

# Host heterogeneity and SARS-CoV-2: from vaccine inequity to variants of concern

Undergraduate Research Symposium (URES)

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# Overview

## Sub-national SARS-CoV-2 vaccine uptake inequalities and their consequences for disease transmission

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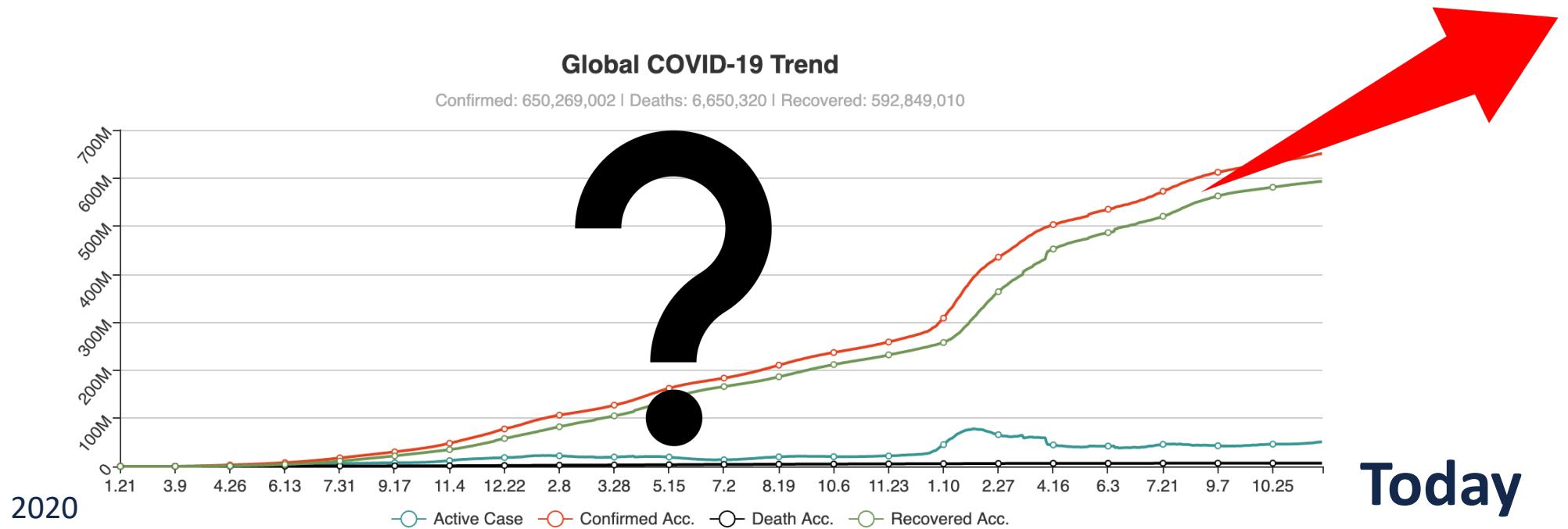
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<sup>5</sup>*Department of Epidemiology, Harvard T.H. Chan School of Public Health, Boston, MA, USA*

