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Incidence, household transmission, and neutralizing antibody seroprevalence of Coronavirus Disease 2019 in Egypt: Results of a community-based cohort

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Abstract

SARS-CoV-2 virus is transmitted in closed settings to people in contact with COVID-19 patients such as healthcare workers and household contacts. However, household personto-person transmission studies are limited. Households participating in an ongoing cohort study of influenza incidence and prevalence in rural Egypt were followed. Baseline enrollment was done from August 2015 to March 2017. The study protocol was amended in April 2020 to allow COVID-19 incidence and seroprevalence studies. A total of 290 households including 1598 participants were enrolled and followed from April to October 2020 in four study sites. When a participant showed respiratory illness symptoms, a serum sample and a nasal and an oropharyngeal swab were obtained. Swabs were tested by RT-PCR for SARS-CoV-2 infection. If positive, the subject was followed and swabs collected on days three, six, nine, and 14 after the first swab day and a serum sample obtained on day 14. All subjects residing with the index case were swabbed following the same sampling schedule. Sera were collected from cohort participants in October 2020 to assess seroprevalence. Swabs were tested by RT-PCR. Sera were tested by Microneutralization Assay to measure the neutralizing antibody titer. Incidence of COVID-19, household secondary attack rate, and seroprevalence in the cohort were determined. The incidence of COVID-19 was 6.9% and the household secondary attack rate was 89.8%. Transmission within households occurred within two-days of confirming the index case. Infections were asymptomatic or mild with symptoms resolving within 10 days. The majority developed a neutralizing antibody titer by day 14 post onset. The overall seroprevalence among cohort participants was 34.8%. These results suggest that within-household transmission is high in Egypt. Asymptomatic or mild illness is common. Most infections seroconvert and have a durable neutralizing antibody titer.

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Author summary

SARS-CoV-2 virus is transmitted via close contact with infected persons, airborne droplets, and contaminated surfaces. People in closed settings with COVID-19 patients, such as healthcare workers and household contacts, were more likely to become infected. Egypt, like all other countries, is currently facing the COVID-19 pandemic. However, data on incidence and seroprevalence in this country is lacking. We followed 290 households participating in an ongoing cohort study of influenza and coronavirus incidence and prevalence. The incidence of COVID-19 was 6.9% and the household secondary attack rate was 89.8%. Transmission within households occurred within two-days of confirming the index case. Infections were asymptomatic or mild with symptoms resolving within 10 days. The majority developed a neutralizing antibody titer by day 14 post onset. The overall seroprevalence among cohort participants was 34.8%. The majority of subjects with three months apart paired-samples continued to be seropositive. Within-household transmission is high. Asymptomatic or mild illness is common. Most infections seroconvert and have a durable neutralizing antibody titer. Increasing awareness among the general public about proper ways of dealing with infections within the household may contribute to decreasing the spread in Egypt and areas with similar cultural background and population structure.

Introduction

The Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) that causes the coronavirus disease 2019 (COVID-19) is now a pandemic and a global crisis. Since 2003, three coronaviruses, the Severe Acute Respiratory Syndrome Coronavirus (SARS-CoV) in 2002–2003, the Middle East Respiratory Syndrome Coronavirus (MERS-CoV) in 2012, and SARS-CoV-2 (2019) emerged to infect human populations causing severe respiratory infections leading to death [1–3].

Since the first detection of SARS-CoV-2 in Wuhan, China, in December 2019, more than 105 million infected cases were reported including over 2.3 million confirmed deaths as of 9 February 2021 [4]. Old age, having chronic diseases such as diabetes, cardiovascular disorders, chronic respiratory illness, hypertension, cancer, and being a health care worker were reported as risk factors to increasing case fatality rate [5].

In Egypt, since the first travel-related case was announced on 14 February 2020, the cumulative infected cases reached more than 120,000 with more than 7,000 deaths [6]. Several interventions were implemented by the Egyptian authorities such as recommending wearing masks, social distancing, isolation of confirmed cases, quarantine of suspected cases and exposed people, personal hygiene, and closure of schools and airports. However, cases continued to be reported. The number of cases peaked between May and July 2020 with the observation of religious and cultural celebrations involving extended family gatherings within households associated with the Holy Month of Ramadan and Fitr Islamic Holiday. The Egyptian Ministry of Health shifted to household isolation for mild cases in May 2020 as compared to hospital isolation for all cases practiced earlier. By July, the number of cases started to decrease and several restrictions were lifted. Case counts remained low but started to increase in November yet remained below 500 cases per day (S1 Fig). Cases concentrated in greater Cairo and other cities but were also reported in rural areas across the country.

