### LETTER TO THE EDITOR

# Possible Pleiotropic Effect of *SRY* Gene May Increase Male Susceptibility to COVID-19

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**GRAPHICAL ABSTRACT** 



To the Editor: We have read with interest the manuscript entitled "SARS-CoV-2 Receptor ACE2 Gene Is

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© American Journal of Hypertension, Ltd 2021. All rights reserved. For Permissions, please email: journals.permissions@oup.com Associated With Hypertension and Severity of COVID 19: Interaction With Sex, Obesity and Smoking," published in January 2021.

The authors<sup>1</sup> make extremely important contributions, especially on genetic predisposition to SARS-CoV-2. They investigated possible associations between genetic polymorphisms of ACE2 and patients with severe coronavirus disease. However, other genetic indicators may indicate a potential susceptibility mechanism for the severity of COVID-19 infection in men. Hormonal contributions can be partly responsible for sex disparities, but additional evidence points to sex chromosome contributions.

Importantly, the *SRY* (sexdetermining region Y) gene in the Y chromosome (Yp11.2) works as a transcription factor for several genes. Studies of Y chromosome in animal models first suggested a *SRY* contribution to elevate blood pressure. *SRY* gene was found to be a positive regulator of the angiotensinogen promoter and a negative regulator of the ACE2 promoter.<sup>2</sup> ACE2 is located on the X chromosome (Xp22.2) and is one of the genes escaping X inactivation.

These findings indicate that *SRY* gene in the animal model modulates the renin-angiotensin-aldosterone system, a critical regulator of blood pressure, and the *ACE2* gene, indicating a sex-dependent activity. Moreover, *SRY* upregulates the renin promoter, while its homolog in the X chromosome, *SOX3* (SRY-related HMG-box 3) downregulates this promoter, thus affecting the renin-angiotensin-aldosterone system pathway.<sup>3</sup>

Interestingly, the results highlighted by Hamet *et al.*<sup>1</sup> indicate an association with ACE2 risk alleles and hypertension, which predispose males to severe COVID-19. Although further studies in humans can delineate whether such activity occurs in other species, several pleiotropic actions of the human *SRY* gene have been reported in the literature in the last decade.

Recent studies described that smoking is a risk factor for cardiovascular disease and COVID-19 severity. Besides being implicated in sexual dimorphism, SRY gene is also involved in gonadal differentiation and male hormone production. Differences between sex hormones affect nicotine metabolism. some studies have shown that women metabolize nicotine faster than men. It may elevate ACE2 and transmembrane serine protease 2 (TMPRSS2) expression,<sup>4</sup> indicating gene-environment interactions, as described by Hamet et al.1 Intriguingly, hormonal changes in adult men may result from hypogonadism, which contributes to various metabolic dysfunctions and can induce overweight.

ACE2 is widely expressed in multiple organs including lungs, heart, kidney, and also the adipose tissue. Obese men, especially those with exceeding visceral adipose tissue, could host higher viral load, contributing to development of more severe COVID-19. Early studies in male and female mice revealed sex differences in gene expression in adipose tissue. By recognizing genetic variabilities, in the future, we can better understand these differences in adipose pathophysiology. However, more studies are necessary to investigate the relationship between obesity and *SRY* gene.

In this context, research applying 3-dimensional structures or gene sequence analysis may contribute to indicate whether *SRY* gene exhibit variations in inhibition, modulating the renin–angiotensin–aldosterone system and whether gene expression imbalance predisposes SARS-CoV-2 interaction in males. Then, the study of sex differences may help to find appropriate therapies for all patients.

#### DISCLOSURE

The authors declared no conflict of interest.

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