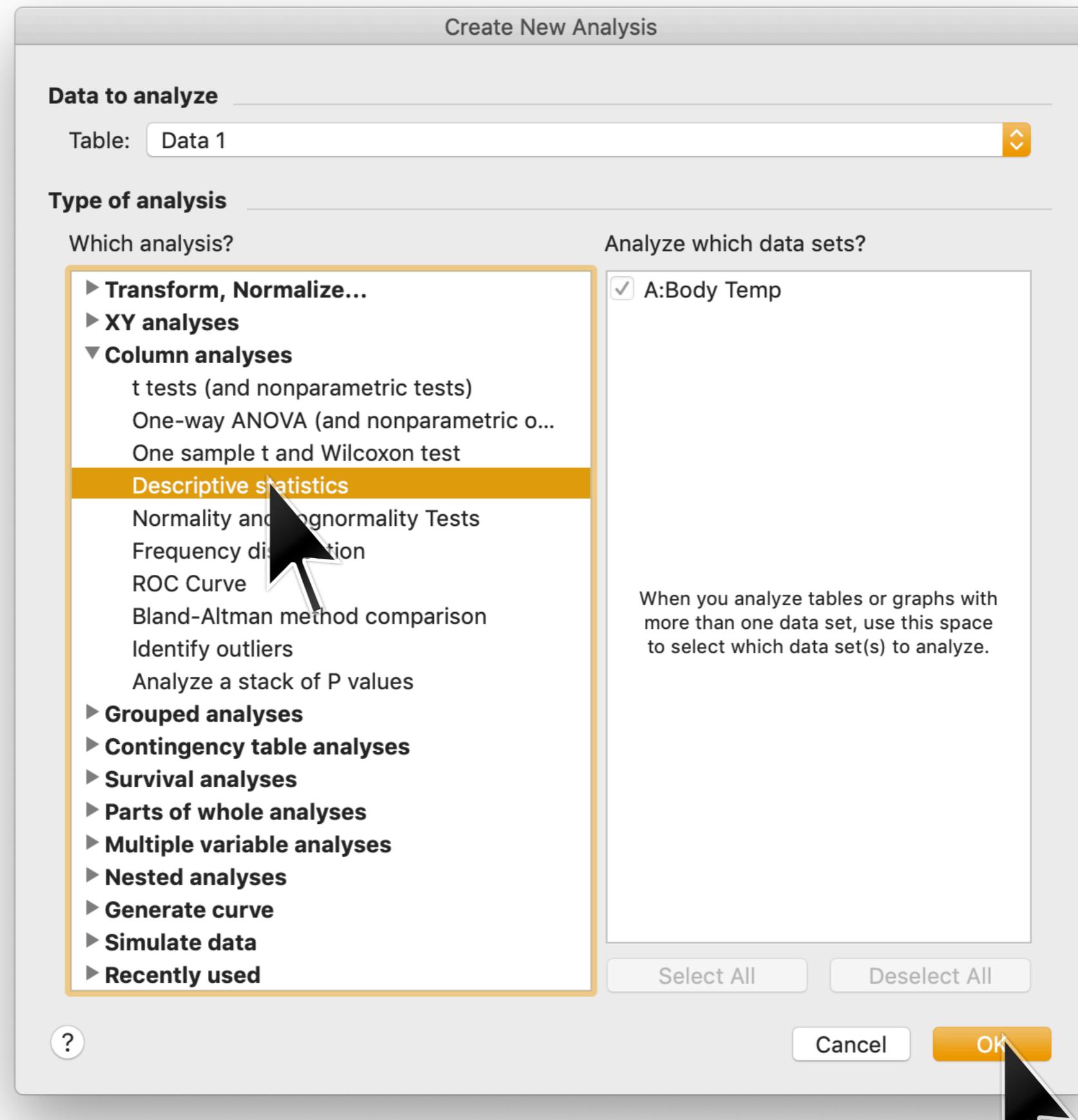
# How to do this in PRISM



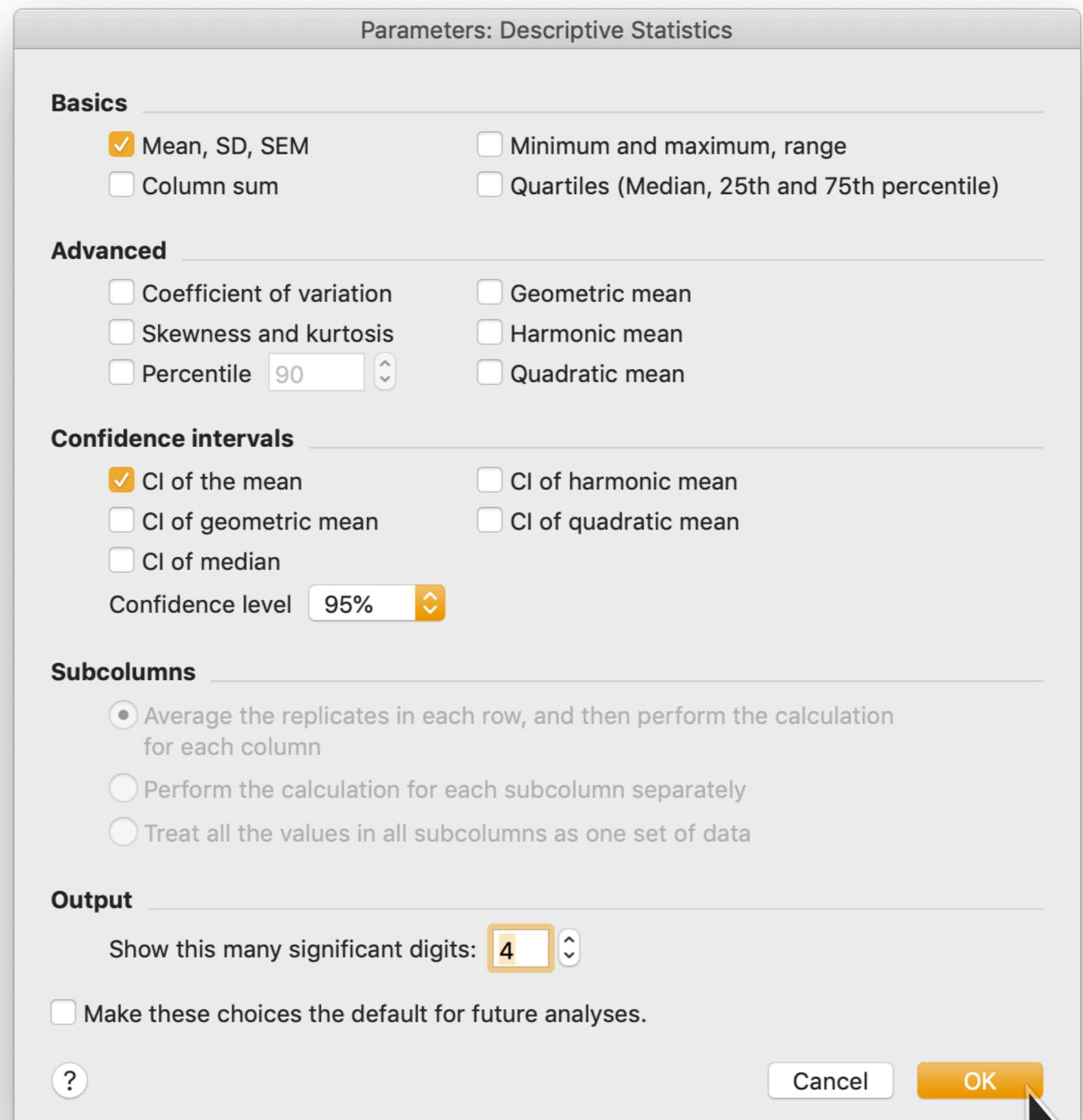
## How to do this in PRISM



# How to do this in PRISM



# How to do this in PRISM



## How to do this in PRISM

The screenshot shows the PRISM software interface with the project file "bodytemp.pzfx — Edited". The left sidebar contains sections for Data Tables, Info, Results, and Graphs. Under Results, "Descriptive statistics of Data 1" is selected. The main area displays a table titled "Descriptive statistics" with two columns: "A" and "B". The table rows are numbered 1 through 14. The first row, "Body Temp", has "A" set to "Y" and "B" set to "Y". Rows 2 through 13 show descriptive statistics: Number of values (130), Mean (98.25), Std. Deviation (0.7332), Std. Error of Mean (0.06430), Lower 95% CI of mean (98.12), and Upper 95% CI of mean (98.38). Rows 14 through 17 are empty.

	A	B
1 Body Temp	Y	Y
2		
3 Mean	98.25	
4 Std. Deviation	0.7332	
5 Std. Error of Mean	0.06430	
6		
7 Lower 95% CI of mean	98.12	
8 Upper 95% CI of mean	98.38	
9		
10		
11		
12		
13		
14		
15		
16		
17		

## We assume the temperature of a healthy person is drawn from a normal distribution

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### **data**

$$x_1, x_2, \dots, x_N$$

$x_i$ : temperature of individual  $i$  in Fahrenheit

### **model**

$$x \sim \text{Normal}(\mu, \sigma^2)$$

$\mu$ : average body temperature

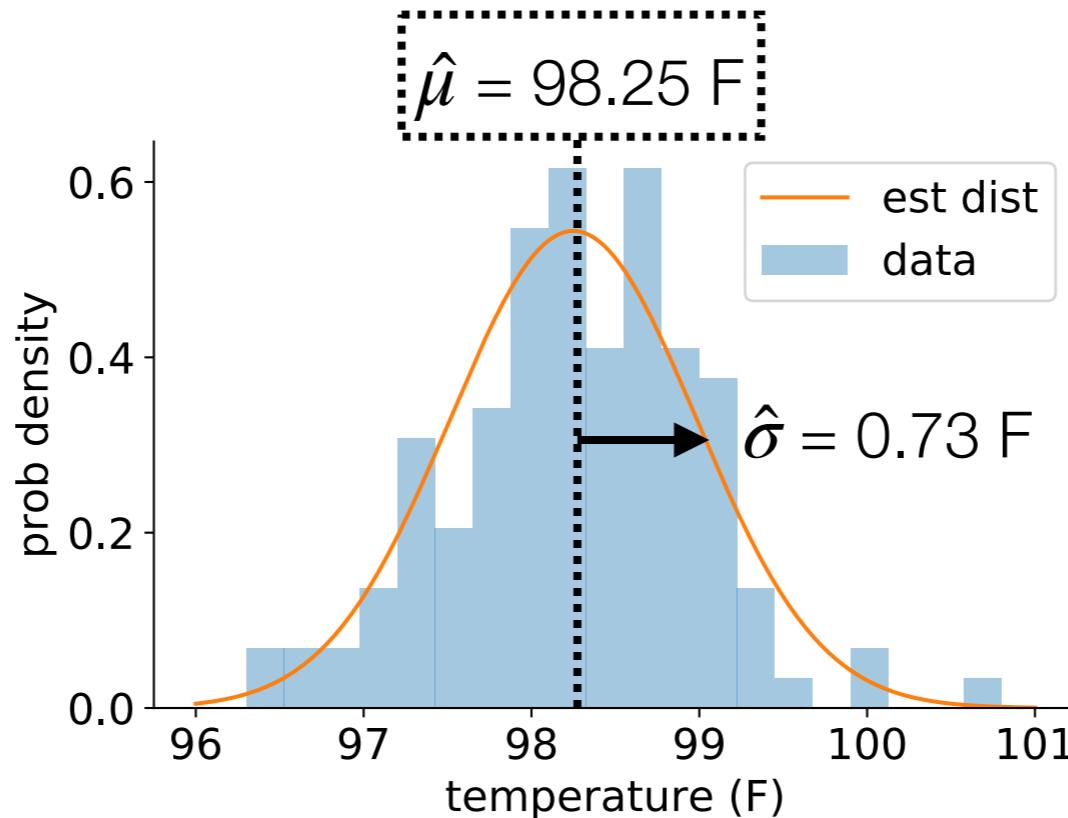
$\sigma$ : standard deviation of temperatures

### **goals**

1. Compute best estimates for both  $\mu, \sigma$
2. Compute confidence intervals for both  $\mu, \sigma$

## We want to infer two parameters from our data

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Here there are two parameters that need to be estimated,  $\mu$  and  $\sigma$

This is unlike with the binomial distribution, where there was only one parameter  $q$ .

