

COVID-19: A Worldwide Gene Polymorphism

COVID-19 created a very harmful and disastrous experience for humans. This experience was associated with very heavy financial and emotional costs (1). Many people lost their lives during this dangerous and deadly pandemic; many suffered short-term or long-term sequelae and many suffered mental injuries as a result of the injuries caused by the disease, which were sometimes irreversible (2, 3). But that was not the whole story. Medical scientists and healthcare workers, both in the clinical field and in the field of basic medical sciences, worked hard to discover the mystery of this disease. Several areas were investigated jointly between basic and clinical sciences and extensive research was conducted on it (4, 5).

One of these areas was the study of gene polymorphism. An attempt was made to investigate the relationship between gene polymorphism and the chance of infection, the possibility of developing dangerous symptoms, the rate of response to treatment, and the rate of complications or the chance of death. One of the genes whose polymorphism has been widely investigated in COVID-19 is the gene is the ACE1 and ACE 2 gene polymorphism (6-8). Several studies have been performed all over the world with significant results demonstrating the significant effects of ACE1 polymorphism on the fate of COVID-19 patients both inside and outside Iran (7-10).

In this issue of the JCMA, the study by Karimi, et al. has demonstrated the role of this gene polymorphism in one of the ethnic groups in Iran (11); a finding in concordance with similar studies (7-9). It seems that gene polymorphism could have a significant effect not only on the geographical-related prevalence of COVID-19 but also on the severity of the disease and the clinical fate of the patients (12, 13). However, the following loci have been proposed as the most probable COVID-19-related ones: ApoE, ACE1, TMPRSS2, CCR5, and HLA loci (14); while ACE1

D/I and C3 polymorphisms have demonstrated the greatest correlation with COVID-19 prevalence/mortality (12-14) which could be supporting the need for studies in Iran as one of the countries most hit by COVID-19, including the study by Karimi, et al. (11). This is another proof for the paradigm shift in medicine (15).

References

1. Dabbagh A. Cellular and Molecular Approaches to COVID-19: "Road to Perdition" or "The Shawshank Redemption". *J Cell Mol Anesth.* 2020;5(4):214.
2. Pourhossein B, Dabbagh A, Fazeli M. Insights into the SARS-CoV2 Outbreak; the Great Global Challenge: A Mini Review. *J Cell Mol Anesth.* 2020;5(1):23-6.
3. Dabbagh A. COVID-19: Apocalypse Now? *J Cell Mol Anesth.* 2020;5(1):1-2.
4. Dorgalaleh A, Dabbagh A, Tabibian S, Baghaeipour MR, Jazebi M, Bahraini M, et al. Patients with Congenital Bleeding Disorders Appear to be Less Severely Affected by SARS-CoV-2: Is Inherited Hypocoagulability Overcoming Acquired Hypercoagulability of Coronavirus Disease 2019 (COVID-19)? *Semin Thromb Hemost.* 2020;46(7):853-5.
5. Safari S, Salimi A, Zali A, Jahangirifard A, Bastanagh E, Aminnejad R, et al. Extracorporeal Hemoperfusion as a Potential Therapeutic Option for Severe COVID-19 patients; a Narrative Review. *Arch Acad Emerg Med.* 2020;8(1):e67.
6. Rajaei S, Dabbagh A. The immunologic basis of COVID-19: a clinical approach. *J Cell Mol Anesth.* 2020;5(1):37-42.
7. Gupta P, Mohapatra E, Patel S, Patnayak LL, Nanda R, Shah S, et al. Effect of the Angiotensin-Converting Enzyme (ACE) (I/D) Polymorphism in COVID-19 Patients and Their Healthy Contacts. *Cureus.* 2023;15(5):e38610.
8. Mir MM, Mir R, Alghamdi MAA, Alsayed BA, Wani JI, Alharthi MH, et al. Strong Association of Angiotensin Converting Enzyme-2 Gene Insertion/Deletion Polymorphism with Susceptibility to SARS-CoV-2, Hypertension, Coronary Artery Disease and COVID-19 Disease Mortality. *J Pers Med.* 2021;11(11).
9. Sarangarajan R, Winn R, Kiebish MA, Bountra C, Granger E, Narain NR. Ethnic Prevalence of Angiotensin-Converting Enzyme Deletion (D) Polymorphism and COVID-19 Risk: Rationale for Use of Angiotensin-Converting Enzyme Inhibitors/Angiotensin Receptor Blockers. *J Racial Ethn Health Disparities.* 2021;8(4):973-80.
10. Aziz MA, Islam MS. Association of ACE1 I/D rs1799752 and ACE2 rs2285666 polymorphisms with the infection and severity of COVID-19: A meta-analysis. *Mol Genet Genomic Med.* 2022;10(11):e2063.
11. Karimi FH, V. Ahmadi, A. Ranjbar Kermani, F. Ferdowsi, S. The

Evaluation of ACE1-I/D Polymorphism in Kurdish Patients With Severe COVID-19. *J Cell Mol Anesth.* 2023;8(3):141-9.

12. Delanghe JR, De Buyzere ML, Speeckaert MM. Genetic Polymorphisms in the Host and COVID-19 Infection. *Adv Exp Med Biol.* 2021;1318:109-18.

13. Verma S, Abbas M, Verma S, Khan FH, Raza ST, Siddiqi Z, et al. Impact of I/D polymorphism of angiotensin-converting enzyme 1 (ACE1) gene on the severity of COVID-19 patients. *Infect Genet Evol.* 2021;91:104801.

14. Dieter C, Brondani LA, Leitão CB, Gerchman F, Lemos NE, Crispim D. Genetic polymorphisms associated with susceptibility to COVID-19 disease and severity: A systematic review and meta-analysis. *PLoS One.* 2022;17(7):e0270627.

15. Sezari P, Dabbagh A. Personalized medicine: the paradigm shift in medicine mandating lifelong learning. *J Cell Mol Anesth.* 2019;4(2):31-2.

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