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Burkholderia infection

Burkholderia - genus of Proteobacteria whose pathogenic members include:



<https://en.wikipedia.org/wiki/Burkholderia>

- *Burkholderia pseudomallei*, causative agent of melioidosis – human pathogen
- *Burkholderia cepacia*, an important pathogen of pulmonary infections in people with cystic fibrosis (CF)
- *Burkholderia cepacia* complex - human pathogen
- *Burkholderia mallei*, responsible for glanders - equine pathogen; zoonotic transfer to humans



<https://www.bangkokpost.com/thailand/general/823108/alert-sounded-over-deadly-melioidosis>

Infection is mainly acquired cutaneously through abrasions in the skin or inhalation of contaminated particles.

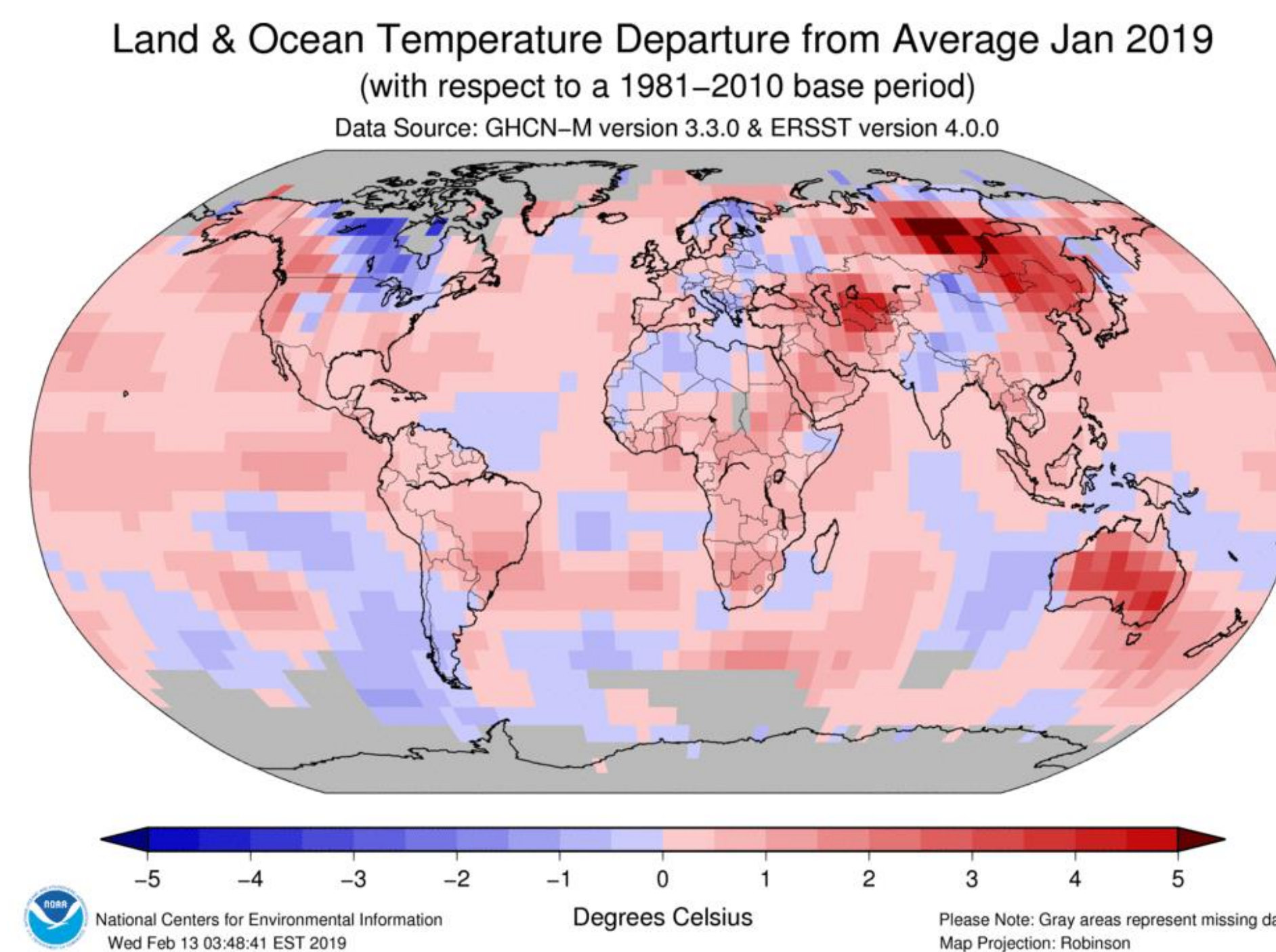
Melioidosis is classically characterized by:

- Pneumonia
- Multiple abscesses
- Septicemia
- Mortality rate of up to 40% or more

Predisposing factors include:

- Diabetes
- Chronic kidney disease
- Immunosuppression

Climate change



<https://www.ncdc.noaa.gov/sotc/service/global/map-blended-mntp/201901.gif>

In addition to increased migration, rising global temperatures and flooding may contribute to creating climates that support better growth of *Burkholderia*.

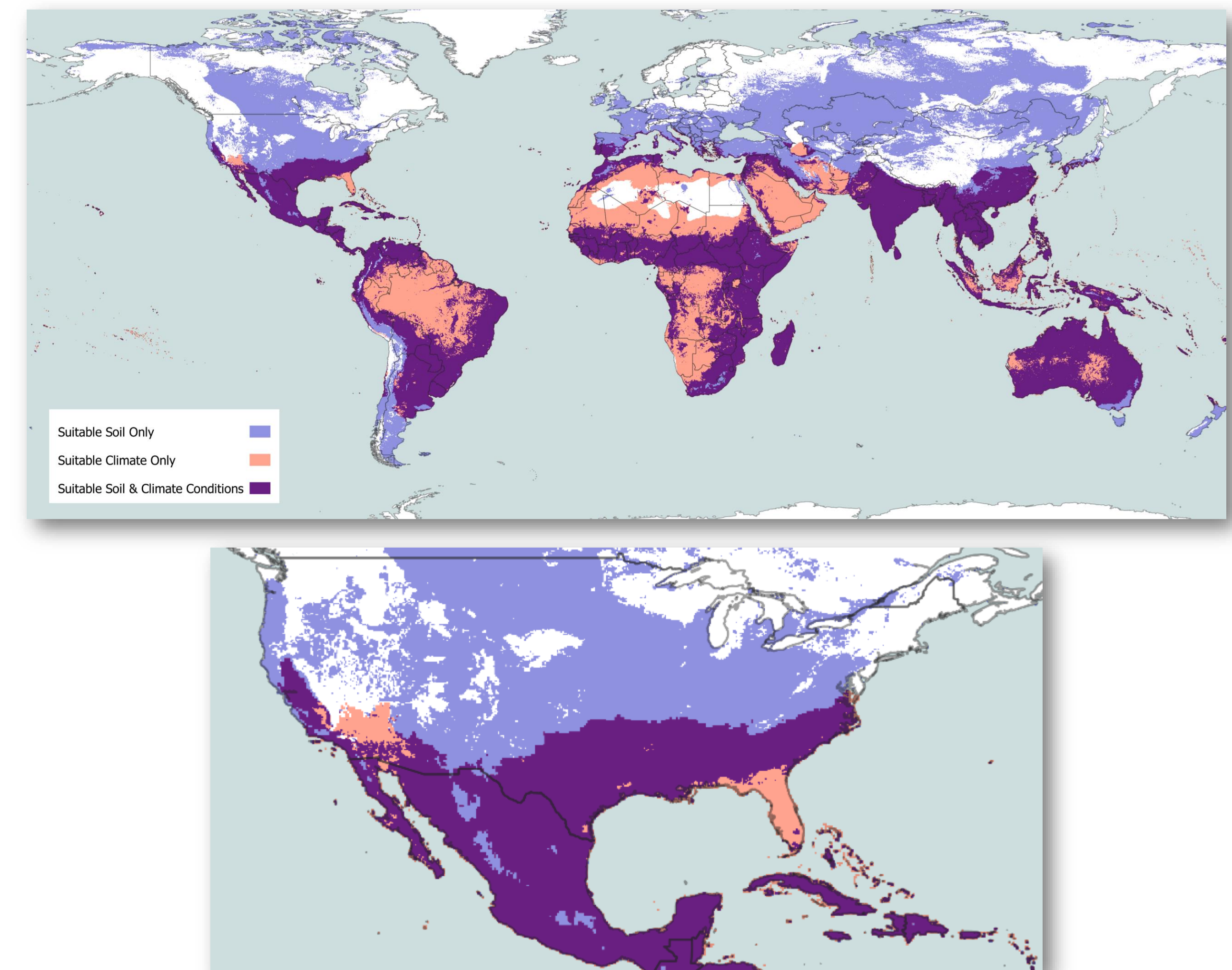
Aims & Methodology

- Explore existing data about soil, water, weather, land use and melioidosis risk factors
- Generate geospatial data and maps to identify ecological niches for *B. pseudomallei* and related soil organisms
- Future Work: Investigate DataX to improve performance and add additional layers to our analysis. DataX enables researchers to access, share, understand, and analyze critical data using the high performance computational resources at the Texas Advanced Computing Center (TACC).

Acknowledgements

We acknowledge support for this project from the Grand Challenges at the University of Texas at Austin Planet Texas 2050 Initiative and the University of Cambridge, UK Global Alliance Fund and Returning Careers Scheme.

Geospatial analysis

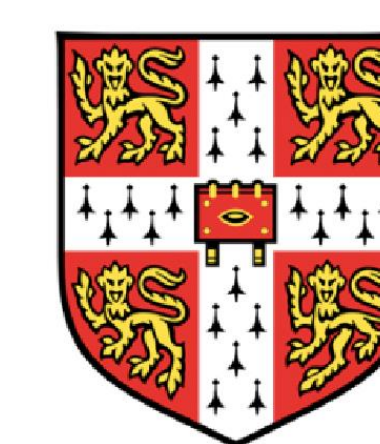


This predicted potential range of *Burkholderia pseudomallei* was based on a dataset^[2] of documented worldwide soil and water occurrences between 1933 and 2013 and global datasets of soil^[6] and climatic^[7] conditions. Suitability for climate factors was based on annual mean temperature, minimum temperature of the coldest month, mean annual precipitation, and mean precipitation of the wettest month while soil suitability was based on cation exchange capacity, percent clay composition, pH, and organic carbon content at depths of 30cm, 60cm, and 100cm for all soil variables.

