

# **Malaria Molecular Surveillance Study Design Workshop**

## **Module 4: Statistical power**

## Remember ...

		Conclusion about $H_0$	
		Fail to reject	Reject
Truth about $H_0$	True	True negative $1 - \alpha$	False positive $\alpha$
	False		

$\alpha$  sets the **false positive rate** of a test. Using  $\alpha$  we can control how often we incorrectly conclude that there is a real effect when there is none.

In power analysis, we also specify an **alternative hypothesis**

**$H_0$ : The population prevalence equals  $p_0$**

**$H_1$ : The population prevalence equals  $p$ , which is different from  $p_0$**

In power analysis, we also specify an **alternative hypothesis**

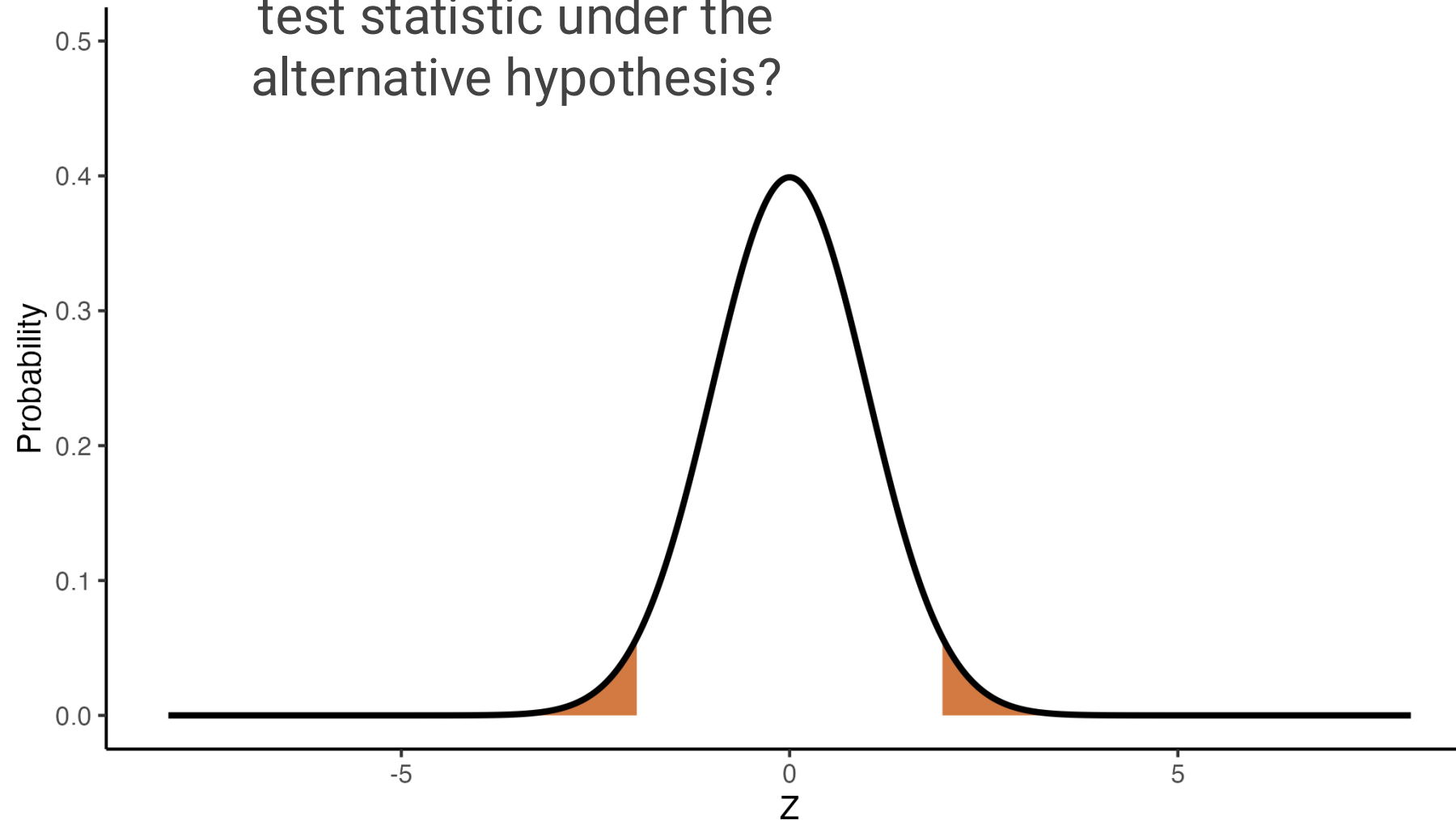
**$H_0$ : The population prevalence equals  $p_0$**

**$H_1$ : The population prevalence equals  $p$ , which is different from  $p_0$**

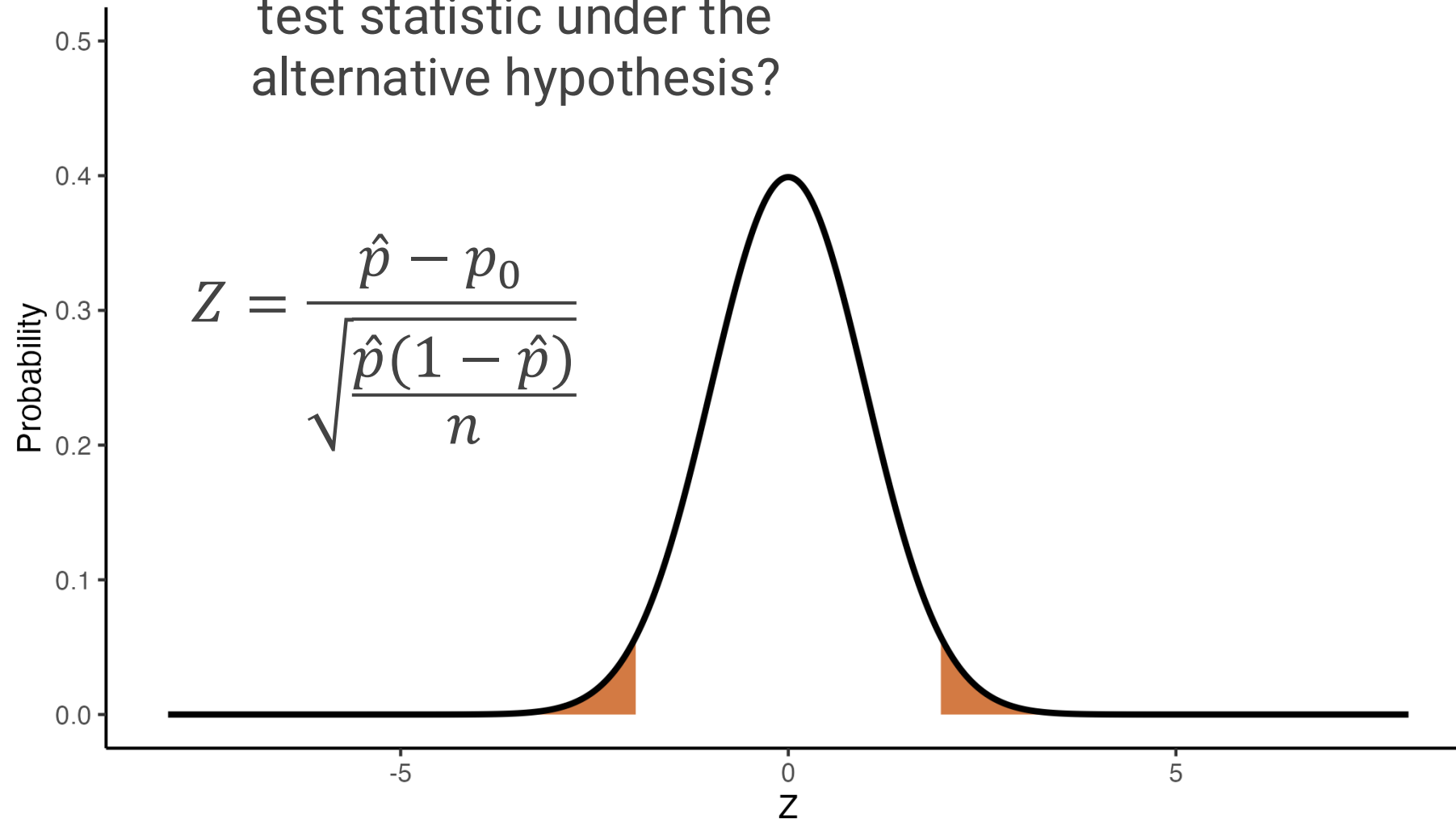
**For example...**

I want to test if the prevalence of *pfprt* K76T mutations is significantly different from 10%. When powering this test, I assume the true prevalence of K76T mutations is 15%.

