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#### **ORIGINAL ARTICLE**

# Renin–Angiotensin–Aldosterone System Inhibitors and Risk of Covid-19

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### **Abstract**

#### BACKGROUND

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There is concern about the potential of an increased risk related to medications that act on angiotensin-aldosterone system in patients exposed to coronavirus disease 2019 (Covid-19)

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blockers, beta-blockers, calcium-channel blockers, or thiazide diuretics and the likelihood of a positive or negative result on Covid-19 testing as well as the likelihood of severe illness (defined as intensive care, mechanical ventilation, or death) among patients who tested positive. Using Bayesian methods, we compared outcomes in patients who had been treated with these medications and in untreated patients, overall and in those with hypertension, after propensity-score matching for receipt of each medication class. A difference of at least 10 percentage points was prespecified as a substantial difference.

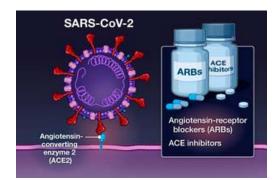
#### **RESULTS**

Among 12,594 patients who were tested for Covid-19, a total of 5894 (46.8%) were positive; 1002 of these patients (17.0%) had severe illness. A history of hypertension was present in 4357 patients (34.6%), among whom 2573 (59.1%) had a positive test; 634 of these patients (24.6%) had severe illness. There was no association between any single medication class and an increased likelihood of a positive test. None of the medications examined was associated with a substantial increase in the risk of severe illness among patients who tested positive.

#### CONCLUSIONS

We found no substantial increase in the likelihood of a positive test for Covid-19 or in the risk of severe Covid-19 among patients who tested positive in association with five common classes of antihypertensive medications.

Introduction



#### **QUICK TAKE**

Inhibitors of the Renin–Angiotensin–Aldosterone System and Covid-19

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EVERE ACUTE RESPIRATORY SYNDROME CORONAVIRUS 2 (SARS-COV-2), THE VIRUS that causes coronavirus disease 2019 (Covid-19), can infect host cells by means of interaction with membrane-bound angiotensin-converting enzyme 2 (ACE2) on respiratory er ACE2 is part of the renin–angiotensin–aldosterone system (RAAS) and its neurohormonal treatment with RAAS inhibitors can increase tissue expression of ACE2 and its presentation.

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observation that hypertension may be associated with an increased risk of death among patients with Covid-19. A large, multicenter study on hypertension and risk of Covid-19 indicated that 24% of patients with severe disease had hypertension, as compared with 14% of patients with nonsevere disease, although that analysis was not adjusted for other clinical features.<sup>9</sup>

However, ACE2 is protective in animal models of acute lung injury, and pretreatment with ACE inhibitors, ARBs, or beta-blockers may reduce the extent of experimentally induced lung injury and improve outcomes, an effect mediated by inhibition of the RAAS.<sup>2,10-15</sup> Thus, others have hypothesized that these medications could theoretically be beneficial, reducing the risk of severe disease among patients with Covid-19.<sup>2</sup>

Owing to the high global prevalence of hypertension (estimated to be 46% among adults in the United States), the relation between antihypertensive medications and Covid-19 outcomes is very important to public health.<sup>2,16</sup> These considerations led the Heart Failure Society of America, the American College of Cardiology, and the American Heart Association to issue a joint statement calling for immediate research into this issue.<sup>17</sup> We sought to estimate the association between the use of antihypertensive medications and the likelihood of a positive test for Covid-19 as well as the likelihood of severe Covid-19 (defined as intensive care, mechanical ventilation, or death) in a cohort of patients in a large health care network in New York City, an epicenter of the global Covid-19 pandemic.

Methods

#### PATIENT POPULATION

We identified all the patients in the New York University (NYU) Langone Health electronic health record who had Covid-19 test results recorded from March 1 to April 15, 2020, including tests sent to commercial laboratories, tests performed at our local laboratory, and tests ordered by NYU Langone Health providers and conducted at the New York City or State Department of Health. Patients were deemed to be Covid-19—positive if any test was positive for SARS-CoV-2 RNA and Covid-19—negative if all tests were negative.

#### MEDICATION AND HISTORY ASSESSMENT

For each identified patient with Covid-19 test results, we extracted medical history from the basis of discretely documented diagnostic codes that had been entered into the medical his problem list sections of the medical record or on the basis of encounter diagnoses, and we con-

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available with the full text of this article at NEJM.org). We extracted medication data for each patient, on the basis of the history in the electronic health record, and defined previous treatment with an antihypertensive medication as use of the medication within the preceding 18 months, provided that there was no evidence that the medication was discontinued at least 1 month before the Covid-19 test. The criteria for medication use included clinical documentation in the chart of prescribed medications. Medication lists and problem lists are updated frequently in our electronic health record by clinicians, who are routinely prompted by the system to cross-reference the medication list with pharmacy fill records and to ask patients about adherence.

