

# Malaria Molecular Surveillance Study Design Workshop

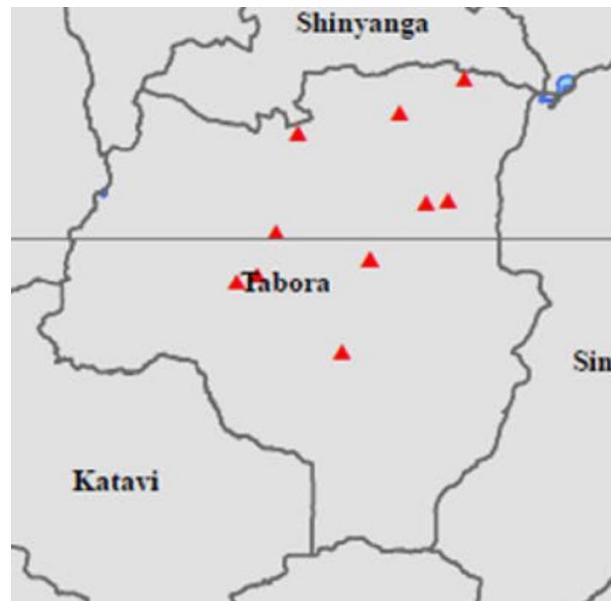
**Module 5:** Dealing with over-dispersion in multi-cluster studies

# What is a multi-cluster study?

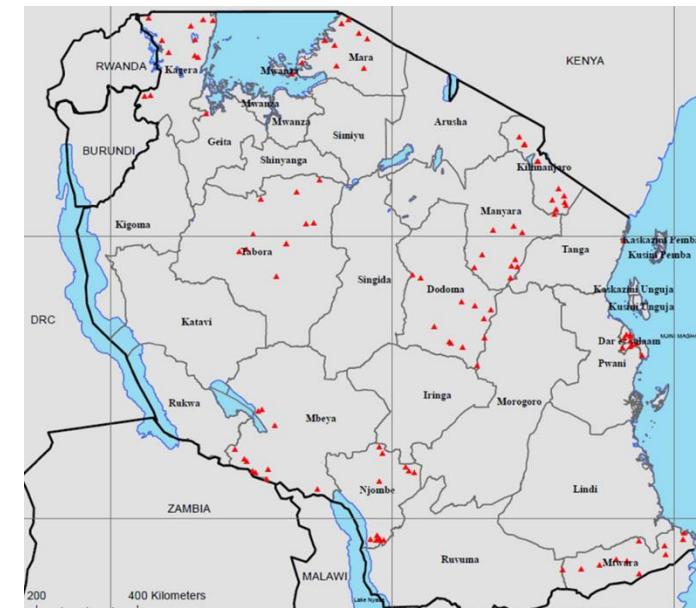
In a multi-cluster study...

- We conduct the study at several sites (clusters)
- **We aim to draw conclusions at a higher level than the site**

**Regional level**



**Country level**



## We can combine information across sites

- Regional-level estimates aim to draw conclusion about the wider population
- Interventions are often delivered at regional level

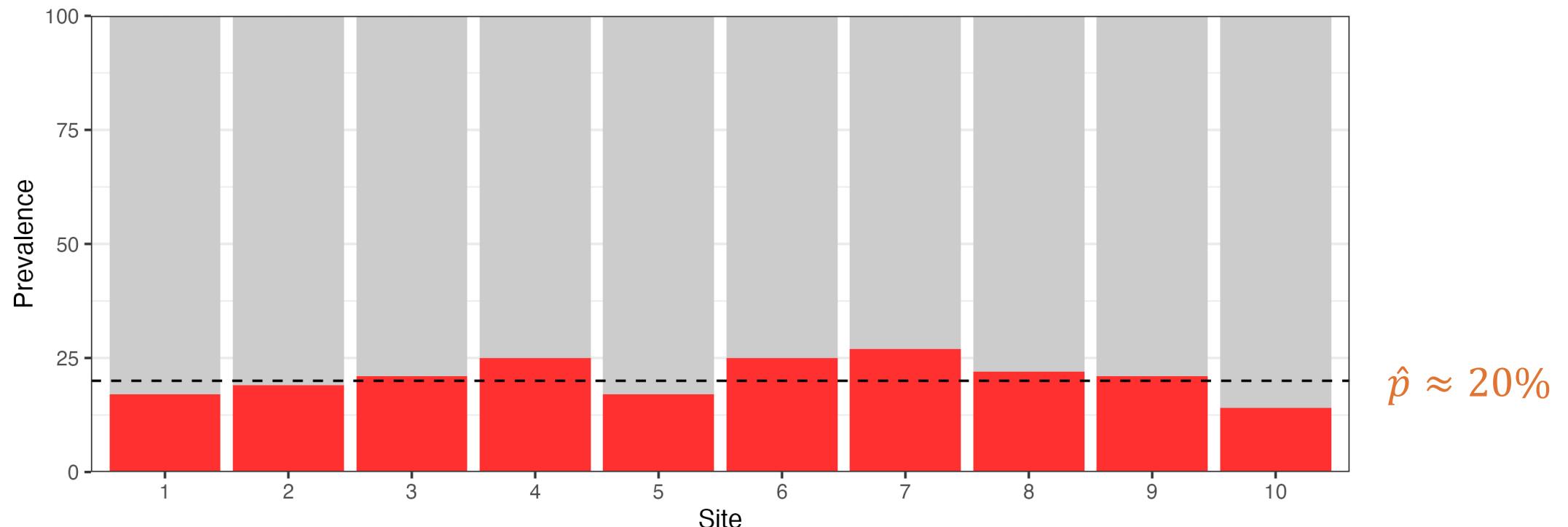
## Or we can explore differences between sites

