Antihypertensive drugs in COVID-19 infection

Hypertension predisposes a poor outcome of COVID-19.¹ It has been speculated that specific antihypertensive drugs may underlie this association.² In particular, angiotensin-converting enzyme inhibitors (ACE-Is) and angiotensin II receptor blockers (ARBs) may up-regulate ACE2—the receptor used by SARS-CoV-2 to enter host cells—which has been suggested to lead to an increased risk of infection,² however, by limiting the effects of angiotensin II, ACE2 could also protect against a more severe COVID-19 infection.³

We studied patients diagnosed with COVID-19 (by PCR, CT scan, or both), hospitalized between 1 March and 8 April 2020 in six Munich hospitals and 4 medical centres across the Netherlands (characteristics available upon request).

A mild course of COVID-19 was defined as documented discharge from the hospital within 10 days without treatment at an intensive care unit (ICU). A severe course was defined as any treatment at an ICU, or death. Patients still hospitalized on 8 April 2020 on a regular ward and missing data of ICU admission or death or number of days hospitalized were excluded from the analysis.

Requirement for informed consent was waived by the Medical Ethics Commission of the Amsterdam UMC on 20 March 2020 (registration number 2020-156) and by the ethics committee of the University Hospital rechts der Isar, Technische Hochschule München on 4 April 2020 (registration number 183/20S).

Data on the course of the disease and medication use were extracted from electronic medical records.

We used least absolute shrinkage and selection operator (lasso) logistic regression on the German patient data to identify comorbidities and patient characteristics predictive for the clinical outcome, including the following regressors: obesity, hypertension, diabetes mellitus, coronary heart disease, congestive heart failure, chronic obstructive pulmonary disease, asthma, chronic kidney disease, cancer, human immunodeficiency virus infection, chronic liver disease, age, and sex. Five-fold cross-validation was then used to find a suitable value for the regularization parameter, using the one-standard deviation rule. Multivariate logistic regression was used to analyse the effect of each individual antihypertensive medication on disease severity. Covariates were chosen by best subset selection, comparing all 16 models that include the five drugs as predictors and combinations of the variables age, sex, hypertension, and diabetes. The best-performing model both in terms of Akaike and Bayesian information criteria included only the variables age, sex, and diabetes in addition to the five antihypertensive medications [Figure 1; additional parameters and standard errors: age 0.028 (0.0057); sex, male 0.66 (0.16); diabetes 0.58 (0.19); intercept: -2.5 (0.4)]. We used SPSS version 26 and MATLAB version 2018a.

We checked the resulting model for collinearity based on condition number and variance decomposition analysis.⁴ The largest condition index was 15.0, indicating mild to moderate collinearity. The variance decomposition analysis further showed that the detected collinearity is between age and the intercept variable (which both still have point estimates significantly different from zero); no further collinearity was detected.

We included n = 1134 COVID-19 patients, admitted to a hospital in Munich (n = 311) or the Netherlands (n = 823). We excluded 254 patients from the analysis because of an undefined outcome. The remaining 880 patients were grouped into severe outcome (n = 415)and mild outcome (n = 465). A total of 414 (47%) were treated before hospital admittance with at least one antihypertensive agent. In the lasso logistic regression analysis, hypertension was found to have the greatest importance among a number of relevant comorbidities in the German cohort. Following suitable regularization, additional variables with non-zero coefficients were: diabetes, age, sex, obesity, chronic obstructive pulmonary disease, asthma, and chronic kidney disease. The observed relationship between hypertension and severity of disease may not only be caused by hypertension per se, but could be caused by the underlying antihypertensive drugs used. Multivariate analysis showed a nominally significant association with poor outcome in patients treated

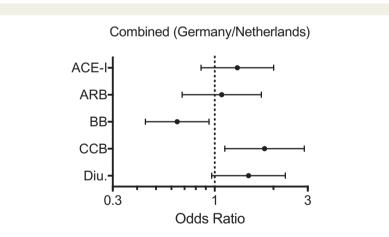


Figure I Association between antihypertensive drugs used and outcomes in the two cohorts combined, corrected for age, sex, and diabetes. An odds ratio >1 indicates a higher chance of treatment in an intensive care unit or death (error bars indicate 95% confidence limits; ACE-I, ACE inhibitor; ARB, angiotensin receptor blocker; BB, beta-blocker; CCB, calcium channel blocker; Diu, diuretic).

© The Author(s) 2020. Published by Oxford University Press on behalf of the European Society of Cardiology.

This is an Open Access article distributed under the terms of the Creative Commons Attribution Non-Commercial License (http://creativecommons.org/ licenses/by-nc/4.0/), which permits non-commercial re-use, distribution, and reproduction in any medium, provided the original work is properly cited. For commercial re-use, please contact journals.permissions@oup.com In conclusion, we find no evidence for adverse outcomes in severely affected COVID-19 patients that used ARBs prior to admission. The use of beta-blockers was associated with a significantly better outcome, whereas the use of calcium channel blockers was associated with poorer outcomes.

Interestingly, recent comparable studies in patients admitted for COVID-19 likewise observed a neutral role for prior use of ARBs^{5,6} and a beneficial role for beta-blockers⁵ as we describe here. However, given the known caveats of these observational studies, there is an urgent need for prospective studies on this subject.

Acknowledgements

We thank Marcel Aries, Pieter Kubben, Mark Janssen, Koos Zwinderman (AMC), and Marije Hoos (MUMC) for their help with this study, and Professor Dr Martin Paul, University of Maastricht, for his conceptual contribution.