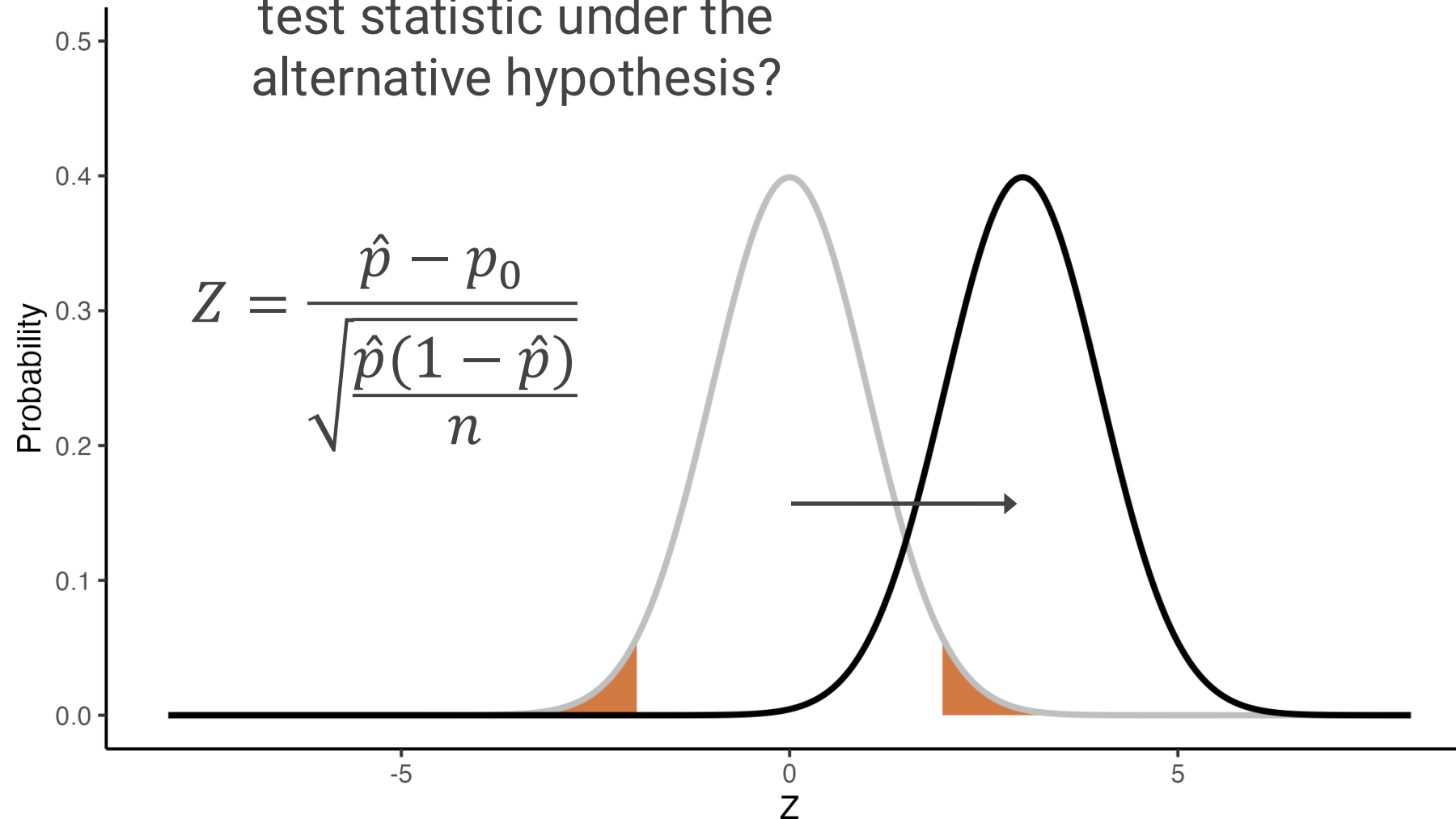
What is the distribution of my  
test statistic under the  
alternative hypothesis?



