

Malaria Molecular Surveillance Study Design Workshop

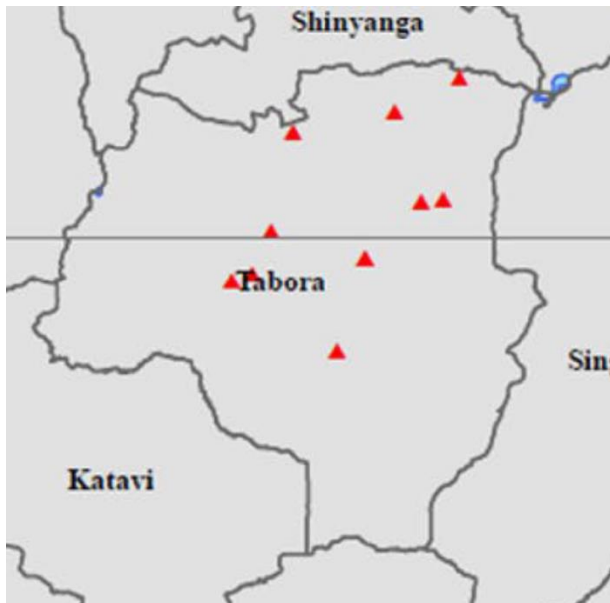
Module 4: Dealing with over-dispersion in multi- cluster studies

What is a multi-cluster study?

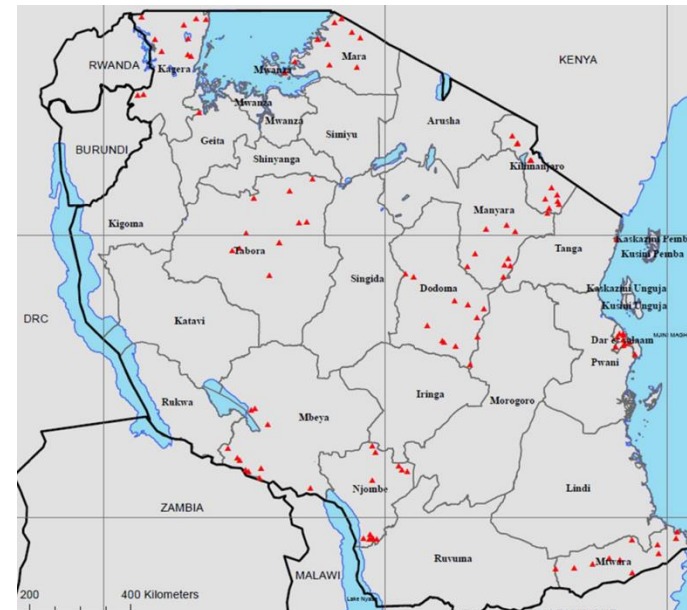
In a multi-cluster study...