## The lingering uncertainty in $\mu$ is described by a t-distribution

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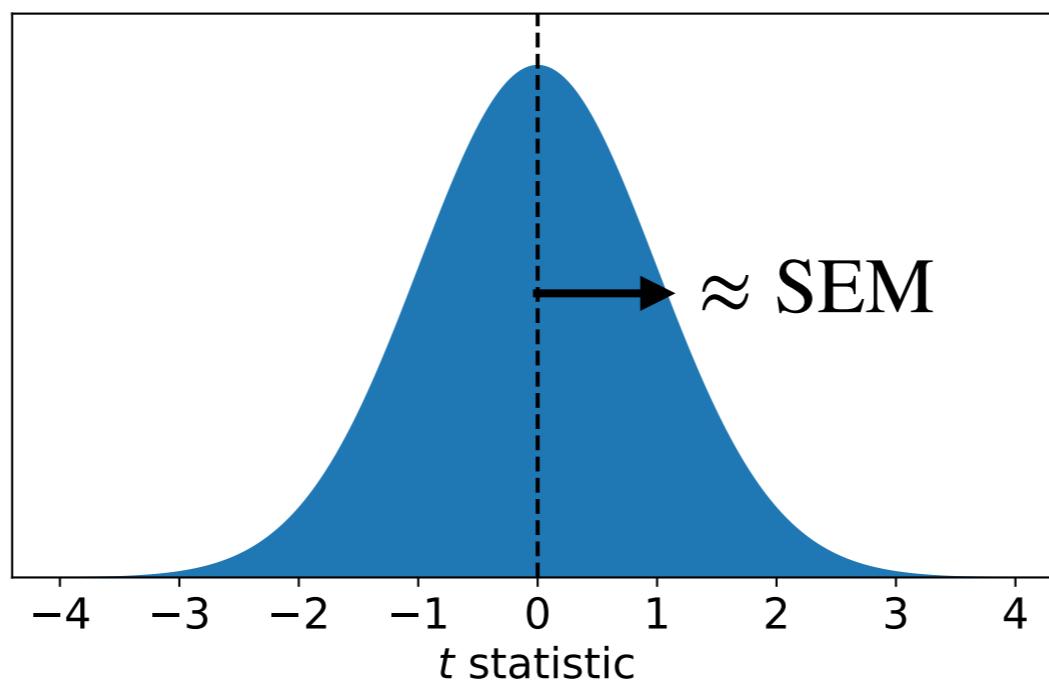
The standard error of the mean (SEM) is given by

$$\text{SEM} = \frac{\hat{\sigma}}{\sqrt{N}}$$

A t-statistic is then used to indicate how strongly  $\mu$  deviates from  $\hat{\mu}$ :

$$t = \frac{\mu - \hat{\mu}}{\text{SEM}}$$

The t-statistic follows a t-distribution  
(almost a normal distribution, but not quite)

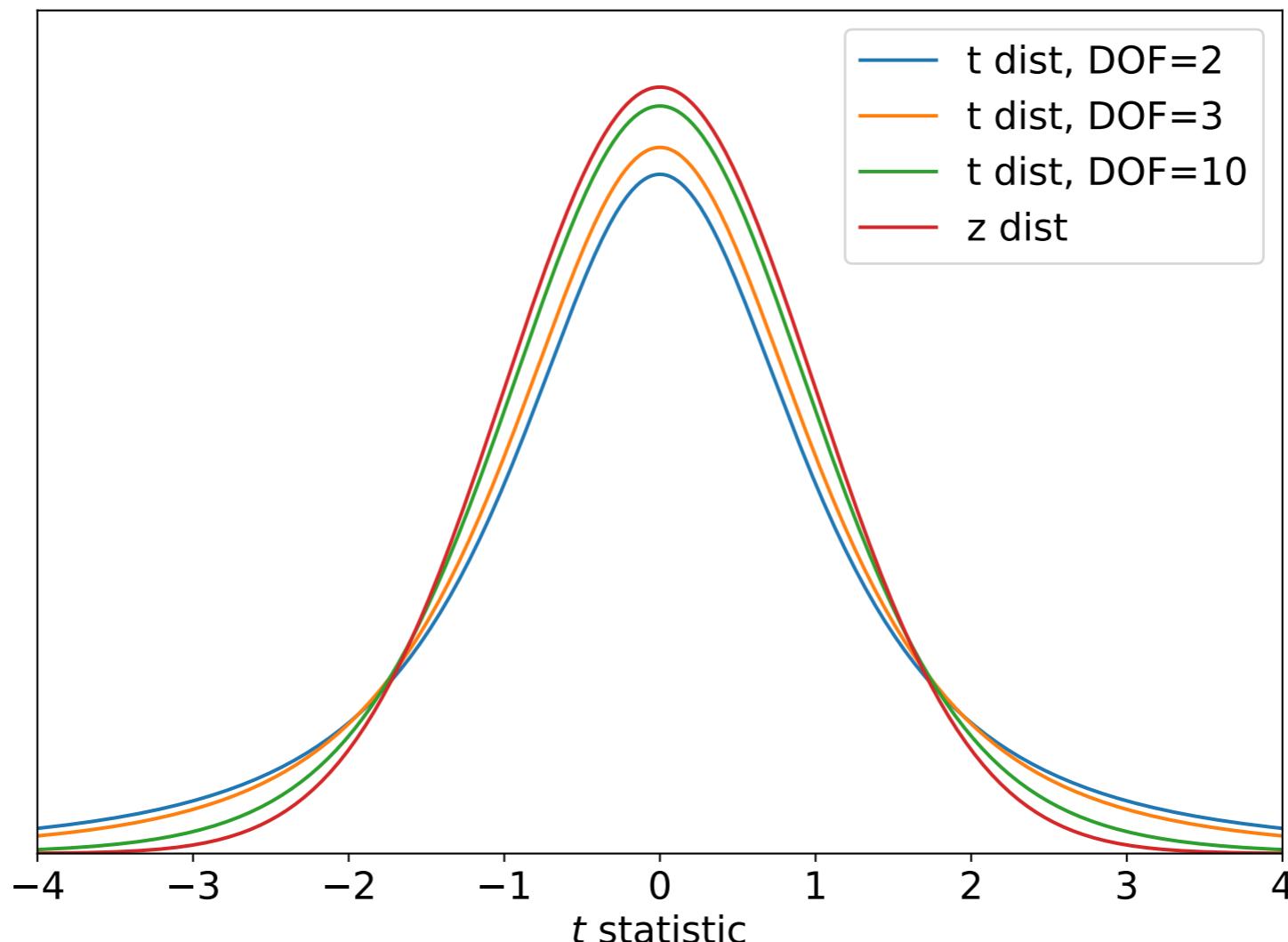


## The shape of the t-distribution is affected by the number of degrees of freedom (DOF)

In this case, we use a t-distribution with DOF given by

$$\text{DOF} = N - 1$$

This is almost indistinguishable from a normal (z) distribution when  $\text{DOF} \gtrsim 10$ .

