



## Michelle Evans

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Identifying climate-drivers of three common infectious diseases in Madagascar via participatory modeling



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Models are not interpretable or credible to stakeholders

Global models may not match the local context

Software tools are not maintained

# Traditional modeling process is linear and siloed



Initial discussion between  
researchers and health actors



## Model Formulation

- Identify project goals
- Identify relevant indicators
- Identify potential input variables

## Model Building

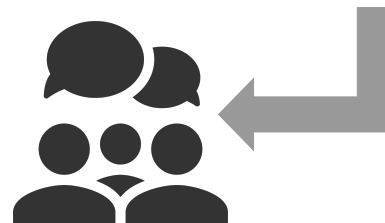
- Collect and process data
- Evaluate data quality
- Construct empirical model

## Model Evaluation

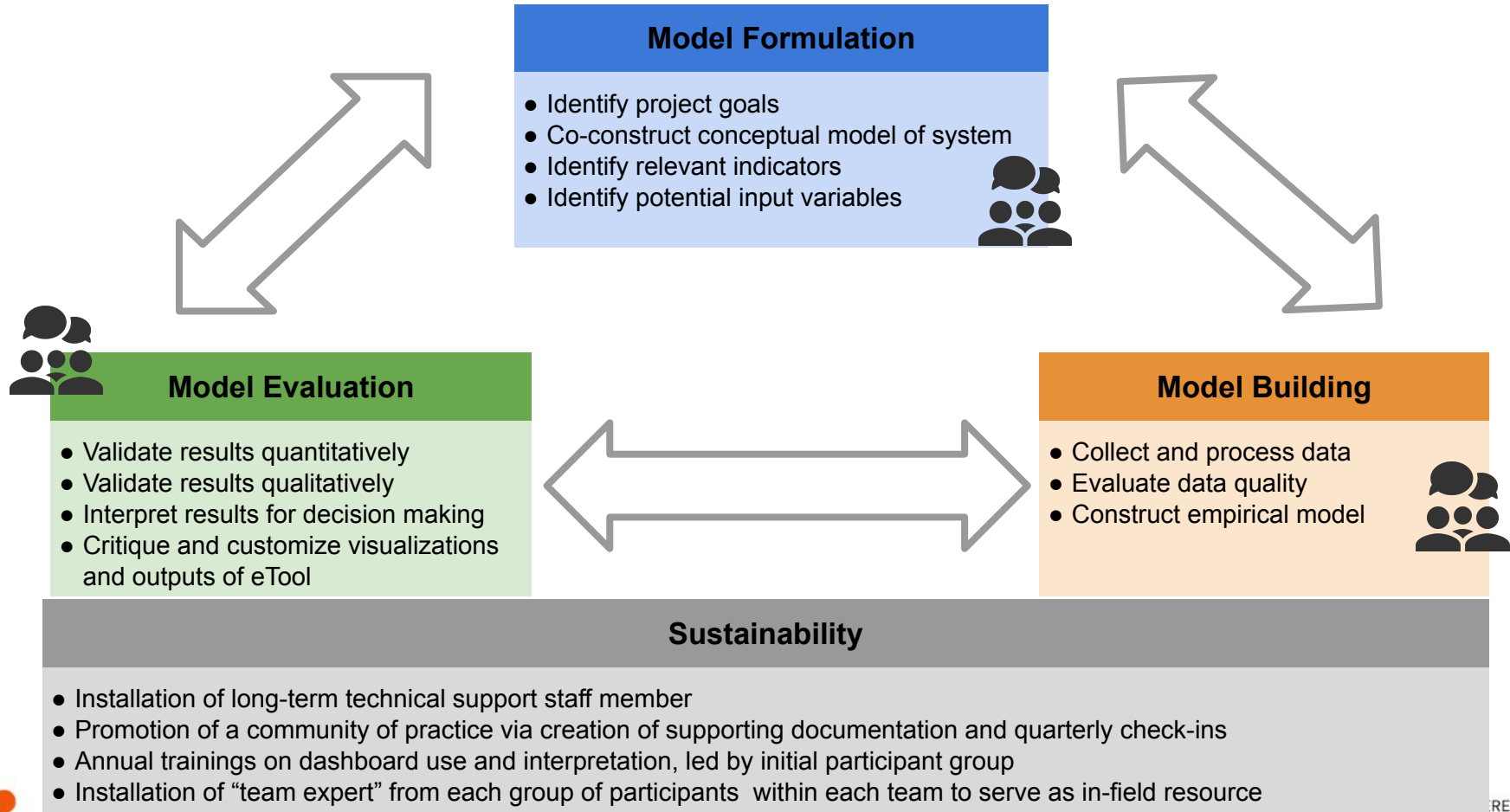
- Validate results quantitatively



Final presentation of results  
to health actors

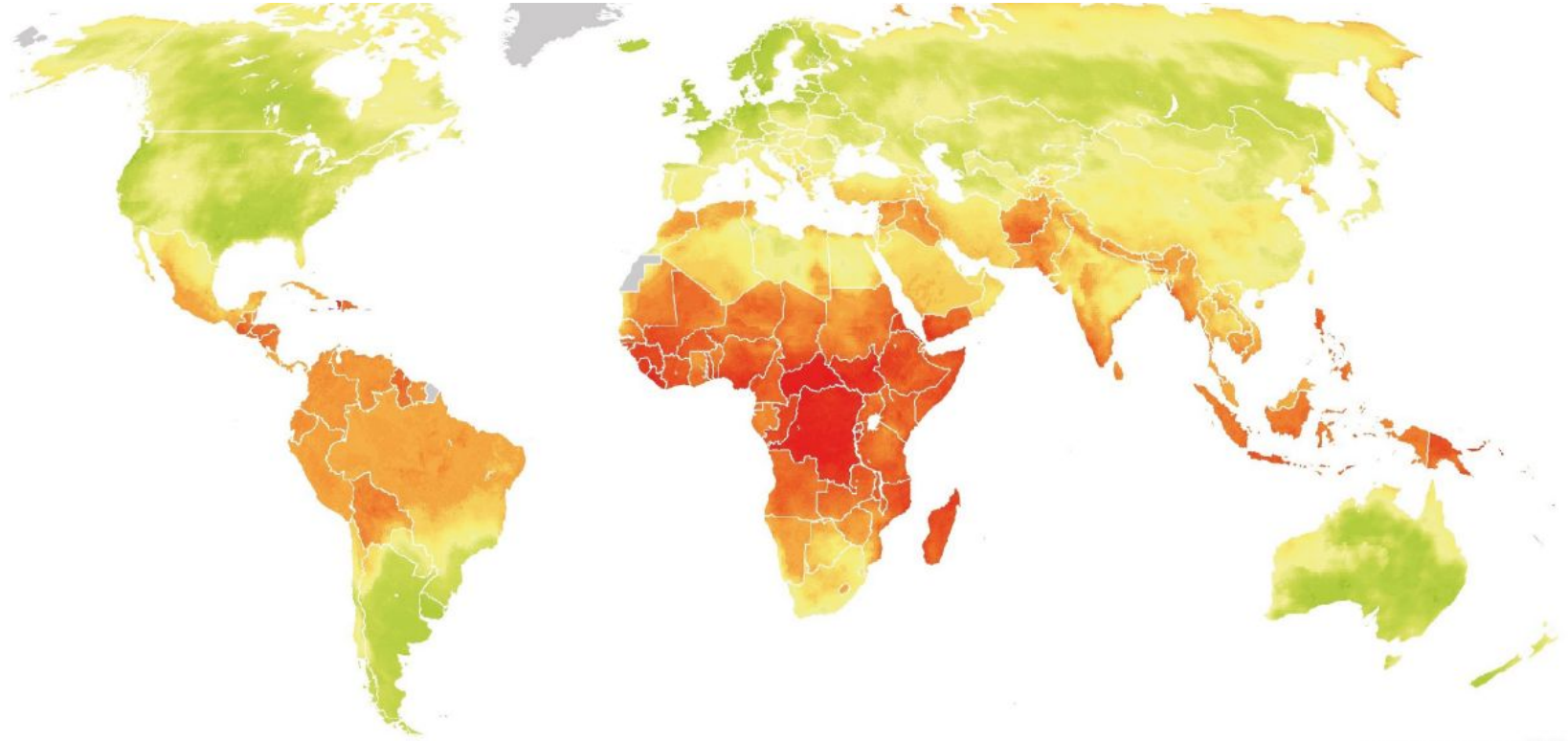


