

1 “Do patients infected with human coronavirus before the Covid-19 pandemic have
2 less risk of being infected with Covid-19? “

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5 **Abstract**

6 **Background/Aim:** Although seasonal human coronaviruses (HCoV) have been known
7 as respiratory tract viruses for many years, the newly identified SARS-CoV-2 caused a
8 pandemic associated with severe respiratory failure. We aimed to evaluate the incidence
9 of COVID-19 infection in patients diagnosed in three tertiary teaching hospitals with
10 and without prior confirmed HCoV infection and compare these cohorts in terms of
11 COVID-19 contraction.

12 **Materials and methods:** In our study, HCoV PCR positive cases obtained
13 retrospectively between June 2014 and March 2020 in three University Hospital
14 Microbiology Laboratories (Cohort 1) and PCR negative patients detected in the same
15 PCR cycle of PCR positive cases (Cohort 2) were examined. Subgroups of HCoV
16 positive cases were evaluated.

17 **Results:** There was no difference between Cohort 1 vs 2 ($p=0.724$). When previous
18 HCoV subgroups of COVID-19 positive patients were examined, no significant
19 difference was found between the betacoronavirus and alphacoronavirus subgroups
20 ($p=0.822$) and between the four groups ie NL63, 229E, OC43, HKU-1 ($p=0.207$) and
21 OC43 subgroup vs others ($p=0.295$).

1 **Conclusion:** Being previously infected with HCoV did not provide protection against
2 COVID-19 in our study group. We suggest evaluating the possible effect of previous
3 OC43 infection on COVID-19 contraction in larger cohorts.

4 **Keywords:** HCoV, Covid-19, OC43, multiplex PCR, prevention, epidemiology

5 **1. Introduction**

6 Humancoronaviruses (HCoV) are large enveloped, positive-stranded RNA viruses
7 divided into four groups. The globally endemic subtypes are HCoV 229E, NL63, OC43
8 and HKU1 [1]. These viruses, are called non-SARS CoV. They may cause up to 1/3 of
9 adult community-acquired upper respiratory tract infections. However, SARS-CoV or
10 MERS-CoV present often with acute respiratory distress syndrome (ARDS) and both
11 cause epidemics with high mortality [2].

12 The SARS-CoV-2 (COVID-19) virus, which was firstly identified in December 2019,
13 spread rapidly all over the World [3]. COVID-19 has a spectrum of asymptomatic
14 infection to mild/moderate pneumonia and/or severe respiratory syndrome with fatal
15 outcomes [1,4,5]. All HCoV including SARS-CoV-2 may activate both innate and
16 adaptive immune responses in infected patients. Theoretically the antigenic similarity
17 (as well as antibodies or immunity to those antigens) between COVID-19, and other
18 HCoV may cause cross protection. Herein, we aimed to evaluate the incidence of
19 COVID-19 infection in patients with and without prior confirmed HCoV infection.

20 **2. Material and methods**

1 This retrospective cohort study was reported according to STROBE Criteria¹ was
2 conducted in three different tertiary-care educational hospitals located in two different
3 cities populated 4.320.519² and 5.663.322³.

4 In this study we compared the incidence of COVID-19 in two different cohorts;
5 Cohort 1 inclusion criterion was to be diagnosed with HCoV infection by respiratory
6 specimen PCR (polymerase chain reaction) between January 2014 and 10 March 2020
7 (the first COVID-19 case in Turkey was seen on March 11, 2020) in the study centers.
8 Exclusion criteria were dying before the COVID-19 outbreak and being under 18 years
9 of age (Figure).

10 Cohort 2 inclusion criterion was to have negative PCR detected in the same PCR cycle
11 of cases in Cohort 1 in the study centers. Whenever possible each Cohort 1 patient was
12 matched with one Cohort 2 patient. Exclusion criteria were dying before the COVID-19
13 outbreak and being under 18 years of age (Figure).