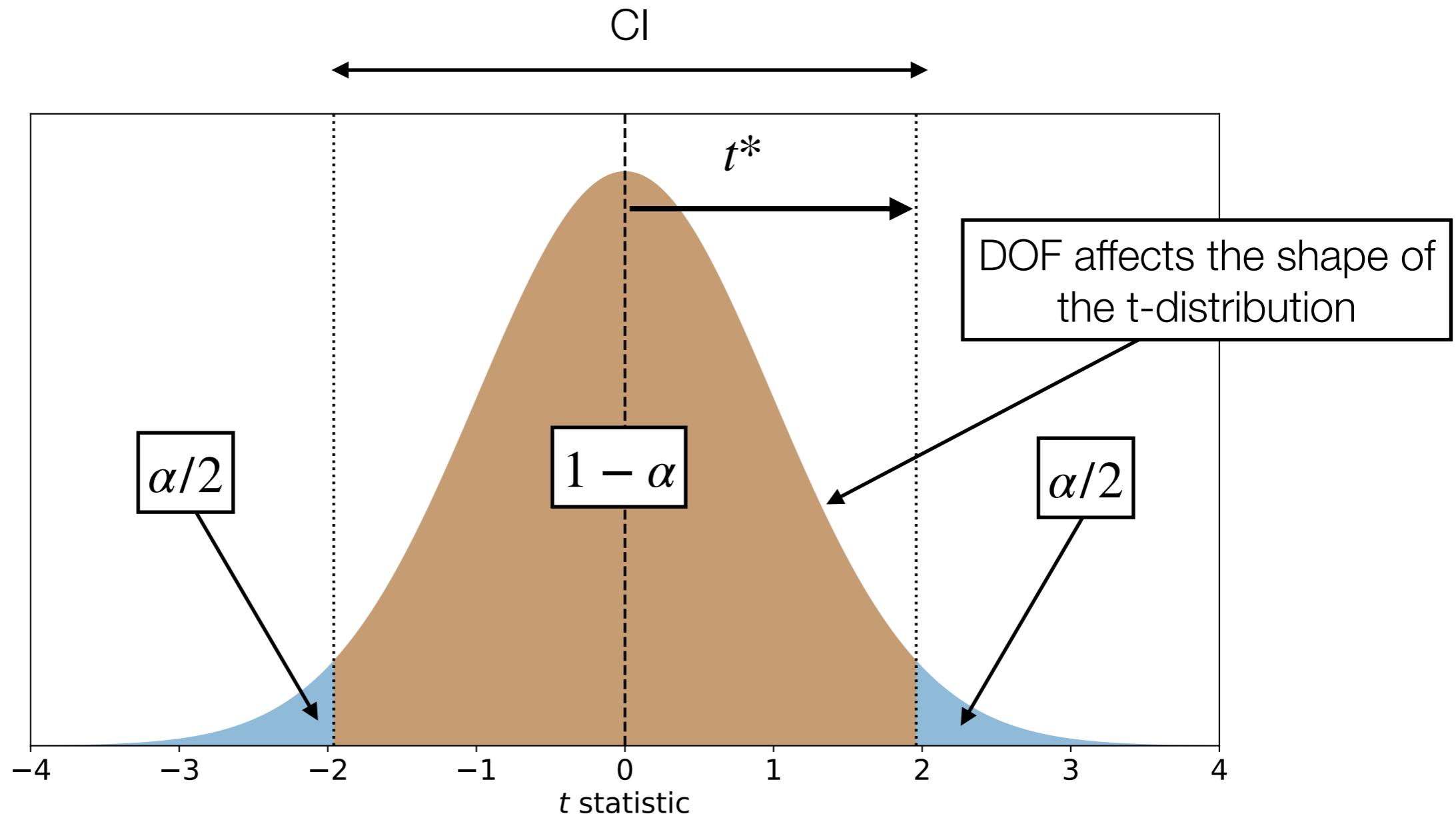


## The t-distribution is used to compute a t-statistic cutoff, which determines the confidence interval

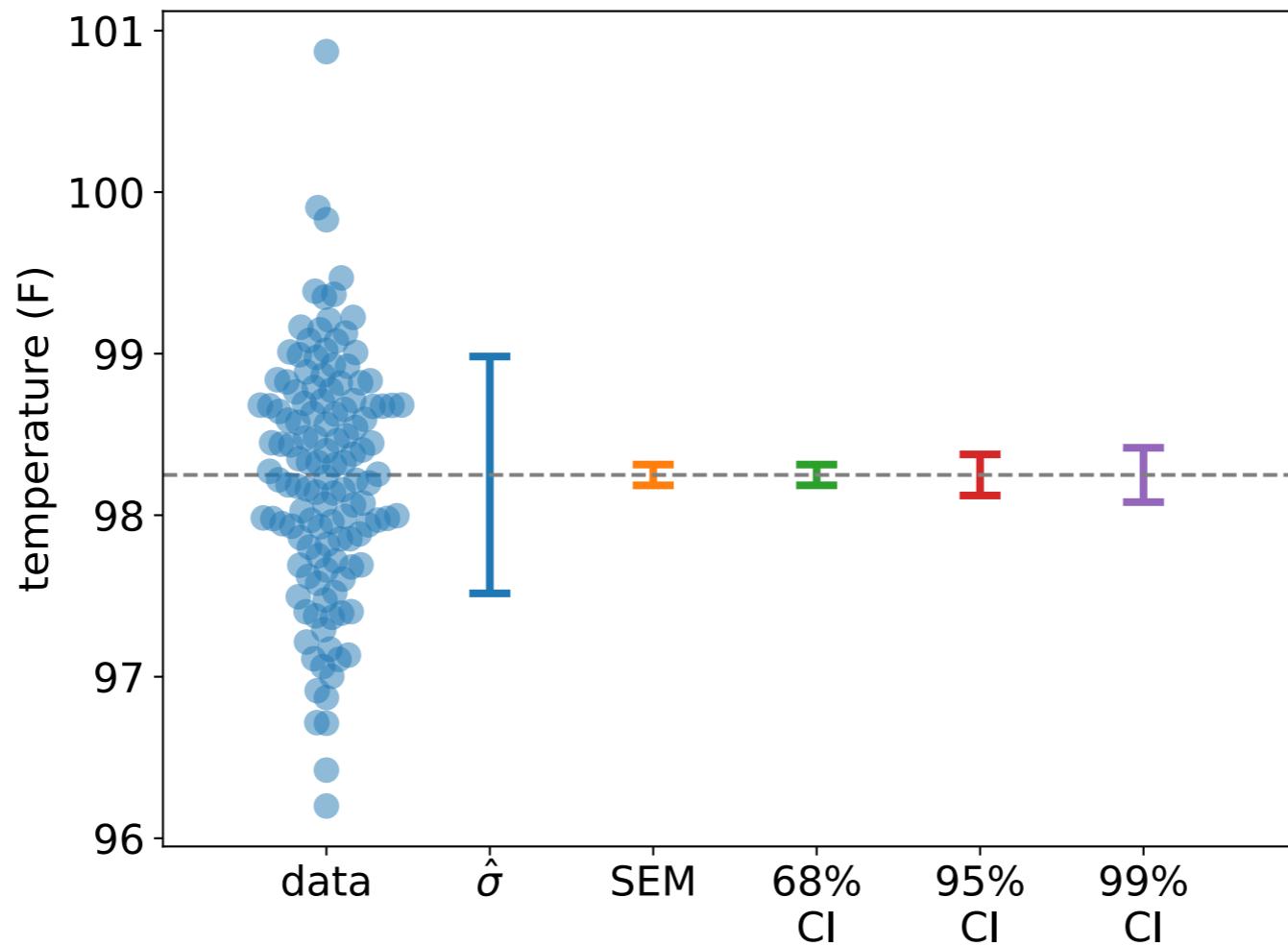
The t-statistic cutoff,  $t^*$ , is determined by both  $\alpha$  and the DOF.

$$\text{margin of error: } W = t^* \cdot \text{SEM}$$

$$\text{confidence interval: } \hat{\mu} \pm W$$



## Confidence intervals (CIs) and standard errors of the mean (SEMs) quantify how uncertain a parameter



SEMs and CIs of the mean quantify the uncertainty in  $\mu$ ,  
not the width of the sampling distribution ( $\hat{\sigma}$ ).

SEMs and CIs decrease in size as the amount of data increases.

CIs increase in size if the required confidence level increases (i.e.,  $\alpha$  decreases)

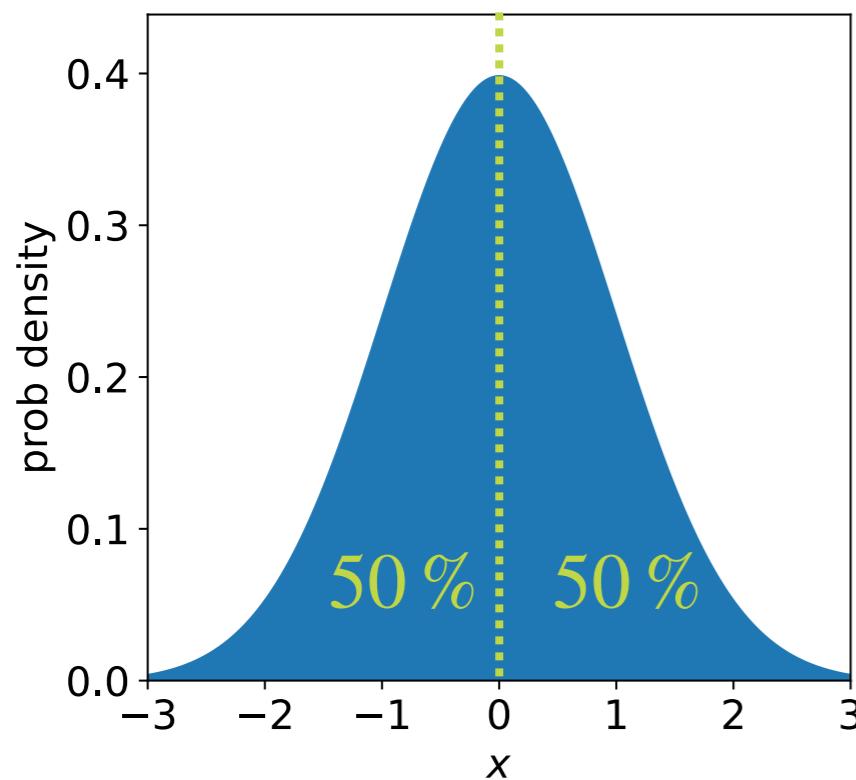
## The median is the standard nonparametric estimate of a distribution's center

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For data: sort the data  $x_1, x_2, x_3, \dots, x_N$  in ascending order. The median is then defined as:

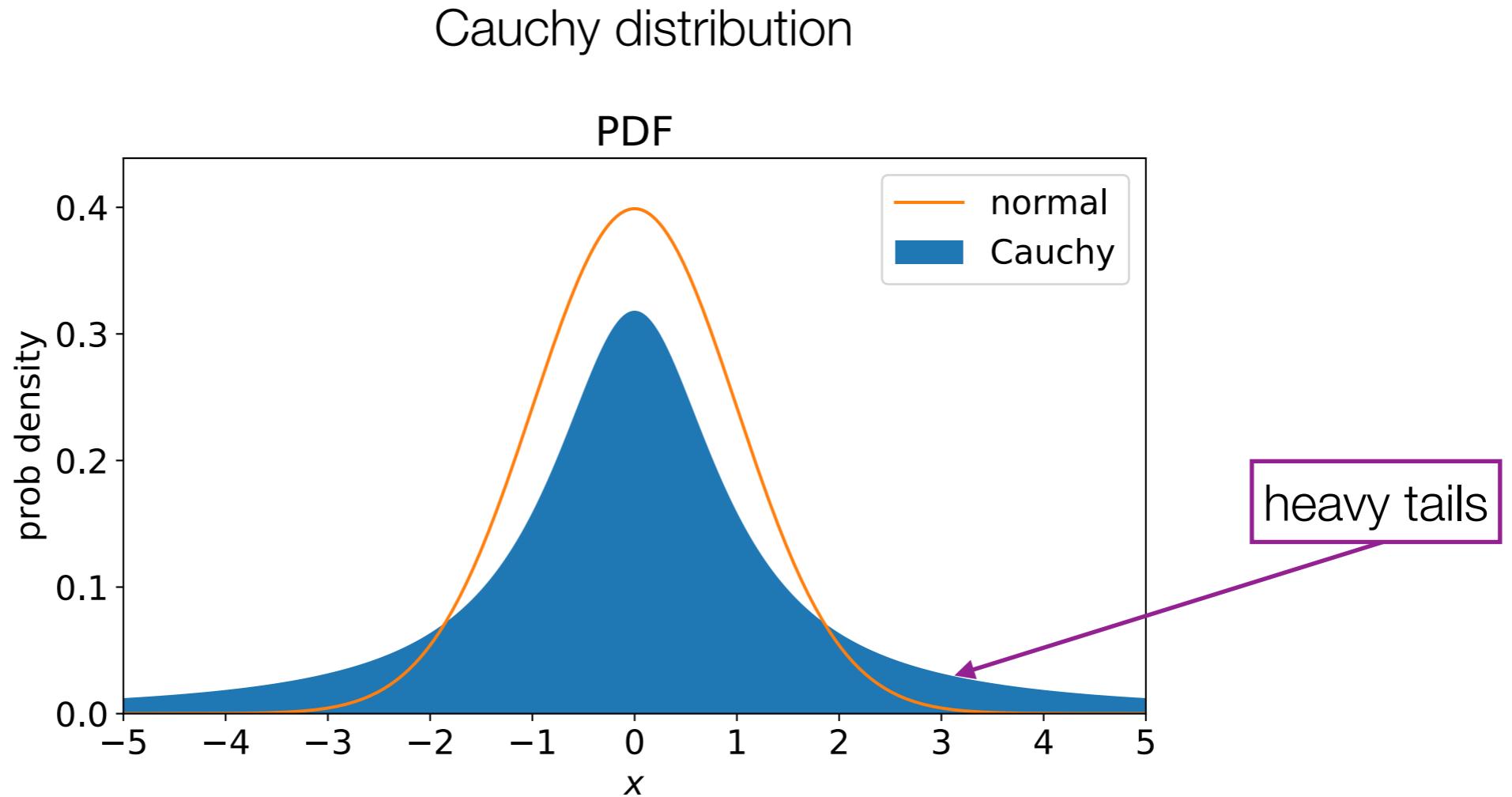
$$\text{median} = q_{50} = \begin{cases} x_{\frac{N+1}{2}} & \text{if } N \text{ odd} \\ \frac{1}{2} \left( x_{\frac{N}{2}} + x_{\frac{N+2}{2}} \right) & \text{if } N \text{ even} \end{cases}$$

For a distribution: the median is the value of  $x$  that separates half the distribution's mass from the other.



