

Suppressing Covid-19: Public Health Policy and Effective Mass-Testing Rates

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Department of Engineering, U of Cambridge

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Joint work with



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Karolinska Institutet

Outline

△ Covid-19 tests

- ▷ Antibody tests
- ▷ PCR tests
- ▷ Antigen tests

~~> Rapid, cheap, at-home tests

△ Epidemiological models

- ▷ SIR models on random population networks
 - ▶ Erdős-Rényi graphs
 - ▶ Random graphs with given degree distribution
- ▷ SIR epidemics with mass testing

△ Necessary testing rates for suppression

- ▷ Rigorous results for a broad class of models
- ▷ Explicit numerical examples

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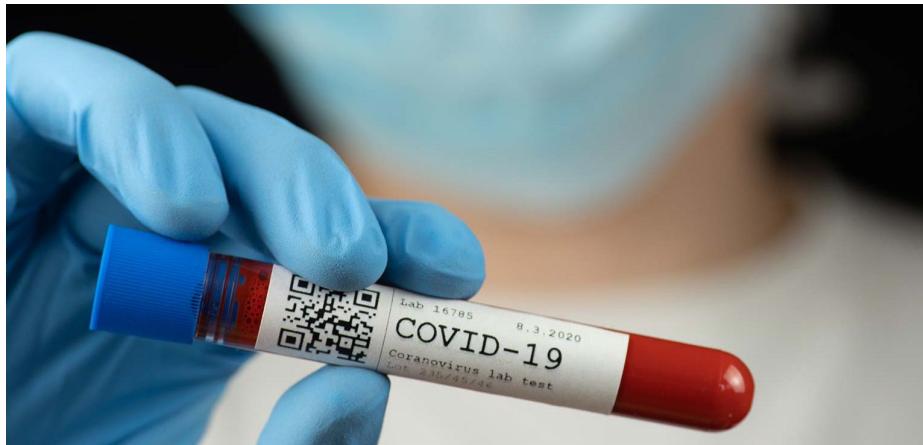
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Covid-19 testing: Why?



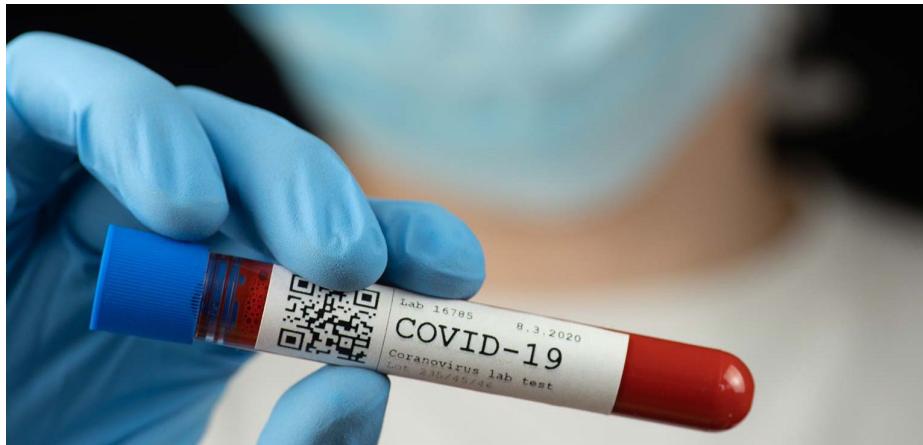
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Currently: ≈ 200 Covid-19 FDA-approved tests in the US
(+many more in development) using different technologies . . .

Antibody tests

Purpose

Detect previously infected
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⇒ Not relevant for our purposes



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Problems

- ▷ A lot of unknowns
 - ～ Do antibodies confer immunity? For how long?
 - ～ Relationship between antibody level (titer) and degree and persistence of immunity?
 - ～ Antibody test accuracy?
- ▷ More than 90 tests on the US market without FDA review
- ▷ Inappropriate use of results as “immunity passports”

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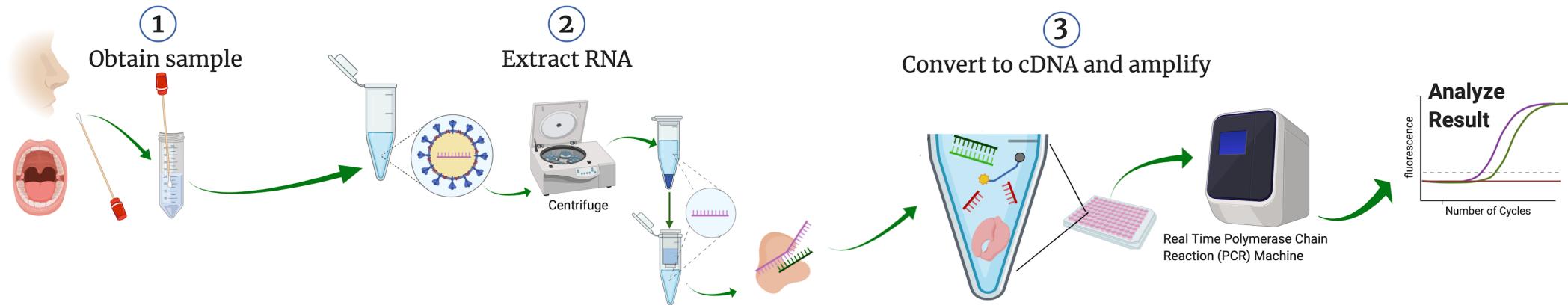
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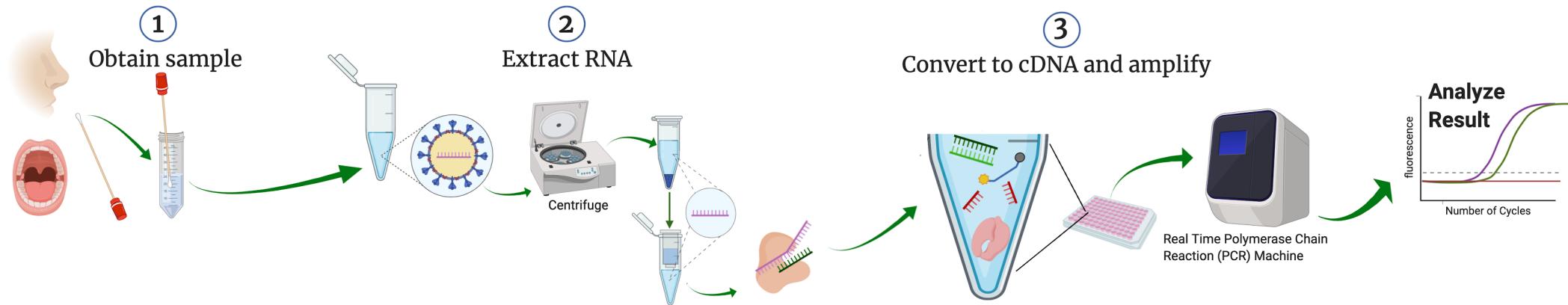
But

- ▷ Important and possibly becoming much more common soon

PCR tests [reverse-transcription polymerase chain reaction tests]

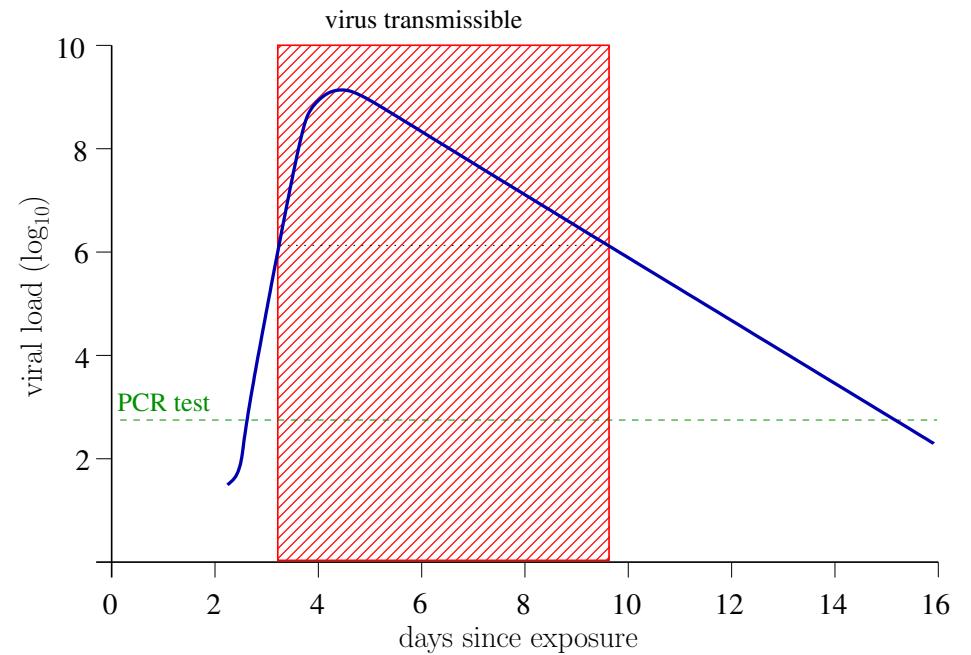


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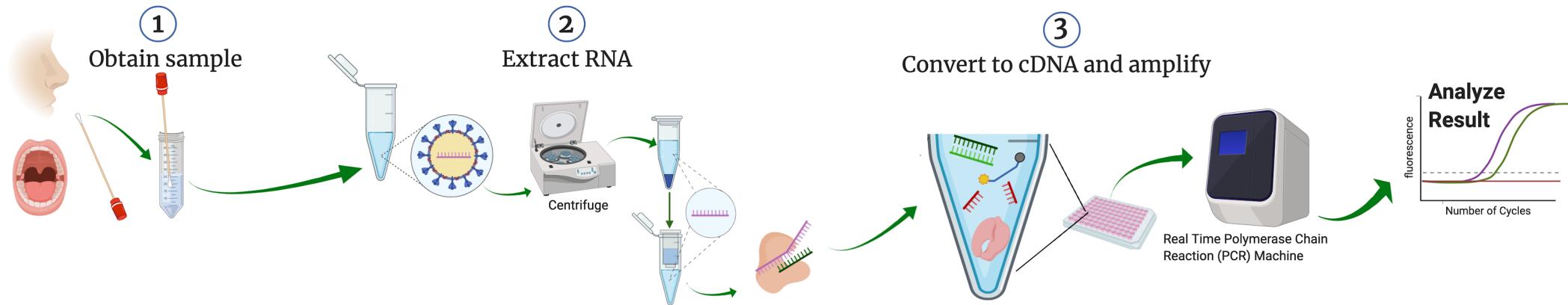


The gold standard

- ▷ They take only a few hours
- ▷ Detect as few as 100-1000 copies of viral RNA in 1ml of sample
- ▷ Sensitivity close to 100%



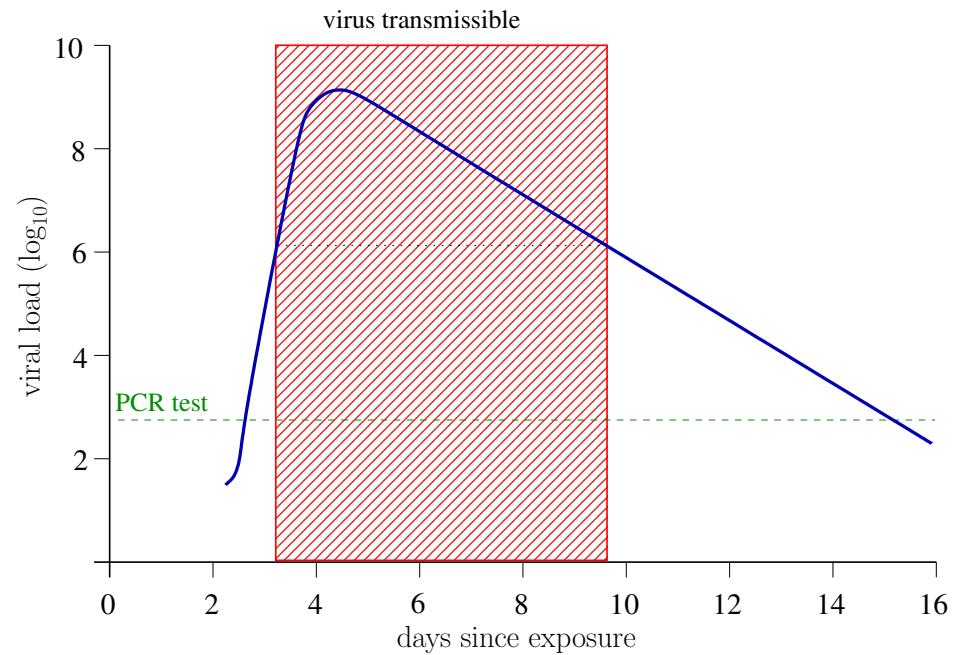
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- ▷ Indeed, **the** FDA test standard
- ▷ **But:** Not designed for an out-of-control pandemic



PCR tests

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- ▷ Very **slow** turnaround times (3-14 days in most places in the US)
~~> they miss the most infectious period
- ▷ **Not enough** of them: Bottlenecks:
Chemical reagents, lab supplies, PCR machines
- ▷ **Expensive** [\$35-200], very tightly regulated, require specialized personnel and equipment [⇒ social inequality issues]
 - ▷ [Ct values not reported]