14 COVID-19 contraction data of the Cohort 1 and 2 were retrieved from the National
15 Hospital Health Management System (HSYS-www.hsys.saglik.gov.tr) on 10 December
16 2020. We chose the 10th December 2020 as the cut off analysis time since COVID-19
17 vaccination started by that date.

18 We further performed a subgroup analysis to evaluate the possible effect of more recent
19 immune response. In order to evaluate the more recent immune response, COVID-19
20 contraction results of 112 patients, who were found to be HCoV positive between

¹ STROBE (2023) Checklist [online] Website <https://www.strobe-statement.org/checklists/> [accessed 20 Aug 2023].

² Turkish Statistical Institute (TSI) (2021) İstatistiklerle İzmir [online]. Website <http://www.izmir.gov.tr/istatistiklerle-izmir> [accessed 13 Aug 2021].

³ The Governorship of Ankara (2021) İlçelerimiz. [online]. Website <http://www.ankara.gov.tr/ilcelerimiz> [accessed 13 Aug 2021]

1 March 2019 and March 2020 were compared with control cases corresponding to the
2 definition in Cohort 2.

3 This study was approved by the Scientific Research Platform of the Ministry of Health
4 (2021-04-29T00_04_05) and Ege University Medical Research Institutional Review
5 Board (2023-1200 23-7.1T/7).

6 Since there were no similar studies asking the same research question at the time we
7 planned and collected the study data, we could not calculate the sample size, but a
8 retrospective power analysis was performed (www.OpenEpi.com). Nevertheless, by
9 considering number of exposed: 304, risk of disease among exposed: 8.9%, number of
10 non-exposed: 297 and risk of disease among non-exposed: 8.1%, the power of the study
11 was 5% which may be considered as low.

12 IBM Statistical Package for Social Sciences (SPSS), version 25 for Windows was used
13 for statistical analysis. Comparisons were made with Chi-square test. A (two-sided)
14 *p* value lower than .05 was accepted as statistically significant.

15 **3. Results**

16 A total of 786 adult cases fulfilled the criteria to be included in Cohort 1 or 2. However,
17 185 cases were excluded from the study because they died before the COVID-19
18 pandemic.

19 Cohort 1 comprised 304 HCoV positive patients (159 females, aged 47.97 ± 18.13 and
20 Cohort 2 included 297 negative control cases (145 females, aged 49.96 ± 19.03). Age and
21 gender did not differ significantly between the cohorts (Table 1). (Figure).

1 When the subgroups of HCoV positive patients in cohort 1 were examined, the
2 alphacoronavirus (229E, NL63, 229E / NL63) rate was 61.8% (n = 188) while the
3 betacoronavirus (OC43, HKU1) rate was 31.9% (n = 97). The subgroup was not
4 determined in 6.3% (n = 19) (Table 1). Overall 8.9% (n = 27) of Cohort 1 were found to
5 be COVID-19 PCR positive. When the HCoV subgroups of the patients with COVID-
6 19 PCR positive were evaluated, 63% (n = 17) were alphacoronavirus, 29.6% (n = 8)
7 were betacoronavirus, and 7.4% (n = 2) did not have any subgroups (Table 2).

8 Cohort 2 comprised 297 patients (a negative control case was not present in all PCR
9 cycles) who were evaluated as the comparison group. Overall 8.1% (n = 24) of them
10 had COVID-19 PCR-positivity. There was no difference between Cohort 1 vs 2
11 (p=0.724, Table 1).

12 When previous HCoV subgroups of COVID-19 positive patients were examined, no
13 significant difference was found between the betacoronavirus and alphacoronavirus
14 subgroups (p=0.822, Table 2) and between the four groups ie NL63, 229E, OC43,
15 HKU-1 (p=0.211, Table 2) and OC43 subgroup vs others (5.2%-3/58 vs. 10.6%-22/227,
16 p=0.277, Table 2). Although there was no statistically significant difference between
17 coronavirus subgroups, the lowest incidence of COVID-19 was found in the OC43
18 subgroup with 5.2%, which was lower than Cohort 2 (Table 2).

