



# Version 2 ▼

### Mar 23, 2020

## Suitability Map of COVID-19 Virus Spread V.2

### Gianpaolo Coro<sup>1</sup>

<sup>1</sup>CNR

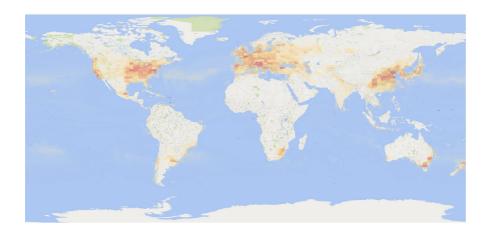
1 Works for me

dx.doi.org/10.17504/protocols.io.bd25i8g6

Coronavirus Method Development Community

Gianpaolo Coro CNR

ABSTRACT



This image reports a Maximum Entropy model that estimates *suitable* locations for COVID-19 spread, i.e. places that could favour the spread of the virus just in terms of environmental parameters.

The model was trained just on locations in *Italy* that have reported a rate of new infections higher than the geometric mean of all Italian infection rates. The following environmental parameters were used, which are correlated to those used by other studies:

- Average Annual Surface Air Temperature in 2018 (NASA)
- Average Annual Precipitation in 2018 (NASA)
- CO2 emission (natural+artificial) averaged between January 1979 and December 2013 (Copernicus Atmosphere Monitoring Service)
- Elevation (NOAA ETOPO2)
- Population per 0.5° cell (NASA Gridded Population of the World)

The model file (in ASC format) and all parameters used are attached.

A higher resolution map and also the model file (in ASC format) and all parameters are available at the external link (Zenodo).

The model indicates highest correlation with *infection rate* for CO2 around 0.03 gCm^-2day^-1, for Temperature around 11.8 °C, and for Precipitation around 0.3 kg m^-2 s^-1, whereas Elevation and Population density are poorly correlated with *infection rate*.

One interesting result is that the model indicates, among others, the Hubei region in China as a high-probability location, and Iran (around Teheran) as a suited location for virus' spread, but the model was not trained on these regions, i.e. it did not know about the actual spread in these regions.

Files

protocols.io

03/23/2020

Citation: Gianpaolo Coro (03/23/2020). Suitability Map of COVID-19 Virus Spread. https://dx.doi.org/10.17504/protocols.io.bd25i8g6

Name (Size)

1\_covid\_suitability\_preview.png (5.7 MB)

md5:dea4e66a1c66d0dfc3b0872adfaa020f

2\_covid\_suitability\_v2\_Hi\_Resolutionv2.png (47.3 MB)

md5:069727a6c5656d276c475606c9b96d47

Altitude.asc (1.9 MB)

md5:ca91c4d56654b77bf572eef1a42af7a5

CO2.asc (5.1 MB)

md5:0ed217e20ab32aad4ab96e5403670ee4

MaxEnt\_Temperature\_Precipitation\_Elevation\_CO2.asc (2.8 MB)

md5:79639fd3540c68450d86fde288edb264

Population.asc (4.6 MB)

md5:57aa6c172b3fc036c08d0560f01436ba

Precipitation.asc (5.5 MB)

md5:3ab587ea0e0fbe3fcbd9ea6b7844271a

Temperature.asc (4.7 MB)

md5:7ea930f59e5ff627a18383f02737f78d



MD5 checksum: This is the file fingerprint, which can be used to verify the file integrity.

### References



Coro, G., Panichi, G., Scarponi, P., & Pagano, P. (2017). Cloud computing in a distributed e-infrastructure using the web processing service standard. Concurrency and Computation: Practice and Experience, 29(18), e4219.

EXTERNAL LINK

https://zenodo.org/record/3719184

THIS PROTOCOL ACCOMPANIES THE FOLLOWING PUBLICATION

Gianpaolo Coro. (2020). Suitability Map of COVID-19 Virus Spread (Version 4) [Data set]. Zenodo. http://doi.org/10.5281/zenodo.3722606

MATERIALS TEXT

This experiment was done using the DataMiner cloud computing system of the D4Science e-Infrastructure and the BiodiversityLab Virtual Reseach Environment. (https://services.d4science.org/group/biodiversitylab/)

Citation: Gianpaolo Coro (03/23/2020). Suitability Map of COVID-19 Virus Spread. https://dx.doi.org/10.17504/protocols.io.bd25i8g6



Gianpaolo Coro. (2020). Suitability Map of COVID-19 Virus Spread

3