PCR tests

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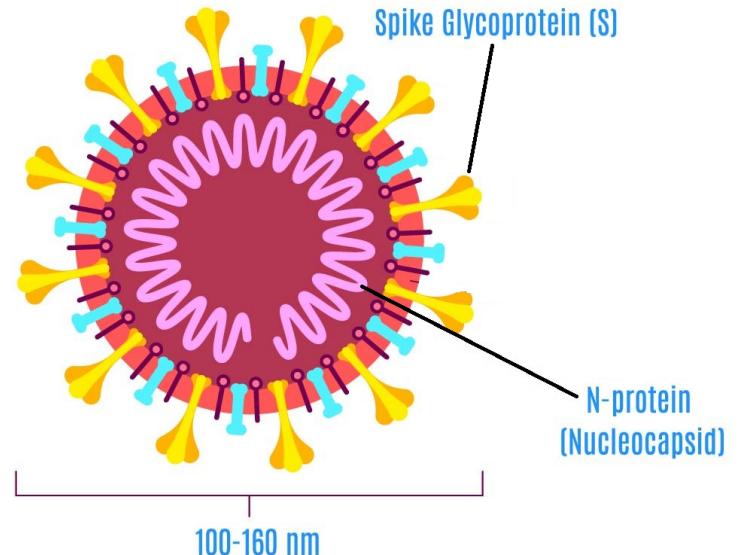
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Partial fixes

- ▷ Pooling or “**group testing**”: Still expensive and not fast enough
- ▷ Saliva-based tests with Ginkgo Bioworks’s **Illumina sequencing machines** instead of PCR. Factor of 6 faster, still slow:
Samples must be shipped centralized locations
- ▷ Saliva-based modified-PCR laboratory tests: UIUC’s **I-COVID**, Yale’s **SalivaDirect**. Results in 2-6 hours, cost \$10-20.
UIUC story highlights the need for even more, cheaper tests

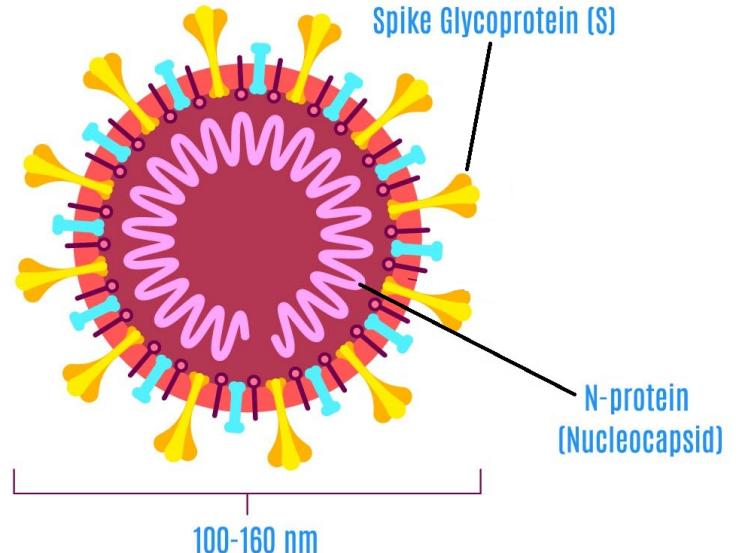
Antigen tests: I lab-based

They do not identify virus RNA
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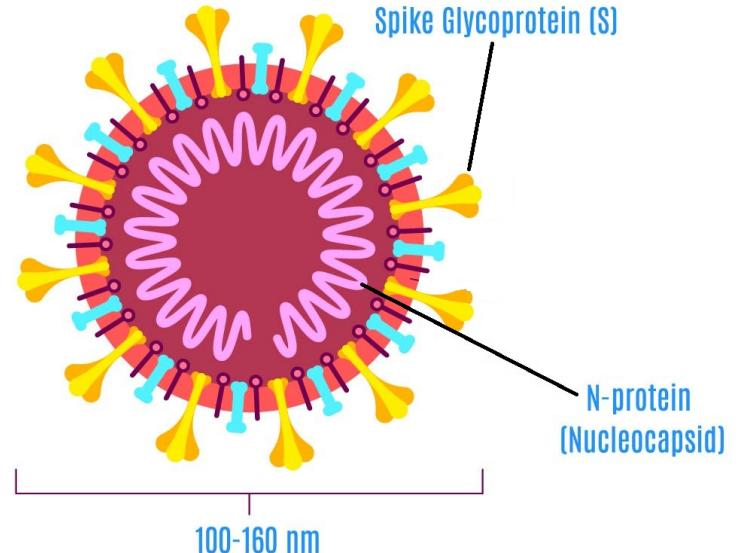
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- E.g., tests made by *Quidel* and *Becton-Dickinson* (US) detect the **nucleocapsid** (N) protein in nasal/throat swab samples
 - Tests cost is \approx half of PCR, give results in 15 mins
 - Can be administered at a point-of-care location
 - Will be making 14 million tests/month by end of September

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Drawbacks

- They only work with a proprietary reader but companies cannot produce it at same scale
- Since nucleocapsid is *inside* the virus they need reagents to break down its outer membrane

Antigen tests: II rapid tests

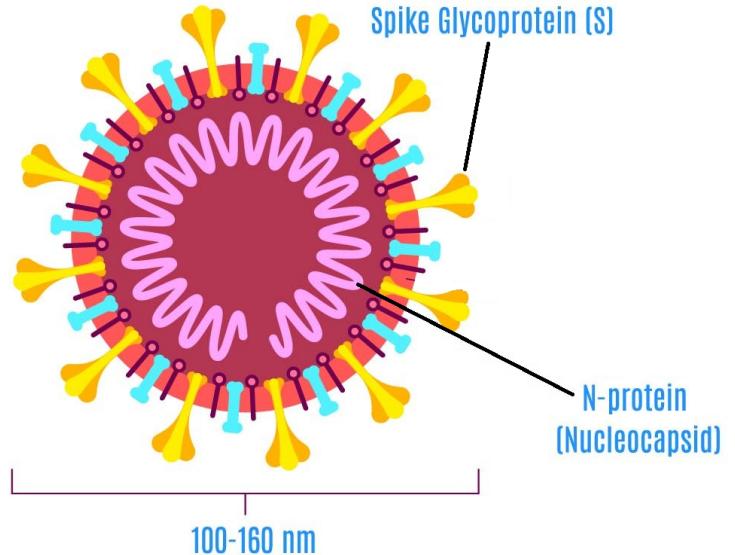
Made by US-based companies including
e25 Bio, Sona Nanotech, Iceni Diagnostics,
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Characteristics

Cheap: \$1-2

Results in 15 minutes

Home tests, no equipment: Saliva + saline solution + small cup



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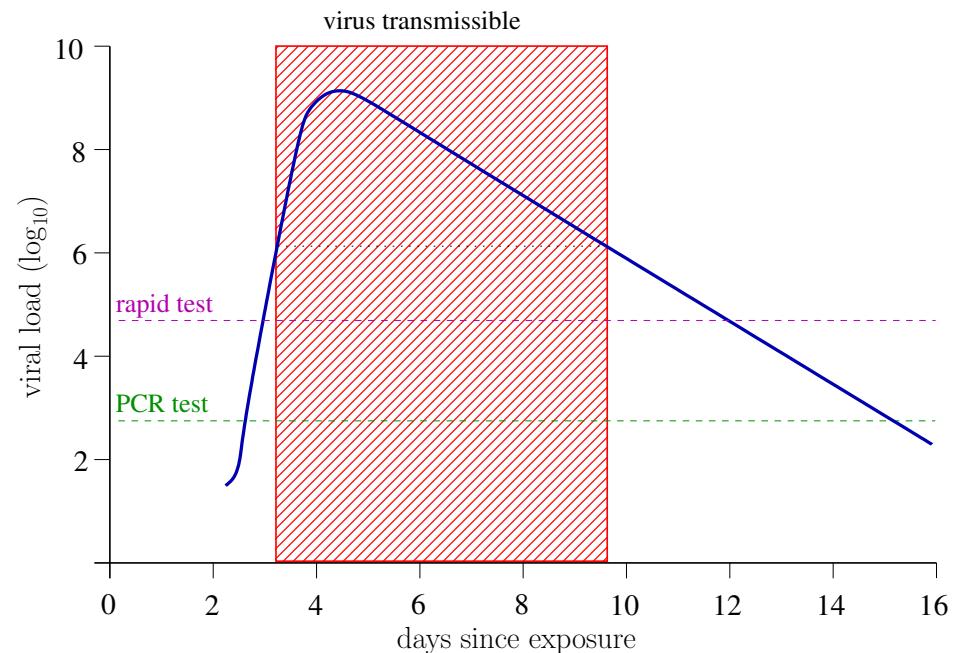
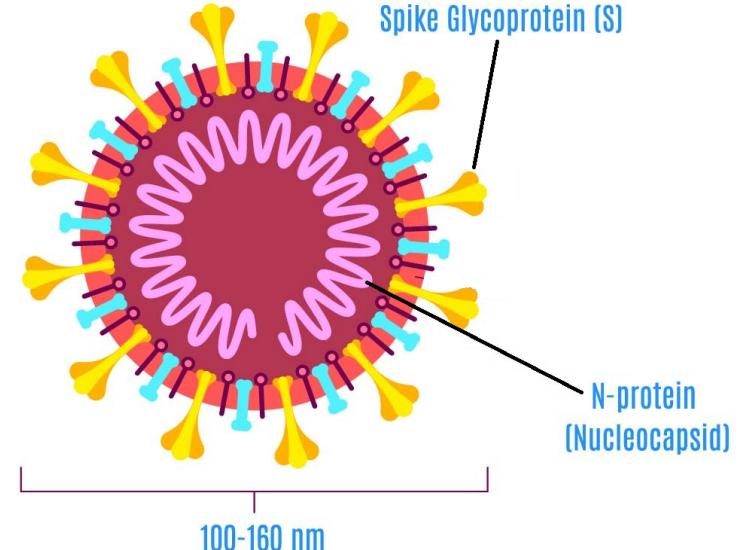
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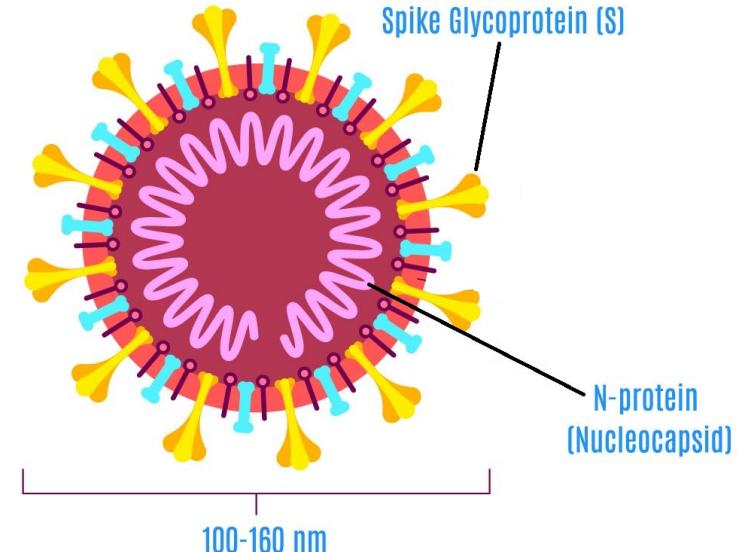
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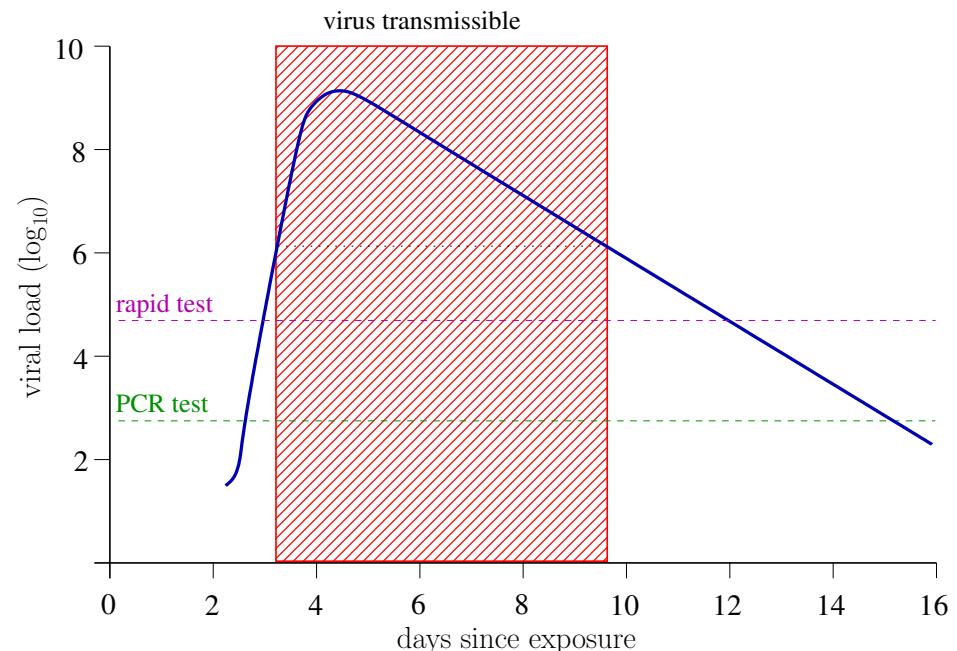


e25 Bio test

Paper strip no larger than 1x5 in

Looks for the **spike** (S) protein
on the *outside* of the virus

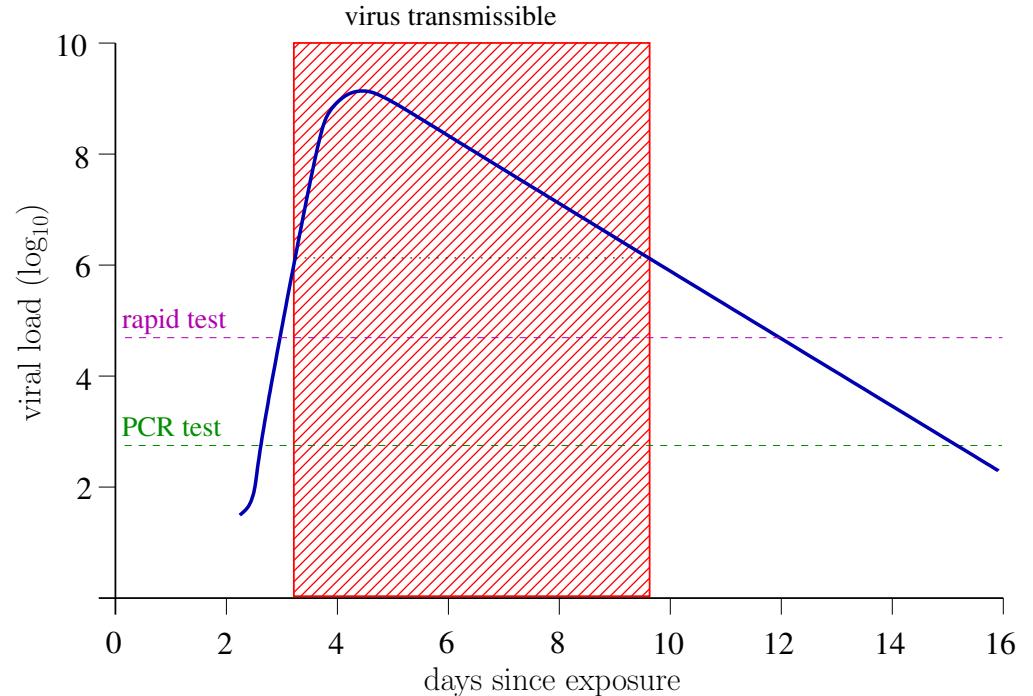
Sensitivity: 60-80%



PCR vs rapid tests

For \approx 24 hours in the beginning
the tests give different results,
rapid tests give false negatives