The virus is transmitted via close contact with infected persons, airborne droplets, and contaminated surfaces [7,8]. People in closed settings with COVID-19 patients, such as healthcare workers and household contacts, were more likely to become infected [5,9]. Evidence suggested that symptomatic people may spread COVID-19 through direct close contact among individuals rather than asymptomatic carriers [10]. Transmission studies conducted in China [11–14] and the USA [15,16] confirmed the transmissibility within households from primary to secondary cases. Yet, SARS-CoV-2 household person-to-person transmission studies are limited. Here, we followed 290 households participating in an ongoing cohort study of influenza and coronavirus incidence and prevalence. We show the clinical features, household secondary attack rate, and seroconversion among household residents.

Results

Incidence and household transmission

As of April 2020, a total of 1598 participants in 290 households at four study sites (Gharbiyah, Kafr El Sheikh, Qalyubiyah, and Fayyoum) were followed up. The median household size was five individuals (range 1–19). The demographic characteristics of those participants are shown in <u>S1 Table</u>. The mean age of the participants was 24 years and the median was 19 years. Around 44% were female and more than half were single/never married. Around a third did not report receiving any formal education, a third completed elementary school, 16% completed intermediate schooling, 6.5% completed schooling, and 9.7% attended college. Around 7% reported smoking, 2% had chronic breathing problems, and 10% reported having non-respiratory chronic diseases.

Fig 1A shows the monthly reported infections among cohort participants between April and September 2020. Most infections (n = 36) were detected in June but decreased in July and started increasing again in August and September. Additionally, two and three influenza A infections were detected in August and September, respectively.

Fig 1B summarizes the incidence of COVID-19 in the cohort. By the end of September 2020, a total of 801 participants (50.1%) developed clinical criteria that met our COVID-19 case definition (suspected infections). Of those, 23 subjects from 23 different households were confirmed by Real-time reverse-transcription polymerase chain reaction (RT-PCR) to be infected with SARS-CoV-2 and were hence considered index infections. Those index infections had 98 household contacts (range of 2–14 contacts per household) of whom 52 (53%) developed symptoms. Of the contacts, 66 (67%) were RT-PCR confirmed to be infected and



Fig 1. A: Number of infections by month in the cohort and in Egypt. B: Suspected, index, and secondary infections of COVID-19 in the cohort. https://doi.org/10.1371/journal.ppat.1009413.g001

63 of them seroconverted. Twenty-two of the RT-PCR negative contacts seroconverted. Thus, the total number of contact infections was 88 (66 RT-PCR confirmed and 22 seroconversions) and the total number of COVID-19 infections in the cohort was 111 (23 index infections and 88 contacts) (S2 Fig). The overall incidence rate in the cohort for the period April-September 2020 was 6.9% (95% confidence interval (CI): 5.8–8.3), the infection rate among suspected infections was 13.8% (95% CI: 11.6–16.4), and the household secondary attack rate was 89.8% (95% CI: 82.2–94.3). Demographic and general health characteristics of household infections did not statistically differ from non-infected cohort participants.

Transmission within household from index to contact infections was mostly within two days after confirmation of index case infection as 50 of the 88 contacts were RT-PCR positive at the first day of sampling (Fig 2). Five additional contacts became RT-PCR on day three, six, and nine of sampling. Only one contact became RT-PCR positive on day 14. The median age of index infections was 33 years (range 11–71 years) while the median age of the infected contacts was 22 years (range 1–80 years). The age difference between infected index and contact infections was not statistically significant. Among the index infections, 14 (60.9%) were females compared to 44 (50%) females among infected contacts (p-value >0.05).

All index infections had symptoms as compared to 52 contacts (Table 1). The most reported symptoms (>70%) among index infections were fever, cough, fatigue, muscle/body aches, headache, and loss of taste/smell. Muscle or body aches was the most commonly-reported symptom among contacts. Most symptoms resolved by day 10 post onset except for loss of



Fig 2. Transmission of COVID-19 infection among close contact households. Except for the dark blue legend, legends indicate the day for the first PCR positive test.

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Symptoms	Index infections (n = 23)		Contact infections (n = 52)	
	No. (%)	Median no. of days (range)	No. (%)	Median no. of days (range)
Fever	22 (95.6)	4 (1-18)	32 (61.5)	3 (1-14)
Cough	18 (78.3)	8 (2–25)	31 (59.6)	7 (2–28)
Shortness of breath	14 (60.9)	6 (1-21)	19 (36.5)	3 (1-25)
Fatigue	20 (87.0)	7 (2–25)	31 (59.6)	7 (2-30)
Muscle/Body aches	21 (91.3)	8 (2–28)	39 (75.0)	7 (3–22)
Headache	18 (78.3)	5 (2-11)	30 (57.7)	4 (1-22)
loss of taste/smell	20 (87.0)	14 (3–26)	28 (53.8)	10 (2-30)
Sore throat	16 (69.6)	5 (1-14)	31 (59.6)	4 (1-10)
Congestion/Runny nose	9 (39.1)	7 (3-14)	23 (44.2)	3 (1-15)
Nausea/Vomiting	12 (52.2)	7 (2-14)	13 (25.0)	4.5 (1-11)
Diarrhea	11 (47.8)	3 (1-8)	18 (34.6)	3 (1-7)
Blurred vision	8 (34.8)	4 (3-10)	11 (21.2)	4 (1-6)
Anorexia	16 (69.6)	9 (2-28)	22 (42.3)	7 (2-30)

Table 1. Occurrence and duration of symptoms among index and contact infections.