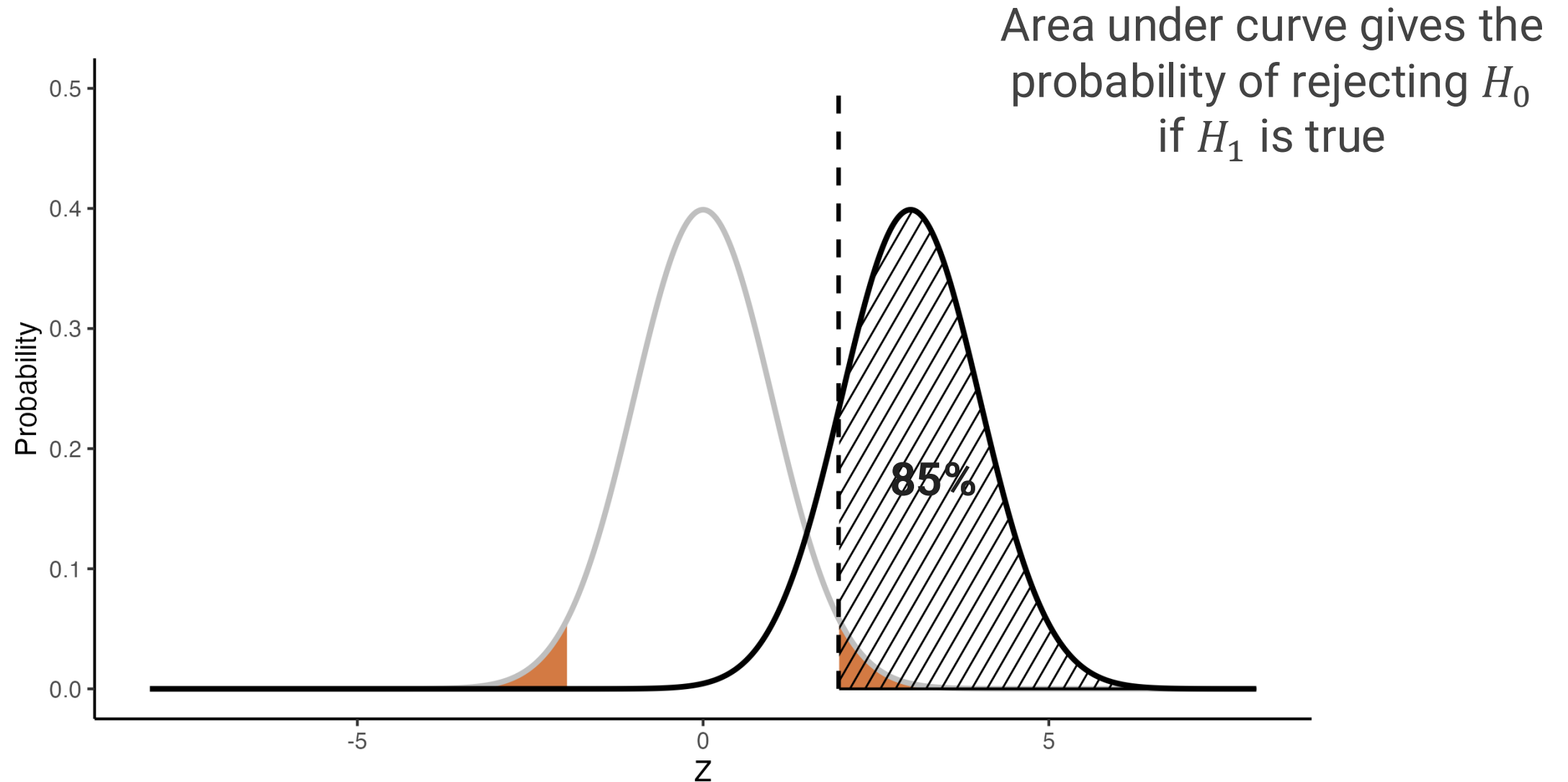
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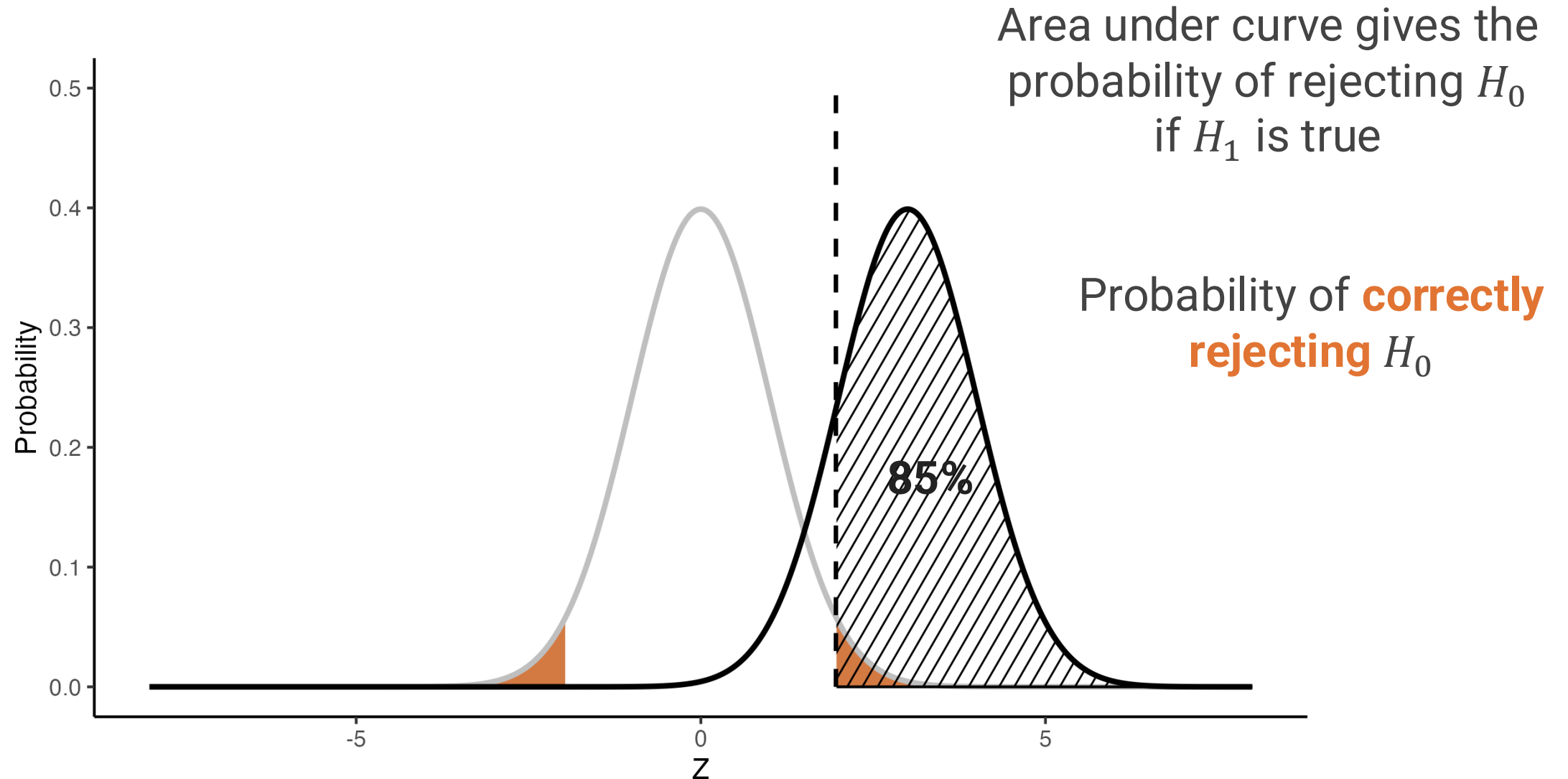
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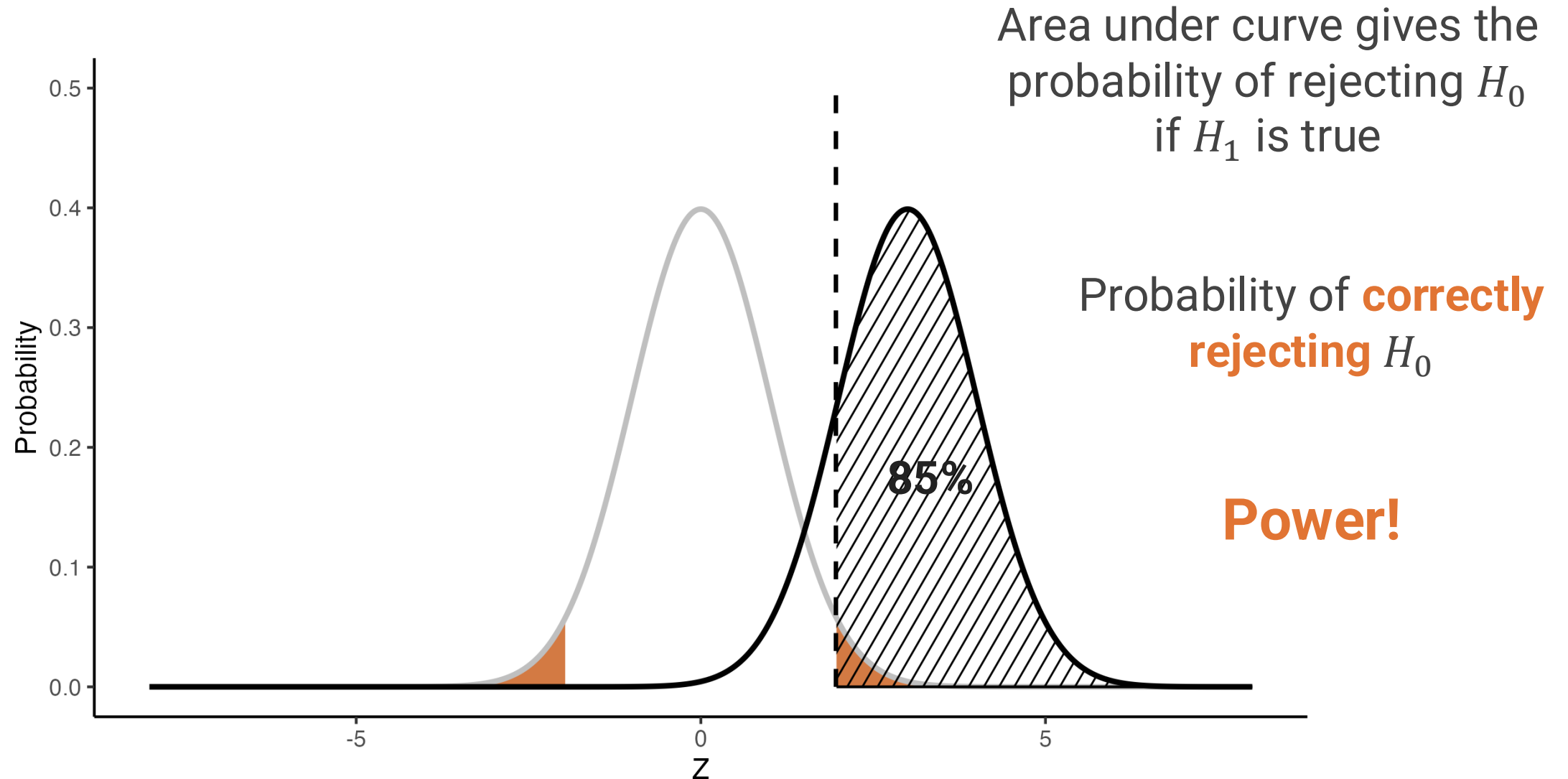
# Power







