

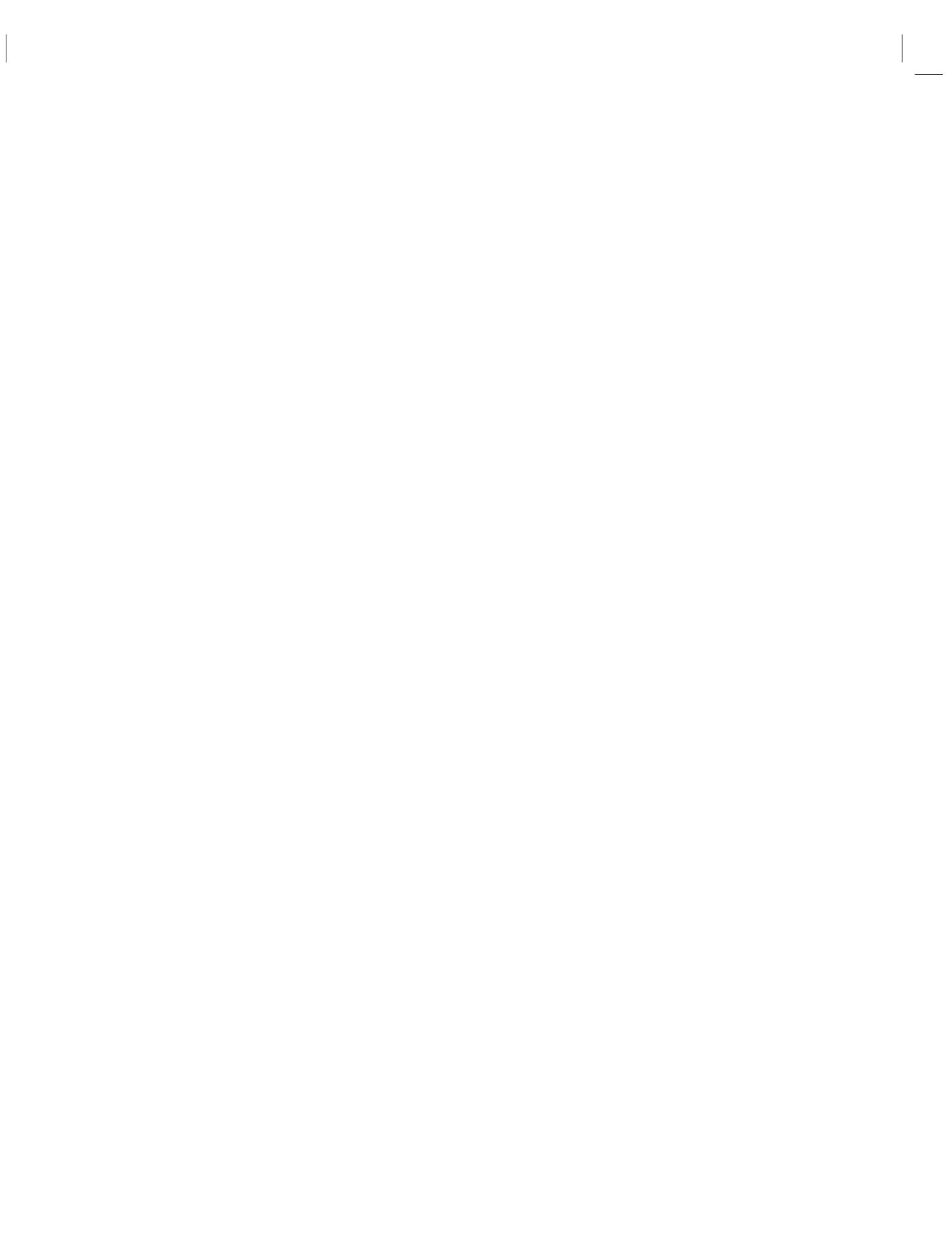
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Climate And Statistics



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Preface

Author: Henri Funk



As the world faces the reality of climate change, natural hazards and extreme weather events have become a major concern, with devastating consequences for nature and humans. The quantification and definition of climate change, extreme events and its implications for life and health on our planet is one of the major concerns in climate science.

This book explains current statistical methods in climate science and their application. We do not aim to provide a comprehensive overview of all statistical methods in climate science, but rather to give an overview of the most important methods and their application. This book is the outcome of the seminar “Climate and Statistics” which took place in summer 2024 at the Department of Statistics, LMU Munich.



FIGURE 1: Creative Commons License

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¹<http://creativecommons.org/licenses/by-nc-sa/4.0/>



Technical Setup

The book chapters are written in the Markdown language. To combine R-code and Markdown, we used rmarkdown. The book was compiled with the bookdown package. We collaborated using git and github. For details, head over to the book's repository².

²https://github.com/henrifnk/Seminar_ClimateNStatistics



1

DL Classification of Weather Patterns over Europe

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Degree: Master

1.1 Abstract

Daily weather in the mid-latitudes is dominated by large-scale atmospheric circulation types (CTs), which influence regional climate variability and extremes. Accurate classification of these CTs is crucial for diagnosing long-term climate dynamics and analysing future changes. This study implements and extends a novel deep learning-based classification method for European CTs, building on the Hess & Brezowsky (HB) framework.

Using a convolutional neural network (CNN) architecture (inspired by applications in climate science and optimization techniques from), we trained the model on high-resolution ERA5 reanalysis data (1950–1980), with sea-level pressure (SLP) and geopotential height at 500 hPa (z500) fields as key predictors. To ensure robustness, a nested cross-validation approach was employed, achieving an overall accuracy of 53.17% and a macro F1 score of 55.14%, outperforming traditional classification methods.

Applied to the CanESM2 ensemble projections, which simulate data with assumptions that climate conditions were consistent with history, our results show significant future frequency shifts in CTs, notably an increase in Icelandic High, Cyclonic (HNZ) occurrences during both summer and winter half-years. This methodological advancement not only enhances classification accuracy but also offers robust ensemble analyses critical for climate-informed decision-making.

1.2 Introduction

In the mid latitudes, daily weather is dominated by large-scale atmospheric circulation patterns, defined by the position of high and low pressure centres (Häckel, 2021; Mittermeier et al., 2022). Due to steep temperature and pressure gradients along the frontal zone between polar and tropical air masses (around 45° latitude), the jet stream is formed in about 10 km altitude. It is characterized by mean wind speeds of 300 km/h and the long-term average wind direction is westerly (Häckel, 2021). Here, dynamic high and low pressure systems develop (Häckel, 2021; Mittermeier et al., 2022). However, the westerly jet stream is not stationary, but deflected with varying amplitude creating the so-called Rossby waves, which influence the position of high and low pressure systems (Häckel, 2021). While an infinite number of atmospheric conditions would be conceivable, recurrent weather patterns with similar meteorological features are observed in practice. Therefore, it is possible to classify weather patterns according to synoptic properties (Bissolli, 2001).

The first concept for a classification of different weather patterns was introduced by [Baur et al. \(1944\)](#). They used surface pressure charts from 1881 to 1939 and assigned the respective weather situation to a specific class for each day based on the spatial distribution of atmospheric pressure and the position of the frontal zones. This lead to a classification into 21 Grosswetterlagen for Europe and East Atlantic ([Baur et al., 1944](#)). In 1963 Baur defined Grosswetterlage as the mean spatial air pressure distribution of a large area, at least the size of Europe, over a period of several successive days. This classification has been revised and improved several times. In 1977, Hess and Brezowsky published their third edition of a revised catalogue of Grosswetterlagen. Here, the authors determined that a Grosswetterlage should only be classified if it can be recognised on at least three successive days. Furthermore, the number was extended to 29 different Grosswetterlagen. In addition to the surface pressure chart, the classification was also based on the geopotential height in 500hPa ([Bissolli and Dittmann, 2001](#); [Bissolli, 2001](#); [Werner and Gerstengarbe, 2010](#)). In the following, Werner and Gerstengarbe published seven editions of a catalogue of the Grosswetterlagen in Europe with updated data. The last edition (2010) covers the years 1881 to 2009 and provides daily information on the Grosswetterlage over Europe. Today, the German Weather Service (DWD) still constantly updates the catalogue of European Grosswetterlagen and publishes the results monthly. Since 1944, the classification of Grosswetterlagen is conducted by experts, hence it is subjectively biased ([Hess, 2005](#)). In the following, the classification according to Hess & Brezowsky will be referred to as HB CT.

HB CTs can be attributed to certain weather situation at certain locations in Europe ([Werner and Gerstengarbe, 2010](#)). Hence in the past, the classification of HB CTs has been used to identify regularities in frequency and duration of occurrences of certain weather conditions. [Bissolli and Dittmann \(2001\)](#) describes a relationship between frequency and duration of the HB CTs and mean annual air temperature and precipitation. Moreover, extreme weather events—such as heavy rainfall, floods, and heat waves—can often be linked to specific HB CTs ([Mittermeier et al., 2022](#)). Such kinds of application could not only be useful to analyse past climate. Future climate projections could also be analysed using the HB CTs, allowing conclusions to be drawn about future developments. However, analyzing future CTs requires many model ensembles to account for internal variability ([Wyser et al., 2021](#)). Manual classification of HB CTs for many ensemble members is no longer possible. It is therefore necessary to consider the possibility of automatization ([Mittermeier et al., 2022](#)). [Mittermeier et al. \(2022\)](#) have found a way to classify HB CTs automatically using a deep learning classifier.

1.3 Data

1.3.1 Data Sources

This section describes in detail the datasets utilized for model training, validation, and testing, providing comprehensive explanations of their characteristics, purposes, and roles within the study. Due to data availability constraints, we replaced the original study's ERA-20C ([Poli et al., 2016](#)) and SMHI-LENS ([Wyser et al., 2021](#)) data with ERA5 ([European Centre for Medium-Range Weather Forecasts, 2017](#)) (higher resolution, 1950–1980) for training and CanESM2 ([Hua et al., 2015](#); [Government of Canada, 2025](#)) (use only 12 of 50 ensemble members) for projections.

1.3.1.1 Historical Data - Training and Validation

We use reanalysis dataset, which is the historical atmospheric data reconstructed by combining observational data and model simulations, for training and validation. Data is provided by European Centre for Medium-Range Weather Forecasts (ECMWF).

We only use two variables mentioned above:

1. SLP: Atmospheric pressure measured at Earth's surface, vital for identifying weather systems such as cyclones and anticyclones.
2. z500: Height of the 500 hPa pressure level in the atmosphere, indicative of atmospheric wave patterns and mid-tropospheric dynamics.

Due to reanalysis data's high-quality, accurate, and detailed historical atmospheric representation, it is enabled to get a robust and precise training of the deep learning model.

The difference of ERA5 (Figure 1.1) and ERA-20C (Appendix Figure 1.7) lies on:

TABLE 1.1: Difference between two reanalysis dataset

Reanalysis Dataset	Temporal Coverage	Spatial Resolution
ERA-20C (paper)	1900–1980	Approximately 125 km
ERA5 (implementation)	1950–1980	High-resolution, about 31 km

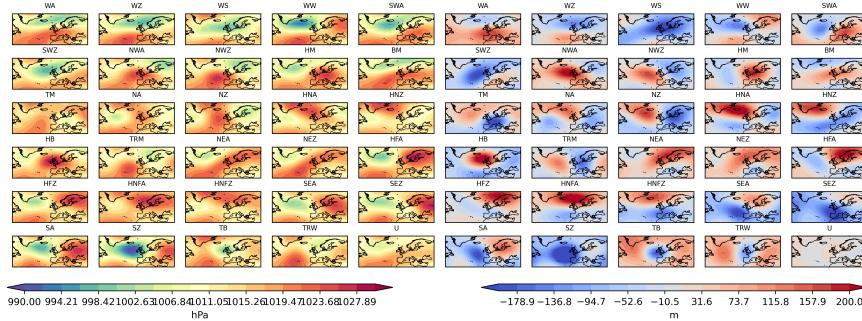


FIGURE 1.1: Synoptic patterns of the 29 HB CTs created using ERA5 data averaged over the period 1950 - 1980, showing SLP on the left and z500 on the right side.

1.3.1.2 Future Climate Projections - Change Analysis

We use ensemble datasets which projected future climate conditions under specific scenarios, to evaluate future changes in atmospheric circulation, considering internal variability, essential for accurate and reliable climate change projections. The data should be preprocessed with the same variables, scale, domain, resolution as training data.

1. CanESM2 ([Government of Canada, 2025](#))
 - Name: CanESM2 (Canadian Earth System Model version 2)
 - Provider: Canadian Centre for Climate Modelling and Analysis (CCCma)
 - Model Used: Climate Model Intercomparison Project phase 5 (CMIP5)
 - Ensemble Members: 50 (due to downloading limitation, we only used 12 ensemble)
 - Scenario: historical, simulate the condition as observed past climate from 1850 to at least 2005 ([Friedlingstein et al., 2008](#)).
 - Temporal Coverage:
 - Historical Reference: 1971–2000
 - Future Projections: 2031–2060
 - Spatial Resolution: Around 2.8°
2. SMHI-LENS ([Wyser et al., 2021](#))

- Name: SMHI-LENS (Swedish Meteorological and Hydrological Institute Large Ensemble)
- Provider: Swedish Meteorological and Hydrological Institute
- Model Used: EC-Earth3 (Climate Model Intercomparison Project phase 6, CMIP6)
- Ensemble Members: 50 (to effectively capture internal variability)
- Scenario: SSP3-RCP7.0, representing a high GHG emission pathway
- Temporal Coverage:
 - Historical Reference: 1991–2020
 - Future Projections: 2071–2100
- Spatial Resolution: Around 0.7°

As a result with higher greenhouse gas emission under SSP3-RCP7.0 scenario, and with a more distant future in the paper:

1. Future warming and circulation shifts might be more pronounced than our implementation.
2. Frequency changes in HB CTs may appear larger in the paper, simply because the climate change signal is stronger.

1.3.2 Data Preprocessing Steps

1. Reset spatial domain: only use data in Europe and the North Atlantic region (30°N–75°N latitude, 65°W–45°E longitude)
 2. Spatial Regridding: All datasets were interpolated to a common 5° grid resolution, ensuring consistency and computational efficiency
 3. Seasonal Centering: Applied to remove seasonal biases, helping the model better recognize atmospheric patterns independently of seasonal influences.
-

1.4 Models and Training

1.4.1 CNN Architecture

The classification model used in this study is based on a CNN, specifically designed to handle the image-like structure of atmospheric data (SLP and z500). The CNN architecture (Figure 1.2) comprises:

1. Input Layer: Two separate input channels corresponding to SLP and z500.
2. Convolutional Layers: Two convolutional layers applied to extract spatial patterns and features from input atmospheric fields. These layers use convolutional kernels to identify local and spatially coherent patterns.
3. Dropout Layer: Included for regularization to mitigate overfitting by randomly dropping out units during training, thus improving the model's generalization.
4. Fully Connected Layers: Two fully connected layers following the convolutional and dropout layers, used for integrating learned spatial features into final predictions of HB CTs.
5. Output Layer: Outputs softmax probabilities for each of the 29 HB CTs, which are then converted into classifications.

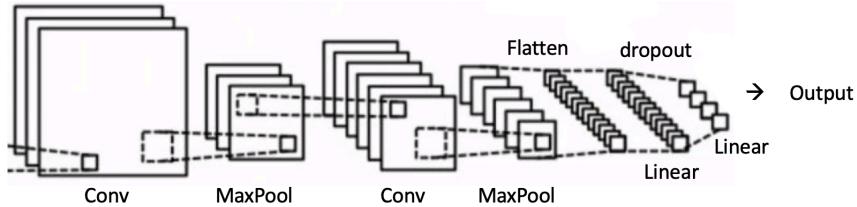


FIGURE 1.2: Schematic diagram of the CNN network

1.4.2 Training Protocol

1.4.2.1 Adam Optimizer

Adam (Adaptive Moment Estimation) ([Mehta et al., 2019](#)) is an optimization algorithm widely used for training neural networks. It combines the benefits of two other methods—AdaGrad ([Ward et al., 2020](#)), which adapts the learning rate for each parameter, and RMSProp ([Zou et al., 2019](#)), which considers the exponentially decaying average of squared gradients—to adjust the learning rates adaptively during training.

Adam optimizer efficiently handles noisy and sparse gradients, common in deep learning tasks, facilitating faster convergence and better handling of large parameter spaces typical of CNNs. This makes Adam particularly suitable for our CNN-based classification model, ensuring effective and efficient training.

1.4.2.2 Bayesian Optimization for Hyperparameter Tuning

To achieve an optimal configuration of hyperparameters for our CNN, we employed Bayesian optimization ([Garnett, 2023](#)), specifically using the Tree-structured Parzen Estimator (TPE) method ([Ozaki et al., 2022](#)) implemented in the Optuna library. Bayesian optimization is particularly suited for deep learning hyperparameter tuning due to its efficiency and ability to intelligently balance exploration and exploitation of the hyperparameter space, thus significantly reducing computational cost compared to grid or random search.

Specifically, the following hyperparameters were optimized:

1. Learning Rate: The learning rate controls the magnitude of updates to the network weights during training. We allowed this hyperparameter to vary logarithmically between 10^{-4} and 10^{-2} , enabling the optimizer to effectively explore both subtle and more substantial updates in network parameters.
2. Weight Decay: Serving as a regularization term to prevent overfitting, the weight decay was also sampled on a logarithmic scale ranging from 10^{-5} to 10^{-3} . This enabled fine control over model complexity and encouraged better generalization.
3. Dropout Rate: Dropout prevents overfitting by randomly dropping neurons during training, forcing the model to learn robust representations. We tuned dropout rate uniformly within a range from 0.2 to 0.6, allowing optimization of the model's regularization intensity.
4. Convolutional Layer Parameters:
 - Number of Output Channels in First Convolutional Layer (out_channels1): Chosen categorically from [4, 8, 16], optimizing the depth and feature extraction capability of initial convolutional operations.
 - Number of Output Channels in Second Convolutional Layer (out_channels2): Selected from [8, 16, 32], further refining the network's capability to learn hierarchical spatial patterns.
 - Kernel Size: Kernel size impacts the spatial context the model considers when learning features. We considered kernel sizes of [3, 5, 7], balancing between capturing detailed local patterns and broader spatial structures.
 - Fully Connected Layer Size: The number of neurons in the fully connected layer was chosen from [32, 64, 128], directly influencing the model's ability to integrate learned spatial features into higher-level abstractions suitable for classification.

The Bayesian optimization was performed over 100 trials, systematically exploring these hyperparameters. Each trial evaluated a distinct combination of parameters based on validation accuracy. The final selection of hyperparameters was determined by the combination that produced the highest validation accuracy, ensuring the CNN model's optimal performance on unseen data while minimizing the computational burden associated with hyperparameter tuning.

1.4.2.3 Nested Cross-Validation

Nested cross-validation ([Zhong et al., 2023](#)) involves two cross-validation loops:

1. Outer Loop: Evaluates the generalization performance of the model on independent test sets.
2. Inner Loop: Used for hyperparameter tuning and model selection within each training fold.

We use 5 folds of outer loop and 5 folds of inner loop.

1.4.2.4 Epochs and Early Stopping

Training was set to a maximum of 35 epochs, with early stopping criteria implemented, patience of 6 epochs without improvement, to prevent overfitting and reduce training time.

1.4.3 Predicting Technique

1.4.3.1 Ensemble Learning

Ensemble learning ([Dietterich et al., 2002](#)) combines predictions from multiple models to enhance predictive performance. In this case, multiple CNN models trained with different initial conditions. A deep ensemble of 30 independently trained CNN models, each initialized with different random weights, is utilized.

Predictions from individual CNN models are aggregated using a weighted averaging method, where each model's contribution is proportional to its validation performance. This enhances robustness, reduces prediction uncertainty, and provides stable and reliable classification results by capturing a wider range of atmospheric variability.