The median of a symmetric distribution is equal to its mean

## The median is less sensitive to outliers than the mean



The standard estimate of the mean  $\hat{\mu}$  will not converge as  $N$  becomes large!

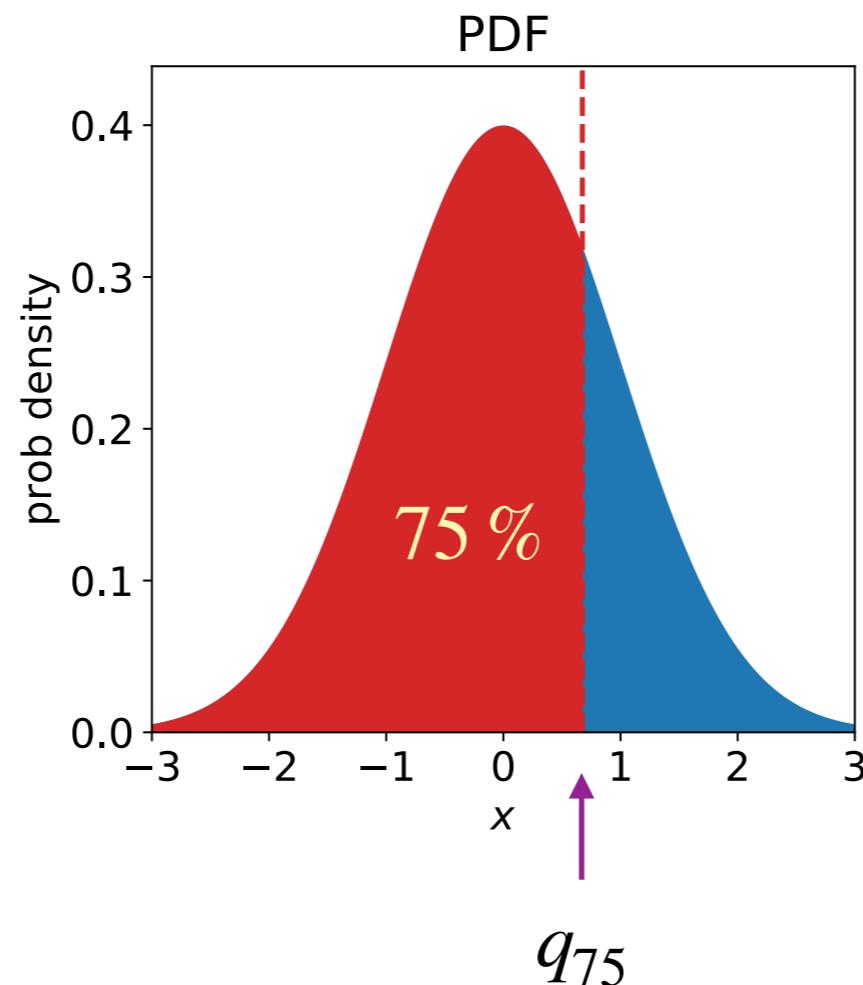
The median  $q_{50}$  does converge, just as quickly as for any distribution.

## Quantiles of a distribution

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More generally, the quantile  $q_K$  of a distribution is the value of  $x$  that bounds  $K\%$  of the distribution's mass.

E.g., the median in the quantile  $q_{50}$

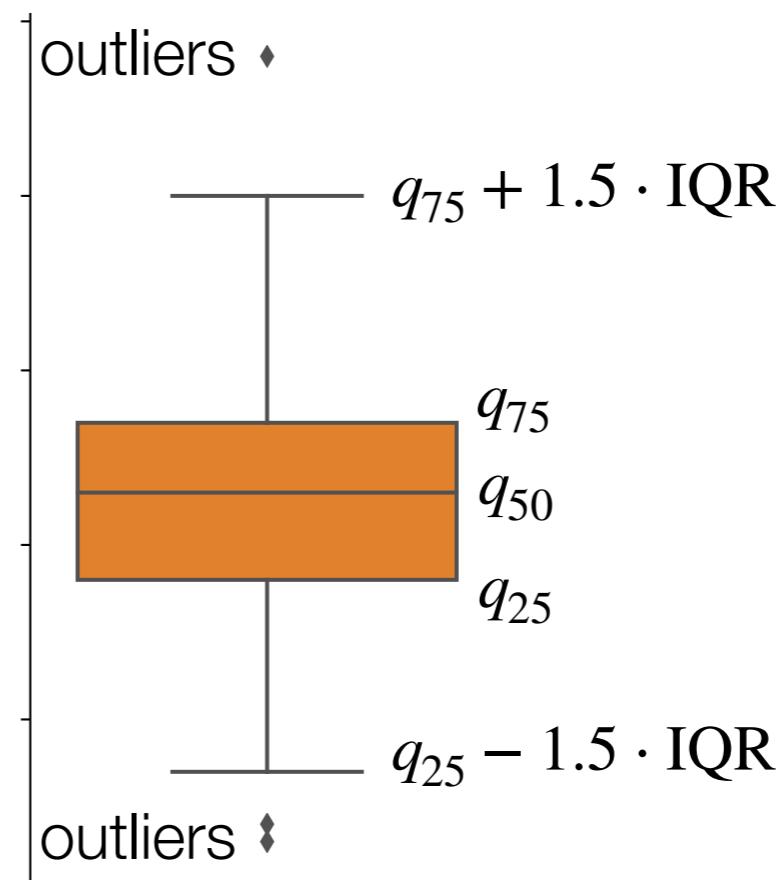
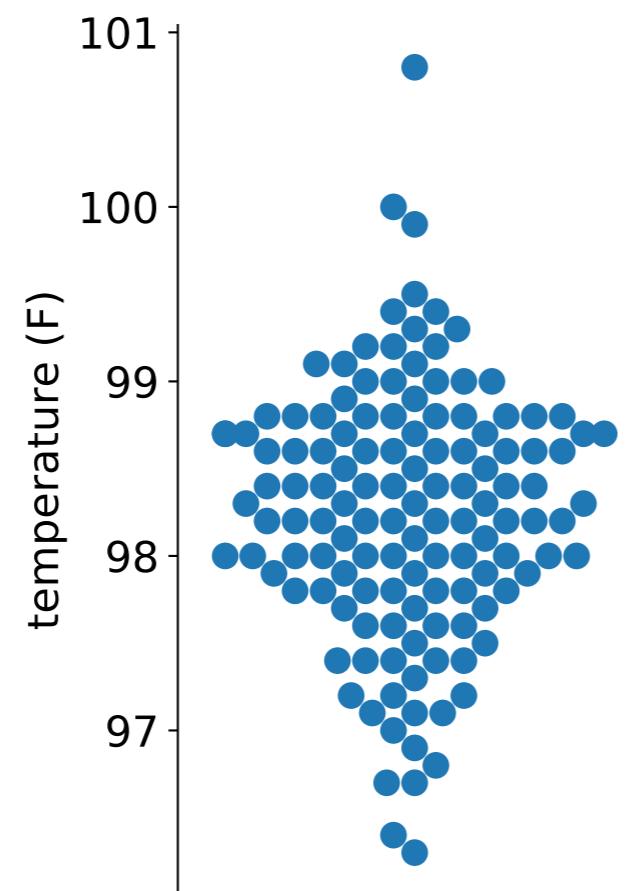


## Box and whisker plots indicate quantiles

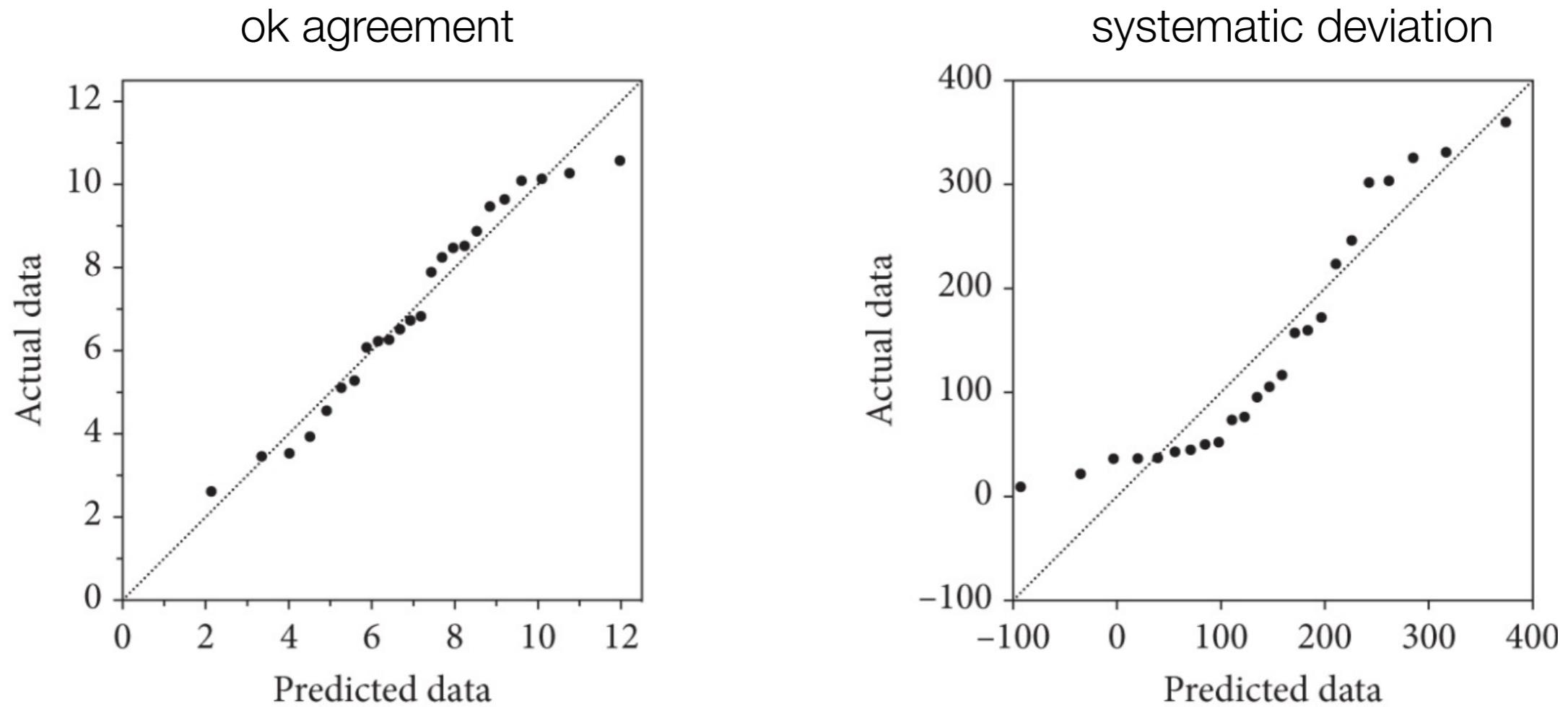
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Interquartile range is defined by

