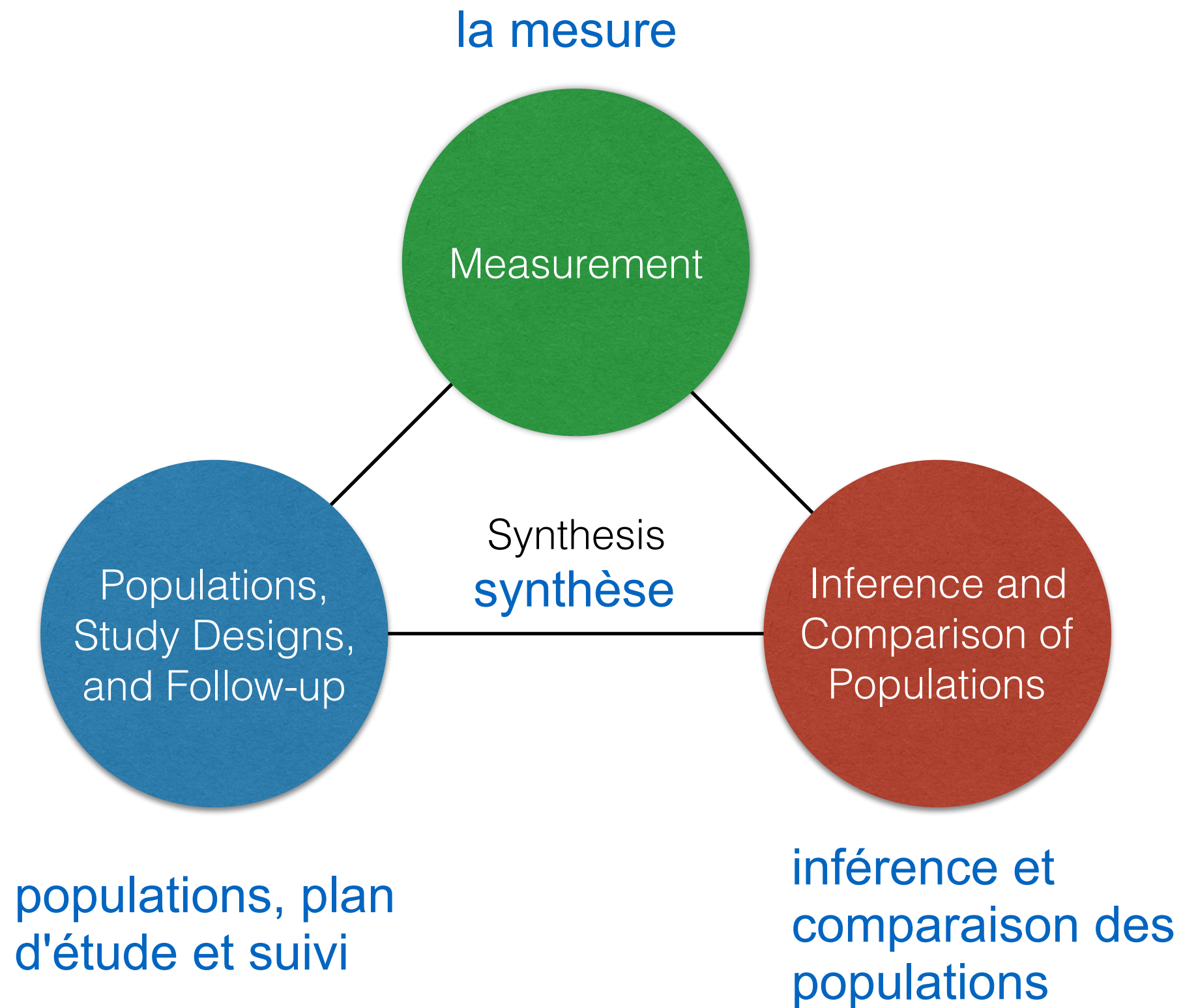


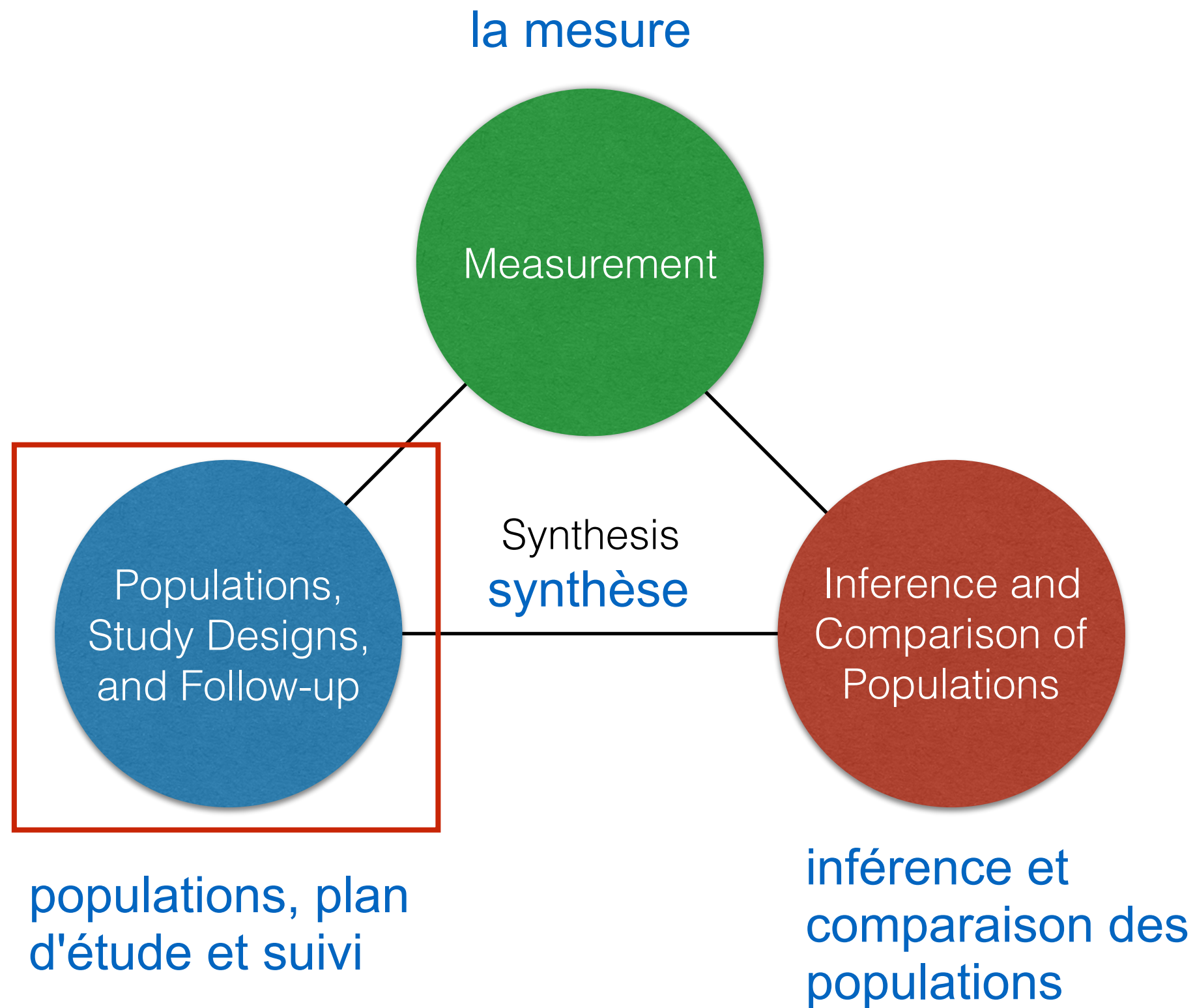
Epidemiological Study Design

Amy Wesolowski
Department of Epidemiology



JOHNS HOPKINS
BLOOMBERG SCHOOL
of PUBLIC HEALTH





Individual A

Time



Individual A



Time



Individual A

Outcome of interest
(disease)

résultat

Time



Epidemiological Questions

What exposure causes the outcome?

Quelle exposition
provoque le
résultat?

What are risk factors?

Quels sont les
facteurs de
risque?

What treatment should be given?

Quel traitement
devrait être
donné?

Individual A

Outcome of interest
(disease)

résultat

Time



Individual A



Time

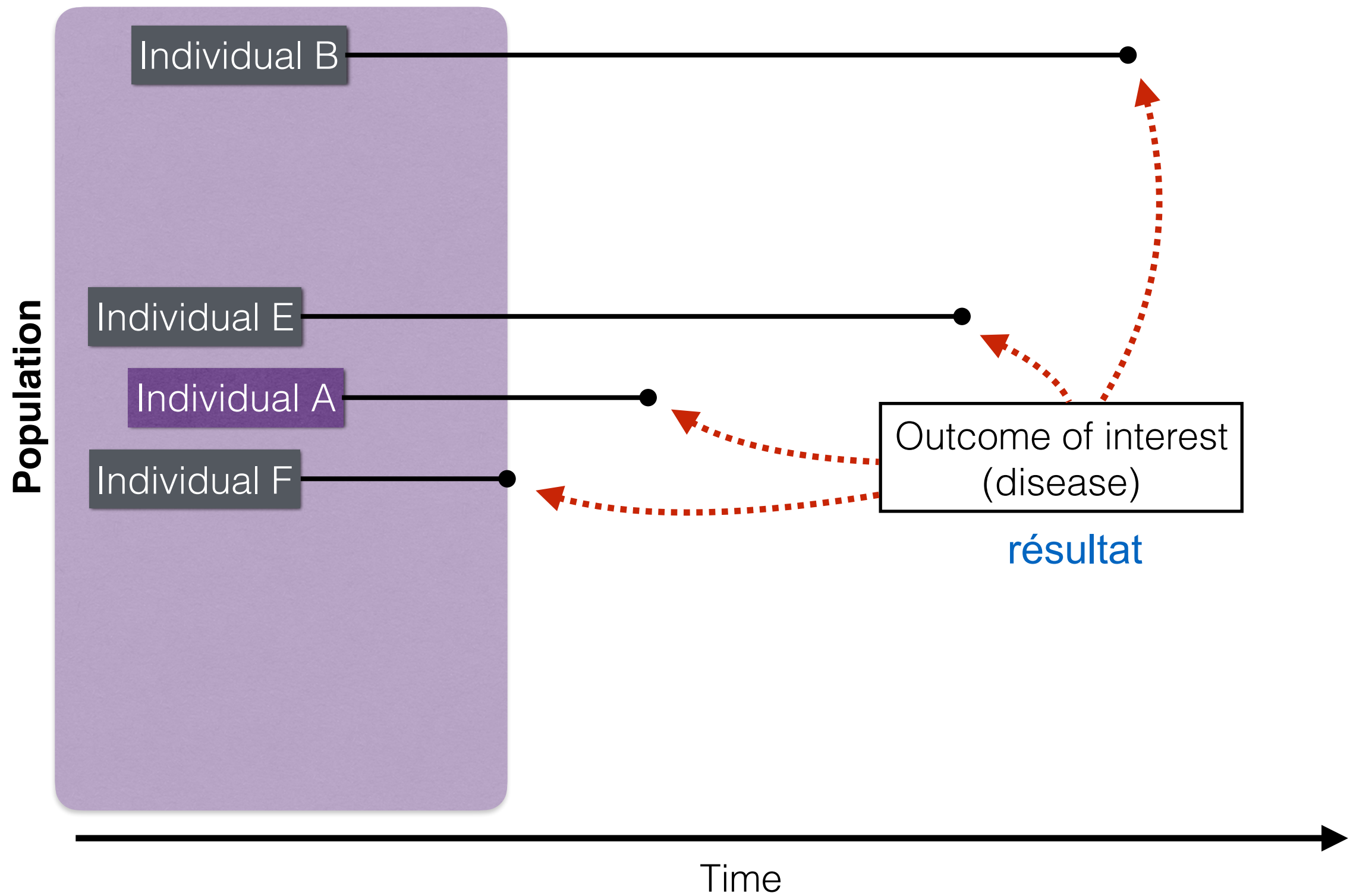


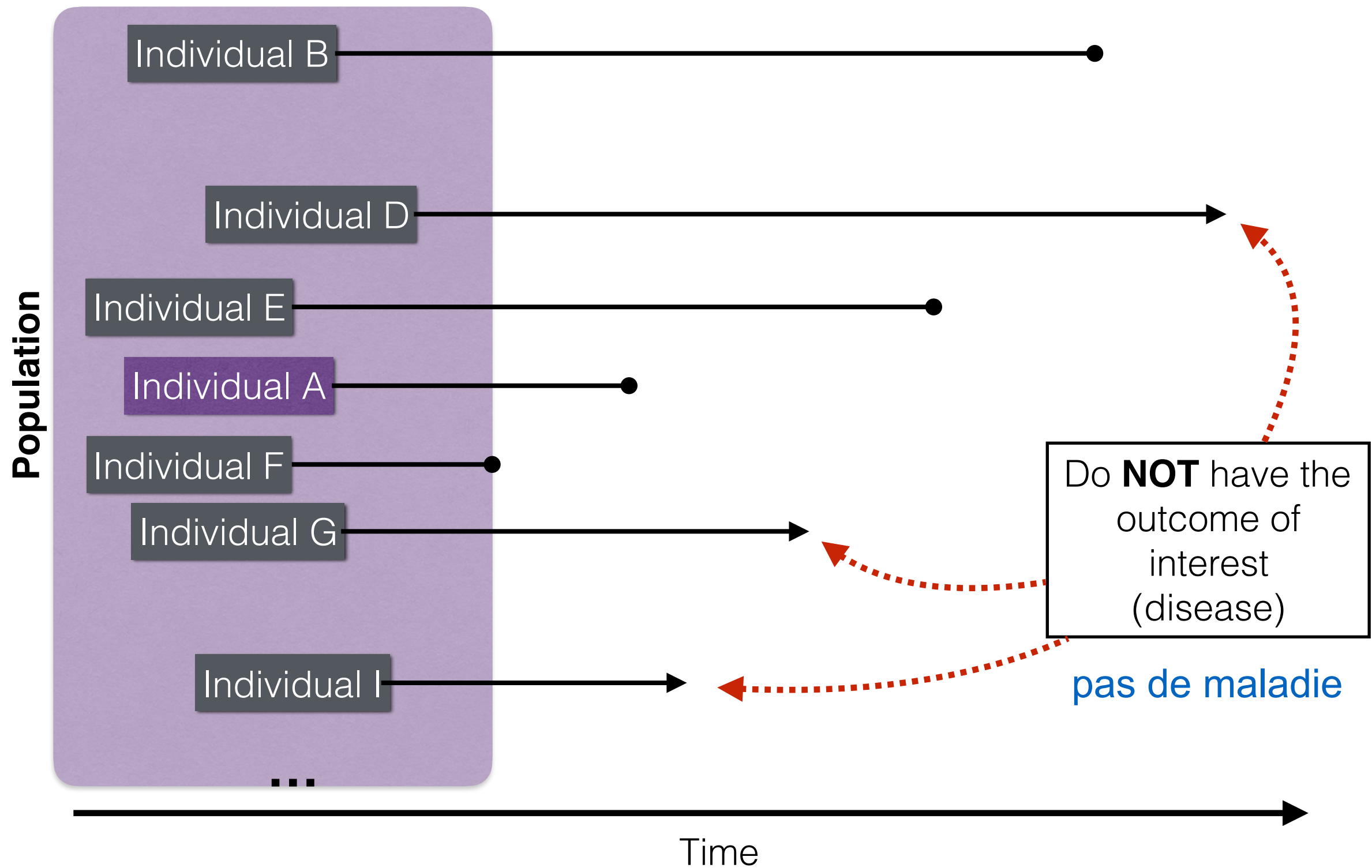
Population

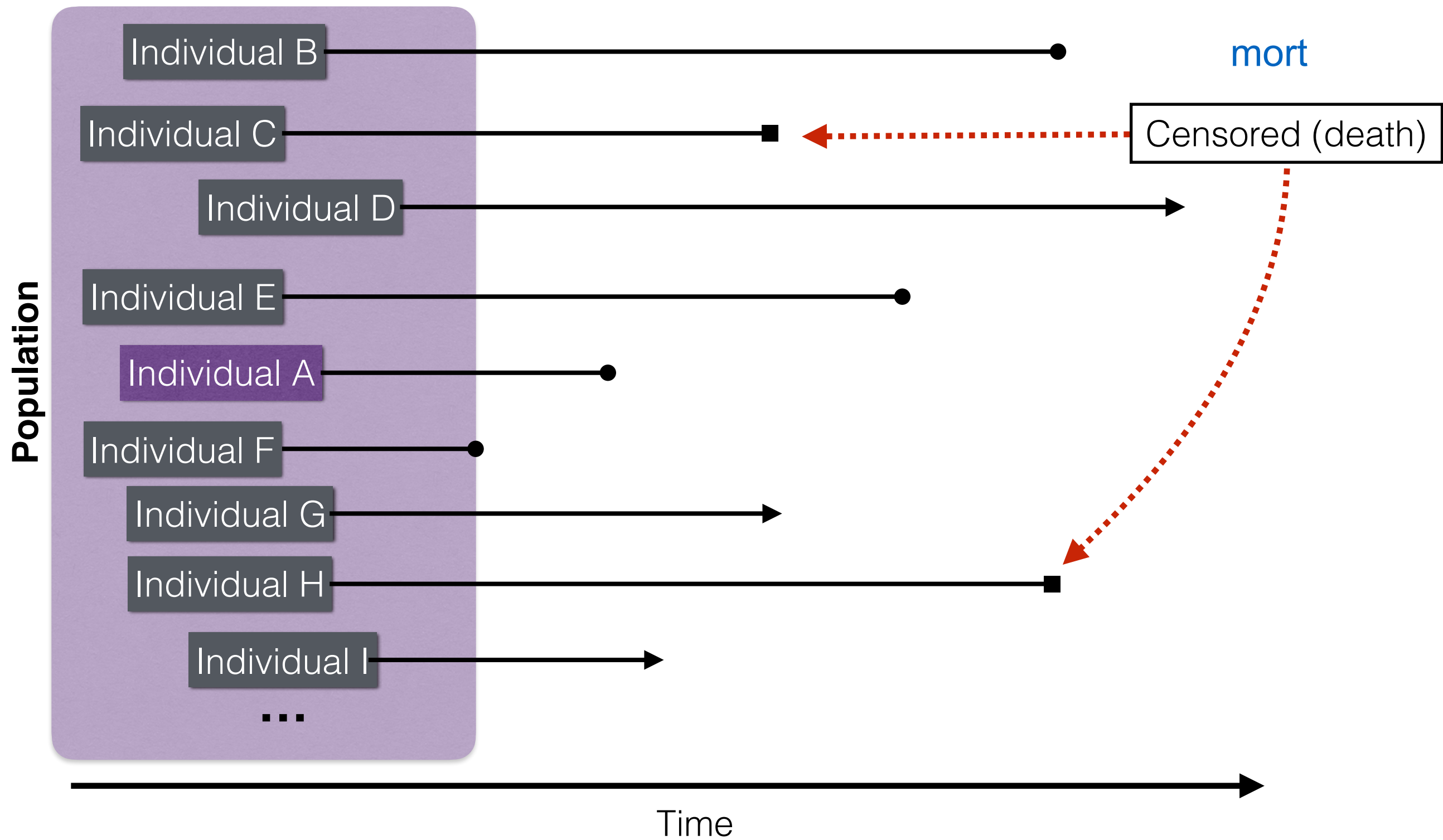
Individual A

Time









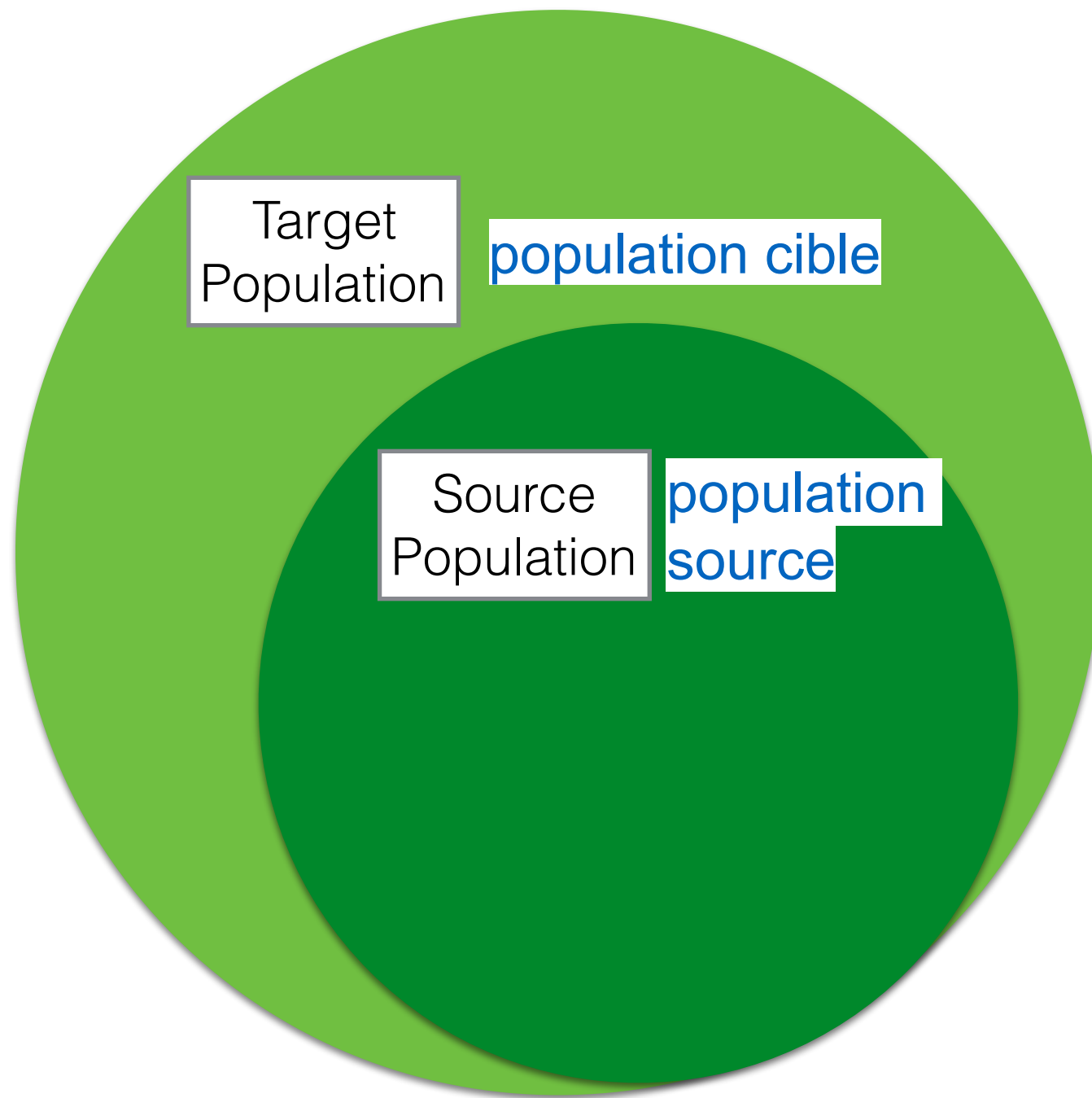


Target
Population

population cible

Target Population:
Want to make inferences about

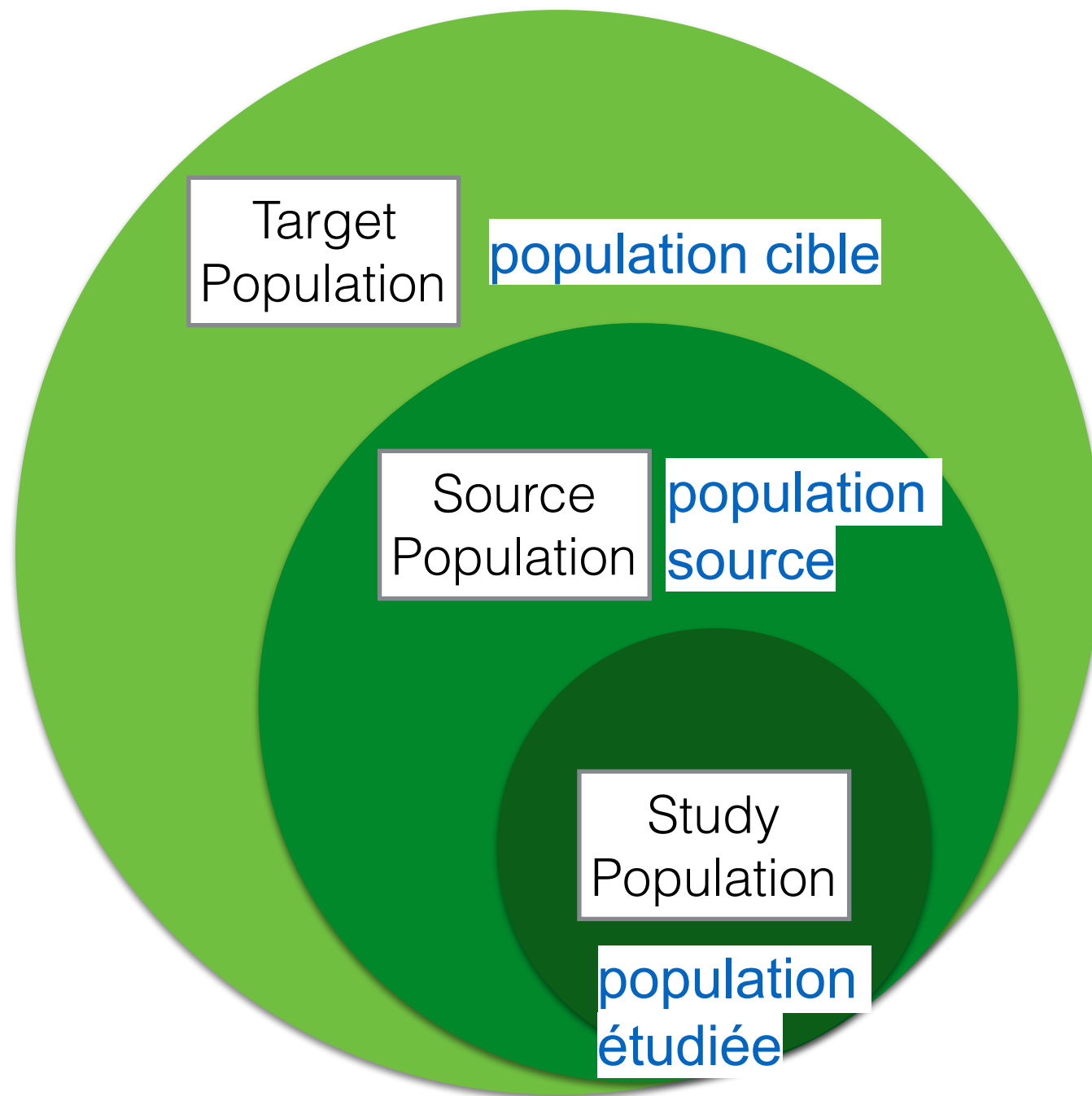




Target Population:
Want to make inferences about

Source Population:
Choosing study population



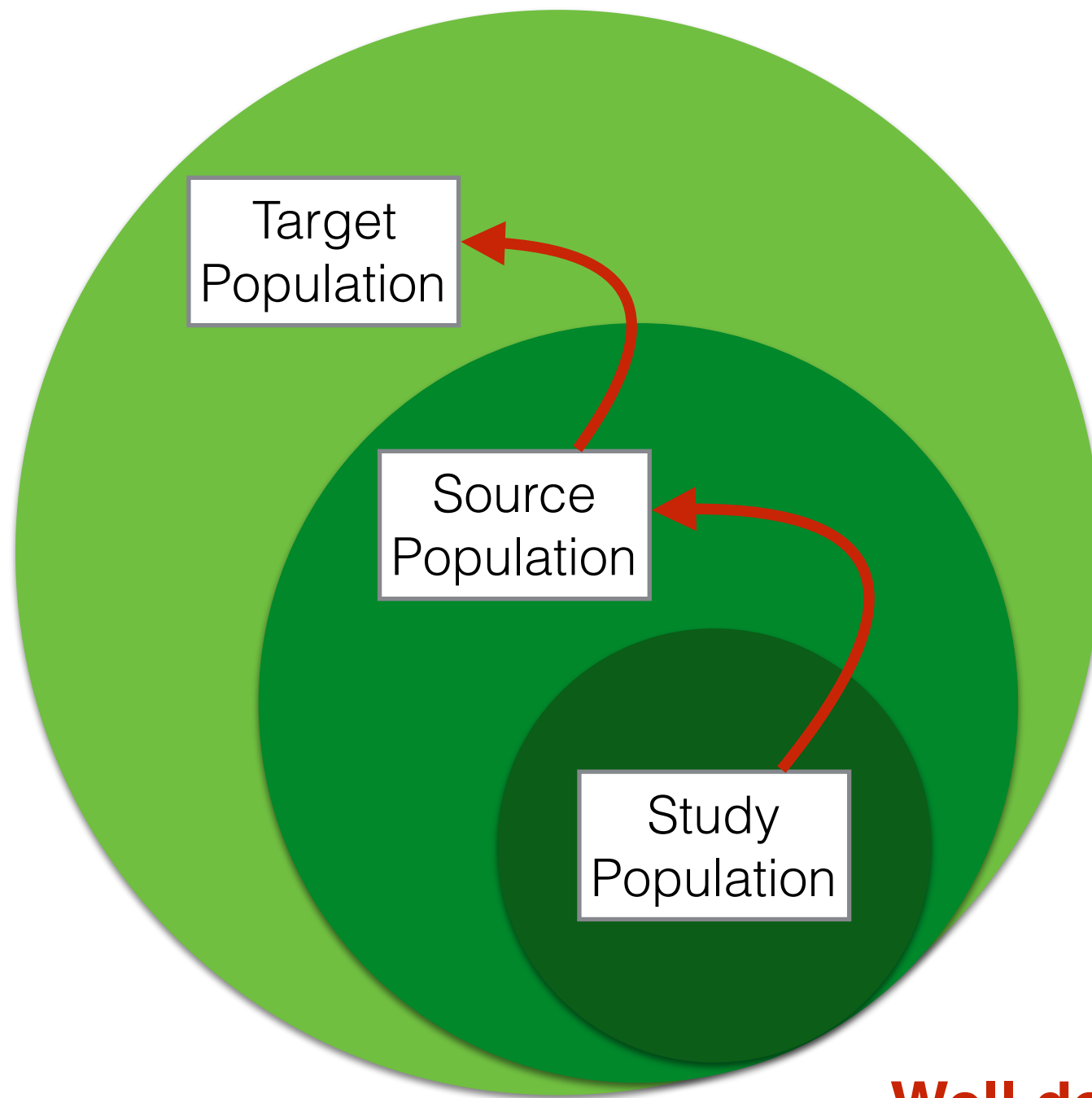


Target Population:
Want to make inferences about

Source Population:
Choosing study population

Study Population:
Enrolled in study





Target Population:
Want to make inferences about

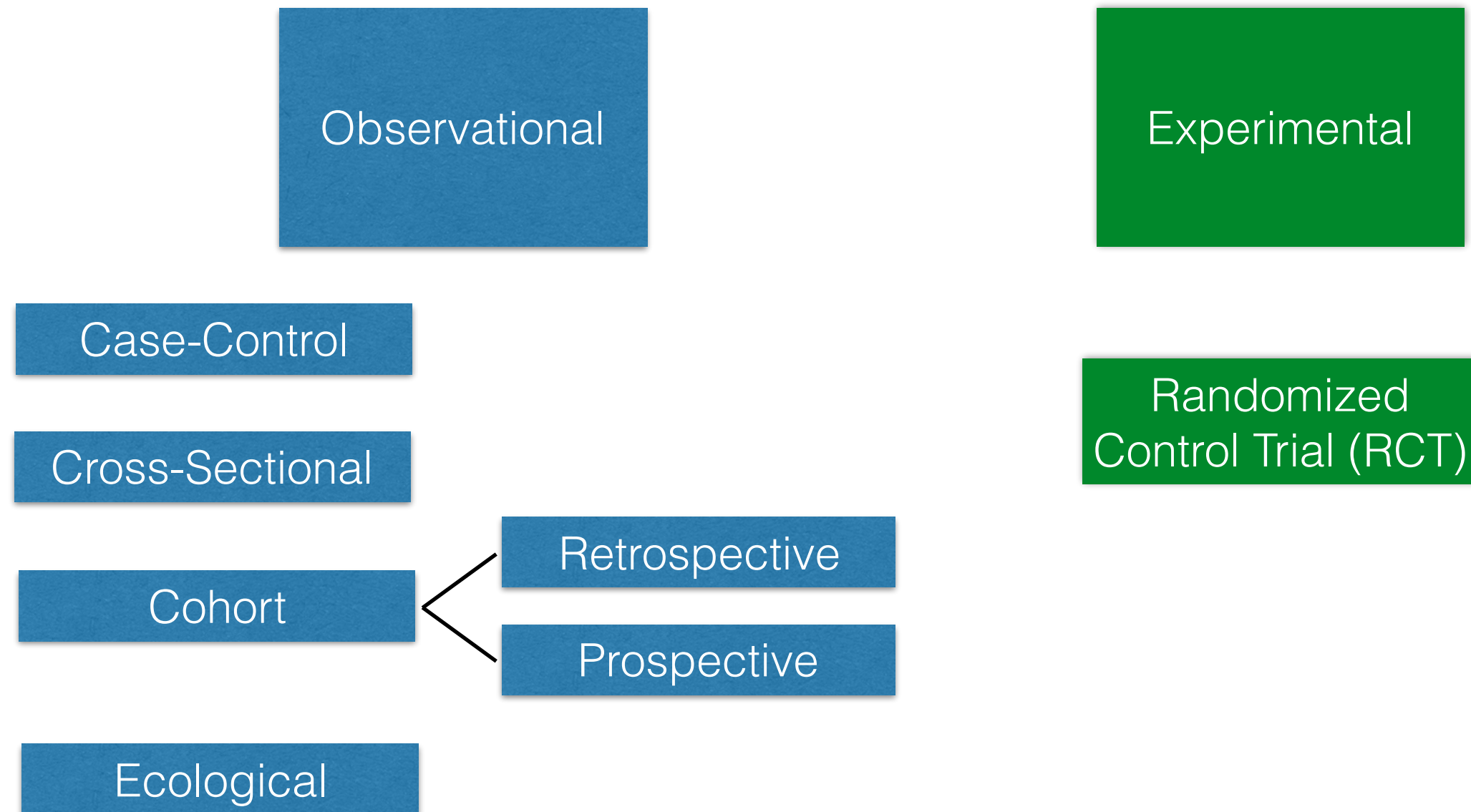
Source Population:
Choosing study population

