Individual’s DNA Repair Capacity and COVID-19: Let’s Take One Step Back to understand it

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Editorial
The outbreak of novel coronavirus from Wuhan, Hubei Province, China has brought turbulence in the entire world. On January 9, 2020; World Health Organization reported that Chinese authorities had identified a novel strain of coronavirus. Each and every country and its authorities are taking major steps to avoid the spread of this pandemic situation. Coronavirus pandemic has disrupted the economy of many countries by hampering business, airline industry, stock market, tourism, and several more. The spread of the virus among humans has demonstrated a wide range of differences in susceptibility. Many individuals get infected by virus or remain asymptomatic or are carriers of it. Immunity, self-hygiene, and civic sense play a pivotal role to avoid getting infected in this scenario of existing overburden in the medical field.

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However, there is a wide variation observed in an individual’s susceptibility towards contracting the COVID-19 infection and their treatment outcomes. Viruses are known to play with the host’s DNA damage responses by either over activating or inhibiting it [1]. There is a need to understand the changes of biological mechanisms undergone on the viral infection and its impact on the host’s DNA damage and repair responses (Figure 1).

![Figure 1: Inter individual variation in DNA repair capacity towards COVID-19 infection outcomes.](image)

Every single minute, humans are exposed to diverse forms of endogenous and exogenous DNA damaging agents which threatens to disrupt the genomic integrity of the cells. Various forms of DNA lesions (single-strand breaks, double-strand breaks, oxidative base damages, abasic sites, mismatch bas-

Understanding the role of DNA damage and repair mechanisms in virus infection and collaborating with the genetic parameters of the individual can shed some light at the end of the dark tunnel. Changes in DNA damage and repair capacity of individuals on getting infected and after treatment can assist to establish the relationship of the underlying potential biological mechanisms. A deeper understanding of the biological basis with genetic factors and environmental exposures can assist to design personalized treatment regime.

References
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