We report here on the following classes of antihypertensive medication — ACE inhibitors, ARBs, beta-blockers, calcium-channel blockers, and thiazide diuretics. We also analyzed the composite of treatment with ACE inhibitors or ARBs, because they act on the same system and are rarely used together. In this study, we considered sacubitril–valsartan to be an ARB.

#### STUDY OVERSIGHT

Our observational study was designed by the authors and approved with a waiver of authorization and informed consent by the institutional review board of the NYU Grossman School of Medicine. The data were analyzed by three of the authors, who vouch for the analysis. Four of the authors vouch for the accuracy of the data. The first author prepared the first draft of the manuscript. Four of the authors had full access to the data and were responsible for the decision to submit the manuscript for publication.

#### **ANALYSIS OVERVIEW**

We calculated the percentages of patients who were Covid-19—negative and who were Covid-19—positive among tested patients, both overall and among patients with a history of hypertension as noted in the electronic health record. Patients with no medical record data other than the Covid-19 test result were excluded from the analysis. Among the patients with a positive Covid-19 test result, we identified patients with severe Covid-19, which was defined as admission to the intensive care unit (ICU), the use of invasive or noninvasive mechanical ventilation, or death. We computed the percentages of patients who met and those who did not meet this definition of severe illness as of April 15, 2020, among patients who tested positive for Covid-19.

To address confounding by indication and other sources of bias arising from the use of observational data, we estimated a propensity score for the likelihood of treatment with each medication class of interest, and we matched patients who had been treated with each medication class to thos not been treated with that medication class using a 1:1 ratio without replacement, accordin estimated propensity scores. We assessed the percentage of patients who tested positive in the likelihood of treatment with each medication class of interest, and we matched patients who had been treated with each medication class to those not been treated with that medication class using a 1:1 ratio without replacement, according to the likelihood of treatment with each medication class of the percentage of patients who tested positive in the likelihood of treatment with each medication class to those not been treated with that medication class using a 1:1 ratio without replacement, according to the likelihood of treatment with each medication class to those not been treated with that medication class using a 1:1 ratio without replacement, according to the likelihood of treatment with each medication class to those not been treated with that medication class using a 1:1 ratio without replacement, according to the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to those not be a second of the likelihood of treatment with each medication class to the likelihood of treatment with each medication class to the likelihood of treatment with each medi

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For a given medication class, we used propensity-score models that adjusted for the following variables — age; sex; race; ethnic group; body-mass index; smoking history; history of hypertension, myocardial infarction, heart failure, diabetes, chronic kidney disease, and obstructive lung disease (e.g., asthma and obstructive pulmonary diseases); and other classes of medication. Propensity-score matching was implemented with the use of a nearest-neighbor strategy. The caliper for the matching was specified in the nearest-neighbor strategy if the unspecified approach did not result in satisfactory balance. For body-mass index, the mean value was imputed for patients with missing data. Our goal was to rule out a clinically meaningful difference in risk, which we defined a priori as an absolute difference of at least 10 percentage points in the likelihood of a positive or negative test for Covid-19 and in the likelihood of the presence or absence of severe Covid-19. Our primary analysis involved patients with hypertension. We also analyzed data from all patients, without regard to hypertension.

Before and after creating the matched sets for each medication class, we computed the difference in the likelihood of the outcome between patients who had been treated and those who had not been treated with each medication class, and we sampled from the posterior distribution of the difference in these proportions using the posterior probabilities conditional on the outcomes in each medication group and a prior distribution. As our prior distribution, we assumed a Beta (a·b) distribution, a conjugate to the binomial distribution, with both shape parameters a and b equal to 1. Because very few data are currently available, these uninformative parameters produce prior probabilities of outcomes that are distributed uniformly between 0 and 1. We estimated the difference as the median of the posterior samples, along with a 95% credible interval for the difference. Association between a medication class and an outcome was considered to be present when the 95% credible interval excluded zero. A substantial difference was considered to be present when the lower boundary of the 95% credible interval exceeded zero and the upper boundary exceeded 10 percentage points. We conducted sensitivity analyses using traditional frequentist approaches rather than Bayesian approaches and using adjusted logistic regression rather than propensity-score matching. Analyses were performed with the use of R statistical software, version 3.6.1 (with the libraries MatchIt and BayesianFirstAid).