Study Population:
Enrolled in study

Well designed studies allow us to make inference about the target population



Overview of different study designs



The Basics

Cross-
Sectional

Case-
Control

Cohort

RCT



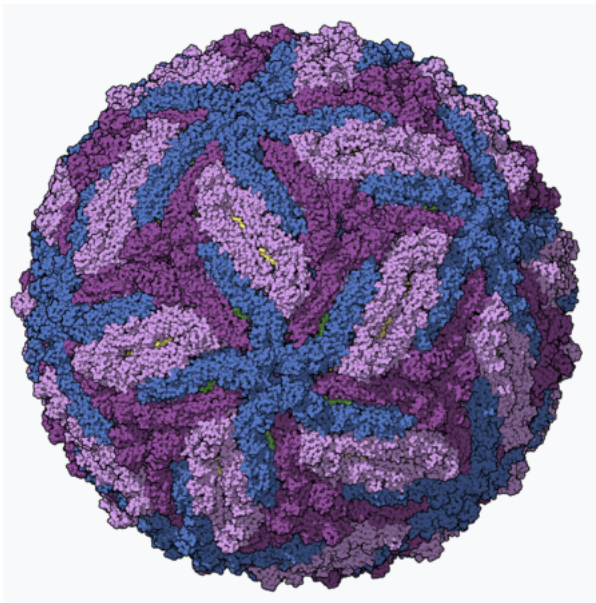
Zika Virus: The Basics

Cross-
Sectional

Case-
Control

Cohort

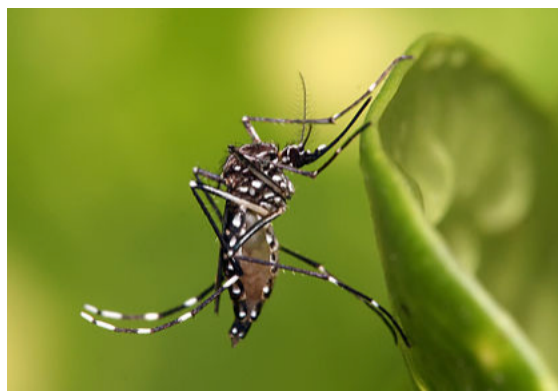
RCT



Virus in the family *Flaviviridae* (related to dengue, yellow fever)

Spread by *Aedes* mosquitoes

Multiple routes of transmission: vector, sexual, vertical



Zika Virus: The Basics

Cross-
Sectional

Case-
Control

Cohort

RCT



Often causes no or mild symptoms
But can spread from pregnant women to
their fetuses and result in microcephaly,
severe brain malformations, other birth
defects

Currently cannot be prevented by
medications or vaccines (current vaccine
trials)



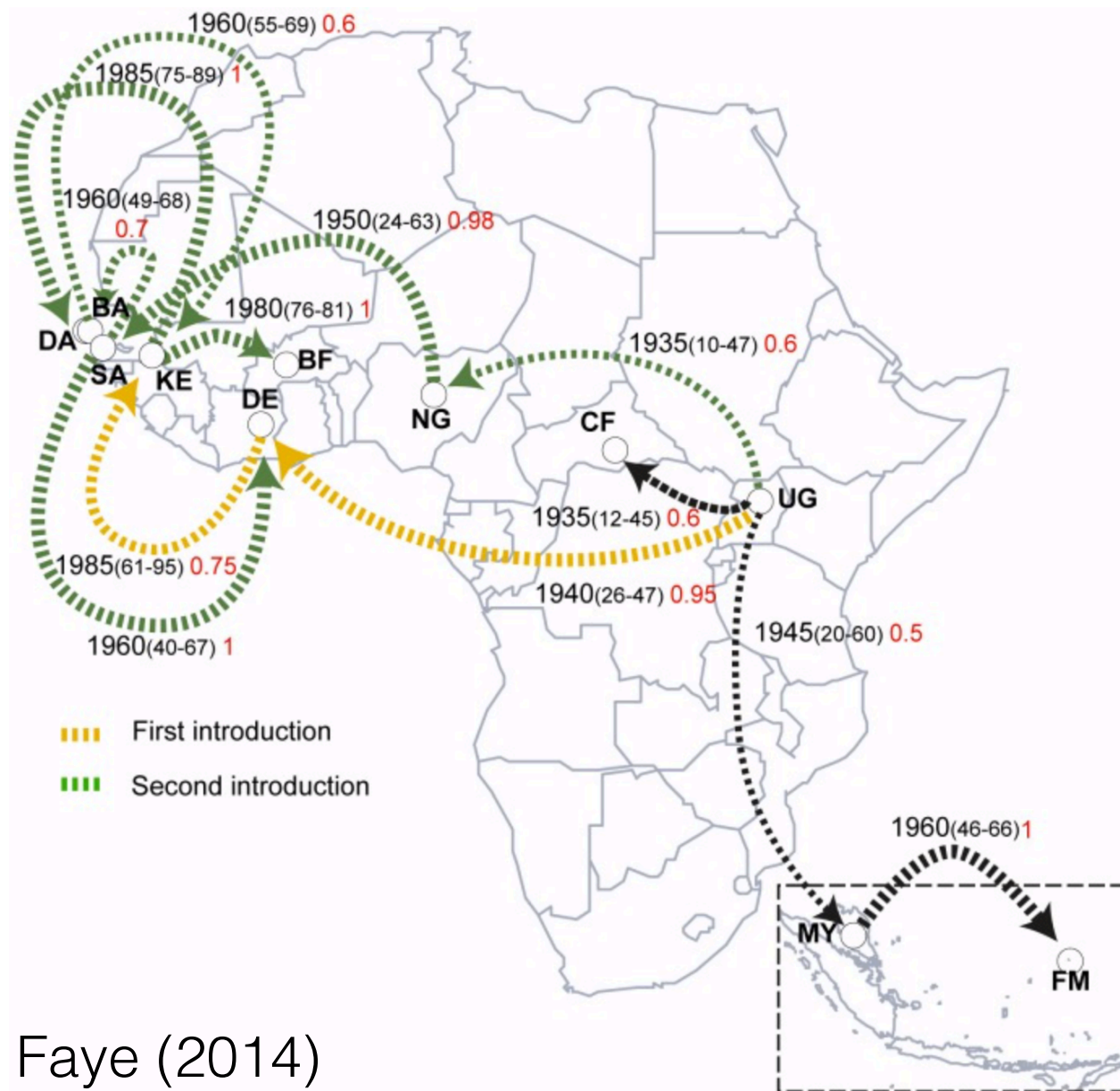
Zika Virus: The Basics

Cross-
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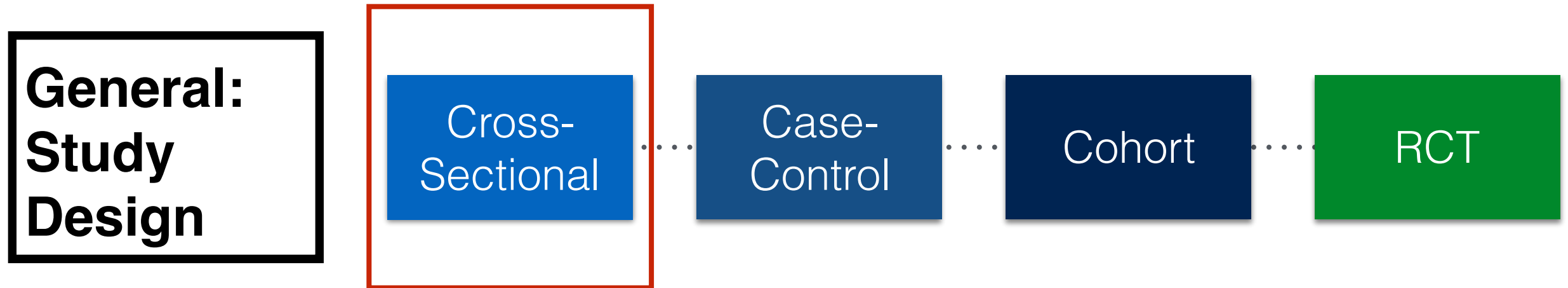


Faye (2014)

Originally isolated in 1947 (Ziika Forest, Uganda)

Sporadic outbreaks in Africa and Asia
Large outbreak in 2015-2016 (Americas, SE Asia, Pacific Islands, Brazil)





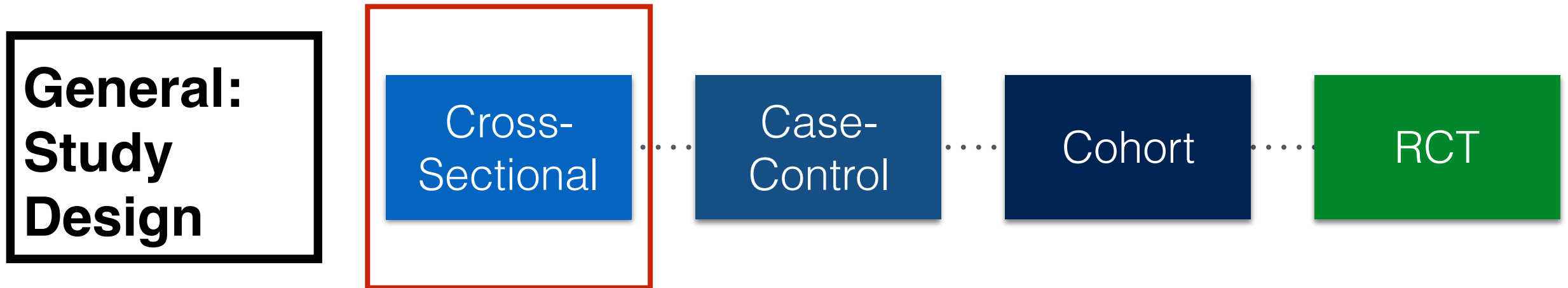
Cross Sectional Study

- Examines the relationship between **diseases** (or other health outcomes) and **other variables of interest** as they exist in a **defined population at one particular point in time**.

Examine la relation entre les maladies (ou d'autres résultats pour la santé) et d'autres variables d'intérêt, telles qu'elles existent dans une population définie à un moment donné.

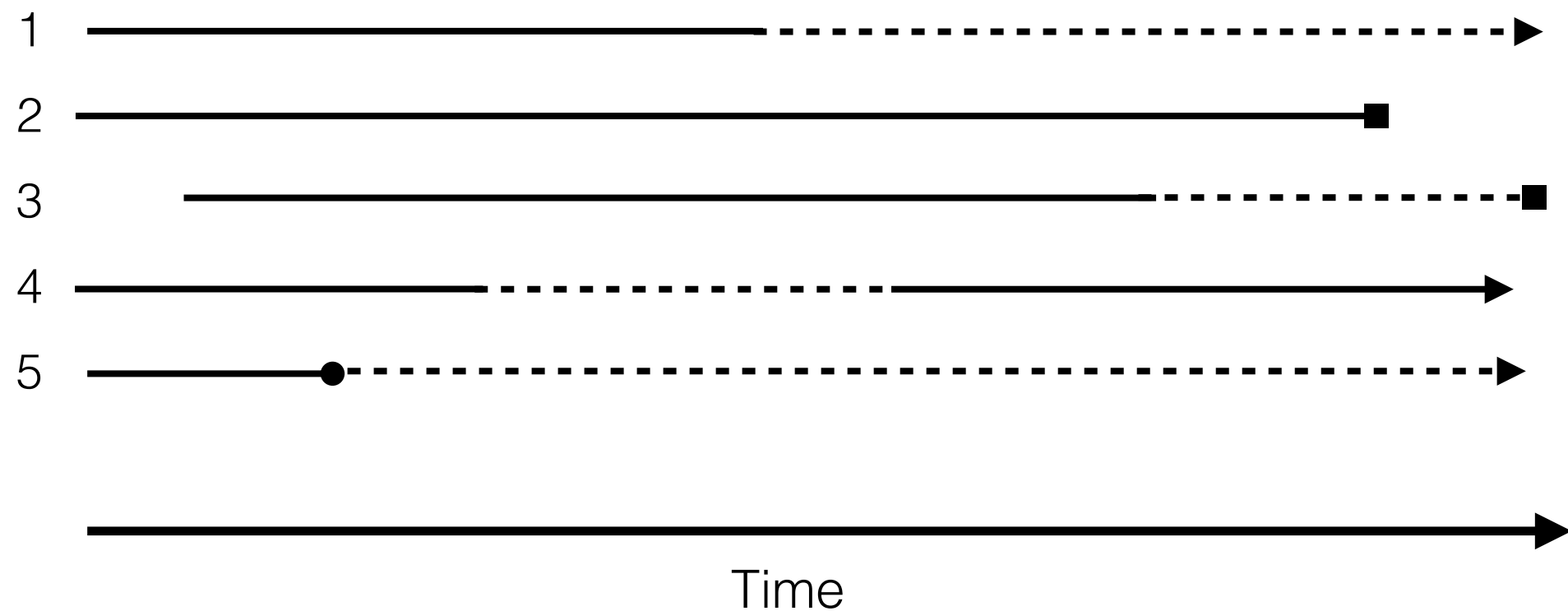
- Presence or absence of the disease are determined for each member of the population.
- Commonly used to estimate disease prevalence rather than incidence.
- **Snapshot of the population at a particular time.**
Instantané de la population à un moment donné.
- **Exposure and outcome are assessed simultaneously.**
L'exposition et les résultats sont évalués simultanément.

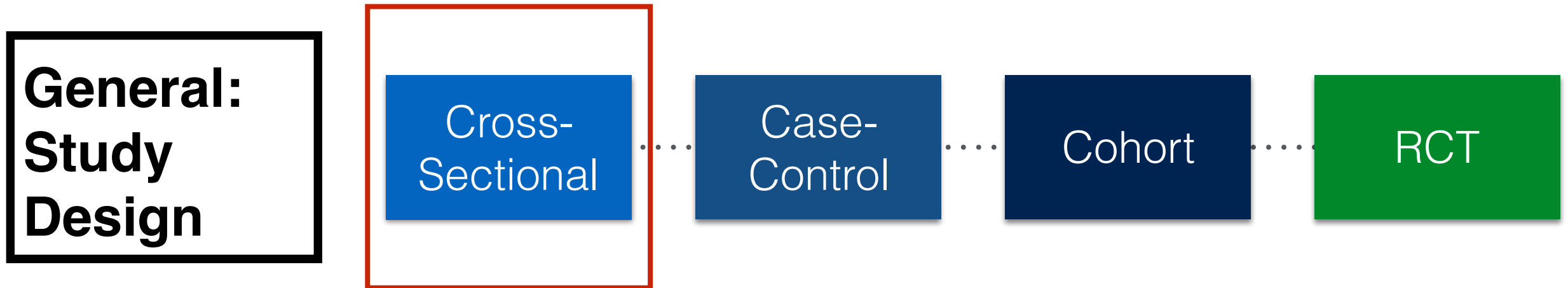




Cross Sectional Study

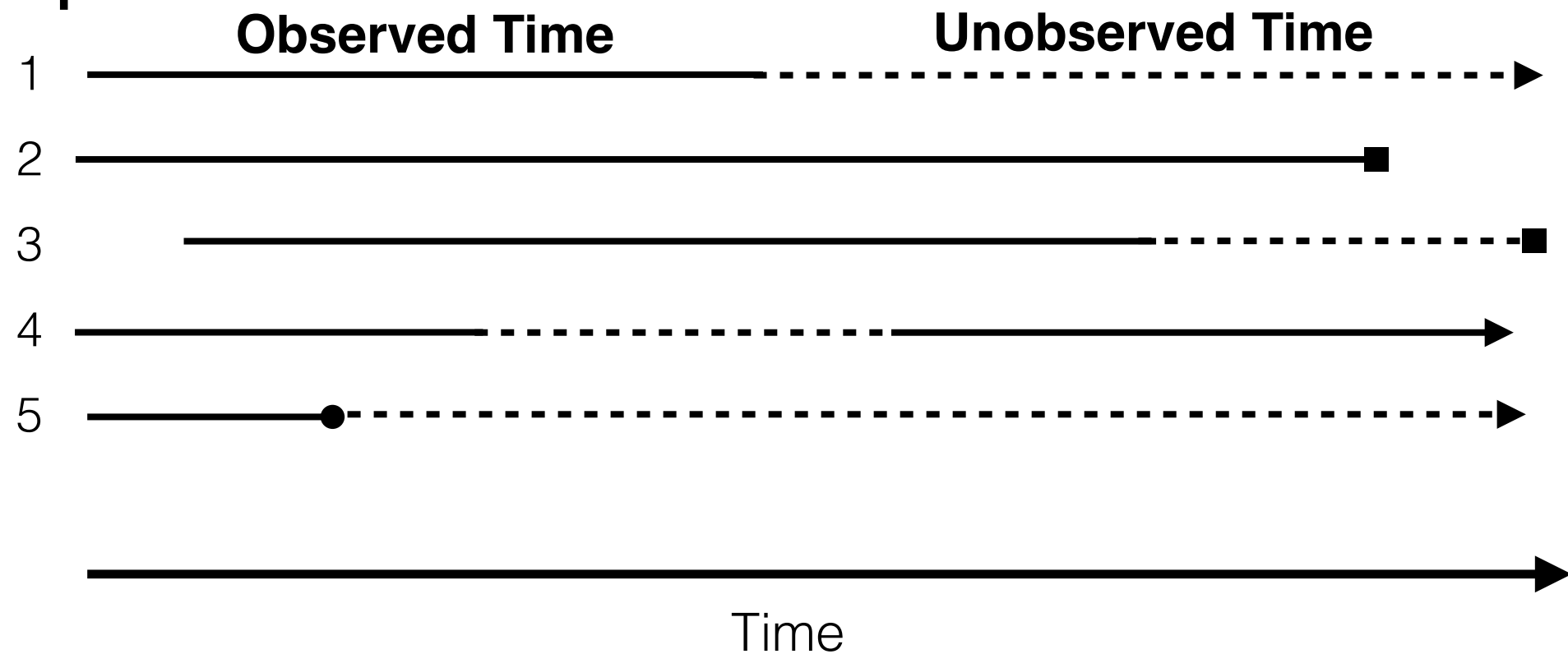
Participant

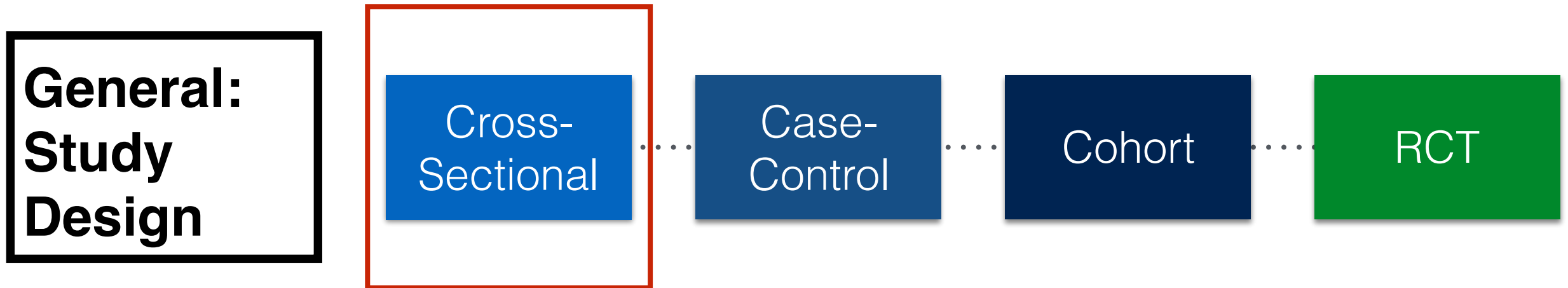




Cross Sectional Study

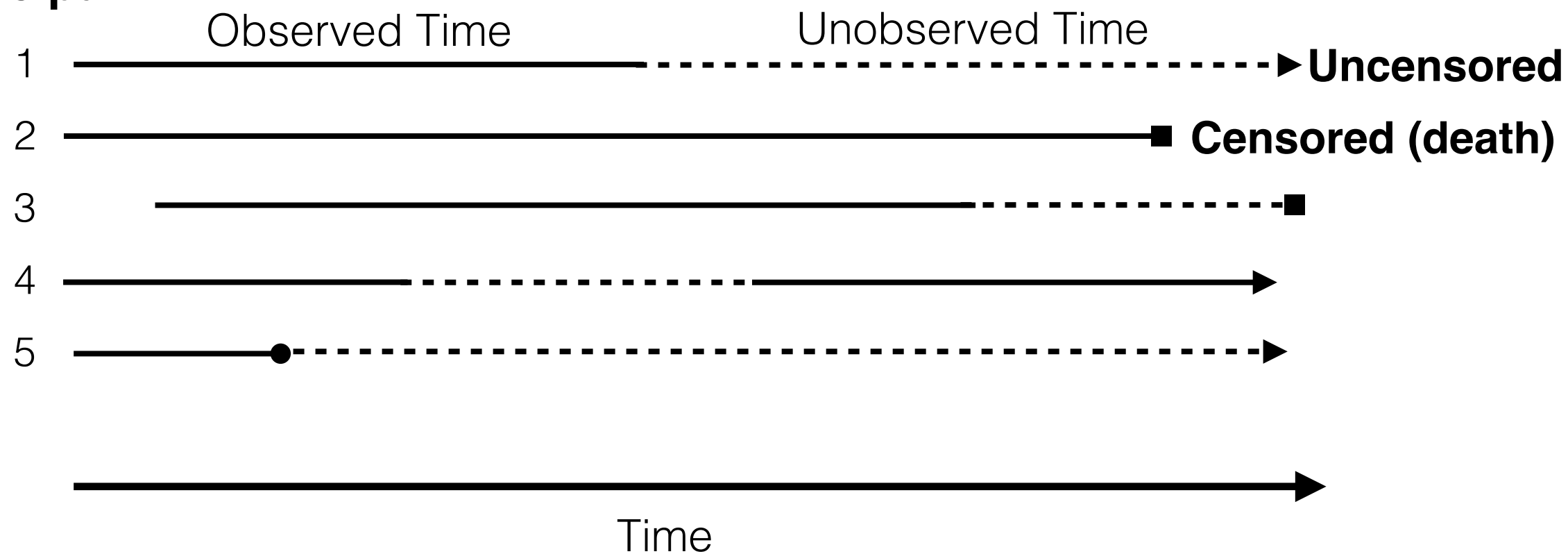
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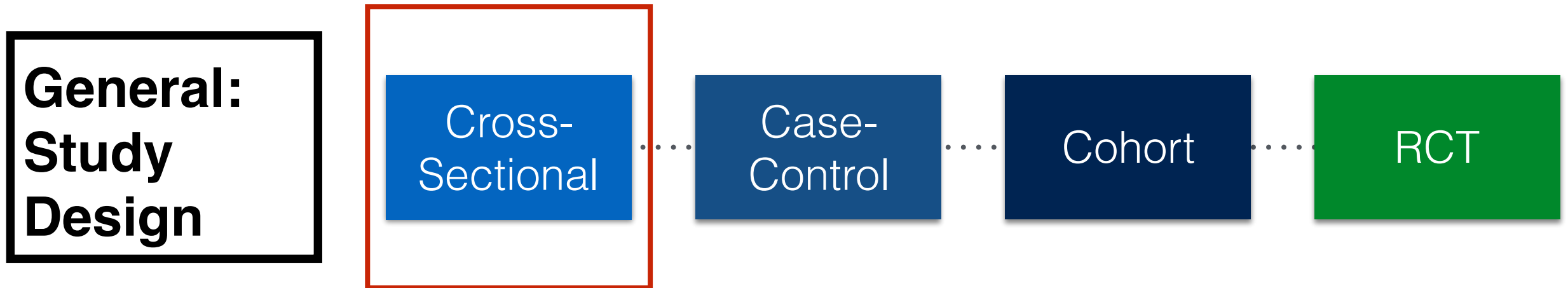