19 The day-30 mortality rate of Cohort 1 and Cohort 2 due to COVID-19 did not differ
20 significantly [3.7% (n=1/27) in Cohort 1 vs. 4.2% (n=1/24) in Cohort 2 (p=0.932)], one
21 patient died on day 1 of the COVID-19 diagnosis and the other on day 6 of COVID-19
22 diagnosis.

1 When we checked the possible effect of the recent HCoV infection on COVID-19, the
2 incidence was 11% (12/109) and not lower in the subgroup of HCoV cases related to
3 March 2019-March 2020 period. Furthermore, cases infected by HCoV during the most
4 recent 3-month and 6-month period before March 2020; were also not low i.e. 11.4%
5 (4/35) in the last 3-month and 13.4% (9/67) in the last 6-month subgroup (Table 3). We
6 did not find any significant difference ($p>0.05$) in the rates of COVID-19 contraction
7 between individuals who had a recent HCoV infection and their matched control cases
8 (Table 3). Finally, since the lowest COVID-19 incidence was in the OC43 HCoV
9 infected subgroup, we analysed the effect of recent infection on COVID-19. COVID-19
10 incidence was 0/4, 0/9 and 1/12 in the subgroup infected with OC43 during the previous
11 3, 6 and 12-month period (comparisons with the incidences of the subgroup infected
12 with other HCoV were not significant $p=0.475$ $p=0.241$ $p=0.776$).

13 **4. Discussion**

14 In this study, we analysed whether HCoV, which act through the same receptors and
15 defense mechanisms as well as have not very high but a level of genetic or antigenic
16 similarity [6-8], have an effect on the prevention from COVID-19. However we
17 determined that previous infection with HCoV was not a protective factor for COVID-
18 19 in our cohort.

19 Anderson et al. [9] conducted a study to evaluate the relationship between seasonal
20 HCoV antibodies and COVID-19 contraction. HCoV antibodies were detected in most
21 of the 431 samples taken in the pre-pandemic period and ~20% of these individuals
22 possessed non-neutralizing antibodies that cross-reacted with SARS-CoV-2 spike and
23 nucleocapsid proteins. They reported that samples with pre-pandemic SARS-CoV-2-

1 reactive antibodies had elevated levels of antibodies against previously circulating
2 betacoronaviruses (especially OC43). However, these antibodies were not associated
3 with protection against SARS-CoV-2 infections or hospitalizations, but boosted upon
4 SARS-CoV-2 infection [9].

5 Similarly, Sagar et al. [10] evaluated the clinical relevance of COVID-19 infection and
6 HCoV infection in 875 previously confirmed HCoV-infected and 15,053 PCR negative
7 controls. SARS-CoV-2 PCR test was performed in 11.4% (n=1812) of a total of 15,928
8 patients, and 25.9% (n=470) of the tested patients were PCR positive. 53.6% (n=252) of
9 the SARS-CoV-2 infected patients were hospitalized, and there was no significant
10 difference in the frequency of hospitalization between the HCoV (+) and HCoV (-)
11 groups. When hospitalized HCoV (+) and HCoV (-) patients were evaluated, it was seen
12 that the HCoV (+) group required less intensive care unit stays (OR 0.1, 95% CI 0.0–
13 0.7) and a lower need for mechanical ventilators (OR 0.0, 95% CI 0.0–1.0). The rates of
14 patients who were hospitalized and died during follow-up were 17.7% in the HCoV(-)
15 group, whereas it was lower with 4.8% in the HCoV(+) group. With these results,
16 unlike our study, they determined that the previously positive group for HCoV was
17 associated with less severe disease and lower mortality rates compared to the HCoV
18 negative group [10]. However, in our study, we compared the all-cause mortality rates
19 rather than disease severity. We found that there was no significant difference in terms
20 of mortality between Cohort 1 and Cohort 2. Nevertheless, lack of difference may be
21 related to the relatively low numbers in both cohort 1 and 2.