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Patient Population.

Table 1.

Characteristic	Patients with Hypertension (N = 4357)	All Patients (N = 12,594)
Median age (interquartile range) — yr	64 (54-75)	49 (14-63)
Female sex no. (%)	2143 (69.2)	7365 (58.5)
Race no. (%) (		
Black	875 (20.1)	1895 (15.0)
Asian	325 (7.5)	1109 (8.8)
White	2098 (48.2)	5862 (46.5)
Other	1019 (24.3)	3728 (29.6)
Hispanic othnic group — no. (%)†	662 (15.2)	2106 (16.7)
>1 Test for Covid-19 — no. (%)	693 (15.9)	1363 (10.8)
Smoking status no: (%)		
Current smicker	231 (5.3)	618 (5.1)
Former smoker	1200 (27.5)	2227 (17.7)
Never smoked	257) (59.1)	8420 (66.9)
Missing data	353 (8.1)	1309 (10.4)
Median body-mass index (interquartile range))	28.7 (24.9-13.1)	27.4 (24.7-21.9
History of hypertension — no. (%)	4357 (100)	4357 (34.6)
History of heart failure no. (%)	703 (16.1)	784 (6.2)
History of myocardial infarction no. (%)	460 (10.4)	325 (4.2)
History of diabetes no. (Ni)	1730 (19.7)	2272 (18.0)
History of chronic kidney disease no. (%)	1090 (25.0)	1214 (9.6)
History of obstructive lung disease — no. (%)	980 (22.5)	1833 (14.6)
Medications no. (%)		
ACE inhibitor	954 (21.9)	1044 (8.3)
ARE	1218 (28.4)	1328 (10.5)
ACE inhibitor or ARE	2141 (49.1)	2319 (18.4)
Beta blocker	1381 (91.7)	1686 (13.4)
Calcium-channel blocker	15.77 (36.2)	1672 (13.3)
Thiazide diuretic	903 (20.7)	989 (7.5)

Characteristics of the Study Population of Patients Tested for Covid-19.

As of April 15, 2020, a total of 12,594 patients had a Covid-19 test result and had an office, emergency department, or inpatient encounter in the NYU Langone Health system. Among these patients, 5894 (46.8%) had a positive test result and 6700 (53.2%) had a negative test result (**Figure 1**). Among the 5894 patients with a positive test, 1002 (17.0%) had severe Covid-19, as indicated by ICU admission (in 726 patients), mechanical ventilation (in 311), or death (in 447) by April 15, 2020. The characteristics of the patients in the study population are shown in **Table 1**.

#### MEDICATION USE AND COVID-19 TEST RESULTS IN PATIENTS WITH HYPERTENSION

Table 2.



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Among the 12,594 patients who underwent testing for Covid-19, a total of 4357 (34.6%) had a history of hypertension; of these patients, 2573 (59.1%) were Covid-19—positive and 1784 (40.9%) were Covid-19—negative. The results of Covid-19 testing stratified according to previous treatment with each medication class in matched patients are shown in Table 2. We ruled out an absolute difference of at least 10 percentage points in the likelihood of a positive test with at least 97.5% certainty for ACE inhibitors, ARBs, beta-blockers, calcium-channel blockers, and thiazide diuretics.

The results of the unmatched analyses are shown in Table S2. In the unmatched analyses, several medication classes were associated with a higher likelihood of testing positive for Covid-19, probably owing to confounding by characteristics of the patients who were treated. Results of propensity-score—matching quality are shown in Tables S4 through S9. The standardized mean differences for each class of medication was greatly reduced after matching.

#### MEDICATION USE AND COVID-19 SEVERITY IN PATIENTS WITH HYPERTENSION

#### Table 3.

Wellston	Matched Patients with Hypertension			All Manhad Fathers		
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Likelihood of Severe Covid-19, According to Treatment with Various Antihypertensive Agents, in Propensity-Score—Matched Patients with a Positive Test for Covid-19, with Hypertension and Overall.

Among the 2573 patients with hypertension and positive results on Covid-19 testing, 634 (24.6%) had severe Covid-19, as indicated by ICU admission (in 422 patients), mechanical ventilation (in 165), or death (in 343) by April 15, 2020. The likelihood of severe Covid-19 among patients who tested positive, matched on propensity score for previous treatment with each medication class, is shown in Table 3. We ruled out a substantial difference (≥10 percentage points) in the likelihood of severe illness with at least 97.5% certainty for all the medication classes. There was a slightly higher likelihood of severe illness associated with the previous use of calcium-channel blockers (median difference between treated patients and untreated patients, 4.4 percentage points; 95% credible interval, 0.5 to 8.2). The results of unmatched analyses are provided in Table S3.