Cross Sectional Study

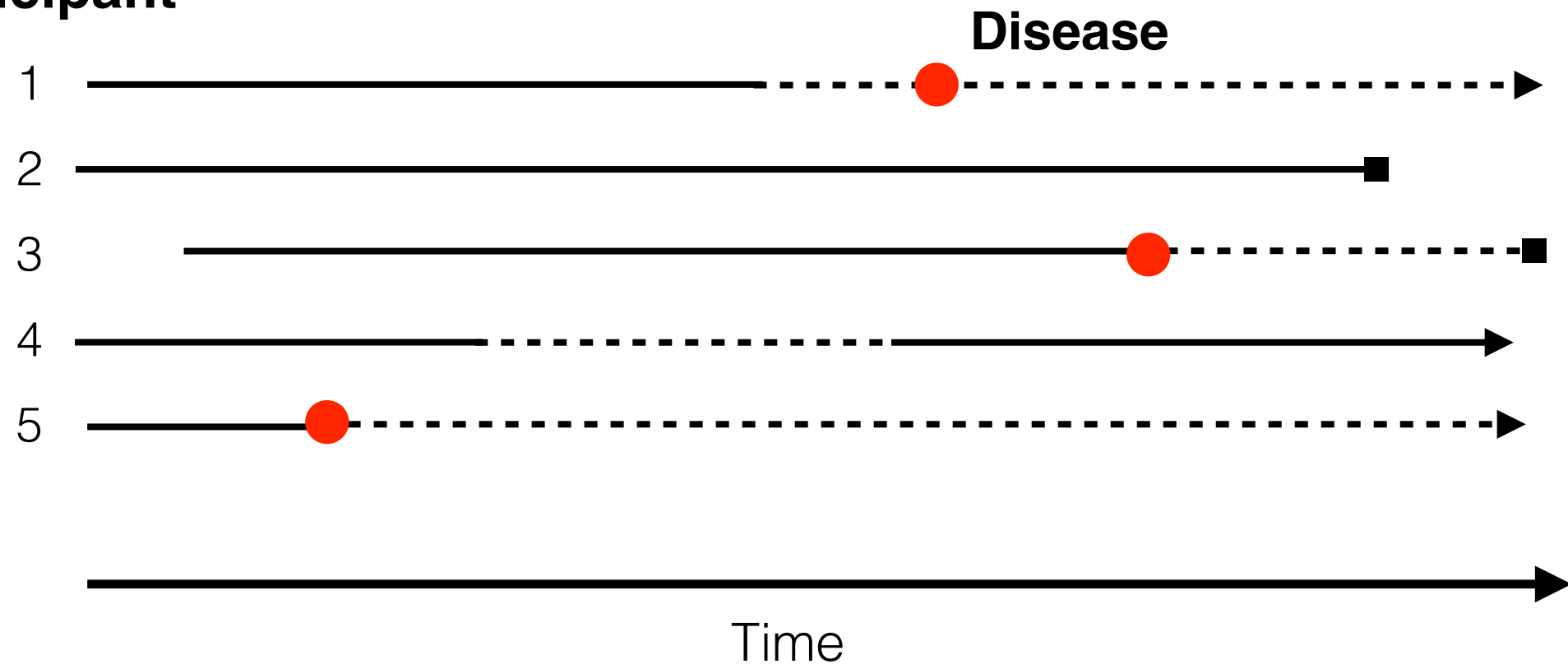
Participant

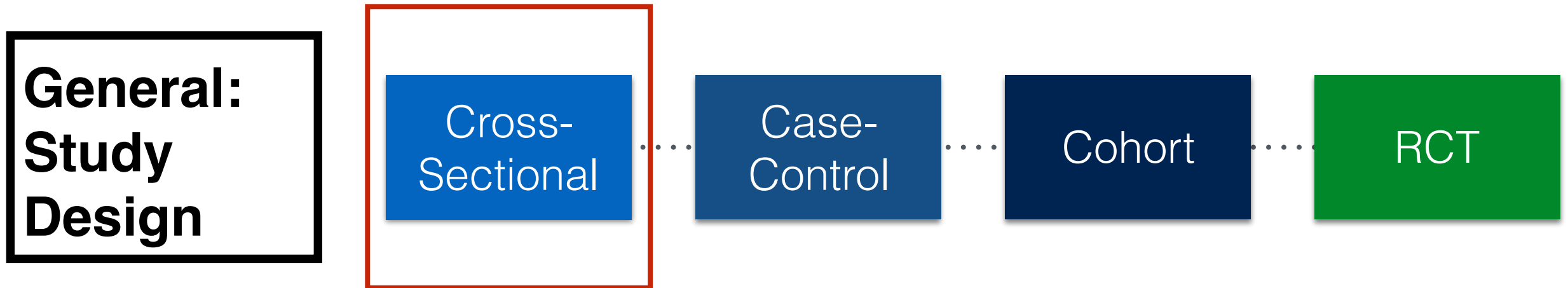




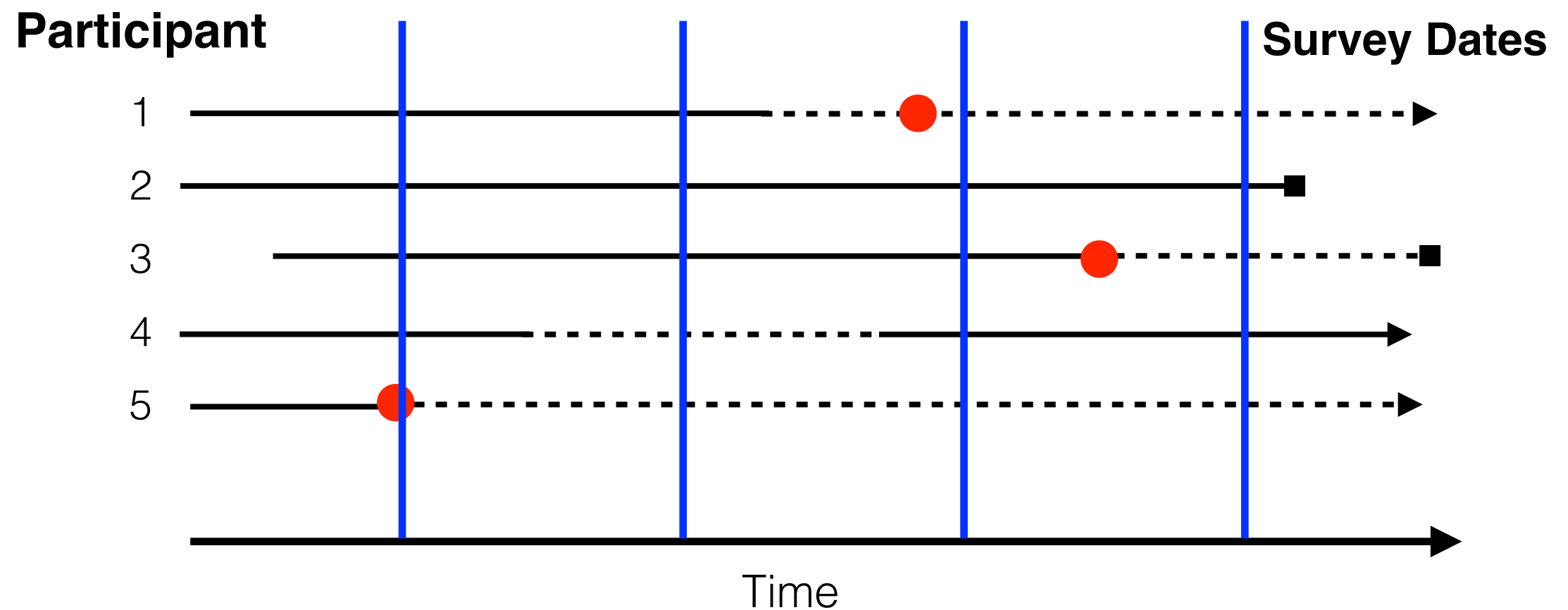
Cross Sectional Study

Participant





Cross Sectional Study



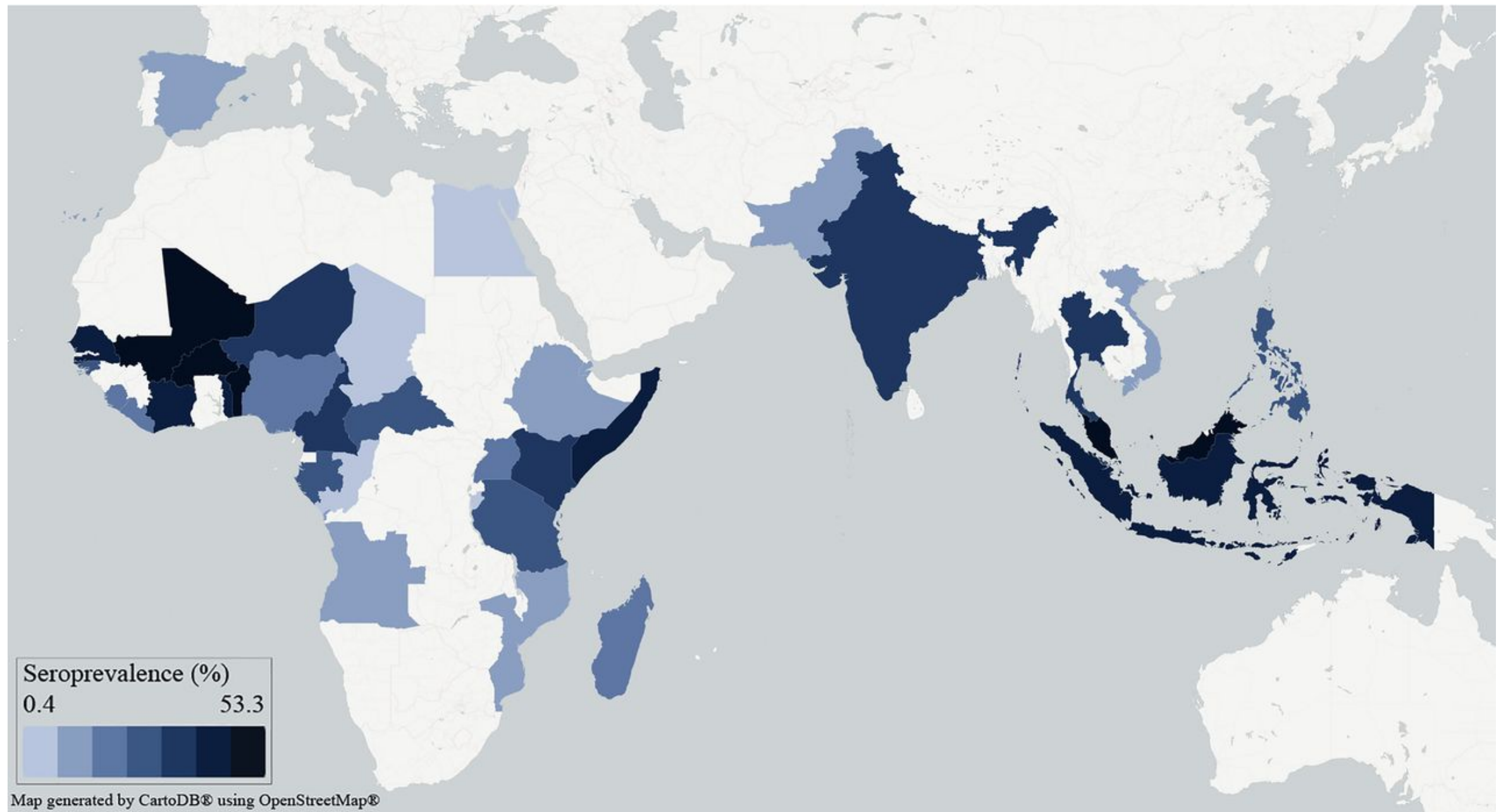
Zika: Study Design

Cross-
Sectional

Case-
Control

Cohort

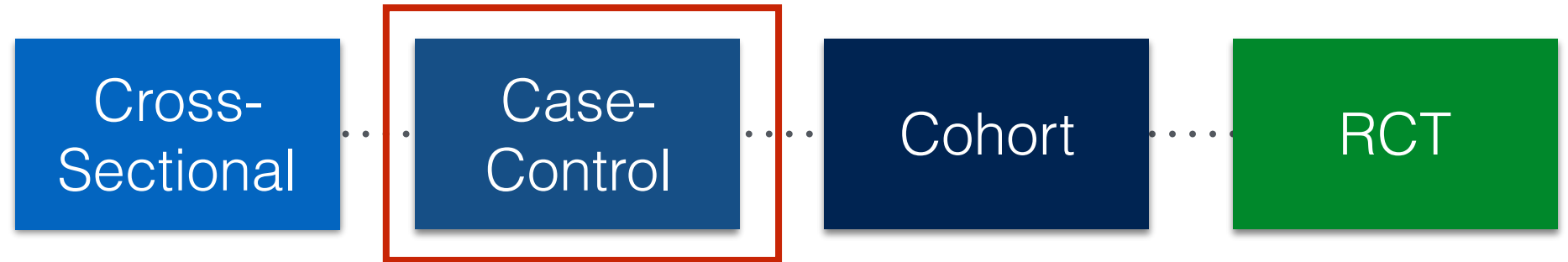
RCT



Posen (2016)



General: Study Design

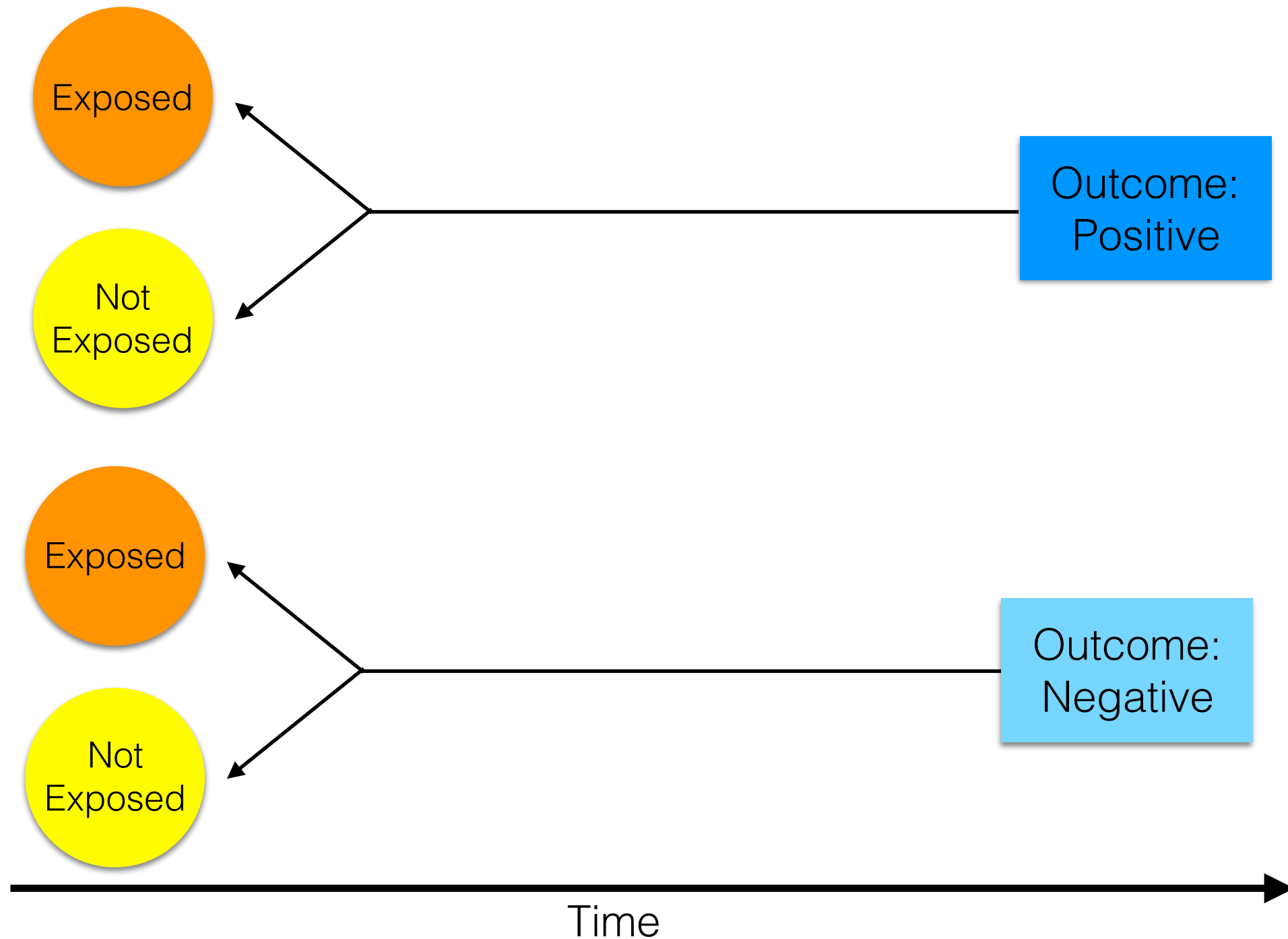
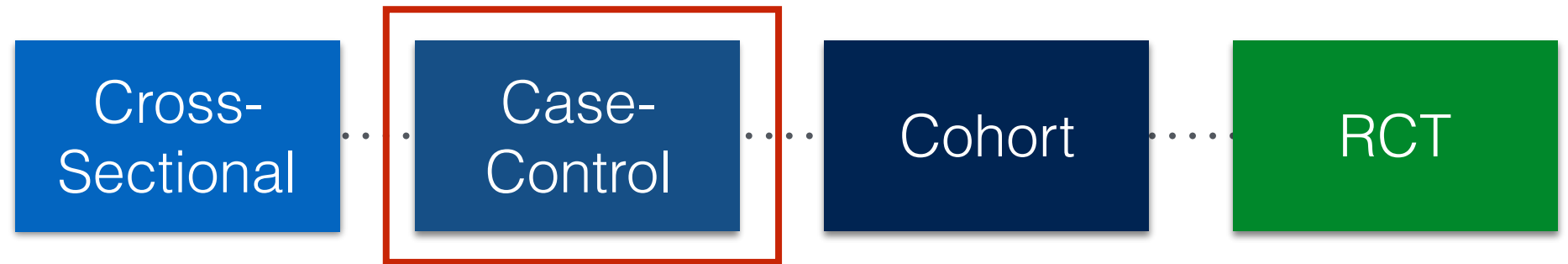


Case-Control Study

- The **observational** epidemiological study of **persons with a disease** (or another outcome variable) of interest and a suitable **control group of persons without the disease**
Étude épidémiologique d'observation sur des personnes présentant une maladie (ou une autre variable de résultat) d'intérêt et sur un groupe de contrôle approprié de personnes non atteintes de la maladie
- Potential relationship of a suspected risk factor or an attribute to the disease is examined by **comparing the the disease and non-diseased subjects** with regard to how frequently the factor or attribute is present in each of these groups.
La relation potentielle d'un facteur de risque présumé ou d'un attribut avec la maladie est examinée en comparant la maladie et les sujets non atteints à la fréquence à laquelle le facteur ou l'attribut est présent dans chacun de ces groupes.



General: Study Design



Zika: Study Design

Cross-
Sectional

Case-
Control

Cohort

RCT

Exposed

Not
Exposed

Exposed

Not
Exposed

Outcome:
Positive

What is the relationship between
microcephaly and Zika?

Outcome:
Negative

Time



Zika: Study Design

Cross-
Sectional

Case-
Control

Cohort

RCT

Maternal
Zika Pos

Maternal
Zika Neg

Child born with
microcephaly

Maternal
Zika Pos

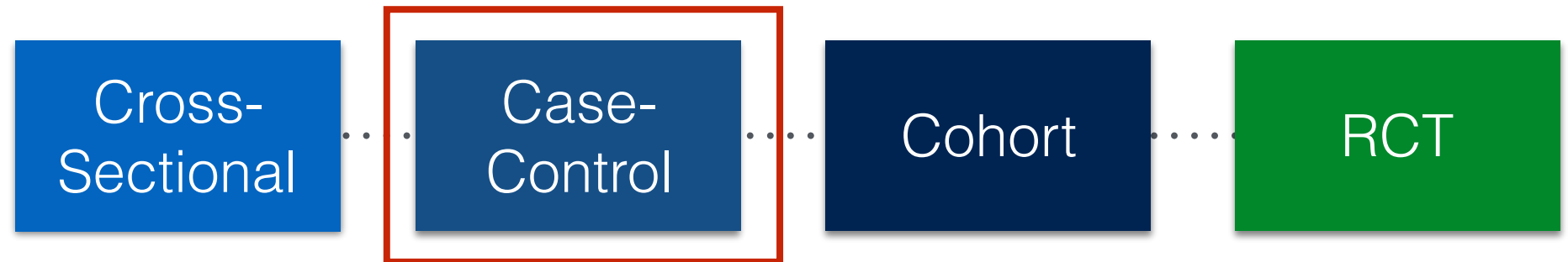
Maternal
Zika Neg

Child born
without
microcephaly

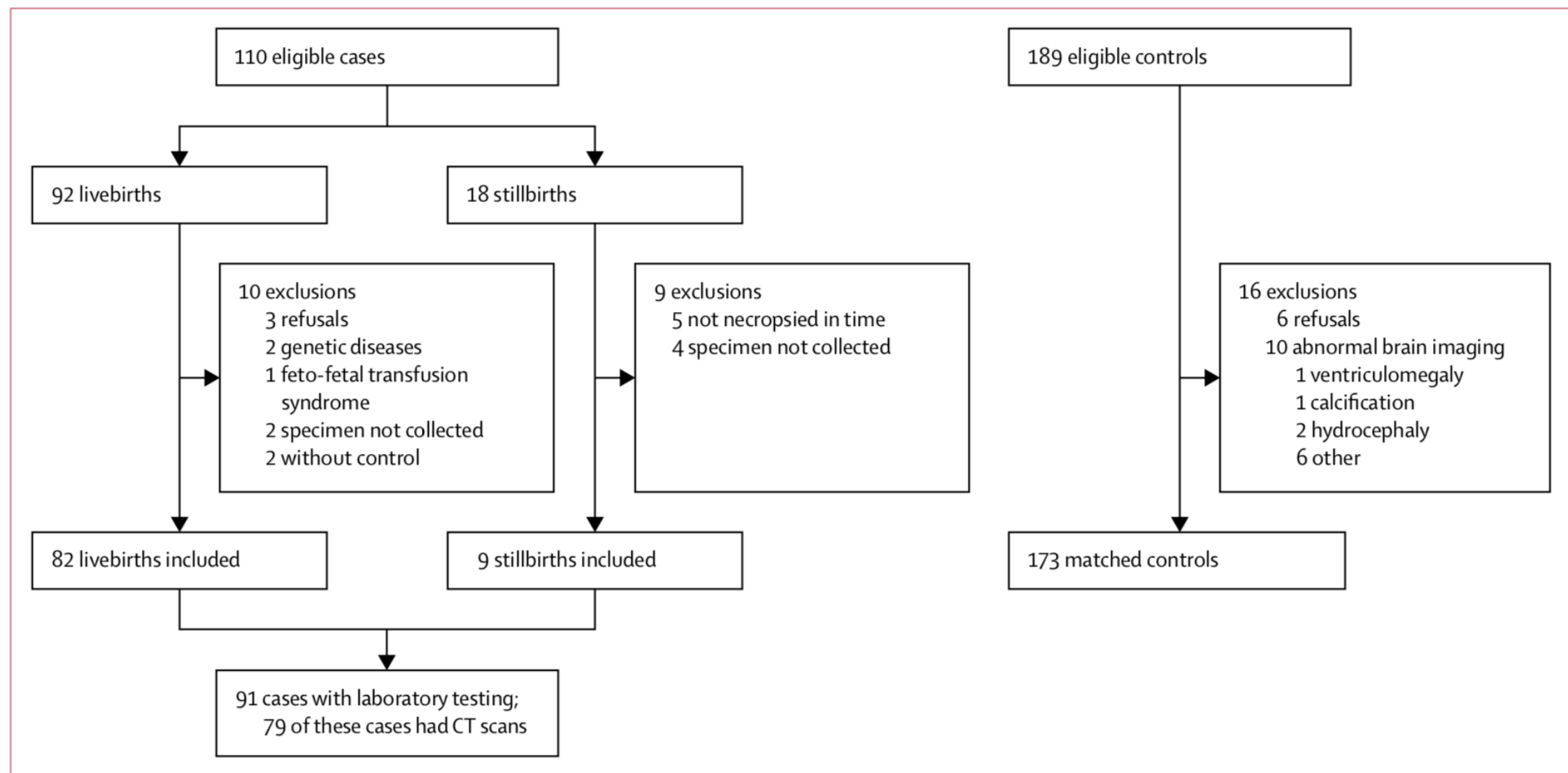
Time



Zika: Study Design



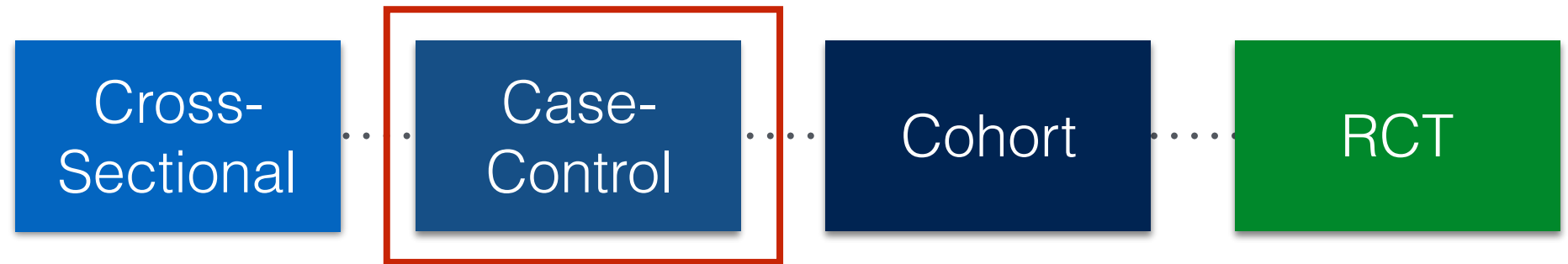
Case-Control study - 8 Brazilian hospitals



de Araujo (2018)



Zika: Study Design



Case-Control study - 8 Brazilian hospitals

	Cases*	Controls*	Matched odds ratio (95% CI)
Serum, CSF samples, or macerated tissue			
Zika-positive, of total cases or controls	32/91 (35%)	0/173	87.0 (15.6–∞)
Zika-positive, of total cases or controls, adjusted†	73.1 (13.0–∞)
Cases, categorised by severity of microcephaly‡			
Severe	19/26 (73%)	0/51	52.4 (9.1–∞)
Not severe	13/65 (20%)	0/122	33.7 (5.6–∞)

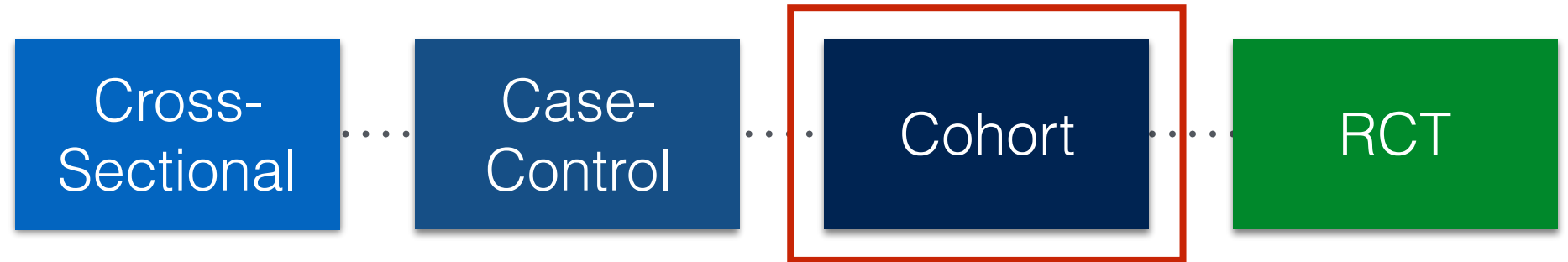
*Data are the number of all cases or controls who were positive for Zika virus, assessed by qRT-PCR or Zika virus-specific IgM/total number of patients (%). †Odds ratio when adjusted by smoking during pregnancy, maternal vaccination against tetanus, diphtheria, and acellular pertussis during pregnancy, and skin colour. ‡Severe is defined as a head circumference of more than 3 SD smaller than the mean for their sex and gestational age.^{10,14} Not severe was defined as a head circumference of 2–3 SD smaller than the mean for their sex and gestational age. Matched odds ratios in this subgroup are crude because of small numbers.

Table 5: Association between microcephaly and Zika virus infection

de Araujo (2018)



General: Study Design



Cohort Study

- **Enrolled group of people** who have a common experience or grouping.
Groupe de personnes inscrites ayant une expérience commune ou un groupe.
 - Age cohort, risk cohort
- General population sample
- Clinic based
- **Prospective or retrospective**
Prospective ou rétrospective