22 Unlike the results of our study, Otlu et al. [11] found a lower incidence of COVID-19 in
23 64 patients with pre-existing HCoV infection compared to the current city (Malatya)
24 incidence during their study period. They used the National COVID-19 surveillance

1 data as we did and showed that four (6.2%) of 64 patients were infected with COVID-
2 19 by the end of 2020, while, simultaneously, the COVID-19 incidence in the city of
3 Malatya ranged from 7.8% (polymerase chain reaction-based diagnosis) to 9.2% (total
4 diagnosis). The differences were reported to be statistically significant (6.2% vs. 7.8%,
5 $p < .01$; 6.2% vs. 9.2%, $p < .001$). In our study, only the OC43 subgroup in Group 1,
6 among all HCoV subgroups, had a COVID-19 incidence of $<7.8\%$ (as reported in the
7 Malatya study) [11].

8 Our study is subject to several limitations. We tried to include all HCoV-positive cases
9 in the study centres after we started using multiplex PCR because the sample size could
10 not be calculated. Since there were no similar studies asking the same research question
11 at the time we planned and collected the study data, we could not calculate the sample
12 size, but a retrospective power analysis revealed the power of the study as 5% which
13 may be considered as low. Thus, our relatively low study sample might have hindered
14 us from demonstrating a potential preventive effect of previous HCoV infection on
15 COVID-19. Nonetheless, given that our data showed a higher incidence of COVID-19
16 in Cohort 1, the likelihood of such an effect appears to be lower. Unlike COVID-19
17 tests, the frequency of testing for seasonal coronaviruses, both pre-COVID and during
18 the pandemic, was typically limited in clinical practice due to the unavailability of
19 related kits (mostly because of reimbursement issues) in hospitals. As a result, this
20 might have significantly impacted the homogeneity of patient populations in the HCoV
21 group. Additionally, we were unable to compare our cohorts to the general population
22 as we do not have exact COVID-19 incidence data for Ankara, İzmir, or Turkey as a
23 whole as of December 10, 2020. We evaluated the patients in Cohort 1 based on their
24 PCR results related to pre-pandemic period. It is possible that some patients in Cohort 1

1 or 2 were infected with common HCoV before or after their HCoV PCR test results in
2 our study. Furthermore, we could not analyze the presence of serological and cellular
3 immunity related to HCoV or relationship of this immunity with the underlying diseases
4 in Cohorts 1 and 2. This means that patients in Cohort 2, who might have been infected
5 with HCoV, couldn't be identified. SARS-CoV and MERS-CoV cases, which are
6 among the betacoronaviruses, are very rare in Turkey [2]. Therefore, the results of these
7 factors are not included among the HCoV positive cases⁴. Another limitation to note is
8 that PCR-negative COVID-19 patients [5] were not included in the study. However, we
9 can confirm that cohort 1 was infected with a human coronavirus other than SARS-
10 CoV-2, and their COVID-19 infection status was analyzed using the most commonly
11 used method worldwide, which is PCR. Despite these limitations, this study is one of
12 the few that investigates whether a prior HCoV infection reduces the risk of contracting
13 COVID-19 [12]. Data were collected from three major university hospitals in two major
14 Turkish cities. To our knowledge, this is the most comprehensive study on this topic in
15 Turkey, a country that had one of the highest numbers of COVID-19 cases globally
16 during the study period. The study also examined the relationship between HCoV and
17 COVID-19, including its subgroups, and the impact of the near-term immunological
18 response.

19 In conclusion our data suggest that being previously infected with HCoV did not
20 provide protection against COVID-19 in our study group. We suggest evaluating the
21 possible effect of previous OC43 infection on COVID-19 contraction in larger cohorts.