$$\text{IQR} = q_{75} - q_{25}$$



**QQ plots are used to visually test whether data follows an expected distribution.**

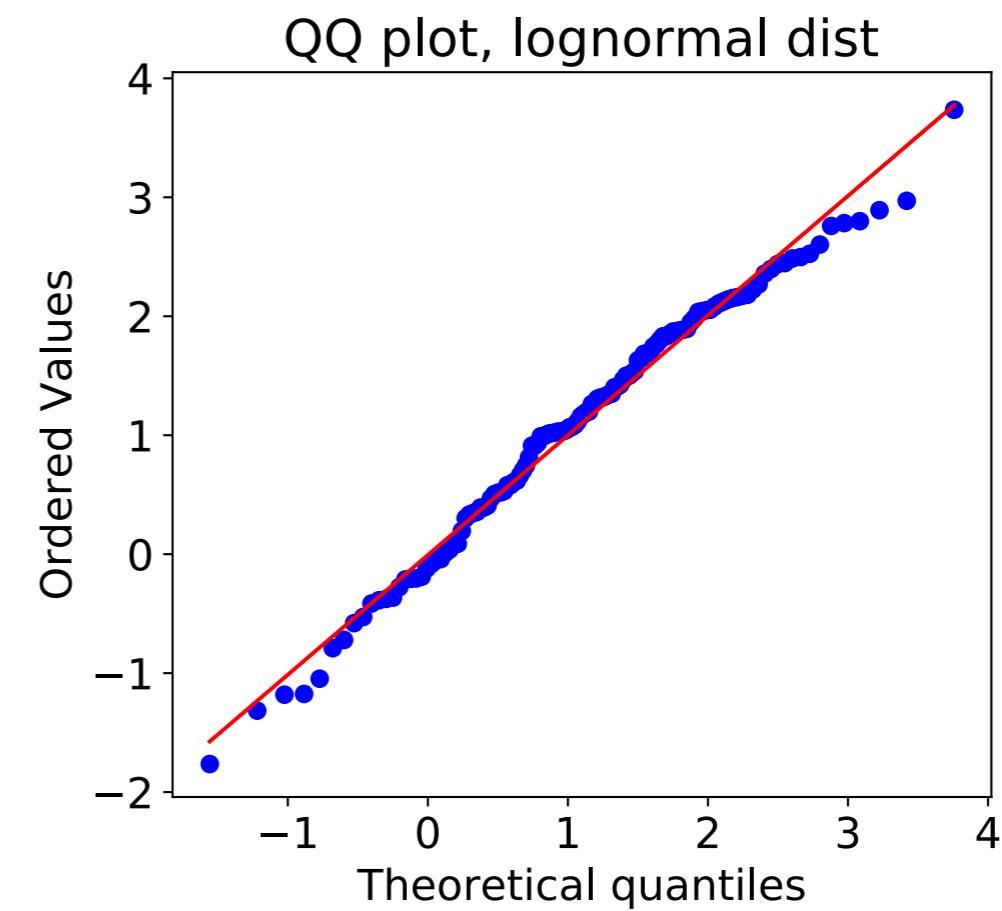
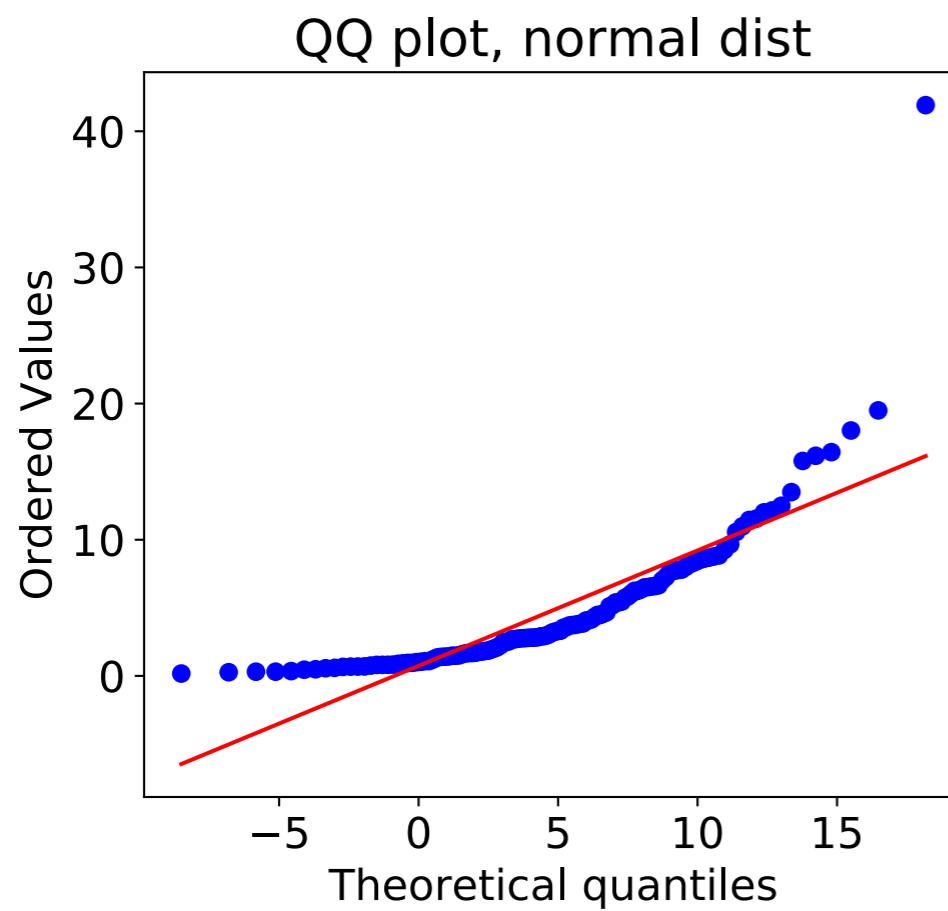
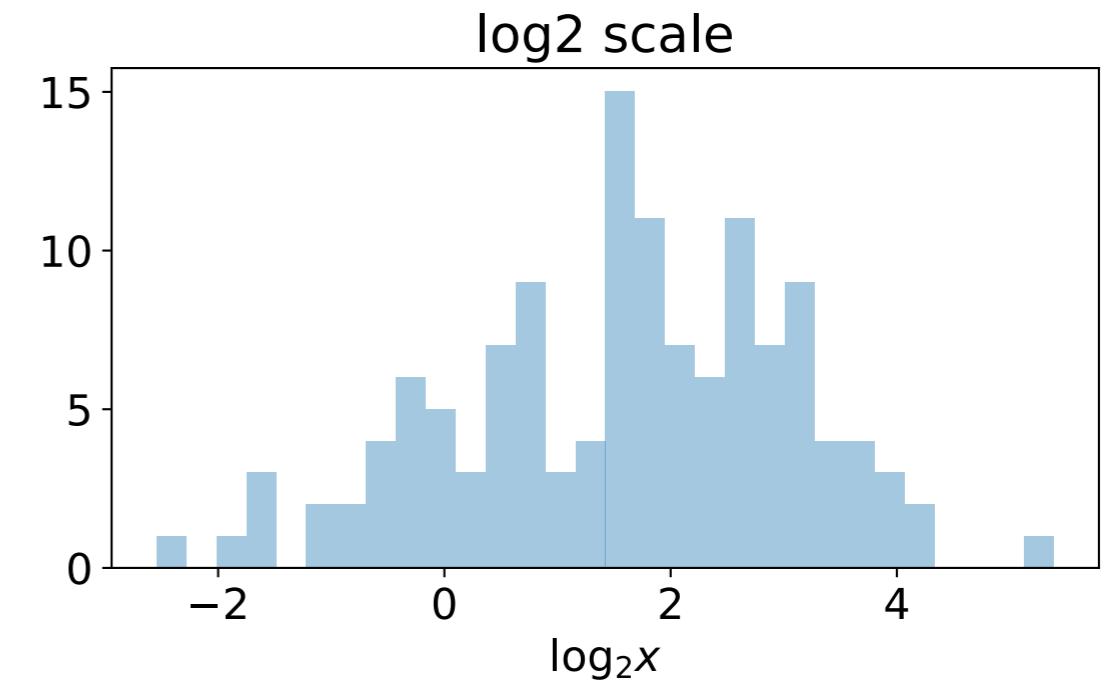
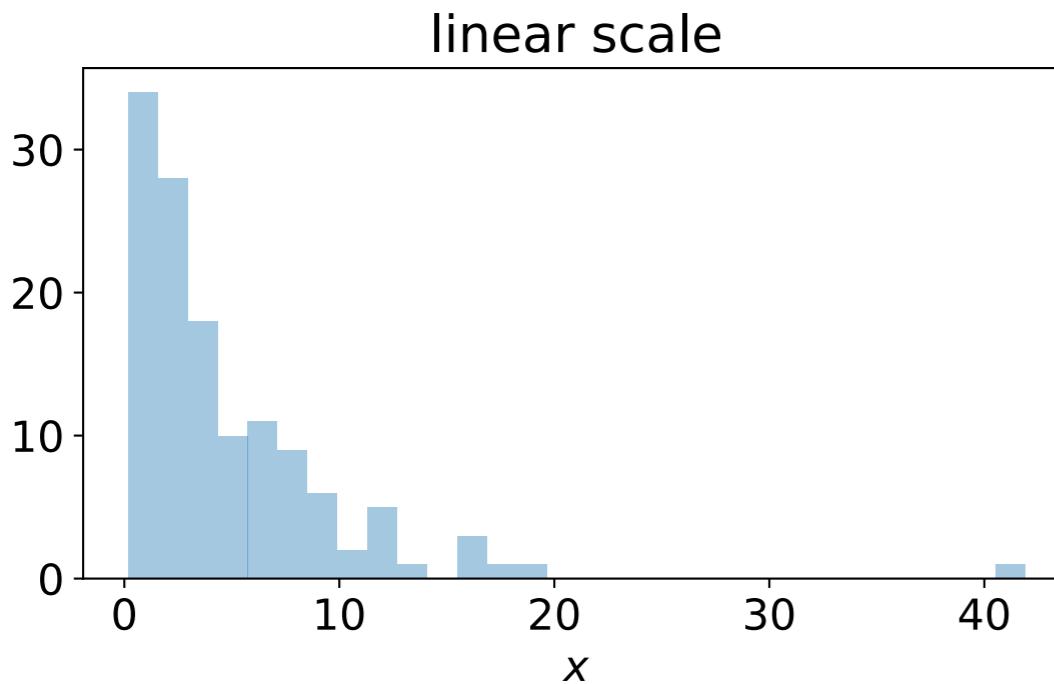


**y axis:** sorted data values  $x_1, x_2, \dots, N$ .

**x axis:** corresponding quantiles  $q_X$  of the inferred distribution, using the percentile values  $X_1, X_2, \dots, X_N$  computed for each data point.

The analysis of a QQ plot is done by eye and making a judgement call.