- Are there geographic trends?
- What is the geographic scale of the threat?
- Can we identify cluster-level covariates?

# Over-dispersion

- Prevalence study over 10 sites
- 100 samples per site
- global prevalence of 20%

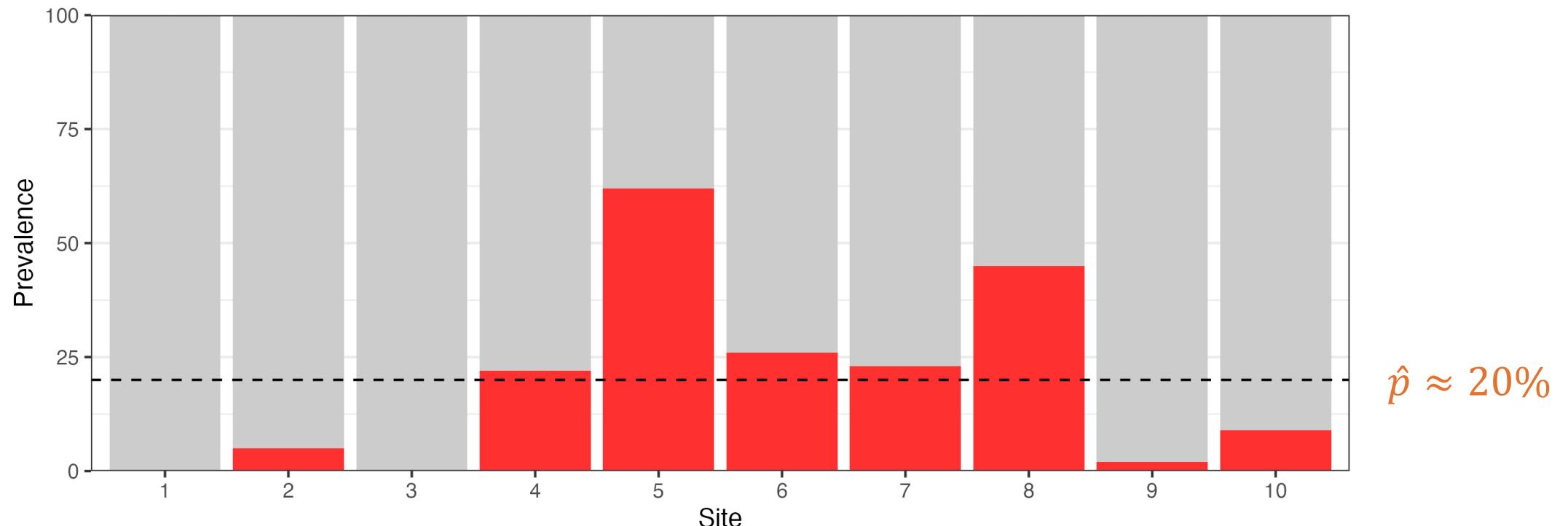
This is what the data spread looks like  
when samples are perfectly **independent**



# Over-dispersion

- Prevalence study over 10 sites
- 100 samples per site
- global prevalence of 20%

This is what data really look like!