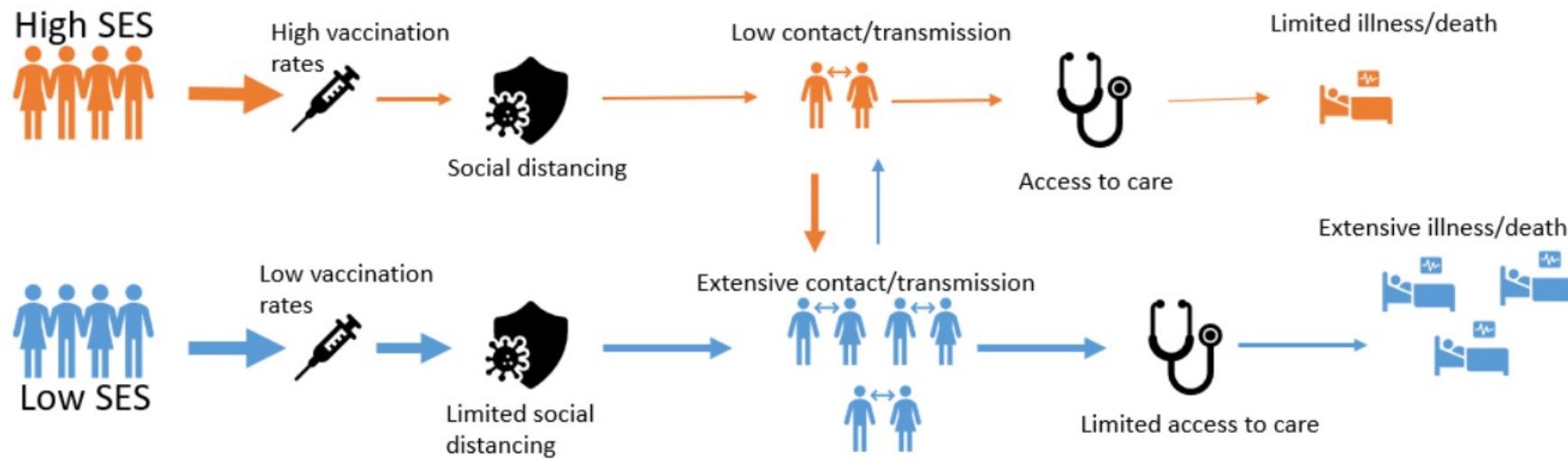
<sup>6</sup>*Department of Demography, University of California, Berkeley, CA, USA*