# Power



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		Fail to reject	Reject
Truth about $H_0$	True	True negative $1 - \alpha$	False positive $\alpha$
	False	False negative	True positive

		Conclusion about $H_0$	
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Truth about $H_0$	True	True negative $1 - \alpha$	False positive $\alpha$
	False	False negative $1 - \text{Power}$	True positive Power

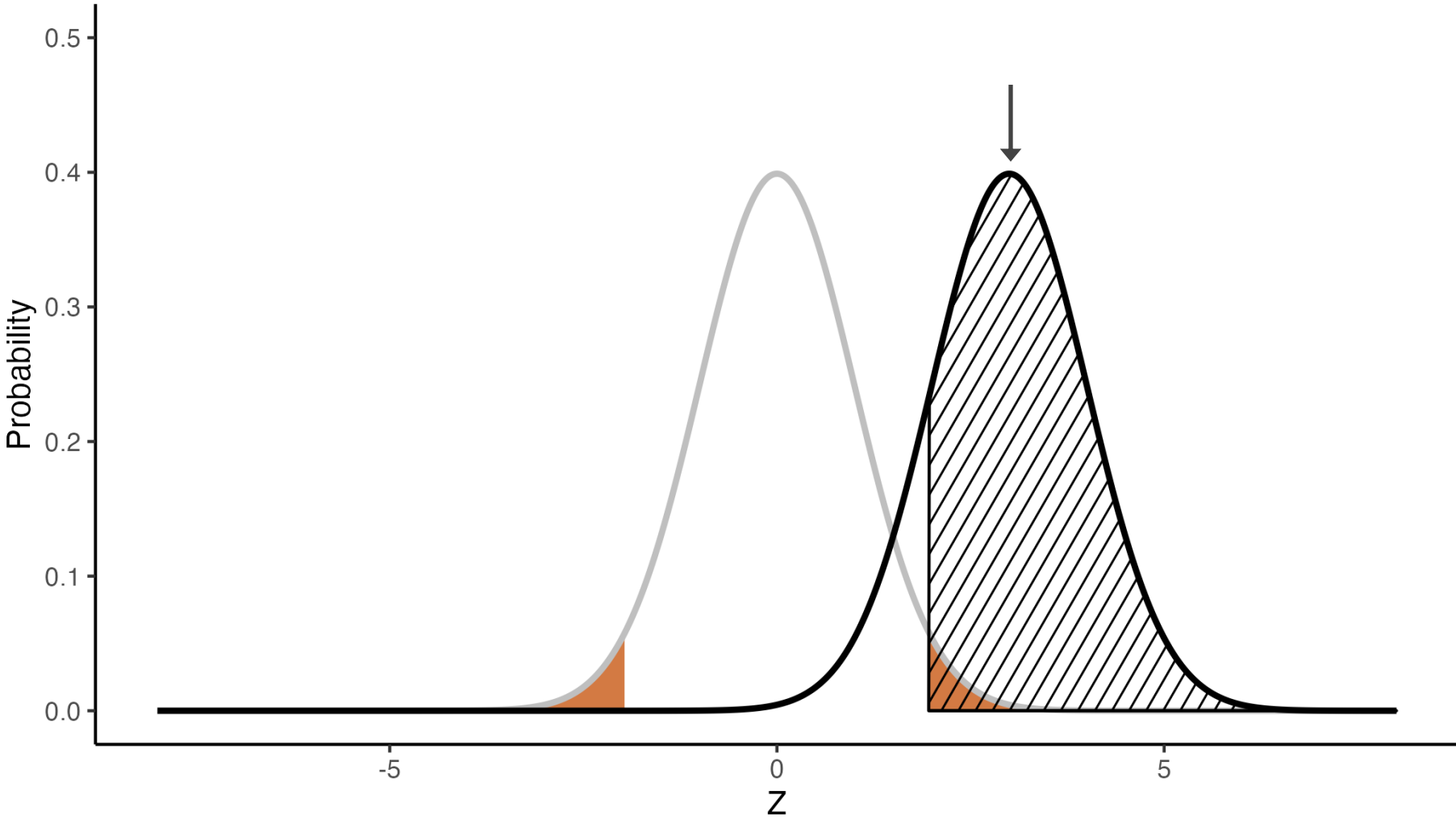
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Truth about $H_0$	True	True negative $1 - \alpha$	False positive $\alpha$
	False	False negative $1 - \text{Power}$	True positive Power

Power is the probability of **correctly rejecting** the null hypothesis. It is the chance that we find something interesting, given that it is there.

		Conclusion about $H_0$	
		Fail to reject	Reject
Truth about $H_0$	True	True negative $1 - \alpha$	False positive $\alpha$
	False	False negative $\beta$	True positive $1 - \beta$