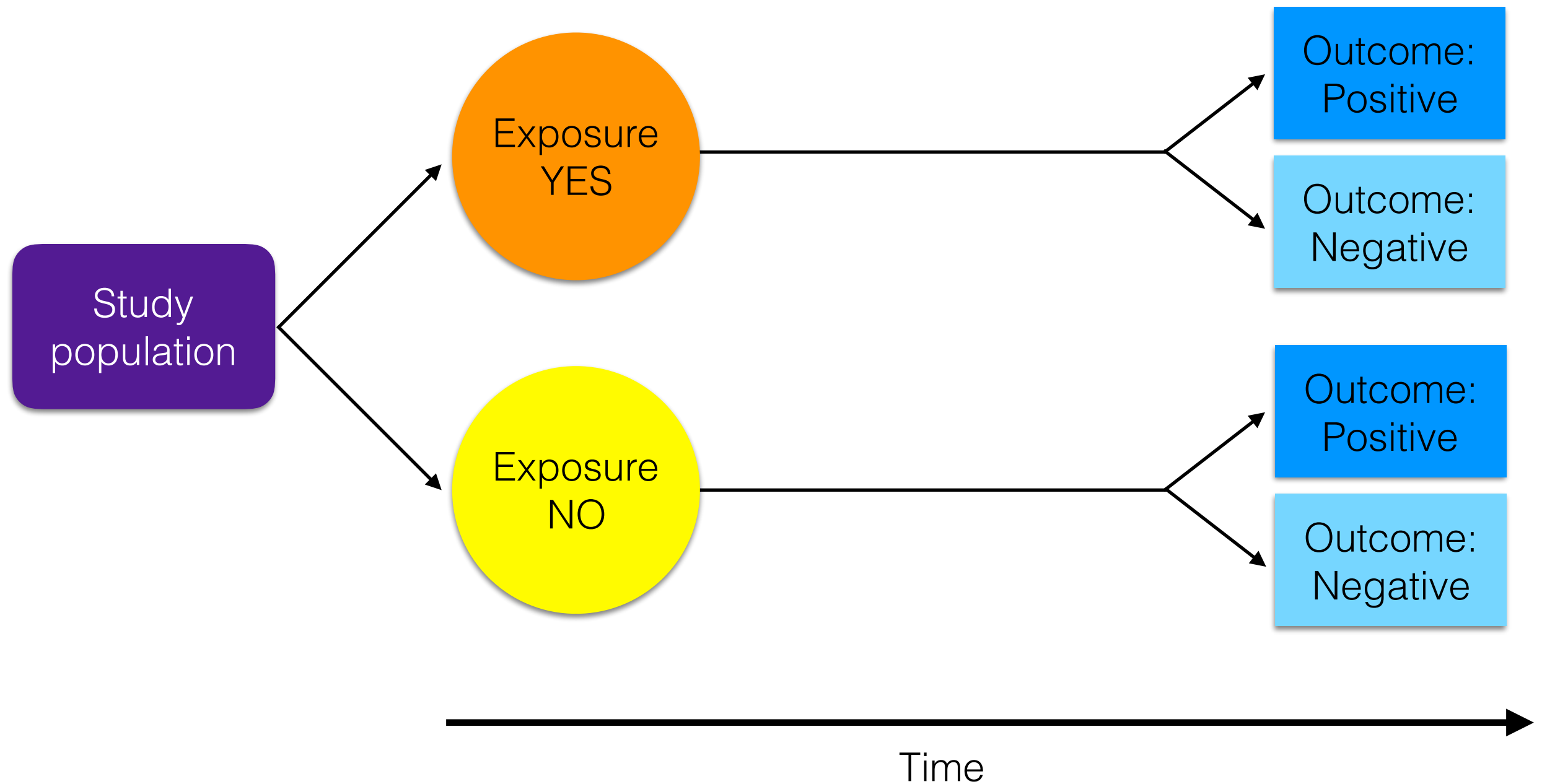
General: Study Design

Cross-
Sectional

Case-
Control

Cohort

RCT



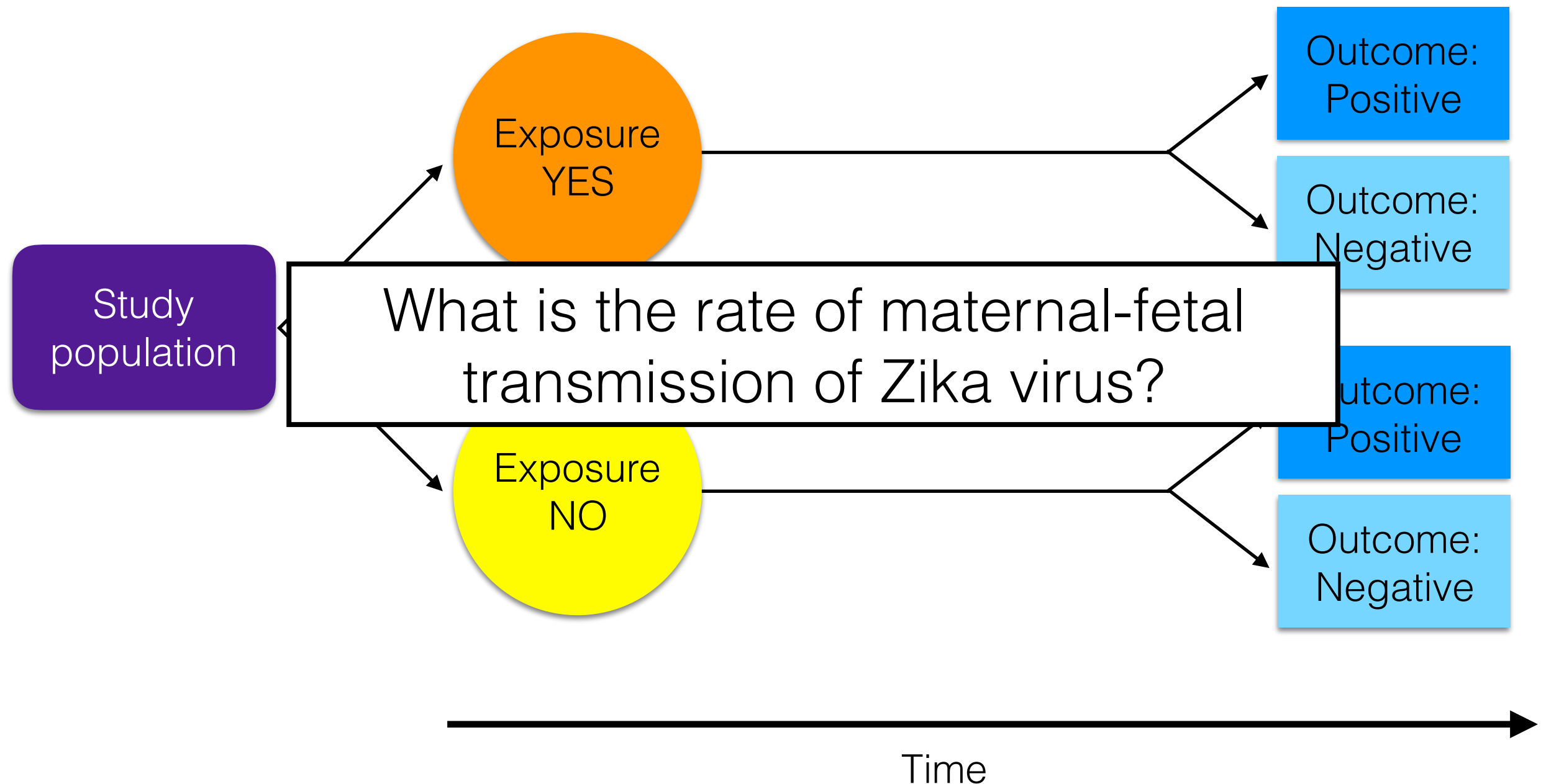
Zika: Study Design

Cross-
Sectional

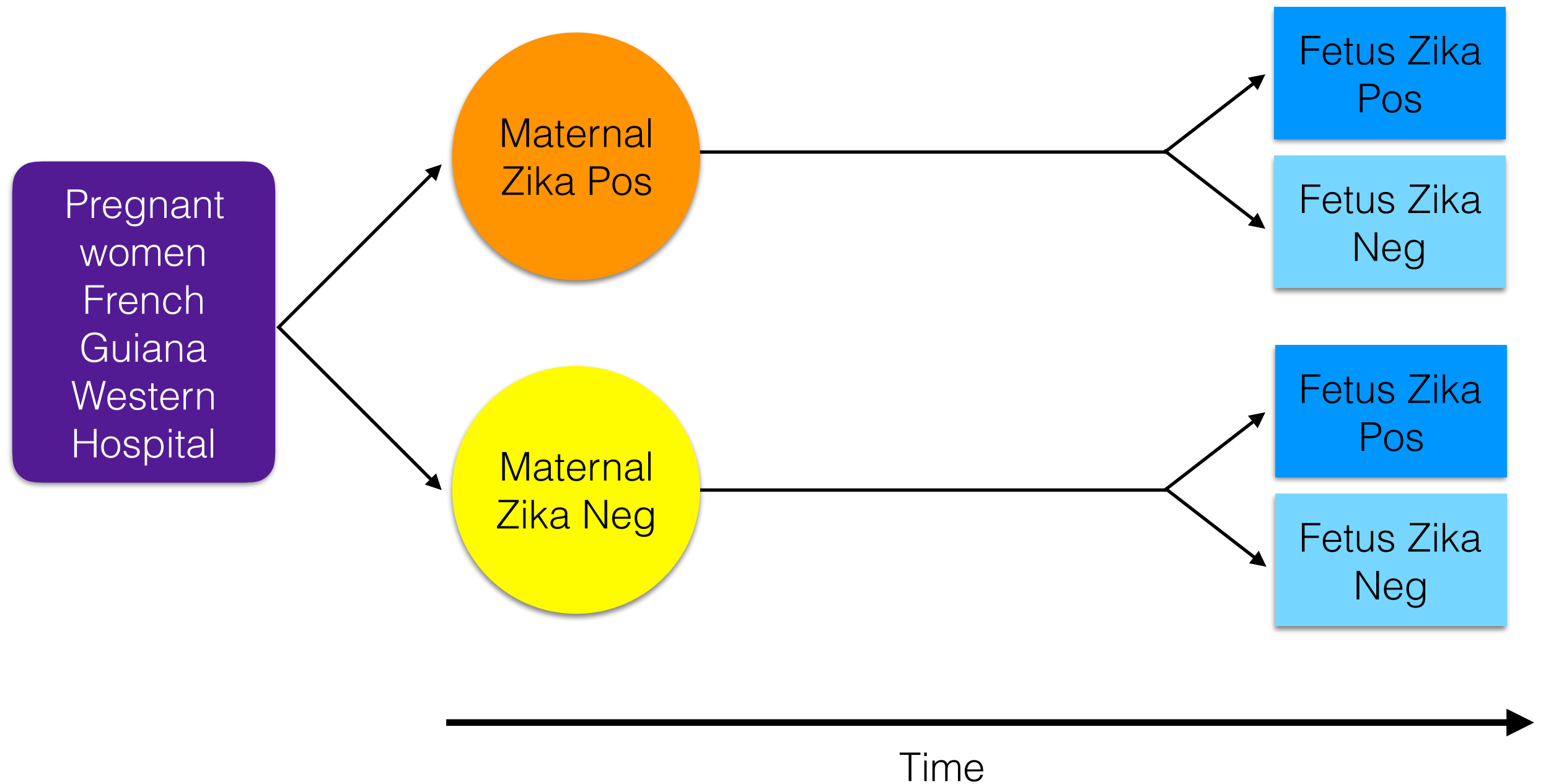
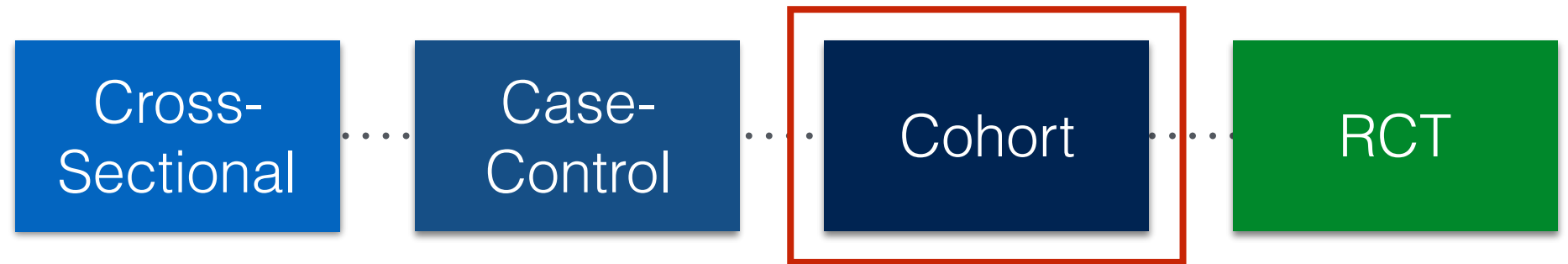
Case-
Control

Cohort

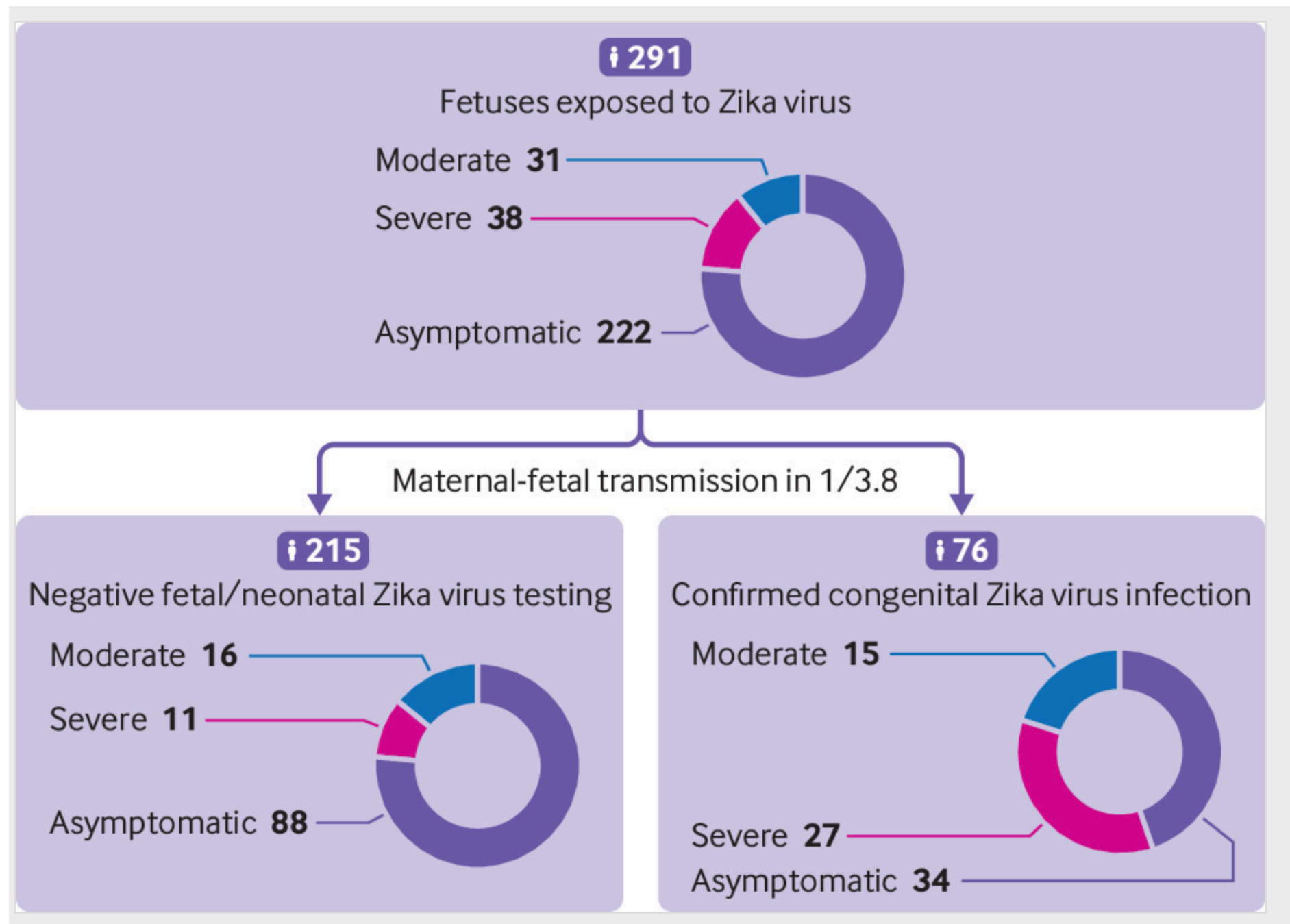
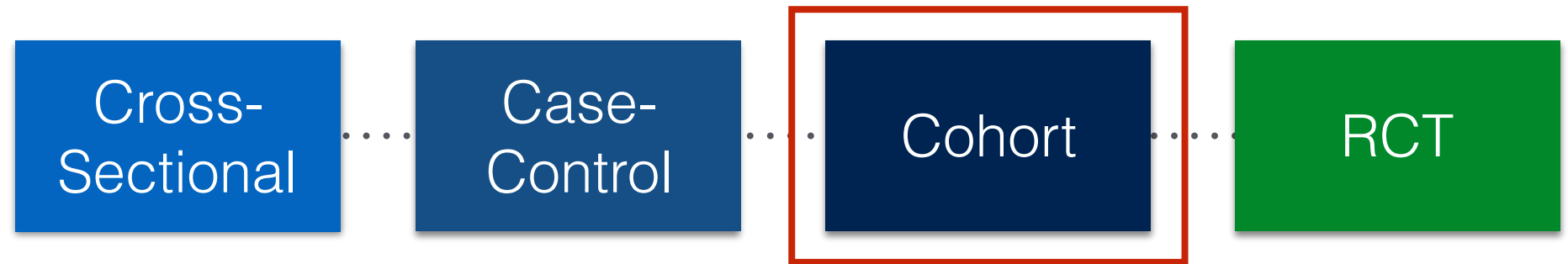
RCT



Zika: Study Design



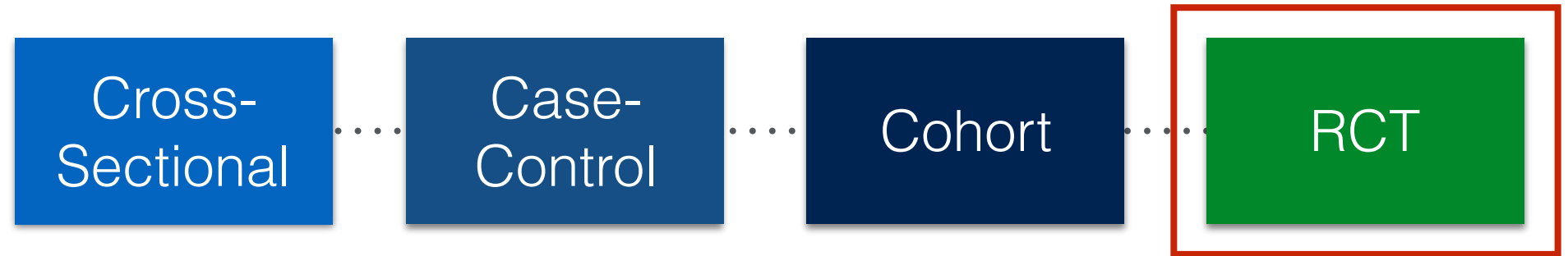
Zika: Study Design



Pomar (2018)



General: Study Design

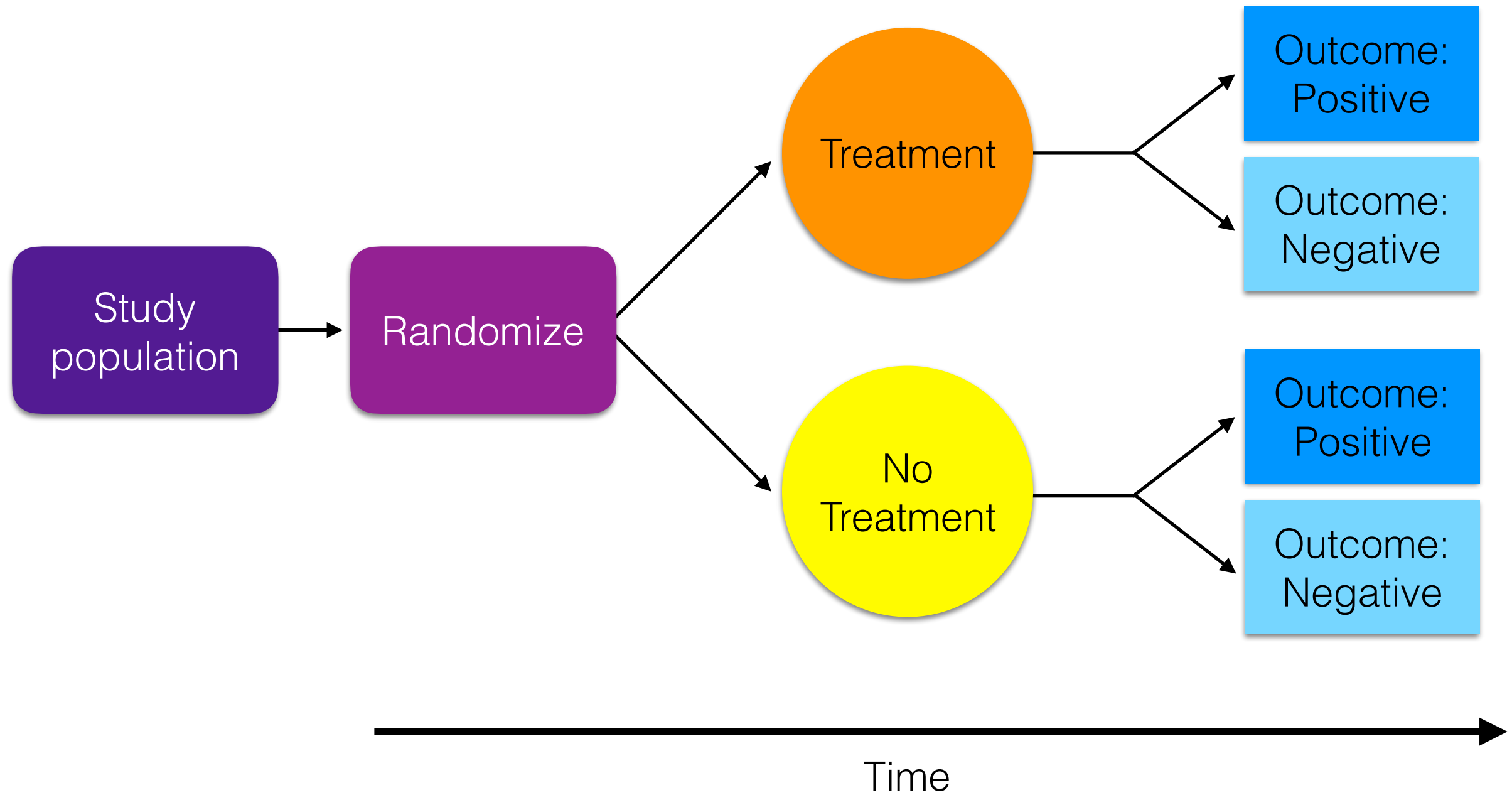
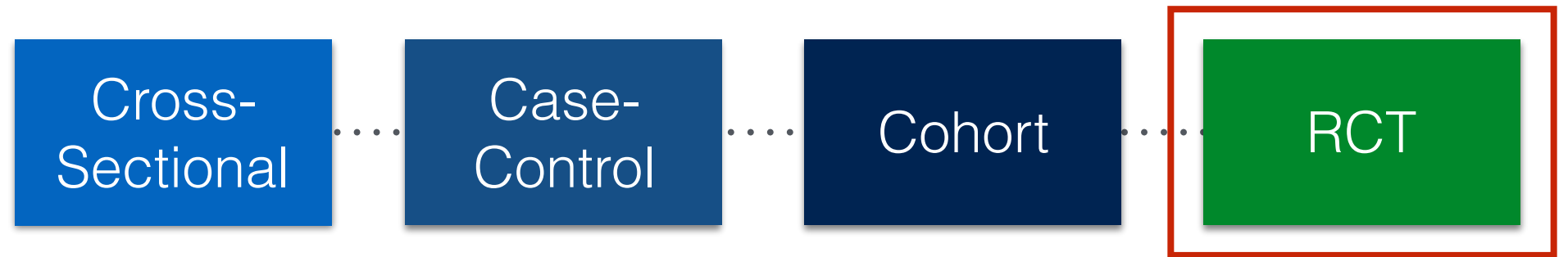


Randomized Control Study

- Experiment in which subjects are **randomly allocated into groups** (test and control that are comparable) to receive or not to receive a preventative or a therapeutic procedure or intervention.
Expérience dans laquelle les sujets sont répartis au hasard dans des groupes (test et contrôle comparables) pour recevoir ou non une procédure ou une intervention préventive ou thérapeutique.
- Results are assessed by comparison of rates of disease, death, recovery, or other outcome in the study groups.
- Generally thought of as the **most rigorous method** of hypothesis testing.
méthode la plus rigoureuse
- **Randomization should be blinded!**
La randomisation devrait être aveuglée!



General: Study Design



Zika: Study Design

Cross-
Sectional

Case-
Control

Cohort

RCT

Treatment

Outcome:
Positive

Outcome:
Negative

Study
population

What is the effectiveness of a Zika
Containing Vaccine?

No
Treatment

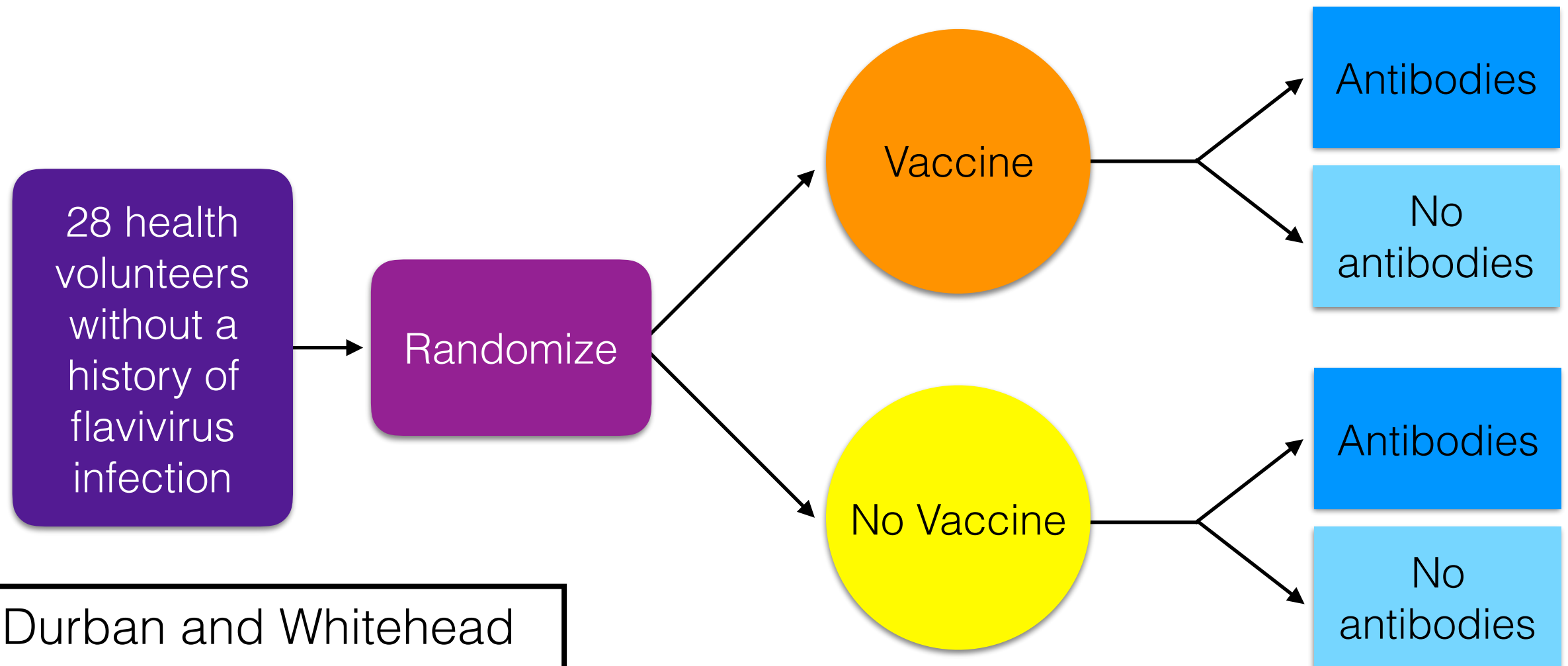
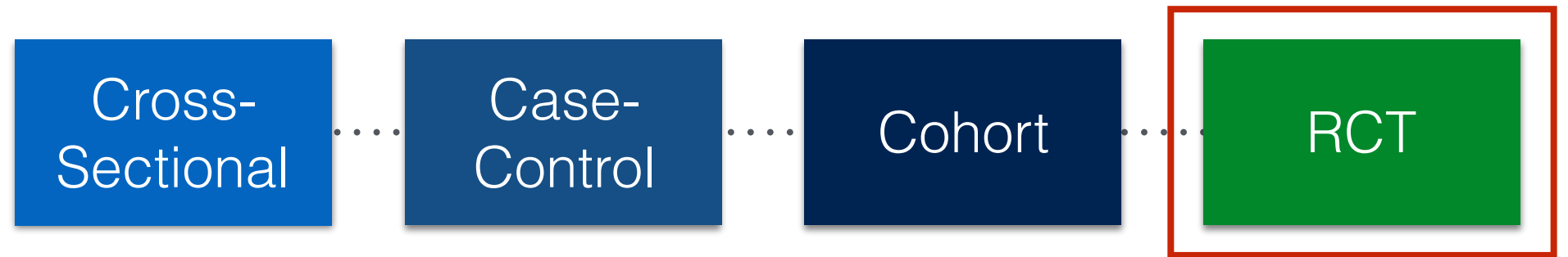
Outcome:
Positive

Outcome:
Negative

Time



Zika: Study Design



Durban and Whitehead
First in-human trial
live attenuated Zika
vaccine
rZIKV/D4Δ30-713

Time



How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)



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The MAHERY-CRS cross-sectional health survey (2017)

Objectives:

1. Perform a cross-sectional sample to determine the prevalence of several diseases and nutritional deficiencies. (*Aiza no tena misy tazomoka, tsy fahampian-dra, tsy fahampian-tsakafo, sns?*)
2. Survey rural communities (*tanàna kely*) in Madagascar: *Ambanivolobe*
3. Compare across different regions (*faritra*) of Madagascar:
 - Vatovavy-fitovinany, Amoron'i Mania, Atsimo Andrefana



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How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)



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How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)

OR

How to balance scientific and non-scientific factors
when designing a study?

How to not be paralyzed (*aza miasa saina*)?

How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)

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How does study design place limits on your analysis?

How to do good science with imperfect studies (*'Science' tsara raha tsy lavorary ny 'study'*)?

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The MAHERY-CRS cross-sectional health survey (2017)



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The MAHERY-CRS **cross-sectional** health survey (2017)



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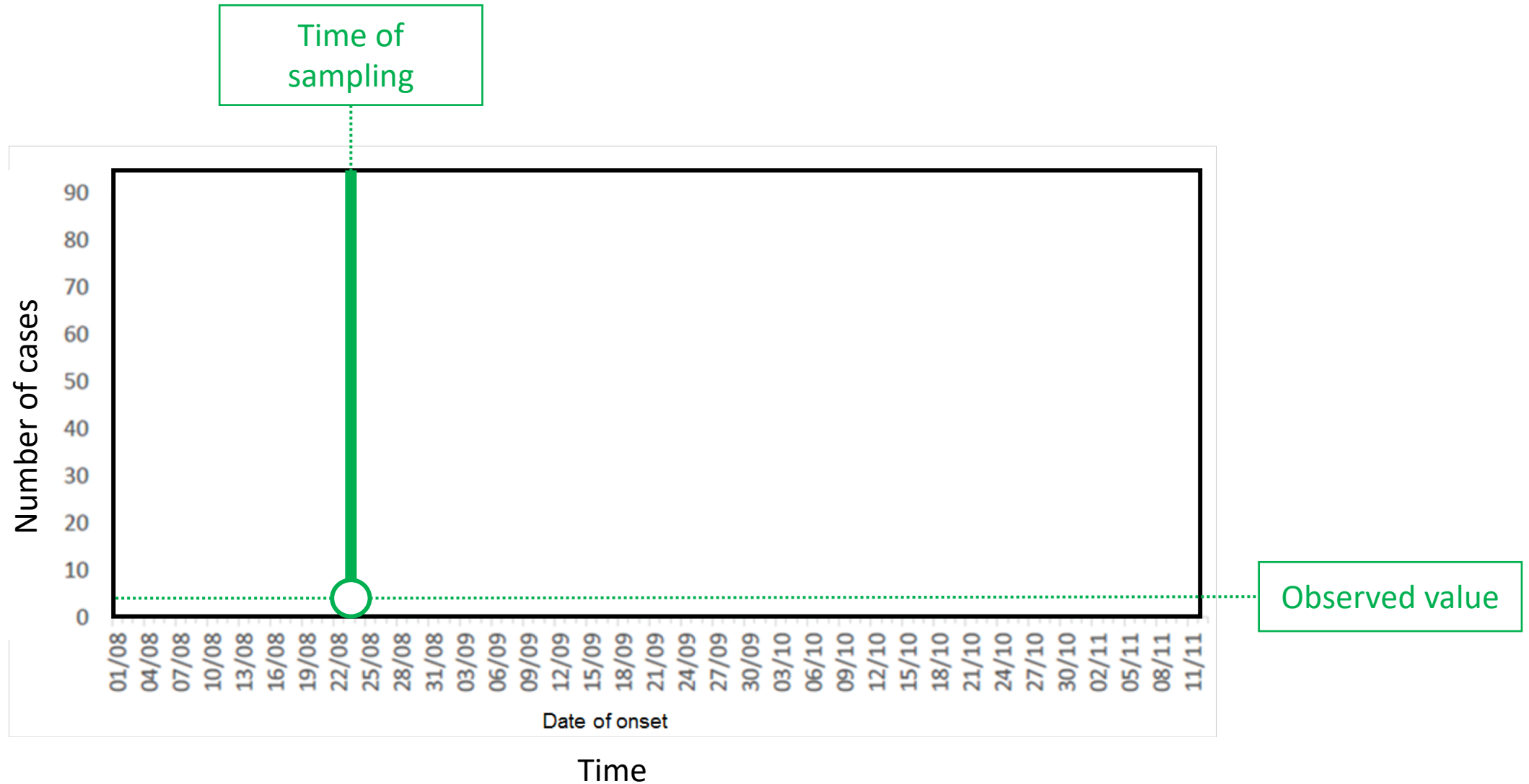
Limitations of cross-sectional surveys

Example: Plague outbreak in Madagascar

What is a major limitation of cross-sectional surveys?
Inona ny olana maveny misakana ny 'cross-sectional surveys'?