#### ANALYSES IN ALL PROPENSITY-SCORE-MATCHED PATIENTS

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difference, -3.8 percentage points; 95% credible interval, -7.1 to -0.4). The results of the unmatched analyses are shown in Tables S2 and S3, and the results of propensity-score—matching quality are shown in Tables S10 through S15.

The likelihood of severe Covid-19 among all the patients who tested positive, matched on propensity score for previous treatment with each medication class, is shown in **Table 3**. We ruled out a substantial difference in the likelihood of severe Covid-19 with at least 97.5% certainty for all the medication classes. Sensitivity analyses with the use of frequentist analyses of propensity-score—matched sets and multivariate logistic regression showed unchanged conclusions in every case (Tables S16 through S19).

Discussion

In this observational analysis in a cohort of more than 12,500 patients who were tested for Covid-19 in a large health network in New York City, previous treatment with medications acting on the RAAS was not associated with a higher risk of testing positive for Covid-19. There was also no substantially higher risk (by ≥10 percentage points) of severe Covid-19 associated with any of the medications studied among the patients with a positive test in our cohort. In most cases, the study sample size permitted us, with a high degree of confidence, to rule out that the risk was higher — even by as little as 6 percentage points — among treated patients than among untreated patients. An effect size smaller than 10 percentage points may be of clinical interest, but we elected to report our findings now before the accrual of a larger sample in order to provide actionable information to the medical community, given the urgency and the rapid progression of the pandemic. Our findings may help to allay concerns on the part of patients and providers regarding the continued use of these agents in patients undergoing testing or receiving treatment for Covid-19.

Our data suggested a modestly lower likelihood of a positive test for Covid-19 among patients taking beta-blockers that was of marginal significance in an analysis that included all matched patients. This finding could be attributable to effects of beta-blockers on the expression or presentation at the cell surface of ACE2, the viral receptor for SARS-CoV-2, or residual confounding in the observational study design. To be relevant in SARS-CoV-2 infection, the effect on ACE2 would need to be present on respiratory epithelium, but relevant studies in animals or humans are limited, so this proposed mechanism remains speculative. Furthermore, we did not observe a similar effect with ACI ARBs, which would be expected to have similar effects on ACE2. Because beta-blockers, AC and ARBs act at different points in the RAAS, their effects on the risk of Covid-19 or the sev

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medications and Covid-19. In a multisite database study that included 78 patients with hypertension who were 65 years of age or older in China, there was no association between previous treatment with any cardiovascular medication class and severe Covid-19, but the risk of severe Covid-19 was lower in a subgroup of 46 patients who had received previous treatment with an ARB than among patients who had not received any of the five drug classes assessed in our study. A single-center study from China showed no significant difference in the likelihood of severe illness related to treatment with ACE inhibitors or ARBs in 42 patients. 20

A limitation of our study is that variations in the diagnostic characteristics for the Covid-19 testing methods that were used in our cohort may have led to misclassification of Covid-19 status. The true sensitivity of testing remains unknown. Some patients in the cohort underwent multiple tests, which increased the likelihood of identifying disease. At this time, many persons with possible Covid-19 in New York City and elsewhere are not tested, particularly those who are not hospitalized. Thus, our study may include an overestimation of the proportion of cases of severe Covid-19. That said, it is unlikely that reduced access to testing would add systematic bias in assessment of the relation between use of antihypertensive medications and Covid-19.

There are several other caveats to consider in interpretation of our analysis. Ascertainment of medication use from the electronic health record may not reflect actual drug exposure. We did not account for socioeconomic status, insurance, or health care access in propensity scores. However, the medication classes that we studied are available as generic drugs, including within the New York City public hospital system, which serves a largely indigent population. Despite propensity-score matching, there may be additional unmeasured confounders affecting our results. Propensity-score matching aims to create overall balance between comparison groups. Our confidence in the evidence from the primary Bayesian analysis is also supported by the sensitivity analyses that used frequentist methods. Some patients in the NYU Langone Health system may have been tested at other health care systems, and we did not have access to those test results.

We identified no adverse effects of use of medications that act on RAAS pathways on the likelihood of a positive test for Covid-19 among patients who were tested or on the likelihood of severe Covid-19 among patients with a positive test.

# **Funding and Disclosures**

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Supplementary Material		~
Supplementary Appendix	PDF	351KB
Disclosure Forms	PDF	319KB

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