## Overdispersion

Sites are more **different** than we would expect on average



## Intra-cluster correlation

People within sites are more **similar** than we would expect on average

# What causes intra-cluster correlation

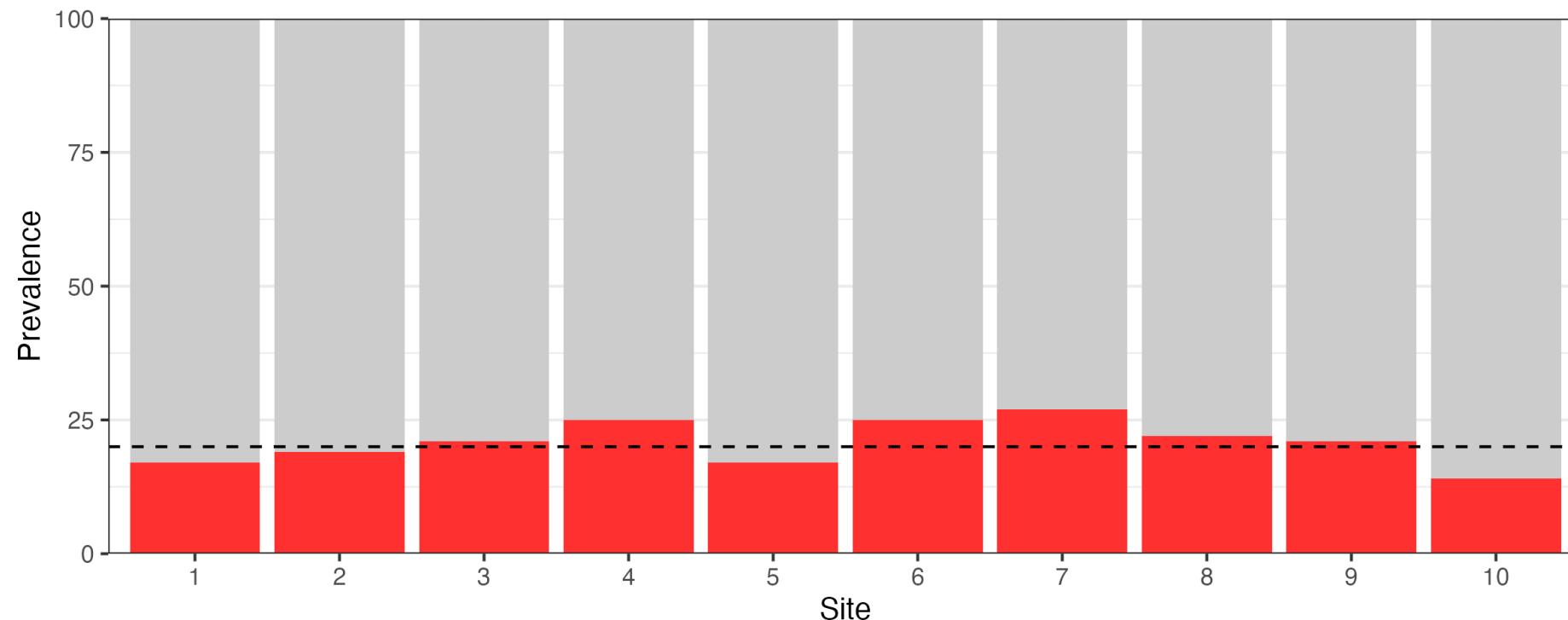
- Similar behaviours/customs
- Similar occupations
- Shared vector reservoirs
- Genetic similarities
- Similar access to healthcare
- Local transmission and outbreaks



# Detecting over-dispersion

There is a level of variability between sites that we expect:

$$\hat{p} \pm \sqrt{\frac{\hat{p}(1 - \hat{p})}{n_i}}$$

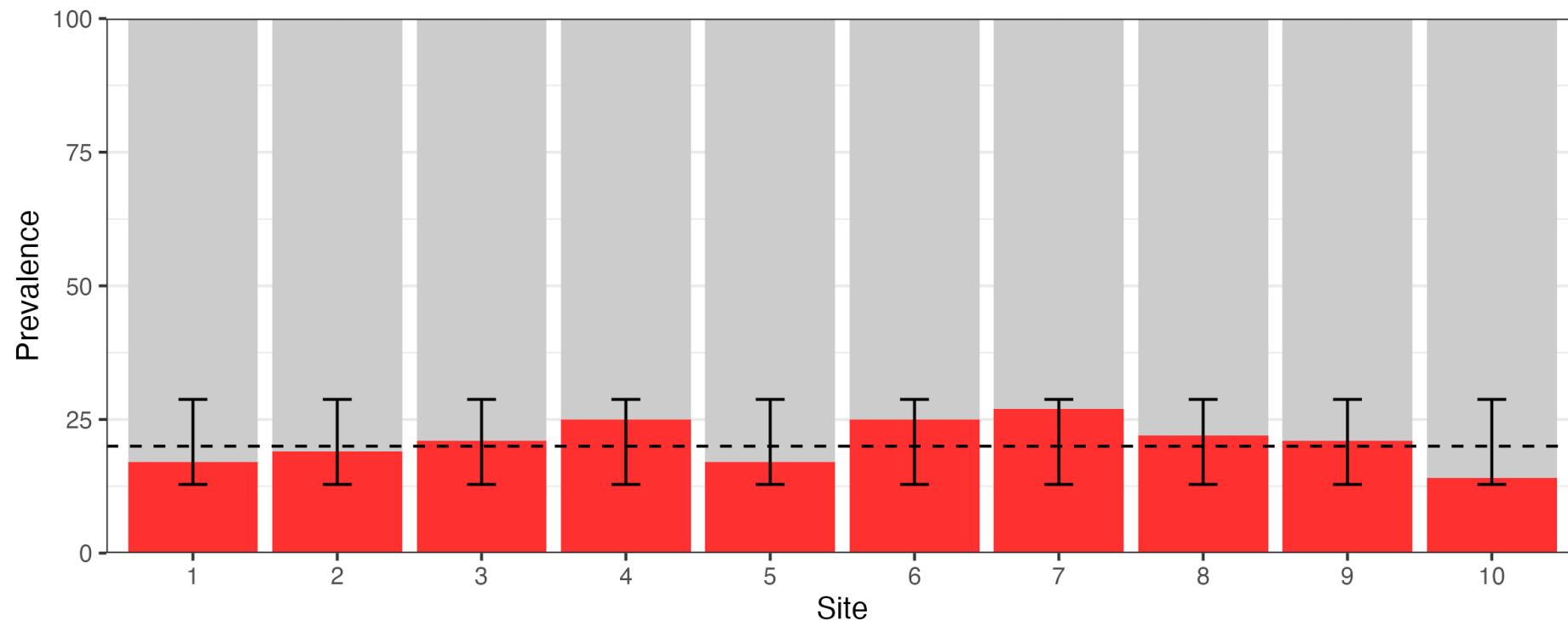


# Detecting over-dispersion

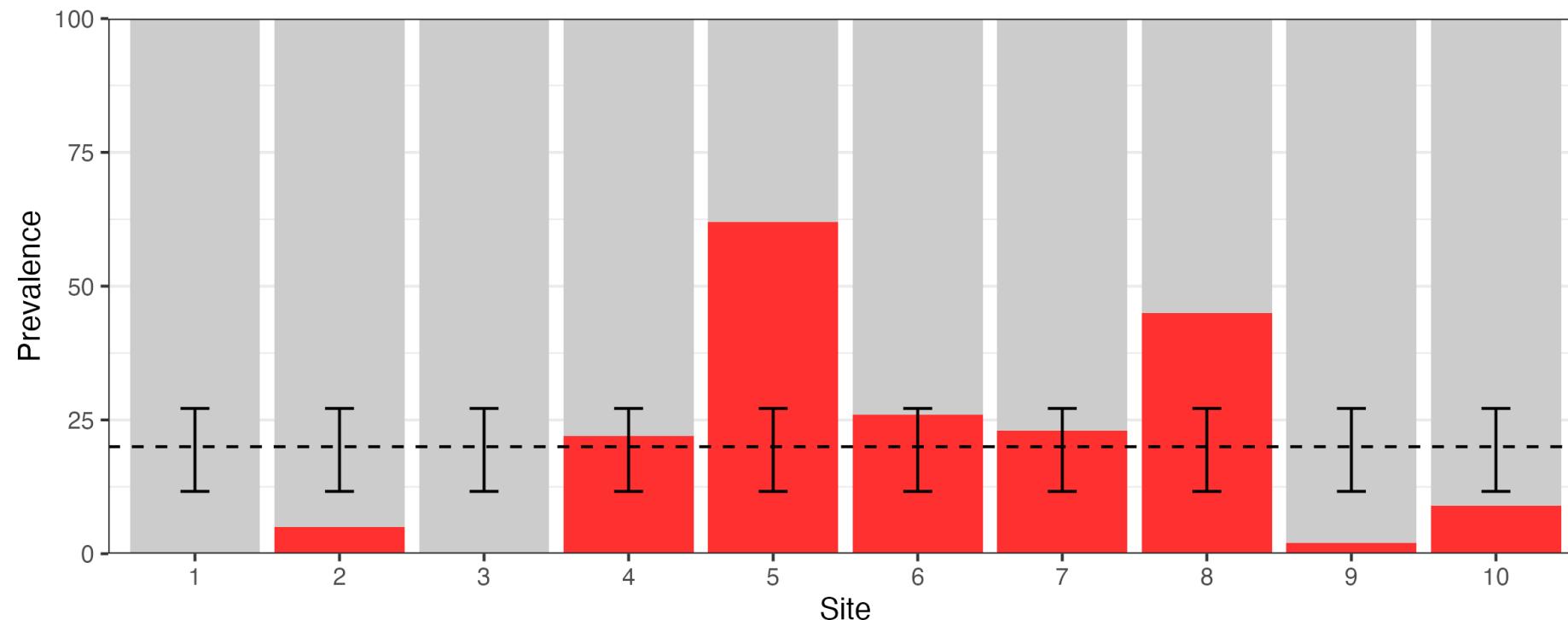
There is a level of variability between sites that we expect:

$$\hat{p} \pm \sqrt{\frac{\hat{p}(1 - \hat{p})}{n_i}}$$

95% of sites within this range

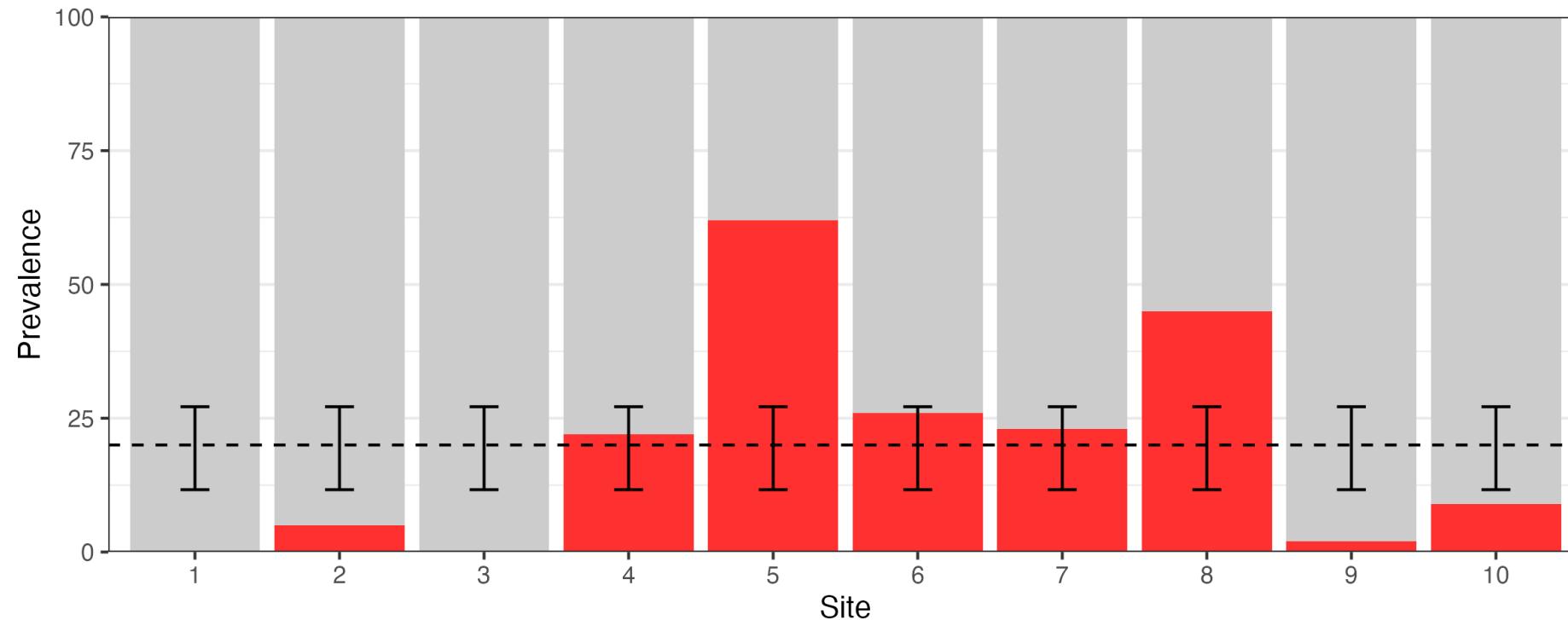


If more than about 10% of sites are outside the range, there is likely some **systematic** overdispersion



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**70% outside the range!**



1. Design effect
2. Effective sample size
3. Intra-cluster correlation coefficient

$$D_{\text{eff}} = \frac{\text{What is the variance of my data?}}{\text{What variance would I expect if samples were independent?}}$$

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The design effect is a measure statistical **inefficiency**. A value of  $D_{\text{eff}} = 1$  is gold standard (although  $D_{\text{eff}}$  can be less than 1).