1.4.3.2 Transitional Smoothing

According to the definition, a HB CT must last at least three days ([Hess and Brezowsky, 1952](#)) to represent a stable atmospheric pattern. Without enforcing this constraint, the raw neural network predictions may contain short, noisy transitions that are meteorologically implausible. Thus, transition smoothing is necessary to:

1. Remove unrealistic, very short-lived classifications.
2. Produce physically consistent and stable time series of HB CTs.
3. Ensure that predictions align with domain-specific expectations.

According to the original paper:

1. Step 1: Identify all transitions where the predicted HB CT lasts fewer than three days.
2. Step 2: Check neighborhood consistency:
 - If the HB CT before and after the short transition is the same, assign this type to the transition days.

3. Step 3: If different types occur before and after:

- Compare the predicted probabilities (confidence scores) for each neighboring type.
- Assign the transition days to the neighbor with the higher predicted probability (i.e., stronger membership).

This method ensures that short transitions are corrected in a physically meaningful way, based on both label consistency and model confidence. Since the original code is not available, our transition_smoothing function follows the paper's logic but introduces minor adaptations for greater robustness and flexibility:

TABLE 1.2: Transitional smoothing implementation

Aspect	Paper Description	Our Code Implementation	Reason for Change
Minimum duration	3 days fixed	min_duration parameter (default=3)	Makes the method flexible for other settings.
Neighbor block length trust	Not specified clearly	min_neighbor_run_length parameter (default=2)	Ensures smoothing only when neighboring segments are stable enough.
Handling first and last segments	Implicitly ignored	Explicitly handled	Avoids out-of-bound errors at sequence edges.
Probability averaging	Single-point comparison	Average probability across transition segment	More robust by considering entire transition region, not a single timestep.
Fallback behavior when neighbors are not strong	Not detailed	Fallback to the only trusted neighbor if one side is strong enough	Avoids wrong smoothing when one neighbor is unreliable.

1.5 Results

1.5.1 Classification Performance

1.5.1.1 F1-score

The classification performance of our model was evaluated using a confusion matrix, F1-scores per class, macro F1-score ([Opitz and Burst, 2019](#)), and overall accuracy ([Alberg et al., 2004](#)), following the approach used in the reference study ([Mittermeier et al., 2022](#)). The overall accuracy achieved by our model on the cross-validation test set is 53.17%, and the macro F1-score is 55.14%. Among all HB CTs (Appendix Figure 1.9), the best performing classes in terms of F1-score are WZ (70.21), HM (63.7), and HB (62.78). These types exhibit distinct spatial patterns, aiding CNN identification. The lowest F1-scores are observed for classes like NA (33.61) and SEZ (44.37). These are likely due to class imbalance caused by rare occurrence. Generally, accuracy for each type is quite high, which may imply overfitting.

The confusion matrix (Appendix Figure 1.17) shows that most misclassifications occur among HB CTs with similar dynamical structures, consistent with expectations from atmospheric science. Diagonal dominance is visible but moderate, reflecting the challenging nature of this classification task.

Compared to the paper results, which reported an overall accuracy of 41.1% and a macro F1-score of 38.3% (Mittermeier et al., 2022), our model achieves slightly higher overall accuracy and macro F1-score.

TABLE 1.3: Metric comparison

Metric	Paper (%)	Our Model (%)
Macro F1-score	38.3	55.1
Overall accuracy	41.1	53.2

The boxplots of F1-score under each HB CT (Figure 1.6; Appendix Figure 1.13) confirm our conclusion:

1. Original study has considerably lower overall F1-scores (around 0.3–0.5), indicating challenges in reliable classification.
2. Our model demonstrates much higher stability with F1-scores generally between 0.80–0.95, suggesting improved reliability and robustness in identifying circulation patterns.

1.5.1.2 RMSE

To further evaluate the quality of the predictions, the Root-Mean-Square Error (RMSE) between the predicted HB CT composites and the true label composites was calculated for variable SLP.

Equation for the calculation of the RMSE with I being the predicted image (in our case: signature plot of the deep learning classifier) and K being the reference image (in our case: signature plot of the labels). M are number of rows and N the number of columns of the pictures to compare. The RMSE thus compares the pixel-wise values of two images. A value of zero indicates a perfect match (Mittermeier et al., 2022; Mueller et al., 2020).

$$\text{RMSE} = \sqrt{\frac{1}{M \times N} \sum_{i=0, j=0}^{M-1, N-1} [I(i, j) - K(i, j)]^2}$$

Additionally, RMSEs for false positives and false negatives were computed separately.

The average RMSE for our implementation's predictions across all HB CTs is 0.88, while the false positives and false negatives yielded RMSEs of 1.08 and 1.23, respectively.

TABLE 1.4: RMSE comparison

RMSE Category	Paper	Implementation
Prediction RMSE	0.89	0.74
False positives RMSE	1.09	1.45
False negatives RMSE	1.28	0.69

Our implementation achieves a lower prediction RMSE (0.74) compared to the original paper (0.89) (Mittermeier et al., 2022). This indicates that, on average, the spatial patterns of the correctly predicted HB CTs are more similar to the true labels in our model than in the reference study.

However, the false positives RMSE is much higher (1.45) than in the paper (1.09) (Mittermeier et al., 2022). This suggests that when our implementation wrongly predicts a HB CT, the resulting spatial pattern is less similar to the true pattern than in the paper.

Our implementation shows a lower false negatives RMSE (0.69) compared to the paper (1.28) (Mittermeier et al., 2022). This means that when the model misses a true HB CT, the spatial signature remains relatively closer to the correct pattern than in the paper.

The overall pattern suggests that:

1. Our implementation is better at maintaining correct structures when making predictions or missing true types.
2. However, when making incorrect predictions (false positives), our implementation model tends to produce worse distortions than the model used in the paper.

1.5.2 Signature Plot Analysis

Figure 1.3 shows the signature plots for four HB CTs selected based on our RMSE results:

1. Two HB CTs with lowest RMSE: WZ and TRW (best performance)
2. Two HB CTs with highest RMSE: NA and HNFA (worst performance)

For each HB CT, four different composite plots are displayed:

1. Labels: Composite of true labels.
2. Predictions: Composite of model predictions.
3. False Positives: Days predicted as the HB CT but labeled differently.
4. False Negatives: Days labeled as the HB CT but predicted differently.

RMSE values are shown below each panel, comparing each composite to the true label composite.

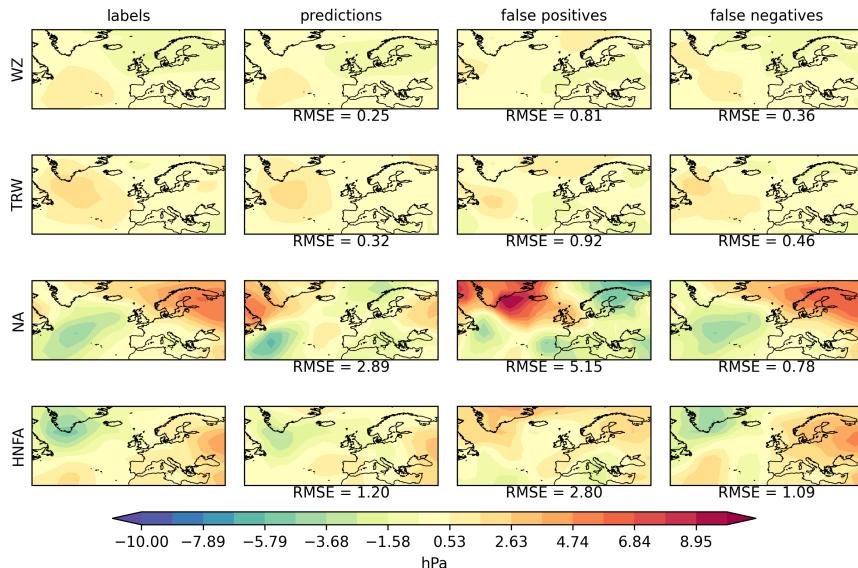


FIGURE 1.3: Signature plots of four selected circulation patterns at slp. Column 1: labels showing the indicated HB CT, Column 2: deep ensemble predictions showing the indicated HB CT, Column 3: signature pattern, when the deep ensemble predicts the indicated HB CT while labels state differently, Column 4: labels stating the indicated HB CT while deep ensemble predicts differently. The RMSE values are calculated by comparing the respective signature plot to the signature plot of the labels (column 1).

Based on our results, we can observe that:

1. WZ:

- The predicted composite closely matches the label pattern (RMSE = 0.25).
- False positives and false negatives show moderate RMSE (0.81 and 0.36), indicating stable identification of WZ.

2. TRW:

- Predictions for TRW are also accurate (RMSE = 0.32).
- False positives (0.92) and false negatives (0.46) suggest TRW is generally recognized reliably.

3. NA:

- Predictions for NA have high RMSE (2.89), indicating difficulty in capturing the correct structure.
- False positives (5.15) show very large errors, and false negatives (0.78) suggest missing NA patterns is common.

4. HNFA:

- Predictions for HNFA also show high RMSE (1.20).
- False positives (2.80) and false negatives (1.09) are large, implying HNFA is a difficult type for the model to classify.

Differences from the original paper (Appendix Figure 1.10) arise primarily from data variations, model differences, and random initialization.

Although the specific CTs selected differ, the general trend remains consistent with the findings of the paper:

1. Some HB CTs are easier to predict (low RMSE), others are harder (high RMSE).
2. False positives tend to have higher RMSE than correct predictions.
3. A larger discrepancy in the color distribution indicates a higher complexity of HB CT, and false negatives vary depending on HB CT complexity.

Thus, while specific HB CTs differ, our results fundamentally agree with the conclusions of the paper that certain patterns are systematically easier or harder to classify, and signature plots are effective to visualize these tendencies.

1.5.3 Frequency Distribution of HB CTs

Figure 1.4 shows the frequency distribution of the 29 HB CTs, expressed as the average number of days per year for the period 1950–1980. For each HB CT, the frequency derived from the true labels (blue bars) is compared to the frequency derived from the network predictions (yellow bars).

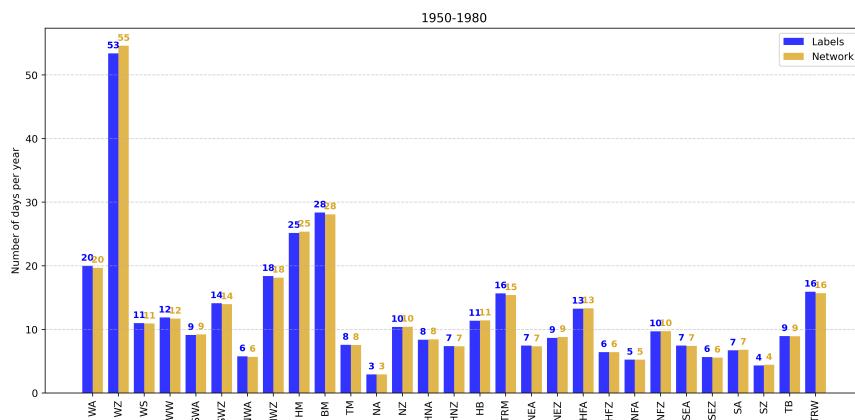


FIGURE 1.4: Frequency distribution of the 29 HB CTs in number of days per year for the training period 1950–1980 (blue) and for predictions of the CNN (yellow).

Overall, the CNN reliably reproduces observed HB CT frequencies:

1. Dominant HB CTs (e.g., WZ, HM) are captured accurately, reflecting robust model learning.
2. Rarer HB CTs (e.g., NA, SZ) exhibit minor deviations, suggesting challenges in accurately learning patterns due to fewer examples, which may indicate slight overfitting.

In comparison with the original study (Appendix Figure 1.11), despite differences in dataset periods (our study: 1950–1980; original: 1900–1980), frequency patterns remain consistent. Slight discrepancies in rarer HB CTs are likely due to dataset differences, emphasizing the importance of extended or more balanced training datasets for better generalization.

The agreement between the label frequencies and the network predictions, as well as the similarity to the paper’s findings, indicates that our model effectively learns the relative occurrence rates of the different HB CTs. Overall, the frequency distribution results support the conclusions of the original study, confirming that the network can reliably reproduce the occurrence rates of various HB CTs.

1.5.4 Future Change

Figure 1.5 and Figure 1.6 show the relative changes in the frequency of occurrence of the 29 HB CTs between the future period (2031–2060) and the historical reference period (1971–2000), for the entire year, winter half-year, and summer half-year, based on 12 climate model realizations. Figure 1.5 is the conclusion directly from predicted results, while Figure 1.6 is obtained after transition smoothing. Positive values indicate an increase in occurrence; negative values indicate a decrease. The lower panels show the F1-scores of the classification model ensemble. Higher F1-scores indicate better classification stability and reliability. The grouping by wind direction allows visual comparison of performance across different HB CT categories.

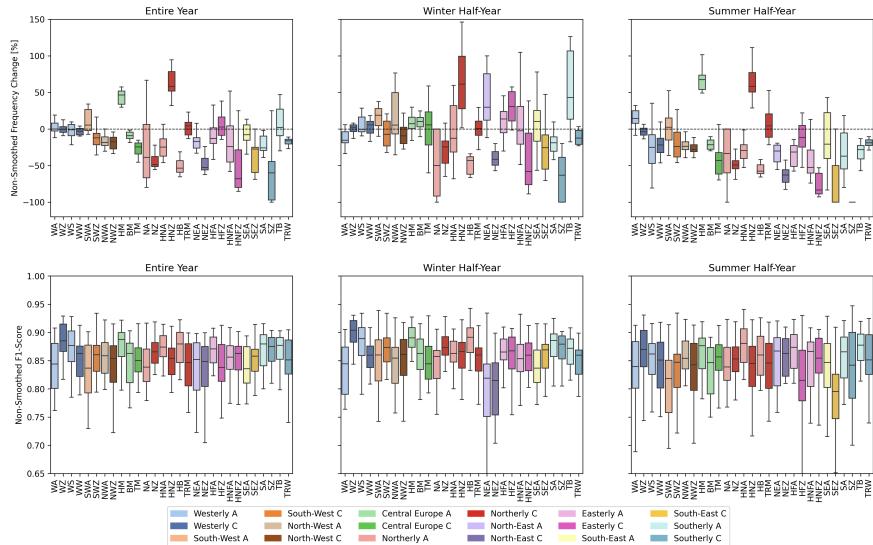


FIGURE 1.5: Boxplots of frequency change and f1 score without transitional smoothing. Upper plots show the change in the relative frequency of occurrence (%) of the HB CTs between the future 2031–2060 and the reference period 1971–2000 for the entire year, the winter half-year (ONDJFM) and the summer half-year (AMJAS). Lower plots illustrate the spread of F1-scores.

Without transition smoothing (Figure 1.5):

1. The boxplots show relatively smaller spreads for most HB CTs.
2. Most relative changes are within $\pm 50\%$.
3. Fewer extreme outliers are observed, and the whiskers are relatively short.
4. The overall signal is dominated by noise, making it hard to detect consistent climate change trends.

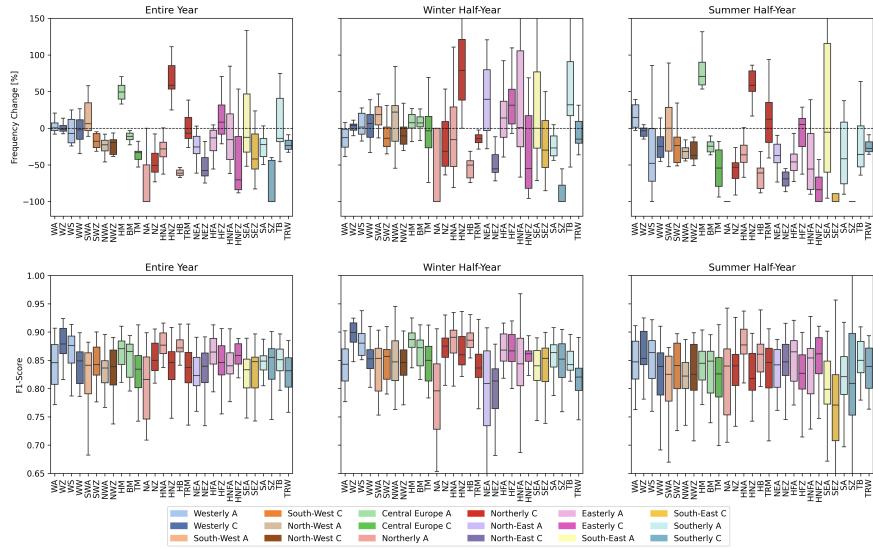


FIGURE 1.6: Boxplots of frequency change and f1 score after transitional smoothing. Upper plots show the change in the relative frequency of occurrence (%) of the HB CTs between the future 2031–2060 and the reference period 1971–2000 for the entire year, the winter half-year (ONDJFM) and the summer half-year (AMJJAS). Lower plots illustrate the spread of F1-scores.

With transition smoothing (Figure 1.6):

1. The boxplots show larger spreads, particularly for rare HB CTs.
2. Relative changes frequently extend beyond $\pm 50\%$, and even approach $\pm 100\%$ in some cases.
3. More extreme outliers appear.
4. F1-score distributions are lower and broader, indicating less stable classification performance.

Overall, transition smoothing, while theoretically beneficial, introduces higher internal variability in our implementation, complicating clear signal identification. Variability differences from the original study stem from methodological factors (ensemble size, smoothing implementation) and underlying climate model characteristics.

Our findings reaffirm the value of a multi-ensemble approach and careful consideration of methodological choices when projecting future circulation trends, underscoring the need for robust ensemble analysis techniques to better discern climate-driven circulation changes from internal variability.

The original study (Appendix Figure 1.13) shows relatively controlled variability, mostly within $\pm 50\%$ (Mittermeier et al., 2022), and fewer extreme outliers, indicating relatively stable and moderate shifts in HB CTs. Our results (Figure 1.5) exhibit larger variability, particularly notable in specific HB CTs such as NA, SEA, HNFZ, and SZ, with frequency changes frequently exceeding $\pm 50\%$, even reaching $\pm 100\%$.

1. Entire Year:

- Original study shows moderate shifts centered around zero, with clear signals for types like WA and SEA (Mittermeier et al., 2022).
- Our implementation shows notably higher variability with clear signals (e.g., strong increase in HNZ, HM and significant decreases in HB and NEZ).

2. Winter Half-Year:

- Original study depicts clear moderate increases for WW and HFA with relatively smaller variability (Mittermeier et al., 2022).

- Our data demonstrate significant changes with pronounced variability and stronger extremes, especially in HNZ and rare CTs like NA and SZ.

3. Summer Half-Year:

- Original data indicate mild variability and clear, moderate increases and decreases in specific types (e.g., decreases in HNFZ, increases in WA) ([Mittermeier et al., 2022](#)).
- Our results again illustrate increased variability with substantial increases (e.g., HNZ, HM) and decreases (e.g., HB, NEZ).

In both cases, the summer half-year has a greater influence on the entire year, despite the winter half-year showing higher variability. Rare CTs such as NA and SZ are observed with significant changes. It would be interesting to analyze how rare conditions evolve, however, due to the unbalanced data, the results are not very reliable.

Key CT Differences:

1. WA (Westerly Anticyclonic):

- Our results show only slight increases, with less pronounced signals for WA, especially compared to significant changes in other types.
- Original study consistently shows WA increasing clearly across all seasons ([Mittermeier et al., 2022](#)).

2. HNZ (Icelandic High Cyclonic):

- In our results, HNZ consistently shows marked increases, particularly pronounced in the summer and winter half-years.
- Original study presents less dramatic decreasing changes for HNZ, reflecting a critical methodological or scenario-driven difference ([Mittermeier et al., 2022](#)).