# Participatory modeling is iterative and inclusive



Participatory modeling offers a way to operationalize disease forecasting models into usable software designed with and for decision makers within the health system

# Madagascar is one of the most vulnerable countries to climate change



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Legend

Extreme Risk

High Risk

Medium Risk

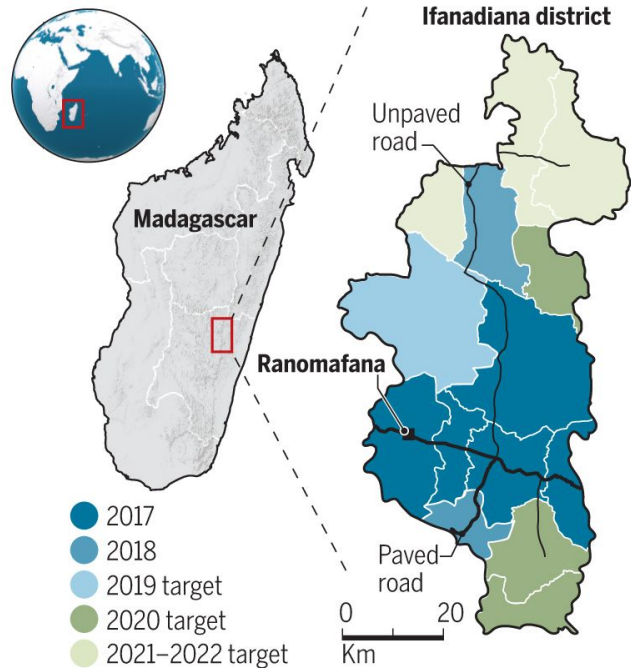
Low Risk

■ No data

VCE 2024



# A health-system strengthening partnership



Began in 2014, in the District of Ifanadiana, Vatovavy.

Now serves over 1 million

Focuses on universal health coverage and community health programs

Existing partnership facilitates a participatory approach



Co-creation of disease forecasting system via participatory modeling to be integrated into a DHIS2 application

Targets last-mile healthcare delivery at the community level (<5km resolution)

Multi-pathogen



Malaria

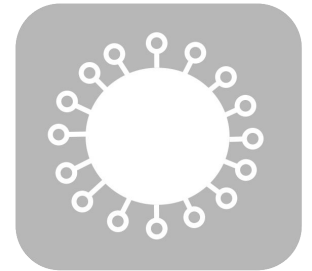


Diarrheal  
Disease



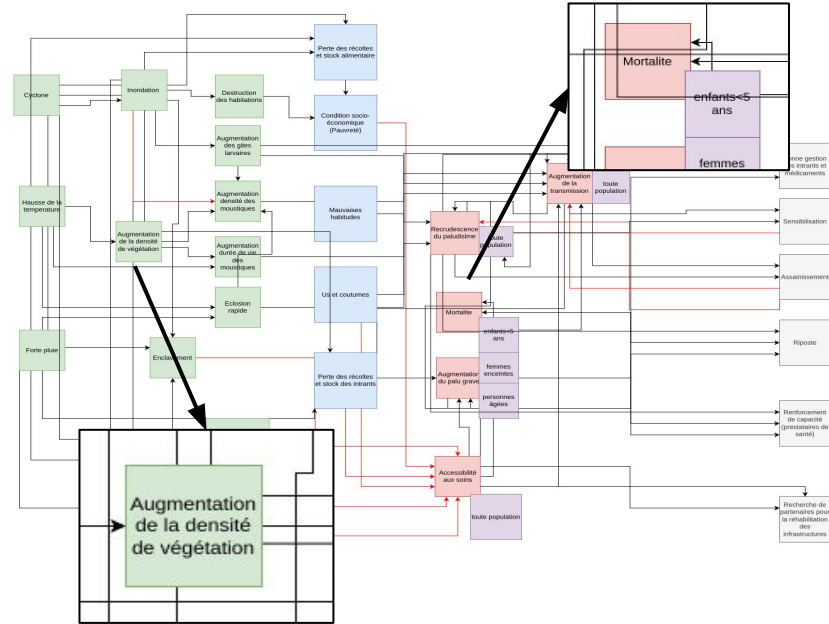
Acute  
Respiratory  
Inf.