# The Design Effect



$$D_{\text{eff}} = \frac{\text{Var}_{\text{obs}}}{\text{Var}_{\text{SRS}}}$$

# The Design Effect

$$D_{\text{eff}} = \frac{\text{Var}_{\text{obs}}}{\text{Var}_{\text{SRS}}} = \frac{s^2}{\frac{1}{c} \sum_{i=1}^c \frac{\hat{p}(1 - \hat{p})}{n_i}}$$

$s^2$  = sample variance

$\hat{p}$  = global prevalence estimate

$c$  = number of clusters

$n_i$  = sample size in  $i^{th}$  cluster

## The Design Effect – worked example

Site	Sample size	Prevalence
1	60	0.00
2	80	0.05
3	70	0.00
4	100	0.22
5	40	0.62
6	60	0.26
7	50	0.23
8	90	0.09

See the Excel file  
[Overdispersion\\_example.xlsx](#)  
to work through steps  
(available on course website)

## The Design Effect – worked example

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$$\text{Var}_{\text{obs}} = 0.0420$$

$$\text{Var}_{\text{SRS}} = 0.0024$$

$$D_{\text{eff}} = 17.73$$

That's great...but what does a value  $D_{\text{eff}} = 17.73$  really mean?

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$N_{\text{eff}}$  is the number of completely independent samples you would need to achieve the same level of precision as your more complex study design

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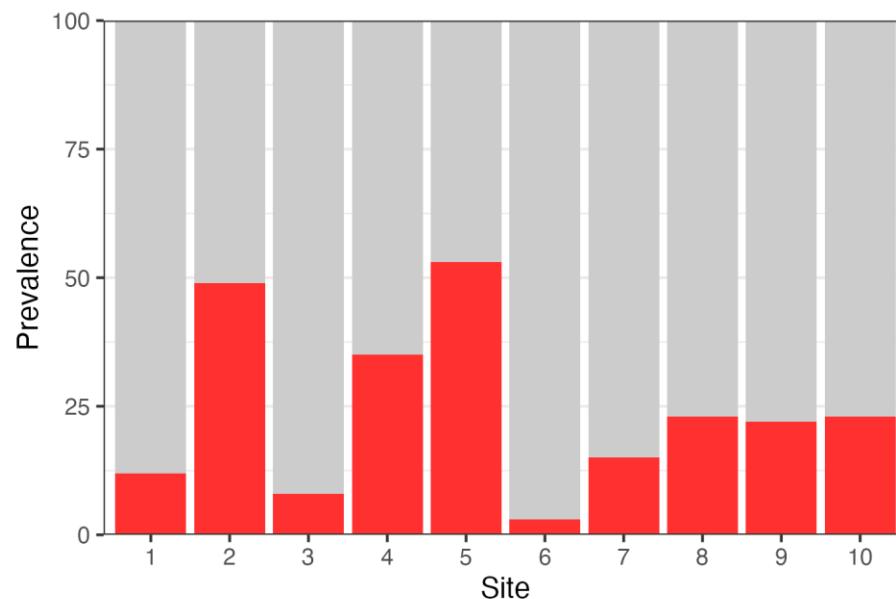
Back to [Overdispersion\\_example.xlsx](#)