3. NA and SZ (Northerly Anticyclonic and Southerly Cyclonic):

- Our results exhibit significantly larger variability and extreme reductions in NA, while SZ shows extreme negative changes particularly in summer.
- Original study maintains relatively modest decreasing changes for these types ([Mittermeier et al., 2022](#)).

The differences in projected frequency changes between our results and the original study primarily stem from:

1. Time period difference (2031-2060 vs. 2071-2100).
2. Scenario differences: CanESM2 model (historical) in our study vs. original SMHI-LENS (SSP3-RCP7.0).
3. Model and methodological distinctions: Implementation differences (e.g., transitional smoothing, dataset variations, and resolution).
4. Ensemble size effects: Our smaller ensemble size (12 vs. original 50) could amplify internal variability signals.

These differences underline the sensitivity of CT analysis to methodological choices, highlighting the importance of transparent documentation of scenario assumptions and model details to ensure the accurate interpretation and comparability of future climate studies.

1.6 Conclusion and Outlook

In this study, we successfully implemented a CNN-based deep learning approach to classify large-scale CTs over Europe, building upon the established HB framework. Using high-resolution ERA5 data for model

training, our method achieved an overall accuracy of 53.17% and a macro F1-score of 55.14%, clearly outperforming traditional classification approaches.

The CNN model effectively learned dominant circulation patterns, yet faced challenges classifying rare or inherently complex CTs. These limitations indicate potential issues of class imbalance and inherent labeling uncertainties associated with manual classifications. One potential improvement to the statistical analysis would be reducing the number of subdivisions of CTs. Subdividing CT according to the wind direction of the jet stream, for example, could increase the robustness of the results. Additionally, the original study found that the most frequent misclassifications occurred between pairs of anticyclonic and cyclonic circulation types ([Mittermeier et al., 2022](#)), which could be addressed through refined classification methods. Furthermore, future research should be conducted to quantify human-level errors in labelling to better understand the impacts of manual classification biases.

Besides the primary training session and results shown in this report, we conducted another model training experiment using a narrower range of hyperparameters. The results, which are illustrated in the presentation, demonstrated noticeably different classification performance compared to the setup we introduced here. In particular, I suspect that the size of the fully connected linear layers is the main reason for this difference. Since in earlier experiments with narrower range of hyperparameters, we set the fully connected layer size as fixed 50 dimensions, however, in the latest optimization round, where more extensive hyperparameter tuning were applied, the model consistently favored a dimension size of 128, which leads to high accuracy for each HB CTs. Therefore, the model shows good performance on the current dataset, it is important to note that this does not necessarily guarantee the absence of overfitting.

Due to limited computational resources, we were unable to perform a more thorough investigation to validate these observations or further optimize the model architecture. Future work should focus on systematically assessing the impact of fully connected layer sizes and further regularization strategies to ensure model robustness and generalization.

The analysis of frequency changes in both our implementation and original paper reveals that for most HB CTs, the changes fall within a range of ± 5 days (relatively $\pm 50\%$) ([Appendix Figure 1.12](#); [Figure 1.6](#); [Appendix Figure 1.14](#); [Appendix Figure 1.13](#)). In the original study, the most significant changes in frequency were observed in the WA circulation type ([Appendix Figure 1.13](#)) ([Mittermeier et al., 2022](#)), while in the own-built model, the HM and HNZ circulation types experienced the largest shifts in frequency. Notably, our results differ from previous studies, primarily due to distinct methodological choices such as dataset selection, scenario assumptions, and smoothing techniques. These variations underline the sensitivity of HB CTs analysis and stress the importance of transparent methodological reporting.

When focusing only on the frequency of HB CTs, without considering their duration, it is important to highlight that the duration of a HB CT significantly influences the amplitude of weather extremes. This underscores the need to integrate both frequency and duration for a more comprehensive understanding of weather patterns. Duration of a HB CT impacts the amplitude of a weather extreme.

To further evaluate the uncertainties in frequency changes, it would be beneficial to combine multi-model and single-model ensembles under different forcing scenarios. This approach could help assess the uncertainties arising from various climate models and the assumptions in different forcing scenarios.

For future studies, we recommend quantifying human-level errors in labeling to understand better the impacts of manual classification biases. Additionally, models that explicitly capture temporal continuity, such as Deep Hidden Markov Models ([Yu et al., 2015](#)) or temporal-aware CNN architectures (e.g., ConvLSTM ([Moishin et al., 2021](#))), could enhance prediction accuracy by directly modeling the persistence characteristic inherent in circulation patterns ([Mittermeier et al., 2022](#)).

1.7 Appendix

1.7.1 Statement

The implementation was carried out using Python version 3.9.18.

1.7.2 Figures

Synoptic patterns of the 29 HB CTs using ERA20C (Figure 1.7) ([Mittermeier et al., 2022](#))

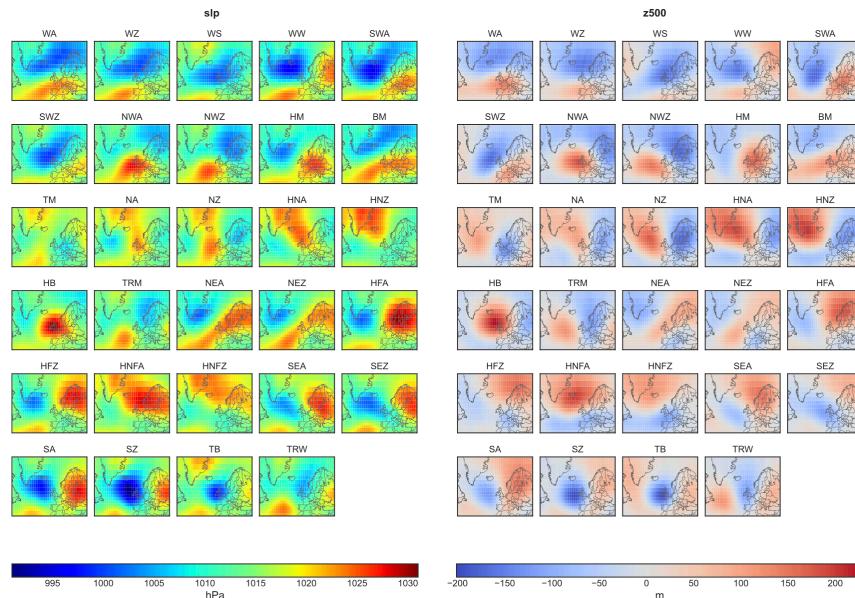


FIGURE 1.7: Synoptic patterns of the 29 HB CTs created using ERA-20C data averaged over the period 1900 -1980, showing SLP on the left and z500 on the right side.

HB CTs (Figure 1.8)

F1 score comparison table (Figure 1.9)

Signature plots of paper - SLP (Figure 1.10) ([Mittermeier et al., 2022](#))

Frequency distribution of paper (Figure 1.11) ([Mittermeier et al., 2022](#))

Boxplot of absolute future change and F1 score of our implementation (Figure 1.12)

Boxplot of future change and F1 score of paper (Figure 1.13) ([Mittermeier et al., 2022](#))

Boxplot of absolute future change and F1 score of paper (Figure 1.14) ([Mittermeier et al., 2022](#))

RMSE table of our implementation - SLP (Figure 1.15)

RMSE table of paper - SLP (Figure 1.16)

Confusion matrix of our implementation (Figure 1.17)

Confusion matrix of paper (Figure 1.18) ([Mittermeier et al., 2022](#))

Full signature plot of our implementation - SLP (Figure 1.19)

Full signature plot of paper - SLP (Figure 1.20) ([Mittermeier et al., 2022](#))

Acronym	Original name (German)	Translated name (English)
WA	Westlage, antizyklonal	Anticyclonic Westerly
WZ	Westlage, zyklonal	Cyclonic Westerly
WS	Südliche Westlage	South-Shifted Westerly
WW	Winkelförmige Westlage	Maritime Westerly (Block E. Europe)
SWA	Südwestlage, antizyklonal	Anticyclonic North-Westerly
SWZ	Südwestlage, zyklonal	Cyclonic South-Westerly
NWA	Nordwestlage, antizyklonal	Anticyclonic North-Westerly
NWZ	Nordwestlage, zyklonal	Cyclonic North-Westerly
HM	Hoch Mitteleuropa	High over Central Europe
BM	Hochdruckbrücke (Rücken) Mitteleuropa	Zonal Ridge across Central Europe
TM	Tief Mitteleuropa	Low (Cut-Off) over Central Europe
NA	Nordlage, antizyklonal	Anticyclonic Northerly
NZ	Nordlage, zyklonal	Cyclonic Northerly
HNA	Hoch Nordmeer-Island, antizyklonal	Icelandic High, Ridge C. Europe
HNZ	Hoch Nordmeer-Island, zyklonal	Icelandic High, Trough C. Europe
HB	Hoch Britische Inseln	High over the British Isles
TRM	Trog Mitteleuropa	Trough over Central Europe
NEA	Nordostlage, antizyklonal	Anticyclonic North-Easterly
NEZ	Nordostlage, zyklonal	Cyclonic North-Easterly
HFA	Hoch Fennoskandien, antizyklonal	Scandinavian High, Ridge C. Europe
HFZ	Hoch Fennoskandien, zyklonal	Scandinavian High, Trough C. Europe
HNFA	Hoch Nordmeer-Fennoskandien, antizykl.	High Scandinavia-Iceland, Ridge C. Europe
HNFZ	Hoch Nordmeer-Fennoskandien, zyklonal	High Scandinavia-Iceland, Trough C. Europe
SEA	Südostlage, antizyklonal	Anticyclonic South-Easterly
SEZ	Südostlage, zyklonal	Cyclonic Southerly
SA	Südlage, antizyklonal	Anticyclonic Southerly
SZ	Südlage, zyklonal	Cyclonic Southerly
TB	Tief Britische Inseln	Low over the British Isles
TRW	Trog Westeuropa	Trough over Western Europe

FIGURE 1.8: HB CTs

	WA	WZ	WS	WW	SWA	SWZ	NWA	NWZ	HM	BM	TM	NA	NZ	HNA	HNZ
Implementation	57.19	70.21	59.03	55.88	51.34	57.18	50.17	53.26	63.70	55.54	52.25	33.61	58.27	60.89	51.54
Paper	44.60	47.08	45.39	37.70	35.36	30.86	38.88	37.07	51.24	47.29	37.23	24.85	44.32	45.57	27.11

	HB	TRM	NEA	NEZ	HFA	HFZ	HNFA	HNFZ	SEA	SEZ	SA	SZ	TB	TRW	Macro
Implementation	62.78	52.30	52.02	51.16	60.57	51.99	55.17	57.32	47.69	44.37	60.27	57.72	59.81	55.87	58.09
Paper	50.99	27.86	41.44	33.12	45.32	24.81	33.35	34.02	38.09	37.93	39.84	38.19	42.11	29.34	38.30

FIGURE 1.9: Macro F1

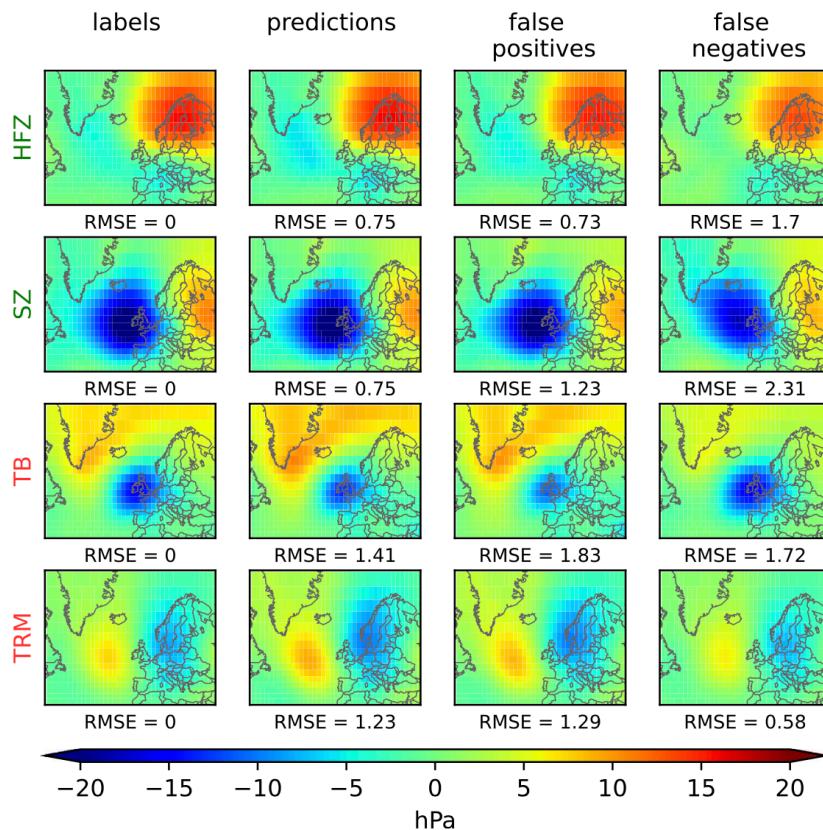


FIGURE 1.10: Signature plots of four selected circulation patterns at slp. Column 1: labels showing the indicated HB CT, Column 2: deep ensemble predictions showing the indicated HB CT, Column 3: signature pattern, when the deep ensemble predicts the indicated HB CT while labels state differently, Column 4: labels stating the indicated HB CT while deep ensemble predicts differently. The RMSE values are calculated by comparing the respective signature plot to the signature plot of the labels (column 1). The four HB CTs are chosen as positive (green) and negative (red) examples.

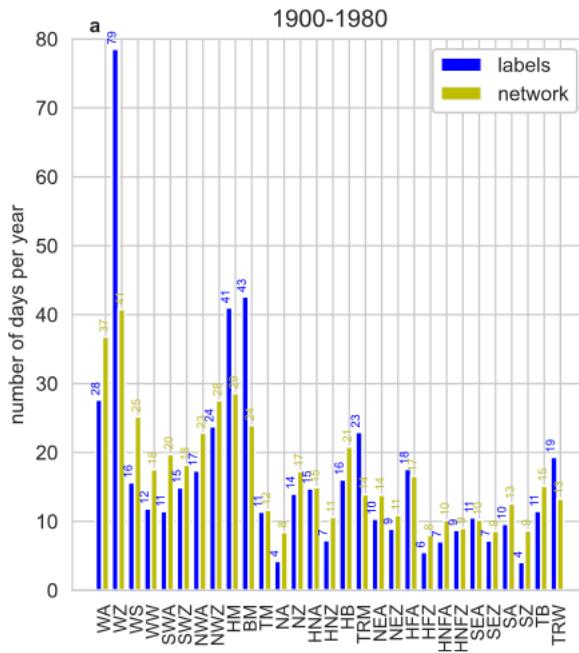


FIGURE 1.11: Frequency distribution of the 29 HB CTs in number of days per year for the training period 1900–1980 (blue) and for predictions of the CNN (yellow).

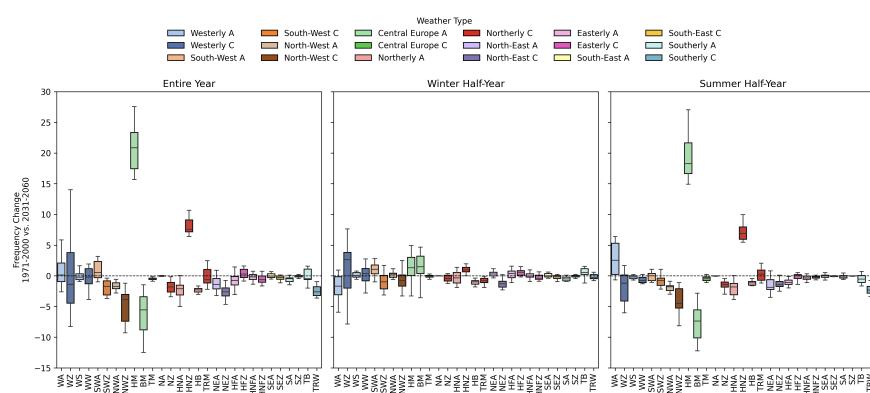


FIGURE 1.12: Boxplots of frequency change and f1 score. Upper plots show the change in the absolute frequency of occurrence (days) of the HB CTs between the future 2031–2060 and the reference period 1971–2000 for the entire year, the winter half-year (ONDJFM) and the summer half-year (AMJJAS). Lower plots illustrate the spread of F1-scores.

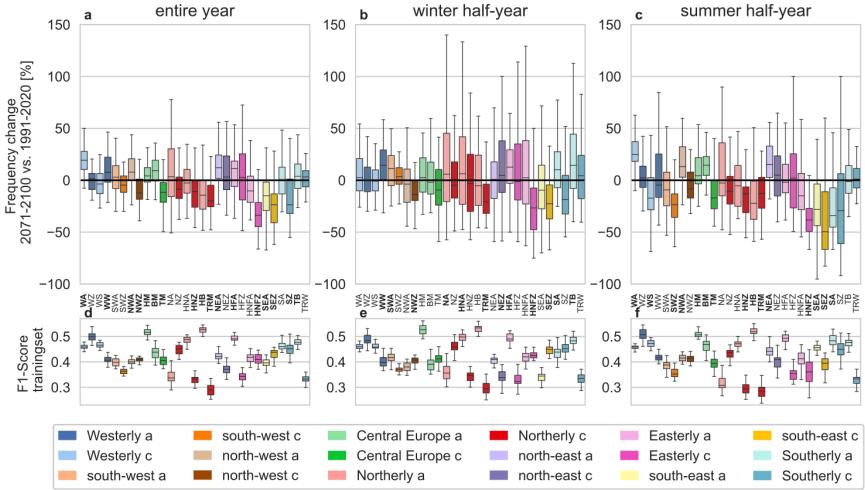


FIGURE 1.13: Boxplots of frequency change and f1 score. Upper plots show the change in the relative frequency of occurrence (%) of the HB CTs between the future 2031–2060 and the reference period 1971–2000 for the entire year, the winter half-year (ONDJFM) and the summer half-year (AMJJAS). Lower plots illustrate the spread of F1-scores.

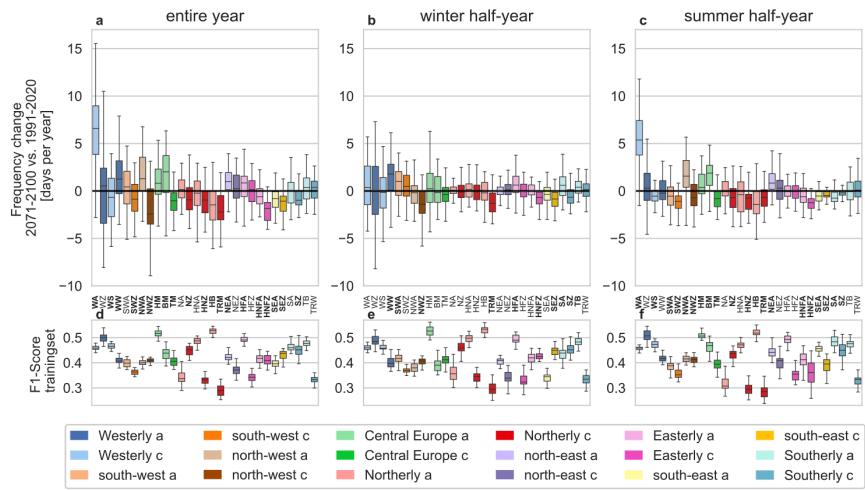


FIGURE 1.14: Boxplots of frequency change and f1 score. Upper plots show the change in the absolute relative frequency of occurrence (days) of the HB CTs between the future 2031–2060 and the reference period 1971–2000 for the entire year, the winter half-year (ONDJFM) and the summer half-year (AMJJAS). Lower plots illustrate the spread of F1-scores.