# PRIDE-C



Predicting Infectious Diseases  
via Environment and Climate

# Model co-creation: conceptual to statistical



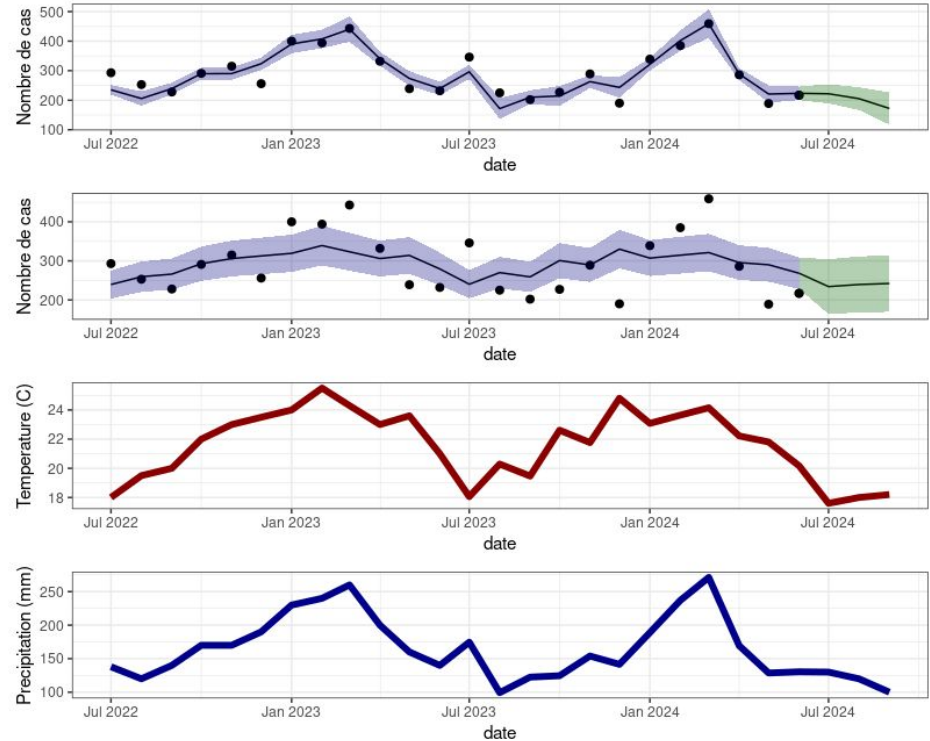
Used participatory research methods to identify environmental conditions and hazards relevant for each disease