- 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

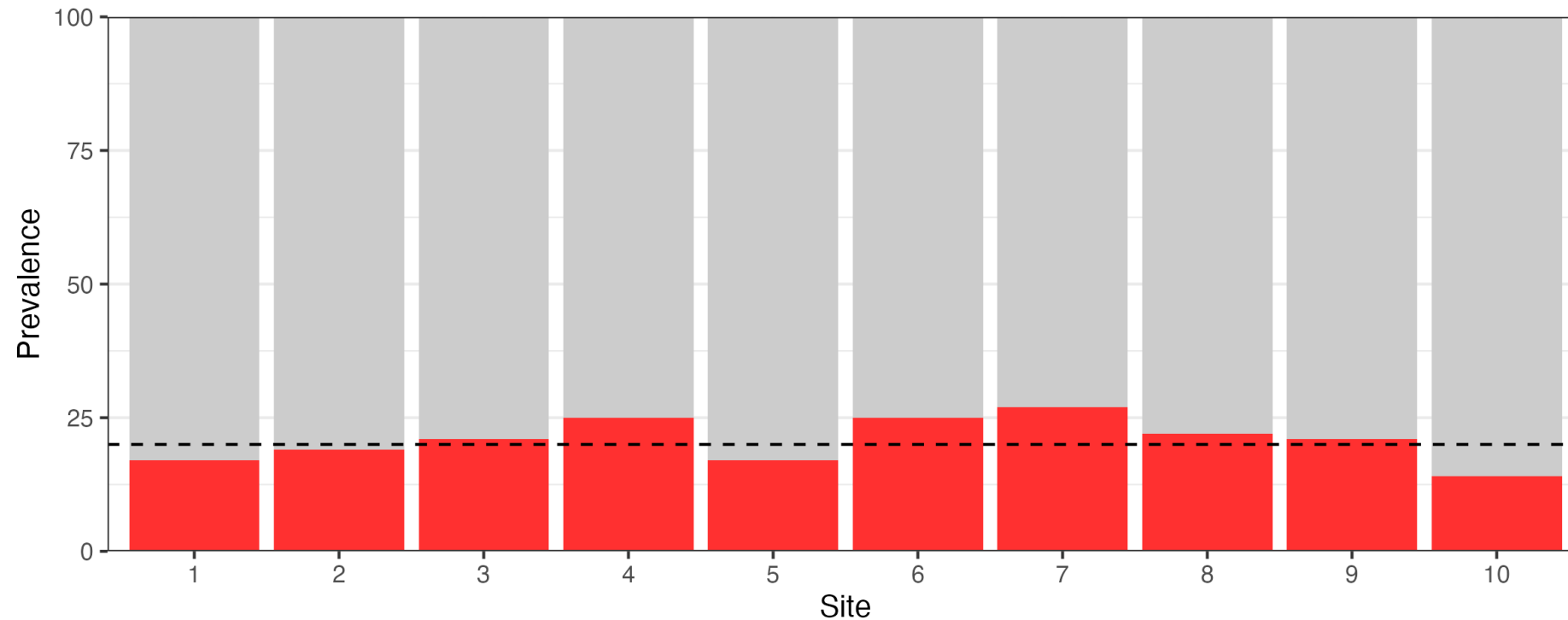
Or we can explore differences between sites