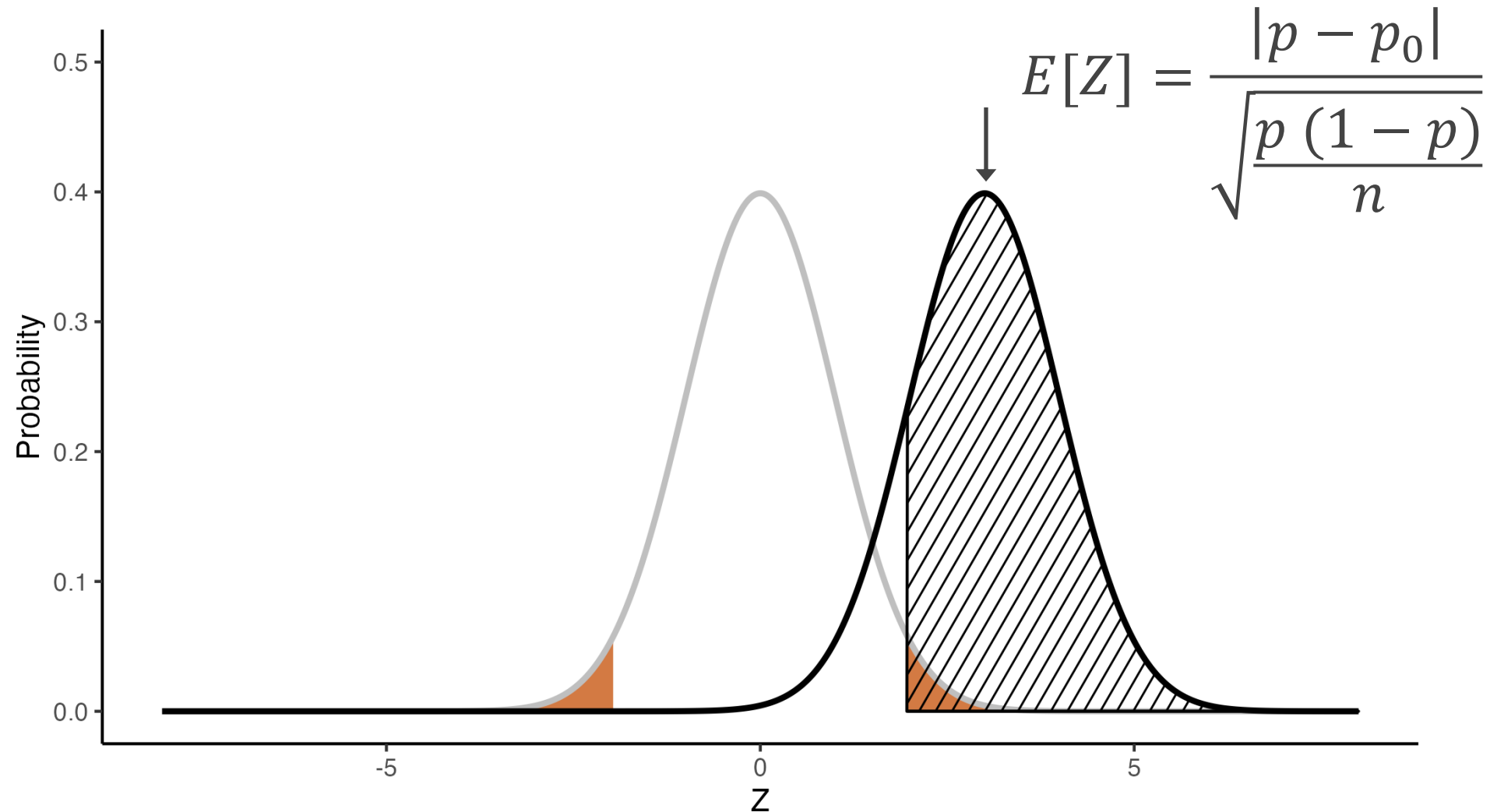
Power is the probability of **correctly rejecting** the null hypothesis. It is the chance that we find something interesting, given that it is there.