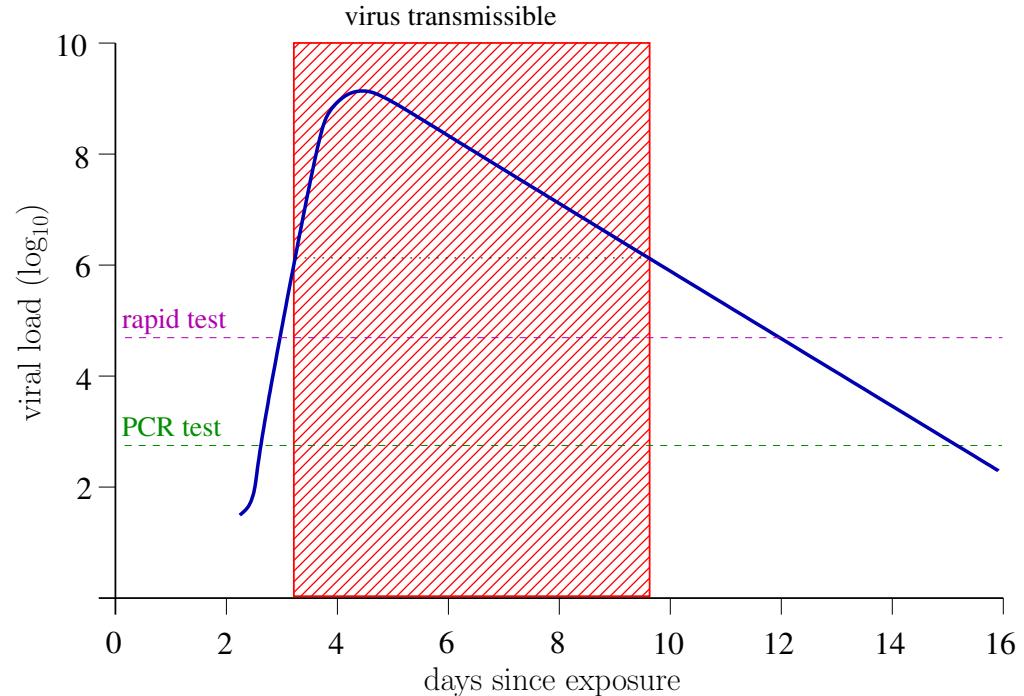
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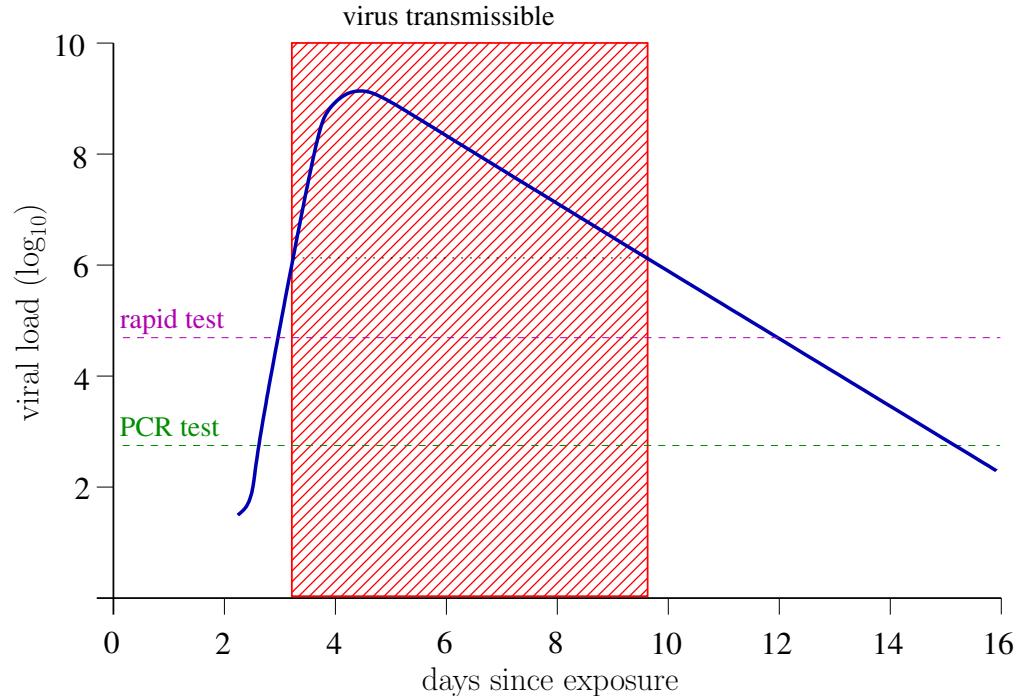
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- FDA will not change its regulations to approve antigen tests but could re-frame them as *transmission-detecting tests* or *surveillance tools*
- The FDA recently stated they would consider less sensitive tests as part of a high-frequency testing plan

Population-scale daily testing

△ Growing movement advocating 10s-100s of millions of rapid tests/day

Idea: **Test every individual before every major social contact:**

Work, school, cinema, shopping, etc



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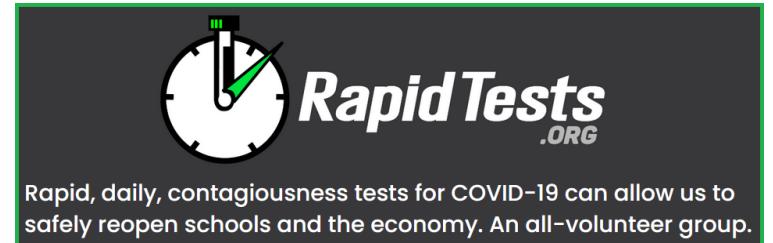
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- ▶ it will stop the virus in three weeks
- ▶ normal life will resume completely
- ▶ only the government can do it!

Cost: Even 500 million tests/day, total cost < 5% of the \$3 trillion Congress already spent on Covid-related support for the economy



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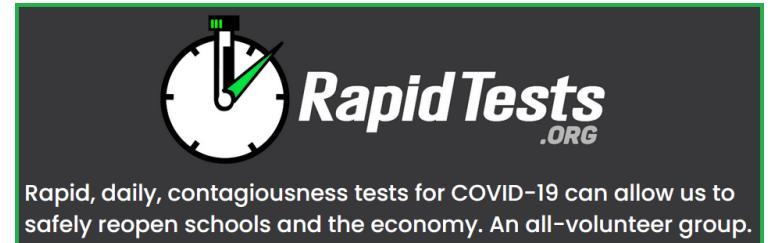
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- ▶ massive production capacity
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- ▶ PCR for the rich/rapid tests for the poor?



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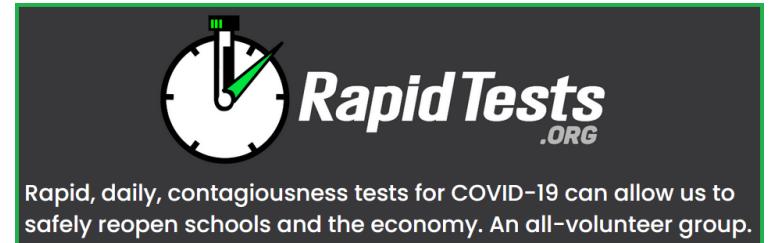
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～ **How much testing is needed for this to work?**



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△ Epidemiological models

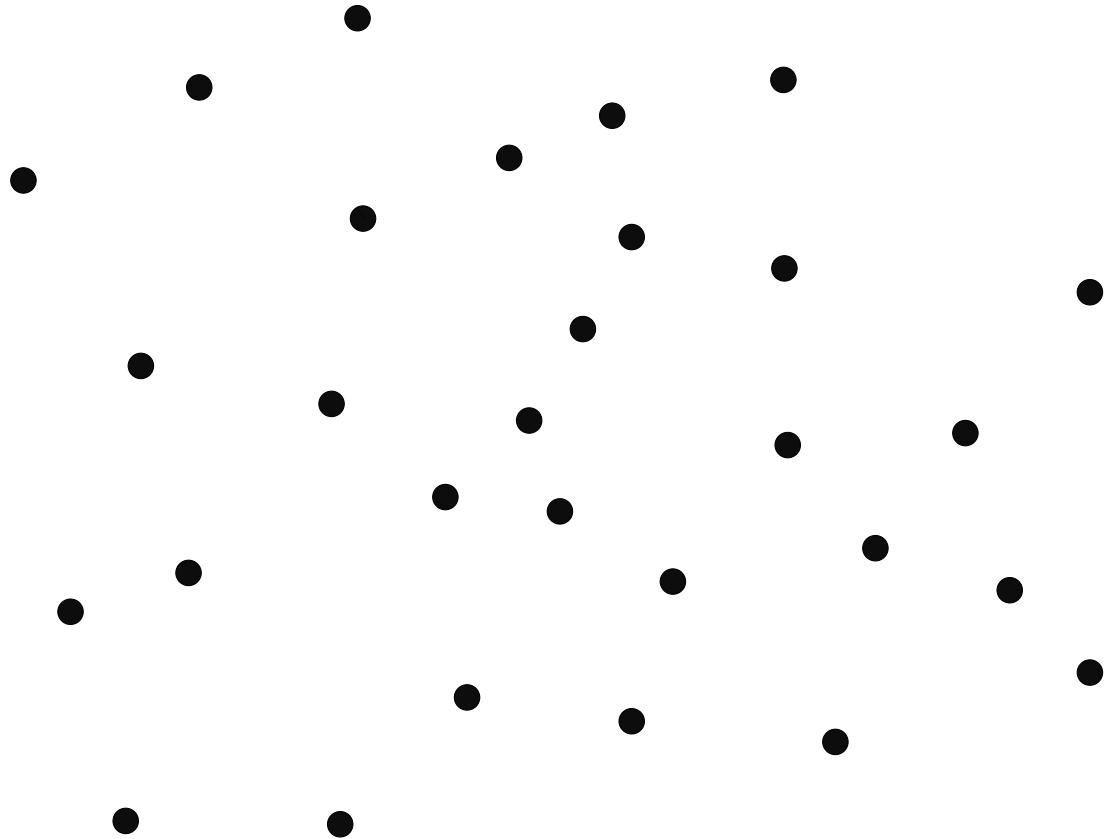
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SIR epidemic on an Erdős-Rényi graph

