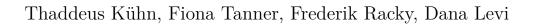
# Temperature sensitivity of dengue in Thailand



Bioinformatics Project group 4 team 4 climate sensitive infectious diseases Summer Term 2023

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#### The full abstract goes here

### Contents

1	Introduction	2
2	Material and Methods	3
3	Results 3.1 Result for Method A	4
4	Discussion	4

# Abbreviations

**DHF** Dengue Hemmorhagic Fever

ARIMA ...

**GAM** ...

### 1 Introduction

The Dengue virus is a vector borne virus, consisting of four serotypes (DENV 1-4). It is transmitted to humans by mosquitoes, more specifically by Aedes aegypti and Aedes albopictus (Phanitchat et al., 2019). The resulting infection goes asymptomatic in many cases, but can also cause Dengue Hemmorhagic Fever (DHF), characterized by symptoms like fever, headache, joint and muscle pain as well as (internal) bleeding and bruising (Gubler, 1998). Dengue is an emerging public health issue, with half of the worlds population at risk of infection. 390 million cases per year are estimated worldwide, with most of them not showing symptoms and thus not being reported. Affected areas range from sub-tropical to tropical regions, with south-east Asia, including Thailand being one of the most seriously affected regions. Newly affected areas also include Europe (WHO, 2023). The disease control of dengue is challenging due to the absence of an effective treatment or vaccine, which leaves only the treatment of symptoms with painkillers like paracetamol (WHO, 2023). Responsible institutions in affected areas currently focus on prevention, vector control, case control and prediction of possible future outbreaks (Phanitchat et al., 2019).

An important factor in predicting the epidemiological dynamics of Dengue is the climate: climate fluctuations due to recurring weather phenomenons and climate change are shown to influence *Aedes* biology and infections (Descloux *et al.*, 2012; Phanitchat *et al.*, 2019). In several studies, maximal temperature has an effect on dengue transmission (Descloux *et al.*, 2012), being associated with higher incidence (Phanitchat *et al.*, 2019) between 27 °C and 29,5 °C. Extreme global climate events like el Niño have been shown to affect disease outbreaks like Dengue as well (Anyamba *et al.*, 2019).

In Thailand, significant climate factors are monsoon (hot and rainy) between may and october and dry season between november and april. It has been shown that in northern districts of Thailand, there has been a detectable rise in temperature since the mid 20th century (Masud et al., 2016).

In this analysis, we are going to investigate the correlation between temperature and Dengue in Thailand from 2006 to 2020 to assess the significance of climate change, recurring climate fluctuations and extremes and geographical factors on Dengue infections. This will allow us to model a forecast for the development of Dengue infections in Thailand in the near future.

# 2 Material and Methods

ERA5 data (climate) The ERA5 database is a global climate database by the European Centre for Medium-Range Weather Forecasts (ECMWF), covering the earth in a 31 km horizontal grid up to 80 km in the atmosphere in the time period from 1950 to present. It was generated from measurements of various climate variables combined with a reanalysis of existing data and past reanalysis data to accurately model and complete the dataset in the given resolution (Hersbach et al., 2020). For this project, monthly temperature data 2m above ground for every province in Thailand in the timeframe of 2006 - 2020 was extracted.

Dengue data The Dengue case numbers were retrieved from annual infectious disease reports published by the Thailand ministry of health. Monthly case numbers of DHF for every province in the timeframe of 2006 - 2020 were used in this project.

Both datasets were used with a resolution at province level. As of 2011, Thailand has a total of 77 provinces, but had 76 provinces before 2011, as Bueng Kan was split from Nong Khai in 2011. For better compatibility of the data before and after 2011, the two new provinces were merged into a province equivalent to Nong Khai before 2011.

Descriptive analysis Linear regression is a method to describe a linear relationship between variables, using the minimum sum of squares between regression line and data points and identify possible trends (Schneider *et al.*, 2010).

**ARIMA** 

**GAM** 

### 3 Results

#### 3.1 Result for Method A

Results go here as subsections.

## 4 Discussion

Your Discussion goes here!

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