# The effective sample size

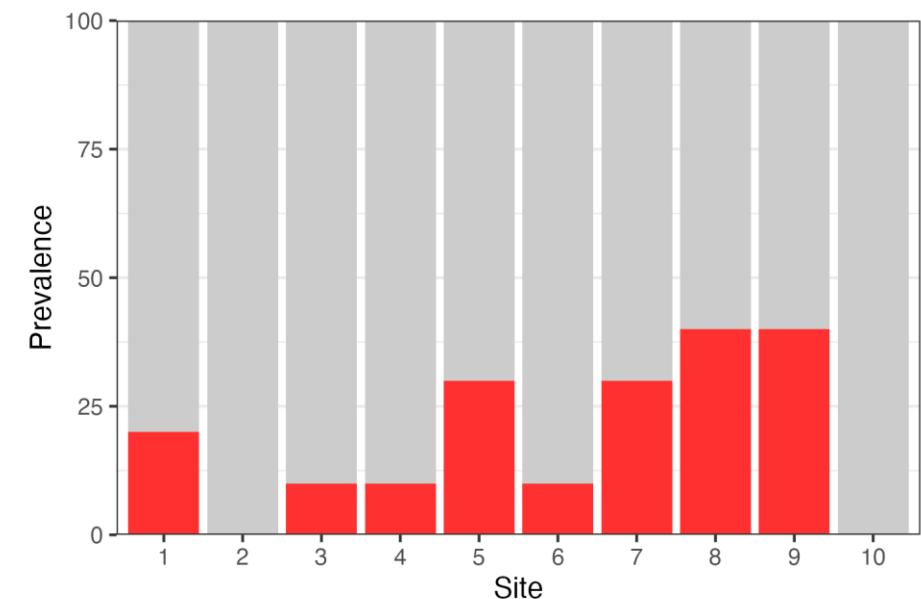
One of these was generated with  $N = 100$   
the other with  $N = 1000$  but  $N_{\text{eff}} = 100$

## Which one is which?

?



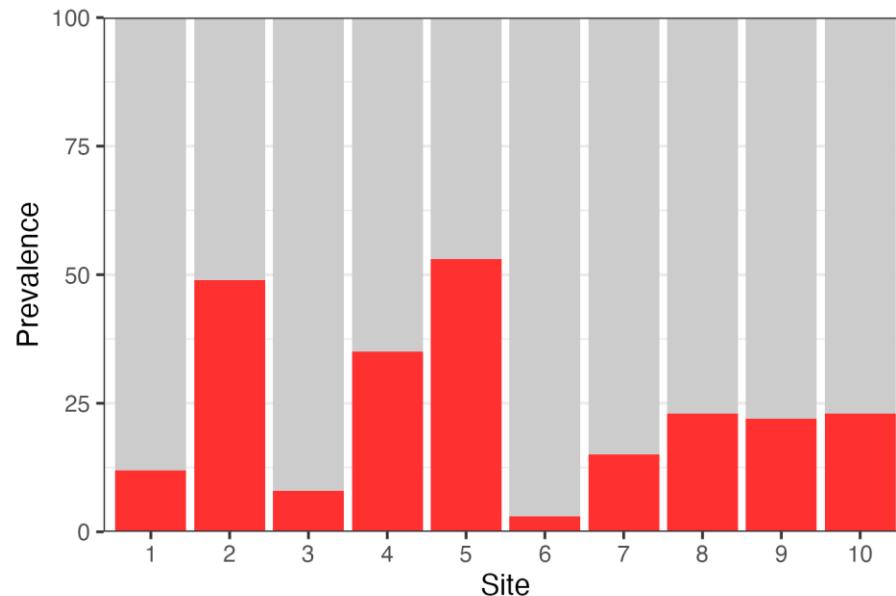
?



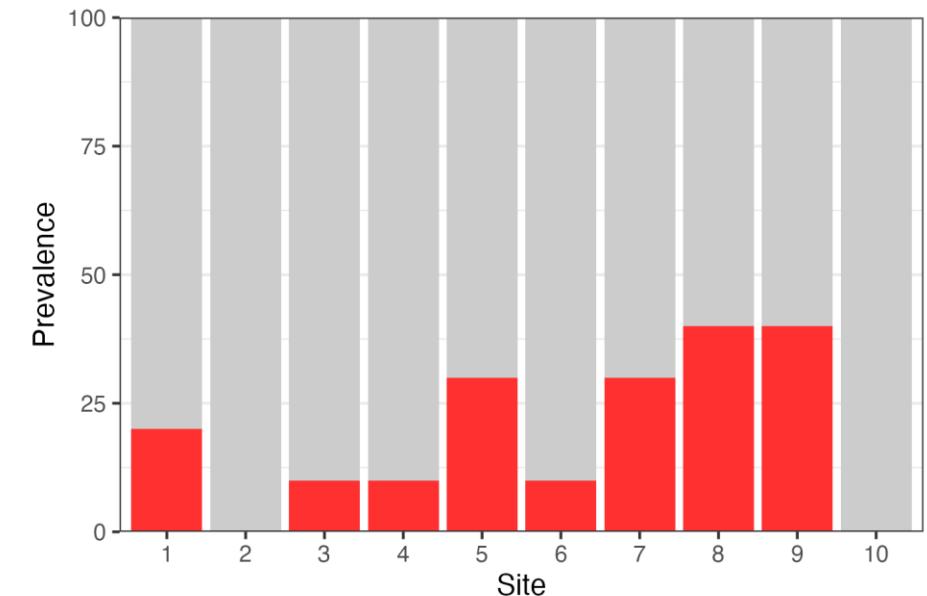
One of these was generated with  $N = 100$   
the other with  $N = 1000$  but  $N_{\text{eff}} = 100$