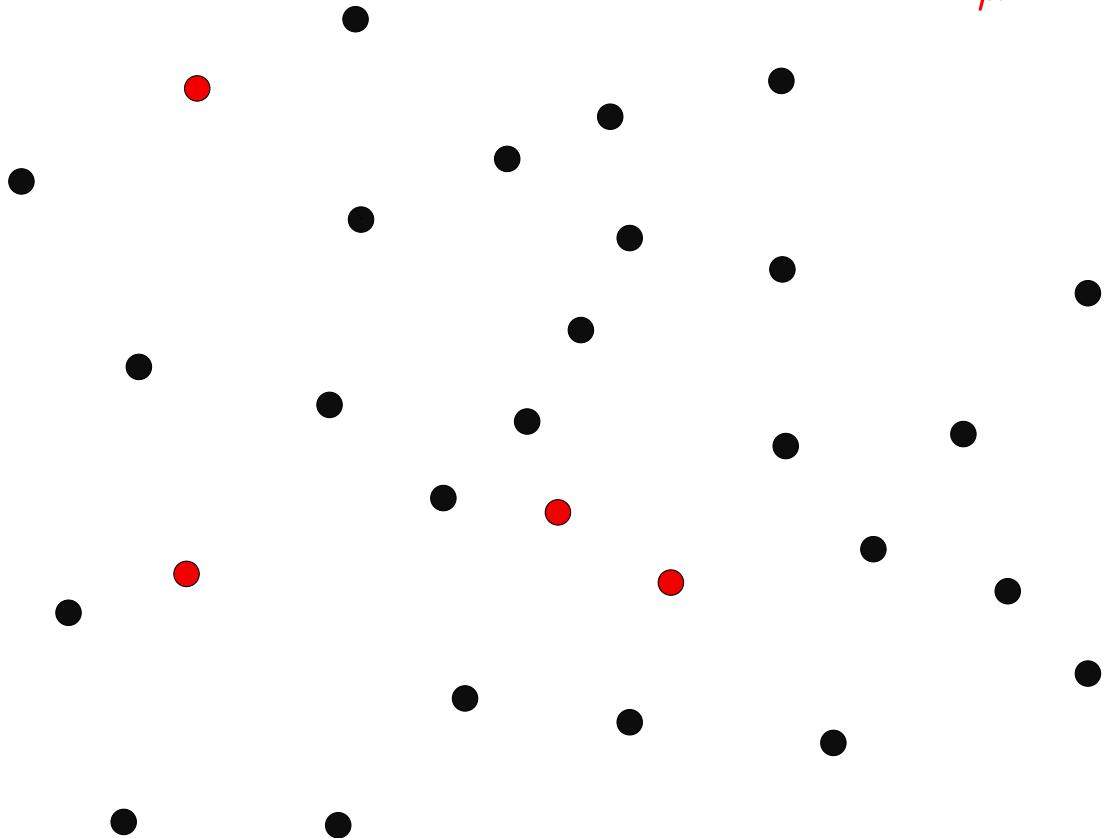
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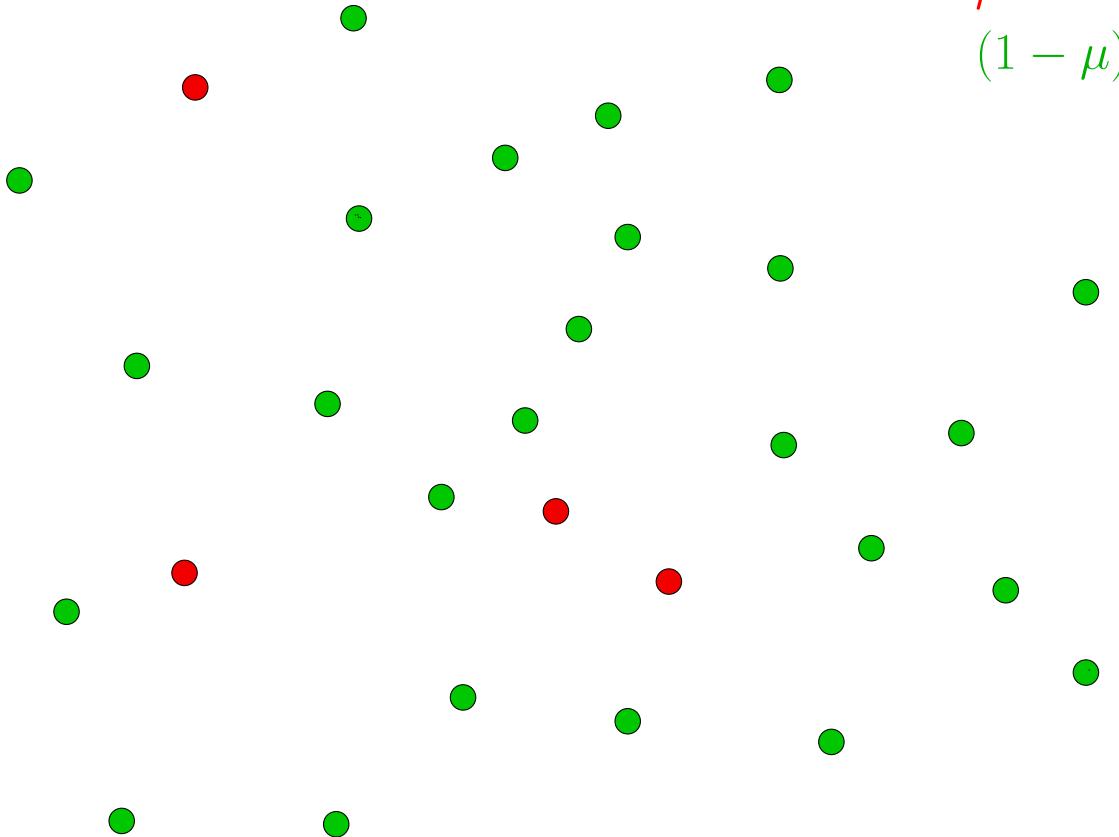
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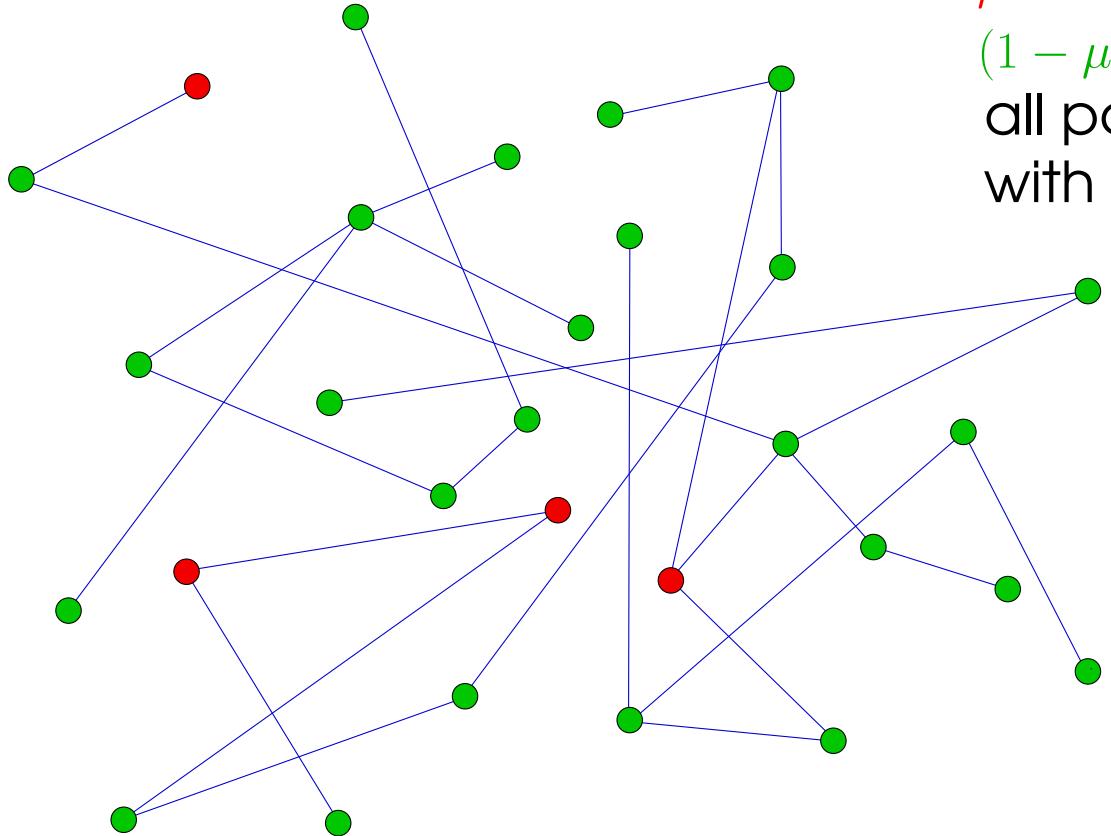


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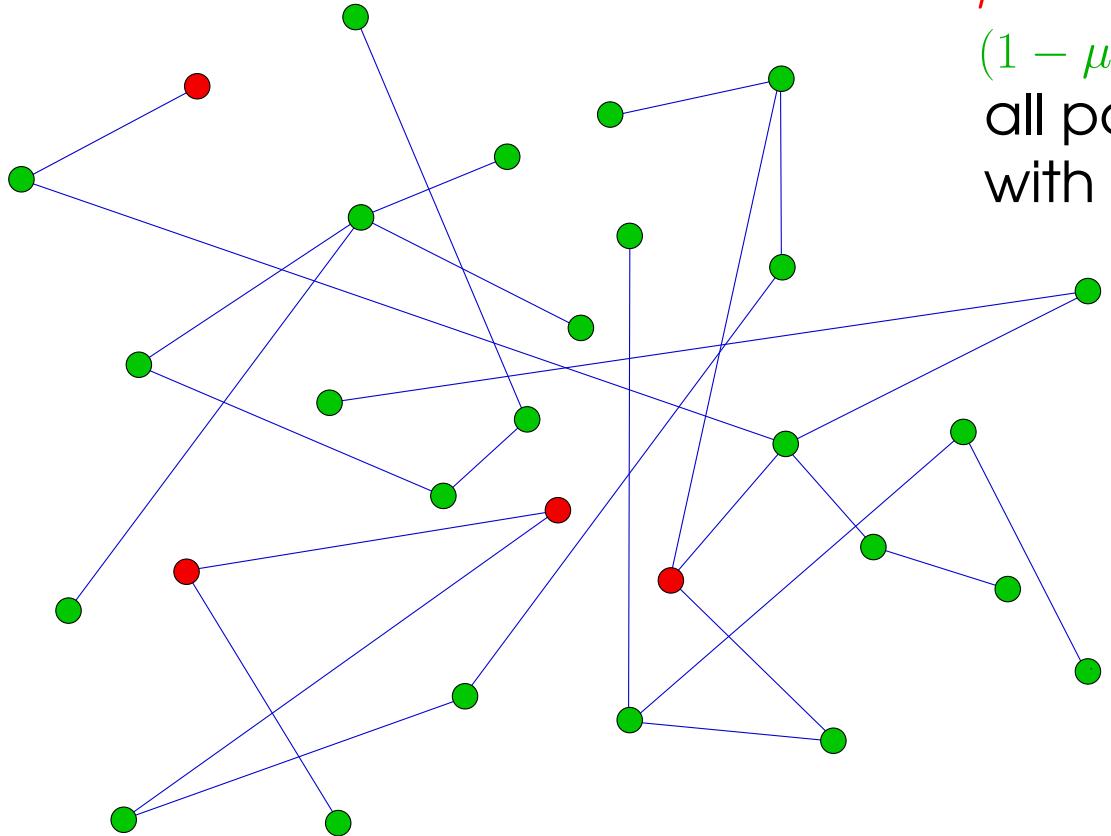


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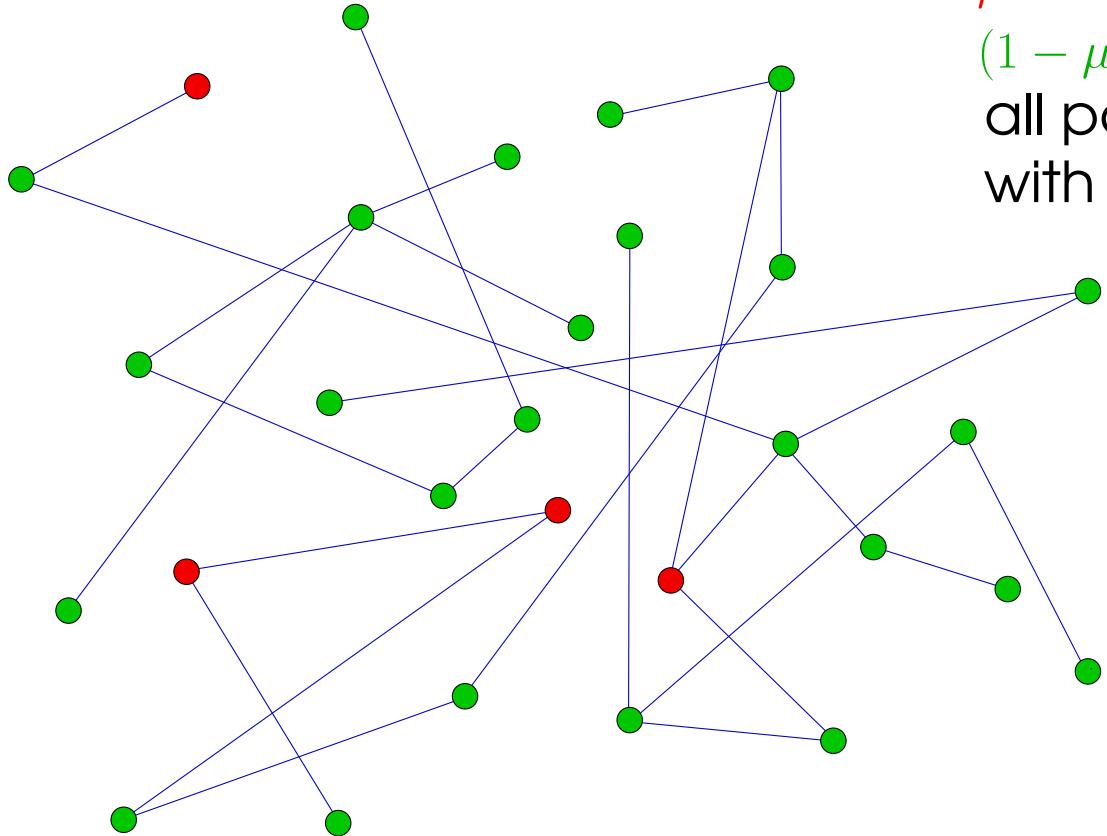
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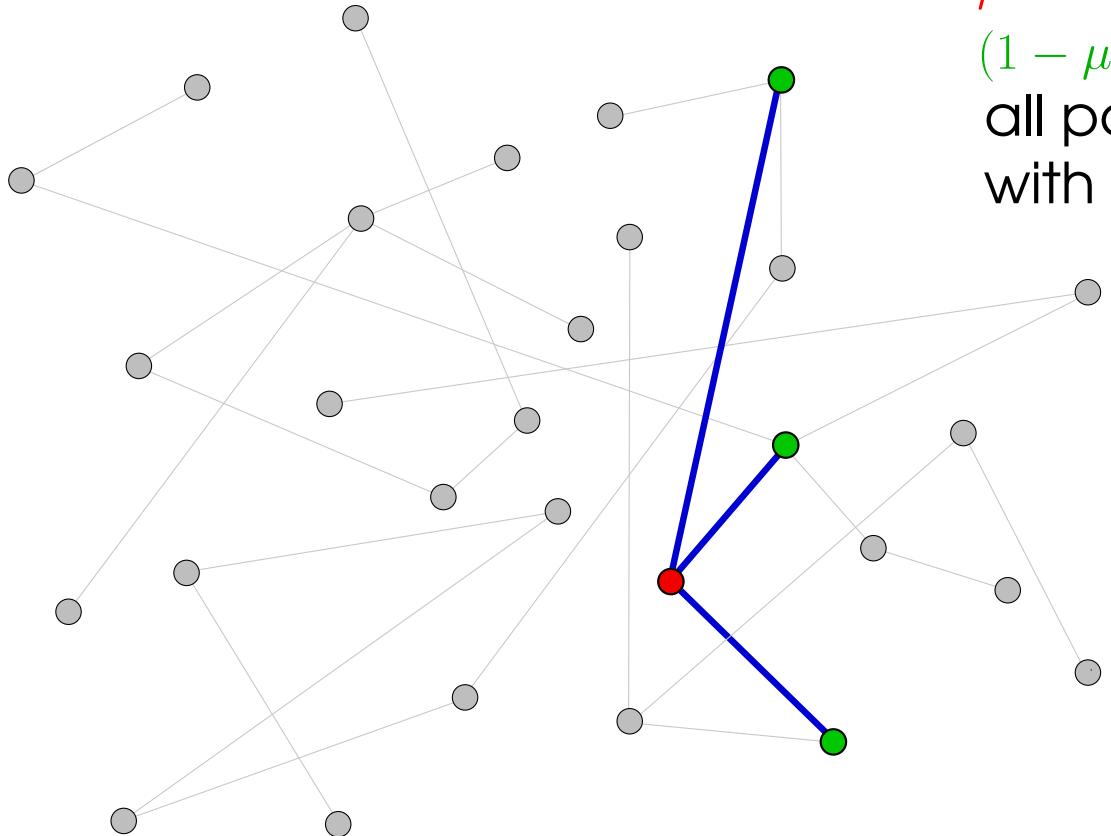
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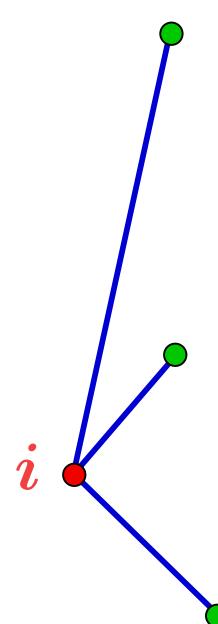
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- ▷ Each infected individual makes infectious contact with their acquaintances at random times