RMSE	WA	WZ	WS	WW	SWA	SWZ	NWA	NWZ	HM	BM	TM	NA	NZ	HNA	HNZ
Pred	0.39	0.25	0.47	0.54	0.48	0.51	1.17	0.51	0.40	0.52	0.90	2.89	0.54	0.56	0.70
FP	0.88	0.81	1.49	0.91	0.83	1.28	1.34	1.00	0.92	0.93	1.54	5.15	0.96	1.04	1.22
FN	0.50	0.36	0.73	0.55	0.39	0.83	1.01	0.49	0.36	0.34	0.83	0.78	0.66	0.92	0.71
RMSE	HB	TRM	NEA	NEZ	HFA	HFZ	HNFA	HNFZ	SEA	SEZ	SA	SZ	TB	TRW	\emptyset
Pred	0.50	0.57	0.72	1.07	0.69	0.86	1.20	0.68	0.70	1.10	0.71	1.01	0.64	0.32	0.74
FP	1.30	1.04	1.33	1.13	1.55	1.88	2.80	1.36	1.25	1.85	1.64	2.42	1.20	0.92	1.45
FN	0.62	0.33	0.54	1.16	0.50	0.71	1.09	0.88	0.76	0.56	1.04	1.10	0.83	0.46	0.69

FIGURE 1.15: RMSE - slp

RMSE	WA	WZ	WS	WW	SWA	SWZ	NWA	NWZ	HM	BM	TM	NA	NZ	HNA	HNZ
Pred	0.93	0.82	1.00	0.88	0.77	0.82	0.73	1.06	0.78	1.04	0.91	0.87	0.39	0.85	1.29
FP	0.78	1.11	1.21	1.08	1.23	1.00	0.83	1.06	0.80	0.84	0.92	0.92	0.89	1.41	1.41
FN	1.25	0.84	2.10	1.42	1.62	1.24	0.83	1.36	0.87	1.12	1.03	1.50	0.80	1.04	1.30

RMSE	HB	TRM	NEA	NEZ	HFA	HFZ	HNFA	HNFZ	SEA	SEZ	SA	SZ	TB	TRW	Ø
Pred	0.66	1.23	0.58	0.89	0.89	0.75	0.74	0.66	0.99	0.91	1.11	0.75	1.41	1.14	0.89
FP	1.11	1.29	0.76	0.96	1.41	0.73	0.84	0.94	0.98	1.22	1.69	1.23	1.83	1.27	1.09
FN	1.57	0.58	1.00	0.99	1.25	1.70	1.99	1.00	1.31	1.58	1.32	2.31	1.72	0.61	1.28

FIGURE 1.16: RMSE - slp

WA	WZ	WS	WW	SWA	SWZ	NWA	NWZ	HM	BM	TM	NA	NZ	HNA	HNZ	HA	HN	TRM	NEA	NEZ	HFA	HFZ	HNFA	HNFZ	SEA	SEZ	SA	SZ	TB	TRW	Σ	Precision		
WA	350.0	1010.0	0.0	3.0	11.0	3.0	10.0	19.0	29.0	71.0	0.0	1.0	0.0	3.0	0.0	1.0	8.0	2.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	620.0	0.58		
WZ	60.0	1260.0	36.0	28.0	15.0	33.0	2.0	59.0	19.0	35.0	3.0	0.0	8.0	1.0	3.0	0.0	36.0	0.0	2.0	1.0	1.0	0.0	2.0	0.0	0.0	0.0	0.0	0.0	1.0	7.0	23.0	1655.0	0.65
WS	0.0	830.0	183.0	2.0	1.0	16.0	0.0	6.0	2.0	1.0	9.0	0.0	1.0	1.0	3.0	0.0	2.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	341.0	0.66		
WW	2.0	54.0	4.0	209.0	5.0	9.0	0.0	2.0	5.0	16.0	1.0	0.0	1.0	2.0	0.0	0.0	5.0	0.0	4.0	9.0	6.0	1.0	3.0	2.0	5.0	1.0	4.0	8.0	10.0	268.0	0.55		
SWA	15.0	37.0	1.0	5.0	134.0	22.0	0.0	0.0	35.0	13.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	4.0	1.0	0.0	0.0	0.0	0.0	5.0	5.0	283.0	0.56			
SWZ	6.0	74.0	6.0	14.0	28.0	259.0	0.0	5.0	10.0	8.0	0.0	0.0	1.0	0.0	2.0	0.0	2.0	0.0	0.0	0.0	3.0	0.0	2.0	0.0	1.0	3.0	7.0	20.0	438.0	0.6			
NWA	20.0	5.0	0.0	0.0	0.0	74.0	19.0	14.0	23.0	0.0	1.0	3.0	2.0	0.0	15.0	1.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	179.0	0.64			
NWZ	20.0	103.0	2.0	2.0	0.0	2.0	9.0	302.0	8.0	24.0	0.0	0.0	28.0	0.0	1.0	15.0	38.0	3.0	4.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	8.0	570.0	0.54		
HM	22.0	25.0	0.0	3.0	14.0	8.0	3.0	4.0	522.0	90.0	1.0	1.0	0.0	13.0	2.0	17.0	3.0	2.0	5.0	25.0	0.0	1.0	1.0	6.0	0.0	4.0	2.0	4.0	780.0	0.61			
BM	66.0	25.0	2.0	9.0	11.0	9.0	3.0	28.0	66.0	549.0	5.0	0.0	5.0	6.0	1.0	23.0	14.0	10.0	13.0	12.0	2.0	1.0	2.0	2.0	0.0	0.0	0.0	2.0	14.0	880.0	0.5		
TM	0.0	4.0	8.0	3.0	0.0	0.0	0.0	3.0	0.0	4.0	116.0	0.0	4.0	1.0	7.0	0.0	20.0	2.0	14.0	2.0	8.0	1.0	14.0	2.0	5.0	0.0	0.0	6.0	11.0	235.0	0.56		
NA	1.0	4.0	3.0	0.0	0.0	2.0	4.0	14.0	6.0	0.0	20.0	3.0	11.0	5.0	8.0	2.0	1.0	0.0	1.0	2.0	3.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	91.0	0.71		
NZ	1.0	13.0	0.0	0.0	0.0	2.0	42.0	1.0	9.0	7.0	0.0	178.0	7.0	7.0	12.0	23.0	2.0	5.0	0.0	0.0	5.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	7.0	322.0	0.62		
HNA	0.0	3.0	1.0	0.0	0.0	0.0	2.0	1.0	11.0	7.0	2.0	1.0	5.0	165.0	12.0	21.0	0.0	4.0	1.0	2.0	0.0	1.0	16.0	4.0	0.0	0.0	0.0	0.0	1.0	2.0	260.0	0.59	
HNZ	0.0	8.0	6.0	0.0	0.0	5.0	0.0	3.0	8.0	7.0	0.0	20.0	12.0	109.0	3.0	6.0	1.0	3.0	2.0	0.0	3.0	23.0	1.0	0.0	0.0	0.0	1.0	5.0	22.0	352.0	0.56		
HN	3.0	0.0	0.0	0.0	0.0	0.0	7.0	10.0	14.0	32.0	1.0	2.0	5.0	19.0	1.0	237.0	5.0	7.0	5.0	1.0	0.0	2.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	352.0	0.59		
TRM	6.0	61.0	5.0	6.0	5.0	0.0	39.0	6.0	25.0	17.0	0.0	12.0	0.0	9.0	9.0	295.0	1.0	12.0	1.0	1.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	2.0	18.0	486.0	0.53	
NEA	2.0	2.0	0.0	3.0	0.0	0.0	0.0	2.0	9.0	33.0	0.0	1.0	2.0	1.0	1.0	13.0	1.0	10.0	14.0	4.0	25.0	3.0	4.0	5.0	1.0	3.0	0.0	0.0	0.0	3.0	231.0	0.62	
NEZ	0.0	2.0	0.0	5.0	0.0	0.0	1.0	4.0	1.0	37.0	4.0	0.0	3.0	0.0	5.0	16.0	8.0	10.0	132.0	9.0	9.0	1.0	13.0	1.0	3.0	1.0	0.0	0.0	0.0	2.0	269.0	0.53	
HFA	1.0	2.0	0.0	12.0	3.0	2.0	0.0	1.0	25.0	25.0	0.0	0.0	0.0	3.0	0.0	1.0	2.0	8.0	5.0	275.0	8.0	6.0	8.0	8.0	17.0	4.0	3.0	0.0	0.0	0.0	411.0	0.55	
HFZ	0.0	1.0	1.0	7.0	0.0	1.0	0.0	0.0	2.0	7.0	4.0	0.0	0.0	0.0	0.0	0.0	2.0	3.0	6.0	31.0	98.0	4.0	9.0	7.0	7.0	1.0	2.0	0.0	6.0	1.0	199.0	0.55	
HNFA	0.0	1.0	0.0	0.0	0.0	2.0	1.0	0.0	4.0	7.0	0.0	0.0	0.0	8.0	3.0	4.0	0.0	4.0	2.0	22.0	1.0	80.0	14.0	4.0	0.0	3.0	0.0	2.0	1.0	163.0	0.63		
HNFZ	0.0	1.0	7.0	3.0	0.0	1.0	0.0	1.0	5.0	3.0	13.0	1.0	3.0	13.0	10.0	6.0	1.0	2.0	7.0	11.0	7.0	11.0	1.0	184.0	2.0	3.0	1.0	0.0	2.0	3.0	301.0	0.54	
SEA	0.0	1.0	4.0	6.0	1.0	0.0	0.0	26.0	7.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0	33.0	4.0	8.0	8.0	98.0	5.0	17.0	5.0	0.0	1.0	231.0	0.54					
SEZ	0.0	3.0	1.0	17.0	0.0	3.0	0.0	1.0	0.0	7.0	2.0	0.0	0.0	1.0	1.0	0.0	5.0	0.0	5.0	13.0	12.0	1.0	12.0	13.0	65.0	2.0	0.0	5.0	6.0	1.0	175.0	0.55	
SA	0.0	1.0	0.0	9.0	10.0	1.0	0.0	0.0	26.0	7.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	14.0	3.0	0.0	1.0	8.0	4.0	113.0	4.0	2.0	4.0	208.0	0.68			
SZ	0.0	2.0	2.0	11.0	0.0	10.0	0.0	0.0	2.0	1.0	0.0	0.0	0.0	1.0	1.0	0.0	1.0	0.0	0.0	1.0	1.0	0.0	1.0	8.0	5.0	7.0	71.0	6.0	3	134.0	0.63		
TB	0.0	14.0	4.0	11.0	2.0	15.0	0.0	1.0	3.0	4.0	5.0	0.0	0.0	3.0	3.0	0.0	2.0	0.0	3.0	1.0	3.0	0.0	1.0	3.0	1.0	2.0	11.0	154.0	26.0	278.0	0.65		
TRW	9.0	44.0	3.0	12.0	4.0	11.0	0.0	8.0	6.0	30.0	11.0	0.0	3.0	1.0	7.0	0.0	32.0	0.0	3.0	2.0	4.0	0.0	6.0	4.0	4.0	3.0	18.0	269.0	494.0	0.57			
Σ	604.0	1934.0	279.0	380.0	238.0	398.0	116.0	564.0	859.0	1097.0	206.0	28.0	289.0	262.0	194.0	403.0	470.0	165.0	247.0	497.0	178.0	127.0	341.0	180.0	118.0	167.0	112.0	237.0	469.0	11183.0	0.59		
Recall	0.56	0.76	0.54	0.47	0.55	0.41	0.53	0.67	0.62	0.49	0.22	0.55	0.63	0.48	0.67	0.51	0.45	0.49	0.87	0.49	0.49	0.61	0.42	0.37	0.54	0.53	0.55	0.54	0.53				

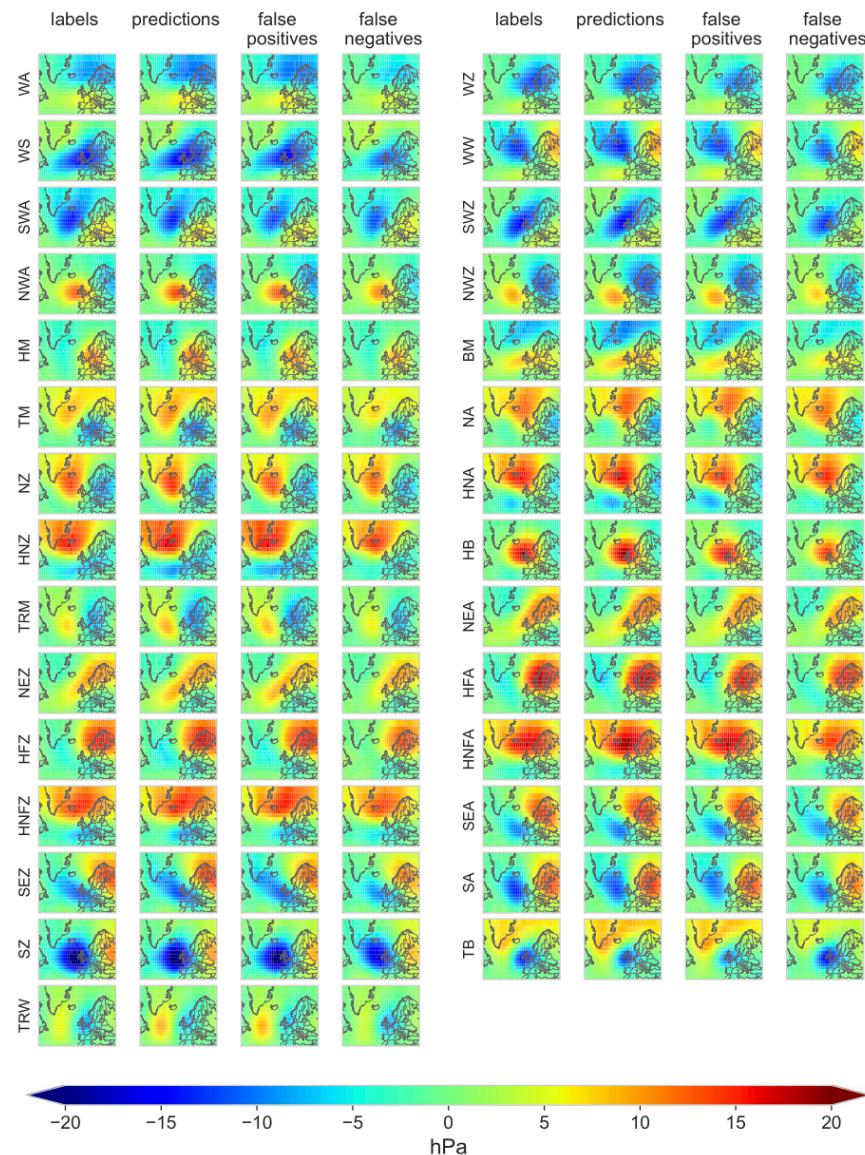
FIGURE 1.17: Confusion matrix

Table A1. Confusion matrix of our proposed smoothed approach, averaged over the test sets of nested cross-validation. Correctly classified classes are highlighted in bold.

	Labels																		Precision												
	WA	WZ	WS	WW	SWA	SWZ	NWA	NWZ	HM	BM	TM	NA	NZ	HNA	HNZ	HB	TRM	NEA	HNFZ	SEA	SEZ	SA	TB	TRW	Σ						
WA	102	62	1	3	4	2	10	8	22	26	0	1	0	0	1	3	1	0	0	0	0	0	1	3	253	0.40					
WZ	13	195	12	5	2	6	1	11	4	2	0	1	2	0	0	6	1	0	0	0	0	0	2	5	272	0.72					
WS	1	51	67	3	0	8	0	4	1	1	4	0	0	0	0	6	0	0	0	1	0	3	0	1	6	5	170	0.40			
WW	4	24	3	42	2	4	1	3	6	6	1	0	0	0	0	3	2	3	2	1	0	0	2	2	6	125	0.33				
SWA	14	18	1	4	34	11	0	0	25	7	0	0	0	0	0	1	1	0	0	0	0	0	0	4	0	1	2	125	0.27		
SWZ	4	36	9	5	8	30	0	1	4	1	0	0	0	1	0	0	2	0	0	0	0	0	0	2	1	4	5	115	0.26		
NWA	14	8	0	1	0	0	58	18	12	20	0	1	4	2	1	12	3	2	2	0	0	0	0	0	0	1	1	159	0.37		
NWZ	10	56	2	2	0	0	0	15	67	2	4	2	1	7	1	0	1	21	1	0	0	0	0	1	0	1	3	198	0.34		
HM	8	4	1	1	4	1	5	1	142	23	0	1	0	2	0	4	0	2	1	7	1	1	0	2	0	3	0	1	216	0.66	
BM	15	5	0	3	1	0	7	2	25	104	0	0	1	1	0	2	3	3	3	0	1	0	1	0	2	0	0	2	187	0.56	
TM	0	8	4	1	0	0	0	3	0	1	34	1	6	0	4	0	10	1	3	1	1	1	5	0	1	0	1	4	6	95	0.36
NA	2	4	0	0	0	0	0	3	1	4	1	0	12	7	12	3	3	1	2	2	1	1	0	0	0	0	0	0	59	0.20	
NZ	2	7	0	0	0	0	0	6	12	1	2	6	4	52	4	5	3	15	2	0	0	0	1	0	0	0	0	1	125	0.42	
HNA	1	4	1	0	0	0	2	0	11	3	1	6	3	54	5	10	1	1	1	2	0	3	2	2	0	0	1	0	116	0.47	
HNZ	0	5	0	0	0	1	1	0	1	1	7	2	2	7	10	17	1	3	0	1	0	1	7	1	0	0	0	1	2	72	0.23
HB	1	1	0	0	0	0	19	4	13	11	0	2	8	0	65	2	5	2	5	2	1	1	1	5	0	1	0	0	0	141	0.46
TRM	3	14	2	1	0	1	1	17	1	4	6	0	9	1	0	35	1	2	0	0	0	0	1	0	0	0	0	1	8	109	0.32
NEA	1	2	0	2	0	0	4	1	10	10	1	1	1	3	0	5	1	43	13	10	2	2	1	1	1	0	0	1	114	0.38	
NEZ	1	2	1	1	0	0	5	1	1	5	4	0	2	1	1	2	5	10	26	2	1	2	1	0	0	1	0	1	79	0.33	
HFA	1	1	0	1	1	0	0	21	5	0	0	3	0	1	1	9	2	63	5	4	1	9	2	4	0	0	1	135	0.47		
HFZ	0	1	1	2	0	0	1	1	3	0	0	0	0	1	3	8	12	12	1	4	2	5	1	1	0	1	1	61	0.20		
HNFZ	0	1	0	0	0	0	0	2	3	1	2	1	1	2	1	1	3	12	1	21	8	3	1	0	1	1	1	79	0.27		
HNFZ	0	3	1	0	0	0	0	0	0	1	1	7	0	0	1	0	2	4	2	6	24	3	2	0	0	1	2	74	0.32		
SEA	0	1	1	1	0	0	0	7	2	0	0	3	0	0	0	1	12	2	2	3	32	9	8	1	1	1	1	87	0.36		
SEZ	0	2	3	4	0	0	1	0	1	2	0	0	0	0	2	1	2	1	4	0	3	6	24	2	3	2	3	68	0.35		
SA	1	2	0	3	4	3	0	0	16	5	0	0	1	0	0	0	0	6	0	0	0	7	1	34	4	1	3	94	0.36		
SZ	0	1	5	4	1	2	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	3	4	8	19	7	5	63	0.30		
TB	2	20	7	4	2	5	0	3	3	1	3	0	0	1	0	0	0	1	0	1	1	2	2	44	12	120	0.37				
TRW	4	18	3	3	1	3	1	4	2	4	2	0	1	0	0	12	0	1	0	1	0	0	1	1	1	7	32	104	0.31		
Σ	205	557	126	97	68	81	140	164	340	255	88	35	111	123	51	115	140	94	79	143	36	49	67	78	59	76	36	89	116	3616	—
Recall	0.50	0.35	0.53	0.43	0.50	0.37	0.42	0.41	0.42	0.41	0.39	0.34	0.47	0.44	0.33	0.57	0.25	0.46	0.33	0.44	0.36	0.40	0.41	0.45	0.53	0.49	0.28	0.41	—		

FIGURE 1.18: Confusion matrix

**FIGURE 1.19:** Signature plot

**FIGURE 1.20:** Signature plot



2

Causal Discovery - Constrain uncertainties in climate projections

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2.1 Abstract

Understanding and evaluating the performance of climate models is essential for improving predictions of future climate variability. Traditional evaluation techniques often fall short in identifying deep-seated structural biases in models. This study introduces a novel process-oriented evaluation approach using causal discovery methods, specifically the PCMCI algorithm, to assess global drought teleconnections. By applying the PCMCI algorithm to SPI-12 precipitation indices from both reanalysis (ERA5) and climate model (CSIRO ACCESS ESM 1.5) datasets, we extract causal networks that reveal underlying climate modes and their interactions. A Varimax-rotated principal component analysis (PCA) was used to reduce data dimensionality, and selected components were analyzed to evaluate the consistency between observed and simulated teleconnections. Results highlight significant differences in causal link structures between the datasets, particularly in ocean-dominated climate modes, suggesting that while PCMCI has limitations in physical interpretation, it holds strong potential for comparative climate model diagnostics. Our findings underscore the importance of integrating causal inference tools into the climate model evaluation toolbox to better constrain model uncertainty and improve future projections.