# How do we calculate power?

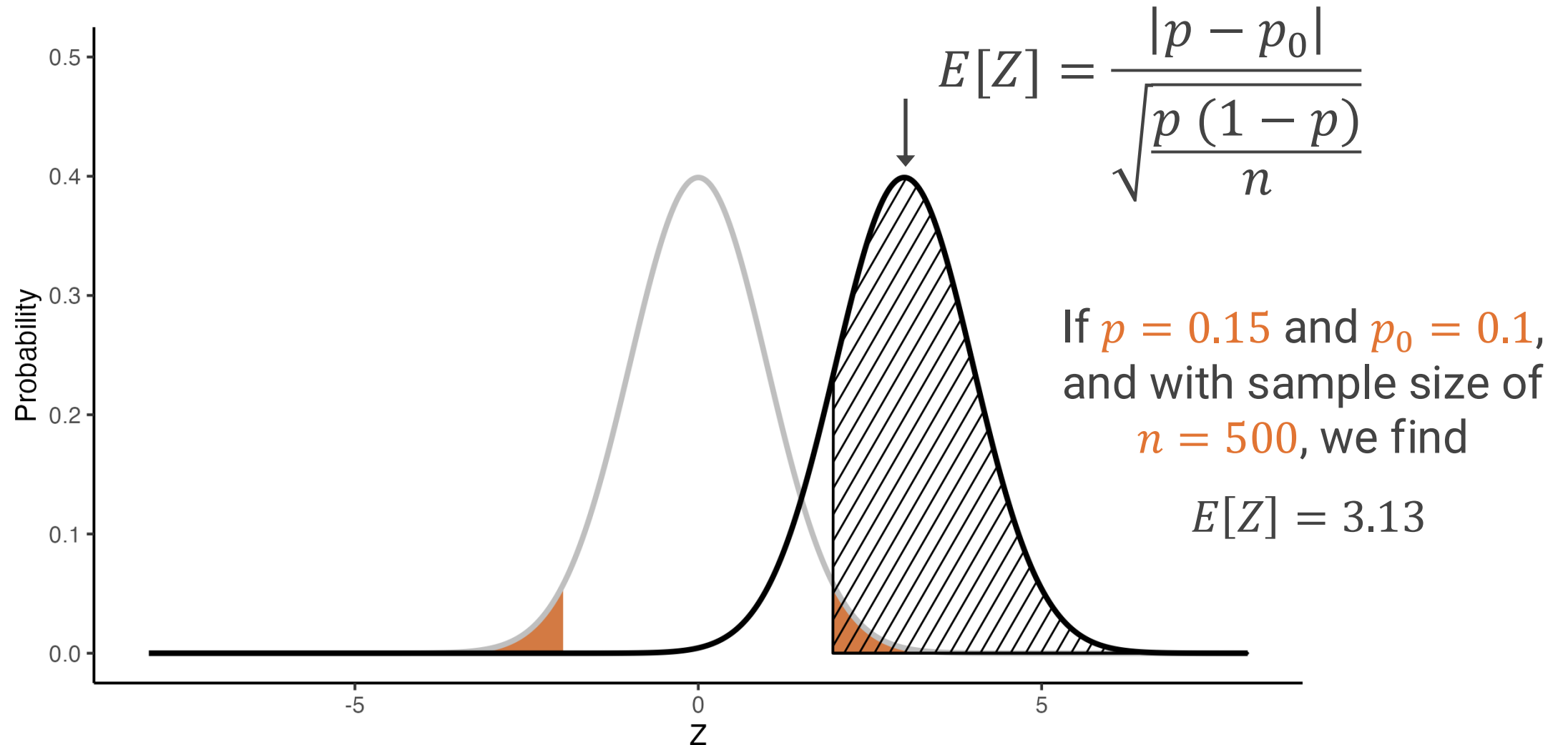




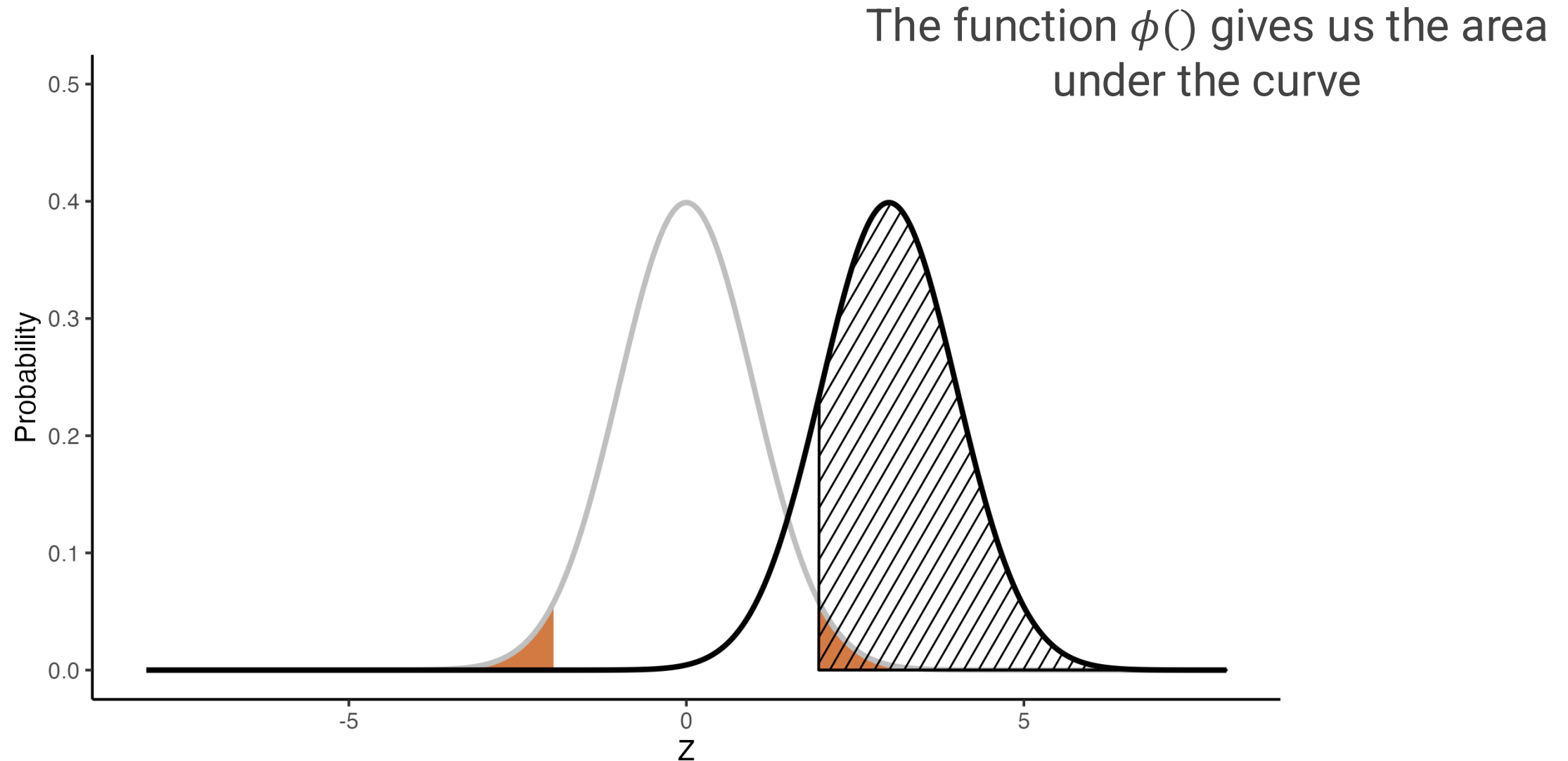
# How do we calculate power?



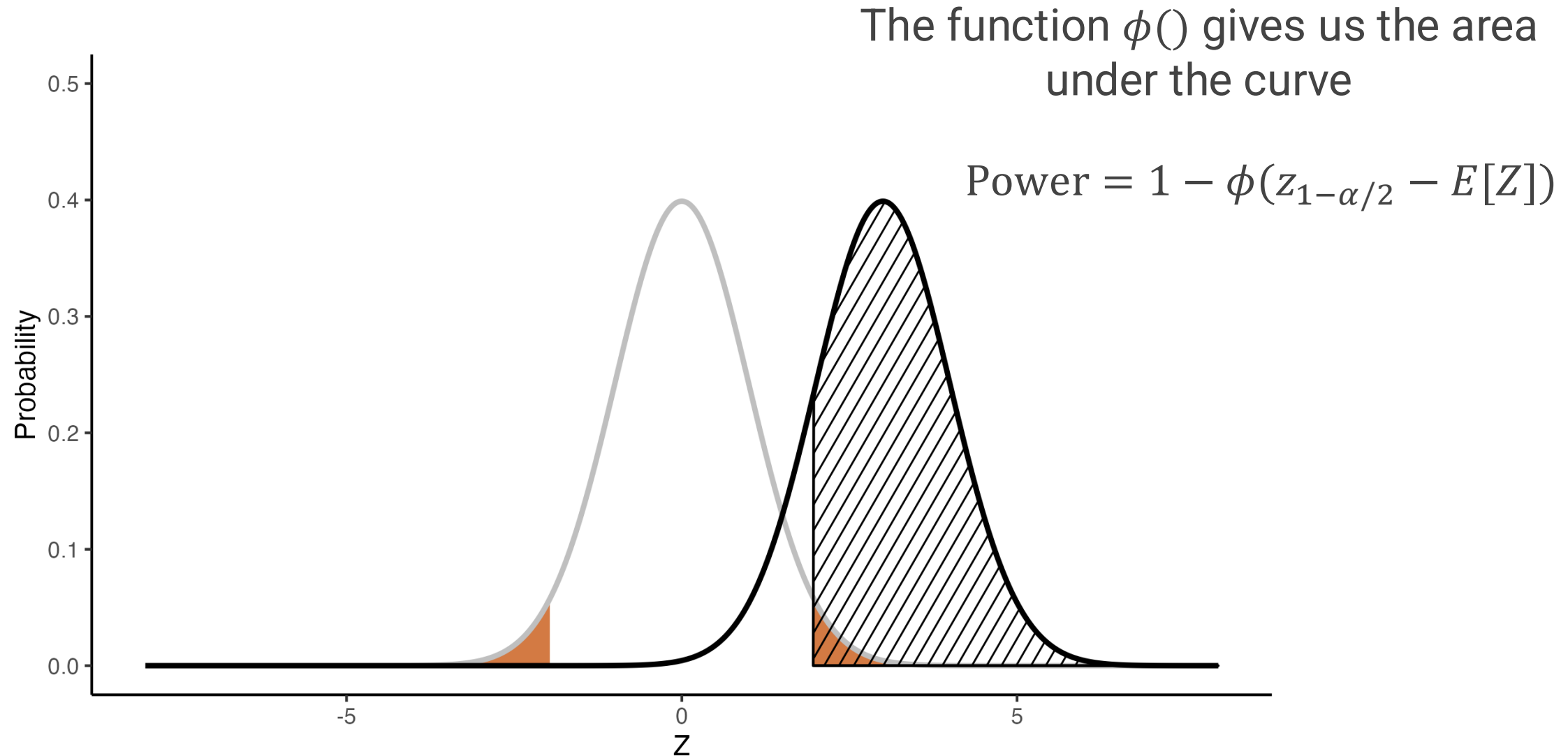
# How do we calculate power?