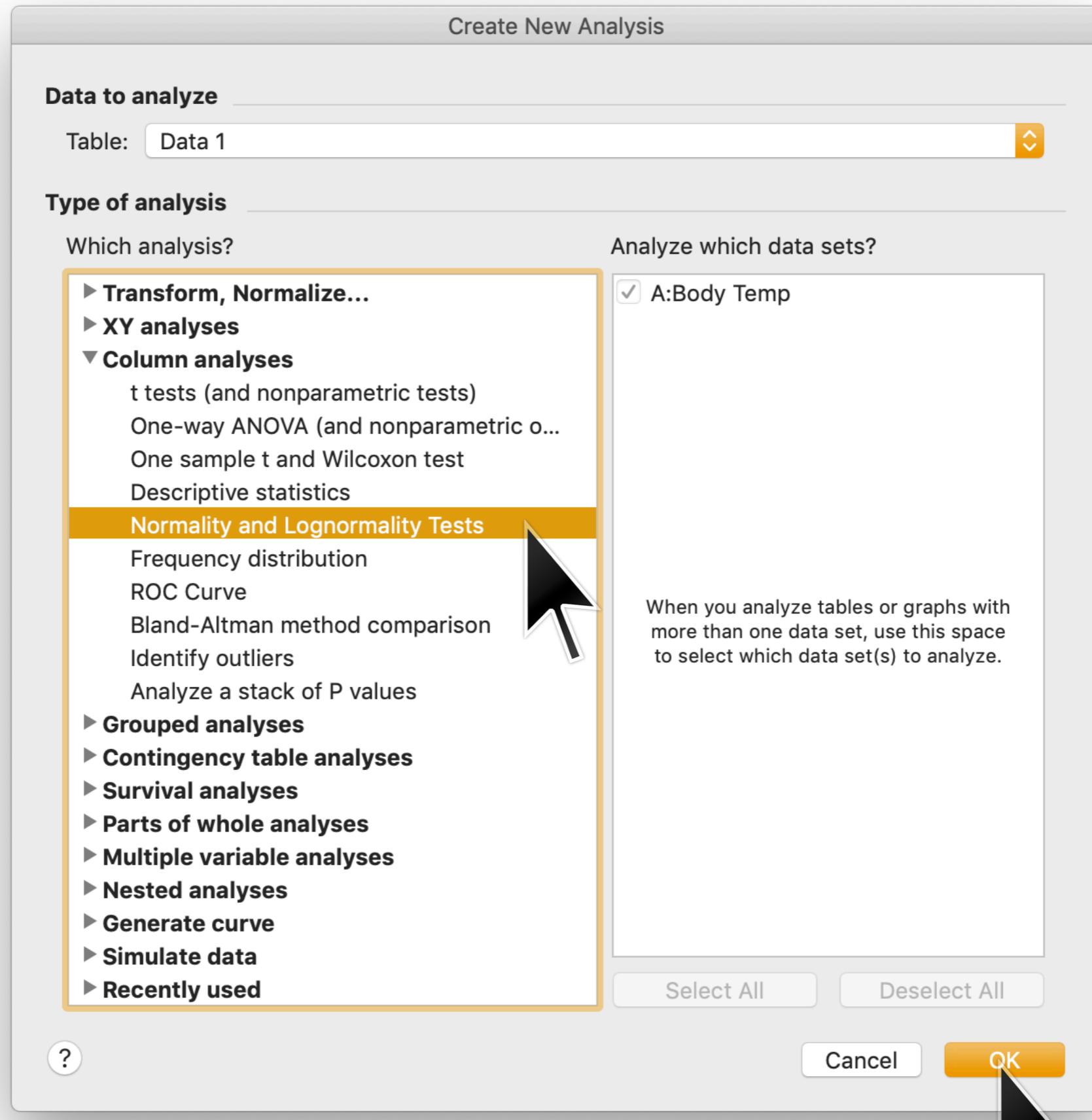
## QQ plot example: simulated lognormal data



## How to do this in Prism

	Group A	Group B	Group C	Group D
	Body Temp	Title	Title	Title
1	96.3			
2	96.7			
3	96.9			
4	97.0			
5	97.1			
6	97.1			
7	97.1			
8	97.2			
9	97.3			
10	97.4			
11	97.4			
12	97.4			
13	97.4			
14	97.5			

# How to do this in Prism



# How to do this in Prism

Parameters: Normality and Lognormality Tests

**Which distribution(s) to test?**

- Normal (Gaussian) distribution
- Lognormal distribution
- Compute the relative likelihood of sampling from a Gaussian (normal) vs. a lognormal distribution (assuming no other possibilities)

**Methods to test distribution(s)**

- Anderson-Darling test
- D'Agostino-Pearson omnibus normality test
- Shapiro-Wilk normality test
- Kolmogorov-Smirnov normality test with Dallal-Wilkinson-Lilliefors P value

**Graphing options**

- Create a QQ plot

**Subcolumns**

- Average the replicates in each row, and then perform the calculation for each column
- Perform calculations on each subcolumn separately
- Treat all the values in all subcolumns as single set of data

**Calculations**

Significance level (alpha)

**Output**

Show this many significant digits (for everything except P values):  ^ v

P value style: GP: 0.1234 (ns), 0.0332 (\*), 0.0021 (\*\*), 0...  ^ v

Make these choices the default for future analyses

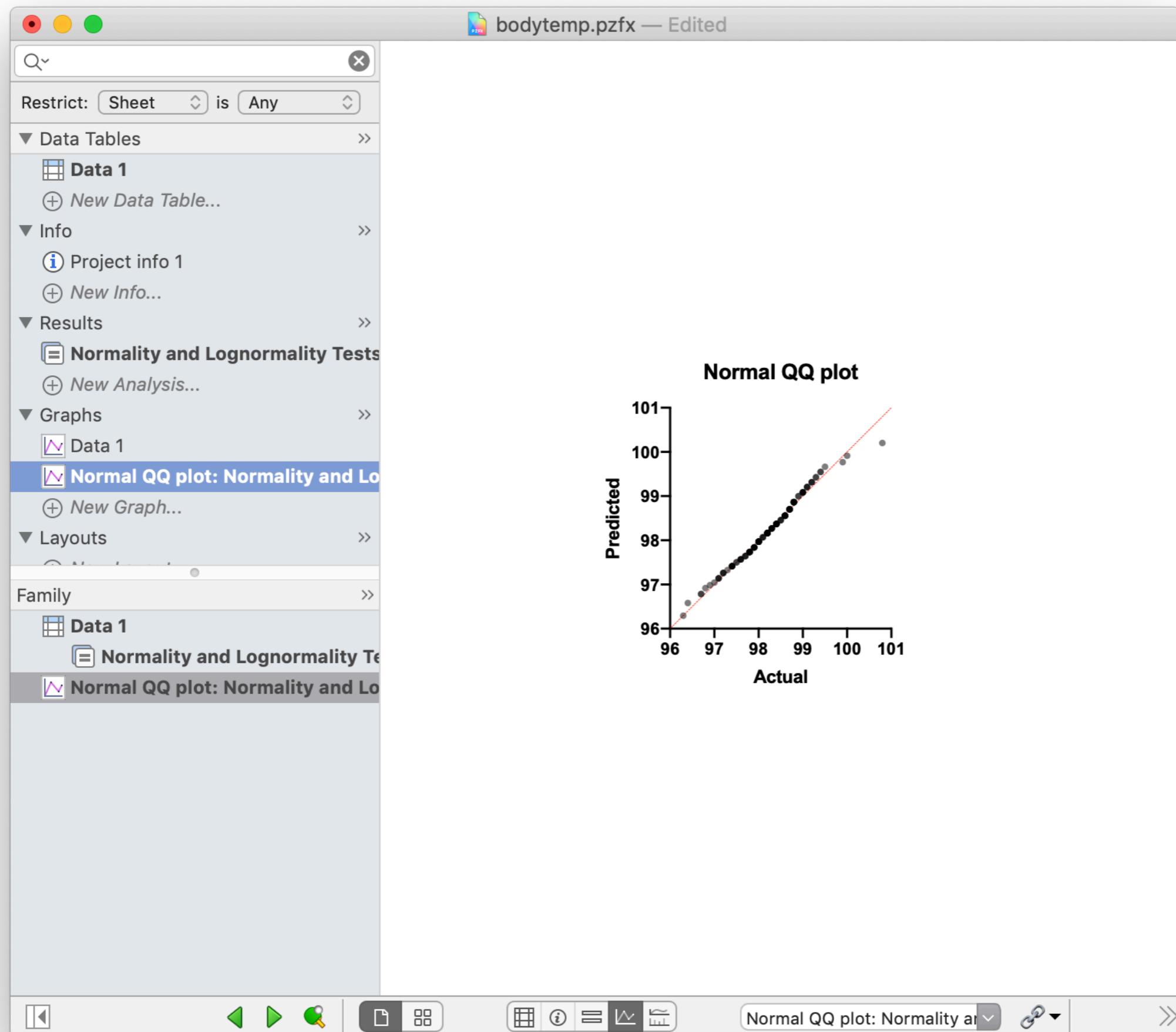
? Cancel OK

# How to do this in Prism

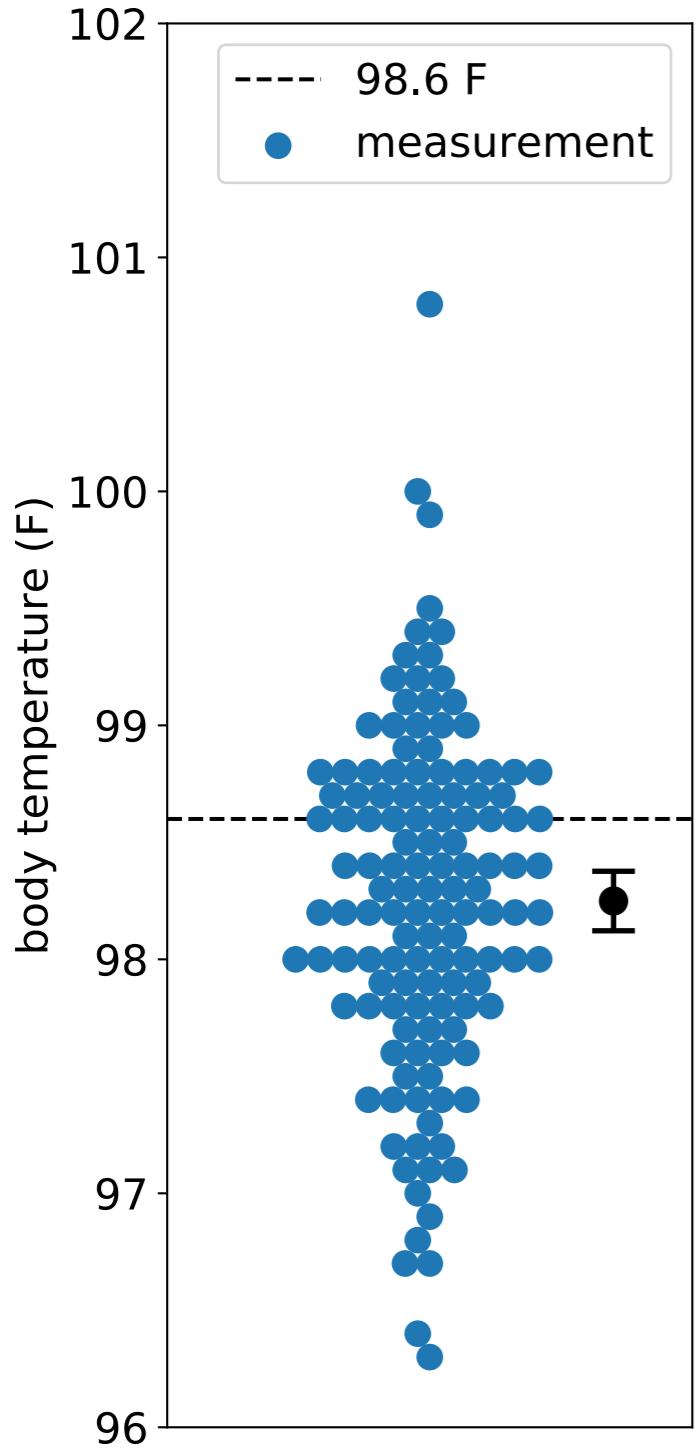
The screenshot shows the Prism software interface with the project file "bodytemp.pzfx — Edited". The left sidebar contains navigation sections like Data Tables, Info, Results, Graphs, and Layouts. The main area displays the "Normality and Lognormality Tests" tabular results. The results table has four columns labeled A, B, C, and D, with "Body Temp" assigned to column A. The table lists four tests: Anderson-Darling test, D'Agostino & Pearson test, Shapiro-Wilk test, and Kolmogorov-Smirnov test. Each test section is highlighted with a red rounded rectangle. The Anderson-Darling test section includes data for A2\*, P value, and normality test results. The D'Agostino & Pearson test section includes data for K2, P value, and normality test results. The Shapiro-Wilk test section includes data for W, P value, and normality test results. The Kolmogorov-Smirnov test section includes data for KS distance, P value, and normality test results. The final row shows the total number of values as 130.