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taste or smell that lasted a median of 14 days among index infections. In general, contacts had fewer symptoms and had their illness resolved quicker than index infections. All infections were mild not requiring hospitalization.

By day 14 post initial RT-PCR confirmation, all index infections had a neutralizing antibody (nAb) titer (geometric mean titer (GMT) 49.4). Seven (30.4%) had a titer \geq 1:80. Among the 88 infected contacts, 85 (96.6%) had an antibody titer (GMT 45.3) of whom 34 (38.7%) had a titer \geq 1:80. The distribution of titers is shown in Fig 3. Three of the index infections had a nAb titer at day 1 of sampling that increased by day 14. Similarly, nine of the contact infections had a titer at day 1 that either increased or remained within 1-fold by day 14 (S2 Table).





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Cohort seroprevalence

All 212 sera collected in April 2020 tested negative by Microneutralization Assay (MN). Of those, 193 individuals were re-sampled in July and 30 of them sero-converted. The distribution of titers among the subjects who were tested in both April and July is shown in S3 Fig. Around 85% of those remained seronegative, 13.5% developed titers between 1:10 and 1:80, and 2% developed titers of 1:160 and above.

The distribution of titers in sera collected in July 2020 is shown in Fig 4. Around 80% were sero-negative, 18% had titers between 1:10 and 1:80, and 2% developed titers of 1:160 and above. The overall GMT was 1:6.9. Age was significantly associated with antibody levels (p-value = 0.017). The mean age of sero-negative participants was 29 years, 27 years for those with antibody titers between 1:80 and 1:160, and 40 years for those with titers greater than 1:160. None of the other variables was associated with antibody titers.

A total of 1260 sera were collected from cohort participants in October 2020. Of those, 438 (34.8%) were seropositive (95% CI: 32.2–37.4) (GMT 61.2). Only 238 (18.5%) participants had a titer \geq 1:80. The distribution of titers is shown in Fig 4. Of the females, 37.2% were positive as compared to 32.6% of the males (p-value >0.05). There was no statistically significant difference between the ages of the seropositive and seronegative participants.

Of participants who were seropositive in July 2020, 200 were resampled in October 2020. Among those, 69 (34.5%) had a stable titer (within 1-fold change) and 76 (38.0%) had an increase in titer (2 or more folds). Four had a decrease in titer (2 or more folds) but remained seropositive while 51 (25.5%) became seronegative. Titer variation among those 200 subjects is shown in S4 Fig.

Discussion

This paper is among the few that describe household transmission of COVID-19 and that show seroconversion and neutralizing antibody titers within households in a community rather than clinical settings. Furthermore, the results presented here are the firsts from a cohort study conducted in low-income country. The six-month incidence of COVID-19 in the cohort was 6.9%. The monthly distribution of new infections matched the distribution of cases in the general Egyptian population for the period April-July 2020 with cases peaking in June then decreasing in July. In Egypt, the first few cases were in March 2020 when the virus was introduced. By June-2020, COVID-19 infections surged potentially due to social activities linked to the last week of the Holy month of Ramadan and Eid-El-Fitr (the religious holiday following Ramadan). Cohort infections increased again in August and September while the number of cases reported in the general Egyptian population remained low. The increase in August coincides with another religious holiday Eid-El-Adha. This difference in incidence between the cohort and the general population may be due to testing only severe cases in the general population compared to testing all symptomatic infections and household contacts in the cohort. Hence, it is likely that the true incidence of disease in Egypt is close to that measured in the cohort. Almost half of the cohort participants reported a related symptom, a relatively high proportion given the time period of the study that is out of the season when respiratory diseases are spread. This may be due to the pandemic scare leading the subjects to become overzealous in self-reporting symptoms.