†*These authors contributed equally to this work*

# Motivation



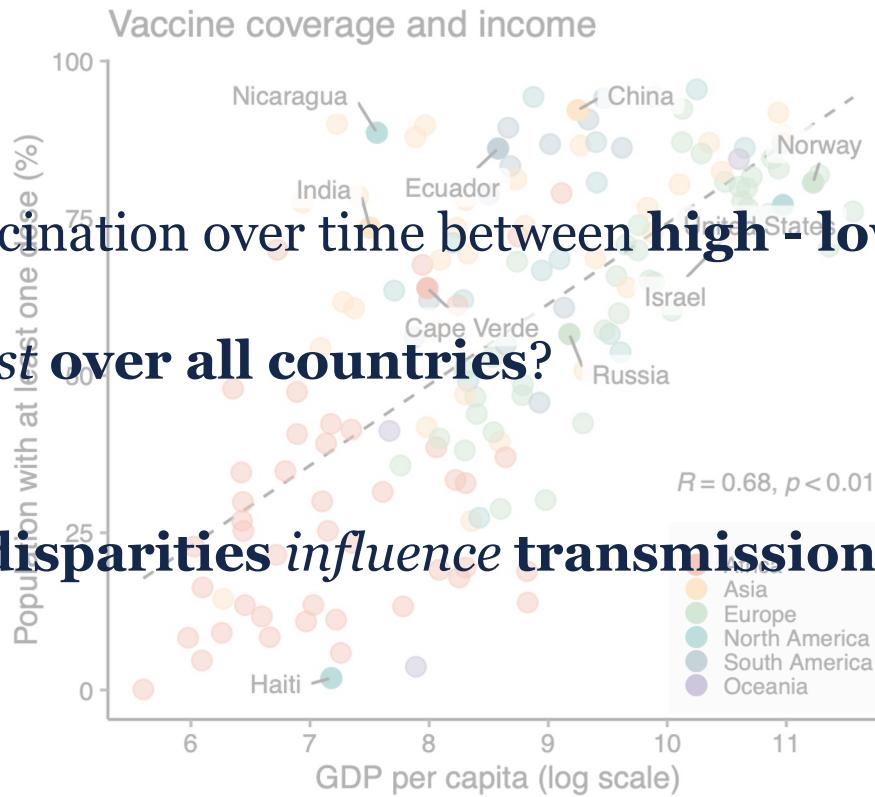
# Motivation



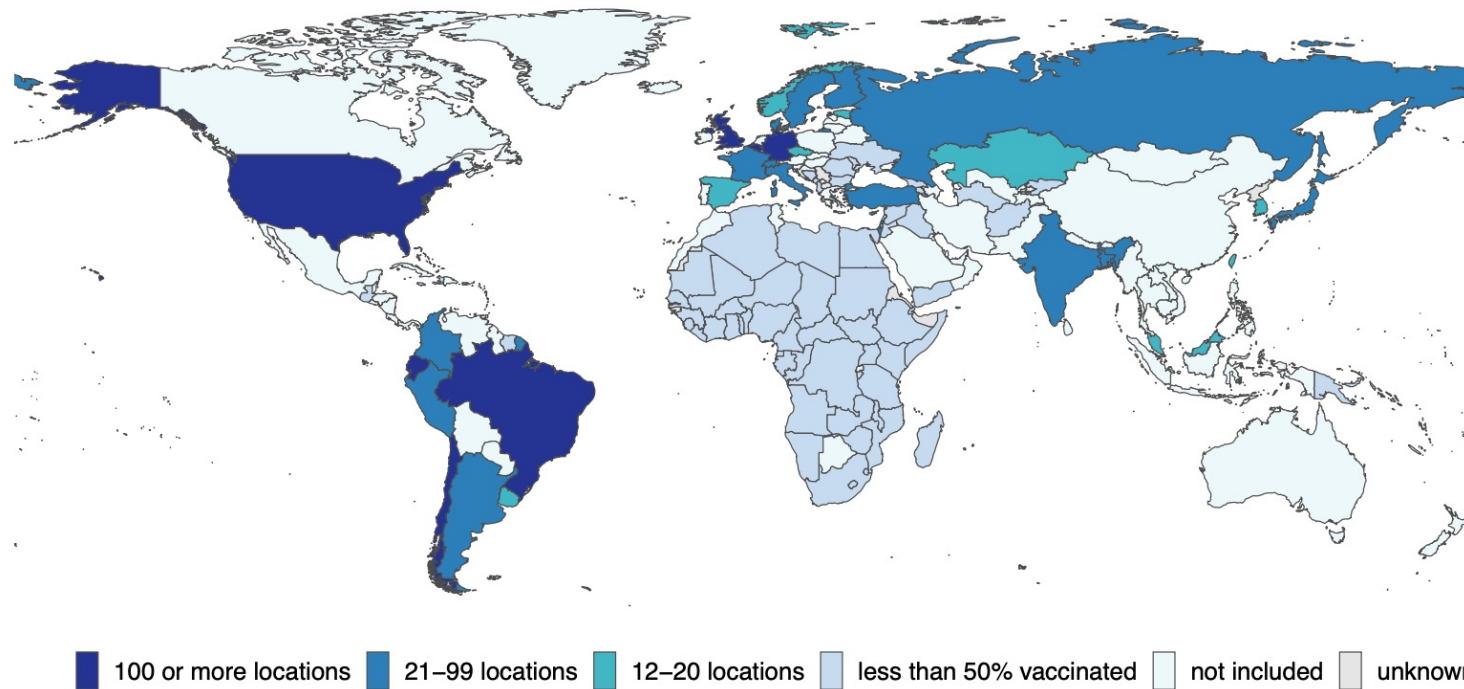
Socioeconomic (**SES**) : relating to or concerned with the interaction of **social** and **economic** factors

# Questions

- What are the **trends** of vaccination over time between **high - low SES groups** within countries?
  - Did these **trends persist over all countries?**
- How do these **trends and disparities influence transmissions** of SARS-CoV-2?