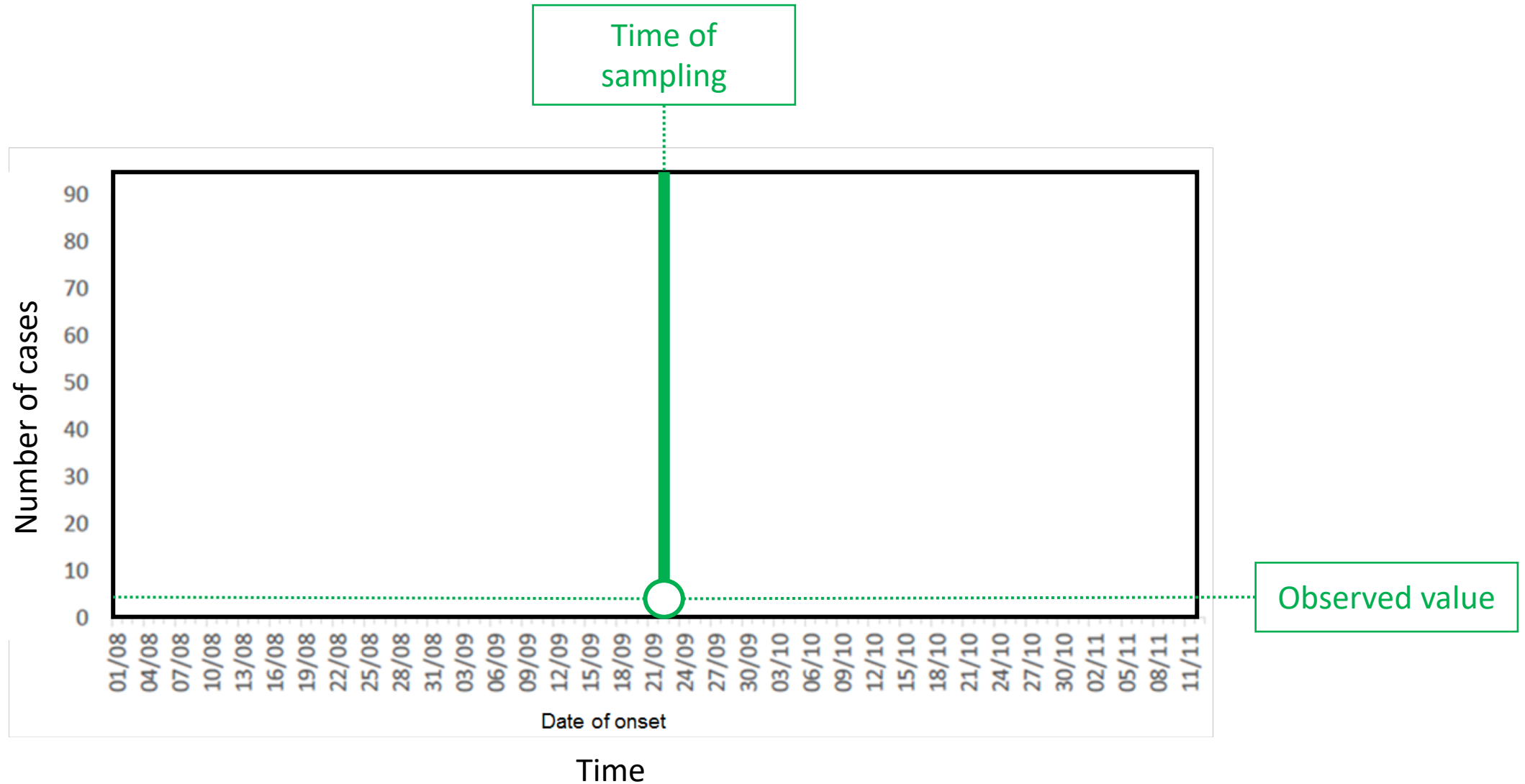
Limitations of cross-sectional surveys

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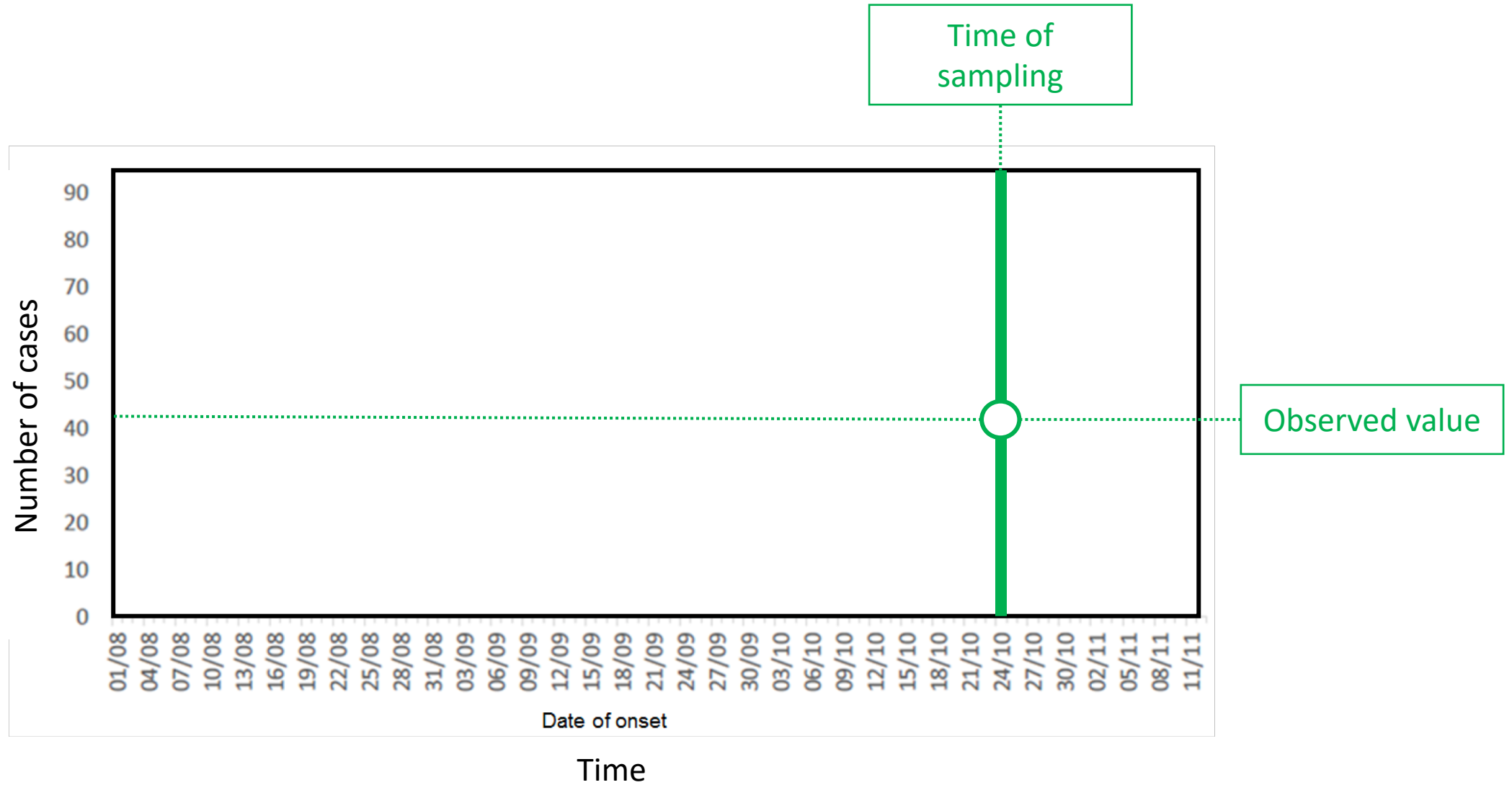
Limitations of cross-sectional surveys

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Limitations of cross-sectional surveys

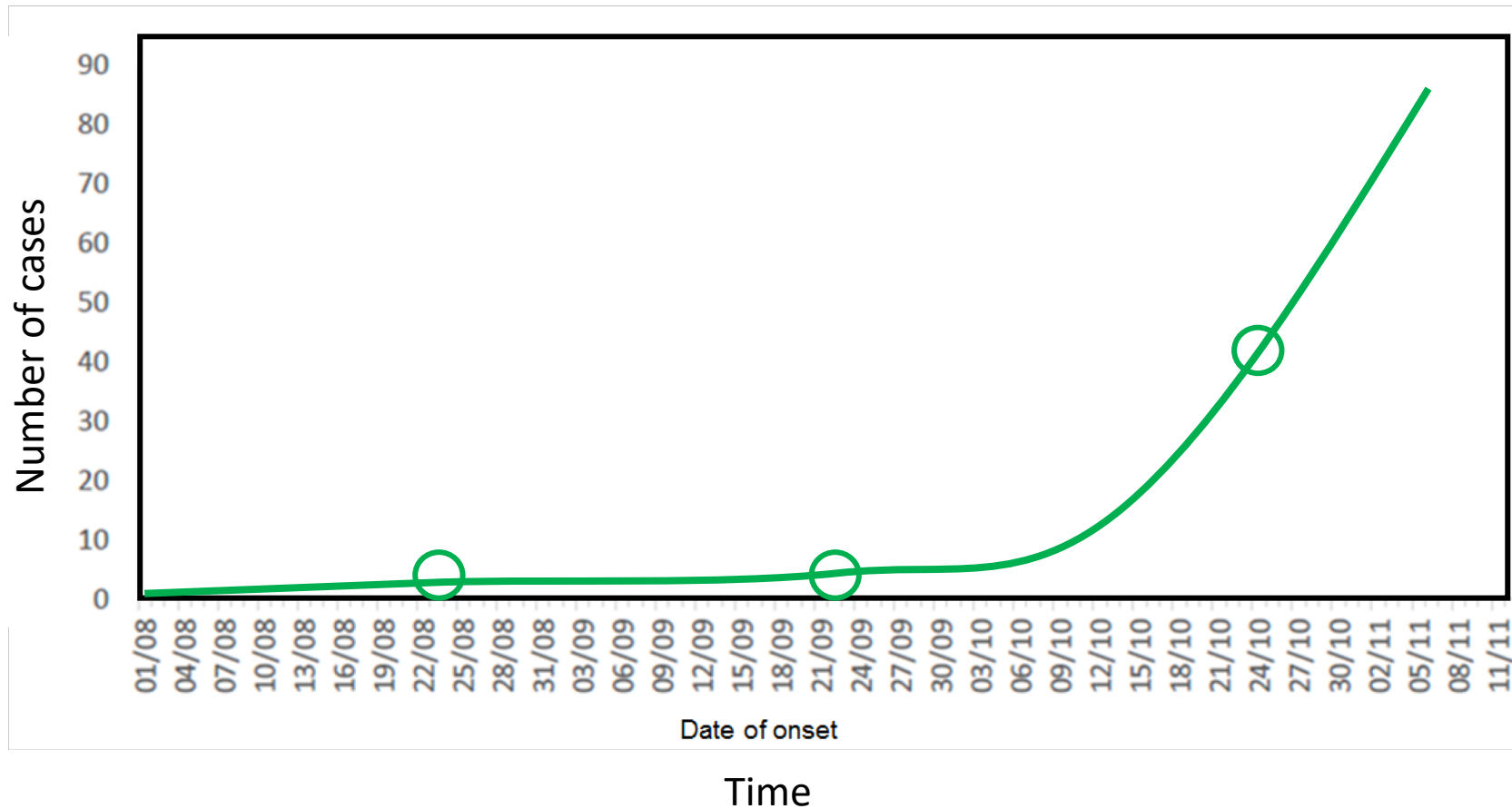
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Limitations of cross-sectional surveys

Example: Plague outbreak in Madagascar

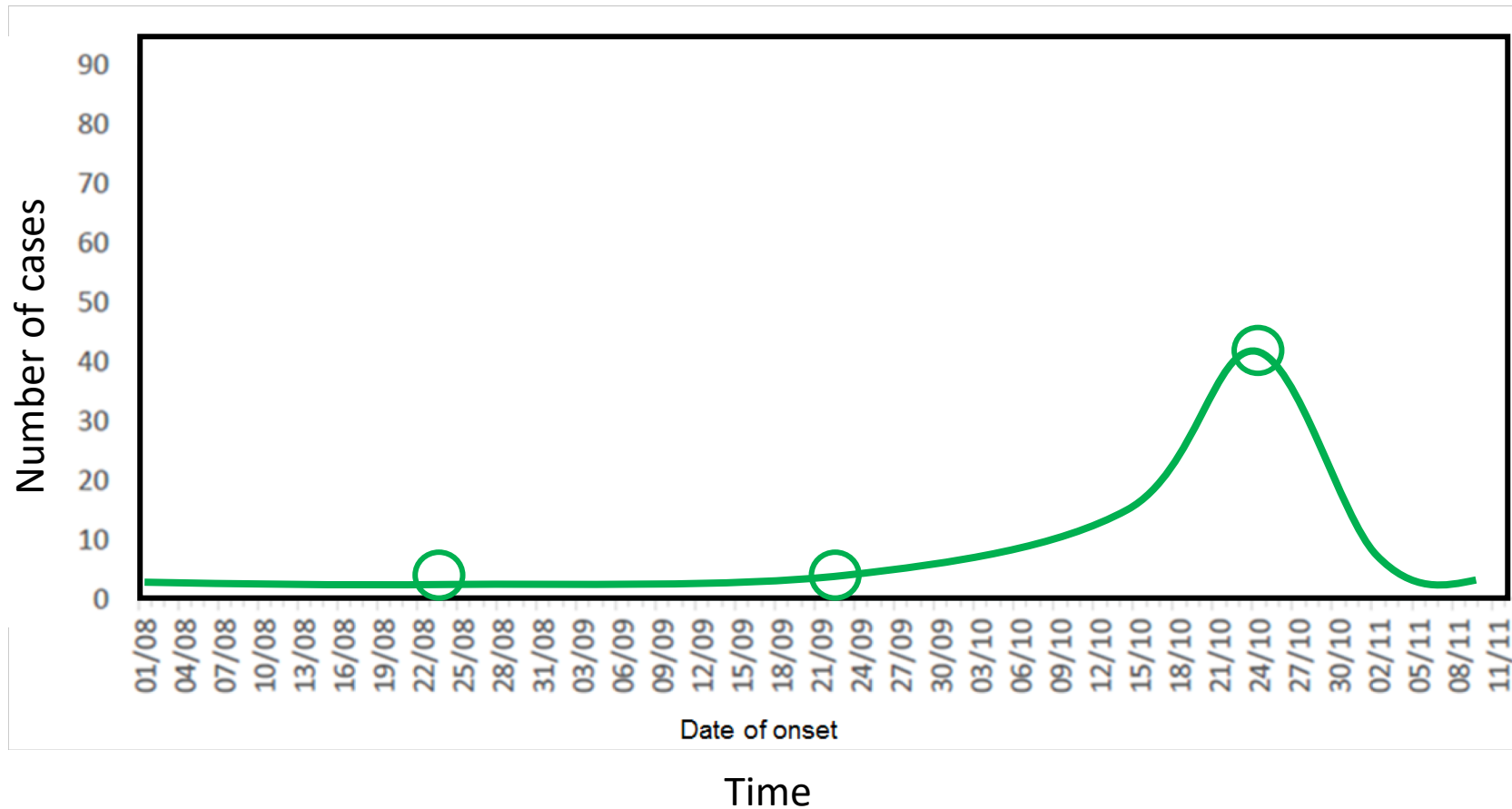
What we might think is happening if we tried to extrapolate from cross-sectional surveys



Limitations of cross-sectional surveys

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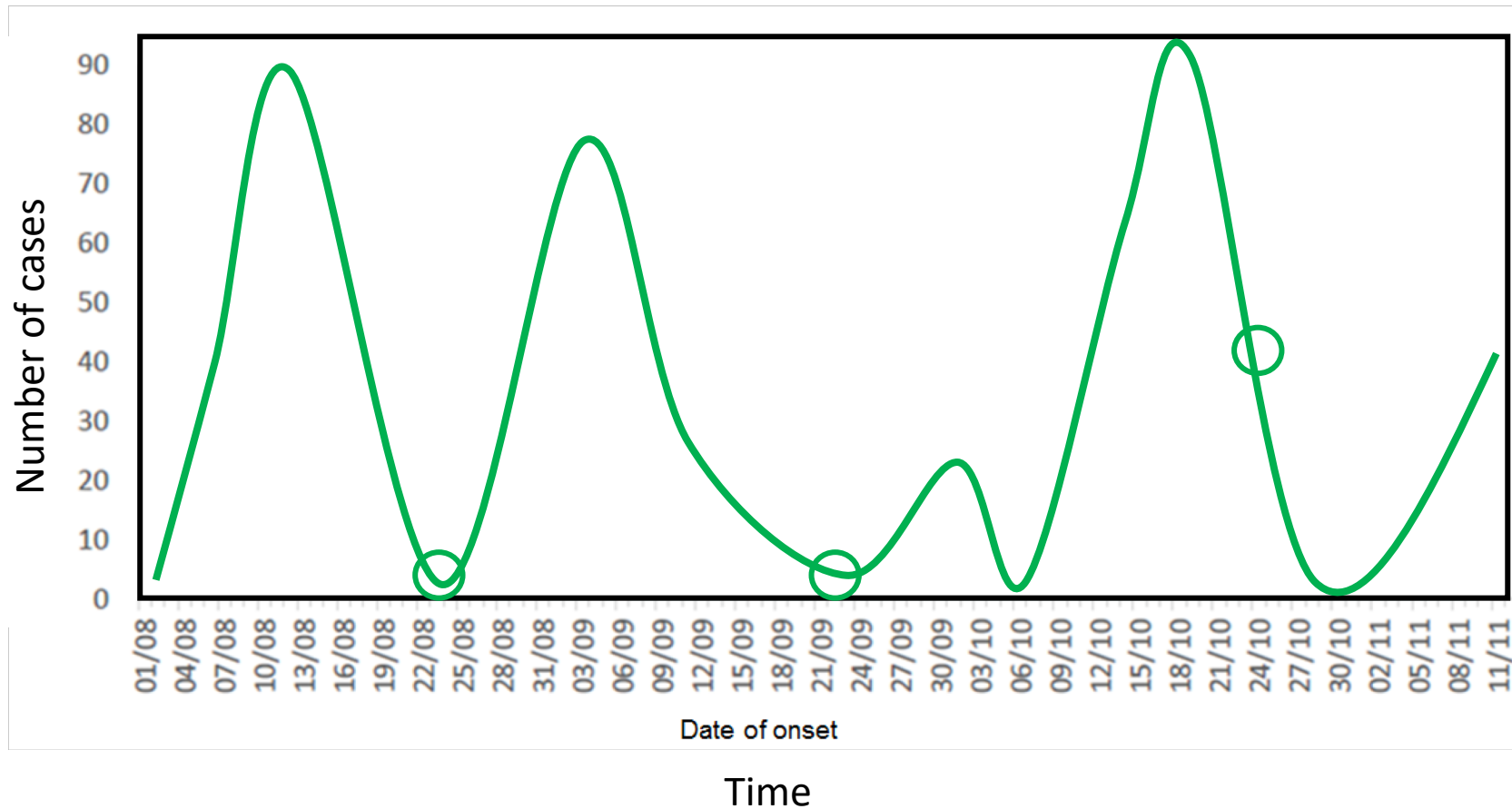
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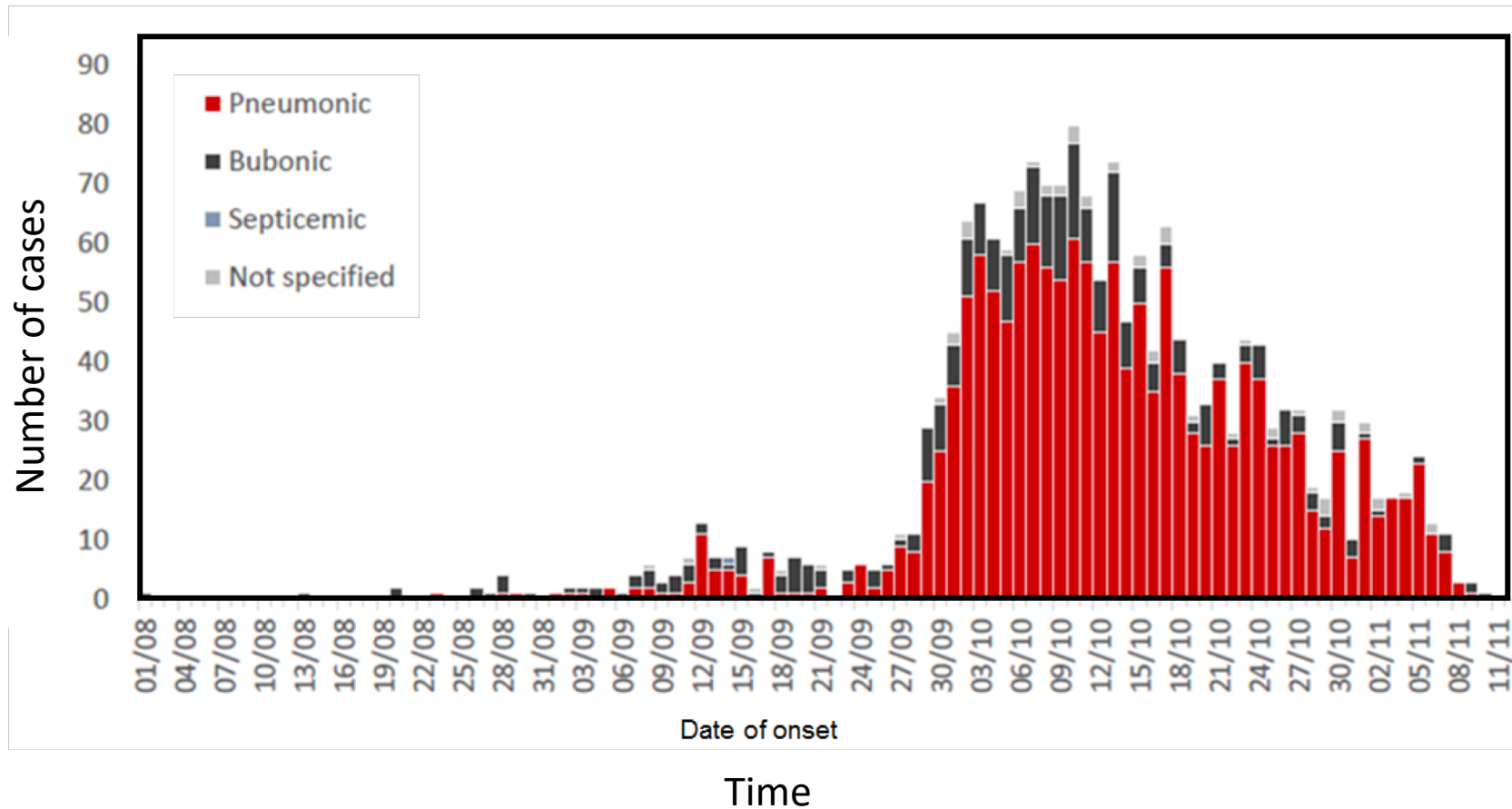
What we might think is happening if we tried to extrapolate from cross-sectional surveys



Limitations of cross-sectional surveys

Example: Plague outbreak in Madagascar

What really happened with the plague



Limitations of cross-sectional surveys

Example: Plague outbreak in Madagascar

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Limitations of cross-sectional surveys

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Difficult to capture changes over time

Sarotra manazava ny fiovana amin'ny fotoana

Limitations of cross-sectional surveys

Example: Plague outbreak in Madagascar

What is a major limitation of cross-sectional surveys?

Inona ny olana maveny no misy ny 'cross-sectional surveys'?

Difficult to capture changes over time

Sarotra manazava ny fiovana amin'ny fotoana

Conclusion: Not all study types can answer all the questions you have

Samy manana ny fanontaniana afaka valiany 'study' tsy iray-iray

How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)

OR

How to balance scientific and non-scientific factors
when designing a study?

How to not be paralyzed (*aza miasa saina*)?

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Limitations of cross-sectional surveys

Example: The MAHERY-CRS cross-sectional health survey (2017)

Problems:

Money is not unlimited
(Budget is \$175,000 USD)

E.g. transportation,
materials, analysis,

Time is not unlimited
(Time window: 1 year)

E.g. my advisor starts
to miss me

Subject participation
is not unlimited

E.g. blood draws are
not fun

Need to balance
multiple objectives
(malaria, viruses, nutrition, etc)

E.g. unknown effect
sizes and statistical
power

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Example: The MAHERY-CRS cross-sectional health survey (2017)

Problems:

Money is not unlimited
(Budget is \$175,000 USD)

E.g. transportation,
materials, analysis,

Time is not unlimited
(Time window: 1 year)

E.g. my advisor starts
to miss me

Subject participation
is not unlimited

E.g. blood draws are
not fun

Need to balance
multiple objectives
(malaria, viruses, nutrition, etc)

E.g. unknown effect
sizes and statistical
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How does study design happen in practice?

Ohatra: The MAHERY-CRS cross-sectional health survey (2017)

OR

**How to balance scientific and non-scientific factors
when designing a study?**

How to not be paralyzed (*aza miasa saina*)?

OR

How does study design place limits on your analysis?

How to do good science with imperfect studies (*'Science' tsara raha tsy lavorary ny 'study'*)?

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How to do good science with imperfect studies (*'Science' tsara raha tsy lavorary ny 'study'*)?

How can R help at every stage from day zero to the day of publication?

(Amin'ny 'study', manomboka andro voalohany hatramin'ny andron'ny 'publication':

Mora mora kokoa ve raha miasa amin'ny R?)

Using R in the MAHERY-CRS Project

Study design

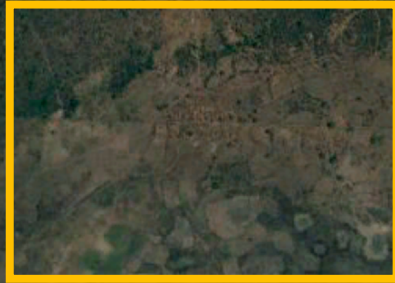
Using R in the MAHERY-CRS Project

Study design


Site and
household selection

```
graph LR; A[Study design] --> B[Site and household selection];
```

A diagram illustrating the relationship between 'Study design' and 'Site and household selection'. A blue box labeled 'Study design' is connected by a line to a black box labeled 'Site and household selection'. A vertical grey line is positioned to the right of the black box.



First, create a list of all *tanana kely* in the study region

An aerial photograph of a village in a dry, brown landscape with scattered green trees. The village consists of several clusters of buildings. Numerous small yellow squares are overlaid on the image, marking specific locations within the village. A blue rectangular box with white text is positioned in the upper right quadrant of the image.

Second, create a list of all
tokantrano in the *tanana kely*

Using R in the MAHERY-CRS Project

Study design

Site and
household selection

```
graph LR; A[Study design] --> B[Site and household selection];
```

A diagram illustrating the flow of information in the MAHERY-CRS Project. It features two rectangular boxes: a blue box on the left labeled 'Study design' and a black box on the right labeled 'Site and household selection'. A thin black line connects the right side of the blue box to the left side of the black box, indicating a sequential relationship. A vertical grey line is positioned to the right of the black box, separating it from the rest of the slide.

Using R in the MAHERY-CRS Project

Study design

Site and
household selection

```
#####  
## Randomly selecting study sites  
#####  
  
# Import a table of study sites  
study_site_table <- read.csv("study_sites.csv", stringsAsFactors = FALSE)  
  
# Load the tidyverse package to access some useful functions  
library(tidyverse)  
  
# Randomly select 6 sites within each region  
## use the group_by() and random() function  
selected_sites_table <- group_by(study_site_table, region_id) %>%  
  mutate(site_random_number = sample(1:length(region_id), length(region_id))) %>%  
  mutate(site_selected_y_n =  
    ifelse(site_random_number <= 6, 1, 0))  
  
selected_sites_table <- selected_sites_table[  
  selected_sites_table$site_selected_y_n == 1, ]
```

R code to randomly select sites and households:
tidyverse package + group_by() function

Using R in the MAHERY-CRS Project

Study design

Site and
household selection

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```

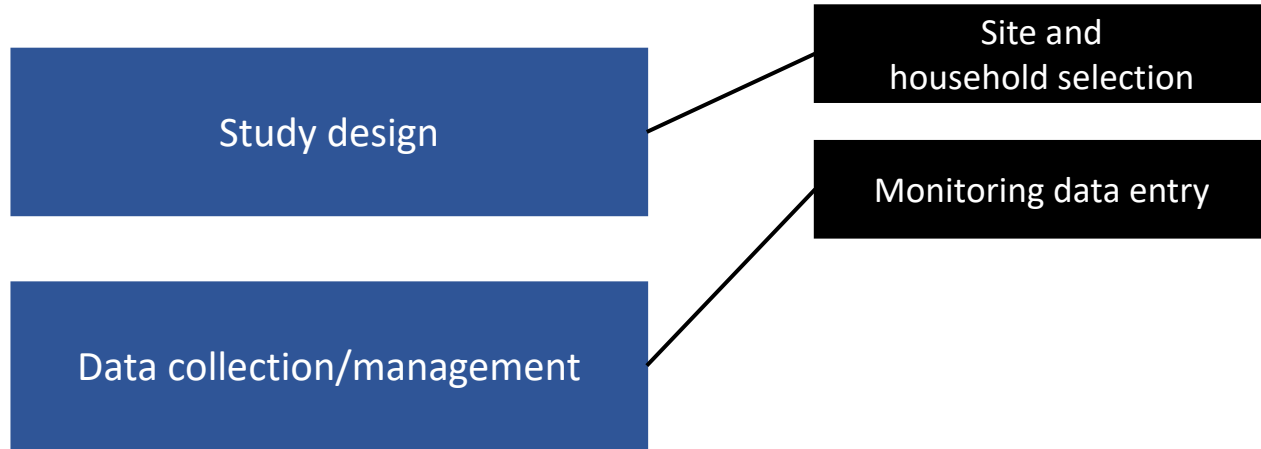
R code to randomly select sites and households:
tidyverse package + group_by() function

```
> selected_sites_table  
# A tibble: 24 x 4  
# Groups:   region_id [4]  
  region_id site_id site_random_number site_selected_y_n  
  <fct>      <int>          <int>          <dbl>  
1 R1         101             5             1  
2 R1         102             1             1  
3 R1         103             6             1  
4 R1         104             3             1  
5 R1         109             2             1  
6 R1         110             4             1  
7 R2         201             4             1  
8 R2         202             6             1
```

Using R in the MAHERY-CRS Project



Using R in the MAHERY-CRS Project



Data Sheet 1

CHECK IN SHEET

Region			Tanana			Andro		Volana		Taona	
R2 R3 R4 R5			V1 V2 V3 V4 V5 V6			2 8		July 2017		2/5	
Enumerators: <u>Hiadana</u>											
Clinic ID			Ananarana			Temperature (Celsius)		HARVARD: Tokonranro		Olo	
#	#	#	#	#	#	#	#	#	#	#	#
1	0	2	6	T				0	1	2	1 4
2	0	2	7	2				0	1	6	0 1
3	0	2	8	N				0	1	6	0 3
4	0	2	9	J				0	1	6	0 4
5	0	3	0	A				0	1	6	0 5
6	0	3	1	R				0	1	6	0 6
7	0	3	2	D				0	2	1	0 1
8	0	3	3	Y				0	2	1	0 2
9	0	3	4	F				0	2	1	0 3
10	0	3	5	T				0	0	3	0 2
11	0	3	6	P				0	0	3	0 3
12	0	3	7	C				0	0	3	0 4
13	0	3	8	G				0	0	3	0 5
14	0	3	9	T				0	0	3	0 6
15	0	4	0					0	0	3	0 7
16	0	4	1					0	3	6	0 1
17	0	4	2					0	3	6	0 3
18	0	4	3	F				0	3	6	0 5
19	0	4	4	T				0	3	6	0 7
20	0	4	5	N				0	1	6	0 2
21	0	4	6	D				0	2	6	0 1
22	0	4	7	T				0	2	6	0 3
23	0	4	8					0	2	6	0 4
24	0	4	9					0	2	6	0 5
25	0	5	0					0	2	8	0 6

Results		#		#		#	
NEG	Pan	INV	PF	NEG	Pan	INV	PF
NEG	Pan	10	2	OK			
NEG	Pan	13	1	OK			
NEG	Pan	13	4	OK			
NEG	Pan	13	3	OK			
NEG	Pan	11	9	OK			
NEG	Pan	11	7	OK			
NEG	Pan	15	4	OK			
NEG	Pan	14	4	OK			
NEG	Pan	10	6	OK			
NEG	Pan	13	7	OK			
NEG	Pan	13	9	OK			
NEG	Pan	14	5	OK			
NEG	Pan	13	3	OK			
NEG	Pan	14	7	OK			
NEG	Pan	14	1	OK			
NEG	Pan	16	2	OK			
NEG	Pan	13	6	OK			
NEG	Pan	12	3	OK			
NEG	Pan	9	8	OK			
NEG	Pan	14	7	OK			
NEG	Pan	14	2	OK			
NEG	Pan	11	9	OK			
NEG	Pan	13	7	OK			
NEG	Pan	12	5	OK			
NEG	Pan	11	7	OK			

Hb = manome

CS.4.02

Data Sheet 2

BLOOD DRAW TIME SHEET

Enumerator: **BEN**

Region: R2 R3 **R4** R5

Tanana: V1 V2 V3 **V4** V5 V6

Andro: **03** June 3/5

Volana (2017): **June 3/5**

#	Tokorano		Olo		Clinic ID		Prise de sang:		Temps		RDT		Hemoglobin		Centrifugation (Temps)		N2 Tank (Temps)		Cryos		
	#	#	#	#	#	#	G.D.P.	Tsila	Heure (#)	Minutes (##)	NEG	INV	Pf	#	#	Heure (#)	Minutes (##)	Heure (#)		Minutes (##)	
1	0	8	5	0	2	1	4	7	<input checked="" type="checkbox"/> G.D.P.	5	48	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	10	2	5	55	6	16	0 1 0
2	0	7	4	0	4	1	4	6	<input checked="" type="checkbox"/> G.D.P.	5	48	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	10	2	5	55	6	16	0 1 0
3	0	8	5	0	3	1	4	8	<input checked="" type="checkbox"/> G.D.P.	5	49	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	1	5	55	6	16	0 1 0
4	0	8	5	0	4	1	4	9	<input checked="" type="checkbox"/> G.D.P.	5	50	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	9	9	5	55	6	16	0 1 0
5	0	8	6	0	3	1	5	1	<input checked="" type="checkbox"/> G.D.P.	5	51	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	2	5	55	6	16	0 1 0
6	0	8	6	0	2	1	5	0	<input checked="" type="checkbox"/> G.D.P.	5	52	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	12	9	6	10	6	29	0 1 0
7	0	8	6	0	4	1	5	2	<input checked="" type="checkbox"/> G.D.P.	5	55	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	9	6	10	6	29	0 1 0
8	0	3	7	0	1	1	5	5	<input checked="" type="checkbox"/> G.D.P.	5	57	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	14	7	6	10	6	29	0 1 0
9	0	8	6	0	5	1	5	3	<input checked="" type="checkbox"/> G.D.P.	5	58	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	8	6	10	6	29	0 1 0
10	0	8	2	0	2	1	5	6	<input checked="" type="checkbox"/> G.D.P.	6	00	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	12	6	6	10	6	29	0 1 0
11	0	8	2	0	4	1	5	7	<input checked="" type="checkbox"/> G.D.P.	6	02	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	7	6	10	6	29	0 1 0
12	0	3	7	0	3	1	6	0	<input checked="" type="checkbox"/> G.D.P.	6	05	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	13	0	6	16	6	29	0 1 0
13	0	7	4	0	3	1	5	4	<input checked="" type="checkbox"/> G.D.P.	6	08	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	8	6	6	10	6	29	0 1 0
14	0	8	2	0	6	1	5	8	<input checked="" type="checkbox"/> G.D.P.	6	12	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	8	7	6	26	6	45	0 1 0
15	0	5	5	0	2	1	6	1	<input checked="" type="checkbox"/> G.D.P.	6	12	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	12	4	6	26	6	45	0 1 0
16	0	8	2	0	7	1	5	9	<input checked="" type="checkbox"/> G.D.P.	6	17	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	13	0	6	26	6	45	0 1 0
17	0	5	5	0	5	1	6	4	<input checked="" type="checkbox"/> G.D.P.	6	20	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	9	8	6	26	6	45	0 1 0
18	0	5	5	0	4	1	6	3	<input checked="" type="checkbox"/> G.D.P.	6	22	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	8	9	6	26	6	45	0 1 0
19	0	5	5	0	3	1	6	2	<input checked="" type="checkbox"/> G.D.P.	6	24	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	10	0	6	26	6	45	2 0 0
20	0	5	5	0	7	1	6	5	<input checked="" type="checkbox"/> G.D.P.	6	24	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	9	3	6	26	6	45	0 1 0
21	0	1	4	0	2	1	6	8	<input checked="" type="checkbox"/> G.D.P.	6	26	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	13	6	6	42	7	02	0 1 0
22	0	6	2	0	8	1	6	7	<input checked="" type="checkbox"/> G.D.P.	6	28	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	12	1	6	42	7	02	0 1 0
23	0	5	5	0	6	1	6	6	<input checked="" type="checkbox"/> G.D.P.	6	29	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	9	9	6	42	7	02	0 1 0
24	0	1	4	0	6	1	7	0	<input checked="" type="checkbox"/> G.D.P.	6	31	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	12	4	6	42	7	02	0 1 0
25	0	1	4	0	5	1	6	9	<input checked="" type="checkbox"/> G.D.P.	6	32	<input checked="" type="checkbox"/> NEG	<input checked="" type="checkbox"/> INV	<input checked="" type="checkbox"/> Pf	11	9	6	42	7	02	0 1 0