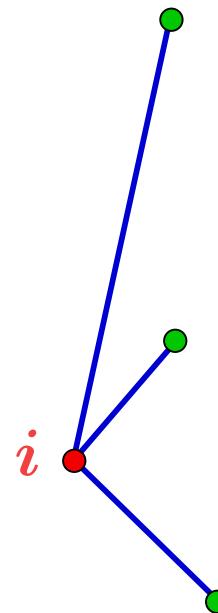
SIR epidemic + random testing



Infections

- ▷ Each infected individual becomes **recovered** at the end of their infection period

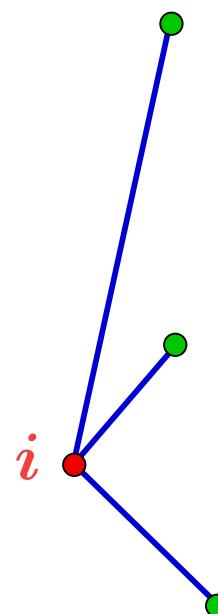
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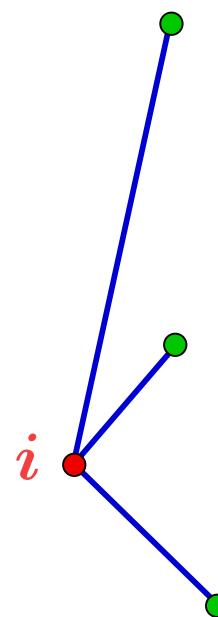
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- ▷ If found **+ve**, individual is quarantined until she becomes **recovered**

SIR epidemic + random testing: Process evolution

Testing parameters

Sensitivity: $1 - \delta$, with δ = probability of false negative

Specificity: assume no false positives

Compliance: q = probability of quarantine compliance

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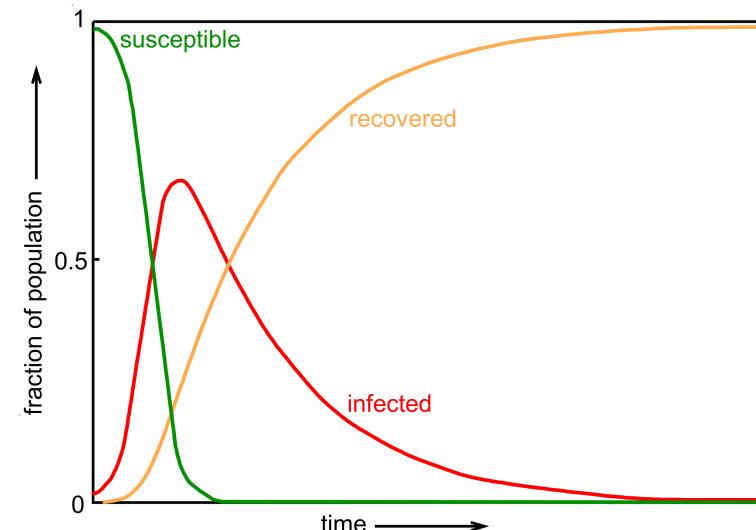
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- ▷ Typical SIR behaviour for large N



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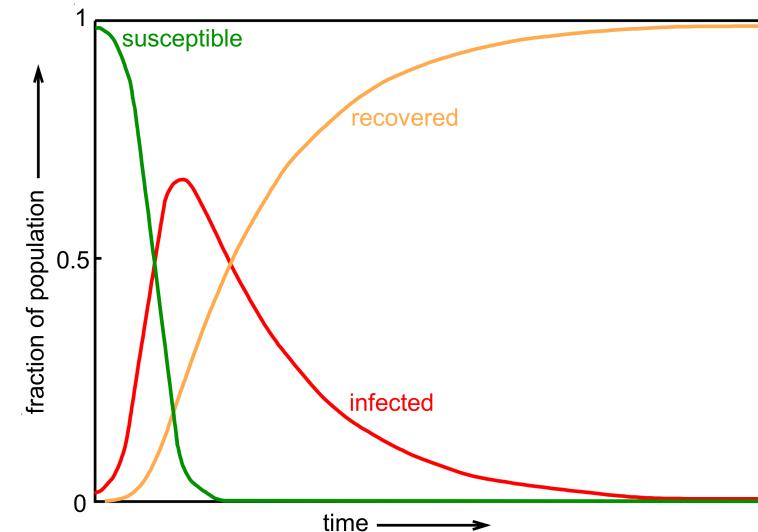
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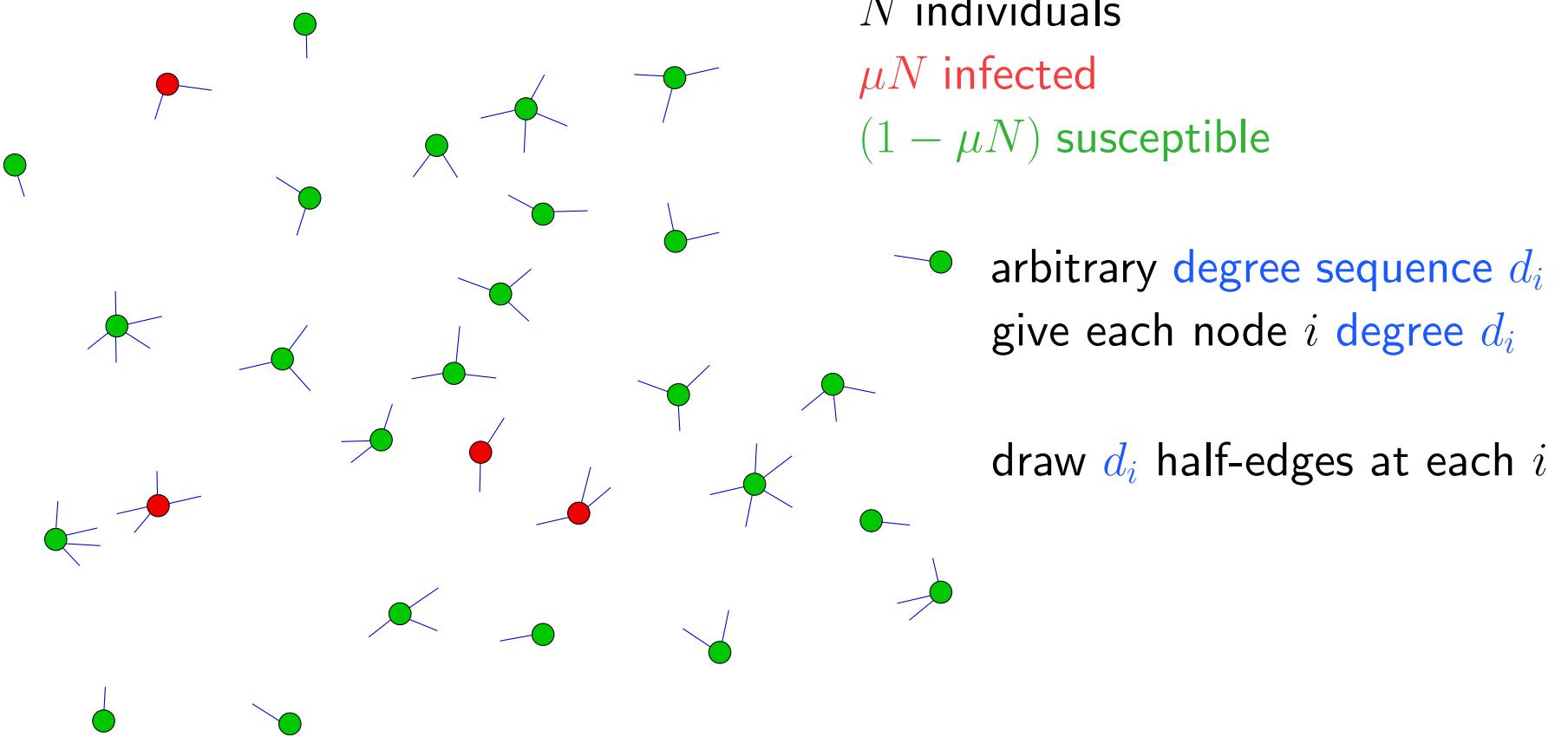
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Assumptions

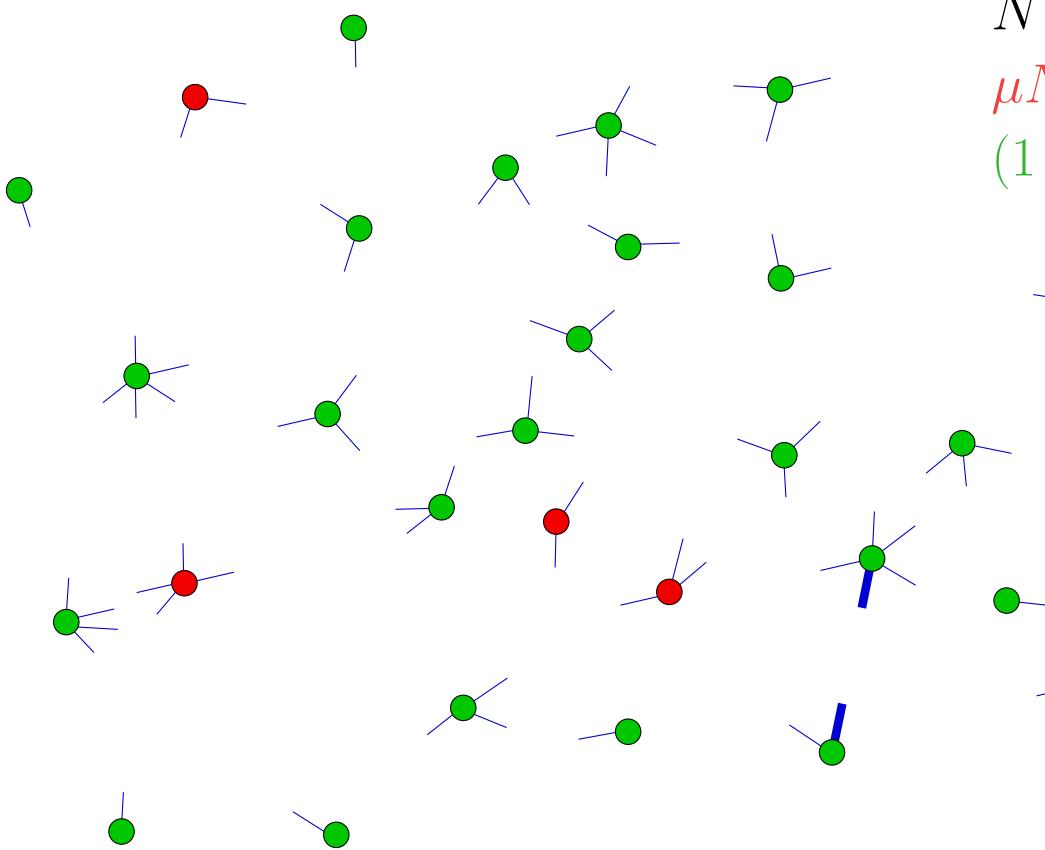
- ~ Model contains several **unrealistic assumptions**:
Poisson degree distr, no false positives, no geographical structure, no disease-specific characteristics, completely random testing
- ~ But these mostly only make our **results more conservative**