# Data Sources



# Data Required

country	resolution	location	pop	date	dose	vaccine	vnum	vcum	vperc
Italy	1	Abruzzo	1281012	2021-08-13	3 or booster	Janssen	0	0	0
Italy	1	Abruzzo	1281012	2021-08-05	partial	Janssen	5	0	0
Italy	1	Abruzzo	1281012	2021-08-05	full	Janssen	0	0	0
Italy	1	Abruzzo	1281012	2021-08-13	partial	Janssen	1	0	0
Italy	1	Abruzzo	1281012	2022-03-19	partial	Moderna	0	0	0
Italy	1	Abruzzo	1281012	2021-12-06	3 or booster	Pfizer	543	0	0
Italy	1	Abruzzo	1281012	2021-08-11	3 or booster	Moderna	0	0	0
Italy	1	Abruzzo	1281012	2021-12-12	partial	Moderna	8	0	0
Italy	1	Abruzzo	1281012	2021-06-30	partial	AstraZeneca	12	0	0

Vaccine Data

Italy	income2	gross disposable income
Republic of Korea	income2	Gross regional income per capita
Spain	income2	Disposable income per capita
UK	income2	GDHI local authority by ITL 1 region
UK	income3	Income - Proportion of LSOAs in poverty
UK	income4	Income deprivation rate
Brazil	income5	POPOULAÇÃO EM POBREZA EXTrema
Kenya	income5	Hardcore poverty (percentage)
Puerto Rico	income5	Median household income (in US dollars)
Republic of Korea	income5	household income
United States	income5	MEDHHINC_2019
France	income6	Poverty rate-Total
Kenya	income6	Overall Poverty (percentage)

SES Data

# Challenges

Data Aggregation

*Varying Structures*

*Incoherent Notations & Naming*

*Too Large to read*

# Methods

Data Aggregation

*Building Rubrics*

*Understanding Messages*

*Optimizing Performances*

# *Varying Structures*

	Covid-19, Dose 1, 85 og over, Troms og Finnmark	Covid-19, Dose 1, 85 og over, Trøndelag	Covid-19, Dose 1, 85 og over, Vestfold	Covid-19, Dose 1, 85 og over, Vestland
05.01.2021	33	221	125	237
06.01.2021	283	322	572	649
07.01.2021	87	254	237	382
08.01.2021	105	58	25	60
10.01.2021	0	0	0	0
11.01.2021	0	0	0	0
12.01.2021	10	102	136	301
13.01.2021	151	283	470	408
14.01.2021	73	209	166	371

# Building Rubrics

```
for (i in seq_along(df)){
  df[208+i] <- ifelse(grepl("Dose 1", colnames(df)[i]), "partial",
                      ifelse(grepl("Dose 2", colnames(df)[i]), "full", "3 or booster"))
  i = i+1
}

i=1
df1 <- county_vaccination

for (i in seq_along(df1)){
  df[416+i] <- ifelse(grepl("Pfizer", colnames(df1)[i]), "Pfizer",
                       ifel
germ2 <- read_csv("Desktop/research/germany/2022-04-14_Deutschland_Landkreise_COVID-19-Impfungen.csv",
                  col_types = cols(Impfdatum = col_date(format = "%Y-%m-%d"),
                                   Impfschutz = col_character()))%>%
  rename(date = 1, location = 2, age = 3, dose = 4, vnum = 5)%>%
  filter(dose == 2)%>%
  group_by(date, location)%>%
  summarise(vnum = sum(vnum))%>%
  inner_join(pop)%>%
  mutate(dose = "full")

i = i+1
}

i=1
df2 <- county_vaccination

for (i in seq_along(df2)){
  df[624+i] <- ifelse(grepl("Agoer", colnames(df2)[i]), "Agoer",
                       ifelse(grepl("oppgett", colnames(df2)[i]), "Ikke oppgett",
                             ifelse(grepl("Innlandet", colnames(df2)[i]), "Hedmark og Oppland",
                                   ifelse(grepl("Romsdal", colnames(df2)[i]), "Møre og Romsdal",
                                         ifelse(grepl("Nordland", colnames(df2)[i]), "Nordland",
                                               ifelse(grepl("Oslo", colnames(df2)[i]), "Oslo (f)",
                                                 ifelse(grepl("Rogaland", colnames(df2)[i]), "Rogaland",
                                                       ifelse(grepl("Svalbard", colnames(df2)[i]), "Svalbard",
                                                         ifelse(grepl("Finnmark", colnames(df2)[i]), "Troms og Finnmark",
                                                               ifelse(grepl("Trøndelag", colnames(df2)[i]), "Trøndelag",
                                                                 ifelse(grepl("Telemark", colnames(df2)[i]), "Vestfold og Telemark",
                                                                   ifelse(grepl("Vestland", colnames(df2)[i]), "Vestland",
                                                                     "Viken)))))))))))))))
```