The household secondary attack rate in our study was more than 89% and is higher than rates reported in other studies. In a study from Wuhan, China, the secondary attack rate was 30% [12]. A similar rate (32%) was calculated in a study in Zhuhai, China [11]. Lower rates were estimated for China in other studies [13,17]. In the USA, secondary attack rates were estimated between 0.5-53% [15,16,18,19]. In Canada, a 14.7% secondary infection rate was calculated [20]. A meta-analysis estimated the rate to be around 17% [21]. Our higher rate may be explained by behavioral factors or by household size. Residents of households of our study may not have isolated themselves from index infections or may not have used proper personal protective equipment as recommended by public health authorities. The median household size in our study was five, potentially larger than that in China or North America, with several individuals sharing same bedrooms. Using protective equipment was shown to reduce secondary attack rates in household settings [22]. Some of the contacts became RT-PCR positive by day 9 or day 14 and hence may potentially be tertiary rather than secondary infections. If those were to be excluded from being secondary, then the secondary attack rate is reduced to 83.7%. Furthermore, there were four contacts that were seropositive at day 0 and did not have an increase in their antibody titer and may have not been related to the household transmission event. If those were excluded, the household secondary attack rate becomes 79.6%. Nonetheless, the household secondary attack rate in our cohort remains high.

Most secondary infections occurred within two days of confirming the index case. This may not be indicative of the true incubation period as the index case was tested when symptomatic. About a third of the secondary infections did not have a positive RT-PCR test probably due to low viral load at the time of sampling or due to improper sampling.

Asymptomatic infection was observed in our study as was detected in other studies with more than 40% of the confirmed secondary infections not showing symptoms [23]. A wide range of symptoms was observed including respiratory and gastro-intestinal manifestations. All the infections had relatively mild disease with symptoms resolving in less than 10 days. Only loss of smell or taste lasted for a median of 14 days. Overall, disease among secondary infections appears to be milder than among index infections with less symptoms reported and shorter duration of illness.

Our data indicate that the majority of the infections we followed (70%) developed nAb titers by day 14. Titers ranged from 1:10 to 1:640. Due to the short time period of follow-up (14 days), it is plausible that detected titers may increase over time and seroconversion may occur in infections that did not seroconvert by day 14. Twelve of the infections we followed had a detectable antibody titer at day of onset. Of those, eight were PCR-confirmed to be infected and eight became symptomatic. The person who did not have a positive PCR did not show any symptoms and had a titer of 1:160 and this titer did not increase over time. This suggests that low-level antibody titers may not fully protect against COVID-19 infection. An alternative explanation is that the detected antibody titer may have been elicited due to infection that has been ongoing before our follow-up commenced or that those infections may have been asymptomatic index infections. Indeed, most of the infections that had a titer at day 1 either stayed within 1-fold or increased by day 14 indicating a progressive antibody response.

Our cohort-wide serological testing revealed that a third of participants were seropositive. This rate increased from zero in April to 20% in July to above 30% in October. This shows that infection is more common than what we detected suggesting that very mild or asymptomatic infections occur frequently. Furthermore, the majority of subjects who were positive three months before the sampling conducted in October continued to have a detectable antibody titer suggesting the durability of neutralizing antibodies over time as indicated by other studies [24]. In a sero-survey among blood donors in Brazil, anti-SARS-CoV-2 antibodies were detected in 4% of the samples [25]. In another population-based study in Brazil, the seroprevalence rate was 0.05% [26]. In the USA, seroprevalence in Los Angeles county was estimated at 4.7% [27]. In a larger study involving several states, calculated seroprevalence rates ranged between 1% and 6.9% [28]. In a population-based study in Geneva, Switzerland, 4.8% of the participants had anti-SARS-CoV-2 IgG [29]. The largest seroprevalence study was conducted in Spain among more than 61,000 participants. Seropositivity was around 5% [30].

In comparison to the above-mentioned reports, our study is the only one that relied on neutralizing antibodies to determine seroprevalence of SARS-CoV-2 antibodies. This could be explained by the presence of COVID-19 case clusters in the villages where the subjects were sampled. However, the measured titers are relatively low as only a small percentage of tested sera had antibody titers greater than 1:160. This could potentially indicate that the majority of those who sero-converted had asymptomatic or mild disease [31]. Another explanation may be related to the fact that we only considered serum dilutions that totally protected the cells from CPE as positive.

Relying on the presence of symptoms in an index case to study within-household transmission is a limitation for this study especially with the commonality of having very mild and asymptomatic COVID-19 patients. This could have led to an underestimation of the calculated overall incidence within the cohort. Additionally, the calculated household secondary attack rates may be biased if the missed asymptomatic index infections did not contribute as much as symptomatic index infections to household transmission. Another limitation is that some contacts may have been asymptomatically infected when the index infection was confirmed hence not meeting the definition of a secondary infection. Data from this study may not be generalizable to the general Egyptian population due to the small sample size and geographic distribution of households. By design, However, our data strongly indicate that household transmission is playing an important role in the spread of COVID-19 in Egypt. It has been previously shown that COVID-19 household transmission and superspreading events may be the main drivers of disease spread [32,33]. Increasing awareness among the general public about proper ways of dealing with cases within the household may contribute to decreasing the spread in Egypt and areas with similar cultural background and population structure.