2.2 Introduction to climate models

Understanding our climate system and how it responds to certain external or internal inputs has always been a key part of scientific research. However, due to the high complexity and nonlinearity of a system as large as the earth that operates on timescales from seconds to decades using experimental methods to understand the earth system is not feasible ([Edwards \(2011\)](#); [Runge et al. \(2019\)](#)). For this reason, models have been used to represent, abstract and simplify the most important drivers of the earth system. The earliest beginnings of conceptual climate models can be traced back to ancient scholars like Ptolemy who distinguished different climatic zones based on the maximal daylength and latitude ([Edwards \(2011\)](#); [Sanderson \(1999\)](#)). Complex mathematical climate models started to emerge in the 19th and 20th centuries through scientists like Milutin Milanković who managed to explain a large part of natural climate variability through periodic cycles of earth's eccentricity, axial tilt and precession ([Edwards \(2011\)](#)). With the rapid technological and scientific breakthroughs since the 20th century the complexity and accuracy of climate models increased tremendously on one hand through better understanding of the underlying principles but also through collection of decades of observational data from satellites and field measurements ([Runge et al., 2019](#)).

Modern global climate models can be divided into two subgroups. The first one are general circulation models (GCMs) which simulate the dynamics of atmosphere (AGCM) and oceans (OGCM) following the laws of fluid

motion, thermodynamics and momentum conservation ([eva \(2013\)](#); [Nowack et al. \(2020\)](#)). In these models atmosphere and oceans are divided into grid cells for which the dynamical equations describing the evolution of variables like temperature or vapor pressure are solved with numerical methods ([cli \(2008\)](#)). An extension to these models are Earth System Models (ESMs) which expand the GCMs by including biogeochemical cycles such as the carbon or nitrogen cycle or atmospheric chemistry ([cli \(2008\)](#)).

Anthropogenic climate change has changed Earth's climate at unprecedented rates. To better understand this change and model future climate pathways the Coupled Model Intercomparison Project (CMIP) was organized by the Working Group on Coupled Modelling (WGCM) in an effort to compare state of the art GCMs and ECMs and tackle important questions regarding climate change ([\(Eyring et al., 2016a\)](#)). Over the last 30 years CMIP went through 6 different Phases, each including more models and addressing a wider range of research questions. With its standardized framework CMIP allows detailed multi-model evaluation which, over the years, revealed model specific systemic differences between individual model groups and observations ([\(Eyring et al., 2019\)](#)). For this reason, rigid climate model evaluation is crucial and has been a rapidly advancing field over the past decades. While more and more routine evaluation metrics and tools like the Earth System Model Evaluation Tool ([\(Eyring et al., 2016b\)](#)) using metrics such as means, variances and trend analysis have been developed, these methods often fail to identify underlying model biases ([\(Nowack et al., 2020\)](#)).

A novel approach to constrain uncertainties in climate models is a process-oriented causal model evaluation (CME) approach introduced by ([Nowack et al., 2020](#)). This method utilizes causal discovery methods developed by ([Runge et al., 2019](#)) (Detecting causal associations in large nonlinear time series datasets, *Sci. Adv.*, 5, eaau4996 2019) to systematically exclude common driver effects and indirect links ([\(Nowack et al., 2020\)](#)), resulting in a network of causal global connections. ([Nowack et al., 2020](#)) applied CME to show that inter-model comparison and comparison to observational data of the resulting causal networks can identify biases in climate models and thus help reducing uncertainties for climate predictions.

In this work we introduce the method proposed by ([Nowack et al., 2020](#)) and show one potential use by applying it to global drought datasets based on reanalysis and global climate model precipitation datasets.

2.3 Process

The use of Causal Networks and especially the PCMCI algorithm, as introduced in the previous section, to help evaluate and better understand large scale climate data timeseries and model outputs has gained some popularity over the last years. One topic where PCMCI algorithms have been applied multiple times are global weather teleconnections. It has been shown multiple times that weather patterns and weather extremes like precipitation and temperature can have significant influence on weather in regions thousands of kilometers away. One example of such a long distance weather teleconnection is the impact of the El Niño Southern Oscillation (ENSO) on North American precipitation and weather patterns ([Ropelewski and Halpert, 1986](#)). An example of a teleconnection between two hydrometeorological extremes is the 2010 floods in Pakistan that were shown to be connected to a heatwave in western Russia ([Lau and Kim, 2012](#)). In both cases the driver behind these teleconnections were atmospheric wave trains (Rossby Waves) that lead to a hydrometeorological connection between the distant regions ([Lau and Kim \(2012\)](#); [Ropelewski and Halpert \(1986\)](#)). Such teleconnections can work in both directions but can also be one-directional as in the ENSO-North-American case (Detecting causal associations in large nonlinear time series datasets, *Sci. Adv.*, 5, eaau4996 2019). The PCMCI algorithm can be used to help discover or confirm suspected teleconnections, however caution when interpreting such results is necessary. One study used PCMCI to show the significant role global teleconnections play in the synchronization of extreme rainfall events ([Boers et al., 2019](#)).

In this work we will apply the PCMCI algorithm to global standard precipitation index (SPI) datasets to assess drought teleconnections. Introduced in 1993, the SPI is a commonly used measure to define droughts by fitting long-term baseline precipitation values to a probability distribution (usually gamma distribution)

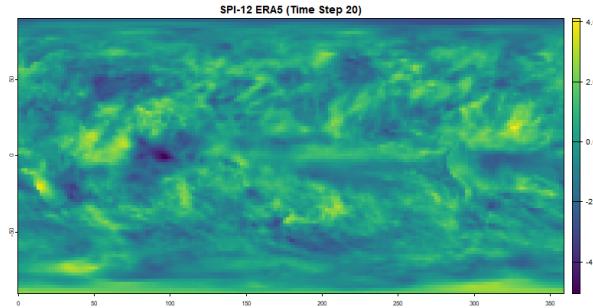


FIGURE 2.1: SPI-12 patterns on ERA5 reanalysis data showing observed drought patterns.

and then transforming the probability values into a standard normal variable with $\mu = 0$ and $\sigma = 1$ (The relationship of drought frequency and duration to time scales 1993). SPI values were calculated based on 12 month cumulative precipitation values (SPI-12) and hence represent long lasting droughts or wet-periods ([Chauhan et al., 2024](#)). In a previous study it was shown that especially oceans play a significant role on modulating global droughts ([Chauhan et al., 2024](#)). Similar to ([Nowack et al., 2020](#)) we applied the PCMCI algorithm to a reanalysis dataset and one climate model dataset. The underlying assumption being that the causal network from the reanalysis dataset represents real world teleconnections. Comparing this to the causal network of the climate model can uncover where the model fails or succeeds at reproducing those teleconnections.

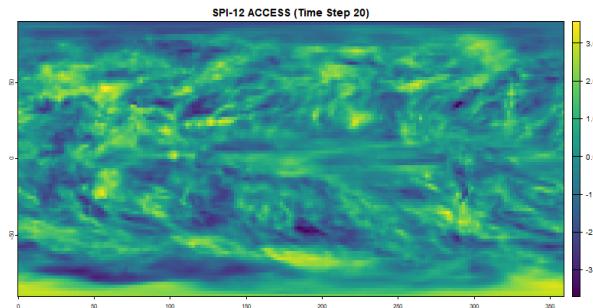


FIGURE 2.2: SPI-12 patterns on ACCESS ESM 1.5 climate model simulation.

2.4 Statistical background

2.4.1 Principal Component Analysis and Varimax Rotation

Our PCA implementation transforms the high-dimensional SPI-12 data into a set of linearly uncorrelated variables that maximize explained variance. This approach effectively identifies coherent drought patterns across geographical regions while substantially reducing computational complexity for subsequent causal analysis. The Kaiser criterion was employed specifically because it provides an objective threshold for component retention by selecting only those components with eigenvalues exceeding unity, which by definition contribute more information than a single original variable.

The Varimax rotation technique redistributes the component loadings to achieve “simple structure,” where each grid cell preferentially loads strongly onto a single component. Mathematically, Varimax maximizes the sum of the variances of squared loadings within each component:

$$V = \sum_k (\sum_i (l_{ik}^2 - \bar{l}_k^2)^2)$$

Where l_{ik} represents the loading of variable i on component k , and \bar{l}_k^2 is the mean of the squared loadings for component k . This optimization produces more spatially distinct patterns compared to unrotated PCA, making it particularly valuable for identifying climatologically meaningful teleconnection patterns. The $|0.4|$ threshold for significant loadings was selected based on conventional practice in climate research, representing a balance between noise reduction and retention of meaningful spatial signals.

2.4.2 PCMCI Algorithm Technical Implementation

The PCMCI algorithm addresses fundamental limitations of traditional correlation analyses by systematically controlling for autocorrelation, common drivers, and indirect causal effects. The PC step implements a condition-selection algorithm where for each time series Y, potential causal parents X are identified through iterative conditional independence testing. The algorithm begins with a full set of potential parents (all time series at all considered lags) and progressively removes links that fail conditional independence tests with increasing conditioning set sizes.

The MCI test then evaluates the conditional independence between each potential cause-effect pair $(X_{t-\tau}, Y_t)$ while controlling for both the past of Y and all other potential common causes, using the formula:

$$\text{MCI} : X_{t-\tau} \perp\!\!\!\perp Y_t | Z_t^Y, Z_{t-\tau}^X$$

Where Z_t^Y represents the parents of Y excluding $X_{t-\tau}$, and $Z_{t-\tau}^X$ represents the parents of $X_{t-\tau}$. This formulation allows PCMCI to distinguish between direct and indirect causal relationships, reducing spurious connections often found in traditional correlation analyses.

In our implementation, we used partial correlation as the conditional independence test with standardized time series data. The time lag parameters were specifically configured with `tau_min = 1` and `tau_max = 5`, allowing us to capture causal relationships occurring between 1 and 5 months. This range is sufficient to detect both relatively rapid atmospheric teleconnections and slower oceanic teleconnection patterns. We deliberately employed a stringent significance threshold with `pc_alpha = 0.0001` to ensure high confidence in the detected causal links, effectively minimizing false positives while accepting a potentially higher rate of false negatives. This conservative approach prioritizes the reliability of identified teleconnections over their quantity, particularly important when comparing model outputs to observational data. The resulting causal networks represent directional relationships between drought patterns, providing insights into the causal mechanisms driving global drought teleconnections and enabling rigorous evaluation of climate model performance in reproducing these teleconnection structures.

2.5 Results

The reanalysis dataset we used is the Copernicus ERA5 post-processed daily statistics on single levels from 1940 to present. CSIRO ACCESS ESM 1.5 data was used as the climate model dataset. For both datasets the years 1950 until (including) 1990 were used. The ERA5 dataset contains daily precipitation values at $0.25^\circ \times 0.25^\circ$ resolution. ACCESS ESM 1.5 data contains daily precipitation rate values at $1.875^\circ \times 1.25^\circ$ resolution. Both datasets were aggregated to total monthly precipitation in mm and regressed to a $2^\circ \times 2^\circ$ resolution using bilinear interpolation.

To reduce the high dimensionality of the datasets we used a Varimax rotated principal component analysis (PCA) to identify large-scale patterns of SPI and thus drought variability. The PCA was applied to monthly SPI-12 values across all grid cells from 1950 to 1990. Thirty principal components (PCs) were retained based on the Kaiser criterion (eigenvalues >1). Varimax rotation was then applied to enhance interpretability by maximizing the variance of squared loadings within each component. The loadings threshold was set at $|0.4|$ to determine significant contributions from individual grid cells. From the rotated components, we selected those with variance contributions exceeding 4% for detailed causal analysis—five for ERA5 and six for ACCESS.

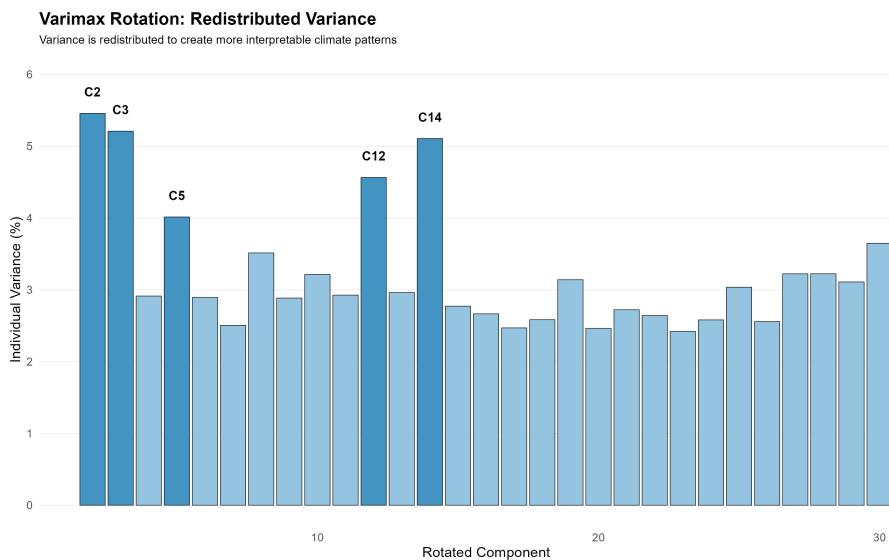


FIGURE 2.3: Varimax rotated principal components for ERA5 dataset.

In the ERA5 reanalysis dataset (Figure 2.3), the variance distribution is characterized by five prominent components (C2, C3, C5, C12, and C14) that individually explain more than 4% of the total variance. Components C2 and C3 show particularly high explanatory power, accounting for approximately 5.4% and 5.2% of the variance respectively. These high-variance components likely represent dominant global drought teleconnection patterns with substantial spatiotemporal coherence. The remaining 25 components each explain approximately 2.5-3.5% of the variance, resulting in a more uniform distribution of explanatory power across the rotated component space. This pattern suggests that after accounting for the major teleconnection modes, the remaining drought variability is distributed across numerous localized or regional patterns of similar importance.

The ACCESS ESM 1.5 climate model (Figure 2.4) exhibits a somewhat different variance structure, with six components (C1, C2, C3, C7, C13, and C21) exceeding the 4% variance threshold. Component C13 displays the highest explanatory power at approximately 5.2%, followed closely by C2 at 5.0%. A notable difference from the ERA5 results is the appearance of C21 as a significant component, suggesting that the climate model

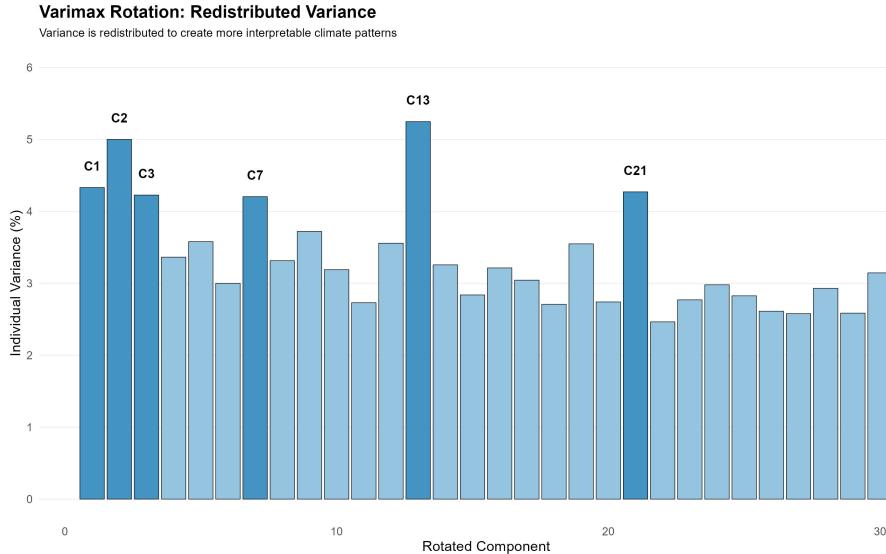


FIGURE 2.4: Varimax rotated principal components for ACCESS dataset.

simulates an important teleconnection pattern that manifests in a higher-order component. This structural difference in the variance distribution between observed and modeled data provides an initial indication that ACCESS ESM 1.5 may represent certain climate processes differently than observed in reanalysis data.

The variance threshold of 4% was selected as an objective criterion to isolate the most influential drought teleconnection patterns while maintaining computational tractability for the subsequent causal analysis. The rotated components exceeding this threshold were selected for detailed causal network analysis using the PCMCI algorithm.

It is important to note, as highlighted by [Hannachi et al. \(2007\)](#), that while Varimax rotation enhances pattern interpretability, the resulting components represent statistical constructs that may not perfectly align with physically coherent climate phenomena. Nevertheless, these rotated components provide valuable insights into the spatial organization of drought variability and offer a robust foundation for subsequent causal analysis of teleconnection structures. The differences in component structure between ERA5 and ACCESS ESM 1.5 datasets suggest potential model biases in representing the spatial organization and relative importance of global drought patterns, a finding that will be further explored through causal network comparison.

2.5.1 Spatial Patterns of Rotated Components

The spatial distributions of the five high-variance ERA5 components (Figure 2.5) reveal distinct geographical footprints that correspond to known ocean-atmosphere coupled systems. Notably, all five components are predominantly oceanic in nature, underscoring the critical role of sea surface temperature patterns in modulating global drought teleconnections.

Component 2 exhibits a clear tropical Pacific signature, with significant loadings concentrated in the central-eastern equatorial Pacific. This pattern strongly resembles the canonical Eastern Pacific El Niño (EP-ENSO) pattern, characterized by maximum sea surface temperature anomalies in the eastern tropical Pacific ([Di Lorenzo et al., 2013](#)). Component 5 displays a broader Pacific footprint extending from the eastern to the central Pacific with notable loadings along the equatorial and subtropical Pacific, suggesting influences from both ENSO and the Pacific Decadal Oscillation (PDO). Component 14 shows a distinctive pattern with significant loadings in both the western-central tropical Pacific and the southeastern Pacific, along with notable signals in the Mediterranean and southeastern Asia, potentially representing a combination of Central Pacific El Niño (CP-ENSO) and Indo-Pacific teleconnection patterns.