# *Incoherent Notations & Naming*

CODGEO	NBMENFISC18	NBPERSMENFISC18	MED18	PIMP18	TP6018	TP60AGE118	TP60AGE218
1	258018	618263	23420	53.6	10.3	14.9	11.8
2	221370	512838	19690	43.6	18.4	29.7	22.8
3	157528	325813	20360	44	15.5	25.3	19.7
4	73620	154535	20580	45.6	16.8	24.9	20.9
5	63474	133391	20880	47.5	13.9	19.9	16.9
6	520415	1099477	22140	55.2	15.8	21.7	17.6

SES Data

Impfdatum	LandkreisId_Impfort	Altersgruppe	Impfschutz	Anzahl
2020-12-27	1003	18-59	1	9
2020-12-27	1003	60	1	19
2020-12-27	1053	18-59	1	13
2020-12-27	1053	60	1	200
2020-12-27	1055	18-59	1	20
2020-12-27	1055	60	1	66
2020-12-27	1056	18-59	1	65
2020-12-27	1056	60	1	7
2020-12-27	1057	18-59	1	12

Vaccine Data

# *Understanding Messages*

CODGEO	NBMENFISC18	NBPERSMENFISC18	MED18	PIMP18	TP6018	TP60AGE118	TP60AGE218
1	258018	618263	23420	53.6	10.3	14.9	11.8
2	221370	512838	19690	43.6	18.4	29.7	22.8
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6	520415	1099477	22140	55.2	15.8	21.7	17.6

SES Data

name	state	wdId	idIBGE
Abadia de Goiás	GO	Q304652	5200050
Abadia dos Dourados	MG	Q582223	3100104
Abadiânia	GO	Q304716	5200100
Abaeté	MG	Q1615444	3100203
Abaetetuba	PA	Q1615298	1500107
Abaiara	CE	Q1796441	2300101
Abaíra	BA	Q1645077	2900108
Abaré	BA	Q1647789	2900207
Abatiá	PR	Q784855	4100103
Abdon Batista	SC	Q1637414	4200051
Abel Figueiredo	PA	Q942803	1500131

Vaccine Data

# *Too Large to read*

 SP_2.csv	16.7 GB
 SP_3.csv	16.7 GB
 SP_1.csv	16.7 GB
 Parana_1.csv	3.8 GB
 Parana_3.csv	3.8 GB
 Parana_2.csv	3.8 GB

user	system	elapsed	Method
24.71	0.15	25.42	read.csv (first time)
17.85	0.07	17.98	read.csv (second time)
10.20	0.03	10.32	Optimized read.table
3.12	0.01	3.22	fread
12.49	0.09	12.69	sqldf
10.21	0.47	10.73	sqldf on S0
10.85	0.10	10.99	ffdf

# Optimizing Performances

```
AC_1 <- read_delim("https://uofi.box.com/shared/static/9h2j58jzgqirxdp75azajhqqytlczcm.csv",
                     ";", escape_double = FALSE, col_types = cols(document_id = col_skip(),
                                                               paciente_id = col_skip()), trim_ws = TRUE)%>%
  select(c(24, 26, 27))

AC_2 <- read_delim("https://uofi.box.com/shared/static/y3dy2xlv911p23danwn3oqtah1cxp1pq.csv",
                     ";", escape_double = FALSE, col_types = cols(document_id = col_skip(),
                                                               paciente_id = col_skip()), trim_ws = TRUE)%>%
  select(c(24, 26, 27))

AC_3 <- read_delim("https://uofi.box.com/shared/static/hdgisfajmtnkda6vc31s39rvbx2hodon.csv",
                     ";", escape_double = FALSE, col_types = cols(document_id = col_skip(),
                                                               paciente_id = col_skip()), trim_ws = TRUE)%>%
  select(c(24, 26, 27))
```