# Model co-creation: conceptual to statistical

INLA models : Bayesian approximation with spatio-temporal structures

Models trained on 4 years of health, social, climate, and environmental data

Disease rates forecast up to 3 months in the future



# Data sources



Digitized health  
clinic registers



DHS-like  
household survey



Routine HMIS  
indicators

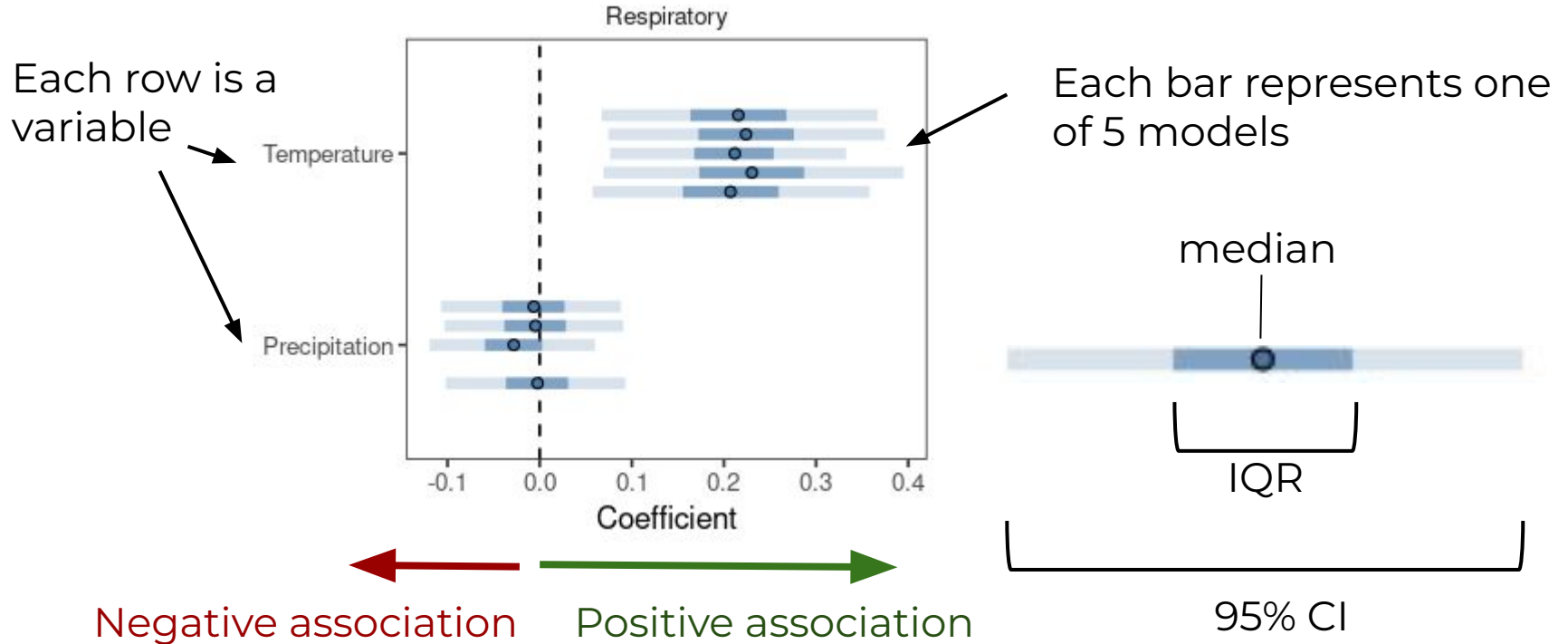


Environmental and climate  
data from satellite imagery

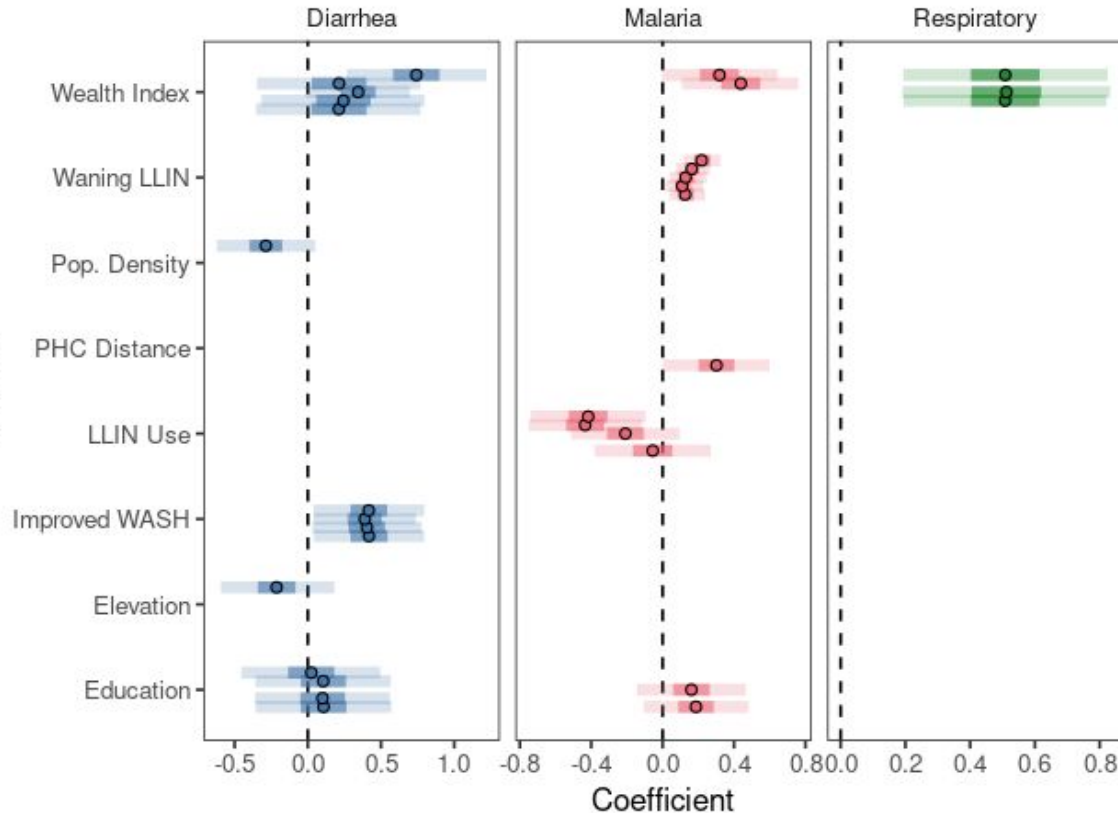


Crowd-sourced dataset of  
buildings and transport  
networks

# Results: How to Interpret



# Social and geographic variables



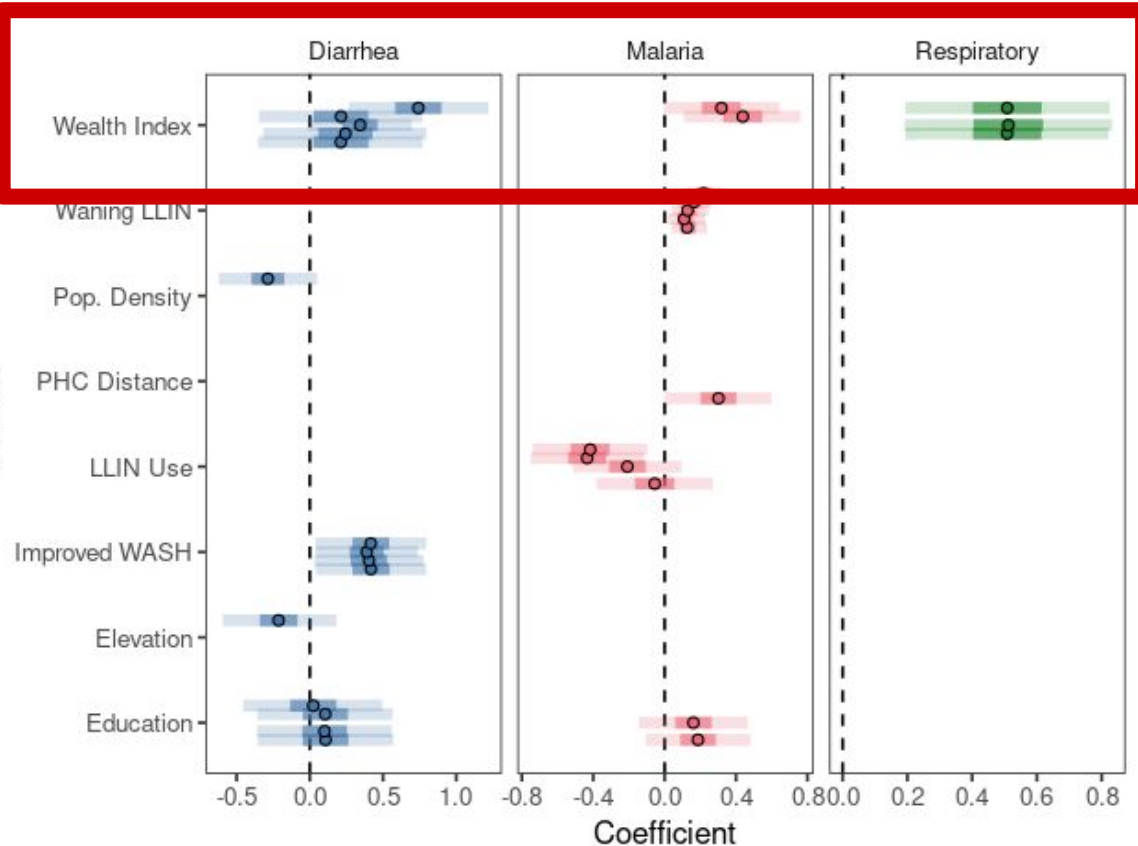
Wealth has a positive association for all diseases

Social variables may represent access to care, rather than exposure

- Need to correct for estimation bias

Importance of controlling for interventions (LLINs)