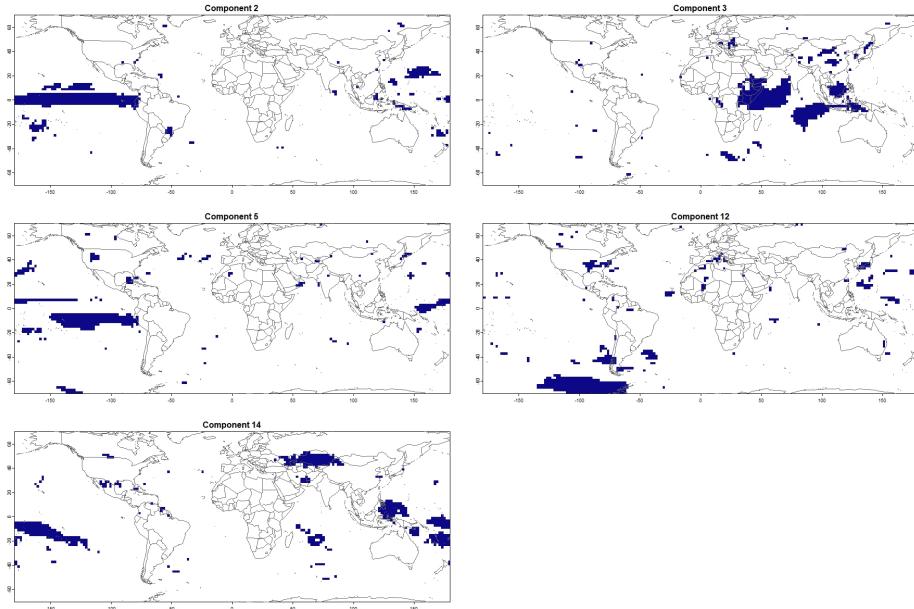


FIGURE 2.5: World map showing the location of the different rotated components from ERA5 dataset

Component 3's spatial distribution extends from the Western Indian Ocean (WIO) across the maritime continent to the Indo-Pacific Warm Pool (IPWP). This pattern likely represents the Indian Ocean Dipole (IOD) in conjunction with IPWP variability, systems known to significantly influence precipitation patterns across the Indo-Pacific region through modulation of the Walker circulation (Zhang and Han, 2020; Newton et al., 2006). Component 12 shows predominant loadings in the Arctic Ocean region, which may partially reflect coordinate projection effects that can amplify variance near the poles in global gridded datasets, as higher latitudes are represented by more grid cells per unit area due to meridional convergence.

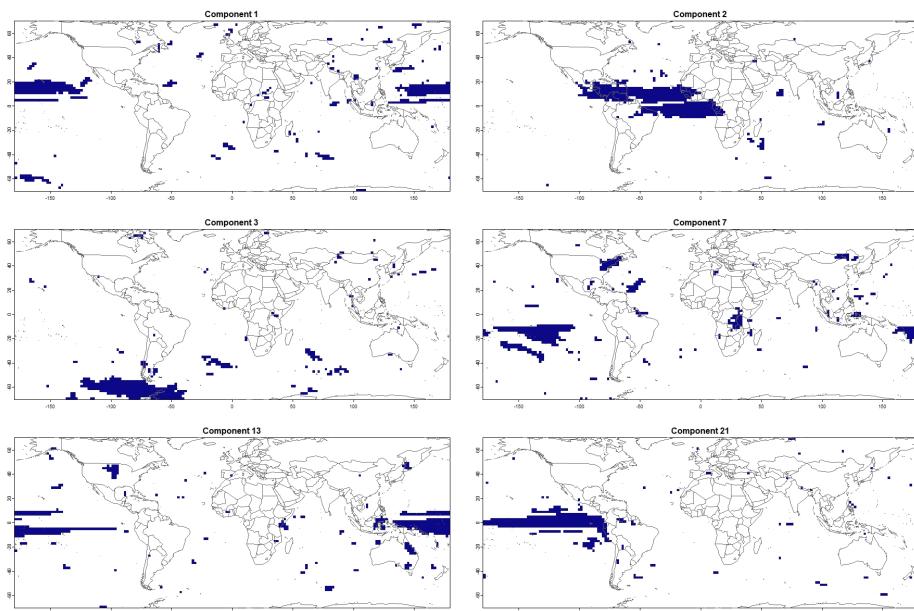


FIGURE 2.6: World map showing the location of the different rotated components from ACCESS dataset

The ACCESS ESM 1.5 dataset (Figure 2.6) reproduces several Pacific Ocean teleconnection patterns similar

to those identified in ERA5, though with notable structural differences. Components 1, 7, and 13 capture various aspects of Pacific variability, with Component 1 showing a strong equatorial Pacific signal comparable to EP-ENSO, Component 7 exhibiting a broader eastern Pacific pattern with extensions into the North Atlantic and Indian Ocean, and Component 13 displaying a pan-Pacific pattern reminiscent of the PDO. Component 21 reveals an intriguing pattern connecting the tropical eastern Pacific with the subtropical North Atlantic, potentially representing an inter-basin teleconnection mechanism.

A critical difference between the datasets emerges in the representation of Indian and Atlantic Ocean teleconnection patterns. While ERA5 identifies a strong WIO-IPWP component (Component 3), ACCESS ESM 1.5 shows no comparable high-variance component in this region. Instead, ACCESS ESM 1.5 exhibits a prominent Atlantic Ocean component (Component 2) that is not evident among the high-variance ERA5 components. This structural difference suggests a potential model bias in representing the relative importance of Indian Ocean versus Atlantic Ocean teleconnection systems, which could significantly impact simulated drought patterns across adjacent continental regions. Component 3 in ACCESS is predominantly concentrated in the Arctic region, similar to Component 12 in ERA5, though with greater extension into northern Eurasia.

Both datasets highlight the dominance of oceanic climate modes in explaining global drought variability, aligning with findings from [Chauhan et al. \(2024\)](#) that oceanic influences play a crucial role in synchronized drought events across multiple regions. The numerous smaller, scattered regions of significant loadings visible in both datasets likely represent mathematical artifacts of the Varimax rotation process rather than physically coherent teleconnections. These secondary signals should be interpreted cautiously as they may not represent robust geophysical connections to the primary climate mode represented by each component.

The differences in spatial patterns between ERA5 and ACCESS ESM 1.5 components provide valuable insights into potential model biases in simulating the spatial structure and relative importance of major climate teleconnection systems.

2.5.2 Causal Network

The PCMCI algorithm was applied to all 30 components from both ERA5 and ACCESS ESM 1.5 datasets, yielding comprehensive causal networks shown in Figures 2.7 and 2.8. Both networks exhibit a dominant pattern of strong autocorrelation, indicated by the self-loops (circular arrows) at each node, consistent with findings from [Nowack et al. \(2020\)](#) in their analysis of sea level pressure datasets. This autocorrelation reflects the inherent memory in climate systems, where drought patterns typically persist over multiple months due to soil moisture feedback mechanisms and the relatively slow evolution of ocean temperature anomalies that drive teleconnection patterns.

Beyond autocorrelation, the full networks reveal complex interconnections between different components, with varying correlation strengths and directionality. The color scale indicates correlation strength and sign, with red representing positive correlations (where an increase in one component leads to a subsequent increase in the connected component) and blue representing negative correlations (where an increase leads to a subsequent decrease). The full networks exhibit a mix of both positive and negative causal links, reflecting the complex feedback mechanisms in global climate teleconnections, where both reinforcing and dampening interactions can occur.

While a comprehensive analysis of all causal links in these full networks is beyond the scope of this study, visual inspection reveals some structural differences between ERA5 and ACCESS ESM 1.5 networks. The ERA5 network appears to have a slightly denser structure of inter-component links compared to ACCESS, particularly in the lower-variance components. This difference suggests that the climate model may not fully capture the complexity of interactions between secondary drought teleconnection patterns, potentially simplifying some of the more subtle causal relationships present in observational data.

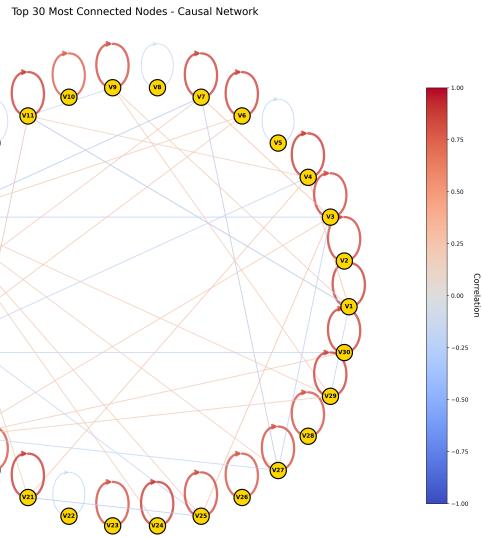


FIGURE 2.7: Causal network of all 30 rotated components from ERA5 dataset showing significant causal links detected by the PCMCI algorithm. Red links indicate positive correlations while blue links represent negative correlations. The prominence of self-loops (autocorrelation) is evident across most nodes.

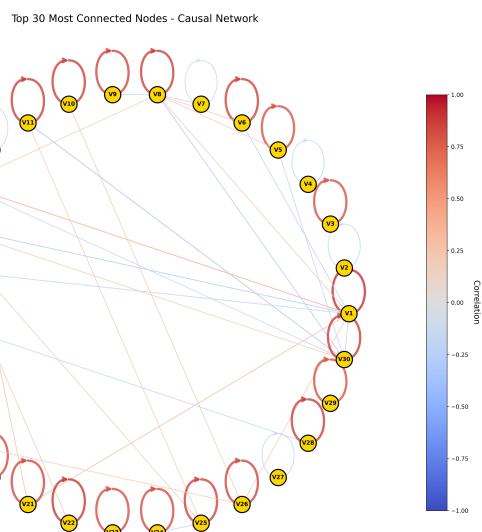


FIGURE 2.8: Causal network of all 30 rotated components from ACCESS ESM 1.5 dataset. Compared to ERA5, the ACCESS model shows a slightly different pattern of inter-component connectivity, though with similarly dominant autocorrelation signals.

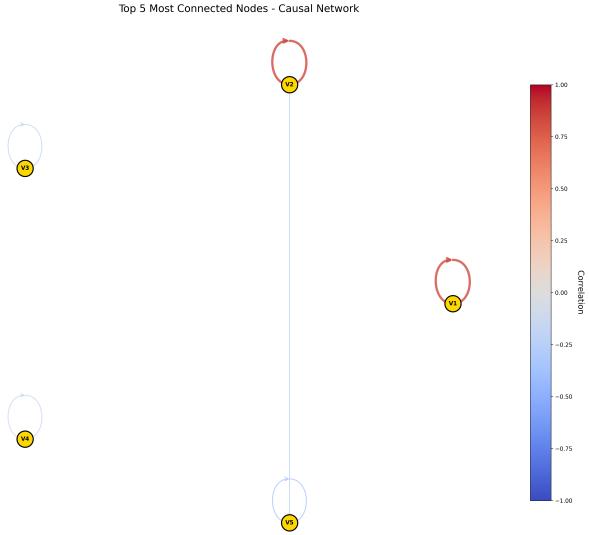


FIGURE 2.9: Causal network of the five selected high-variance components from ERA5 dataset. Node labels correspond to components ($V_1=C_2$, $V_2=C_3$, $V_3=C_5$, $V_4=C_{12}$, $V_5=C_{14}$). Note the single weak negative causal link between the Western Indian Ocean/Indo-Pacific Warm Pool component (V_2) and the Pacific component (V_5).

2.5.3 High-Variance Component Networks

Focusing on the high-variance components (Figures 2.9 and 2.10), we observe notable differences in causal structure between ERA5 and ACCESS ESM 1.5. In the ERA5 network (Figure 2.9), the five selected components ($V_1=C_2$, $V_2=C_3$, $V_3=C_5$, $V_4=C_{12}$, $V_5=C_{14}$) exhibit minimal inter-component causal connectivity, with only one significant causal link detected: a negative correlation from component 3 (V_2 , representing the Western Indian Ocean/Indo-Pacific Warm Pool) to component 14 (V_5 , representing Pacific variability patterns). This negative relationship aligns with known climate dynamics, where warming in the Western Indian Ocean/Indo-Pacific Warm Pool can induce atmospheric wave patterns that influence Pacific circulation, typically manifesting as a dampening effect consistent with the negative correlation observed. This finding corroborates Chauhan et al. (2024), who identified similar teleconnections between these ocean basins in their global drought analysis.

In contrast, the ACCESS ESM 1.5 network (Figure 2.10) exhibits a more interconnected structure among its six high-variance components ($V_1=C_1$, $V_2=C_2$, $V_3=C_3$, $V_4=C_7$, $V_5=C_{13}$, $V_6=C_{21}$). The model simulates a triangular causal structure between three Pacific-dominated components (V_1 , V_5 , and V_6 , corresponding to components 1, 13, and 21), with positive correlations of moderate strength. This interconnected Pacific structure suggests that the climate model simulates stronger intra-basin coupling within the Pacific than is evident in the observational data. Notably, component V_1 (C_1) exerts causal influence on both V_5 (C_{13}) and V_6 (C_{21}), while V_5 and V_6 also share a direct causal link, forming a closed causal triangle that implies potential feedback mechanisms within Pacific climate variability.

The striking difference between these causal structures—ERA5 showing primarily inter-basin teleconnection (Indian-Pacific) versus ACCESS showing stronger intra-basin connections (within Pacific)—reveals a key bias in the climate model's representation of drought teleconnection mechanisms. The model appears to overemphasize Pacific internal dynamics while underrepresenting the crucial teleconnections between the Indian and Pacific Oceans that are evident in observational data. This finding has important implications for the model's ability to simulate drought propagation patterns, particularly for regions influenced by Indian Ocean climate variability.

It is worth noting that while both datasets exhibit strong autocorrelation in their components (self-loops with

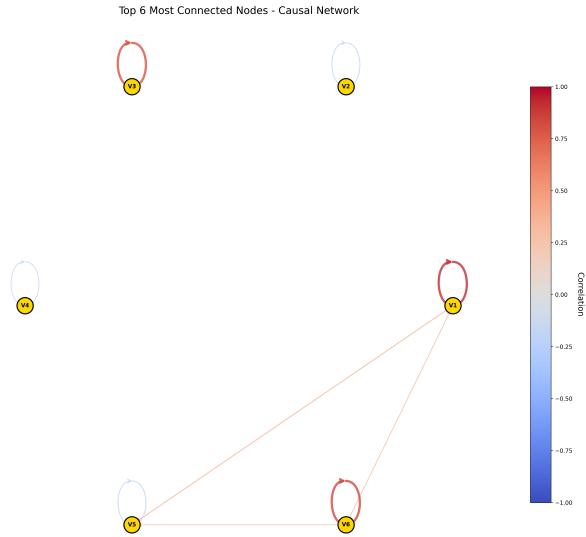


FIGURE 2.10: Causal network of the six selected high-variance components from ACCESS ESM 1.5 dataset. Node labels correspond to components (V1=C1, V2=C2, V3=C3, V4=C7, V5=C13, V6=C21). The model simulates a triangular causal structure between Pacific Ocean components (V1, V5, V6), which differs substantially from the observed ERA5 pattern.

positive correlations), the relative strength of these autocorrelations differs between ERA5 and ACCESS. The model generally simulates stronger autocorrelation in its Pacific components, which may indicate that the model's representation of Pacific climate modes exhibits greater persistence than observed in nature. This bias in temporal dynamics could affect the model's ability to accurately simulate the timing and duration of drought events associated with Pacific variability.

These differences in causal network structure between ERA5 and ACCESS ESM 1.5 highlight the utility of causal discovery methods for climate model evaluation, revealing specific biases in teleconnection representation that might not be apparent from traditional evaluation metrics focused on means, variances, or spatial patterns alone. The detection of these structural differences in causal mechanisms provides valuable insights for targeted model improvement efforts, particularly in the representation of ocean-atmosphere coupling processes that drive global drought teleconnections.

2.6 Limitations

This study has explored the application of the PCMCI algorithm as a causal discovery method for understanding teleconnection structures in global drought patterns. While our results demonstrate the potential of this approach, they also highlight several important methodological challenges that must be addressed for effective implementation in climate science applications.

The application of causal discovery methods to high-dimensional climate datasets necessitates dimensionality reduction techniques such as the Varimax-rotated PCA employed in this study. However, these statistical approaches do not inherently guarantee physically meaningful representations of the climate system. The rotated components, while statistically optimized for variance explanation and interpretability, may combine or separate physical processes in ways that do not align with actual climate dynamics. This creates a fundamental dependency on expert knowledge to either validate the physical relevance of statistically derived

components or to pre-select regions of interest based on established climate science understanding ([Nowack et al., 2020](#)).

Similarly, the causal links identified through PCMCI analysis require careful interpretation within the context of known atmospheric and oceanic processes. A statistically significant causal relationship between two components does not automatically translate to a well-understood physical mechanism. The blue link between Western Indian Ocean/Indo-Pacific Warm Pool and Pacific components in our ERA5 analysis, while statistically robust, requires corroboration from atmospheric dynamics theory to establish its physical validity. This interpretive requirement limits the explanatory power of causal networks as standalone discovery tools, particularly when applied to complex, multi-scale phenomena like global drought teleconnections.

The stringent statistical threshold (`pc_alpha = 0.0001`) applied in our implementation further constrains the detection of weaker but potentially important causal connections, as evidenced by the sparse causal network in our ERA5 analysis. While this conservative approach minimizes false positives, it may suppress the identification of emerging or secondary teleconnection pathways but nevertheless contribute significantly to drought propagation dynamics.

2.7 Conclusions

Despite these limitations as an explanatory method, our findings strongly support the value of causal discovery approaches for comparative model evaluation purposes. The notable differences between ERA5 and ACCESS ESM 1.5 causal structures—particularly the contrasting inter-basin versus intra-basin teleconnection patterns—reveal specific model biases that might remain undetected through traditional evaluation metrics focused on means, variances, or spatial patterns.

As demonstrated by Nowack et al. (2020) and corroborated by our results, causal networks provide a process-oriented diagnostic framework that can identify where models succeed or fail in reproducing the underlying causal mechanisms that drive climate variability. This capability becomes especially valuable when evaluating climate projections across multiple models, where observational validation is not possible. In such contexts, a model's ability to reproduce known causal structures in historical simulations may serve as an important indicator of its reliability in projecting future climate states.

Furthermore, the structural differences detected in our analysis suggest specific areas for targeted model improvement, particularly in the representation of Indian Ocean-Pacific Ocean teleconnections that appear underemphasized in the ACCESS ESM 1.5 model. Such diagnostic insights can guide focused development efforts to enhance the physical representation of key teleconnection mechanisms in climate models.

3

Cold Extremes

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Supervisor: Henri Funk

3.1 Abstract

Extreme cold events have long been associated with severe societal impacts on energy systems, infrastructure, and public health. Therefore, it remains important to explore the potential for such events to occur in the future and develop appropriate measures in advance. In the context of global warming, even cold winters in Central Europe have been affected by rising temperatures. In this study, we investigated whether extremely cold winters—such as the coldest winter in Germany in 1963—could still occur under a warming climate.

We first applied a dynamical adjustment approach combined with elastic net regression to confirm that atmospheric circulation was the main driver of temperature anomalies. This method also captured a decreasing tendency in the frequency and magnitude of cold extremes under global warming conditions. Furthermore, by examining the most extreme cold storylines from the supplementary boosted data provided by ([Sippel et al. \(2024\)](#)), we found that extremely cold winters—such as the event observed in 1963—remain physically plausible despite a warming climate.