## Which one is which?

$N = 1000$



$N = 100$



## Overdispersion

Sites are more **different** than we would expect on average



## Intra-cluster correlation

People within sites are more **similar** than we would expect on average

The ICC ( $r$ ) is a value between 0 and 1 that represents the correlation between individuals in the same site.

We can write the design effect in terms of the ICC:

$$D_{\text{eff}} = 1 + (\bar{n} - 1)r$$

$\bar{n}$  = average cluster size

$r$  = ICC

We can write the ICC in terms of the design effect:

$$r = \frac{D_{\text{eff}} - 1}{\bar{n} - 1}$$

# The intra-cluster correlation coefficient



Back to [Overdispersion\\_example.xlsx](#)

# Why is the ICC useful?



## Scenario

Your study population has a true ICC of  $r = 0.053$  in terms of *mdr1 N86Y* prevalence.  
Assume we do not know the true ICC.

A pilot study is run over 10 clusters using a sample size of  $N = 200$ , divided into  $n = 20$  per cluster. Over-dispersion is measured, and we find a design effect of  $D_{\text{eff}} = 2.0$ .

A follow-up study is now planned with a much larger sample size of  $N = 10,000$ , divided into  $n = 1000$  per cluster. When designing the study, we assume the same design effect as the pilot, meaning we expect an effective sample size of  $N_{\text{eff}} = 5,000$ , which is still very large.

When the results come in, we measure the design effect on the new data. We find it has increased to  $D_{\text{eff}} = 53.9$ ! We now only have an effective sample size of just  $N_{\text{eff}} = 185$ !

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**What happened here!?**

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$$D_{\text{eff}} = 1 + (\bar{n} - 1)r$$

### Pilot study

$$\begin{aligned} \bar{n} &= 20 & \longrightarrow & D_{\text{eff}} = 2.0 \\ r &= 0.053 \end{aligned}$$

## Why is the ICC useful?

$$D_{\text{eff}} = 1 + (\bar{n} - 1)r$$

### Pilot study

$$\begin{aligned}\bar{n} &= 20 \\ r &= 0.053\end{aligned} \quad \longrightarrow \quad D_{\text{eff}} = 2.0$$

### Follow-up study

$$\begin{aligned}\bar{n} &= 1000 \\ r &= 0.053\end{aligned} \quad \longrightarrow \quad D_{\text{eff}} = 53.9$$

# Why is the ICC useful?

$$D_{\text{eff}} = 1 + (\bar{n} - 1)r$$

## Pilot study

$$\bar{n} = 20$$



$$D_{\text{eff}} = 2.0$$

$$r = 0.053$$

## Follow-up study

$$\bar{n} = 1000$$



$$D_{\text{eff}} = 53.9$$

$$r = 0.053$$

- ❖ The design effect is not an **intrinsic** measure of the population.
- ❖ It relates to **our study** of the population. It **depends on sample size** as well as intrinsic factors.
- ❖  $D_{\text{eff}}$  is hard to compare objectively between studies, while  $r$  is easy.

## 1. Design effect

- A simple measure of statistical inefficiency

## 2. Effective sample size

- An intuitive way of measuring efficiency

## 3. Intra-cluster correlation coefficient

- Facilitates comparison between studies

# How can we design multi-cluster studies?



New versions of formulae (precision, power, sample size etc.) that take over-dispersion into account:

## Generalization of Wald interval:

$$\hat{p} \pm \sqrt{\frac{\hat{p}(1 - \hat{p})}{N} D_{\text{eff}}}$$

In the design stage, this means we will have to **assume** a value of the design effect, or the ICC

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

- Work with the NMCP of Tanzania to analyse data from a multi-site *pfhrp2/3* deletion prevalence study
- Detect and quantify over-dispersion in the data
- Plan a new study that accounts for over-dispersion



[Workshop materials](#)

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