# Social and geographic variables



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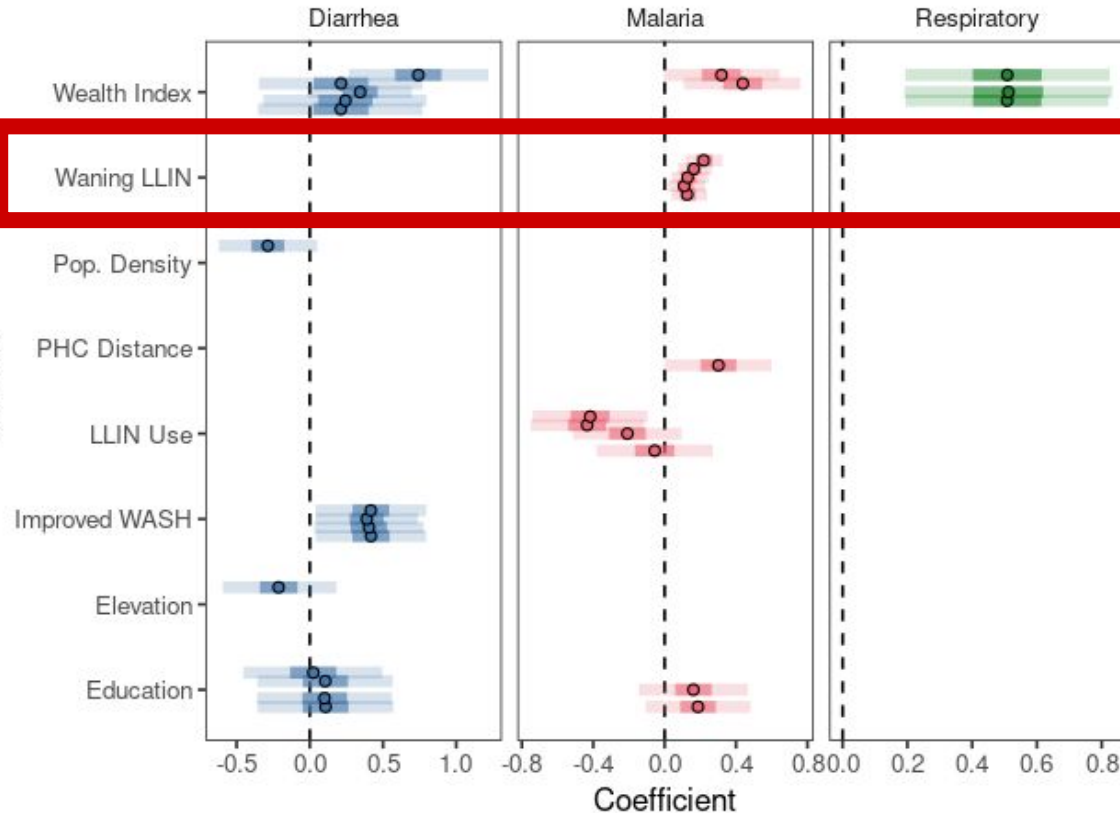
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# Social and geographic variables



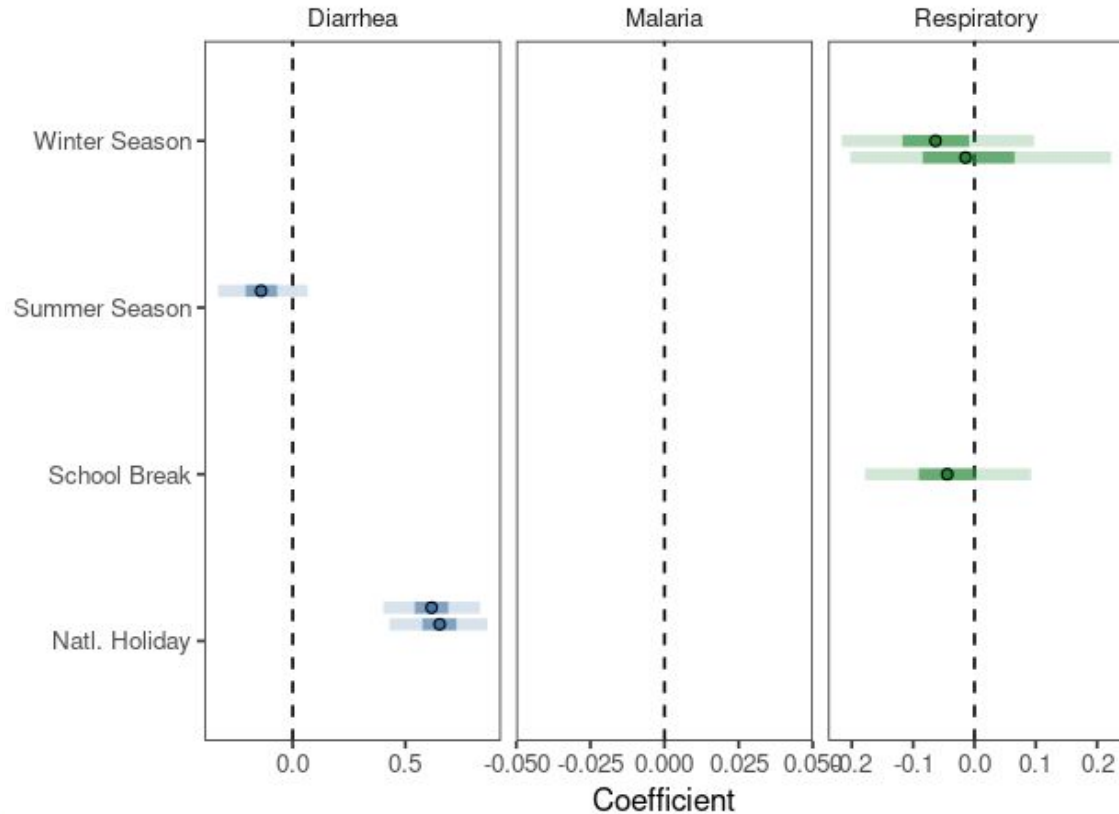
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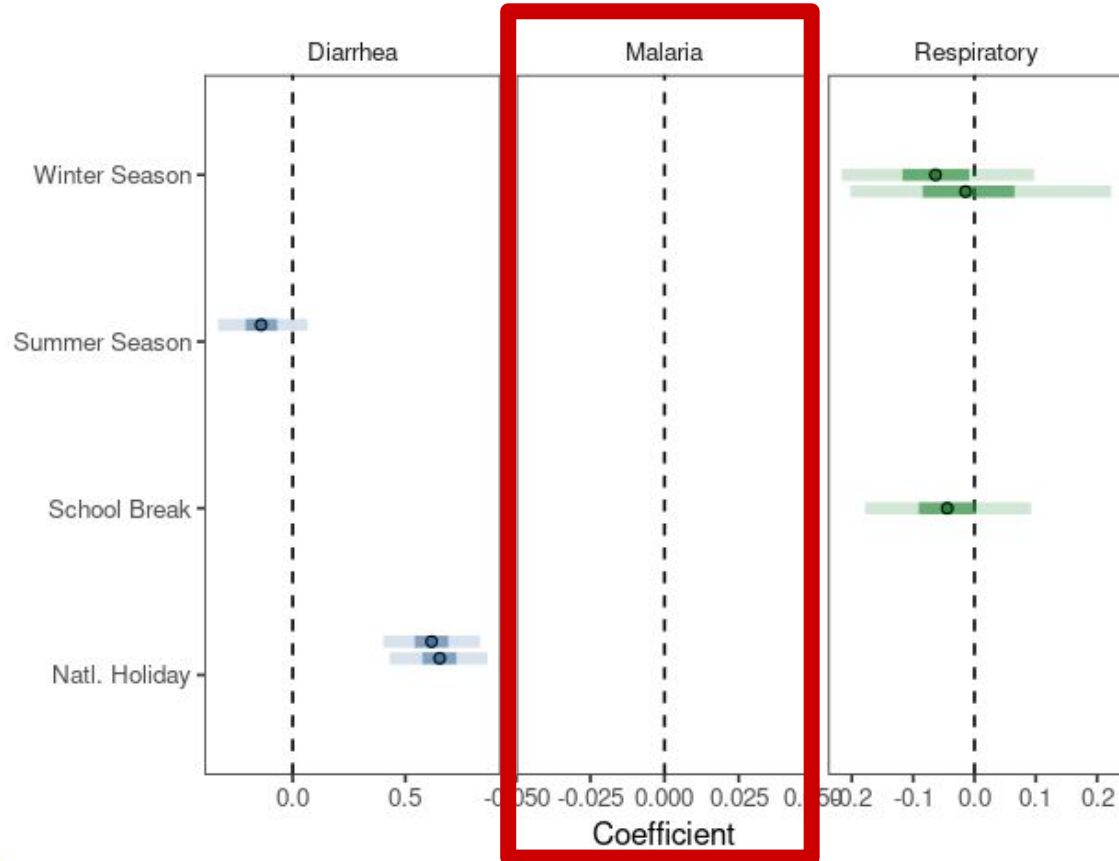
# Calendar variables