	A	B	C	D
Body Temp	Y	Y	Y	Y
<b>Test for normal distribution</b>				
<b>Anderson-Darling test</b>				
A2*	0.5201			
P value	0.1829			
Passed normality test (alpha=0.05)?	Yes			
P value summary	ns			
<b>D'Agostino &amp; Pearson test</b>				
K2	2.704			
P value	0.2587			
Passed normality test (alpha=0.05)?	Yes			
P value summary	ns			
<b>Shapiro-Wilk test</b>				
W	0.9866			
P value	0.2332			
Passed normality test (alpha=0.05)?	Yes			
P value summary	ns			
<b>Kolmogorov-Smirnov test</b>				
KS distance	0.06473			
P value	>0.1000			
Passed normality test (alpha=0.05)?	Yes			
P value summary	ns			
<b>Number of values</b>	130			

# How to do this in PRISM



## Student's $t$ test (one sample)



### Null Hypothesis:

a population is normally distributed with  
a known mean value of  $\mu_{\text{null}}$

### Data:

measurements:  $x_1, x_2, \dots, x_N$

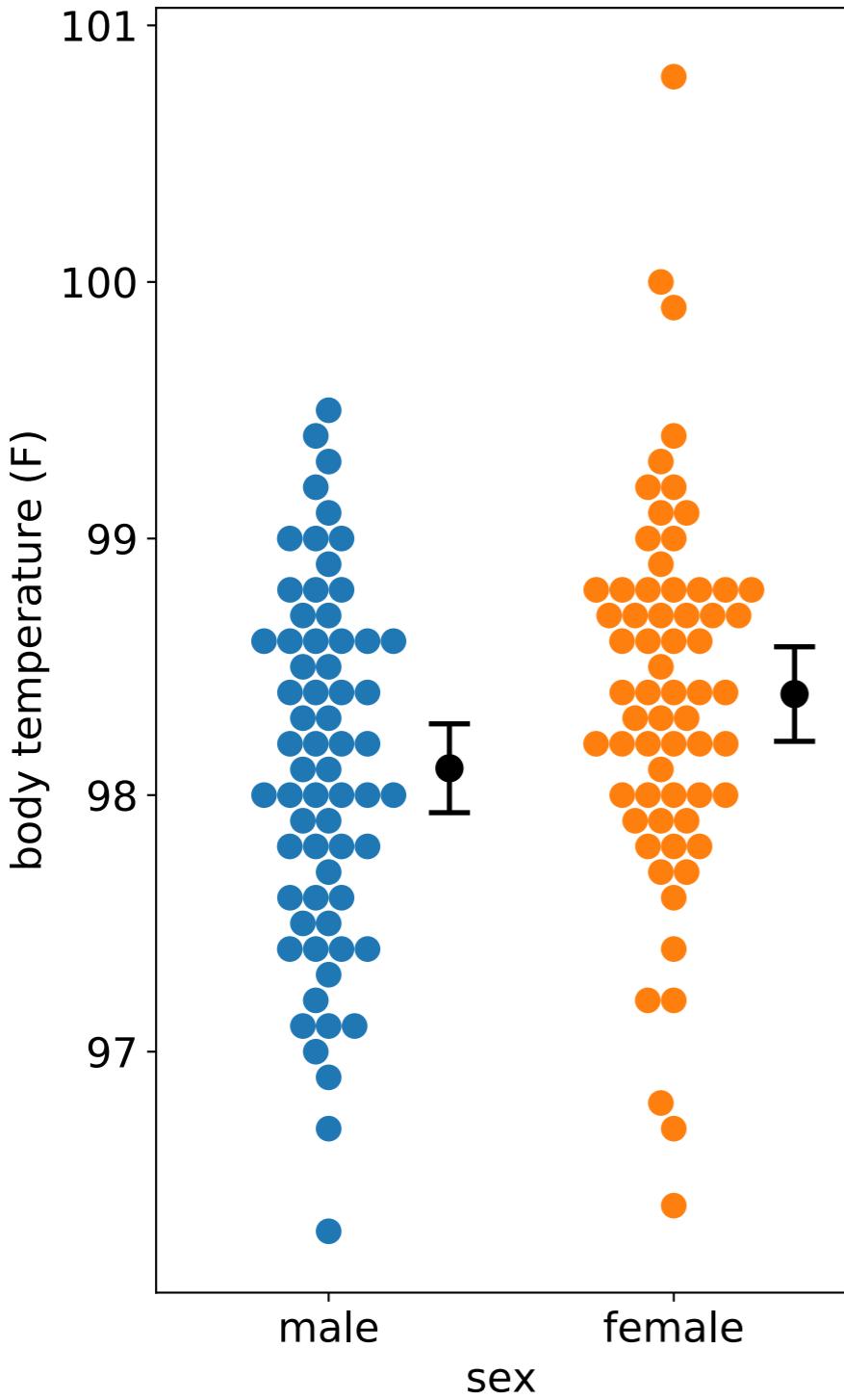
### Test statistic:

$$t = \frac{\hat{\mu} - \mu_{\text{null}}}{\text{SEM}}$$

### Null distribution:

$t$  distribution with DOF =  $N - 1$ .

## Student's $t$ test (two sample, equal variance)



**Null Hypothesis:**

two populations have the same mean

**Data:**

$x_1, x_2, \dots, x_m$  and  $y_1, y_2, \dots, y_n$

**Assumptions:**

the two populations follow normal distributions and have equal variances

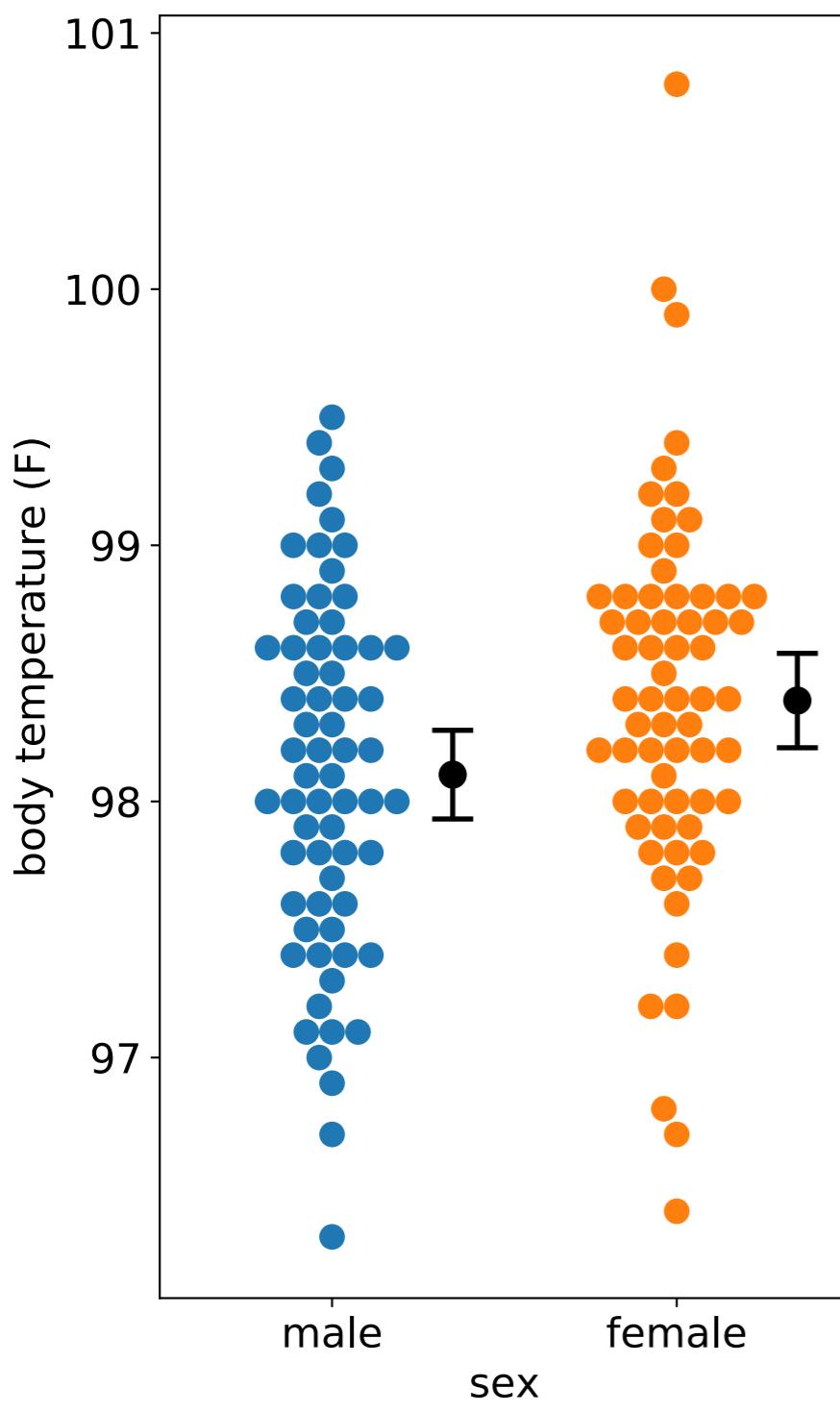
**Test statistic:**

$$t = \frac{\hat{\mu}_x - \hat{\mu}_y}{\hat{\sigma} \sqrt{\frac{1}{m} + \frac{1}{n}}}, \quad \hat{\sigma} = \sqrt{\frac{(m-1)\hat{\sigma}_x^2 + (n-1)\hat{\sigma}_y^2}{m+n-2}}$$

**Null distribution:**

$t$  distribution with DOF =  $m + n - 2$ .