- 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**

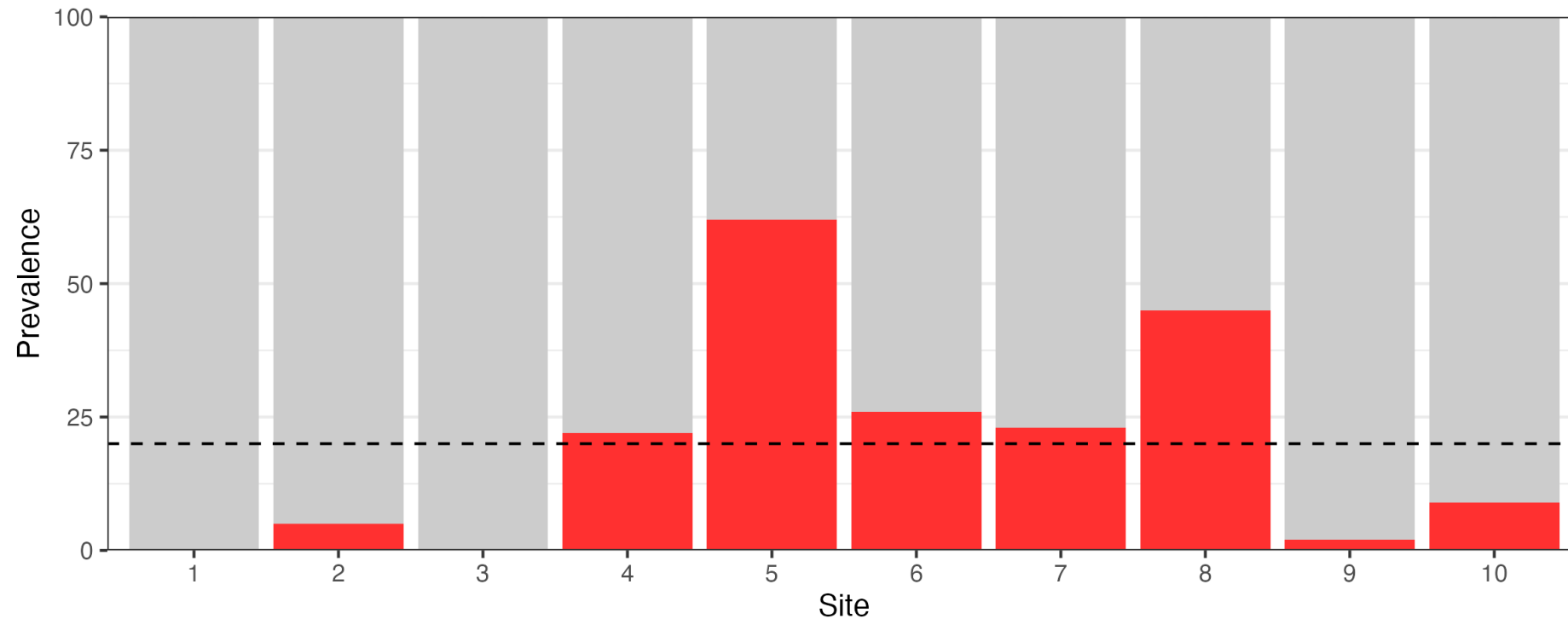


$\hat{p} \approx 20\%$

Over-dispersion

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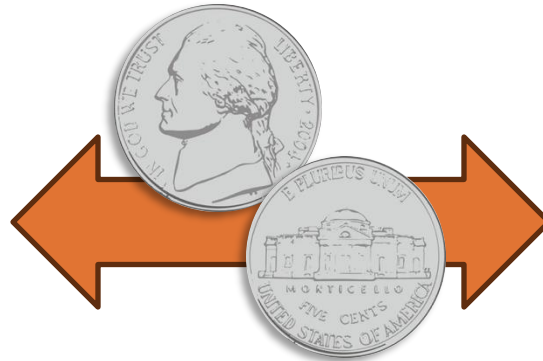
This is what data really look like!