SIR epidemic on the configuration model



SIR epidemic on the configuration model



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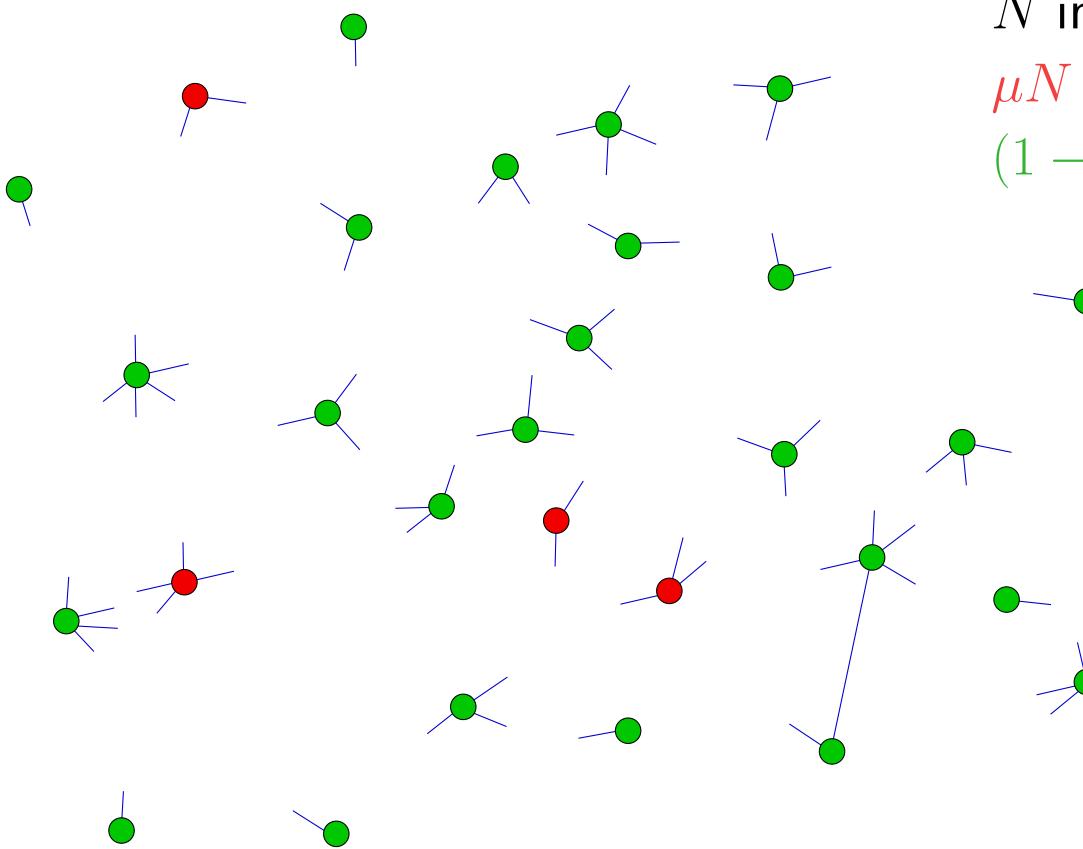
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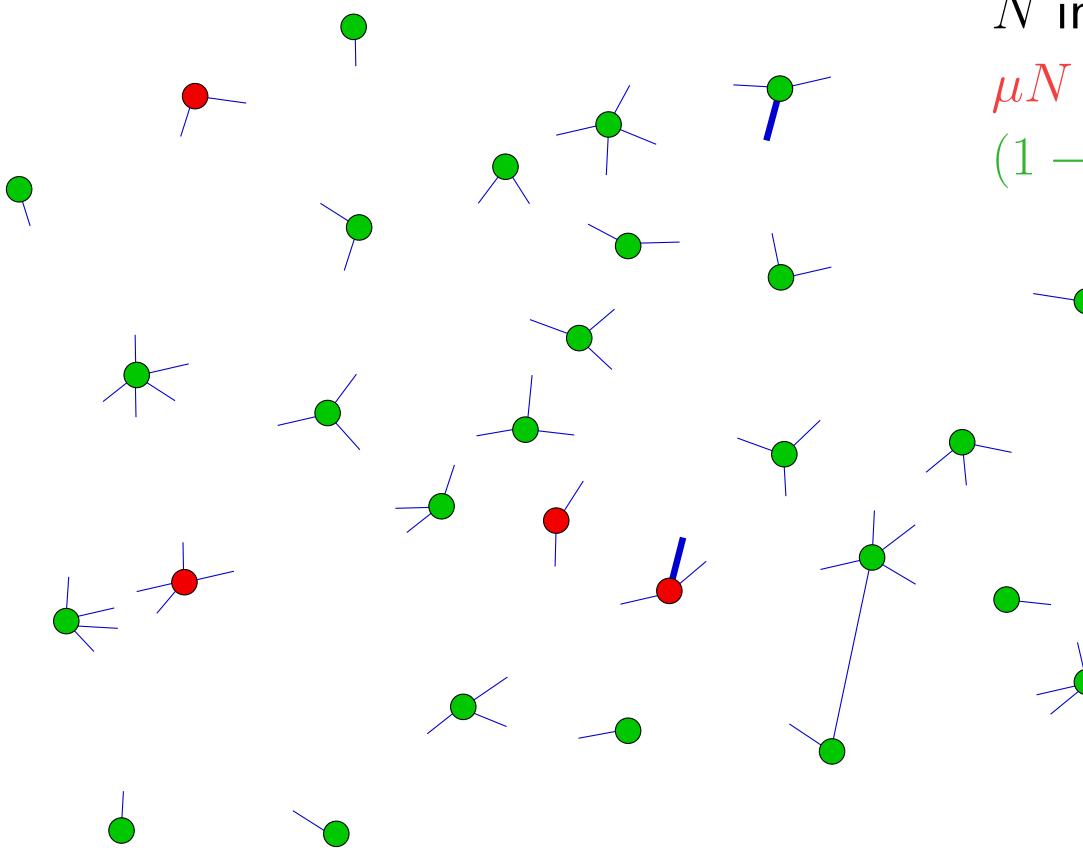
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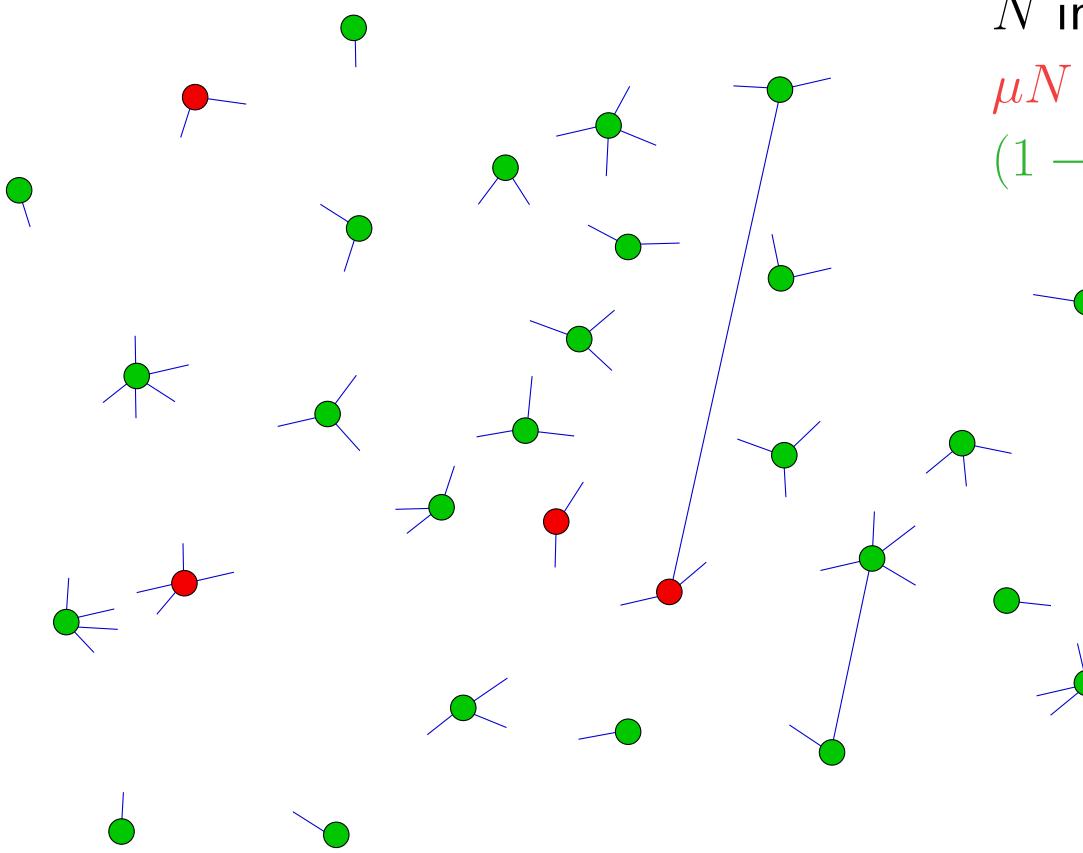
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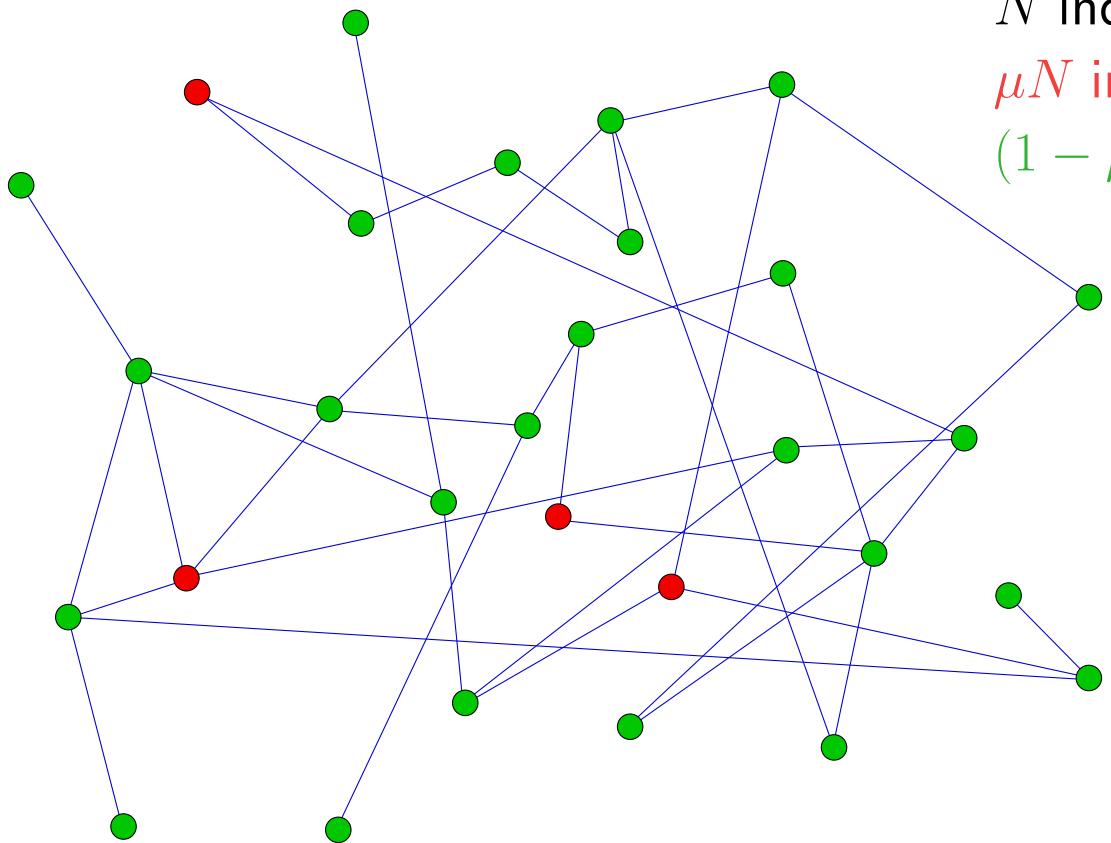
$(1 - \mu N)$ susceptible

arbitrary degree sequence d_i
give each node i degree d_i

draw d_i half-edges at each i

randomly select two
join them
repeat

SIR epidemic on the configuration model



N individuals

μN infected

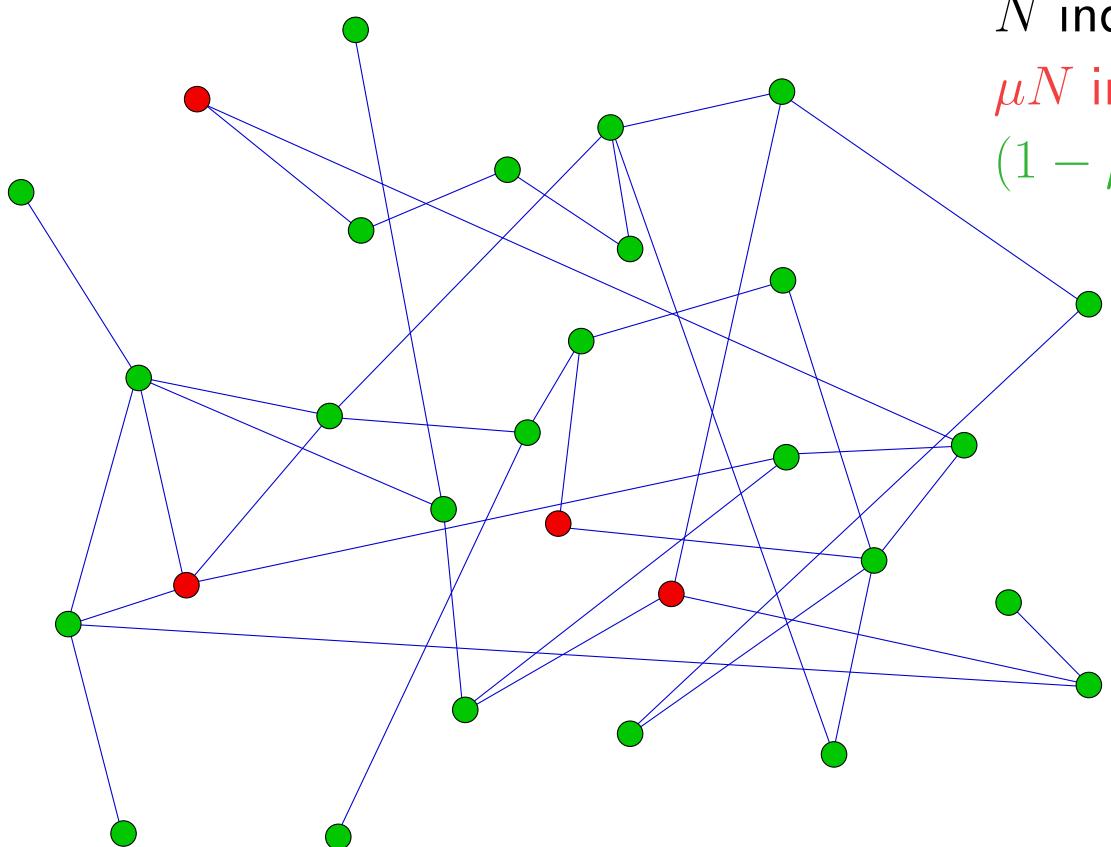
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- ▷ **Possible problem:** There may be self-loops or multiple edges
- ▷ **Easy fix:** Re-do the pairing until there are no self-loops or multiple edges
- ▷ **Result:** Graph uniformly chosen among those with degree sequence $\{d_i\}$

SIR epidemic + testing on the configuration model

Assumptions

- ▶ Degree sequence $\{d_i\}$ can be chosen arbitrarily. But:
- ▶ $\{d_i\}$ usually chosen according to a given degree distr $\{p_k\}$
- ▶ Proportion of individuals with degree k is $\approx p_k$
~~> This holds separately for both infected and susceptible nodes

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Epidemic + testing process

- ▷ Exactly the same as before (once graph is fixed)
- ▷ Again, typical SIR behaviour for large N

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↗ How does the testing rate affect the evolution of the epidemic?