References

- Limmathurotsakul et al. Nature Microbiology, (2016). doi: 10.1038/nmicrobiol.2015.8.
Limmathurotsakul, D. (2015). Data_Melioidosis_1910_2014. doi: 10.6084/m9.figshare.1447097.v1
TACC: <http://tacc.utexas.edu/>
DataX: <https://ptdatax.tacc.utexas.edu/>
Bridging Barriers: <https://bridgingbarriers.utexas.edu/planet-texas-2050/>
Global soil data: <https://files.isric.org/soilgrids/data/recent/>
Global climate data: <http://worldclim.org/bioclim>
QGIS: <https://qgis.org>
Python: <https://python.org>

Geospatial Analysis to Guide Identification of Environments Suitable of Supporting *B. pseudomallei* in Africa



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Burkholderia may be underreported in West Africa

Although considered a tropical disease, the hub of melioidosis infection was previously thought to be within Southeast Asia and Northern Australia (Selected regions in the area circled red below)



<http://ontheworldmap.com/>

A recent environmental suitability model has however predicted the occurrence of the soil-dwelling *B. pseudomallei* in regions in West Africa where melioidosis may be endemic but underreported (Selected regions in the area circled blue above).



<http://www.usccn.org/Pages/default.aspx>

Of the 470 strains listed in the published *global B. pseudomallei* culture collection, only 8 are reported to be African strains [Nat Microbiol, 2017 2:16263].