$\hat{p} \approx 20\%$

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

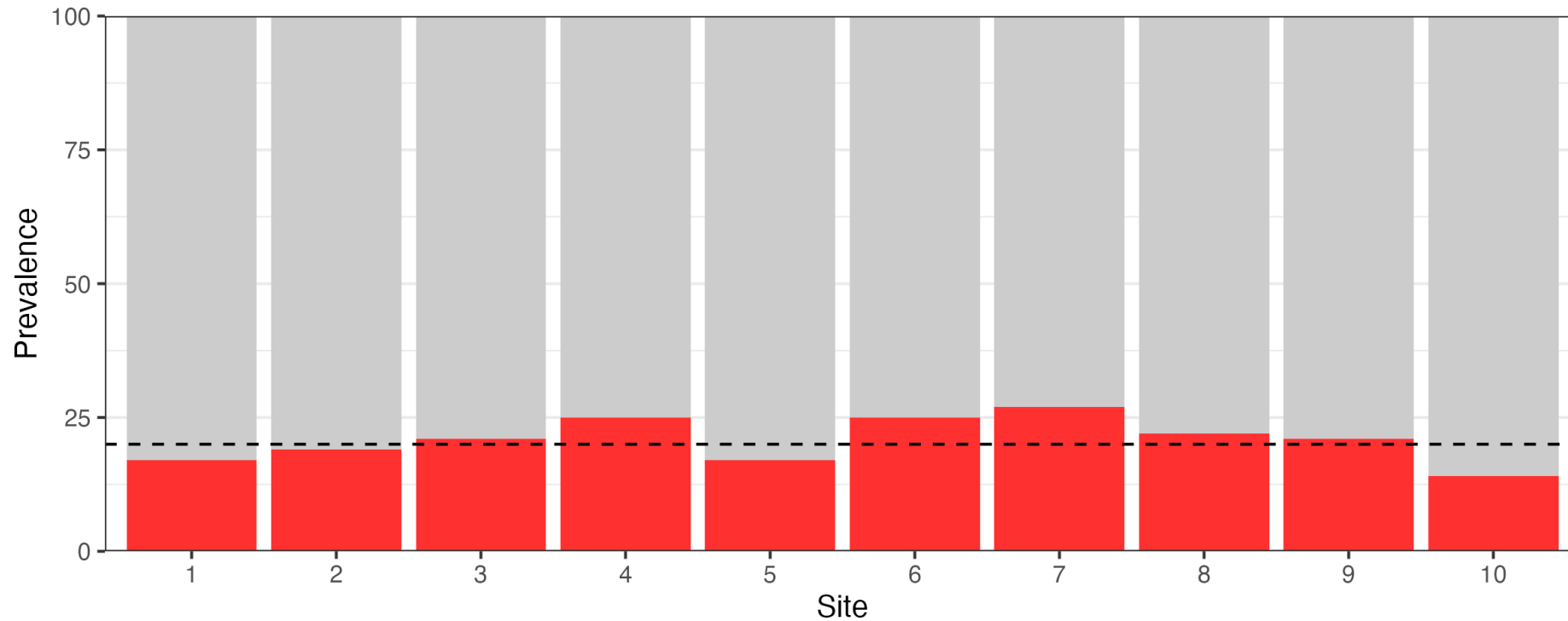
- Similar behaviours/customs
- Similar occupations
- Shared vector reservoirs
- Genetic similarities
- Similar access to healthcare
- Local transmission and outbreaks