Materials and methods

Ethics statement

Ethical approval for the study was granted by the IRBs of St. Jude Children's Research Hospital (USA) (reference number 007079, dated March 20, 2020) and Human Link (Lebanon) (dated

March 23, 2020) as well as the Research Ethics Committee of the National Research Centre (Egypt) (protocol number 14 155, dated March 22, 2020). Written informed consent was obtained from all subjects over 18 years old, written assent was obtained for children between 14 and 17 years old, parental written consent was obtained for all participants less than 18 years old.

Cohort study design

Details of the study design and protocol have been previously published [34]. Briefly, households raising backyard poultry were selected from five villages in four Nile Delta governorates (Sharkiyah, Gharbiyah, Kafr El Sheikh, and Qalyubiyah) and Fayyoum governorate starting August 2015. All individuals within the household who were older than two years were invited to participate. Baseline enrollment was completed in March 2017. A total of 2402 subjects were enrolled from 390 households in the five study sites. The study protocol was amended in April 2020, to allow COVID-19 incidence and seroprevalence studies in four study sites (Fayyoum, Gharbiyah, Kafr El Sheikh, and Qalyubiyah) comprising 290 households.

To determine infection rates, study staff visited enrolled households on a weekly basis to check whether any study participant was reporting respiratory illness symptoms of fever of 38°C or higher, cough, sore throat, or shortness of breath. When a study participant was verified to have symptoms, a serum sample was collected, and a nasal swab and an oropharyngeal swab were obtained and tested by RT-PCR for SARS-CoV-2 infection. If any of the swabs tested positive, the subject was considered an index infection, and the study team obtained nasal and oropharyngeal swabs from the participant on days three, six, nine, and 14 after the first swab day as well as a serum sample on day 14. Furthermore, all previously enrolled subjects residing with the index case were swabbed and blood samples obtained following the same sampling schedule two days after the diagnosis of the index case. A symptoms data were collected by the study staff.

To determine cohort seroprevalence, sera were collected from 212 subjects in April 2020, 1,244 subjects in July, and 1,260 subjects in October to assess seroprevalence among participants.

Study staff interacted with cohort subjects following strict biosafety procedures including wearing N95 masks, gloves, disposable gowns, and hair and shoe covers. Disposed personal protective equipment were placed in biohazard bags and transported to the lab for proper disinfection and disposal. All staff were rigorously trained on all aspects of study protocol and biosecurity measures.

Viral testing

Swab samples were subjected to viral RNA extraction using QIAamp Viral-RNA Kit (Qiagen, Germany) according to manufacturer's instructions. RT-PCR screening assays (E gene, N gene, ORF1b-nsp14, and RdRp gene assays) with gene specific primers and probes [35,36] were conducted using Verso 1-step qRT-PCR Kit (Thermo, USA). A 25 μ l total reaction included 5 μ l template RNA, 12.5 μ l 2x one-step buffer, 1 μ l of each forward and reverse primers (10 μ M), 0.5 μ l probe (10 μ M), 1.25 μ l RT-Enhancer, 0.25 μ l enzyme mixture, and 3.5 μ l ddH₂O. Thermal cycling started with 15 min at 50°C (reverse transcription), 95°C for 15 min (polymerase activation), 45 cycles at 95°C for 15 s (denaturation), then 60°C for 30 s (annealing and extension). The positive control was a synthetic plasmid designed in-house including selected nucleotides that are detected by all used assays.

Serological testing

MN was conducted to measure the nAb titer in human sera using Vero-E6 (ATCC, CRL-1586) cell monolayers using SARS-CoV-2/Egypt/NRC-03/2020 under biological safety level 3 [37]. This virus was isolated from a swab of a confirmed patient in Egypt (passage 0) and cultured again in Vero-E6 cells to increase titers (passage 1). Stocks used for the MN assay were from passage 2 of the virus. Briefly, the collected sera were inactivated at 56°C for 1 hr. Sera were serially diluted two-fold from 1:10 to 1:1280 in DMEM media supplemented with 4% BSA, 1% antibiotic antimycotic (Gibco, USA), then mixed with equal volume of 100 tissue culture infectious dose (TCID₅₀/mL) of SARS-CoV-2/Egypt/NRC-03/2020 isolate and incubated for 1 hr at 37°C. A total volume of 35 µl of the virus–sera mix was inoculated in duplicate to Vero-E6 cell in a 96-well tissue culture plates. After 1 hr of incubation at 37°C, the inoculums were removed. The plates were then incubated for three more days at 37°C in 5% CO₂ in a humidified incubator. A virus back-titration was performed without immune serum to confirm TCID₅₀ viral titer used. Cytopathic effect (CPE) was observed post 72 hrs of infection. The reciprocal of the serum dilution that protected cells from CPE was considered the nAb titer. Negative sera were given a value of 1:5.