Outline

△ Covid-19 tests

- ▷ Antibody tests
- ▷ PCR tests
- ▷ Antigen tests
- ～ Rapid, cheap, at-home tests

△ Epidemiological models

- ▷ SIR models on random population networks
 - ▶ Erdős-Rényi graphs
 - ▶ Random graphs with given degree distribution
- ▷ SIR epidemics with mass testing

△ Necessary testing rates for suppression

- ▷ **Rigorous results for a broad class of models**
- ▷ Explicit numerical examples

Basic reproduction number R_0 for the E-R model

For the SIR epidemic on the Erdős-Rényi graph

Theorem

As $N \rightarrow \infty$:

- i. **no testing** [e.g. Andersson (1999), Neal (2003)]

$$R_0 = \frac{\alpha\beta}{\beta + \gamma}$$

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$$R_0(\theta) = \frac{\alpha\beta}{\beta + \gamma + \theta(1 - \delta)q}$$

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Proof.

Adding testing to the model is exactly equivalent to shortening the mean infection duration:

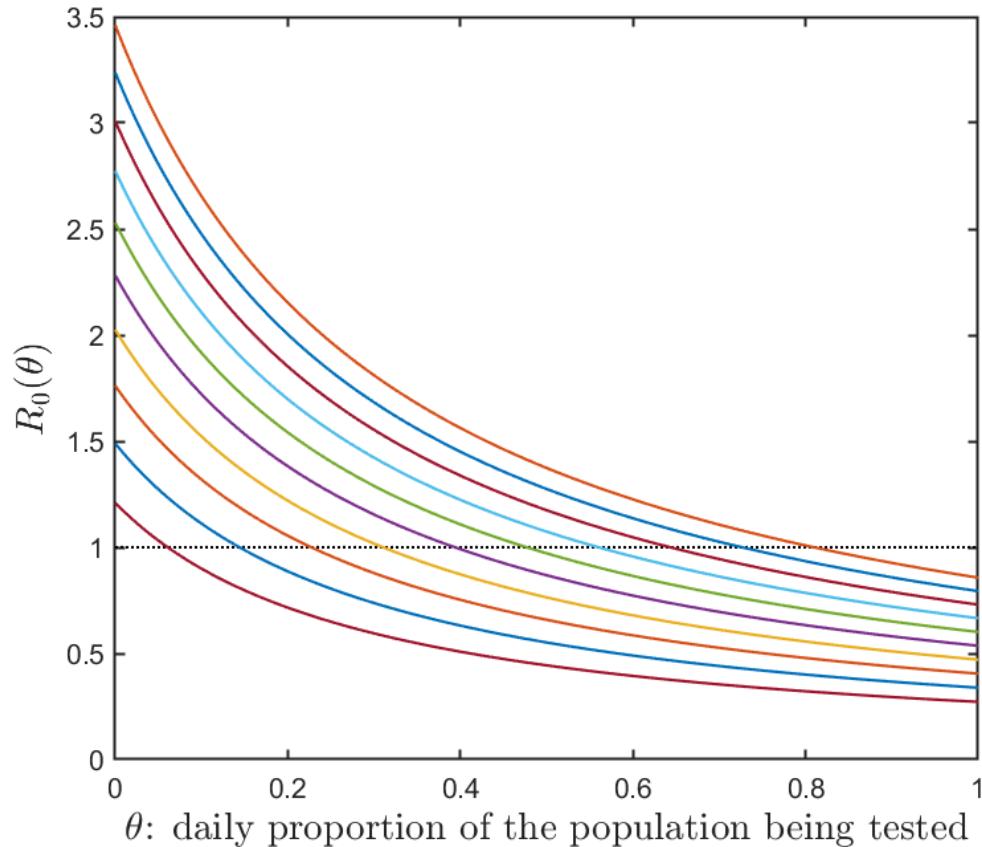
$$\gamma \mapsto \gamma + \theta(1 - \delta)q$$

□

R_0 for the E-R model: Examples

1-in-10000 initially infected
20 acquaintances/individual on average
average infectious period 7 days
contact rate β varies
 $\Rightarrow 1.2 \leq R_0 \leq 3.5$

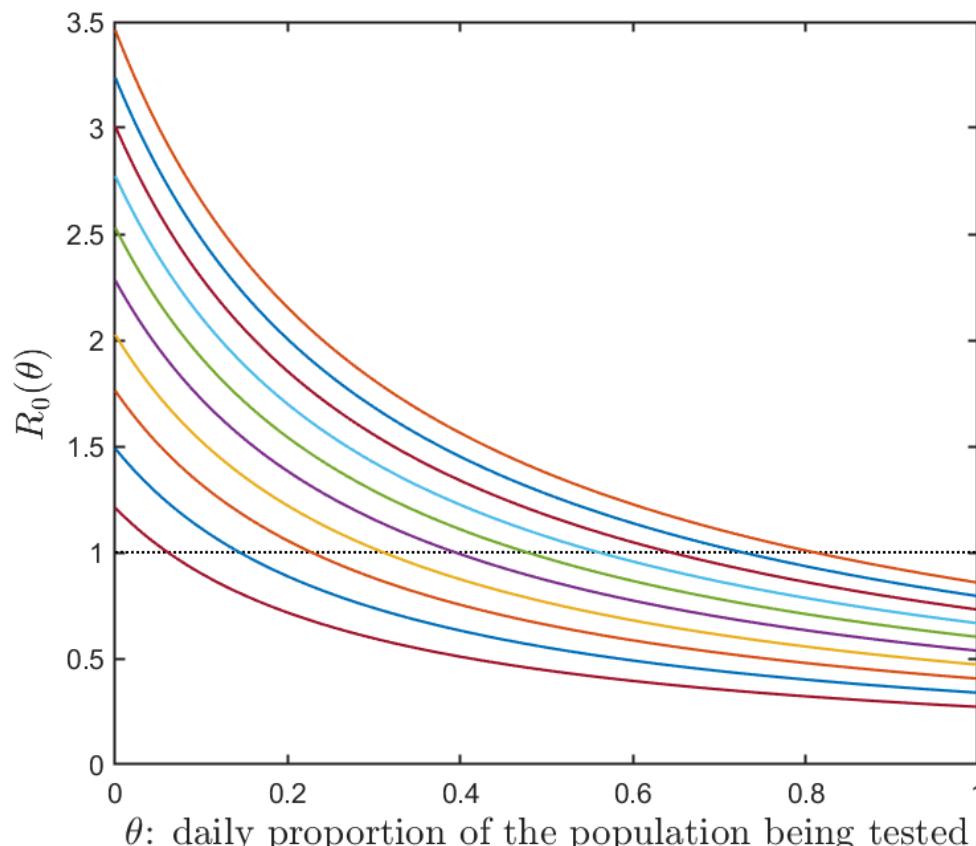
quarantine compliance 75%
test sensitivity 70%
testing rate θ



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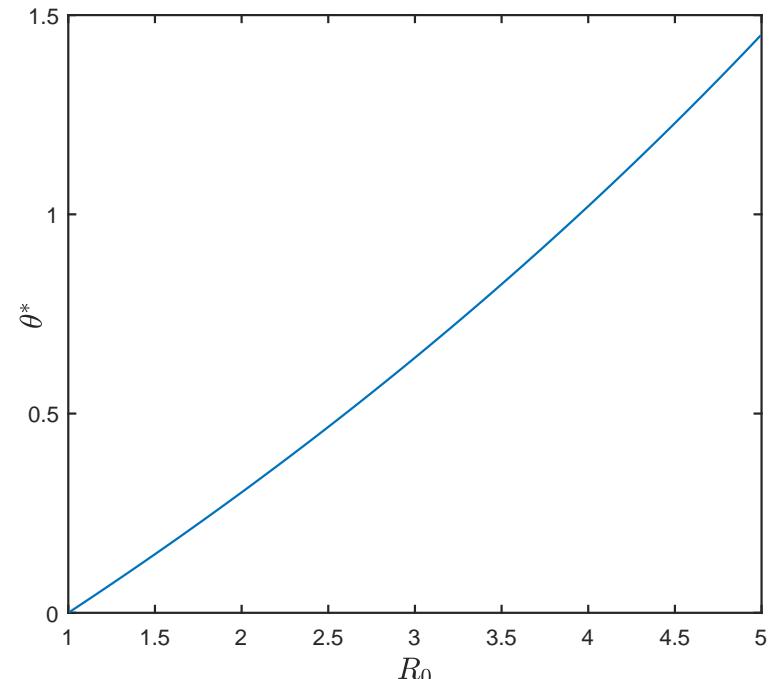


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Corollary

Testing rate required for $R_0(\theta) < 1$:

$$\theta > \theta^* = \frac{\alpha\beta - \beta - \gamma}{q(1 - \delta)}$$



Epidemic size for the E-R model

For the SIR epidemic on the Erdős-Rényi graph, write

$$T_N = \text{total size of the epidemic}$$

and let

$$\begin{aligned}\tau(r, \mu) &= \min\{t > 0 : e^{-rt} = 1 + \mu - t\} \\ s(r, \mu) &= 1 - e^{-r\tau(r, \mu)}\end{aligned}$$

Theorem

As $N \rightarrow \infty$:

- i. **no testing** [e.g. Neal (2003)] $\frac{T_N}{N} \rightarrow s = s(R_0, \mu)$
- ii. **random testing**

$$\frac{T_N}{N} \rightarrow s(\theta) = s(R_0(\theta), \mu)$$

Proof.

Same as before

□

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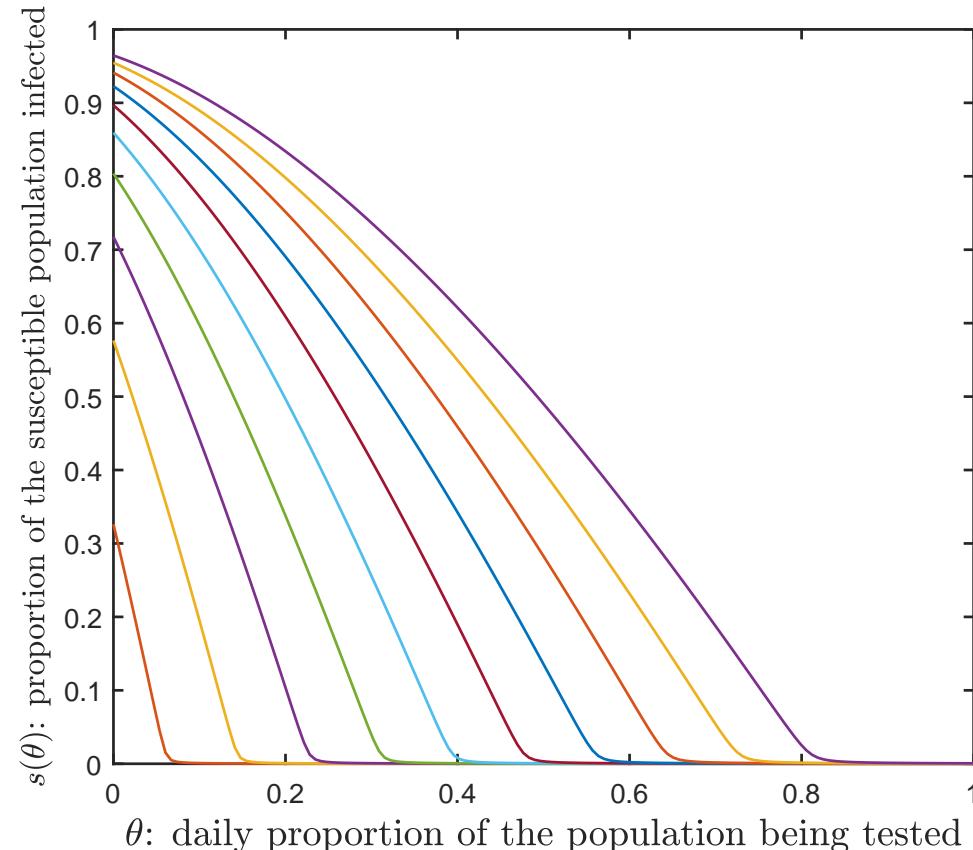
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Small epidemics with the E-R model

For the SIR epidemic on the Erdős-Rényi graph suppose that instead of μN initially infected individuals we only have m of them

Let
$$f(p, \gamma) = \gamma \int_0^\infty \exp \left\{ -\gamma z - \alpha(1-p)(1 - e^{-\beta z}) \right\} dz$$

We say there is a **small epidemic** if $T_N = O(1)$ as $N \rightarrow \infty$

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Theorem

- i. **no testing** [Martin-Löf (1986)]

Suppose $R_0 > 1$

Let $\textcolor{red}{p}$ be the smallest root of $f(p, \gamma) = p$ in $[0, 1]$

\Rightarrow With prob $\textcolor{red}{p}^m$ there is only a **small epidemic**

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ii. random testing

Suppose $R_0(\theta) > 1$

Let $p(\theta)$ be the smallest root of $f(p, \gamma + \theta(1 - \delta)q) = p$ in $[0, 1]$

\Rightarrow With prob $p(\theta)^m$ there is only a **small epidemic**

Proof.