Detecting over-dispersion

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

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

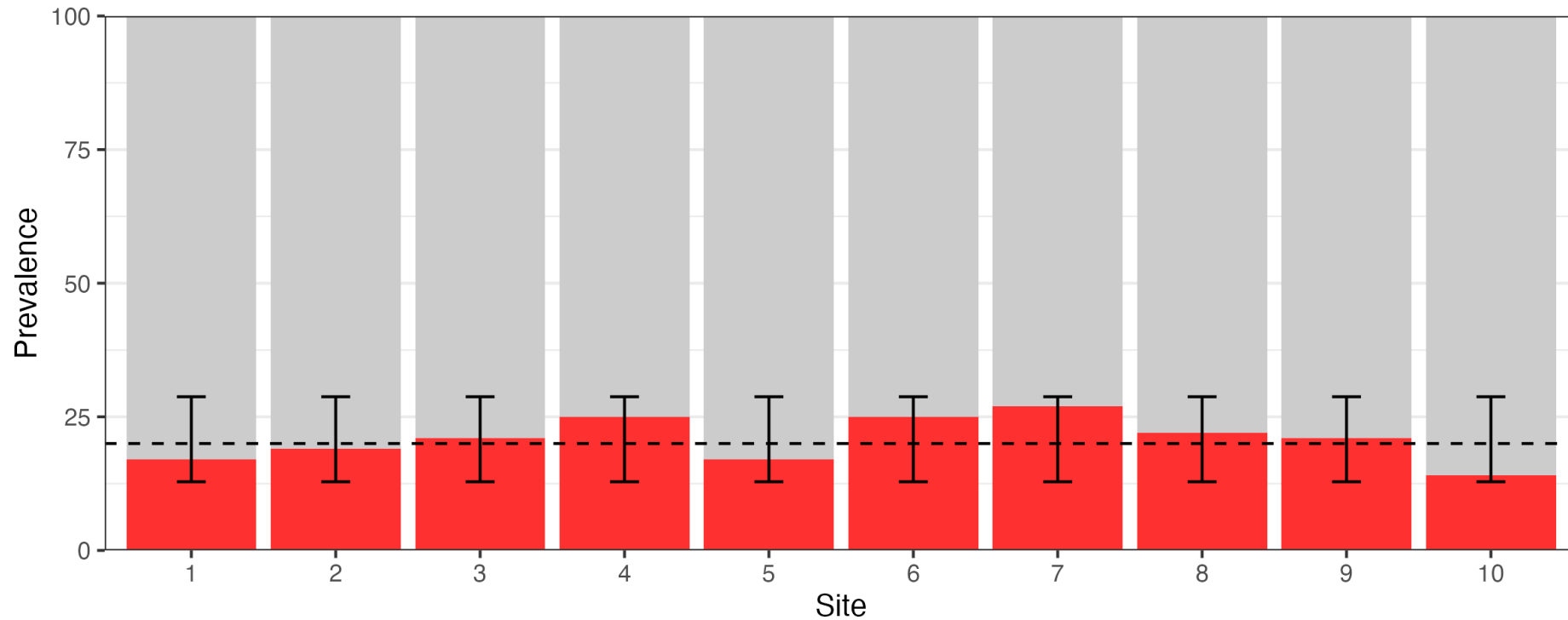


Detecting over-dispersion

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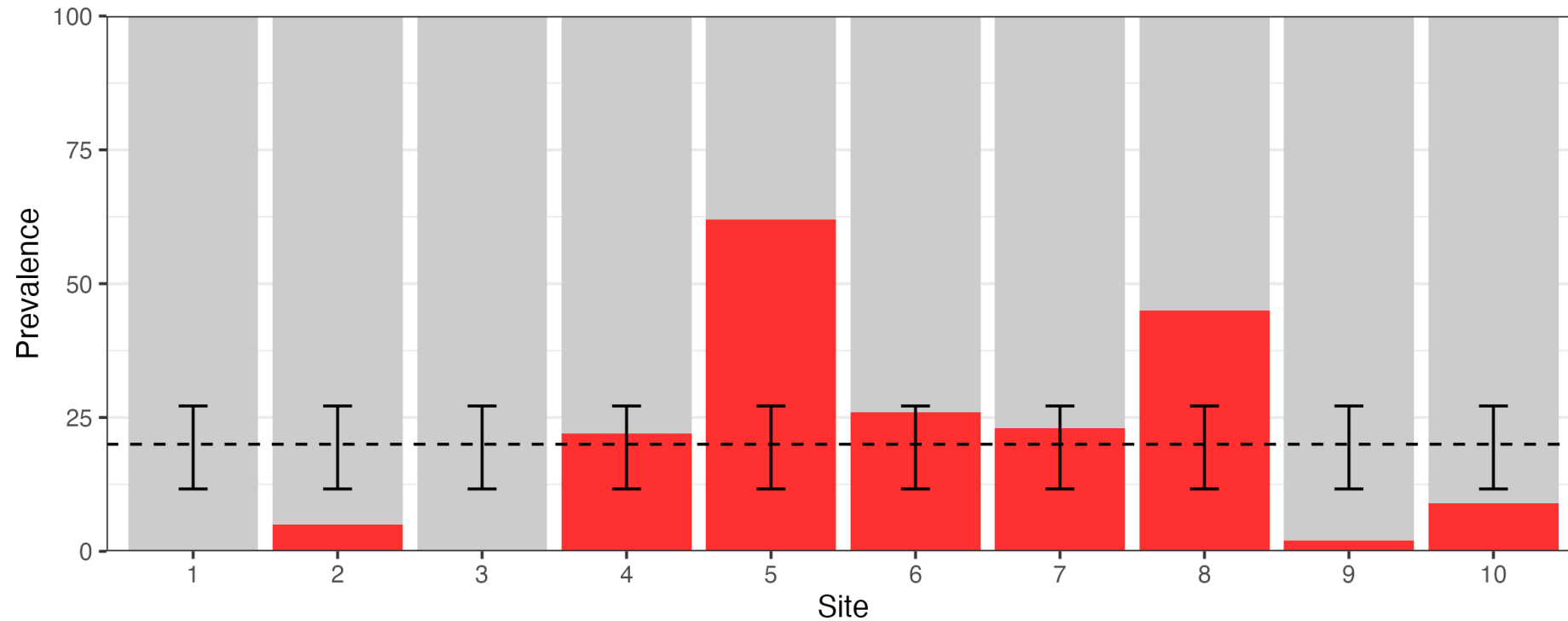
$$\hat{p} \pm \sqrt{\frac{\hat{p}(1 - \hat{p})}{n_i}}$$

