

A new strategy for pooling COVID-19 tests to detect outbreaks early

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Pooling in a school/university setting: saliva or nose samples from students sharing the same classroom/student residence are pooled in a single tube; the virological status of the mixture is assessed by a single molecular test (RT-qPCR or RT-LAMP). A positive test indicates that at least one individual is infected with CoV2-SARS, allowing early action to be taken to contain an outbreak. The screening process can be repeated on a regular basis. Credit: Jean-François Rupprecht, the Centre for Theoretical Physics (CNRS/Aix-Marseille Université/Université de Toulon), 2021



Researchers have proposed a new quantitative strategy for pooling COVID-19 tests in order to monitor spread and detect outbreaks early within closed communities, such as nursing homes or universities. Vincent Brault of Université Grenoble Alpes in Grenoble, France, and colleagues present this method in the open-access journal *PLOS Computational Biology*.

In the face of limited supplies, some communities have achieved largescale COVID-19 testing by grouping individuals' saliva or nasal swab samples into <u>pools</u>. If a pool tests negative, all people in the pool receive a negative result. If it tests positive, follow-up testing shows who in that pool is positive. However, pooling can potentially dilute a positive <u>sample</u>, resulting in a negative result when someone in the pool is in fact positive: a false-negative result.

In the new study, Brault and colleagues used quantitative tools to examine the problem of false negatives in the context of closed communities. While previous research on pooling has often focused on minimizing the number of tests needed to pinpoint a positive case, this study focused on pooling as a way to track the presence of COVID-19 in a community.

The research team developed mathematical models that estimate the risk of false-negative results when individual samples are pooled for COVID-19 testing via the reverse transcription quantitative polymerase chain reaction (RT-qPCR) method. These models drew on large-scale clinical datasets on the concentration of SARS-CoV-2—the virus that causes COVID-19—within individual samples.

Based on the models, the researchers developed a quantitative pooledtesting protocol for determining the prevalence of COVID-19 within a closed community, such as a nursing home or university. While further refinement is needed to prepare the protocol for practical application, it



could set the stage for earlier detection of outbreaks.

"We found that pooling is not only a promising method to break transmission chains at the earliest stage, but that it is also surprisingly efficient at estimating the level of viral penetration in communities," Brault said. "We think pooled testing might become a broader tool to combat asymptomatic spread among youngsters at schools."

More information: Brault V, Mallein B, Rupprecht J-F (2021) Group testing as a strategy for COVID-19 epidemiological monitoring and community surveillance. *PLoS Comput Biol* 17(3): e1008726. <u>doi.org/10.1371/journal.pcbi.1008726</u>

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