Same idea as before

□

Probability of small epidemic: Examples

20 acquaintances/individual on average

quarantine compliance 75%

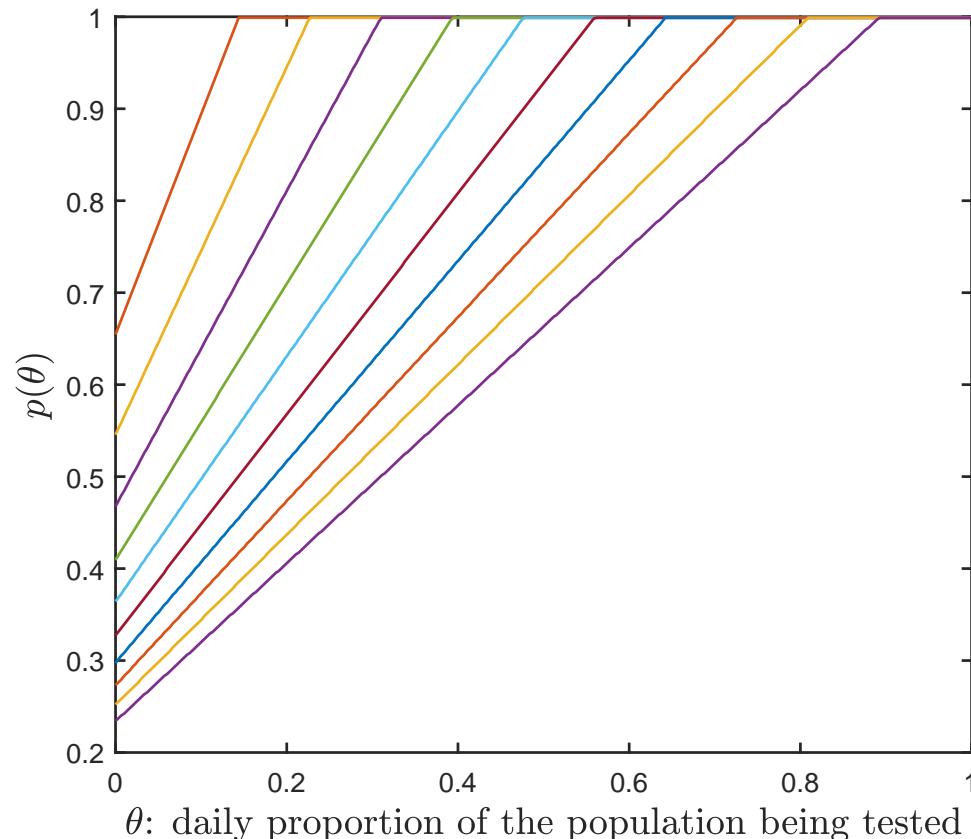
average infectious period 7 days

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contact rate β varies

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Basic reproduction number R_0 for the configuration model

For the SIR epidemic on the configuration model with degree distr $\{p_k\}$ let

$$\lambda = \sum_{k=0}^{\infty} kp_k \quad v^2 = \sum_{k=0}^{\infty} k(k-1)p_k$$

Theorem

As $N \rightarrow \infty$ (under mild conditions):

- i. **no testing** [e.g. Janson-Luczak-Windridge (2014)]

$$R_0 = \left(\frac{\beta}{\beta + \gamma} \right) \left(\frac{(1 - \mu)v^2}{\lambda} \right)$$

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Proof.

Same idea as before

□

R_0 for the configuration model: Examples

1-in-10000 initially infected

degree distr $p_k \propto k^{-1.75} e^{-0.02k}$

$\lambda \approx 3.5$ acquaintances/individual on average

average infectious period 7 days

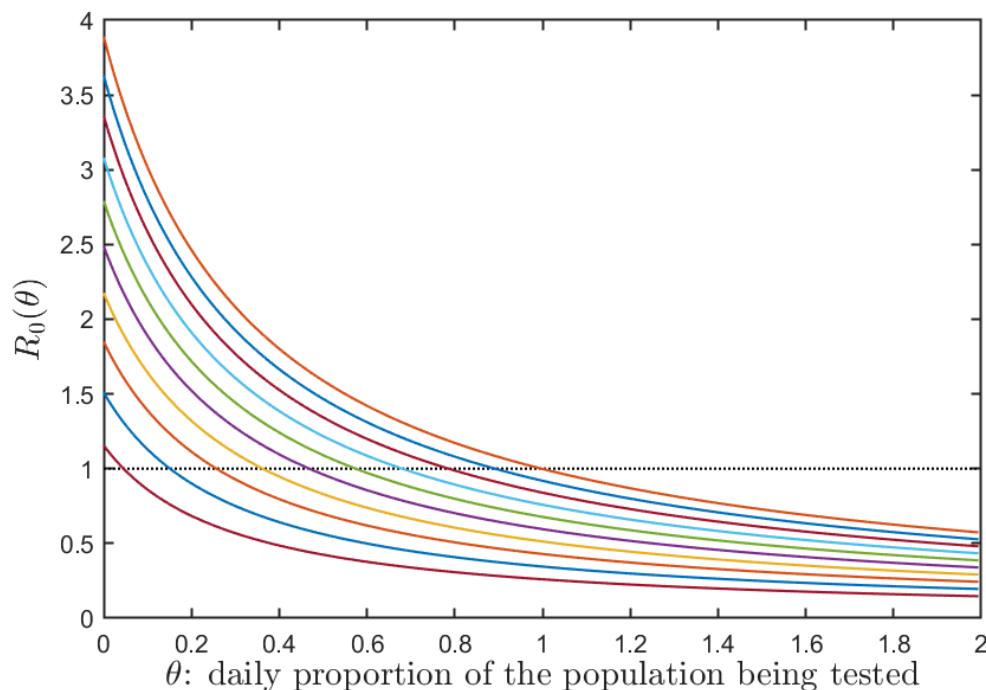
contact rate β varies

$$\Rightarrow 1.15 \leq R_0 \leq 3.9$$

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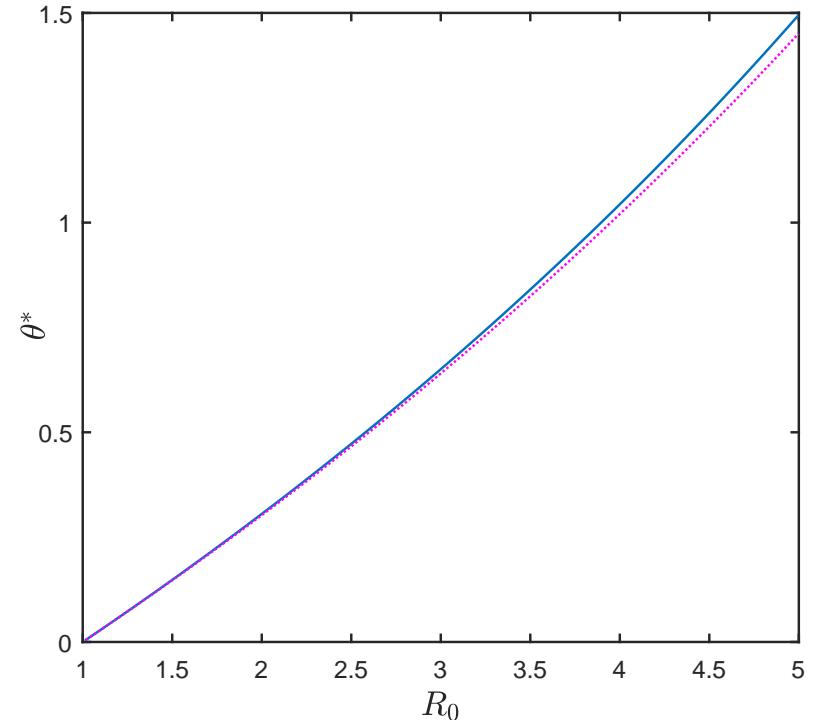
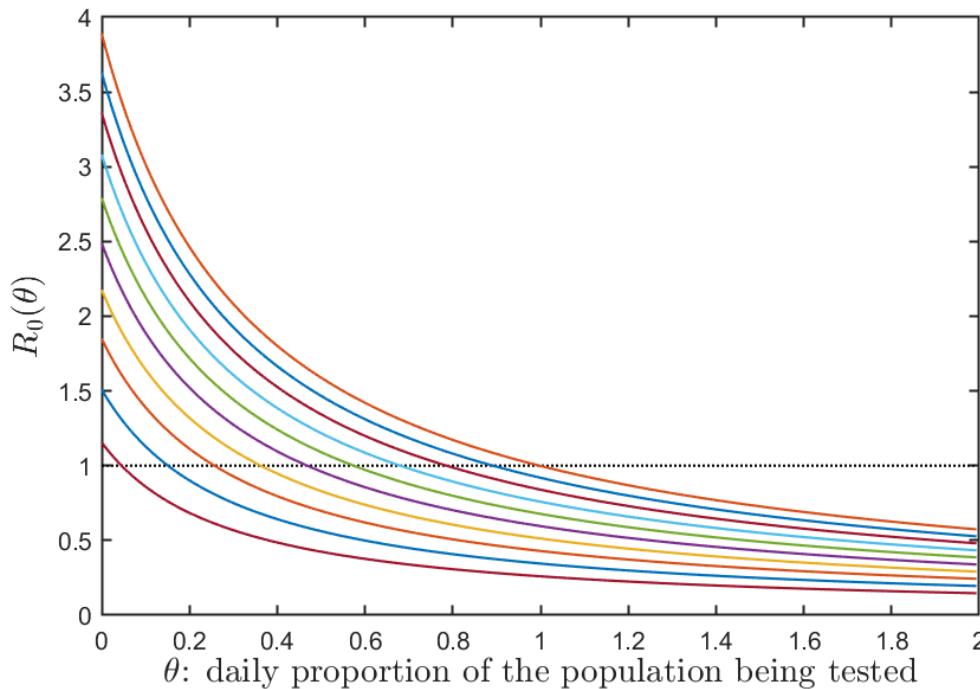
test sensitivity 70%

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Corollary

Testing rate required for $R_0(\theta) < 1$:

$$\theta > \theta^* = \frac{1}{q(1-\delta)} \left[\beta \left(\frac{(1-\mu)v^2}{\lambda} - 1 \right) - \gamma \right]$$



Concluding remarks

~> **A Covid-19 public policy proposal**

- ▷ Rapid, cheap, at-home, saliva-based, paper tests
- ▷ Daily population-scale testing

~> **Epidemiological models suggest**

- ▷ Random testing is effective
 - ▶ Reduces R_0
 - ▶ Decreases epidemic size
 - ▶ Its benefits are additive to other measures

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~> **Epidemiological models suggest**

- ▷ Random testing is effective
 - ▶ Reduces R_0
 - ▶ Decreases epidemic size
 - ▶ Its benefits are additive to other measures
- ▷ Approximately daily testing *may* in fact be sufficient to suppress the pandemic

~> **Precise mathematical results offer**

- ▷ Strong, quantitative evidence of effectiveness
- ▷ Useful, conservative rules of thumb

In recent news

△ The rapid testing proposal is gaining traction

Aug 27: First FDA-approved rapid test: Abbott's \$5, 15-minute test

White House announced \$750 million deal with Abbott

Sept 1: New rapid test by Roche-SD Biosensor partnership
will be made available in Europe and the UK

Sept 9: UK PM announced "Operation Moonshot", likely cost £100bn
aiming for 10 million daily tests by spring

△ Around the world

Italy. Approved 3-minute saliva test the "Daily Tampon"

France. New "antigénique rapide" test used by authorities

Senegal. UK-Senegal partnership developed \approx \$1 home antigen test

India. Authorities switching over to a rapid antigen test

References https://www.dpmms.cam.ac.uk/~ik355/PAPERS/Covid_talk_bib.pdf

Slides https://www.dpmms.cam.ac.uk/~ik355/PAPERS/Covid_talk_slides.pdf

Our paper <https://www.dpmms.cam.ac.uk/~ik355/pubs.html>

Technical assumptions for the configuration model

Assume initially N_I infected individuals s.t. $N_{I,k}$ have degree k and $N_S = N - N_I$ susceptible individuals s.t. $N_{S,k}$ have degree k

Assume $N_I/N \rightarrow \mu$ and $N_S/N \rightarrow (1 - \mu)$ for $\mu \in (0, 1)$, that $\lambda \in (0, \infty)$ and that for all k

$$\frac{N_{S,k}}{N_S} \rightarrow p_k, \quad \frac{N_{I,k}}{N_I} \rightarrow p_k$$
$$\sum_{k=0}^{\infty} k \frac{N_{S,k}}{N_S} \rightarrow \sum_{k=0}^{\infty} kp_k = \lambda, \quad \sum_{k=0}^{\infty} k \frac{N_{I,k}}{N_I} \rightarrow \sum_{k=0}^{\infty} kp_k = \lambda.$$

Two last technical assumptions are required

For $N_k = \text{total no of individuals with degree } k$:

$$\max\{k ; N_{I,k} > 0\} = o(N) \quad \text{and} \quad \sum_{k=0}^{\infty} k^2 N_k = O(N)$$