Comorbidity and Genetic Factors and their Impacts on Patients with COVID-19



Team Members: Rohan Chatterjee, Francisco Contreras, Jimuel Julaton, Luis Gonzalez, Riese Atianzar, Emily Gonzalez, Chen-Ching Lin, vodafone Juan Hernandez, Carlos Hernandez, Ting Fung Ha Faculty Advisor: Dr. Navid Amini **Project Liaison:** Haley Kirk **Department of Computer Science**

College of Engineering, Computer Science, and Technology California State University, Los Angeles



Background

COVID-19, a highly infectious disease caused by the SARS-CoV-2 virus, has had a profound impact on the world, affecting health, economies, and daily life. Despite extensive research, there are still several overlooked aspects related to the disease that have been considered controversial and therefore, have not been thoroughly studied. It is important to consider these areas to gain a complete understanding of the virus and its impact on human health.



The objective of this project is to study overlooked or unconsidered aspects related to COVID-19 to gain a better understanding of the disease. This is being done by studying various areas, including clinical factors such as blood type and its association with disease severity and mortality, as well as the impact of geographical, economical, and social factors on COVID-19 cases, vaccinations, and mortality. Our approach involves segmenting the deliverables into three separate branches.



Internet Access vs Covid Cases

Visualization Application



Blood Type Analysis	
learn	 pandas
Comorbidities and Traits Investigated	
 Blood Group Fever Chills Cough Dyspnea Anosmia Ageusia Loss of Appetite Asthenia 	 Cyanosis Rhinorrhea Sore Throat Diarrhea Muscle Ache Nausea Vomiting Headache Gender.
After preprocessing the d was used to ident	ata the Boruta Algorithm ify core features.
Boruta Core Selected Features for Mortality (All Blood Types)	
Chills	Features were



A comparison using percentage of internet access of different countries and COVID-19 cases in those countries, with the countries having more access also having more cases.





Note:

You can use the following QR codes to access the project's online versions.



Features:

- **Realtime data**
- Easy interaction
- Intuitive Design
- Custom

Dashboard

stability in Fisher's Diarrhea Exact and Chi Squared tests. Cough Sore Throat Next multiple models were created for each, each varying a single feature informing if the sample matches the corresponding blood type. Random Forest All Features Random Forest Core Features - 0.714 - A+ — 0.777 - A-— 0.714 - AB+ - 0.737 - AB-- 0.771 - AB-— 0.695 - B+ — 0.766 - B+ ----- 0.725 - B----- 0.8 - B-0.71 - 0+ ---- 0.754 - O+ ---- 0.726 - 0----- 0.797 - 0-0.4 0.6 False Positive Rate 0.0 0.2 0.6 0.8 10 0.0 0.2 0.4 0.6 0.8 1.0 False Positive Rate **Results:** Identified Core Comorbidities associated with COVID-19 Core Comorbidities helped reduce noise in the models

Cyanosis

further supported

by high rank order

- No significant changes were found when examining blood type
- Blood type likely has no effect on COVID-19 mortality

similar were found for results severity