Statistical analysis

Data analysis was performed using SPSS v23 (IBM, Armonk, NY). Student's t-test was used to compare means and chi-square to compare categories. A p-value ≤ 0.05 was considered statistically significant.

Supporting information

S1 Table. Distribution of demographic and health data of the study participants. (DOCX)

S2 Table. Characteristics of seropositive infections at baseline. (DOCX)

S1 Fig. Cumulative number of COVID-19 cases and deaths in Egypt. (DOCX)

S2 Fig. Household transmission and seroprevalence of COVID-19 infection among close contact households. (DOCX)

S3 Fig. Seroconversion neutralizing antibody titers among cohort participants, April to July 2020.

(DOCX)

S4 Fig. Sero-dynamics of COVID 19 antibodies in 200 subjects who tested positive in July and were resampled in October. (DOCX)

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References

- Drosten C, Gunther S, Preiser W, van der Werf S, Brodt HR, Becker S, et al. Identification of a novel coronavirus in patients with severe acute respiratory syndrome. The New England journal of medicine. 2003; 348(20):1967–76. https://doi.org/10.1056/NEJMoa030747 PMID: 12690091.
- Zaki AM, van Boheemen S, Bestebroer TM, Osterhaus AD, Fouchier RA. Isolation of a novel coronavirus from a man with pneumonia in Saudi Arabia. The New England journal of medicine. 2012; 367 (19):1814–20. https://doi.org/10.1056/NEJMoa1211721 PMID: 23075143.
- Zhu N, Zhang D, Wang W, Li X, Yang B, Song J, et al. A Novel Coronavirus from Patients with Pneumonia in China, 2019. The New England journal of medicine. 2020; 382(8):727–33. Epub 2020/01/25. https://doi.org/10.1056/NEJMoa2001017 PMID: 31978945; PubMed Central PMCID: PMC7092803.
- WHO. COVID-19 Weekly Epidemiological Update 2021 [12/2/2021]. Available from: https://www.who. int/docs/default-source/coronaviruse/situation-reports/20210209_weekly_epi_update_26.pdf?sfvrsn= 836a69b9_3&download=true.
- Wu Z, McGoogan JM. Characteristics of and Important Lessons From the Coronavirus Disease 2019 (COVID-19) Outbreak in China: Summary of a Report of 72314 Cases From the Chinese Center for Disease Control and Prevention. JAMA. 2020. Epub 2020/02/25. https://doi.org/10.1001/jama.2020.2648 PMID: 32091533.
- WHO. Coronavirus disease (COVID-19) Situation report—187, 27 July 2020. 2020 [27–7–2020]. Available from: https://www.who.int/emergencies/diseases/novel-coronavirus-2019/situation-reports/.
- van Doremalen N, Bushmaker T, Morris DH, Holbrook MG, Gamble A, Williamson BN, et al. Aerosol and Surface Stability of SARS-CoV-2 as Compared with SARS-CoV-1. The New England journal of medicine. 2020; 382(16):1564–7. Epub 2020/03/18. https://doi.org/10.1056/NEJMc2004973 PMID: 32182409; PubMed Central PMCID: PMC7121658.
- Xu Y, Li X, Zhu B, Liang H, Fang C, Gong Y, et al. Characteristics of pediatric SARS-CoV-2 infection and potential evidence for persistent fecal viral shedding. Nat Med. 2020; 26(4):502–5. Epub 2020/04/ 15. https://doi.org/10.1038/s41591-020-0817-4 PMID: <u>32284613</u>; PubMed Central PMCID: PMC7095102.
- Pitzer VE, Leung GM, Lipsitch M. Estimating variability in the transmission of severe acute respiratory syndrome to household contacts in Hong Kong, China. Am J Epidemiol. 2007; 166(3):355–63. Epub 2007/05/12. https://doi.org/10.1093/aje/kwm082 PMID: 17493952; PubMed Central PMCID: PMC7110150.
- WHO. Transmission of SARS-CoV-2: implications for infection prevention precautions 2020 [26/7/ 2020]. Available from: https://www.who.int/news-room/commentaries/detail/transmission-of-sars-cov-2-implications-for-infection-prevention-precautions.
- Wu J, Huang Y, Tu C, Bi C, Chen Z, Luo L, et al. Household Transmission of SARS-CoV-2, Zhuhai, China, 2020. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. Epub 2020/05/12. https://doi.org/10.1093/cid/ciaa557 PMID: 32392331.