T4.4.07

Data entry into excel

Ben R2 CheckIn 2017 08 04

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V
1	Clinic ID	Name	Temp	Harvard	RDT	Hb	Enum	Page	Sheet ID	R	V	Date	kg	cm	MUAC	CC						
662	019		35.8	042.03	neg	13.2	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
663	020		36.2	042.04	pan	10.3	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
664	021		35.8	042.05	neg	11.8	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
665	022		35.6	055.01	neg	15.0	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
666	023		35.9	055.02	neg	12.1	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
667	024		35.8	055.03	neg	13.2	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
668	025		29.7	055.04	neg	10.6	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
669	026		28.6	055.05	neg	10.7	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
670	027		31.2	059.01	neg	15.6	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
671	028		27.7	059.02	neg	16.1	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
672	029		27.5	059.03	neg	14.2	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
673	030		31.1	059.04	neg	16.0	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
674	031		29.8	059.05	neg	14.6	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
675	032		31.5	059.06	neg	14.9	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
676	033		32.4	059.07	panpf	11.0	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
677	034		36.2	059.08	neg	9.7	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
678	035		36.3	059.09	pan	7.9	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
679	036		36.5	059.10	neg	10.1	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
680	037		32.8	025.01	neg	13.8	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
681	038		33.6	025.02	panpf	9.2	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
682	039		33.7	025.03	pan	10.6	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
683	040		33.6	025.04	neg	10.7	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
684	041		32.8	020.01	pan	11.5	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
685	042		32.1	020.02	neg	11.5	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
686	043		33.2	020.03	neg	13.3	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						
687	044		32.5	020.04	panpf	9.8	Hervet	1 of 3	C2.4.01	2	4	20-Feb-17	ND	ND	ND	ND						

Data entry into excel

*Fa manana olana hely isika:
Mora mandiso rehefa mampiditra 'data'*

Data entry into excel

Fa manana olana hely isika:

Mora mandiso rehefa mampiditra 'data'

Solution: Ataovy mampiditra 'data' in-telo

Data entry into excel
(Repeat to minimize errors)

[illegible]

Replicate 1

+

[illegible]

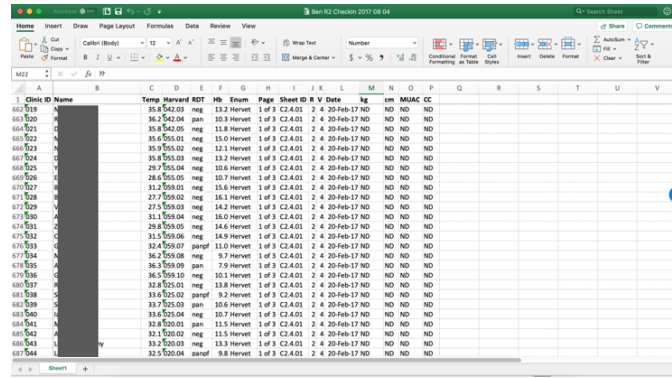
Replicate 2

+

[illegible]

Replicate 3

Data entry into excel
(Repeat to minimize errors)



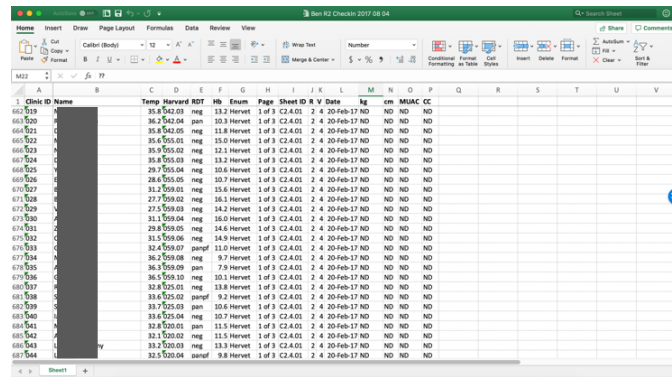
Clinic ID	Name	Temp	Harvard	RDT	Hb	Enum	Page	Sheet ID	R	V	Date	Ig	cm	MMAC	CC
66519		35.8	561.05	neg	11.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66520		36.2	561.04	pan	10.3	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66521		35.8	561.05	neg	11.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66522		35.8	561.05	neg	11.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66523		35.9	561.02	neg	11.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66524		35.8	561.03	neg	11.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66525		29.7	561.04	neg	10.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66526		28.8	561.05	neg	10.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66527		31.2	561.01	neg	15.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66528		27.7	561.02	neg	16.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66529		27.5	561.03	neg	14.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66530		31.1	561.04	neg	16.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66531		29.8	561.05	neg	14.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66532		31.5	561.06	neg	14.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66533		32.4	561.07	pan	11.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66534		36.2	561.08	neg	9.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66535		36.5	561.09	pan	7.9	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66536		36.5	561.10	neg	10.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66537		32.8	561.01	neg	13.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66538		33.6	561.02	pan	9.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66539		33.7	561.03	pan	10.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66540		33.8	561.04	neg	10.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66541		32.8	561.05	pan	11.5	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66542		32.1	561.06	neg	11.5	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66543		33.2	561.03	neg	13.3	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66544		32.5	561.04	pan	9.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND

Replicate 1

Fa hitondra olana hely hafa:
Mila mihaza ny 'errors' / Mila mizaha ny 'errors'
Solution: R code

Replicate 2

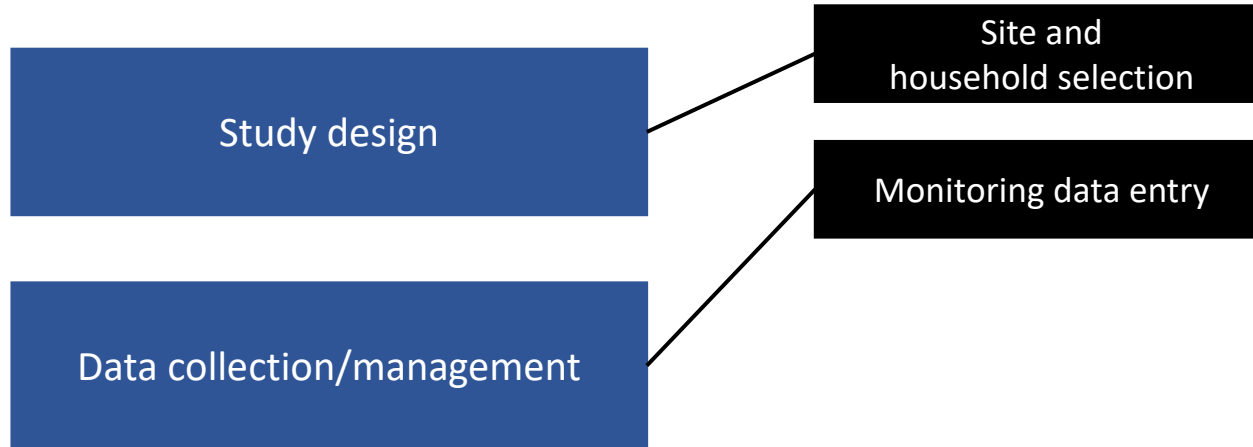
+



Clinic ID	Name	Temp	Harvard	RDT	Hb	Enum	Page	Sheet ID	R	V	Date	Ig	cm	MMAC	CC
66519		35.8	561.05	neg	13.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66520		36.5	561.04	pan	10.3	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66521		35.8	561.05	neg	11.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66522		35.8	561.05	neg	11.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66523		35.9	561.02	neg	11.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66524		35.8	561.03	neg	11.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66525		29.7	561.04	neg	10.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66526		28.8	561.05	neg	10.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66527		31.2	561.01	neg	15.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66528		27.7	561.02	neg	16.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66529		27.5	561.03	neg	14.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66530		31.1	561.04	neg	16.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66531		29.8	561.05	neg	14.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66532		31.5	561.06	neg	14.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66533		32.4	561.07	pan	11.0	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66534		36.2	561.08	neg	9.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66535		36.5	561.09	pan	7.9	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66536		36.5	561.10	neg	10.1	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66537		32.8	561.01	neg	13.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66538		33.6	561.02	pan	9.2	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66539		33.7	561.03	pan	10.6	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66540		33.8	561.04	neg	10.7	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66541		32.8	561.05	pan	11.5	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66542		32.1	561.06	neg	11.5	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66543		33.2	561.03	neg	13.3	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND
66544		32.5	561.04	pan	9.8	Harvet	1 of 3	C2.A01	2	4	20-Feb-17	ND	ND	ND	ND

Replicate 3

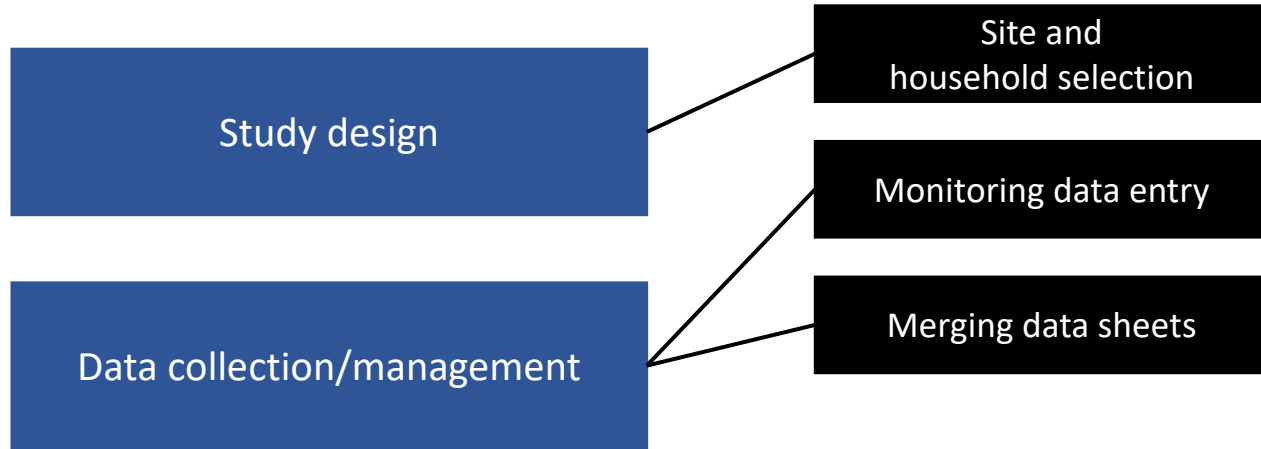
Using R in the MAHERY-CRS Project



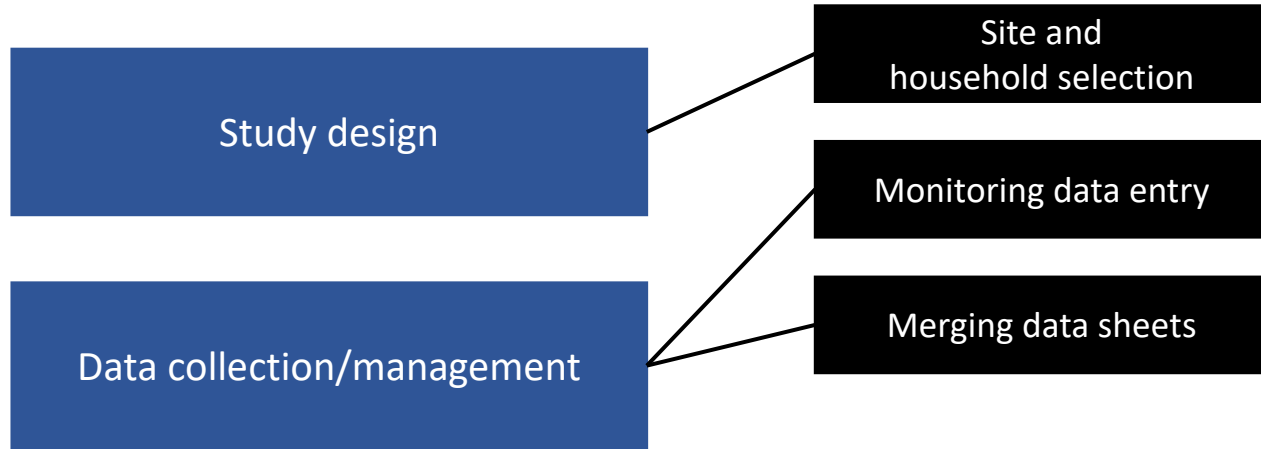
```
#####  
## Monitoring data entry  
#####  
  
# Concatenate data fields in a row into one string using a function and a for loop  
concatenate.function <- function(dataframe){  
  df <- dataframe  
  concatenated_df <- rep(0, length(df[,1]))  
  for(i in 1:length(df[,1])){  
    concatenated_df[i] <- paste(  
      df[i, 1], df[i, 2], df[i, 3], df[i, 4], df[i, 5], sep = ".")  
  }  
  return(concatenated_df)  
}  
  
df1.concatenated <- concatenate.function(df1)  
df2.concatenated <- concatenate.function(df2)  
  
# compare the repeated data entries to see if entries are the same  
compare.function <- function(dataframe1, dataframe2){  
  rows_with_errors <- rep(NA, length(dataframe1))  
  for(i in 1:length(dataframe1)){  
    rows_with_errors[i] <- ifelse(dataframe1[i] == dataframe2[i], 0, 1)  
  }  
  return(rows_with_errors)  
}  
  
compare.function(df1.concatenated, df2.concatenated)
```

R code to compare data tables:
for loops + ifelse() function

Using R in the MAHERY-CRS Project



Using R in the MAHERY-CRS Project



Olana: Mila manambatra 'data files' roa

Clinical data

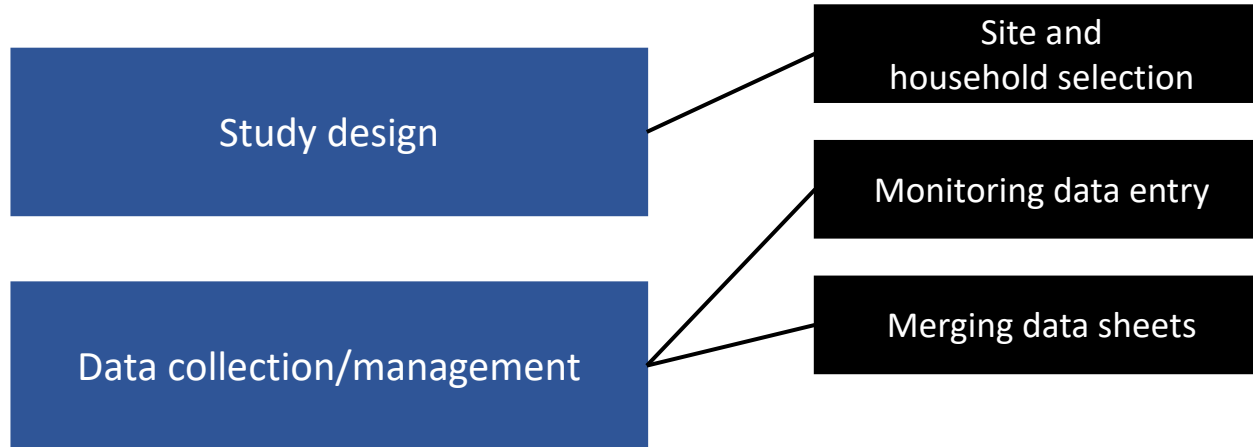
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Survey data

=

Data to analyze

Using R in the MAHERY-CRS Project



Olana: Mila manambatra 'data files' roa

Clinical data

+

Survey data

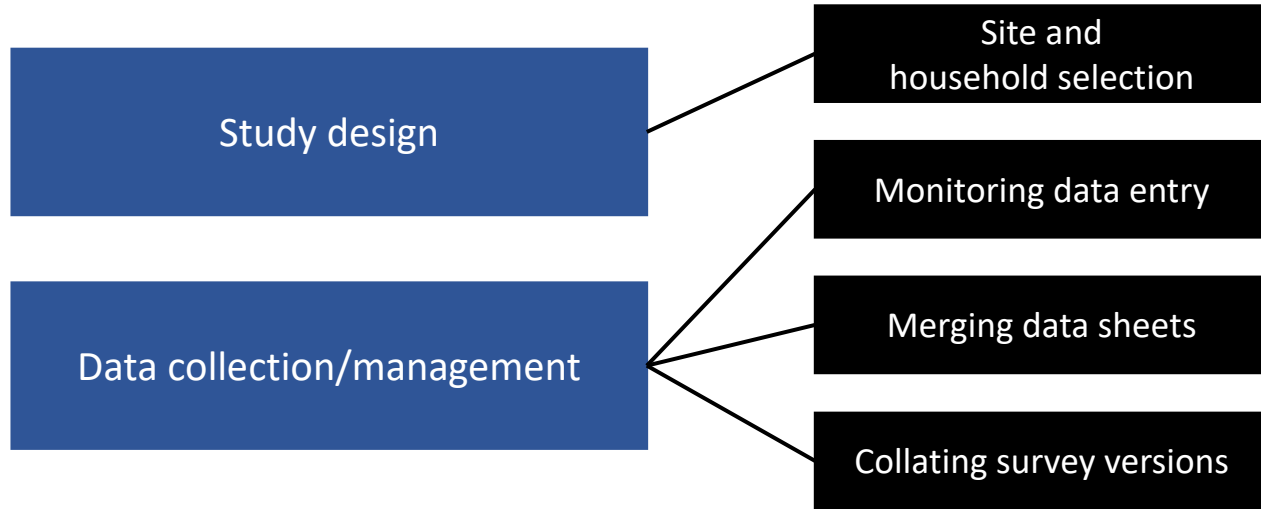
=

Data to analyze

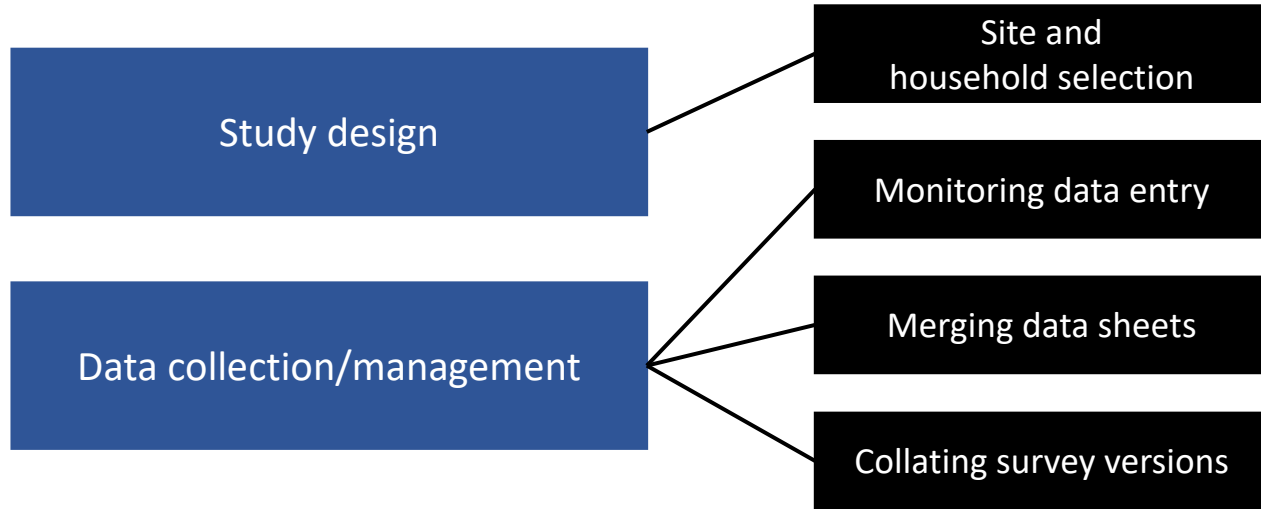
```
#####  
## Merging data sheets  
#####  
  
# Import clinical data and survey data  
clinical_data <- read.csv("clinical_data.csv", stringsAsFactors = FALSE)  
survey_data <- read.csv("survey_data.csv", stringsAsFactors = FALSE)  
  
# Subset the clinical data to keep the relevant variables  
rdt_subset_variables <- c("unique_ind_id", "rdt_result")  
clinical_data_trimmed <- clinical_data[rdt_subset_variables]  
  
# Merge data frames using the full_join() function  
  
# First, need to check for duplicate IDs using anyDuplicated()  
# If the output of anyDuplicated > 0, then there are duplicate IDs  
anyDuplicated(clinical_data_trimmed$unique_ind_id)  
anyDuplicated(survey_data$unique_ind_id)  
  
# Merge dataframes using the full_join() function and store as data_joined  
data_joined <- full_join(clinical_data_trimmed, survey_data, by = "unique_ind_id")  
  
# Check merged sheet  
head(data_joined)  
str(data_joined)
```

Check for duplicate IDs, merge using full_join()

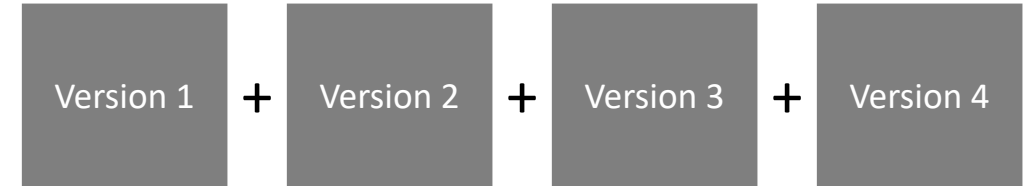
Using R in the MAHERY-CRS Project



Using R in the MAHERY-CRS Project



Olana be vata: Mila manambatra 'versions' efatra

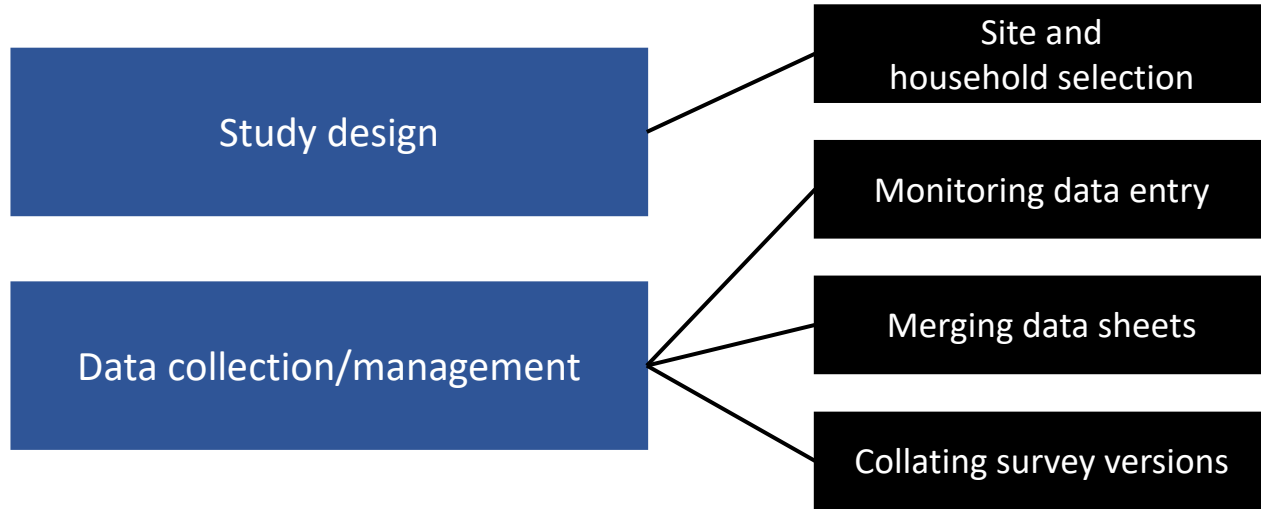


	English question text	Malagasy question text	name_r5v	label_r5v	name_R3V2	label_R3V2	name_R3V1	label_R3V1	name_r2v	label_r2v	name_r3v	label_r3v
	Section 00: Standard Initial Section	Section 00: Standard premiere question	_06_01_001	Section 01: Standard pr	_06_01_001	Section 01: Standard	_06_01_001	Section 01: Standard	_06_01_001	Section 01: Standard	_06_01_001	Section 01: Standard
	Site ID (standardized ID code for sites: Region Code + Site Code)	Site ID	_06_01_001_01	Site ID	_06_01_001_01	Site ID	_06_01_001_01	Site ID	_06_01_001_01	Site ID	_06_01_001_01	Site ID
	Region code (2 = Mananjary, 3 = Toliara, 4 = Morombe, 5 = Amoron'i Mania)	Region	_06_01_001_02	Region	_06_01_001_02	Region	_06_01_001_02	Region	_06_01_001_02	Region	_06_01_001_02	Region
	Village code (1-6)	Village code (1-6)	_06_01_001_03	Village code (1-6)	_06_01_001_03	Village code (1-6)	_06_01_001_03	Village code (1-6)	_06_01_001_03	Village code (1-6)	_06_01_001_03	Village code (1-6)
	Unique Household ID (Region Code + Site Code + HH Code)	Household ID	_06_01_001_04	Household ID	_06_01_001_04	Household ID	_06_01_001_04	Household ID	_06_01_001_04	Household ID	_06_01_001_04	Household ID
	Unique Individual ID (Region Code + Site Code + HH Code + Ind Code)	Individual ID	_06_01_001_05	Individual ID	_06_01_001_05	Individual ID	_06_01_001_05	Individual ID	_06_01_001_05	Individual ID	_06_01_001_05	Individual ID
	Prompt: This first section is for the enumerator to respond (not a question)	Ho an'ny mpanadiady manokana ity section 1 ity	_06_01_002	Ho an'ny mpanadiady r	_06_01_002	Ho an'ny mpanadiady	_06_01_002	Ho an'ny mpanadiady	_06_01_002	Ho an'ny mpanadiady	_06_01_002	Ho an'ny mpanadiady
	Enumerator Name	Mpanadiady	_06_01_003	Mpanadiady	_06_01_003	Mpanadiady	_06_01_003	Mpanadiady	_06_01_003	Mpanadiady	_06_01_003	Mpanadiady
	District	Disctrict	_06_01_004	Disctrict	_06_01_004	Disctrict	_06_01_004	Disctrict	_06_01_004	Disctrict	_06_01_004	Disctrict
	Village Name (Mananjary District)	Village Mananjary	_06_01_005	Village Mananjary	_06_01_005	Village Mananjary	_06_01_005	Village Mananjary	_06_01_005	Village Mananjary	_06_01_005	Village Mananjary
	Village Name (Toliara II District)	Village Toliara II	_06_01_006	Village Toliara II	_06_01_006	Village Toliara II	_06_01_006	Village Toliara II	_06_01_006	Village Toliara II	_06_01_006	Village Toliara II
Data	Village Name (Morombe District)	Village Morombe	_06_01_007	Village Morombe	_06_01_007	Village Morombe	_06_01_007	Village Morombe	_06_01_007	Village Morombe	_06_01_007	Village Morombe
	Village (Amoron'i Mania Faritra)	Village Amoron'i Mania	_06_01_008	Village Amoron'i Mania	_06_01_008	Village Amoron'i Ma	_06_01_008	Village Amoron'i Ma	_06_01_008	Village Amoron'i Ma	_06_01_008	Village Amoron'i Ma
	Enumerator: Are you ready to begin the survey?	Vonona hanomboka?	_06_01_009	Vonona hanomboka?	_06_01_009	Vonona hanomboka?	_06_01_009	Vonona hanomboka?	_06_01_009	Vonona hanomboka?	_06_01_009	Vonona hanomboka?
	Enumerator: Does the interviewee agree to do the questionnaire and do they understand the questionnaire?	Manaiky ny hamaly ny fanontaniana ve ireo olona voakasik'ialy fanadiadiana?	_06_01_010	Manaiky ny hamaly ny	_06_01_010	Manaiky ny hamaly r	_06_01_010	Manaiky ny hamaly r	_06_01_010	Manaiky ny hamaly r	_06_01_010	Manaiky ny hamaly r
	Text of summary of consent/assent form to remind enumerators in case there are questions	Text of summary of consent/assent form to remind enumerators in case there are questions (See to the right)	_06_01_0121	Fanazavana mikasika il	_06_01_0121	Fanazavana mikasika	_06_01_0121	Fanazavana mikasika	_06_01_0121	Fanazavana mikasika	_06_01_0121	Fanazavana mikasika
	Section 01: ID and gender	Section 01: ID des individus et genre	_06_01_0131	Section 02: ID des indiv	_06_01_0131	Section 02: ID des in	_06_01_0131	Section 02: ID des inv	_06_01_0131	Section 02: ID des inv	_06_01_0131	Section 1.1: ID des i
	Gender	Genre	_06_01_013	Genre	_06_01_013	Genre	_06_01_013	Genre	_06_01_013	Genre	_06_01_013	Genre
	Harvard Household ID	Harvard Menage ID (###)	_06_01_014	Harvard Menage ID (##	_06_01_014	Harvard Menage ID (_06_01_014	Harvard Menage ID (_06_01_014	Harvard Menage ID (_06_01_014	Harvard Menage ID
	Harvard Individual	Harvard Individu ID (##)	_06_01_015	Harvard Individu ID (##	_06_01_015	Harvard Individu ID (_06_01_015	Harvard Individu ID (_06_01_015	Harvard Individu ID (_06_01_015	Harvard Individu ID
	DOB: Year	Daty nahaterahana: Taona (####)	_06_01_016	Daty nahaterahana: Ta	_06_01_016	Daty nahaterahana:	_06_01_016	Daty nahaterahana:	_06_01_016	Daty nahaterahana:	_06_01_016	Daty nahaterahana:
	DOB: Month	Daty nahaterahana: Volana	_06_01_017	Daty nahaterahana: Vc	_06_01_017	Daty nahaterahana:	_06_01_017	Daty nahaterahana:	_06_01_017	Daty nahaterahana:	_06_01_017	Daty nahaterahana:
	DOB: Day	Daty nahaterahana: Andro (##)	_06_01_018	Daty nahaterahana: Ar	_06_01_018	Daty nahaterahana:	_06_01_018	Daty nahaterahana:	_06_01_018	Daty nahaterahana:	_06_01_018	Daty nahaterahana:
	Actual or approximate DOB?	TENA Taona na VERS Taona?	_06_01_0181	TENA Taona na VERS T	_06_01_0181	TENA Taona na VERS	_06_01_0181	TENA Taona na VERS	_06_01_0181	TENA Taona na VERS Taona?		
	Calculated age (2017 minus year of birth)	Taonan'ilay olona	_06_01_019	Taonan'ilay olona	_06_01_019	Taonan'ilay olona	_06_01_019	Taonan'ilay olona	_06_01_019	Taonan'ilay olona	_06_01_019	Taonan'ilay olona
	Section 02: Anthropometry and direct measures (recorded in individuals survey for R3V only)	Section 02: Lanja: Anthropometry and direct measures (R3V only)	NA	NA	NA	NA	_06_02_001	Section 02: Antropon	NA	NA	NA	NA
	Height (cm)	Taille (##.# or ###.#)	NA	NA	NA	NA	_06_02_002	Taille (##.# or ###.#)	NA	NA	NA	NA
	Weight (kg)	Poids (##.# or ###.# or ###.#)	NA	NA	NA	NA	_06_02_003	Poids (##.# or ##.# or #	NA	NA	NA	NA
	MUAC (cm)	MUAC (##.#)	NA	NA	NA	NA	_06_02_004	MUAC (##.#)	NA	NA	NA	NA
	Cranial circumference (cm)	Circonference de la tete (CDLT) (##.#)	NA	NA	NA	NA	_06_02_005	Circonference de la te	NA	NA	NA	NA
	Section 03: Development and disability	Section 03: Développement et handicap	_06_023_001	Section 03: Développer	_06_023_001	Section 03: Développ	_06_023_001	Section 03: Développ	_06_02_001	Section 02: Dévelop	_06_02_001	Section 02: Dévelop
	Does this individual have a physical disability that prevents them from moving like others their own age?	Manana fahasembanana ara-batana ve io olona io izay manjary lasa sakana amin'ny fihetsiny raha hoarina amin'ny ireo olona mitovy taona aminy?	_06_023_002	Manana fahasembanan	_06_023_002	Manana fahasemban	_06_023_002	Manana fahasemban	_06_02_002	Manana fahasemban	_06_02_002	Manana fahasemban
	Please describe the disability.	Inona ilay fahasembanana ara-batana?	_06_023_003	Inona ilay fahasemban	_06_023_003	Inona ilay fahasemba	_06_023_003	Inona ilay fahasemba	_06_02_003	Inona ilay fahasemba	_06_02_003	Inona ilay fahasemba