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⁴ European Centre for Disease Prevention and Control (2023) Geographical distribution of confirmed cases of MERS-CoV, by reporting country, April 2012 – June 2023 [online]. Website <https://www.ecdc.europa.eu/en/publications-data/geographical-distribution-confirmed-cases-mers-cov-reporting-country-april-2012-7> [accessed 25 Jul 2023]

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1 **Conflict of interest**

2 The authors have no competing interests to declare.

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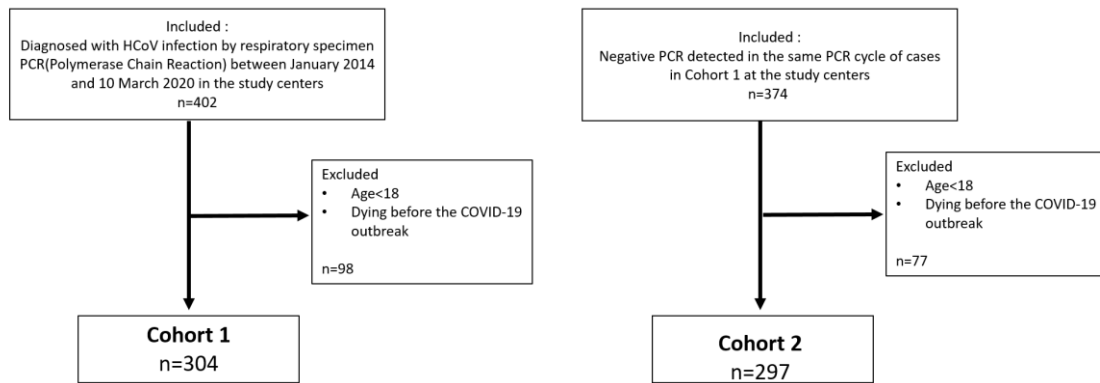
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1 Figure. Study inclusion and exclusion criteria and study population



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1 Table 1: Comparison of Covid-19 incidence in the study cohorts (as of 10 December
 2 2020)

	Cohort 1 (n =304)	Cohort 2 (n =297)	p
Female	159 (52.3%)	145 (48.8%)	0.393
Male	143 (47.7%)	154 (51.2%)	
Age	47.97±18.13	49.96±19.03	0.189
COVID-19 positivity (as of 10 December 2020)	27 (8.9%)	24 (8.1%)	0.724
The day-30 mortality rate due to COVID-19	1/27 (3.7%)	1/24 (4.2%)	0,932

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1 Table 2: Comparison of Covid-19 incidence in the Cohort 1 subgroups (as of 10
 2 December 2020)

		COVID-19 Negative	COVID-19 Positive	p
Alfacoronavirus(61.8%)		171/188 (90.9%)	17/188 (9.1%)	0.822
Betacoronavirus(31.9%)		89/97 (91.8%)	8/97 (8.2%)	
Non subgroup(6.3%)		17/19 (89.5%)	2/19 (10.5%)	
Alfacoronavirus (n=188)	229E/NL63	60/63 (95.2%)	3/63 (4.8%)	0.211
	229E	84/92 (91.3%)	8/92 (8.7%)	
	NL63	27/33 (81.8%)	6/33 (18.2%)	
Betacoronavirus (n=97)	OC43	55/58 (94.8%)	3/58 (5.2%)	
	HKU-1	34/39 (84.6%)	5/39 (15.4%)	

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- 1 Table 3: Comparison of the Covid-19 incidence in the group that contracted HCoV
- 2 during the March 2019-March 2020 period (as of December 10, 2020)

	HCoV Positive	Related controls	p
3-month (10 December 2021-10 March 2022)	4/35 (11.4%)	3/35 (8.6%)	0.690
6-month (10 September 2019- 10 March 2022)	9/67 (13.4%)	3/58 (5.2%)	0.117
12-month (10 March 2019-10 March 2020)	12/109 (11%)	6/97 (6.2%)	0.221

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