

Review of: "A Multi-factor Model of COVID-19 Epidemic in California"

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Potential competing interests: No potential competing interests to declare.

The study introduces a multi-factor model to assess COVID-19 transmission in California counties, correlating factors like population, income, and density with cumulative cases. However, the model's limitations include oversimplification, omitted variables, and limited predictive ability, emphasizing the intricate nature of pandemic dynamics and the need for comprehensive analysis. The study has a good contribution and well written. I have some points to mention:

- a) The model relies on a limited set of factors such as population, income, density, Gini coefficient, and size. There might be other significant factors that were not considered, potentially leading to an incomplete understanding of the COVID-19 spread.
- b) The study emphasizes correlation between factors and outcomes, but it's important to note that correlation does not imply causation. The identified correlations might not necessarily represent direct causal relationships.
- c) The model assumes a linear relationship between factors and outcomes. In reality, the relationships might be more complex and nonlinear, leading to potential inaccuracies in predictions.
- d) While the selected factors are relevant, other important variables might have been omitted from the analysis. For instance, factors related to healthcare infrastructure, travel patterns, public health measures, and cultural practices could play a role in the spread of the virus.
- e) The study considers a fixed time frame from March 2020 to June 2023. The dynamics of the pandemic, including changes in virus variants, vaccination campaigns, and evolving public health responses, are not fully accounted for.
- f) The study acknowledges that the model lacks strong predictive power, especially in forecasting cumulative cases and duration early in the epidemic. This limits its utility for future pandemic planning and response.
- g) The study treats infection rate as a random variable with a normal distribution. In reality, infection rates can be influenced by various complex factors, such as human behavior, healthcare capacity, and public health interventions.
- h) The study briefly mentions the possibility of extending the model using convergence rates of successive approximations, but this is left as future work. The feasibility and accuracy of such extrapolation methods are not explored in depth.
- i) The study emphasizes correlation coefficients and R-squared values as measures of model performance. However,

these metrics might not fully capture the validity and reliability of the model in predicting real-world outcomes.