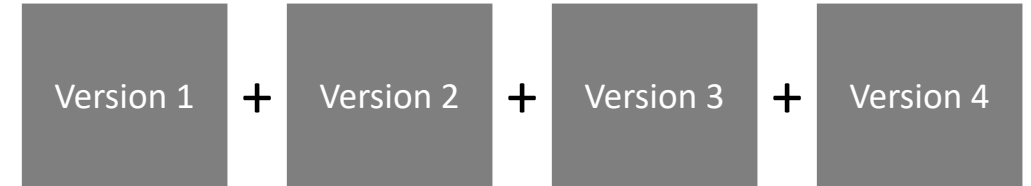
efatra

osion 4

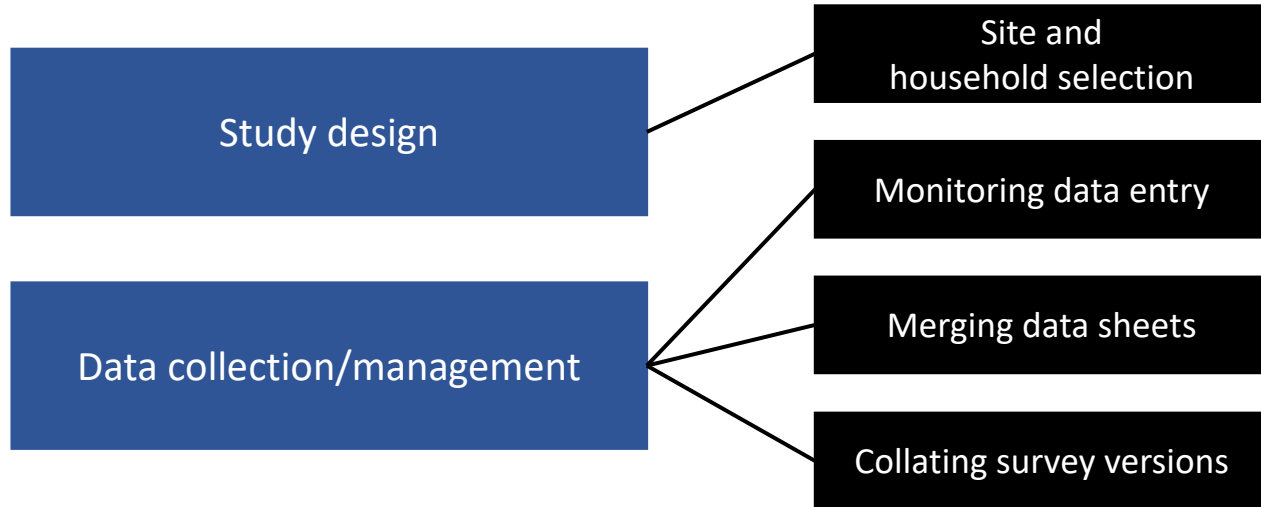
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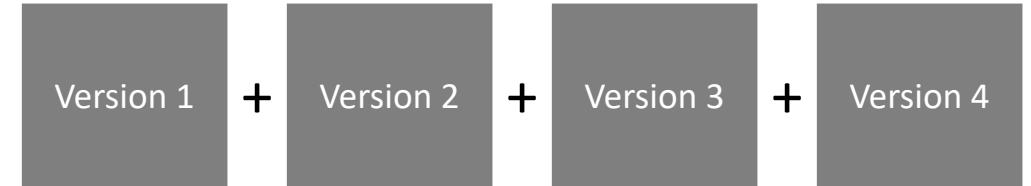
Olana be vata: Mila manambatra 'versions' efatra



Using R in the MAHERY-CRS Project



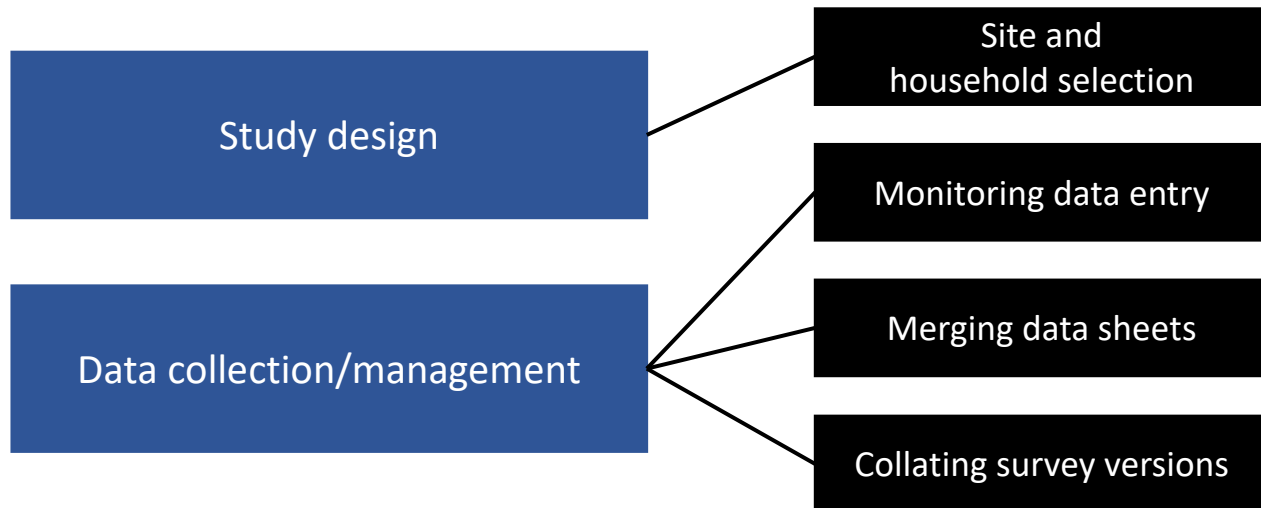
Olana be vata: Mila manambatra 'versions' efatra



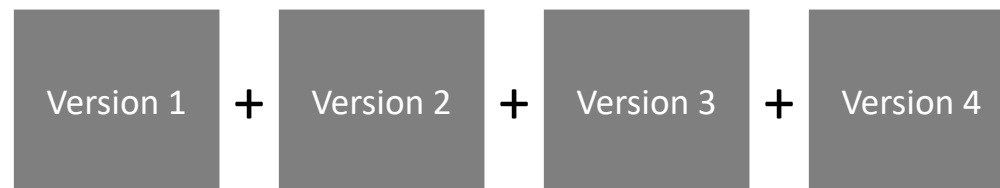
My strategy:

Step 1)

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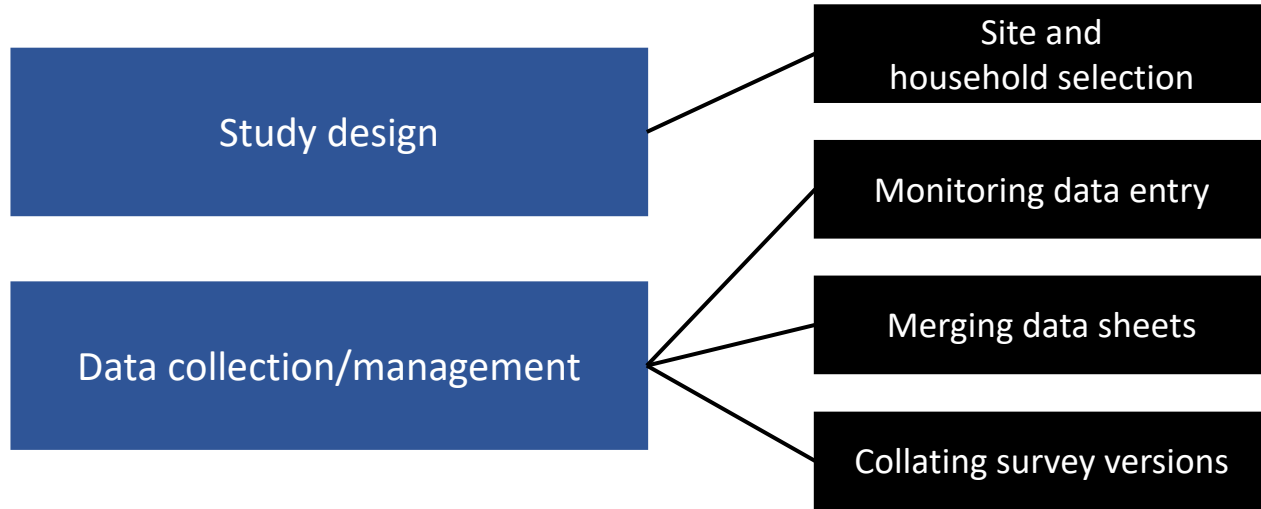
Olana be vata: Mila manambatra 'versions' efatra



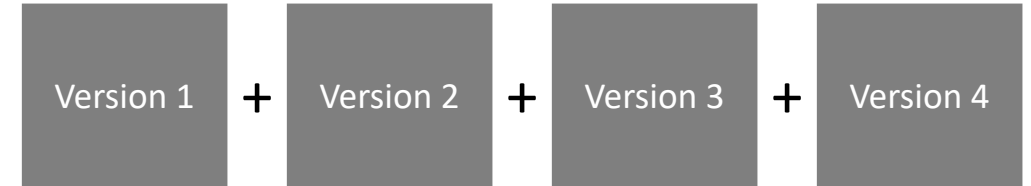
My strategy:

Step 1) Mangataka fanampiana amin'ny Amy Wesolowski sy Amy Winter ary Jessica Metcalf (satria manan-tsaina be izy telo)

Using R in the MAHERY-CRS Project



Olana be vata: Mila manambatra 'versions' efatra

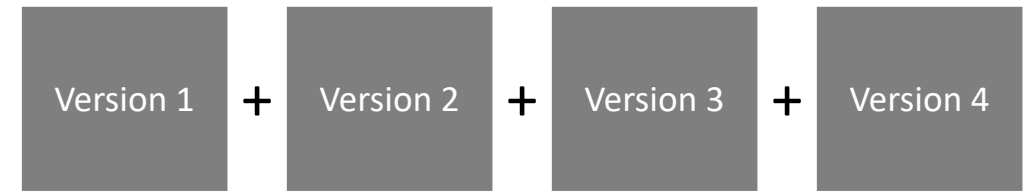
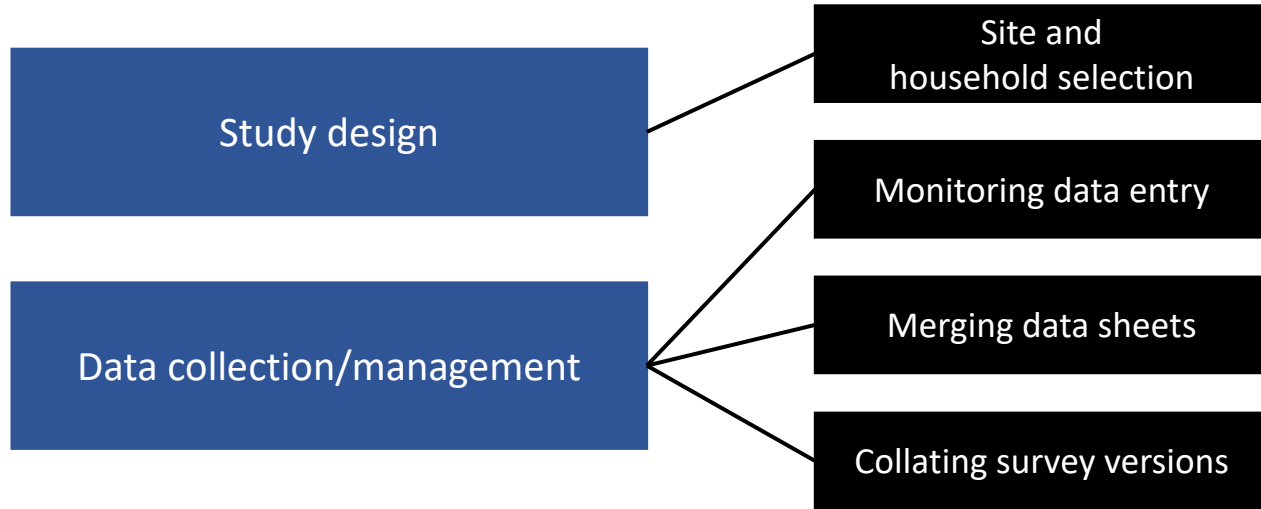


My strategy:

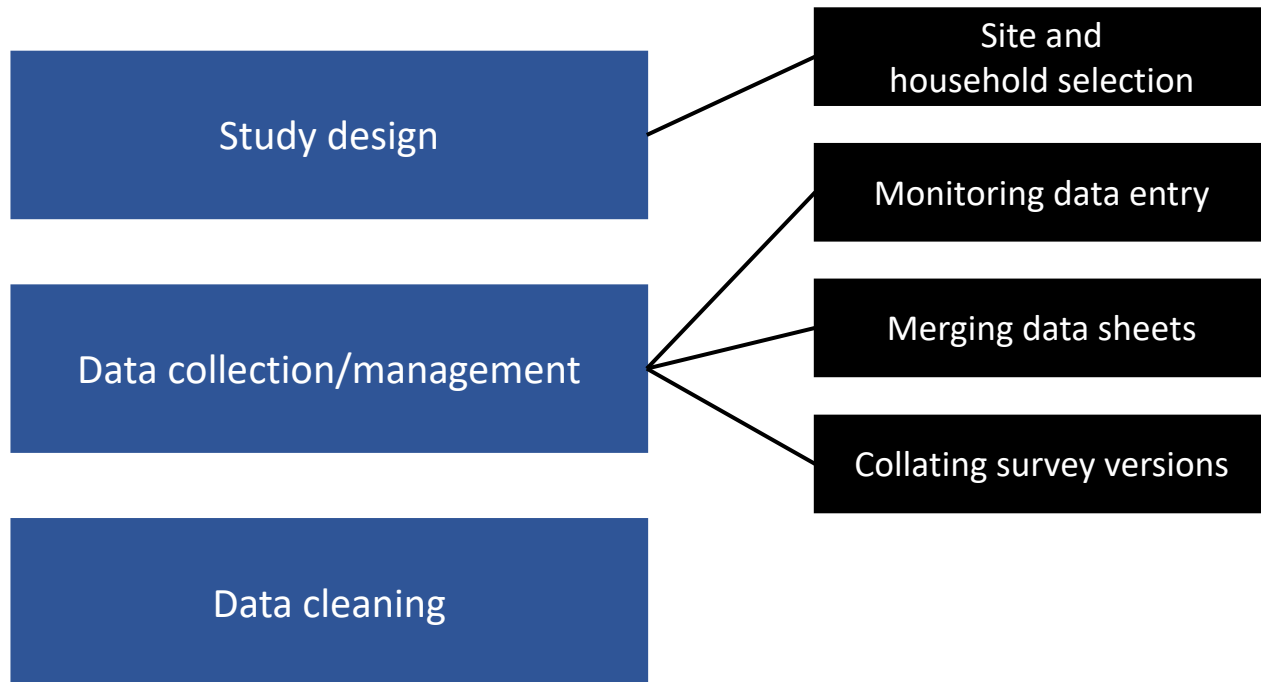
Step 1) Mangataka fanampiana amin'ny Amy Wesolowski sy Amy Winter ary Jessica Metcalf (satria manan-tsaina be izy telo)

Step 2) "Misaotra betsaka"

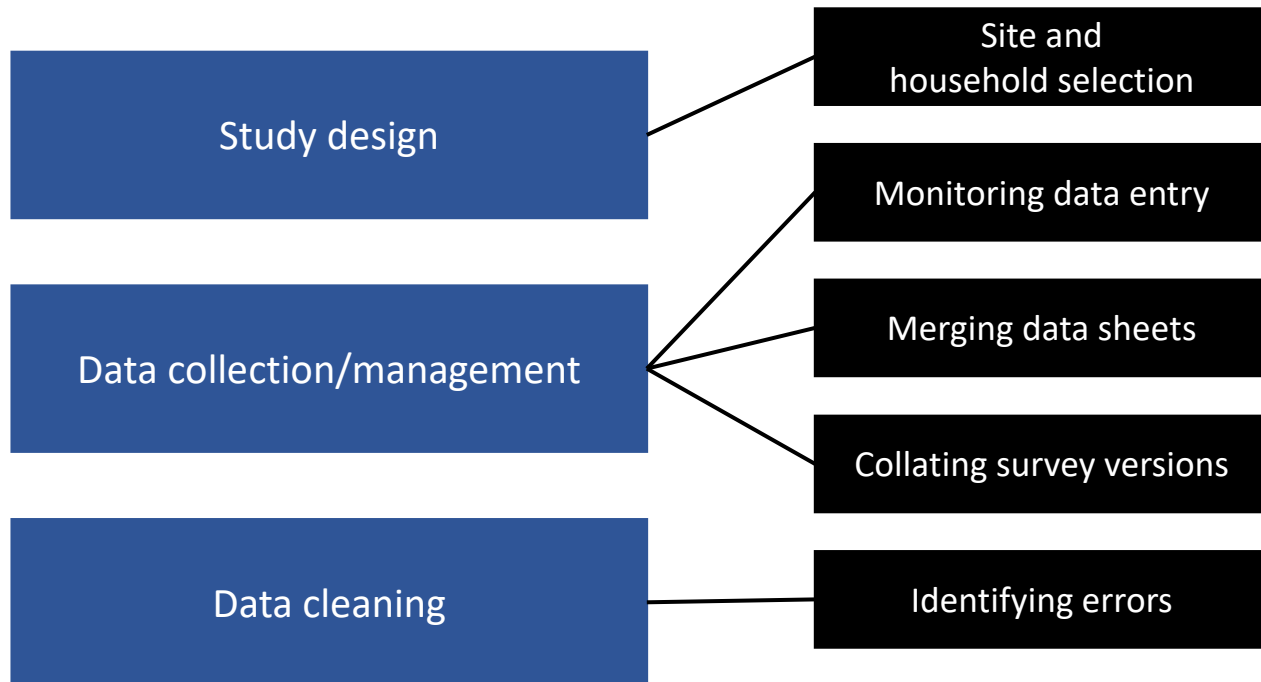
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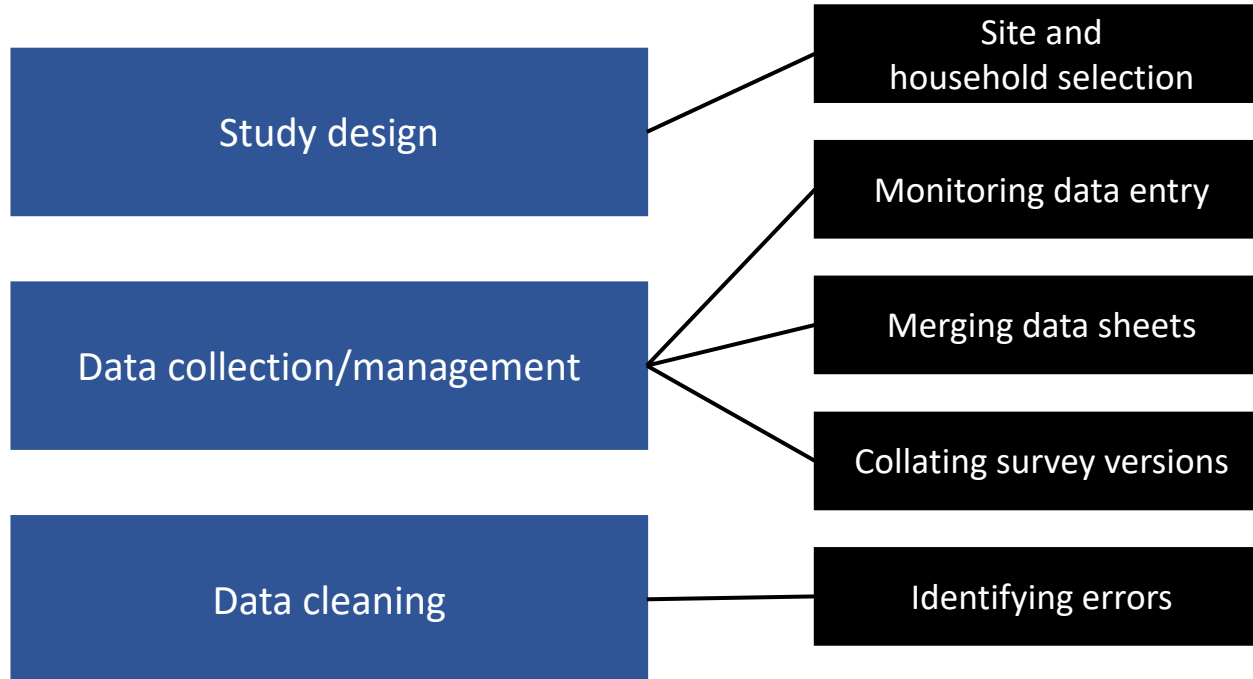
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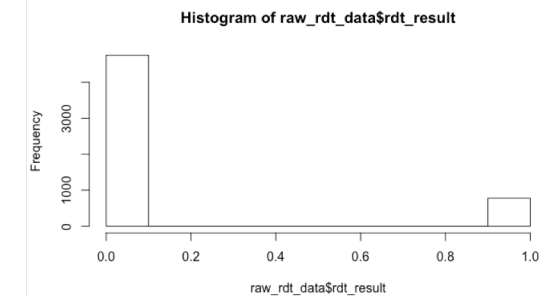
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Using R in the MAHERY-CRS Project

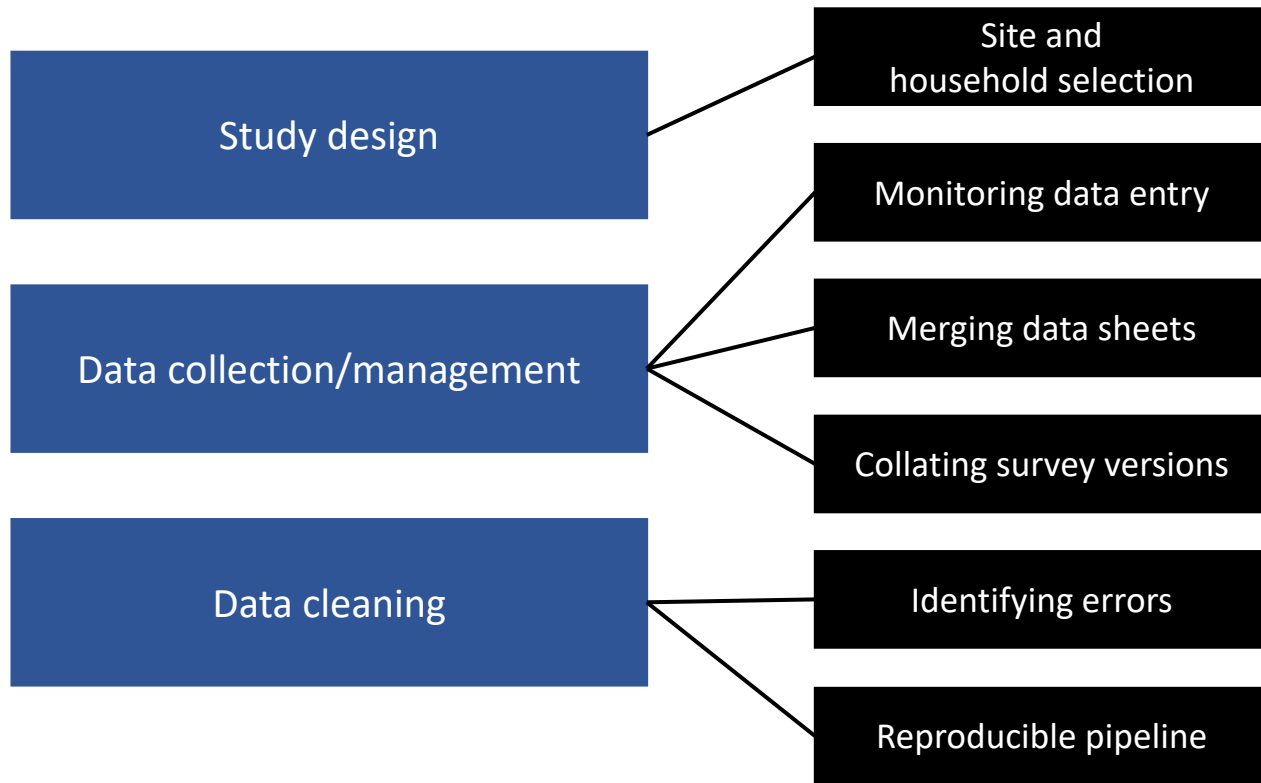


```
#####  
## Reproducible pipeline  
#####  
  
#Recode rdt_result:  
#(1) trim out invalid RDT result individuals  
#(2) n = N = negative = 0; pan or panpf = 1  
  
#use subsetting to keep only the individuals with a valid RDT result  
raw_rdt_data <- raw_rdt_data[!raw_rdt_data$rdt_result == 'i',]  
  
#Check: use a simple barplot to see if "i"s have been removed  
plot_counts <- table(raw_rdt_data$rdt_result)  
barplot(plot_counts)  
  
#recode the rdt_results such that they are 1s (positive) or 0s (negatives)  
raw_rdt_data$rdt_result <- recode(  
  raw_rdt_data$rdt_result, N = 0, n = 0, pan = 1, panpf = 1)  
  
#Check: use a simple barplot to see if the 1, 0 recoding was correct  
plot_counts <- table(raw_rdt_data$rdt_result)  
barplot(plot_counts)  
  
#from str() we see that rdt results are stored as num even though just 1s and 0s.  
#Convert to integer:  
raw_rdt_data$rdt_result <- as.integer(raw_rdt_data$rdt_result)  
  
#check with a histogram  
hist(raw_rdt_data$rdt_result)  
#-----
```