95% of sites within this range



Detecting over-dispersion

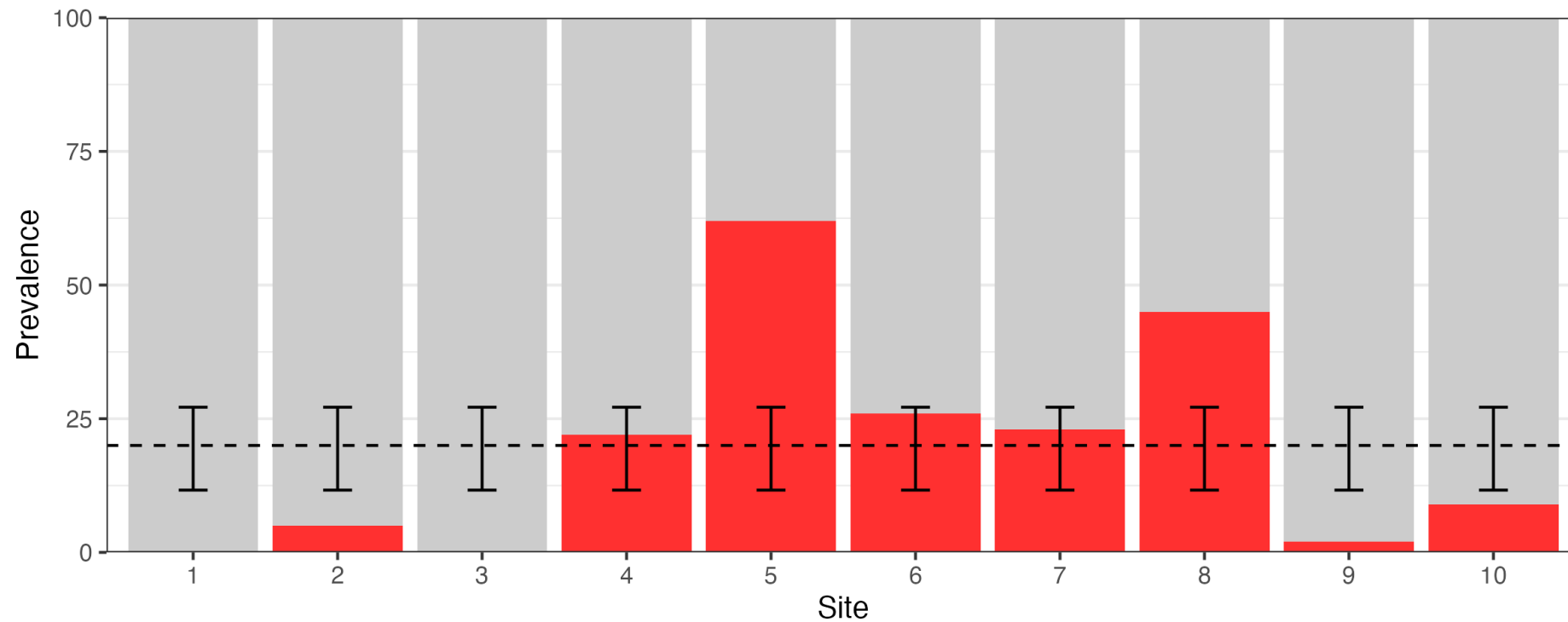
If more than about 10% of sites are outside the range, there is likely some **systematic** overdispersion



Detecting over-dispersion

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

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_{eff} can be less than 1).

