***eLife’s* transparent reporting form**

We encourage authors to provide detailed information *within their submission* to facilitate the interpretation and replication of experiments. Authors can upload supporting documentation to indicate the use of appropriate reporting guidelines for health-related research (see [EQUATOR Network](http://www.equator-network.org/%20)), life science research (see the [BioSharing Information Resource](https://biosharing.org/)), or the [ARRIVE guidelines](http://www.plosbiology.org/article/info%3Adoi/10.1371/journal.pbio.1000412) for reporting work involving animal research. Where applicable, authors should refer to any relevant reporting standards documents in this form.

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**Sample-size estimation**

* You should state whether an appropriate sample size was computed when the study was being designed
* You should state the statistical method of sample size computation and any required assumptions
* If no explicit power analysis was used, you should describe how you decided what sample (replicate) size (number) to use

Please outline where this information can be found within the submission (e.g., sections or figure legends), or explain why this information doesn’t apply to your submission:

This was an observational study.

**Replicates**

* You should report how often each experiment was performed
* You should include a definition of biological versus technical replication
* The data obtained should be provided and sufficient information should be provided to indicate the number of independent biological and/or technical replicates
* If you encountered any outliers, you should describe how these were handled
* Criteria for exclusion/inclusion of data should be clearly stated
* High-throughput sequence data should be uploaded before submission, with a private link for reviewers provided (these are available from both GEO and ArrayExpress)

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This was an observational study.

**Statistical reporting**

* Statistical analysis methods should be described and justified
* Raw data should be presented in figures whenever informative to do so (typically when N per group is less than 10)
* For each experiment, you should identify the statistical tests used, exact values of N, definitions of center, methods of multiple test correction, and dispersion and precision measures (e.g., mean, median, SD, SEM, confidence intervals; and, for the major substantive results, a measure of effect size (e.g., Pearson's r, Cohen's d)
* Report exact p-values wherever possible alongside the summary statistics and 95% confidence intervals. These should be reported for all key questions and not only when the p-value is less than 0.05.

Please outline where this information can be found within the submission (e.g., sections or figure legends), or explain why this information doesn’t apply to your submission:

Results section:

*Characteristics of HCW...*

[Paragraph 2]

Fishers exact test (symptomatic vs asymptomatic / positivity)

Fishers exact test (Clinical area / positivity)

[Paragraph 3, Figure 1]

Mann Whitney test (viral loads / symptoms)

[Paragraph 3, Figure S1]

Mann Whitney test (viral loads / duration of symptoms or clinical risk score)

*Identification of two clusters...*

Fishers exact test (clinical area / ‘asymptomatically’ screened positivity)

Fishers exact test (positivity ward F / other green areas)

Fishers exact test (positivity Ward Q / other red areas)

(For large datasets, or papers with a very large number of statistical tests, you may upload a single table file with tests, Ns, etc., with reference to sections in the manuscript.)

**Group allocation**

* Indicate how samples were allocated into experimental groups (in the case of clinical studies, please specify allocation to treatment method); if randomization was used, please also state if restricted randomization was applied
* Indicate if masking was used during group allocation, data collection and/or data analysis

Please outline where this information can be found within the submission (e.g., sections or figure legends), or explain why this information doesn’t apply to your submission:

Groups were defined as follows: *asymptomatic HCW screening group, symptomatic HCW screening group and symptomatic household contact screening group*. This is described in materials and methods, *staff screening protocols*.

**Additional data files (“source data”)**

* We encourage you to upload relevant additional data files, such as numerical data that are represented as a graph in a figure, or as a summary table
* Where provided, these should be in the most useful format, and they can be uploaded as “Source data” files linked to a main figure or table
* Include model definition files including the full list of parameters used
* Include code used for data analysis (e.g., R, MatLab)
* Avoid stating that data files are “available upon request”

Please indicate the figures or tables for which source data files have been provided:

Figure 1:

Individual CT values by group

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|

|  |  |
| --- | --- |
| HCW asymptomatic group  | HCW symptomatic / symptomatic household contact group  |
| 35 | 23 |
| 36 | 24 |
| 31 | 24 |
| 31 | 21 |
| 23 | 34 |
| 31 | 24 |
| 36 | 28 |
| 25 | 32 |
| 32 | 16 |
| 34 | 26 |
| 23 | 21 |
| 18 | 22 |
| 27 | 32 |
| 30 | 19 |
| 33 | 30 |
| 35 | 33 |
| 32 | 34 |
| 28 | 30 |
| 32 | 35 |
| 36 | 18 |
| 36 | 36 |
| 14 | 35 |
| 36 | 23 |
| 35 | 33 |
| 29 | 24 |
| 31 | 26 |
| 31 | 23 |
| 29 | 24 |
| 32 | 21 |
| 34 | 23 |
| 23 |  |

 |  |

Figure 2:

Central pie chart distribution of HCWs in asymptomatic screen

|  |  |
| --- | --- |
| (i)Asymptomatic | 6 |
| (ii) ≤7 days | 12 |
| (iii) >7 days | 12 |
| Total | 30 |
|  |  |

Left pie chart probability of COVID-19 based on retrospective analysis using severity criteria within those who had experienced symptoms for >7 days

|  |  |
| --- | --- |
| High | 10 |
| Medium | 1 |
| Low | 1 |

Right pie chart probability of COVID-19 based on retrospective analysis using severity criteria within those who had experienced symptoms for ≤7 days

|  |  |
| --- | --- |
| High | 3 |
| Medium | 2 |
| Low | 7 |

Figure 3:

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Ward  | Tests assigned to this location  | Positive tests assigned to this location  | Positive tests from this location as % of all positive tests returned  | Positive tests from this location as a percentage of tests performed at this location  |
| A  | 26 | 0 | 0.00 | 0.00 |
| B  | 44 | 1 | 3.57 | 2.27 |
| C  | 30 | 0 | 0.00 | 0.00 |
| D  | 38 | 0 | 0.00 | 0.00 |
| E  | 54 | 1 | 3.57 | 1.85 |
| F  | 43 | 4 | 14.29 | 9.30 |
| G  | 15 | 0 | 0.00 | 0.00 |
| H  | 12 | 0 | 0.00 | 0.00 |
| I  | 17 | 0 | 0.00 | 0.00 |
| J  | 31 | 0 | 0.00 | 0.00 |
| K  | 54 | 3 | 10.71 | 5.56 |
| L  | 11 | 0 | 0.00 | 0.00 |
| M  | 35 | 1 | 3.57 | 2.86 |
| N  | 20 | 0 | 0.00 | 0.00 |
| O  | 44 | 4 | 14.29 | 9.09 |
| P  | 19 | 0 | 0.00 | 0.00 |
| Q  | 37 | 7 | 25.00 | 18.92 |
| R  | 72 | 3 | 10.71 | 4.17 |
| S  | 59 | 1 | 3.57 | 1.69 |
| T  | 42 | 2 | 7.14 | 4.76 |
| U  | 33 | 1 | 3.57 | 3.03 |
| Total  | 737 | 28 | 100.00 | 63.50 |

Figure 4:

|  |  |  |  |
| --- | --- | --- | --- |
|  | HCW asymptomatic group | HCW symptomatic / symptomatic household contact group | Reactive screening group |
| F | 4 | 2 | 3 |
| Q | 7 | 1 | 1 |

Figure S1:

|  |  |  |
| --- | --- | --- |
| Asymptomatic group | <=7 days | > 7days |
| 23 | 23 | 35 |
| 16 | 26 | 36 |
| 23 | 30 | 31 |
| 21 | 26 | 31 |
| 22 | 28 | 23 |
| 32 | 33 | 31 |
| 33 | 35 | 36 |
| 16 | 24 | 25 |
| 19 |  | 32 |
| 36 |  | 34 |
| 24 |  | 23 |
| 24 |  | 18 |
| 21 |  | 27 |
| 34 |  | 30 |
| 34 |  | 33 |
| 30 |  | 35 |
| 21 |  | 32 |
| 24 |  | 28 |
| 32 |  | 32 |
| 18 |  | 36 |
| 24 |  | 36 |
| 35 |  | 14 |
|  |  | 36 |
|  |  | 35 |
|  |  | 29 |
|  |  | 31 |
|  |  | 31 |
|  |  | 29 |
|  |  | 32 |
|  |  | 34 |
|  |  | 23 |