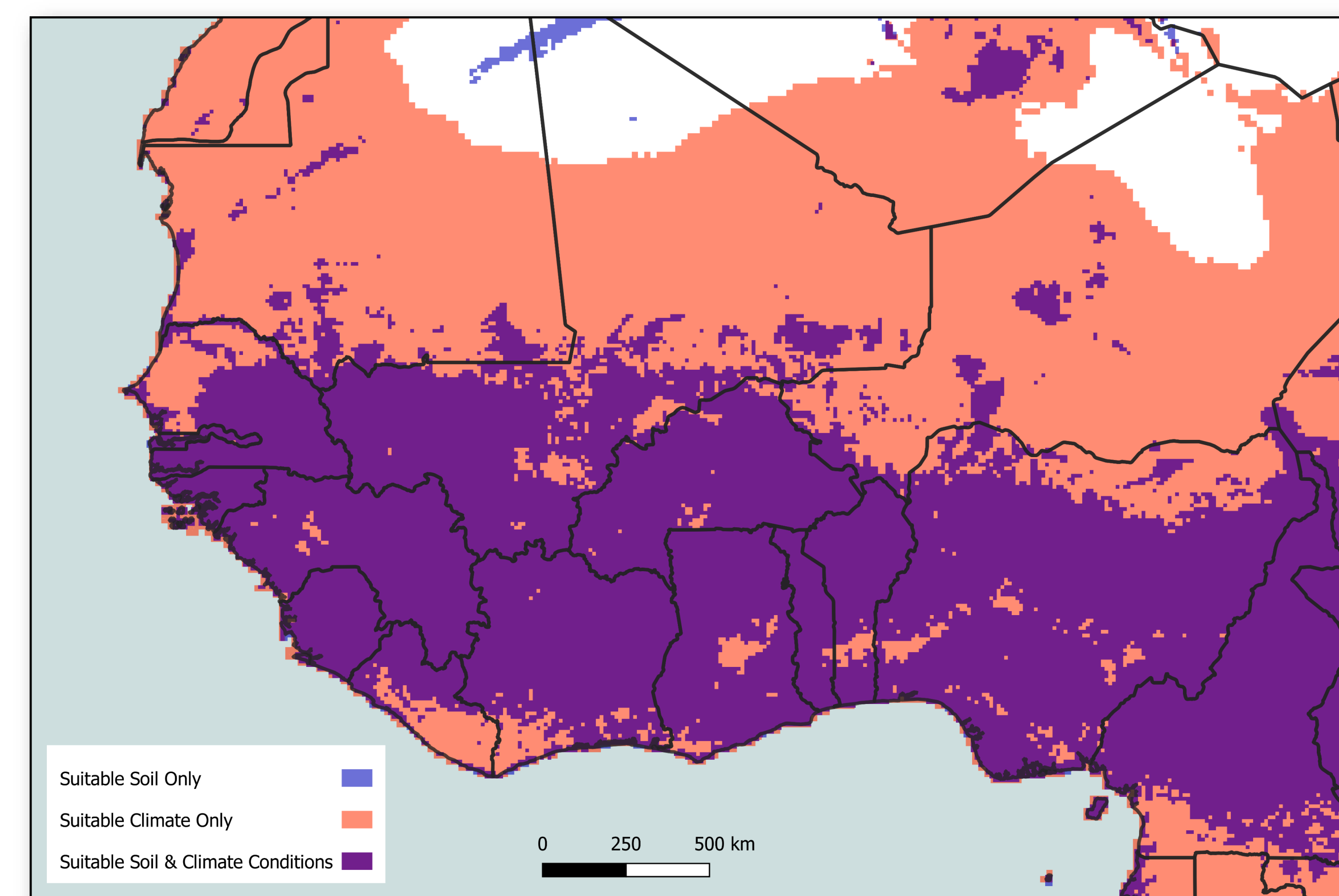
Aims & Methodology

- Explore existing data about soil, water, weather, land use and melioidosis risk factors in West Africa
- Generate geospatial data and maps to improve identification of areas for soil sampling and clinical research of melioidosis in Africa.
- Future Work: Investigate DataX to improve performance and add additional layers to our analysis. DataX enables researchers to access, share, understand, and analyze critical data using the high performance computational resources at the Texas Advanced Computing Center (TACC).