- Wang Z, Ma W, Zheng X, Wu G, Zhang R. Household transmission of SARS-CoV-2. The Journal of infection. 2020. Epub 2020/04/14. <u>https://doi.org/10.1016/j.jinf.2020.03.040</u> PMID: <u>32283139</u>; PubMed Central PMCID: PMC7151261.
- Li W, Zhang B, Lu J, Liu S, Chang Z, Cao P, et al. The characteristics of household transmission of COVID-19. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. Epub 2020/04/18. <u>https://doi.org/10.1093/cid/ciaa450</u> PMID: <u>32301964</u>; PubMed Central PMCID: PMC7184465.
- Chan JF-W, Yuan S, Kok K-H, To KK-W, Chu H, Yang J, et al. A familial cluster of pneumonia associated with the 2019 novel coronavirus indicating person-to-person transmission: a study of a family cluster. Lancet. 2020; 395(10223):514–23. Epub 01/24. https://doi.org/10.1016/S0140-6736(20)30154-9 PMID: 31986261.
- Burke RM, Midgley CM, Dratch A, Fenstersheib M, Haupt T, Holshue M, et al. Active Monitoring of Persons Exposed to Patients with Confirmed COVID-19—United States, January-February 2020. MMWR Morbidity and mortality weekly report. 2020; 69(9):245–6. Epub 2020/03/07. https://doi.org/10.15585/ mmwr.mm6909e1 PMID: 32134909.
- Rosenberg ES, Dufort EM, Blog DS, Hall EW, Hoefer D, Backenson BP, et al. COVID-19 Testing, Epidemic Features, Hospital Outcomes, and Household Prevalence, New York State-March 2020. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. https:// doi.org/10.1093/cid/ciaa549 PMID: 32382743; PubMed Central PMCID: PMC7239264.
- Jing QL, Liu MJ, Zhang ZB, Fang LQ, Yuan J, Zhang AR, et al. Household secondary attack rate of COVID-19 and associated determinants in Guangzhou, China: a retrospective cohort study. Lancet Infect Dis. 2020. https://doi.org/10.1016/S1473-3099(20)30471-0 PMID: 32562601.
- 18. Grijalva CG, Rolfes MA, Zhu Y, McLean HQ, Hanson KE, Belongia EA, et al. Transmission of SARS-COV-2 Infections in Households—Tennessee and Wisconsin, April-September 2020. MMWR Morbidity and mortality weekly report. 2020; 69(44):1631–4. https://doi.org/10.15585/mmwr.mm6944e1 PMID: 33151916; PubMed Central PMCID: PMC7643897 Journal Editors form for disclosure of potential conflicts of interest. Carlos G. Grijalva reports personal consulting fees from Sanofi, Merck, and Pfizer; grants from Sanofi, Campbell Alliance, the National Institutes of Health, the Agency for HealthCare Research and Quality, and a contract from the Food and Drug Administration, outside the submitted work. Natasha B. Halasa reports grants from Sanofi and Quidel and personal fees from Genetech, outside the submitted work. No other potential conflicts of interest were disclosed.
- Lewis NM, Chu VT, Ye D, Conners EE, Gharpure R, Laws RL, et al. Household Transmission of SARS-CoV-2 in the United States. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. https://doi.org/10.1093/cid/ciaa1166 PMID: 33185244; PubMed Central PMCID: PMC7454394.
- Wilkinson K, Chen X, Shaw S. Secondary attack rate of COVID-19 in household contacts in the Winnipeg Health Region, Canada. Can J Public Health. 2020. https://doi.org/10.17269/s41997-020-00451-x PMID: 33205377; PubMed Central PMCID: PMC7671665.
- Fung HF, Martinez L, Alarid-Escudero F, Salomon JA, Studdert DM, Andrews JR, et al. The household secondary attack rate of SARS-CoV-2: A rapid review. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. https://doi.org/10.1093/cid/ciaa1558 PMID: 33045075; PubMed Central PMCID: PMC7665336.
- Wang Y, Tian H, Zhang L, Zhang M, Guo D, Wu W, et al. Reduction of secondary transmission of SARS-CoV-2 in households by face mask use, disinfection and social distancing: a cohort study in Beijing, China. BMJ Glob Health. 2020; 5(5). https://doi.org/10.1136/bmjgh-2020-002794 PMID: 32467353; PubMed Central PMCID: PMC7264640.
- Luo Y, Trevathan E, Qian Z, Li Y, Li J, Xiao W, et al. Asymptomatic SARS-CoV-2 Infection in Household Contacts of a Healthcare Provider, Wuhan, China. Emerg Infect Dis. 2020; 26(8):1930–3. <u>https://doi.org/10.3201/eid2608.201016 PMID: 32330112</u>.
- Wajnberg A, Amanat F, Firpo A, Altman DR, Bailey MJ, Mansour M, et al. Robust neutralizing antibodies to SARS-CoV-2 infection persist for months. Science. 2020. <u>https://doi.org/10.1126/science.</u> abd7728 PMID: 33115920.
- Amorim Filho L, Szwarcwald CL, Mateos SOG, Leon A, Medronho RA, Veloso VG, et al. Seroprevalence of anti-SARS-CoV-2 among blood donors in Rio de Janeiro, Brazil. Rev Saude Publica. 2020; 54:69. https://doi.org/10.11606/s1518-8787.2020054002643 PMID: 32638883; PubMed Central PMCID: PMC7334006.
- Silveira MF, Barros AJD, Horta BL, Pellanda LC, Victora GD, Dellagostin OA, et al. Population-based surveys of antibodies against SARS-CoV-2 in Southern Brazil. Nat Med. 2020. <u>https://doi.org/10.1038/s41591-020-0992-3</u> PMID: 32641783.
- Sood N, Simon P, Ebner P, Eichner D, Reynolds J, Bendavid E, et al. Seroprevalence of SARS-CoV-2-Specific Antibodies Among Adults in Los Angeles County, California, on April 10–11, 2020. JAMA.