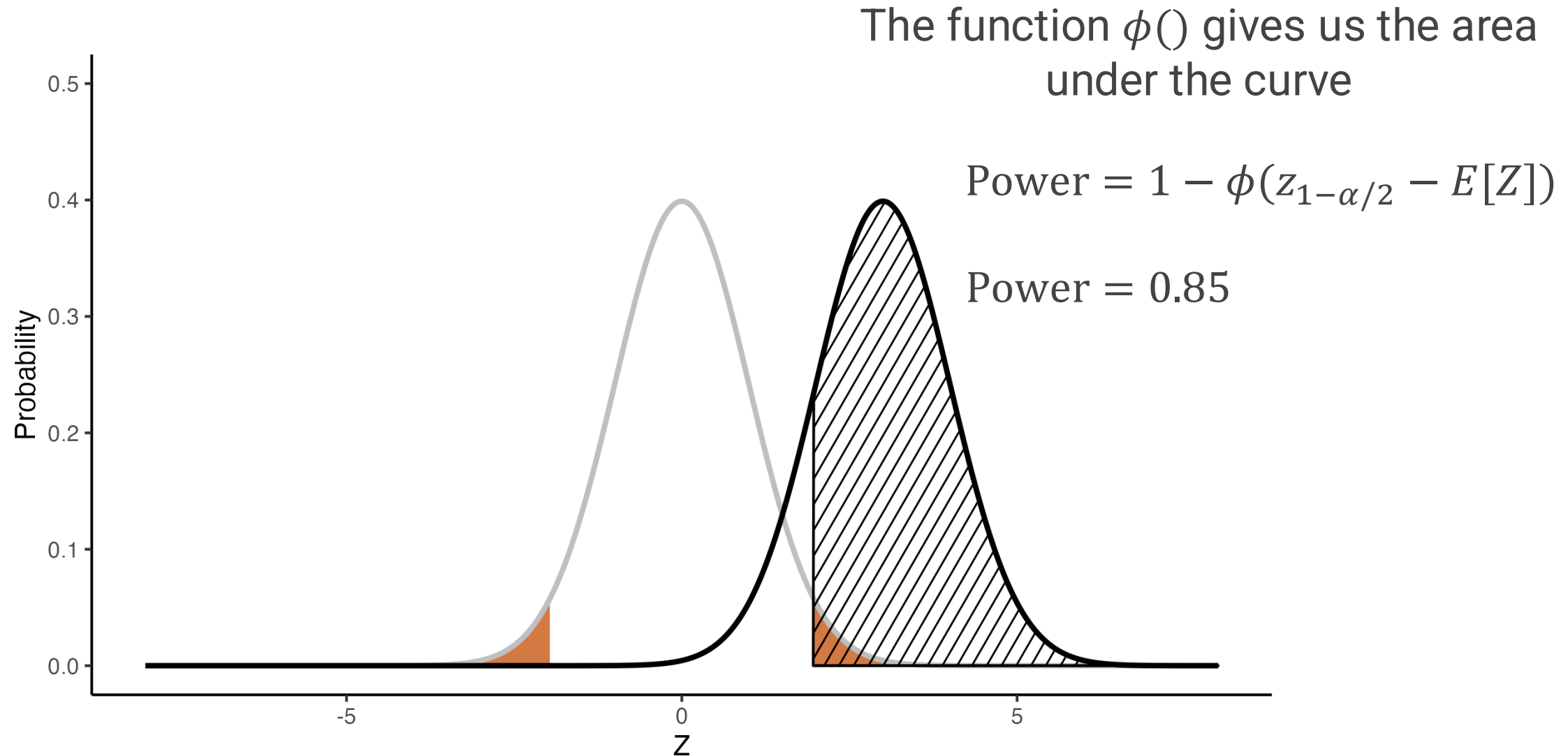
# How do we calculate power?



# How do we calculate power?



# How do we calculate power?



# Power as a function of sample size

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$$\text{Power} = 1 - \phi(z_{1-\alpha/2} - E[Z])$$

# Power as a function of sample size

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$$\text{Power} = 1 - \phi \left( z_{1-\alpha/2} - \frac{|p - p_0|}{\sqrt{\frac{p(1-p)}{n}}} \right)$$

# Power as a function of sample size

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$$\text{Power} = 1 - \phi \left( z_{1-\alpha/2} - \frac{|p - p_0|}{\sqrt{\frac{p(1-p)}{n}}} \right)$$

Power varies as a function  
of sample size

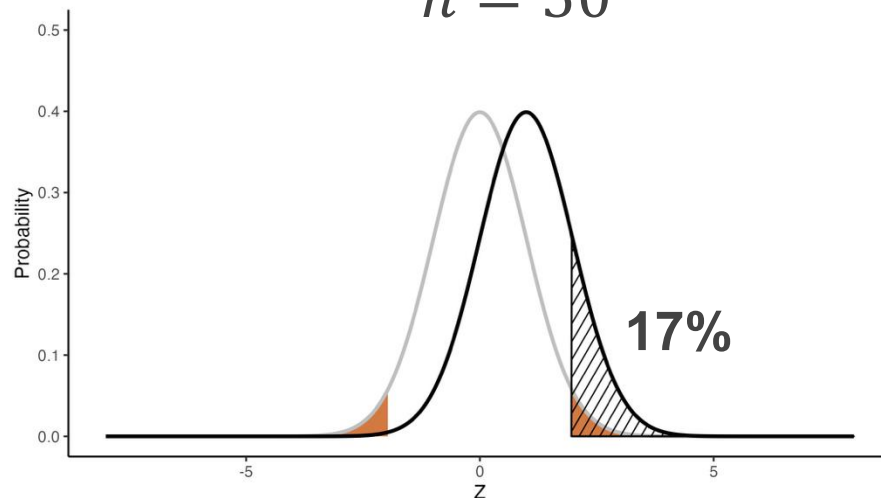


# Power as a function of sample size

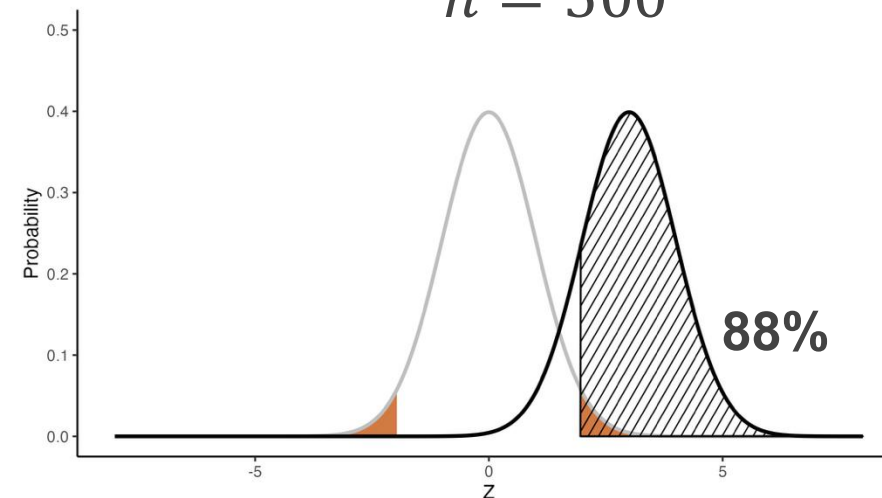
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Power varies as a function of sample size