The Design Effect



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

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

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

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

The effective sample size



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

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

The effective sample size



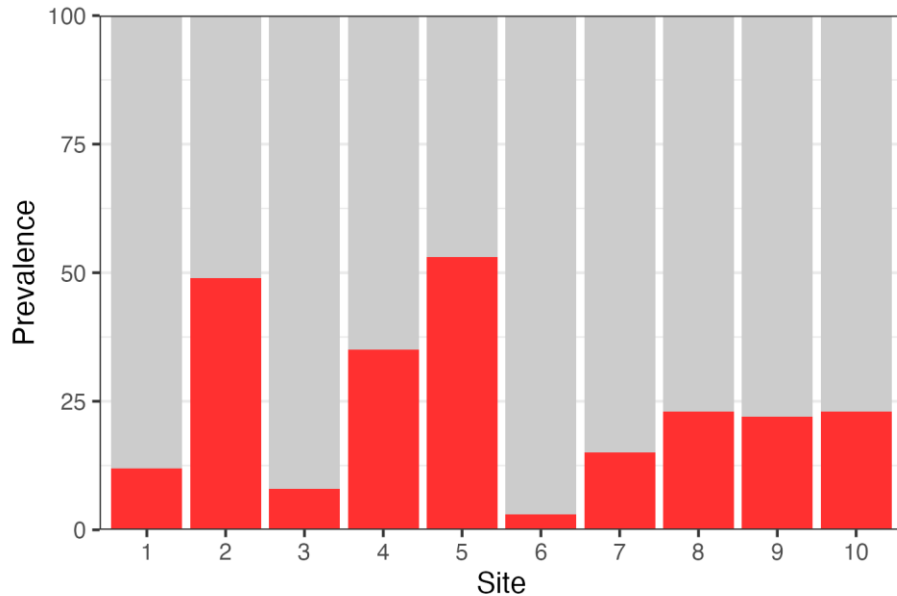
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The effective sample size

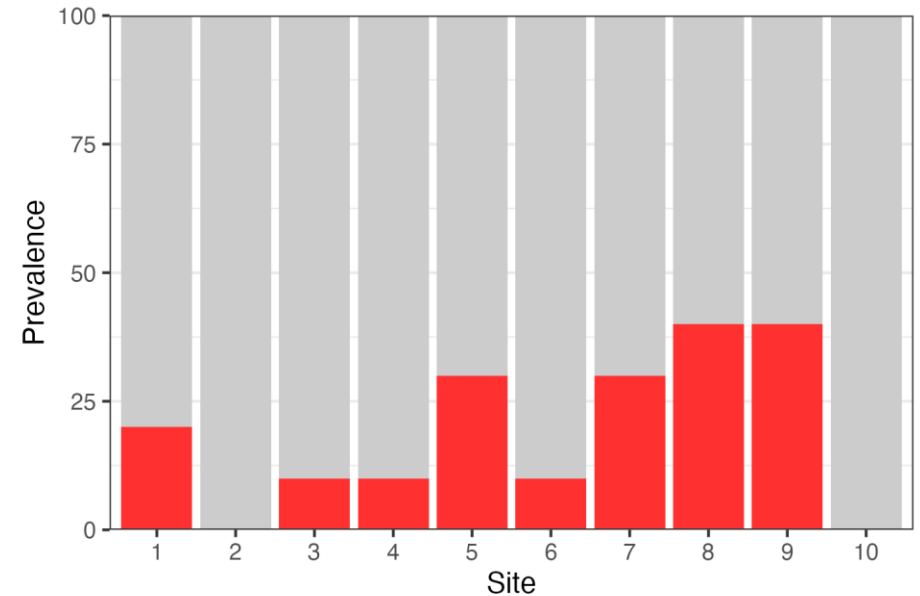
One of these was generated with $N = 100$
the other with $N = 1000$ but $N_{\text{eff}} = 100$

Which one is which?

?



?