2020. https://doi.org/10.1001/jama.2020.8279 PMID: 32421144; PubMed Central PMCID: PMC7235907.

- Havers FP, Reed C, Lim T, Montgomery JM, Klena JD, Hall AJ, et al. Seroprevalence of Antibodies to SARS-CoV-2 in 10 Sites in the United States, March 23-May 12, 2020. JAMA Intern Med. 2020. <u>https:// doi.org/10.1001/jamainternmed.2020.4130</u> PMID: 32692365.
- Stringhini S, Wisniak A, Piumatti G, Azman AS, Lauer SA, Baysson H, et al. Seroprevalence of anti-SARS-CoV-2 IgG antibodies in Geneva, Switzerland (SEROCoV-POP): a population-based study. Lancet. 2020. https://doi.org/10.1016/S0140-6736(20)31304-0 PMID: 32534626; PubMed Central PMCID: PMC7289564.
- Pollan M, Perez-Gomez B, Pastor-Barriuso R, Oteo J, Hernan MA, Perez-Olmeda M, et al. Prevalence of SARS-CoV-2 in Spain (ENE-COVID): a nationwide, population-based seroepidemiological study. Lancet. 2020. https://doi.org/10.1016/S0140-6736(20)31483-5 PMID: 32645347; PubMed Central PMCID: PMC7336131.
- Lynch KL, Whitman JD, Lacanienta NP, Beckerdite EW, Kastner SA, Shy BR, et al. Magnitude and kinetics of anti-SARS-CoV-2 antibody responses and their relationship to disease severity. Clinical infectious diseases: an official publication of the Infectious Diseases Society of America. 2020. https:// doi.org/10.1093/cid/ciaa979 PMID: 32663256.
- Liu Y, Eggo RM, Kucharski AJ. Secondary attack rate and superspreading events for SARS-CoV-2. Lancet. 2020; 395(10227):e47. https://doi.org/10.1016/S0140-6736(20)30462-1 PMID: 32113505; PubMed Central PMCID: PMC7158947.
- Xu X, Liu X, Wang L, Ali ST, Du Z, Bosetti P, et al. Household transmissions of SARS-CoV-2 in the time of unprecedented travel lockdown in China. medRxiv. 2020. https://doi.org/10.1101/2020.03.02. 20029868 PMID: 32511615; PubMed Central PMCID: PMC7276042.
- El Rifay AS, Elabd MA, Abu Zeid D, Gomaa MR, Tang L, McKenzie PP, et al. Household Transmission of Zoonotic Influenza Viruses in a Cohort of Egyptian Poultry Growers. JMIR research protocols. 2015; 4(2):e74. <u>https://doi.org/10.2196/resprot.4331</u> PMID: <u>26099368</u>; PubMed Central PMCID: PMC4526956.
- Chu DKW, Pan Y, Cheng SMS, Hui KPY, Krishnan P, Liu Y, et al. Molecular Diagnosis of a Novel Coronavirus (2019-nCoV) Causing an Outbreak of Pneumonia. Clinical chemistry. 2020; 66(4):549–55. Epub 2020/02/08. https://doi.org/10.1093/clinchem/hvaa029 PMID: 32031583; PubMed Central PMCID: PMC7108485.
- Corman VM, Landt O, Kaiser M, Molenkamp R, Meijer A, Chu DKW, et al. Detection of 2019 novel coronavirus (2019-nCoV) by real-time RT-PCR. Eurosurveillance. 2020; 25(3):2000045. <u>https://doi.org/10.2807/1560-7917.ES.2020.25.3.2000045</u> PMID: 31992387.
- 37. Perera RA, Wang P, Gomaa MR, El-Shesheny R, Kandeil A, Bagato O, et al. Seroepidemiology for MERS coronavirus using microneutralisation and pseudoparticle virus neutralisation assays reveal a high prevalence of antibody in dromedary camels in Egypt, June 2013. Euro Surveill. 2013; 18(36):pii = 20574. https://doi.org/10.2807/1560-7917.es2013.18.36.20574 PMID: 24079378.