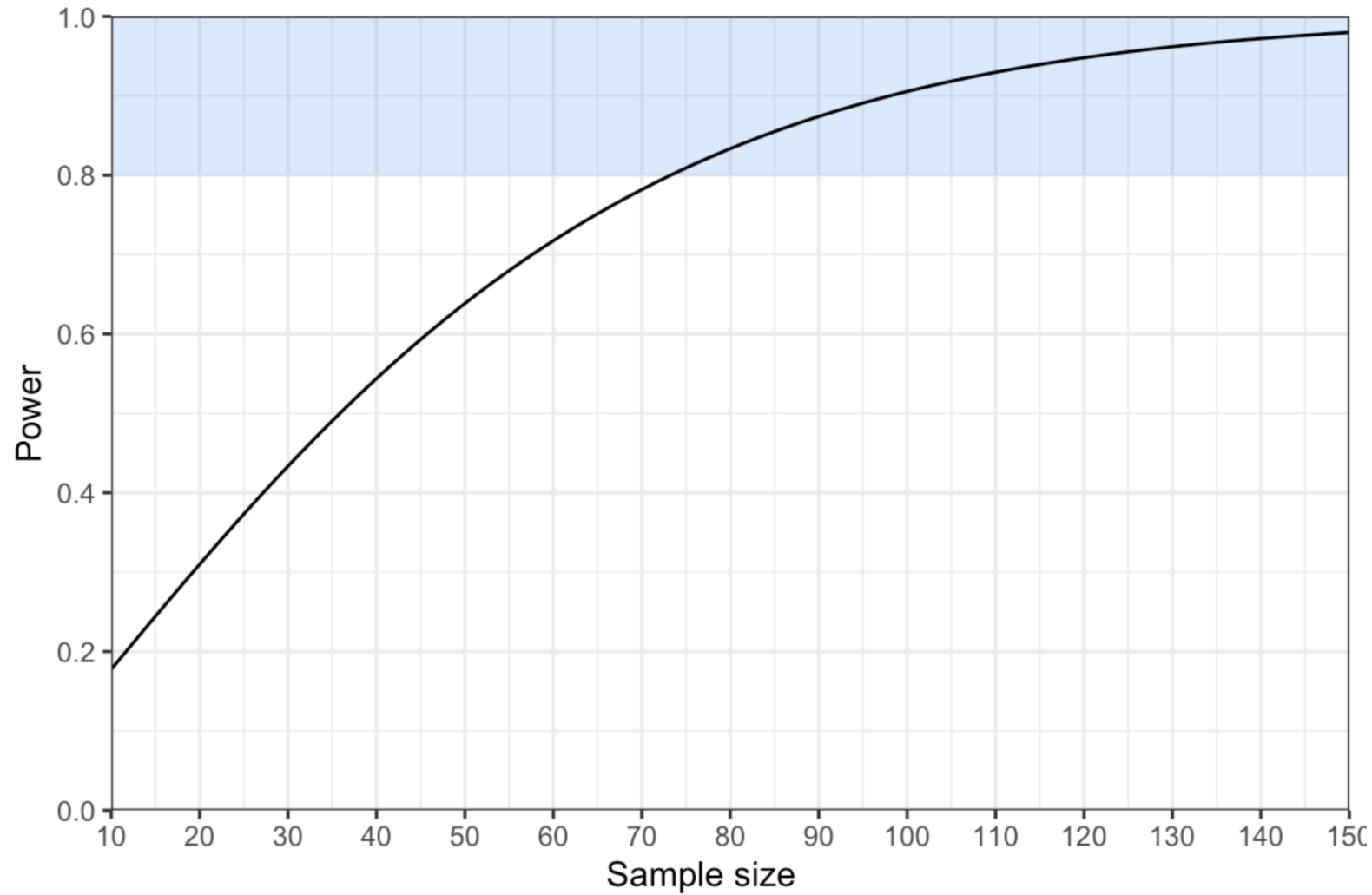
Small sample  
 $n = 50$



Large sample  
 $n = 500$

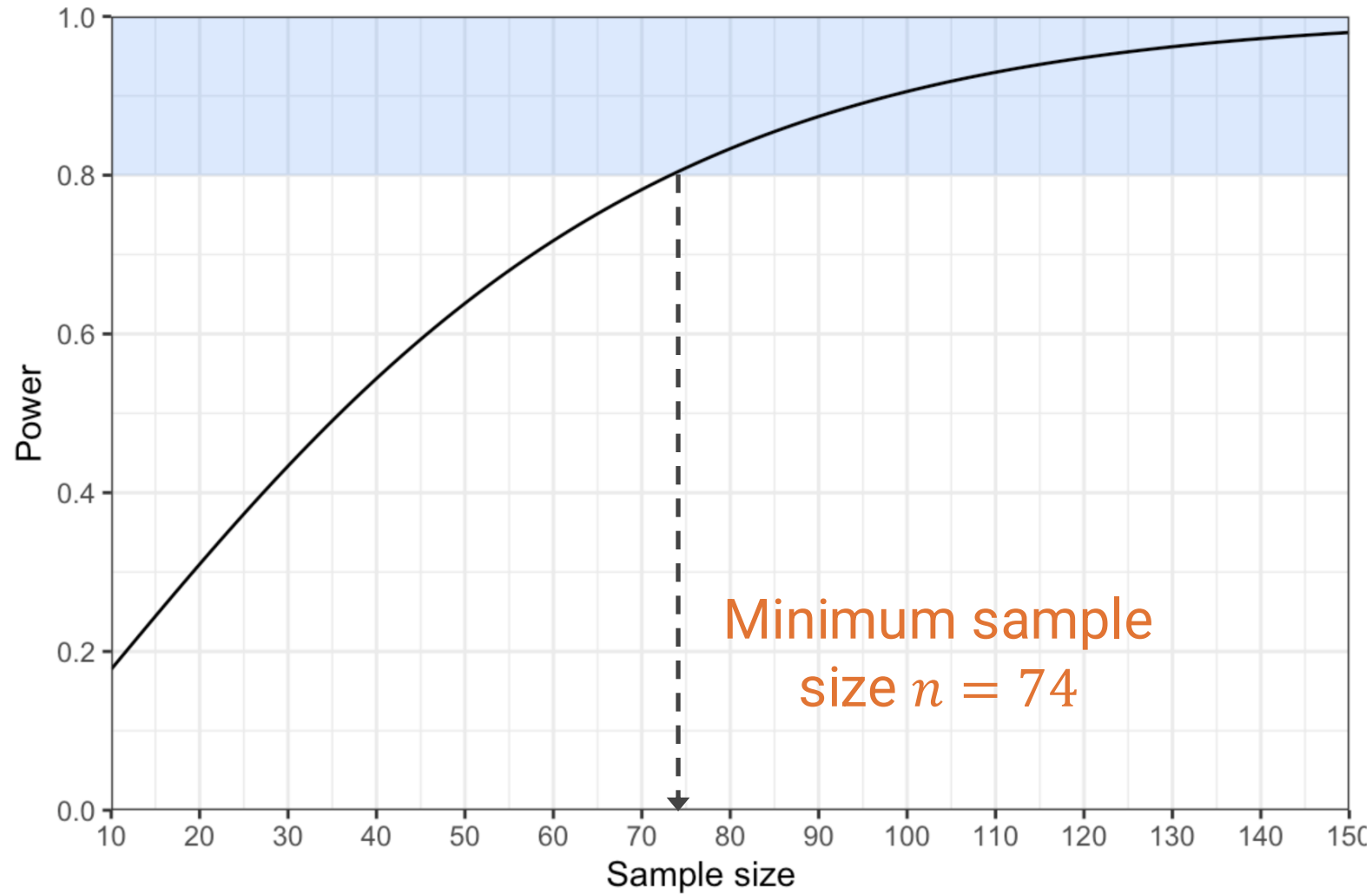


# Power curves



80% power

# Power curves



80% power

Minimum sample  
size  $n = 74$

$$\text{Power} = 1 - \phi \left( z_{1-\alpha/2} - \frac{|p - p_0|}{\sqrt{\frac{p(1-p)}{n}}} \right)$$

Can we reverse-engineer this to find the value of  $n$  that achieves a target power?

$$\text{Power} = 1 - \phi \left( z_{1-\alpha/2} - \frac{|p - p_0|}{\sqrt{\frac{p(1-p)}{n}}} \right)$$

Can we reverse-engineer this to find the value of  $n$  that achieves a target power?

$$n = \left( z_{1-\beta} + z_{1-\frac{\alpha}{2}} \right)^2 \frac{p(1-p)}{(p - p_0)^2}$$

For 80% power, we find  $z_{1-\beta} = 0.84$

- **Power** is the true positive rate. It is the chance of **correctly rejecting the null hypothesis**.
- Power increases with **sample size**. We can use power curves or sample size formulae to choose a value of  $n$

**Format:** Interactive R code, accessed through the web

- Test for change in prevalence
- Using power curves and sample size tables
- Test for detection of rare *pfk13* variant



Workshop materials

[https://mrc-ide.github.io/MMS-SD\\_workshop/](https://mrc-ide.github.io/MMS-SD_workshop/)