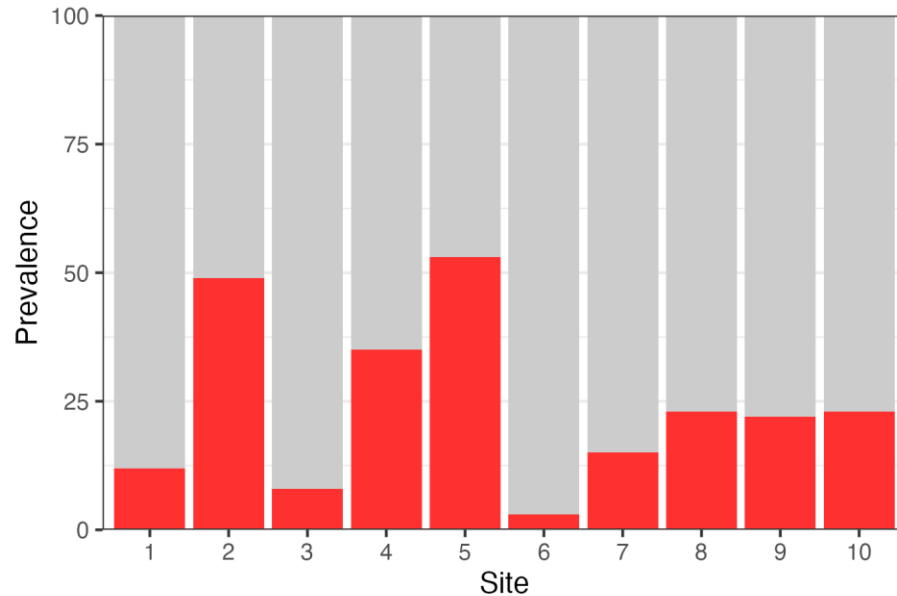


The effective sample size

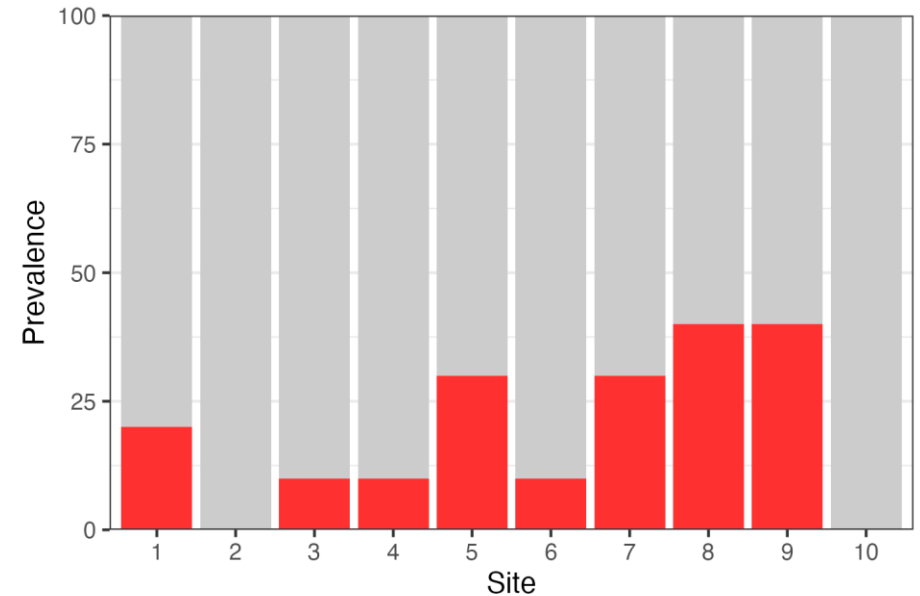
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$N = 1000$

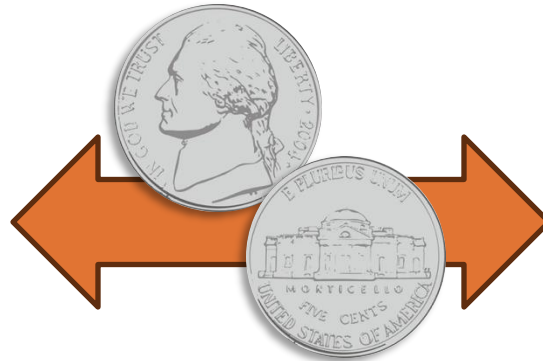


$N = 100$



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 intra-cluster correlation coefficient



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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Pilot study

$$\bar{n} = 20$$

$$r = 0.053$$



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

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Pilot study

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

Follow-up study

$$\begin{array}{l} \bar{n} = 1000 \\ r = 0.053 \end{array} \longrightarrow D_{\text{eff}} = 53.9$$

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Follow-up study

$$\begin{array}{l} \bar{n} = 1000 \\ r = 0.053 \end{array} \longrightarrow D_{\text{eff}} = 53.9$$

- ❖ 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_{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/)

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