3.2 Introduction and Geographic Background

Extreme cold events, or cold waves, are periods of unusually low temperatures that can severely impact society, leading to increased mortality, energy crises, and disruptions to infrastructure, transportation, and agriculture ([Pinto et al. \(2024\)](#)). Europe has experienced several significant cold waves in recent history, including February 2012 in Eastern and Central Europe ([Planchon et al. \(2015\)](#)), January 2017 in Southeastern Europe ([Anagnostopoulou et al. \(2017\)](#)), March 2018 across Northern and Western Europe ([Karpechko et al. \(2018\)](#)), and the winter of 2023, which was exacerbated by energy shortages linked to the Russian-Ukrainian war ([Quesada et al. \(2023\)](#)). Despite long-term warming trends, cold waves remain a major concern due to their unpredictable nature and severe socio-economic consequences.

Cold waves typically arise from persistent atmospheric circulation patterns that direct the cold Arctic or Eurasian air into Europe ([Quesada et al. \(2023\)](#)). Key mechanisms include Scandinavian blocking, sudden stratospheric warming events, North Atlantic sea surface temperature anomalies, and snow-albedo feedbacks, which can amplify and prolong cold conditions. One of the most extreme examples is the winter of 1962/1963, the coldest on record in many Central European countries (([Eichler, 1970](#); [Sippel et al., 2024](#))). This winter was characterized by prolonged high-pressure blocking over Northwestern Europe, which diverted the usual westerly flow and allowed persistent easterly winds to bring frigid air into the continent ([Loikith and Neelin \(2019\)](#)). Extensive snow cover reinforced the cold through high albedo effects, leading to the freezing of major

European rivers and lakes, including the Rhine, Rhône, IJsselmeer, and large parts of the Baltic Sea ([Groisman et al. \(1994\)](#)). The resulting extreme conditions had severe impacts on human health, infrastructure, and energy systems, highlighting the risks posed by such events even in a warming climate ([Eichler \(1970\)](#)).

This study builds on the work of ([Sippel et al. \(2024\)](#)), who investigated the question: “Could an extremely cold central European winter such as 1963 happen again despite climate change?” Their research addressed two key questions: (1) If a winter atmospheric circulation similar to 1963 were to re-occur in present-day climate, what would be the intensity in terms of cold temperatures? and (2) Is a winter as cold as 1963 or colder still possible in Central Europe today? The present study focuses in greater detail on the second question, assessing the potential for such an extremely cold winter in the current climate and its implications for future extreme events.

3.3 Data processing

In order to investigate potential worst-case cold winter conditions in Germany with a particular focus on Bavaria, we utilized the ERA5 reanalysis dataset ([Hersbach et al. \(2020\)](#)) to analyze temperature anomalies during the winter months, specifically from December to February (DJF). To detect anomalies, we applied a 90-day moving average to remove the seasonal cycle, using 1981–2010 as the reference period. Seasonal temperature anomalies were calculated from daily anomalies.

3.4 Methods

3.4.1 Dynamical Adjustment using Elastic Net Regression

Dynamical adjustment is a technique in climate science, that aims to estimate the influence of atmospheric circulation on a target surface climate variable, such as surface air temperature (([Wallace et al., 1995](#); [Smoliak et al., 2015](#); [Deser et al., 2016](#))). Here, we first apply dynamical adjustment to explore the influence of circulation patterns on temperature anomalies and to better understand the results obtained from other methods. Formally, the temperature anomaly at time t , denoted $T(t)$, is expressed as:

$$T(t) = T_{\text{circ}}(t) + T_{\text{resid}}(t),$$

where:

1. **Circulation-induced component** ($T_{\text{circ}}(t)$) : Represents the part of the temperature anomaly that is driven by large-scale atmospheric circulation patterns.
2. **Residual component** ($T_{\text{res}}(t)$) : Captures thermodynamical effects, including externally forced warming and other unknown influences not explained by circulation patterns.

Atmospheric circulation is typically difficult to measure directly, as it is not a single, easily defined quantity. Instead, it influences observable variables such as sea level pressure (SLP) and geopotential height, which are commonly used as proxies for large-scale circulation (([Smoliak et al., 2015](#); [Sippel et al., 2019](#))). These variables capture essential aspects of circulation patterns, including the strength and position of high- and low-pressure systems, the configuration of jet streams, and the occurrence of blocking events. To extract the circulation-induced component, we use sea level pressure (SLP) patterns as a proxy for atmospheric circulation. By applying statistical regression techniques, we estimate the part of temperature variability

that these circulation patterns can explain. However, this method assumes a linear separation between circulation and thermodynamical effects. In reality, climate processes can be more complex, making this a limitation of the approach.

In our study, we pursue two different approaches to dynamical adjustment. Both methods aim to estimate the circulation-induced component of daily mean winter temperature over our study region Bavaria, using a regularized linear regression technique, called “elastic net regression” ([Zou and Hastie \(2005\)](#)). The first approach is based on the ERA5 reanalysis dataset, where an elastic net regression model is trained using sea level pressure (SLP) grid cells as predictors.

We also use a second dynamical adjustment approach, in which the regression model is trained on the CESM2-LE, using the same predictors as in the ERA5-based model. We subtract the domain-average mean trend of geopotential height patterns to account for the long-term column expansion due to warming ([Sippel et al. \(2024\)](#)), which allows for a greater focus on the interannual variability of atmospheric circulation and its impact on temperature. The resulting regression model, trained entirely on CESM2-LE, is independent of the observational data and is subsequently applied to the ERA5 dataset for comparison.

3.4.1.1 Elastic Net Regression

The model estimates the coefficient vector β by minimizing the following penalized least squares objective function:

$$\hat{\beta} = \arg \min_{\beta} \{ \|y - X\beta\|_2^2 + \lambda [\alpha \|\beta\|_1 + (1 - \alpha) \|\beta\|_2^2] \}$$

where:

- y denotes the vector of surface temperature anomalies,
- X represents the matrix of circulation-related predictors (e.g., gridded SLP values),
- $\lambda \geq 0$ is a tuning parameter controlling the overall penalty strength,
- $\alpha \in [0, 1]$ determines the balance between the L1 and L2 penalties.

This formulation combines two regularization methods: Lasso (L1) and Ridge (L2) regression. The L1 penalty results in sparsity by shrinking specific coefficients to zero exactly, which can be interpreted as making variable selection by including only the most important predictors in the model. The L2 penalty shrinks the coefficients more evenly and stabilizes the model if predictors are highly correlated. By adjusting the mixing parameter α , Elastic net blends the advantages of both methods, enabling variable selection while maintaining model stability and predictive accuracy in multicollinearity. These properties make elastic nets particularly suitable for modeling temperature responses to spatially structured and interdependent circulation fields.

3.4.2 Ensemble Boosting

Cold extremes pose significant challenges in climate science due to their substantial socio-economic impacts. Traditional climate models struggle to capture the rare and intense nature of these events, necessitating advanced methodologies such as ensemble boosting. Hence, ([Sippel et al. \(2024\)](#)) explored the principles of ensemble boosting and its application to evaluate whether a worst-case cold winter such as 1963 is still possible. They focused on a 30-member CESM2 initial condition large ensemble (CESM2-ETH) from 2005 to 2035 to generate physically plausible worst-case scenarios of extremely cold winters.

Ensemble boosting is a technique designed to enhance the representation of extreme weather and climate events in model simulations. The core concept involves perturbing an initial state within a climate model,

allowing different yet physically consistent realizations of an extreme event. By systematically re-initializing the model with minuscule perturbations, it becomes possible to explore the tail behavior of the event distribution. In the context of climate modeling, boosting follows a two-step approach:

1. **First-order boosting** – Re-initialization occurs approximately 5-20 days before an identified extreme event using a round-off perturbation. This yields multiple ensemble members, each evolving uniquely but within the constraints of atmospheric dynamics.
2. **Second-order boosting** – After identifying the coldest simulations from the first-order boosted ensemble, additional perturbations are applied to these extreme cases, further refining the representation of worst-case scenarios.

This approach enables a more comprehensive understanding of potential extreme cold events by expanding the dataset of plausible realizations beyond those found in standard climate model ensembles, such as single model initial-condition large ensemble (SMILEs).

3.4.2.1 Data and Methodology of Ensemble Boosting

In the study of ([Sippel et al. \(2024\)](#)), the CESM2-ETH large ensemble, spanning 900 winter seasons (December-January-February, DJF) from 2005 to 2035, serves as the foundational dataset. This dataset follows the CMIP6 historical forcing (2005-2014) and the SSP3-7.0 scenario (2015-2035). Each of the 30 ensemble members originates from a transient historical simulation with a round-off perturbation in atmospheric initial conditions. To analyze extreme cold events, a boosting methodology was applied:

- **First-order boosting:** The coldest December during the 2020s in the CESM2-ETH ensemble was identified. This simulation was then perturbed and re-initialized for each day from December 1-15, generating 50 ensemble members per day. This resulted in a total of 750 simulations, capturing a well-constrained representation of early winter cold conditions.
- **Second-order boosting:** To further explore extreme cold persistence into January, the two coldest simulations from the first-order boosted set were selected. These were subsequently re-initialized daily from January 1-15, with 50 ensemble members per day, leading to 1500 additional simulations.

The perturbation methodology maintained physical consistency by applying small modifications to the specific humidity field (q) at each grid point, with a magnitude of 10^{-13} . These perturbations ensured mass, energy, and momentum conservation up to the precision of a round-off error. The coupled model was then run for 60 days, with ensemble spread remaining small for the first 4-5 days before diverging significantly.

Instead of replicating the exact methods used by ([Sippel et al. \(2024\)](#)), this study focuses on leveraging the provided supplementary boosted data by ([Sippel et al. \(2024\)](#)) to examine the most extreme cold storylines. Specifically, the study analyzed: (i) The three coldest storylines (minimum temperature and average temperature) from the BSSP370cmip6.0480013.zip dataset. (ii) The three coldest storylines (minimum temperature and average temperature) from the BSSP370cmip6.0230013.zip dataset.

These datasets consist of (i) first-order boosting simulations originating from ensemble member 13 of CESM2-ETH, initialized on December 6, 2022, and December 15, 2022, as well as second-order boosting simulations branching off from specific first-order boosted members ensemble member 23 on December 6, 2022, and ensemble member 48 on December 15, 2022). By analyzing these datasets, this study aims to answer the research question whether winters such as 1963 are still possible in today's climate.

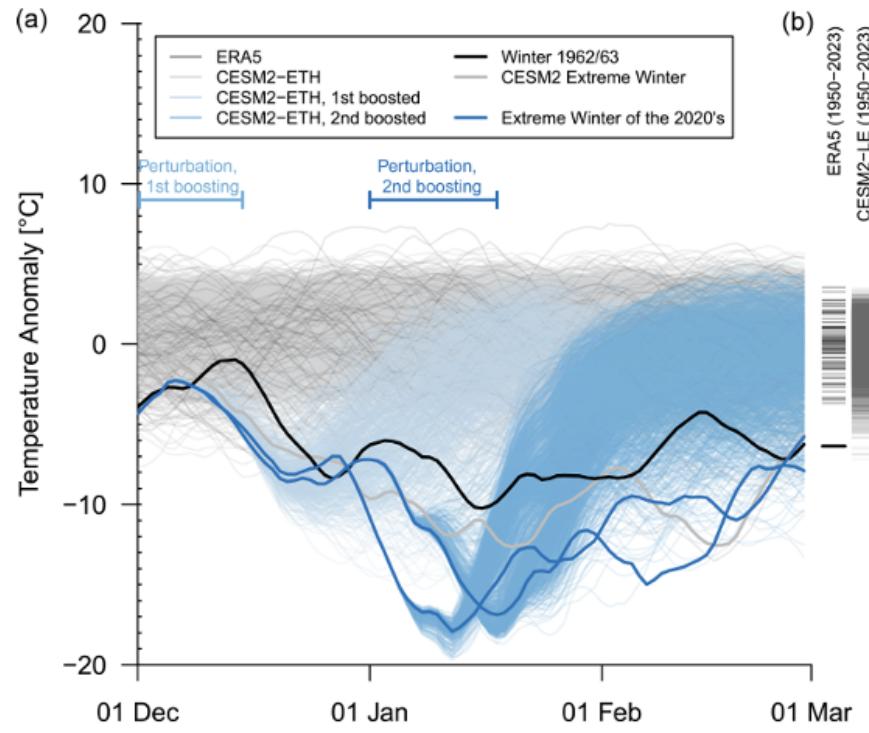


FIGURE 3.1: This figure (a) provides an illustrative example of model boosting, adapted from Sippel et al. (2024)

3.5 Results and Discussion

Here, the key findings derived from the applied methods are presented, followed by a reflection on their scientific implications and a discussion of the methodological limitations and uncertainties involved in the analysis.

3.5.1 Results from Dynamical Adjustment

The circulation-induced component of temperature variability is clearly separated from the residual component, which is not explained by circulation and likely reflects thermodynamical effects (Fig.3.2). The residual time series shows a consistent upward trend (Fig.3.2 bottom), indicating that thermodynamical warming plays a significant role in addition to circulation changes. Besides, the circulation-induced variability shows a strong Pearson correlation of $R = 0.8$ with the observed, detrended DJF temperature anomalies over Bavaria (Fig.3.2 top), thus supporting the conclusion that circulation is the main driver of inter-annual winter temperature variability as suggested by the reference study. ([Sippel et al. \(2024\)](#))

- **Top:** 1951–2024 winter (DJF) temperature anomalies and the contribution of atmospheric circulation (blue line)
- **Bottom:** residual temperature anomaly time series when atmospheric circulation contributions are removed and the trend of this “circulation conditional” residual.

The dark blue line (adjusted using ERA5) shows a clear upward trend, suggesting a decrease in the frequency of cold spells (Fig.3.2 top). This indicates that, in addition to thermodynamical effects, changes in

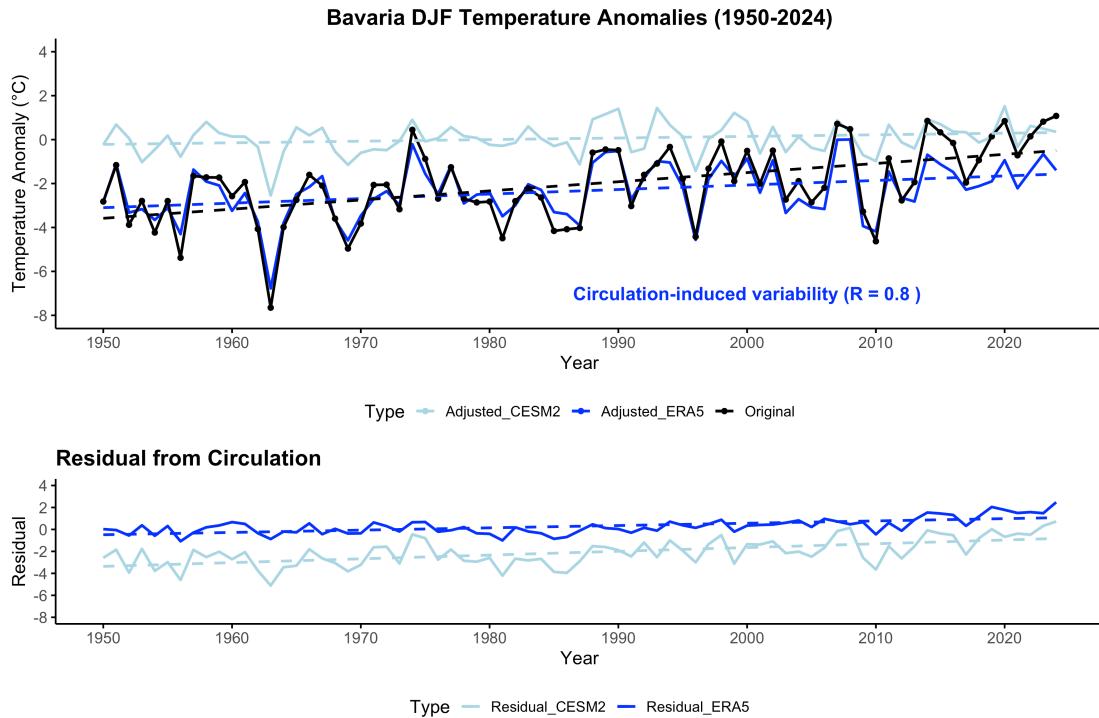


FIGURE 3.2: Winter temperature anomaly time series over Bavaria and long-term trends, dashed lines show linear trends in the original time series (black) and the circulation-induced and residual component (blue).

atmospheric circulation have also contributed to winter warming over Bavaria. However, the CESM2-based light blue line remains relatively flat, showing little to no evidence of strong externally forced changes and suggesting that the future of regional circulation changes under external forcing remains highly uncertain.

This raises a critical uncertainty, as discussed in (Sippel et al. (2024)): whether the circulation trend observed in recent decades represents an externally forced signal or just natural variability. If the trend is indeed forced but not captured by the models, extreme cold winters like 1963 may become less likely in the future. Conversely, if the observed trend is primarily due to natural variability, it could reverse, and similar cold extremes may occur again. This uncertainty remains a key challenge in climate modeling, and understanding it better is crucial for improving future predictions.

3.5.2 Results from Ensemble Boosting

The three coldest storylines, in terms of both minimum temperature and average temperature, were analyzed for ensemble members 23 and 48 from the first-order boosted simulations.

For ensemble member 23, the lowest recorded temperatures in the second-order boosted simulations reached a minimum temperature of -26.3°C , while the average temperature over the winter period from January to March was -12.1°C . Figure 3.3 illustrates the three coldest minimum temperature storylines in the left panel and the three coldest average temperature storylines in the right panel. The ensemble members associated with these extreme conditions—ens010, ens040, and ens042 for minimum temperature, and ens037, ens039, and ens047 for average temperature—exhibit pronounced cold events, highlighting the capacity of the boosting technique to explore the statistical tail of extreme winter conditions.