No calendar variables were identified as important for malaria

Strong role of national holiday for diarrheal disease

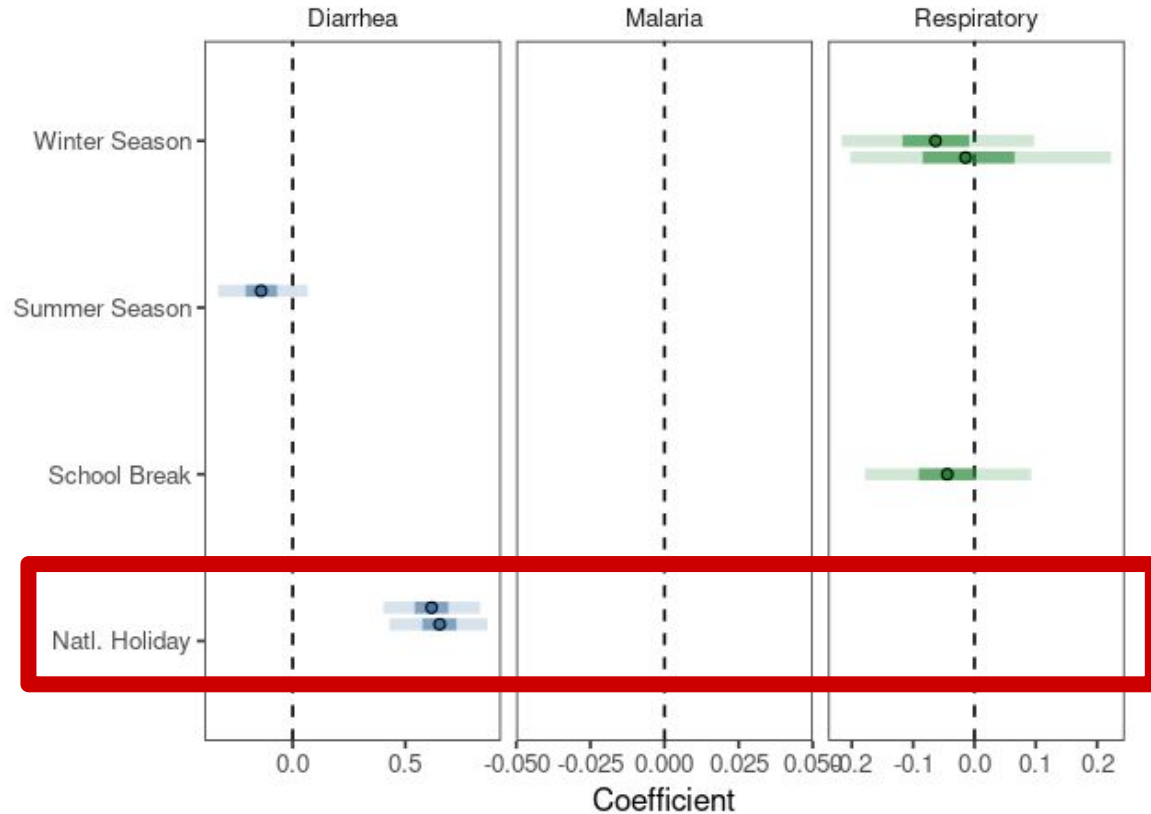
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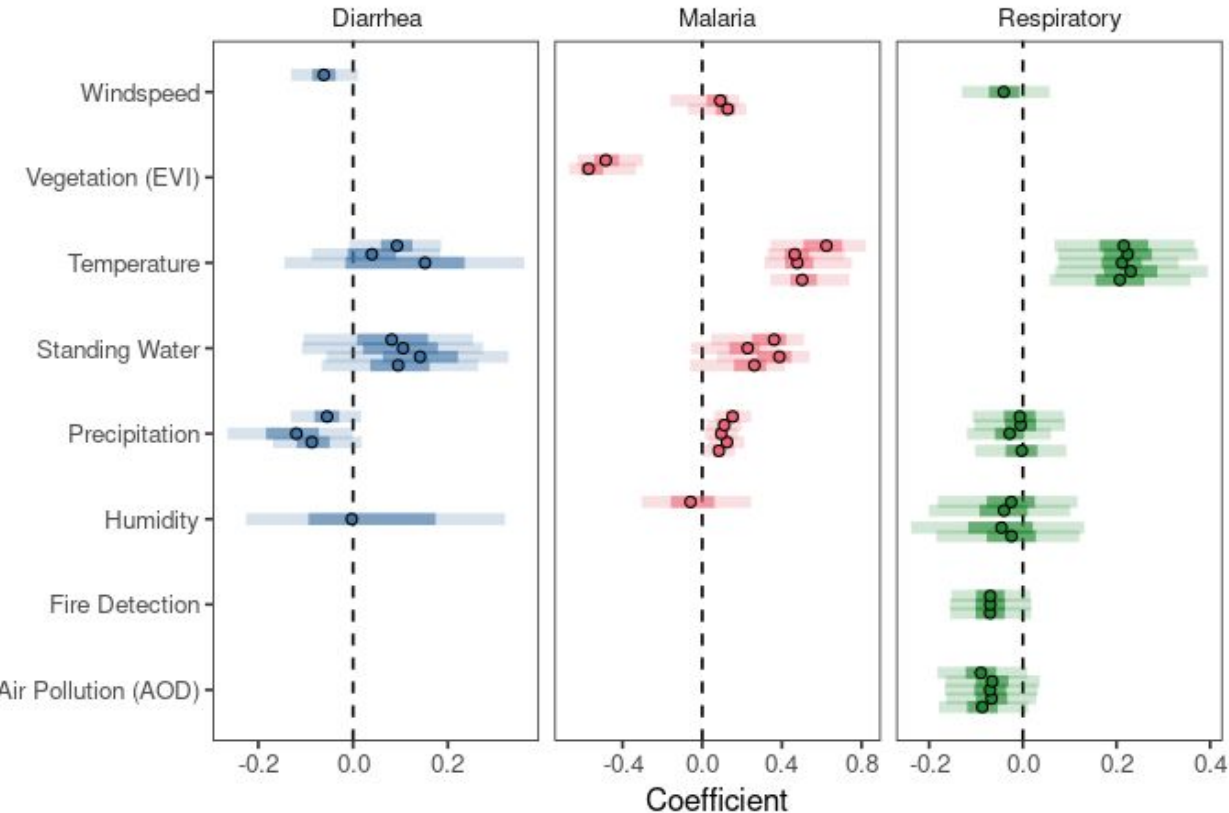
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# Climatic and environmental variables