Funding

This study was supported by the Netherlands Heart Foundation CVON-PRIME (to Y.P.), ERN GUARD-Heart (to H.B.), the AMC foundation by means of an AMC PhD Fellowship (to J.O.), the German Federal Ministry of Education and Research (BMBF) within the framework of FRA-NFT on Cardiovascular Disease (Druggable-MI-genes: 01KL1802), within the scheme of target validation (BlockCAD: 16GW0198K), the German Centre of Cardiovascular Research (DZHK) Munich Heart Alliance, within the framework of the e:Med research and funding concept (AbCD-Net: 01ZX1706C; Quan-T-cell: 01ZX1505), and as a co-applicant for the British Heart

Foundation (BHF)/German Centre of Cardiovascular Research (DZHK) collaboration. Further support has been received from the German Research Foundation (DFG) as part of the Sonderforschungsbereich SFB 1123 (B02) and the Sonderforschungsbereich SFB TRR 267 (B05). The Bavarian State Ministry of Health and Care, furthermore funded this work within its framework of DigiMed Bayern (grant no, DMB-1805-0001). The work has also been supported by the German Federal Ministry of Economics and Energy in its scheme of ModulMax (grant no. ZF4590201BA8).

Conflict of interest: none declared.

References

- Guan W-J, Ni Z-Y, Hu Y, Liang W-H, Ou C-Q, He J-X, Liu L, Shan H, Lei C-L, Hui DSC, Du B, Li L-J, Zeng G, Yuen K-Y, Chen R-C, Tang C-L, Wang T, Chen P-Y, Xiang J, Li S-Y, Wang J-L, Liang Z-J, Peng Y-X, Wei L, Liu Y, Hu Y-H, Peng P, Wang J-M, Liu J-Y, Chen Z, Li G, Zheng Z-J, Qiu S-Q, Luo J, Ye C-J, Zhu S-Y, Zhong N-S. Clinical Characteristics of Coronavirus Disease 2019 in China. N Engl J Med 2020;**382**:1708–1720.
- Fang L, Karakiulakis G, Roth M. Are patients with hypertension and diabetes mellitus at increased risk for COVID-19 infection? *Lancet Respir Med* 2020;8: e21.
- Imai Y, Kuba K, Rao S, Huan Y, Guo F, Guan B, Yang P, Sarao R, Wada T, Leong-Poi H, Crackower MA, Fukamizu A, Hui C-C, Hein L, Uhlig S, Slutsky AS, Jiang C, Penninger JM. Angiotensin-converting enzyme 2 protects from severe acute lung failure. *Nature*. 2005;**436**:112–116
- Belsley DA, Kuh E, Welsh RE. Regression Diagnostics: Identifying Influential Data and Sources of Collinearity. New York: John Wiley & Sons, Inc.; 1980.
- Reynolds, HR, Adhikari, S, Pulgarin C, Troxel AB, Iturrate E, Johnson SB, Hausvater A, Newman JD, Berger JS, Bangalore S, Katz SD, Glenn GI, Kunichoff D, Chen Y, Ogedegbe G, Hochman JS. Renin–angiotensin–aldosterone system inhibitors and risk of Covid-19. 2020;doi: 10.1056/NEJMoa2008975
- Mehra MR, Desai SS, Kuy S, Henry TD, Patel AN. Cardiovascular disease, drug therapy and mortality in Covid-19. N Eng J Med 2020;doi: 10.1056/NEJMoa2007621.

Sara-Joan Pinto-Sietsma¹[†], Michael Flossdorf^{2†}, Veit R. Buchholz², Joost Offerhaus¹, Hidde Bleijendaal¹, Martijn Beudel¹, Paul G.A. Volders³, Rachel M.A. ter Bekke³, Tom Dormans⁴, Peter-Paul Zwetsloot⁵, Peter de Jager⁵, Steffen Massberg⁶, Patrick Rämer⁷, Clemens Wendtner⁷, Ellen Hoffmann⁸, Kathrin Rothe², Susanne Feihl⁹, Thorsten Kessler¹⁰, Yigal M. Pinto^{1‡}, and Heribert Schunkert^{10‡}

¹University of Amsterdam, Amsterdam, The Netherlands, ²Institute for Medical Microbiology, Immunology and Hygiene, Technische Universität München, Deutsches Zentrum für Infektionsforschung, München, Germany, ³Maastricht University, Maastricht, The Netherlands, ⁴Zuyderland Hospital, Heerlen, The Netherlands, ⁵Ieroen Bosch Hospital, 's-Hertogenbosch, The Netherlands, ⁶Department of Cardiology. Klinikum der Ludwig-Maximilians-Universität and Deutsches Zentrum für Herz-Kreislauf-Forschung, München, Germany, ⁷Department of Hospital Hygiene and infection prevention, Munich Municipal Hospital Group, München, Germany, ⁸Department of Cardiology and Internal Intensive Care Medicine, Heart Center Munich-Bogenhausen, Munich Municipal Hospital Group, Englschalkinger Str. 77, Munich, Germany, ⁹Klinikum Rechts der Isar, Technische Universität München, München, Germany; and , ¹⁰Department of Cardiology, Technische Universität München, Deutsches Zentrum für Herz-Kreislauf-Forschung, Munich Heart Alliance; München, Germany

*Corresponding author. Department of Experimental Cardiology, Amsterdam Universitair Medische Centra, meibergdreef 9, 1105AZ Amsterdam, The Netherlands: Tel: +31 205664688, Fax: +31 205664687, Email: y.pinto@amsterdamumc.nl

[†]These authors contributed equally to this work. [‡]These author should be considered as joint senior authors.