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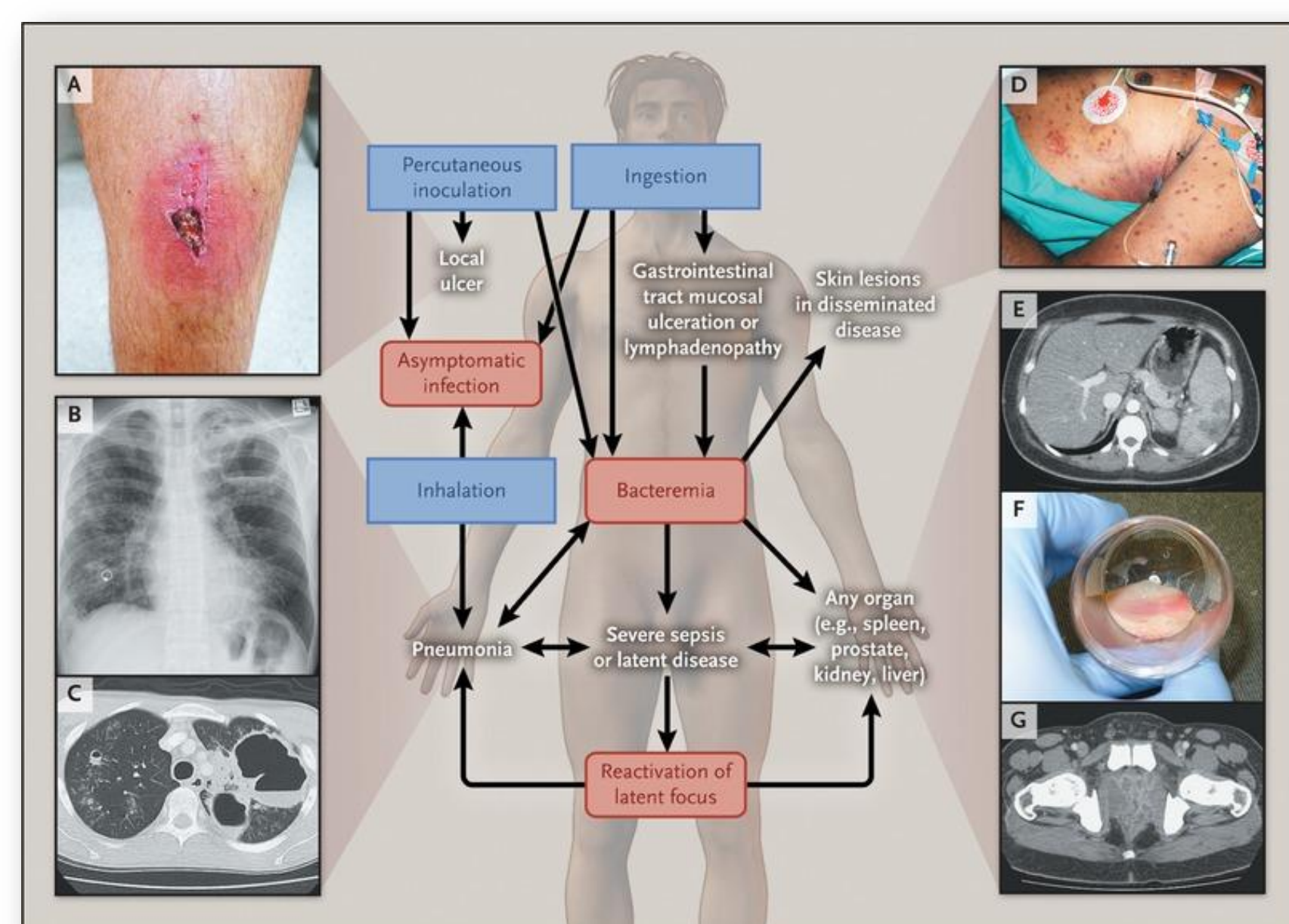
Geospatial analysis



Predicted potential range of *Burkholderia pseudomallei* based on a dataset^[2] of documented worldwide soil and water occurrences between 1933 and 2013 and global datasets of soil^[7] and climatic^[8] conditions. Suitability for climate factors was based on annual mean temperature, minimum temperature of the coldest month, mean annual precipitation, and mean precipitation of the wettest month while soil suitability was based on cation exchange capacity, percent clay composition, pH, and organic carbon content at depths of 30cm, 60cm, and 100cm for all soil variables.

References

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 Limmathurotsakul, D. (2015). Data_Melioidosis_1910_2014. doi: 10.6084/m9.figshare.1447097.v1
 Chewapreecha et al. (2017) Nature Microbiology. doi: 10.1038/nmicrobiol.2016.263.
<http://tacc.utexas.edu/>
<https://ptdatax.tacc.utexas.edu/>
<https://bridgingbarriers.utexas.edu/planet-texas-2050/>
 Global soil data: <https://files.isric.org/soilgrids/data/recent/>
 Global climate data: <http://worldclim.org/bioclim>
 QGIS: <https://qgis.org>
 Python: <https://python.org>



<https://www.nejm.org/doi/full/10.1056/NEJMra1204699>

The low reported cases in West Africa may be due to the diverse clinical manifestations of the disease as well as inadequate availability of identification methods