## Welch's $t$ test



### Null Hypothesis:

two populations have the same mean but  
not necessarily the same standard deviation

### Data:

$x_1, x_2, \dots, x_m$  and  $y_1, y_2, \dots, y_n$

### Advantage:

Fewer assumptions than standard unpaired  $t$  test

### Disadvantage:

Less power than standard unpaired  $t$  tests

### Test statistic:

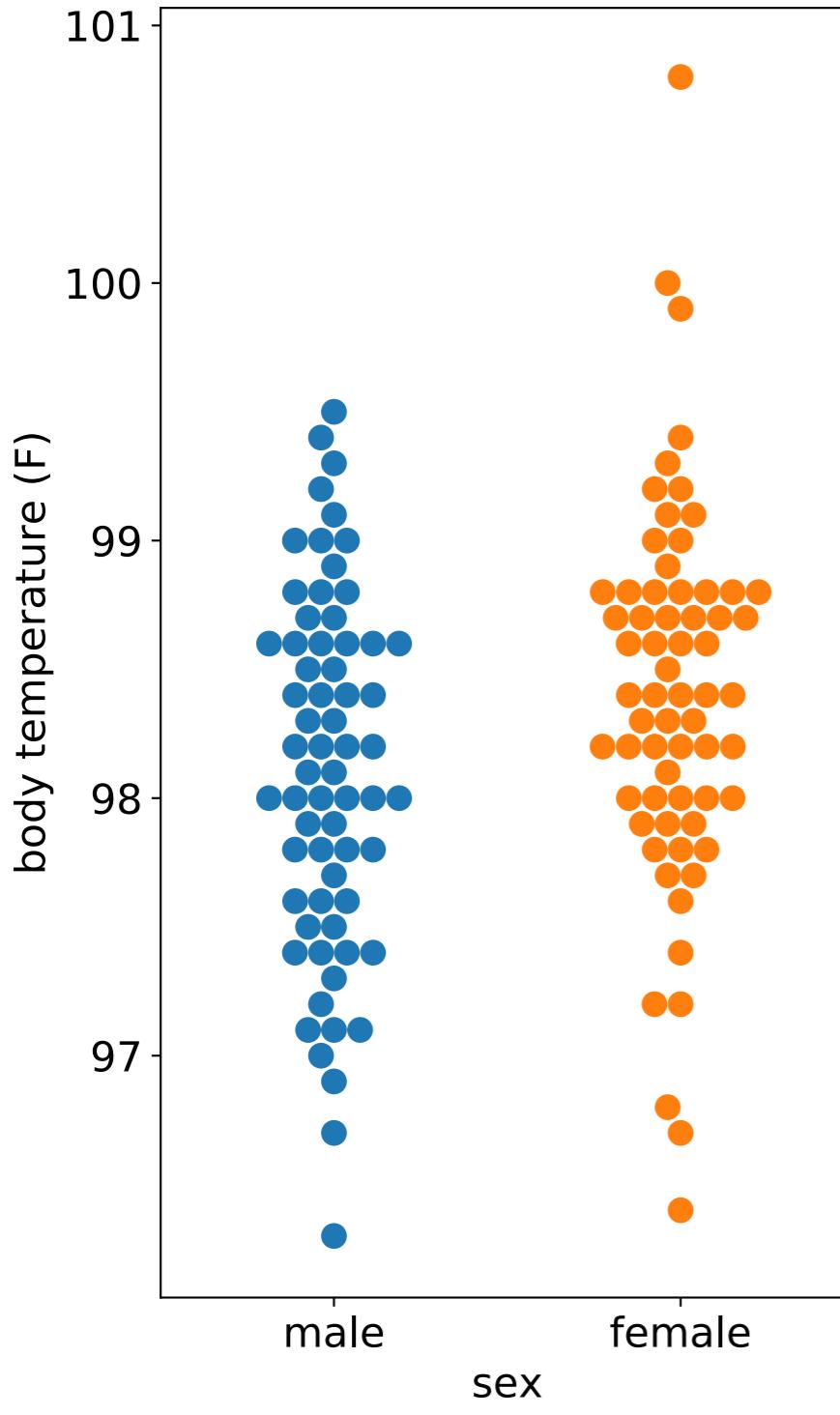
$$t = \frac{\hat{\mu}_x - \hat{\mu}_y}{\sqrt{\frac{\hat{\sigma}_x^2}{m} + \frac{\hat{\sigma}_y^2}{n}}}$$

### Null distribution:

Student's  $t$  distribution with

$$\text{DOF} = \frac{\left( \frac{\hat{\sigma}_x^2}{m} + \frac{\hat{\sigma}_y^2}{n} \right)^2}{\frac{(\hat{\sigma}_x^2/m)^2}{m-1} + \frac{(\hat{\sigma}_y^2/n)^2}{n-1}}$$

## Mann Whitney U test (Wilcoxon rank-sum test)



### Null Hypothesis:

If  $x$  is sampled from population 1 and  $y$  is sampled from population 2,  
 $p(x > y) = p(x < y)$

### Data:

$x_1, x_2, \dots, x_m$  and  $y_1, y_2, \dots, y_n$

### Advantage:

No assumptions about the mathematical form of  $p(x)$  and  $p(y)$ .

### Disadvantage:

Somewhat less powerful than Student's  $t$  test

### Test statistic:

$U$  (based on rank-order of  $xs$  and  $ys$ )

temp\_by\_sex.pzfx

	Group A male	Group B female	Group C Title	Group D Title
1	96.3	96.4		
2	96.7	96.7		
3	96.9	96.8		
4	97.0	97.2		
5	97.1	97.2		
6	97.1	97.4		
7	97.1	97.6		
8	97.2	97.7		
9	97.3	97.7		
10	97.4	97.8		
11	97.4	97.8		
12	97.4	97.8		
13	97.4	97.9		
14	97.5	97.9		

## Create New Analysis

### Data to analyze

Table: Data 1

### Type of analysis

Which analysis?

- ▶ Transform, Normalize...
- ▶ XY analyses
- ▼ Column analyses
  - t tests (and nonparametric tests)**
  - One-way ANOVA (and nonparametric o...)
  - One sample t and Wilcoxon test
  - Descriptive statistics
  - Normality and Lognormality Tests
  - Frequency distribution
  - ROC Curve
  - Bland-Altman method comparison
  - Identify outliers
  - Analyze a stack of P values
- ▶ Grouped analyses
- ▶ Contingency table analyses
- ▶ Survival analyses
- ▶ Parts of whole analyses
- ▶ Multiple variable analyses
- ▶ Nested analyses
- ▶ Generate curve
- ▶ Simulate data
- ▶ Recently used

Analyze which data sets?

- A:male
- B:female

Select All

Deselect All

?

Cancel

OK

## Parameters: t Tests (and Nonparametric Tests)

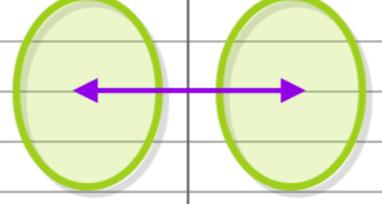
Experimental Design

Residuals

Options

**Experimental design** Unpaired Paired

		Group A	Group B
		Control	Treated
1	Y	Y	
2			
3			
4			
5			

**Assume Gaussian distribution?**

- Yes. Use parametric test.
- No. Use nonparametric test.

**Choose test**

- Unpaired t test. Assume both populations have the same SD
- Unpaired t test with Welch's correction. Do not assume equal SDs

?

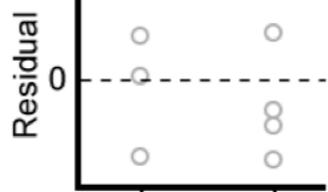
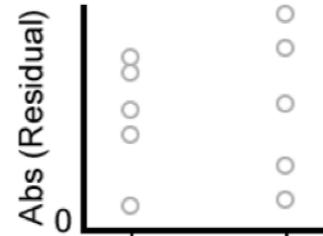
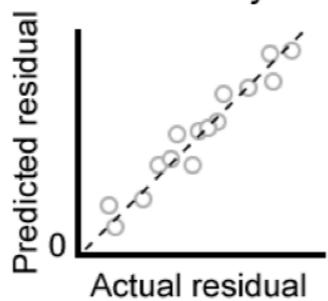
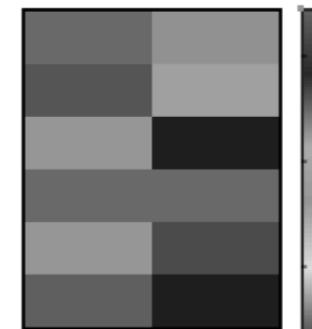
Cancel

OK

Experimental Design

Residuals

Options

**What graphs to create?***Correct model?* Residual plot*Equal variance?* Homoscedasticity plot*Normality?* QQ plot Heatmap plot**Diagnostics for residuals** Are the residuals Gaussian?

Normality tests of Anderson-Darling, D'Agostino, Shapiro-Wilk and Kolmogorov-Smirnov.

 Make options on this tab be the default for future tests.

?

Cancel

OK

## Parameters: t Tests (and Nonparametric Tests)

Experimental Design   Residuals   Options

### Calculations

P value:  One-tailed  Two-tailed (recommended)

Report differences as: female - male

Confidence level: 95%

Definition of statistical significance: P < 0.05

### Graphing options

- Graph differences (paired)
- Graph ranks (nonparametric)
- Graph correlation (paired)
- Graph CI of difference between means

### Additional results

- Descriptive statistics for each dataset
- t Test: Also compare models using AICc
- Mann-Whitney: Also compute the CI of difference between medians

Assumes both distributions have the same shape.
- Wilcoxon: When both values on a row are identical, use method of Pratt

If this option is unchecked, those rows are ignored and the results will match prior version of Prism

### Output

Show this many significant digits (for everything except P values): 4

P value style: GP: 0.1234 (ns), 0.0332 (\*), 0.0021 (\*\*), 0.0002 (\*\*\*), <0.000... N= 6

Make options on this tab be the default for future tests.



Cancel

OK

temp\_by\_sex.pzfx — Edited

Search

Data Tables >

Data 1  
New Data Table...

Info >

Project info 1  
New Info...

Results >

Unpaired t test of Data 1  
New Analysis...

Graphs >

Data 1  
QQ plot: Unpaired t test of Data 1  
Mean diff. CI plot: Unpaired t test  
New Graph...

Layouts >

New Layout...

Family >

Data 1  
Unpaired t test  
QQ plot: Unpaired t test of Data 1  
Mean diff. CI plot: Unpaired t test

Tabular results

Unpaired t test  
Tabular results

	Table Analyzed	
1	Data 1	
2		
3	Column B	female
4	vs.	vs.
5	Column A	male
6		
7	<b>Unpaired t test</b>	
8	P value	0.0239
9	P value summary	*
10	Significantly different ( $P < 0.05$ )?	Yes
11	One- or two-tailed P value?	Two-tailed
12	t, df	t=2.285, df=128
13		
14	<b>How big is the difference?</b>	
15	Mean of column A	98.10
16	Mean of column B	98.39
17	Difference between means (B - A) :	$0.2892 \pm 0.1266$
18	95% confidence interval	0.03882 to 0.5396
19	R squared (eta squared)	0.03921
20		
21	<b>F test to compare variances</b>	
22	F, DFn, Dfd	1.132, 64, 64
23	P value	0.6211

Unpaired t test of Data 1

Row 1, Column A

temp\_by\_sex.pzfx — Edited

**Tabular results**

**Unpaired t test**  
Tabular results

20					
21	<b>F test to compare variances</b>				
22	F, DFn, Dfd	1.132, 64, 64			
23	P value	0.6211			
24	P value summary	ns			
25	Significantly different ( $P < 0.05$ )?	No			
26					
27	<b>Normality of Residuals</b>				
28	<b>Test name</b>	<b>Statistics</b>	<b>P value</b>	<b>Passed normality test (alpha=0.05)?</b>	<b>P value summary</b>
29	Anderson-Darling (A2*)	0.3633	0.4359	Yes	ns
30	D'Agostino-Pearson omnibus (K2)	2.467	0.2913	Yes	ns
31	Shapiro-Wilk (W)	0.9906	0.5264	Yes	ns
32	Kolmogorov-Smirnov (distance)	0.05178	0.1000	Yes	ns
33					
34	<b>Data analyzed</b>				
35	Sample size, column A	65			
36	Sample size, column B	65			
37					
38					

Unpaired t test of Data 1

Row 1, Column A