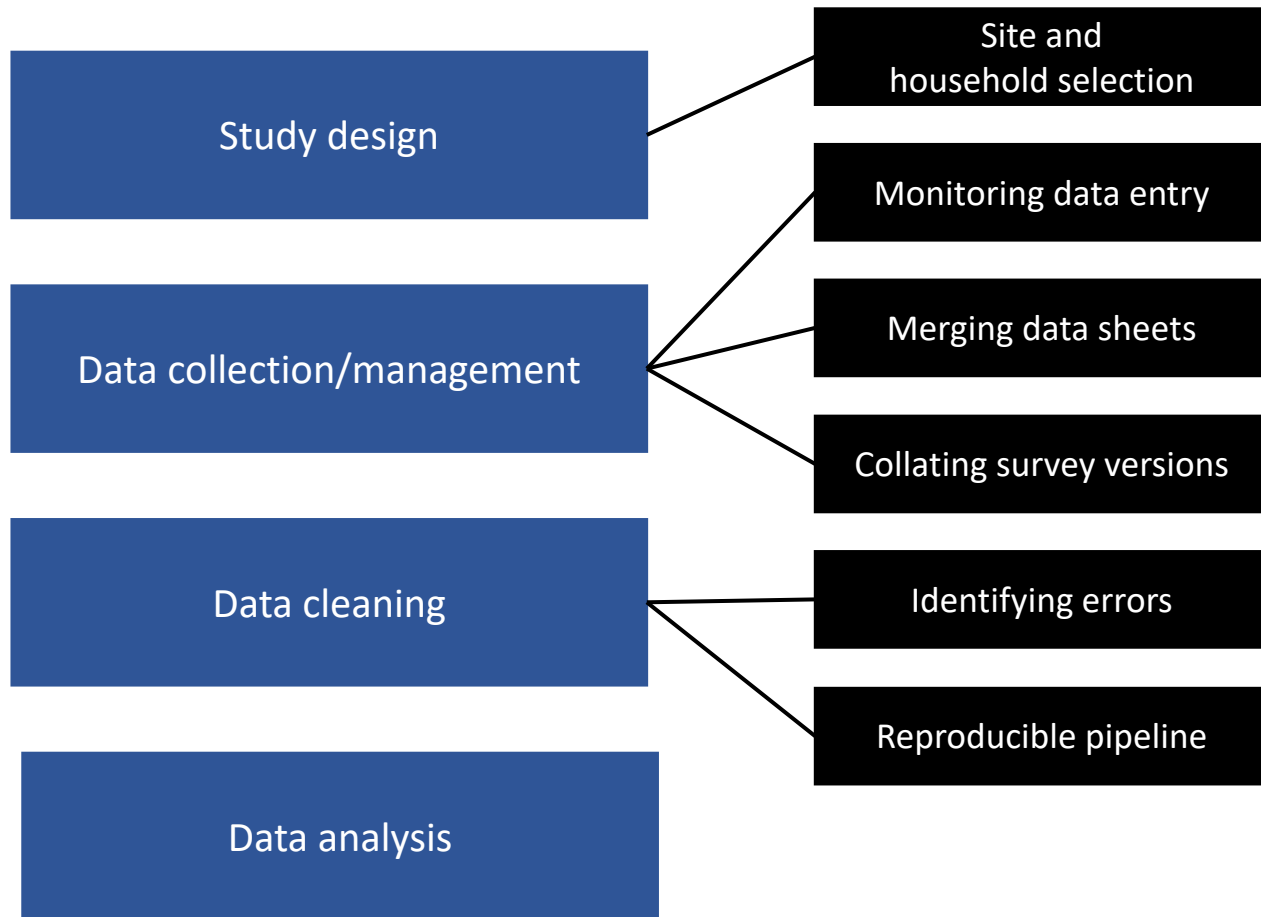


Using histograms to check for incorrect data values

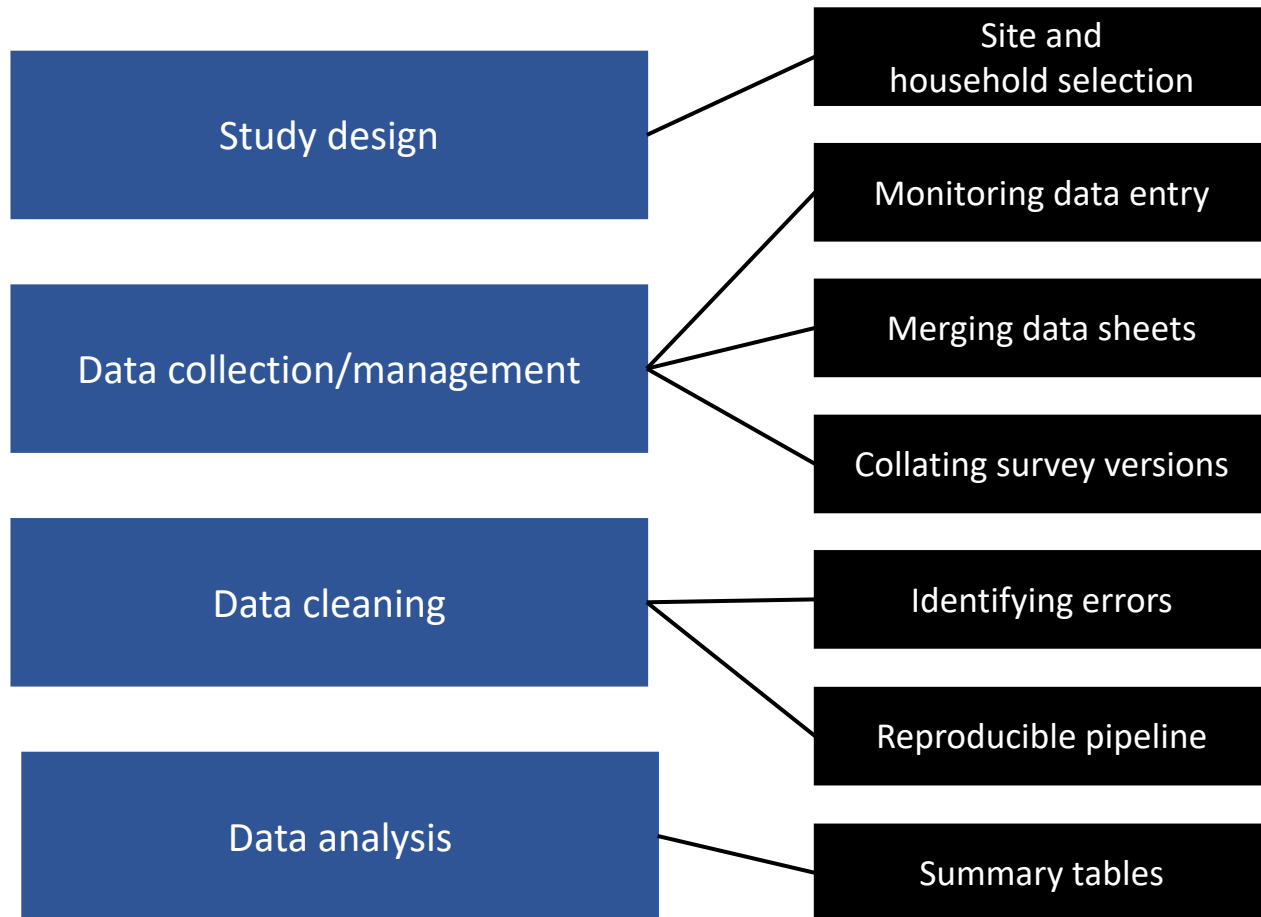
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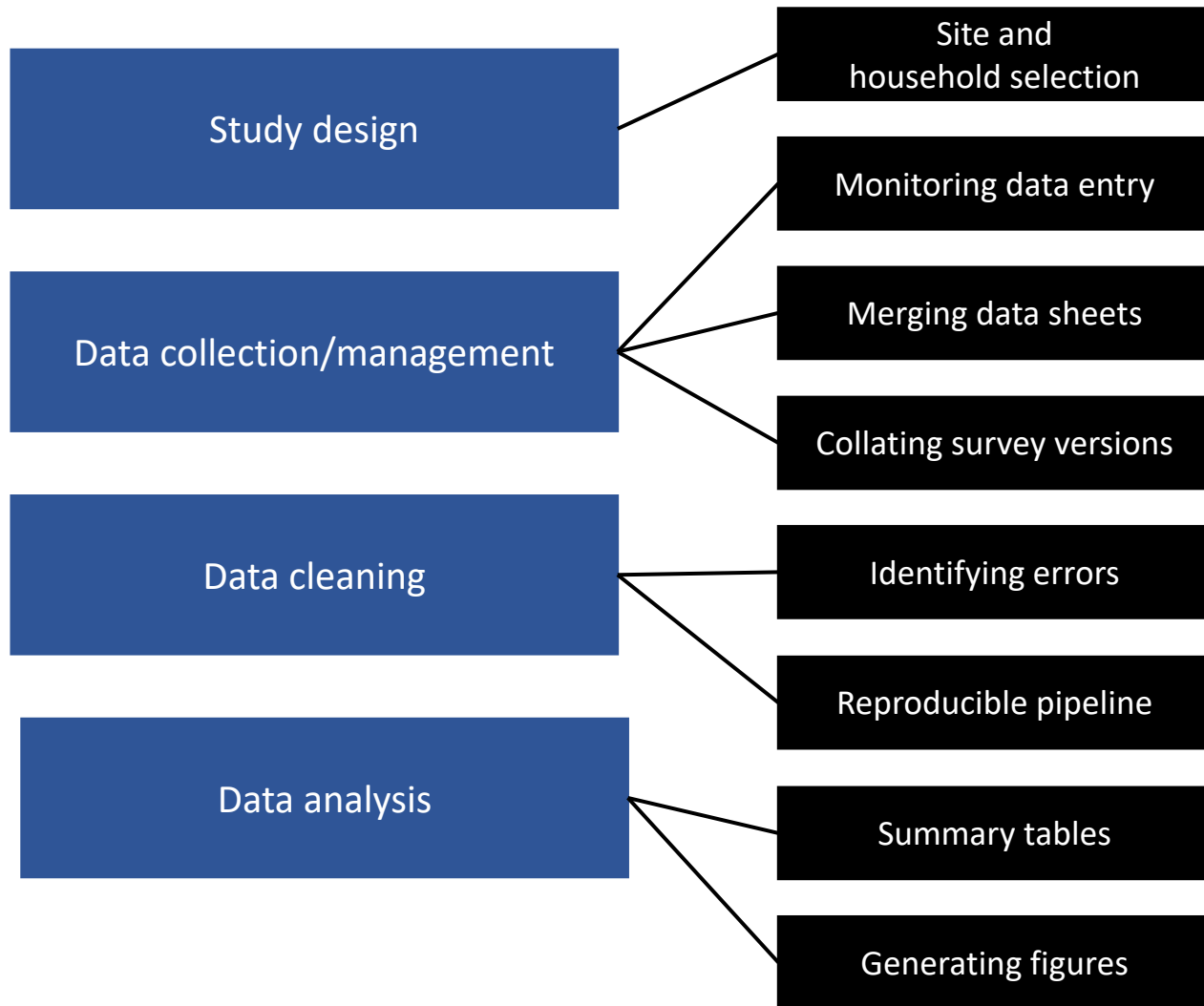
Using R in the MAHERY-CRS Project



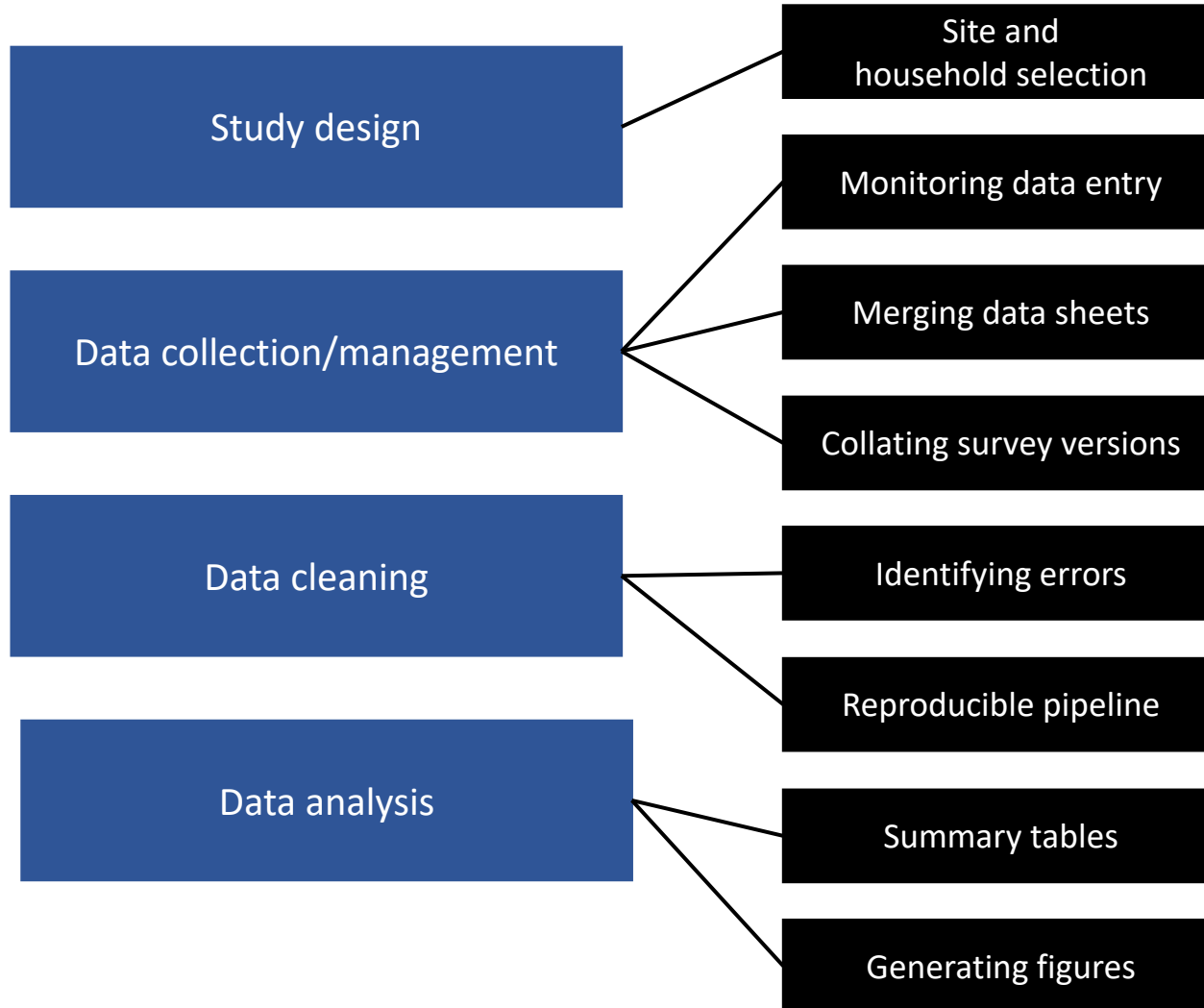
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Code to generate age-structure figures

```
#####  
## Generate figures: Age structure of malaria infections  
#####  
  
p.region2 <- ggplot(data = melted.plot_age_structure_region2,  
  aes(x=Age_bins,  
      y=ifelse(test = Sex == "Male",  
               yes = -value, no = value),  
      fill=variable,  
      color=variable,  
      alpha = variable)) +  
  geom_bar(stat="identity", position = "identity") +  
  coord_flip() +  
  scale_colour_manual(values=c("dimgray", "tomato")) +  
  scale_fill_manual(values=c("dimgray", "tomato")) +  
  scale_alpha_manual(values=c(1, 0.2)) +  
  labs(y = "Frequency", x = "Age") +  
  scale_y_continuous(labels = abs, limits = 0.25 * c(-1,1))
```


Figure 3

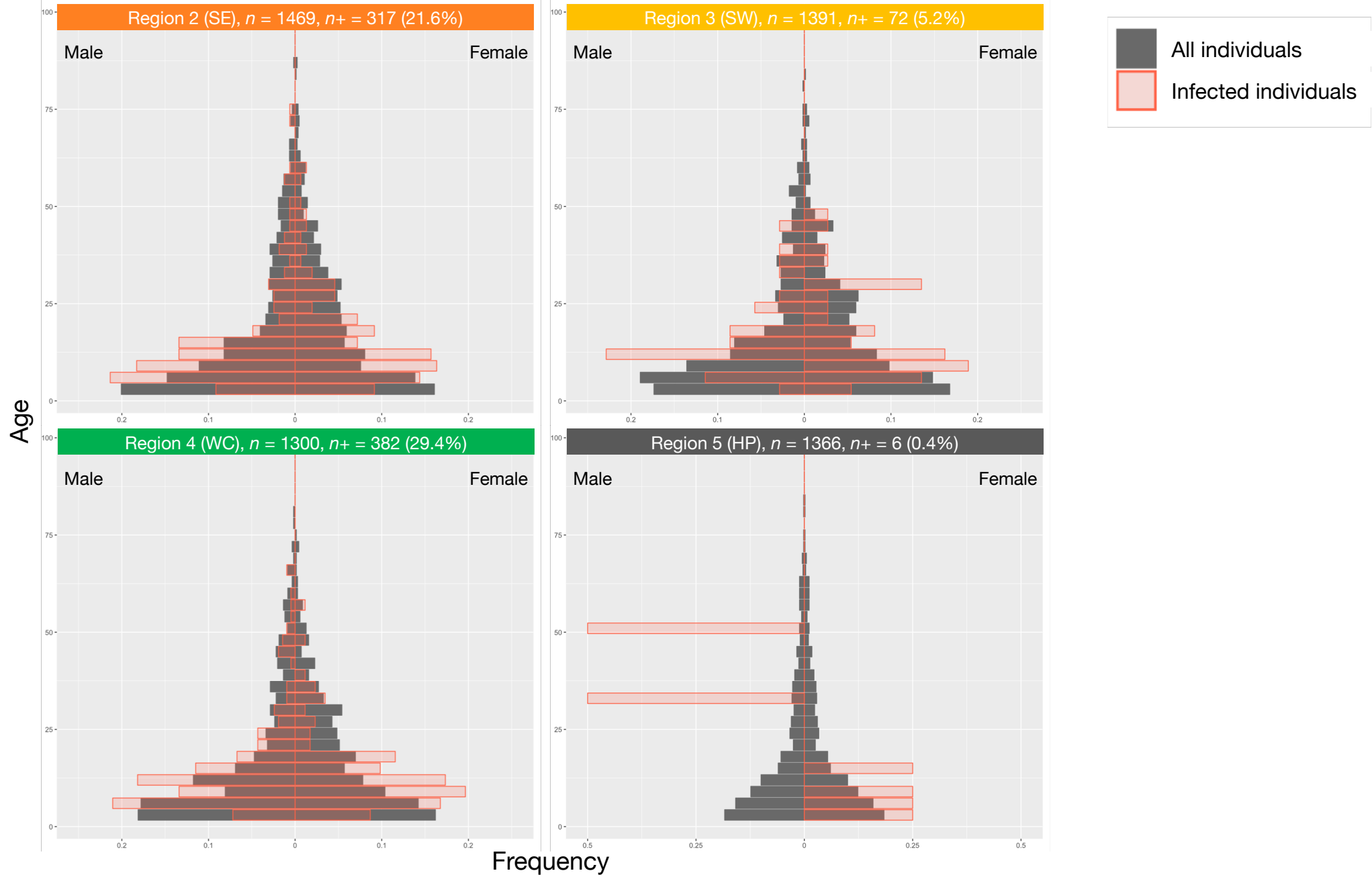
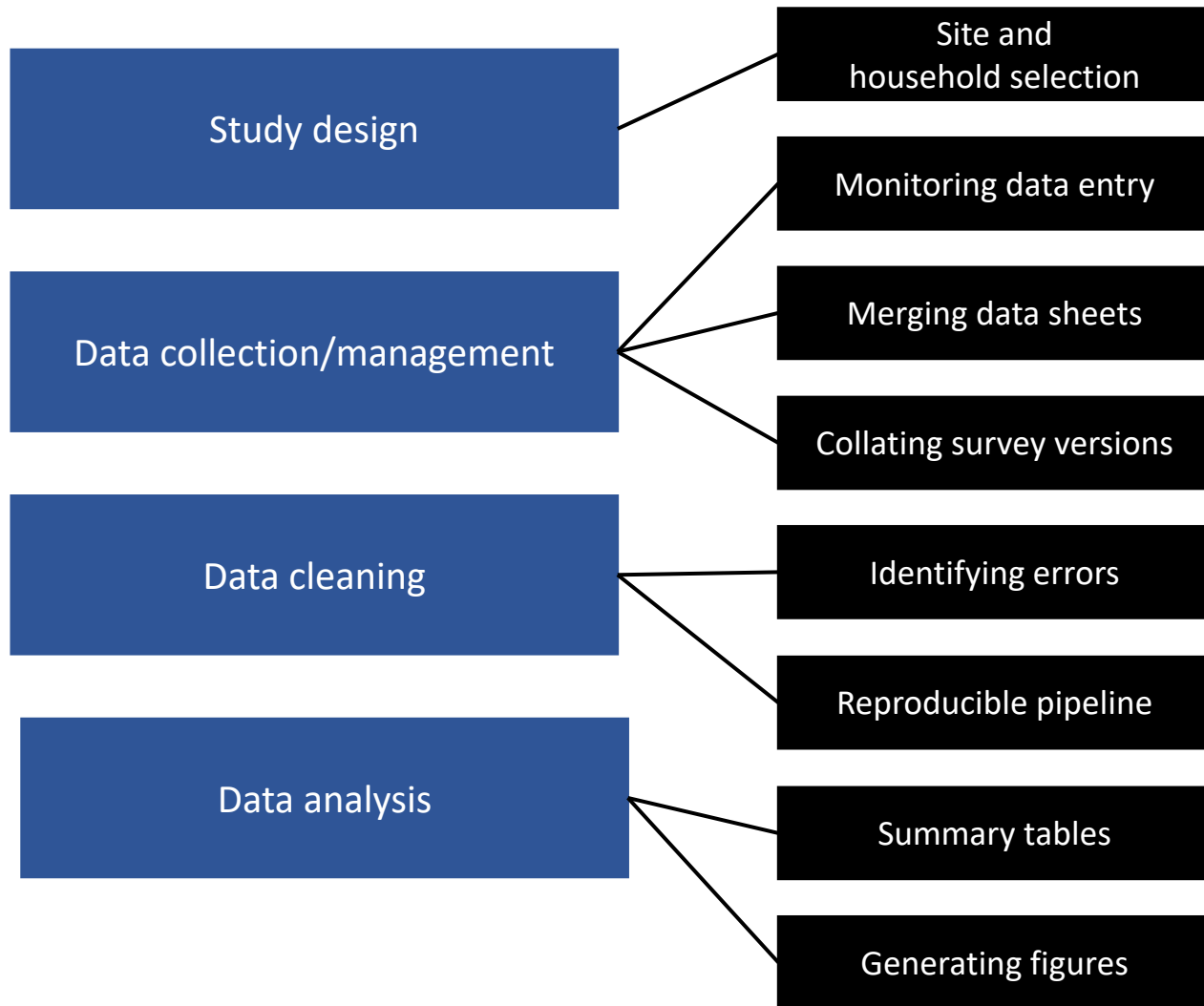
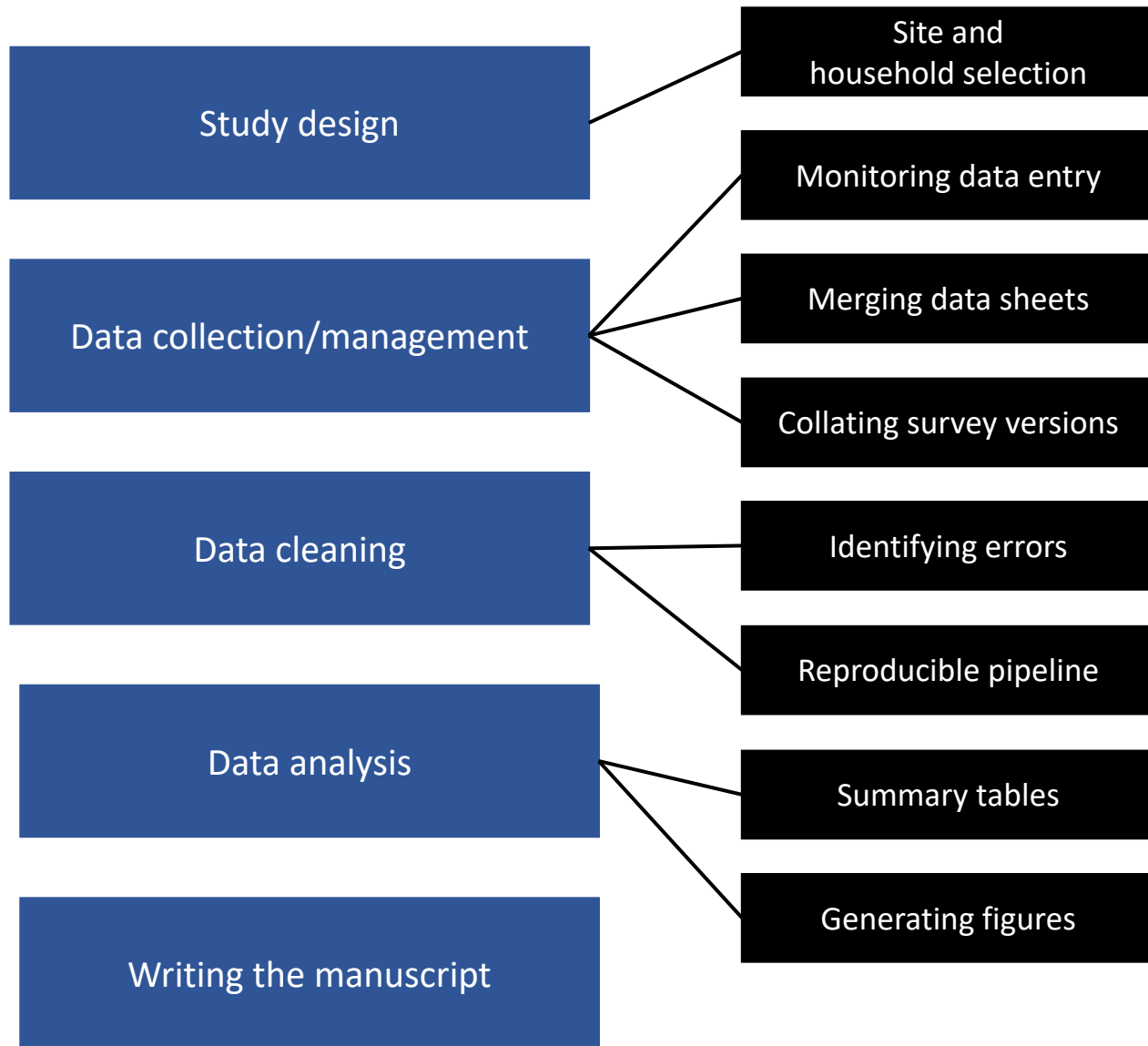


Figure 3: Age distribution of all individuals (gray) and infected individuals (red) shown by region. The proportion of individuals within a 3 year age bin is shown mirrored by sex with males are on the left and females on the right. Sample size, n , and the number of individuals positive, n_+ , are shown for each region (southeast, SE; southwest, SW; west coast, WC; high plateau, HP).

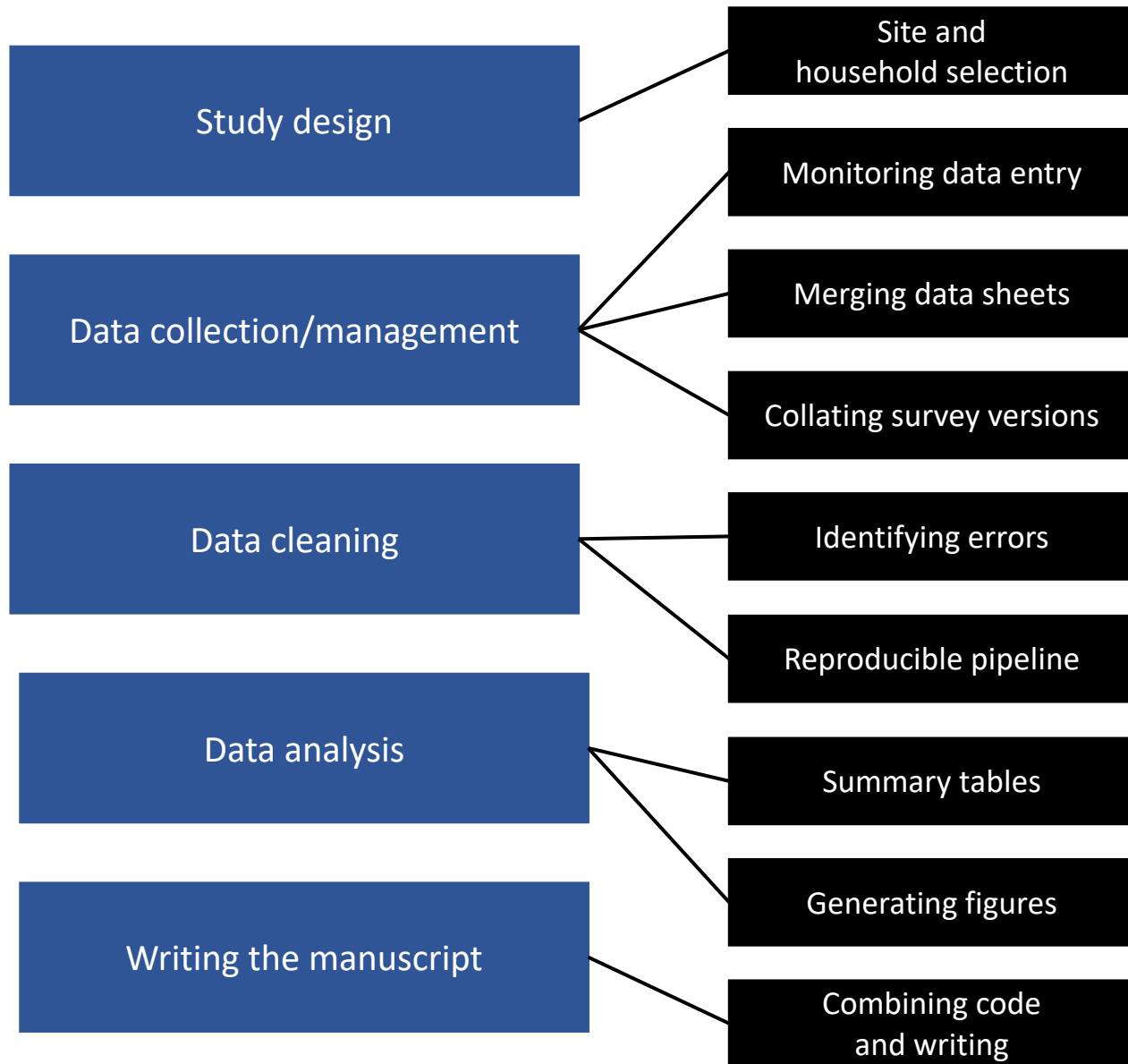
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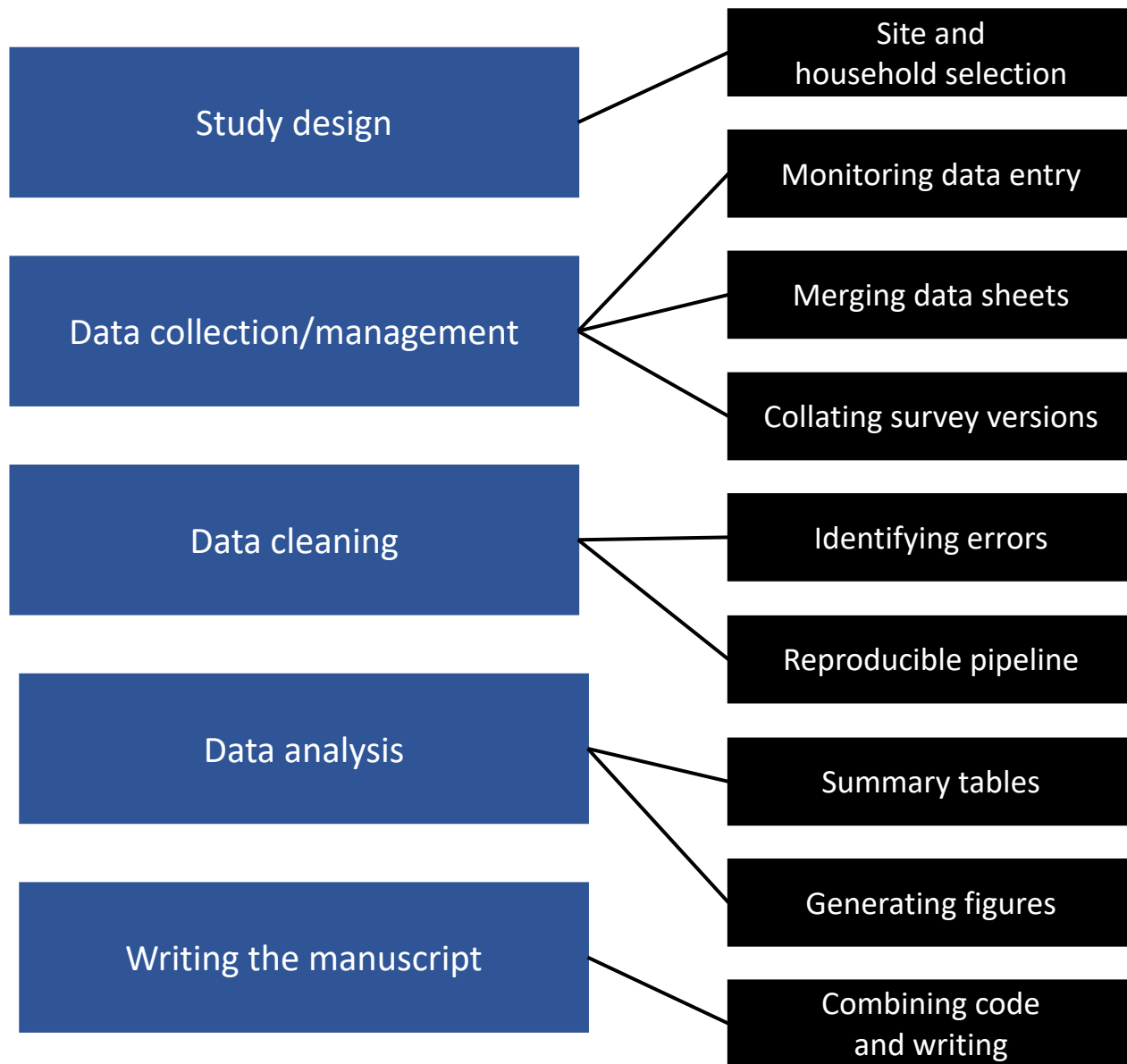
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R Markdown
<https://rmarkdown.rstudio.com/>

How can R help at every stage from day zero to the day of publication?

(Amin'ny 'study', manomboka andro voalohany hatramin'ny andron'ny 'publication':

Mora mora kokoa ve raha miasa amin'ny R?)