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Hypotheses and facts for genetic factors related to severe COVID-19

Kotsev SV *et al.* Genetics and severe COVID-19

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Abstract

Genomewide association analysis allows the identification of potential candidate genes
involved in the development of severe coronavirus disease-2019 (COVID-19). Hence, it
seems that genetics matters here, as well. Nevertheless, the virus's nature, including its
RNA structure, determines the rate of mutations leading to new viral strains with all
epidemiological and clinical consequences. Given these observations, we herein

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New findings by scientists at the National Institutes of Health and their collaborators help explain why some people with COVID-19 develop severe disease. The findings also may provide the first molecular explanation for why more men than women die from COVID-19. The researchers found that more than 10% of people who develop severe COVID-19 have misguided antibodies—autoantibodies—that attack the immune system rather than the virus that causes the disease. Another 3.5% or more of people who ...			
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Jul 03, 2020 · These risk **factors**, however, do not fully explain why some have no or mild symptoms while others become seriously ill. Thus, **genetic risk factors** are being investigated. An early study (Ellinghaus et al. 2020) identified two genomic regions associated with **severe** COVID-19: one region on chromosome 3 containing six genes and one region

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New insights into genetic susceptibility of COVID-19: an ...

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Jul 15, 2020 · We further discussed that polymorphisms in **ACE2** or **TMPRSS2** could guide effective treatments (i.e., hydroxychloroquine and camostat) for COVID-19. This study suggested that ACE2 or TMPRSS2 DNA polymorphisms were likely associated with genetic susceptibility of COVID-19, which calls for a human genetics initiative for fighting the COVID-19 pandemic.

Major Genetic Risk Factor for Severe COVID-19 Is Inherited ...

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Oct 05, 2020 · By Karolinska Institutet October 5, 2020. A study published in Nature shows that a segment of DNA that causes their carriers to have an up to three times higher risk of developing **severe COVID-19** is inherited from Neanderthals. The study was conducted by researchers at Karolinska Institutet and Max Planck Institute for Evolutionary Anthropology. COVID-19 affects some people ...

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The 2019 novel coronavirus, 2019-nCoV, officially named severe acute respiratory syndrome coronavirus 2, SARS-CoV-2, is a positive-sense, single-stranded RNA coronavirus. The virus is the causative agent of coronavirus disease 2019, COVID-19, and is contagious through human-to-human transmission.

The virus is thought to have a zoonotic origin, and comparisons of the genetic sequences of this virus and other virus samples have shown similarities to SARS-CoV and bat coronaviruses which makes an ultimate origin in bats likely, although an intermediate reservoir such as a pangolin is thought to be involved.

Wikipedia

Help & information

US updates: Centers for Disease Control and Prevention (CDC)

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Author: Salmo Raskin **Publish Year:** 2020

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Cited by: 1 **Author:** Ingrid Fricke-Galindo, Ramcés Falfán-Valen...

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The major genetic risk factor for severe COVID-19 does not ...

<https://www.nature.com/articles/s41598-021-91711-4>

Jun 11, 2021 · The COVID-19 Host **Genetics** Initiative a global initiative to elucidate the role of host **genetic factors** in susceptibility and severity of the SARS-CoV-2 virus pandemic. Eur. J.

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