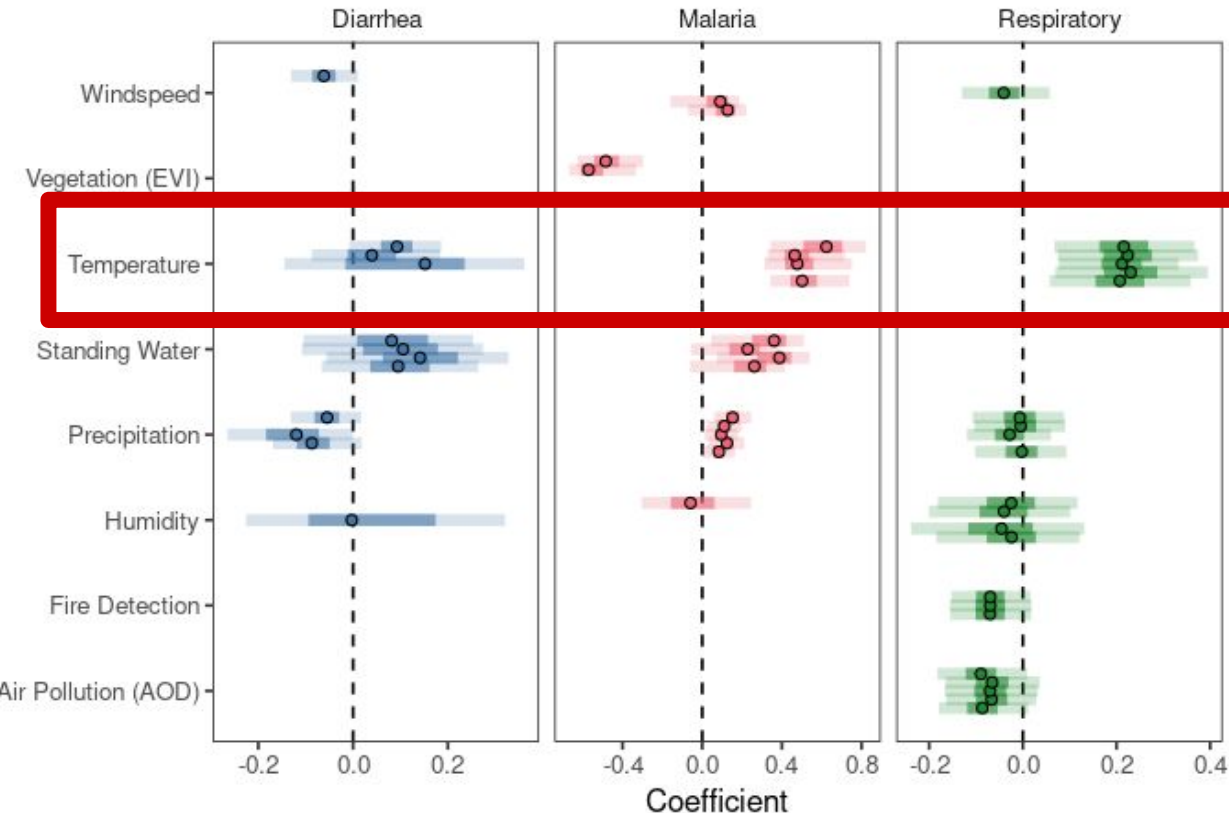


Many shared variables across diseases

Temperature is positively associated with all diseases

Variables with clear seasonality have larger coefficients  
→ Role of stochasticity