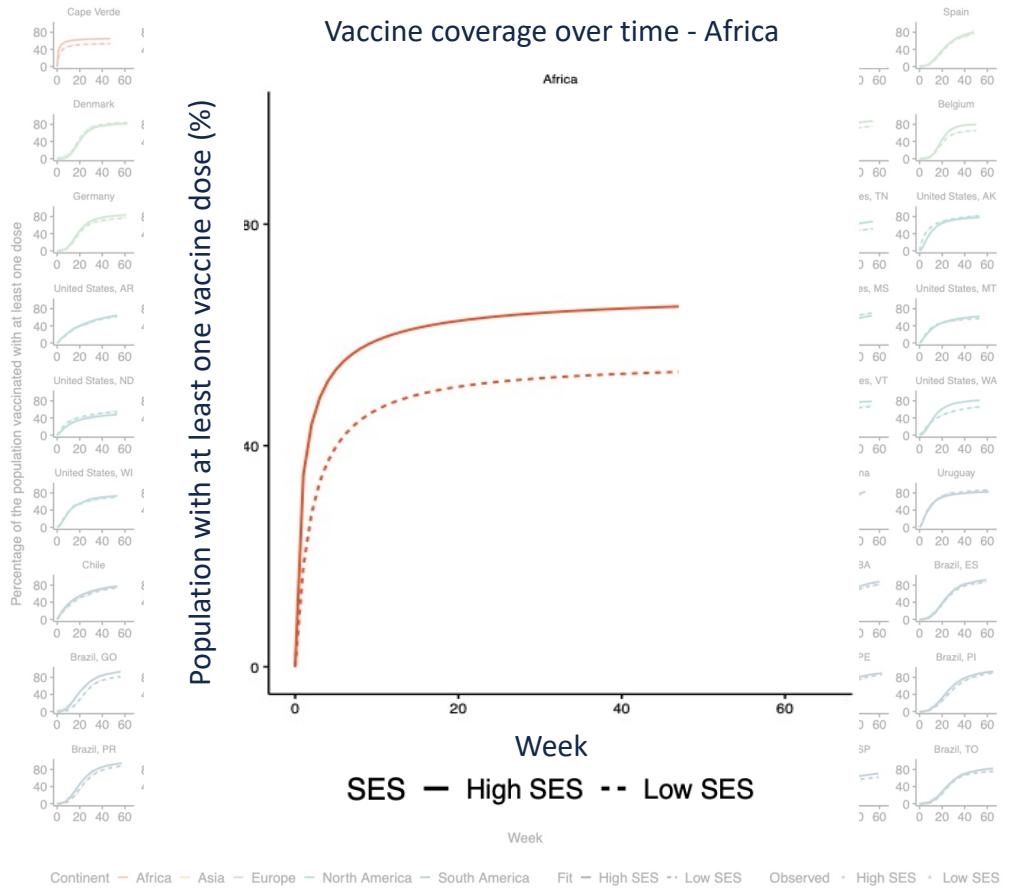
```
AC_1 <- fread("https://uofi.box.com/shared/static/9h2j58jzgqirxdp75azajhqqytlczcm.csv", select = c(
  "estabelecimento_municipio_codigo", "vacina_fabricante_nome", "vacina_dataAplicacao", "vacina_descricao_dose"
), sep = ";")

AC_2 <- fread("https://uofi.box.com/shared/static/y3dy2xlv911p23danwn3oqtah1cxp1pq.csv", select = c(
  "estabelecimento_municipio_codigo", "vacina_fabricante_nome", "vacina_dataAplicacao", "vacina_descricao_dose"
), sep = ";")

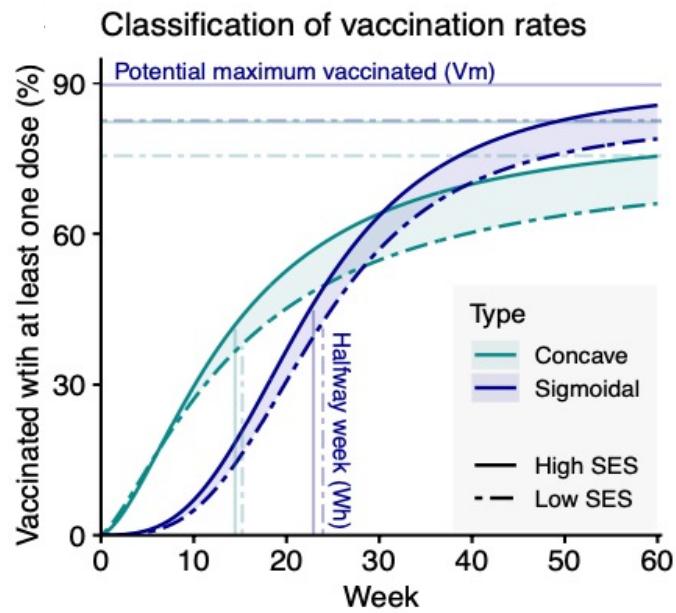
AC_3 <- fread("https://uofi.box.com/shared/static/hdgisfajmtnkda6vc31s39rvbx2hodon.csv", select = c(
  "estabelecimento_municipio_codigo", "vacina_fabricante_nome", "vacina_dataAplicacao", "vacina_descricao_dose"
), sep = ";")
```