# Climatic and environmental variables

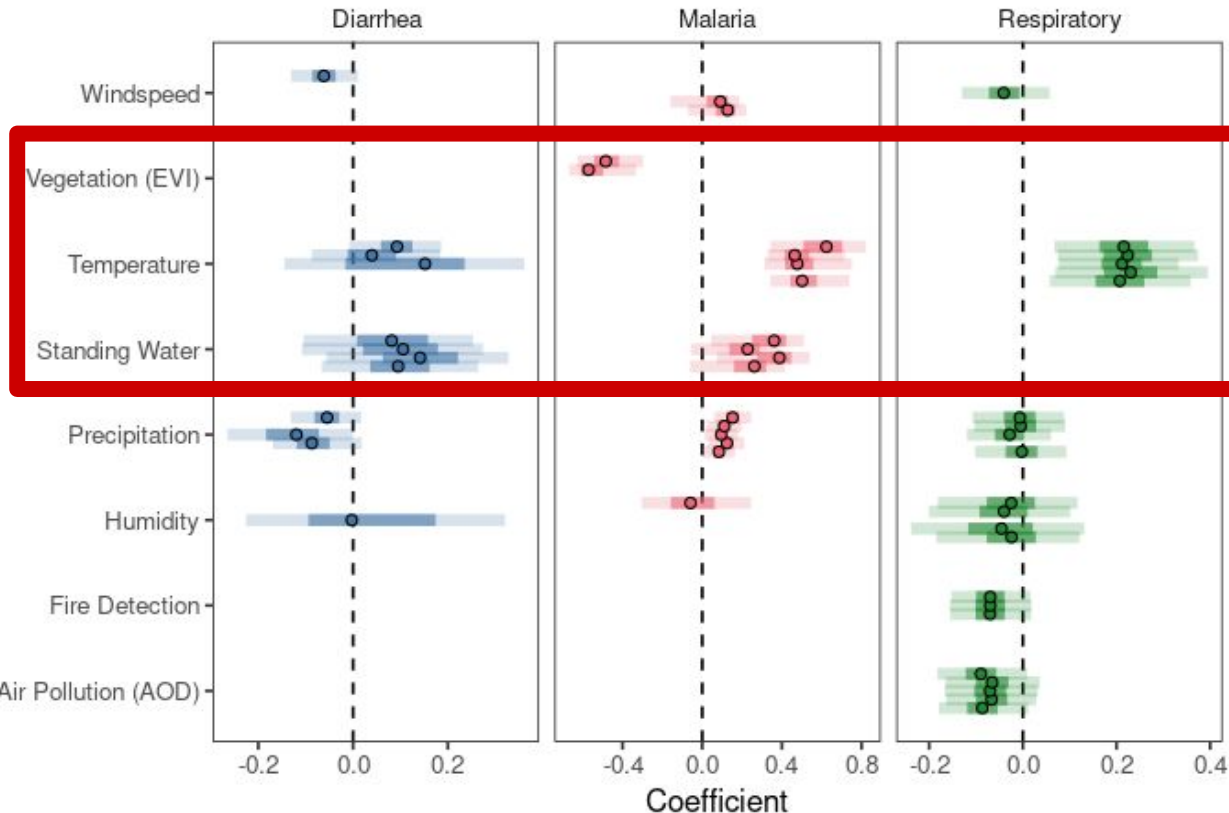


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# Climatic and environmental variables



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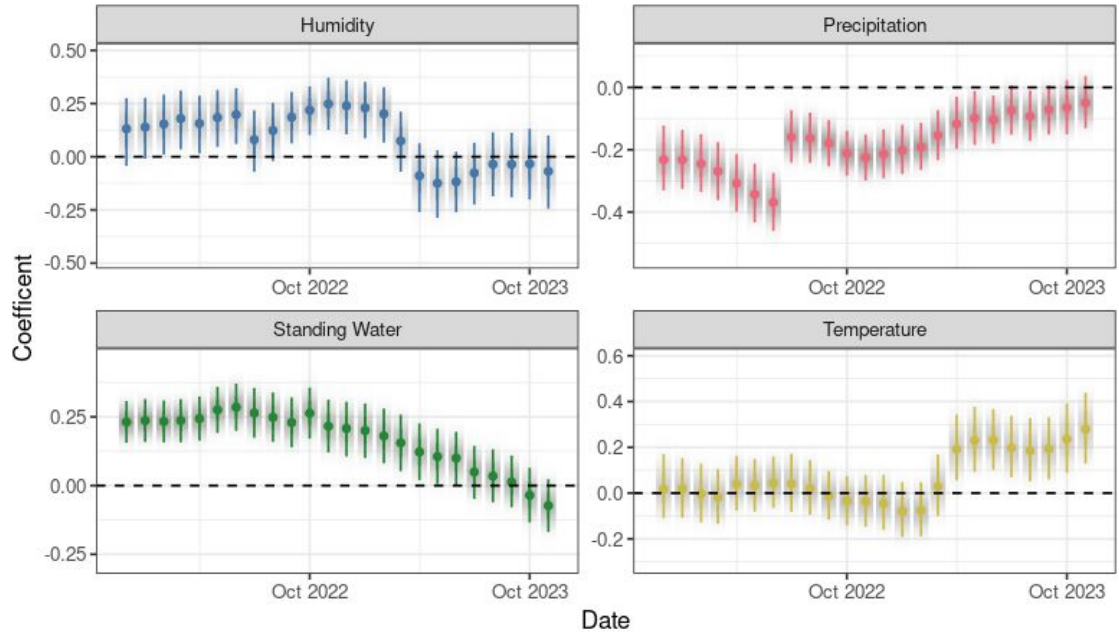
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# Variable coefficients change over time

Models trained on historical data must be updated regularly

Due to changing climate or interaction with changing social context (e.g. HSS)?





# Next steps

Creating ensembles of models using different modeling frameworks (ARIMA, Random Forest, GAM)

Second validation of models with health actors in November 2024

Deployment of beta version of DHIS2 application for district-wide use

Release of R package containing modeling workflow for use with DHIS2 data



## PRIDE-C Team



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Questions?

Vous avez des  
questions?

Misy fantanina?



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