BR_Set_1_MG.csv
BR_Set_1_MS.csv
BR_Set_1_MT.csv
BR_Set_1_PA.csv
BR_Set_1_PB.csv
BR_Set_1_PE.csv
BR_Set_1_Pl.csv
BR_Set_1_PR.csv
BR_Set_1_RJ.csv
BR_Set_1_RJ1.csv
BR_Set_1_RJ2.csv
BR_Set_1_RN.csv
BR_Set_1_RO.csv
BR_Set_1_RR.csv
BR_Set_1_RS.csv
BR_Set_1_SC.csv
BR_Set_1_SE.csv
BR_Set_1_SP.csv
BR_Set_1_TO.csv

# Data Visualization



# Results



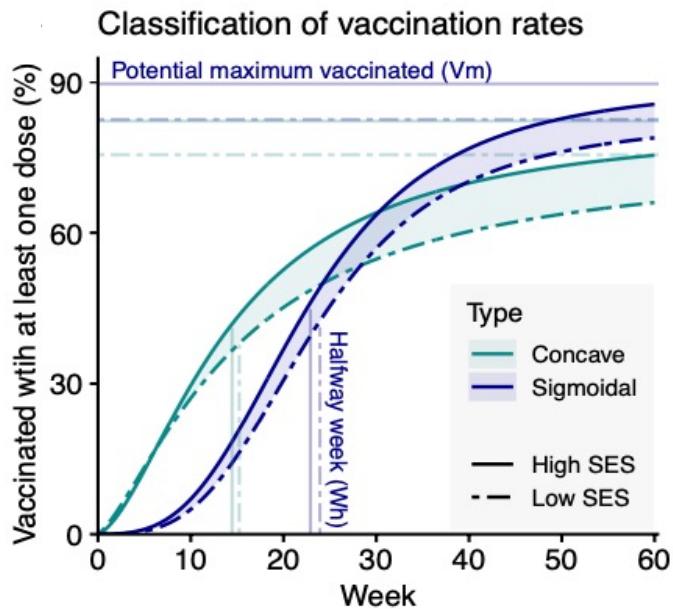
Concave strategy

A rapid initial roll-out  
Quickly reaching half of the vaccination potential

Sigmoidal strategy

Slow to begin  
Reaches a steady state more rapidly

# Results



## Concave strategy

A rapid initial roll-out  
Quickly reaching half of the vaccination potential

The **sigmoidal strategy** results in  
a **substantially higher burden** of disease

Speed of the initial roll-out > the level of coverage

Modifying the roll-out strategy for **low SES groups** for a reduction in deaths not attainable by modifying the high SES group alone



Thank you!  
Questions?