For ensemble member 48, the coldest recorded temperatures in the second-order boosted simulations included a minimum temperature of -25.9°C , which consistently occurred at the start of the initialization, and an

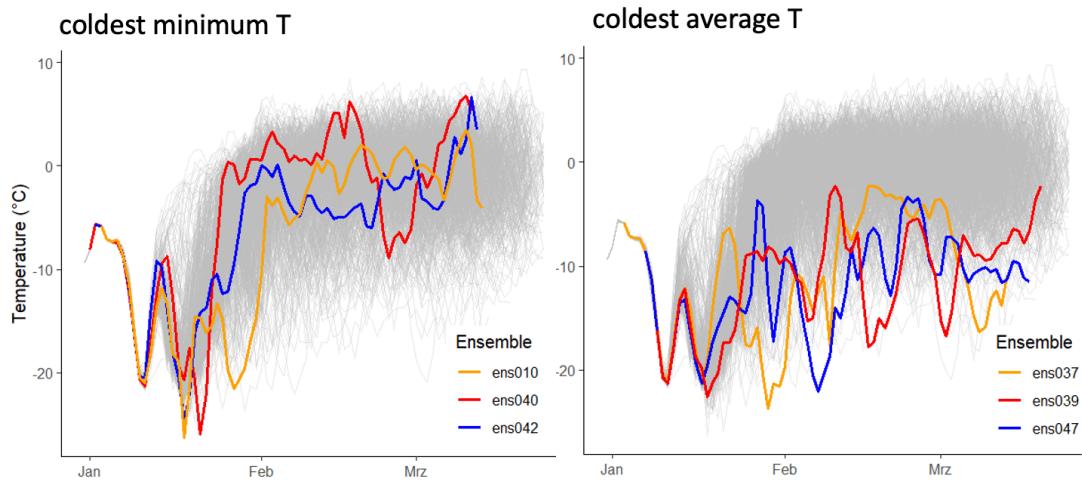


FIGURE 3.3: The three coldest storylines (based on minimum and average temperature) are derived from ensemble member 23. The data are extracted from the file BSSP370cmip6.0230013.zip, provided by Sippel et al. (2024)

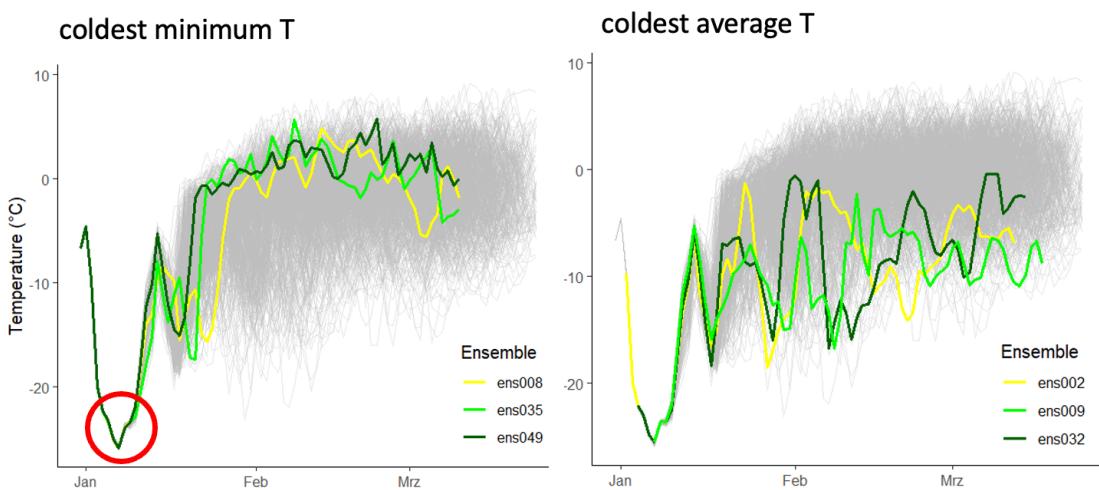


FIGURE 3.4: The three coldest storylines (based on minimum and average temperature) are derived from ensemble member 48. The data are extracted from the file BSSP370cmip6.0480013.zip, provided by Sippel et al. (2024)

average temperature of -10.2°C over the winter period from January to March (Fig.3.4). The fact that the lowest minimum temperatures always appeared at the beginning of the initialization phase suggests potential influences of edge conditions, temporal dependencies, or initialization bias. To address these factors, early January was excluded from the analysis. When focusing on mid-to-late winter, the lowest minimum temperature was observed in simulation ens037, reaching -23.1°C .

The ensemble boosting results demonstrate that extremely cold temperatures, such as in the winter of 1962/1963 or colder, can still be reached today under an SSP3-7.0 scenario. This contrasts with the findings by (Quesada et al. (2023)), who observed a general decline in the frequency and severity of cold events across Europe. However, the presented results align with other studies suggesting that under certain conditions—such as a weakening Atlantic Meridional Overturning Circulation (AMOC)—cold extreme intensities may increase in the future (Meccia et al. (2023)). This apparent contradiction illustrates the complex and multifaceted nature of cold event dynamics. It also reflects the low seasonal predictability identified in recent research, which attributes this uncertainty to chaotic atmospheric forcings, such as variability in westerly winds (Kautz et al. (2021)). The finding that extreme cold events remain possible is further supported by (Brunner et al. (2018)), who showed that approximately 70% of central European cold extremes coincide with atmospheric blocking between 60°W and 30°E —highlighting the continued relevance of large-scale weather patterns as a dominant driver.

Last, there are important limitations to consider. The distribution and frequency of “boosted” cold events are sensitive to the specific events selected for enhancement, which may influence the representativeness of the results. Additionally, re-initializing the model with different atmospheric conditions could potentially generate even colder outcomes, indicating that the full range of extreme winter possibilities might not be fully captured.

4

Flood Frequency Analysis

Author: Hannes Grün, Robin Schüttpelz

Supervisor: Henri Funk

Suggested degree: Master

Abstract

Floods are among the most impactful natural hazards in Bavaria, causing significant economic and ecological damages. Traditional univariate flood frequency analyses, which estimate return periods based solely on peak discharge, often fall short in capturing the complexity of real-world flood events characterized by interdependent features such as volume, duration, and peak. This chapter investigates multivariate modeling approaches using copula theory to better describe the joint behavior of these variables. Building on the foundational work of [Grimaldi and Serinaldi \(2006\)](#), the study evaluates the applicability of nested Archimedean copulas and extends the modeling framework to vine copulas to accommodate asymmetric dependencies and varying tail behaviors. Empirical data from 21 hydrological stations across the Isar and Danube rivers are analyzed, revealing 3 distinct dependence structures that cannot be adequately captured by symmetric or nested copulas alone. This work compares copula model performances and discusses implications for estimating multivariate return periods. The findings underscore the importance of flexible dependence modeling in flood risk assessment.

4.1 Introduction

Floods rank among the most severe natural hazards in Bavaria, both in terms of ecological and socio-economic impact. Structural damages in Bavaria typically occur to private residential buildings, agricultural facilities, public infrastructure, and critical transport systems such as roads and bridges. In 2024 alone, flood events in Bavaria caused an estimated €4 billion in damages. Environmental impacts, such as contamination from leaked heating oil, also pose serious consequences but are often difficult to quantify in monetary values ([für Umwelt \(LfU\)](#)). These significant damages emphasize the need for robust risk assessment tools to support decision-making in floodplain management and infrastructure design. Traditional flood frequency analysis in hydrology has largely focused on univariate approaches, where the relationship between flood peak and its return period is estimated independently of other hydrological variables ([Khajehali et al. \(2025\)](#)). While such analyses are suitable for preliminary risk assessments, they have limited capability to represent the complexity of real-world flood events, especially when applied to the design of hydraulic structures or the planning of flood protection measures ([Grimaldi and Serinaldi \(2006\)](#)). Flood events are characterized not only by their peak discharge but also by their volume and duration. These variables are interdependent, and their joint behavior plays a critical role in determining the severity of flooding impacts. For instance, a moderate peak may still result in significant inundation if the flood duration is long or if the volume exceeds

reservoir capacity. To model these dependencies, multivariate statistical techniques have been introduced, enabling the analysis of joint cumulative distribution functions and probability density functions across multiple variables (Grimaldi and Serinaldi (2006)). Copula theory provides a robust statistical framework for conducting multivariate flood analyses. By separating the marginal distributions from the dependence structure, the joint behaviour of hydrological variables can be studied more flexibly.

In particular, asymmetric copula functions—such are highly effective in flood frequency analysis due to their ability to model varying strengths of dependence between pairs of variables. This is particularly relevant in hydrology, where empirical evidence suggests that the flood peak can strongly influence the dependence between volume and duration, introducing asymmetry into the joint distribution (Grimaldi and Serinaldi (2006)). The work of Grimaldi and Serinaldi (2006) represents a significant contribution in this field. They proposed a trivariate flood event analysis using nested Archimedean copulas to jointly model flood peak, volume, and duration. However, this work points out the shortcomings of nested Archimedean copulas and extends the author's approach by vine copulas. The present thesis addresses three main research questions: (1) How can copula models be applied to characterize flood events in Bavaria, similarly to the approach by Grimaldi and Serinaldi (2006)? (2) How do nested Archimedean copulas compare to more flexible structures such as vine copulas in representing interdependencies? (3) How does the interpretation of return periods differ when using univariate versus multivariate approaches? By addressing these questions, this work aims to contribute to a more nuanced understanding of flood risk, ultimately improving predictive accuracy and informing better flood management strategies in Bavaria.

4.2 Data

Grimaldi and Serinaldi (2006) used the variables peak, volume and duration of the most severe flood event within a year. These variables are derivable from yearly hydrological discharge data. The discharge data we use during our analysis is provided by the Bavarian Environmental Agency's hydrological service ((GKD)) which is data from multiple measurement station along the Isar and the Danube. Based on this, the following gives a brief description of the data, discusses possible flood event detection methods, derives the variables of interest based on the flood definition and ends with a display of the crucial aspects of the obtained data.

After removing stations with too little observations, the data contains discharge values in 15 minute steps for 21 stations along the Isar and Danube from different starting time points, but always up to 31.12.2024. Of these stations, 12 are along the Isar and 9 along the Danube where every station had at least 44 years of observations. As seen towards the end of this section, the alpine river Isar and the low-lying Danube have contrasting hydrological characteristics, enabling a meaningful comparison of flood dynamics in Bavaria. The exact spatial distribution of the considered station displayed plot 4.1.

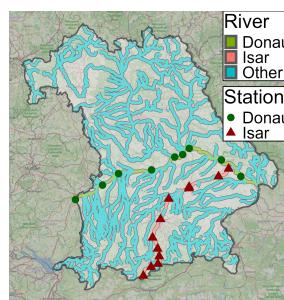


FIGURE 4.1: The location of the selected measurement stations for the Isar and the Danube in Bavaria.

Given the annual discharge data for all these stations, we require to identify the most severe flood event within each year which defined as the event with the largest discharge peak. To stabilize event detection, the following is based on daily average discharge values we calculated based on the 15 minute time intervals in

the original data.

The flood detection approach proposed by [Grimaldi and Serinaldi \(2006\)](#) of using the straight-line method based on a fixed threshold was found to be highly unreliable, but so was a quantile-based straight-line method. Both approaches exhibit significant uncertainties in identifying flood events, particularly, they tend to overestimate flood duration. Instead, we applied the baseflow methods proposed and implemented by [Wasko and Guo \(2025\)](#). This method relies on the baseflow index (BFI) which is the ratio of the baseflow volume to the volume of streamflow. A default BFI threshold of 0.5 was used to distinguish events dominated by rapid runoff contributions typically associated with rainfall- or melt-induced flooding. Exemplary, figure 4.2 shows the hydrograph for the station in Munich in 2024.

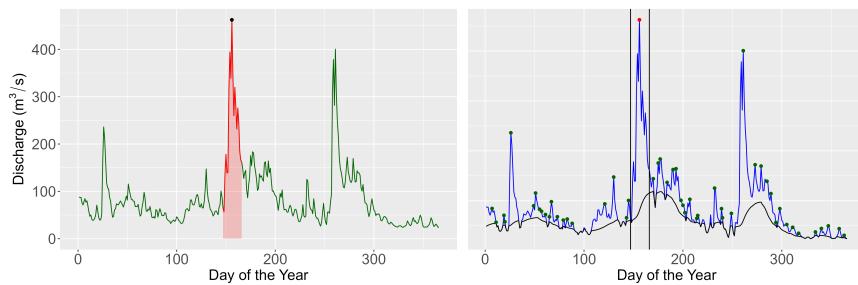


FIGURE 4.2: An exemplary hydrograph of the Munich station in 2024 (left) with the applied flood detection method (right).

Both subplots in the figure display the same hydrograph, but the right describes the identification of the most severe flood. First, all flood events are selected. Then, the flood with the largest peak is identified. The flood duration is based on the previously described process. Finally, the variables of interest are determined. That is, flood peak is the maximum discharge value occurring within the event, flood duration is the time span measured in days between the start and end of the event, as determined by the BFI threshold crossings. Flood volume is the cumulative streamflow over the flood duration, representing total discharge volume in m^3 . These are neatly displayed on the left subplot.

Now, we come to the most crucial aspect in our data and show that this structure is also found in [Grimaldi and Serinaldi \(2006\)](#). That is, fig 4.3 displays the rank correlation coefficient Kendall's τ of every possible combination between the 3 variables separated by river.

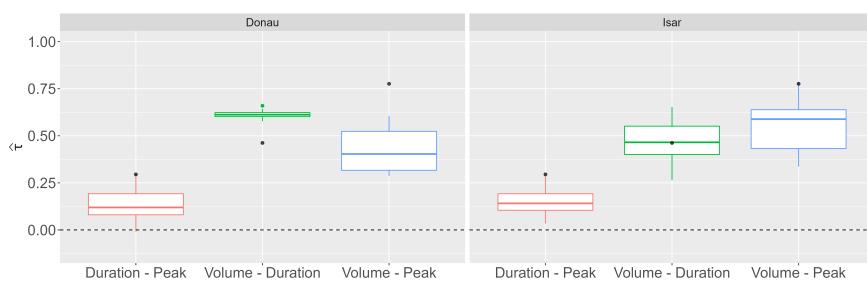


FIGURE 4.3: The Kendall's tau coefficients for the three dependency structures of volume, duration and peak for the Isar and the Danube. Black dots mark the respective results found in [Grimaldi \(2006\)](#).

The boxplots in figure 4.3 are based on the 9 and 12 stations along each river, respectively, and depict the τ values for the corresponding variable combination seen on the x -Axis. The black dots refer to the τ values observed by [Grimaldi and Serinaldi \(2006\)](#). Most important here is that none of the boxplots align horizontally. That is, the strength of dependence differs between all pairs of variables. Thereby, our data suggests 3 distinct dependencies. This finding is most crucial and, as we will see later on, renders [Grimaldi and Serinaldi \(2006\)](#) approach infeasible. Because, as seen from the black dots, not only our data suggest 3

separate dependence structures, but also the river [Grimaldi and Serinaldi \(2006\)](#) considered.

Also interesting is the exact order of correlation values by each river. For both rivers, duration and peak always had the lowest correlation value. For the Danube, volume and duration are always the variables with the highest correlation with an exception of only one station. Nevertheless, all these values are quite similar as seen from the width of the boxplot in figure 4.3. For the Isar, on the other hand, we observe not only more variation in the correlation values, but here the most correlated pair tends to be volume and peak. Of the 12 stations, 8 had volume and peak to be the pair with the highest correlation. This emphasizes the aforementioned contrasting hydrological characteristics which are highly relevant for copula modelling and, thereby, for our analysis.

Finally, the analysis section utilizes return periods of flood peaks to derive average discharge values conditioned on a certain peak. To ensure these are comparable among stations, they are normalized by the station specific mean and standard deviation. To now characterize a conditional distribution using our data, we ordered all flood events within each station by their peak values and then selected the quantiles corresponding to the return periods 2, 5, 10, 20 and 50 years. Thereby, we obtained 21 average discharge values for each return period. Consider figure 4.4 for a visualization.

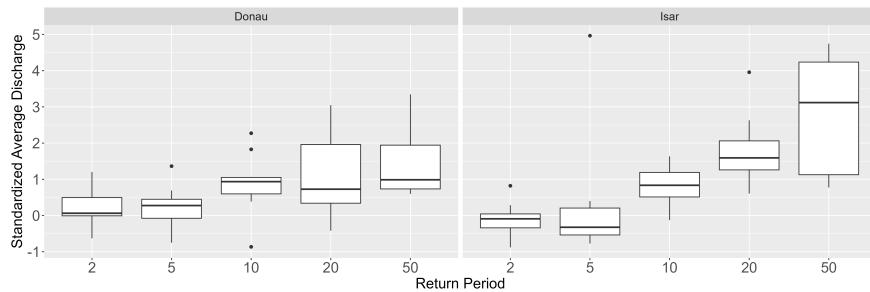


FIGURE 4.4: The standardizes average discharge values by peak quantiles of the GKD data.

First of all, note that the average discharge values are on the y-axis ranging from -1 to 5 . This is due to the standardization process within each station. The x-axis denotes the return periods of the peak. To account for the different structure between rivers, the figure considers them separately. Thereby, each boxplot in the Danube column is based on 9 data points and 12 for the Isar where each data point corresponds to a station. For both subplots, the average discharge increases with an increase in the return period. However, while the subplot of the Danube suggests a moderate increase in average discharge values, the Isar has a stark increase. But by the size of the boxplot at return period 50, this increase does not hold for all stations.

4.3 Methods

To address the dependence structures identified in the previous section, this chapter extends the approach of [Grimaldi and Serinaldi \(2006\)](#) by incorporating vine copulas. This extension is necessary because a simpler approach is insufficient to capture the full correlation pattern observed in the data. The following introduces the foundational theory of copulas, the family of Archimedean copulas as well as nested Archimedean and vine copula models. In addition, methods for copula fitting and model selection are briefly discussed. Then, some applied, but non-essential methods are briefly established. Together, these elements form the theoretical framework on which this paper is based. Finally, a few words to the implementation of these methods and used packages.

4.3.1 Copulas

Zhang and Singh (2019) (p. 62) describe a copula as a cumulative distribution function (CDF) with standard uniform margins. The dimension d of a copula denotes the number of random variables it relates and, hence, a copula is at least bivariate ($d \geq 2$). To give a mathematical definition, consider the vector $u = (u_1, \dots, u_d) \in \mathbb{R}^d$ where $u_j \in [0, 1]$ for $j = 1, \dots, d$. Then, a d dimensional copula is defined by Durante and Sempi (2016) (p. 14) as function $C : [0, 1]^d \rightarrow [0, 1]$ if, and only if, the following conditions hold:

- i) $C(u_1, \dots, u_d) = 0$ if $u_j = 0$ for at least one $j \in \{1, \dots, d\}$.
- ii) $C(1, 1, \dots, 1, u_j, 1, \dots, 1) = u_j$.
- iii) C is d -increasing.

According to Nelsen (2006) (p. 9), condition i) shows that copulas are grounded. In this context, grounded means that plugging in 0 for just one of the variables yields a copula value of 0, independent of the other variables' value. The author also mentions that, using condition ii), the margins of the function C with respect to a certain variable are obtained by plugging in 1 for all other variables. Finally, the condition of C to be d -increasing is cumbersome to map out in higher dimensions, which is why the following is restricted to the $d = 2$ case. According to Nelsen (2006) (p. 8), the copula function C is 2-increasing if for all $u_1, u_2, v_1, v_2 \in [0, 1]$ with $u_1 \leq u_2$ and $v_1 \leq v_2$:

$$C(u_2, v_2) - C(u_2, v_1) - C(u_1, v_2) + C(u_1, v_1) \geq 0$$

Simply put, 2-increasing means that the volume under the copula density function over the rectangle $[u_1, u_2] \times [v_1, v_2]$ is non-negative. This interpretation follows from the fact that copula functions are defined as CDF and holds for higher dimensions, too.

The next section introduces the central theorem in copula theory and also derives the already mentioned copula density.

4.3.2 Sklar's Theorem

Sklar's Theorem is central to the theory of copulas as it proves that any multivariate distribution can be constructed using copulas (Nelsen (2006) p. 17, Durante and Sempi (2016) p. 42). Thereby, this theorem allows to separate the representation of the dependence structure and marginal distribution functions. The theorem is given by Nelsen (2006) (p. 18):

Let $F_{1, \dots, d}$ be a d -dimensional joint distribution function with univariate margins F_1, \dots, F_d . Then, there exists a d -dimensional copula C such that

$$F_{1, \dots, d}(x_1, \dots, x_d) = C(F_1(x_1), \dots, F_d(x_d)) = C(u_1, \dots, u_d)$$

where $u_i = F_i(x_i)$. Also, C is unique if F_1, \dots, F_d are continuous. Equation (4.3.2) allows 2 important conclusion: One, any multivariate CDF may be expressed as a composition of a copula function C and the univariate margins F_1, \dots, F_d . Thereby, Zhang and Singh (2019) (p. 66) conclude that C connects the multivariate CDF to its margins which allows to separately consider marginal and joint behavior of variables. That is, the problem of determining any multivariate CDF is reduced to determining the copula. And two, the marginal distributions do not need to be of the same family because Sklar's theorem holds regardless.

The aforementioned copula density function is given by (see Zhang and Singh (2019), p. 66):

$$c(u_1, \dots, u_d) = \frac{\partial C(u_1, \dots, u_d)}{\partial u_1 \dots \partial u_d} = \frac{f(x_1, \dots, x_d)}{\prod_{i=1}^d f_i(x_i)}$$

where $f(x_1, \dots, x_d)$ denotes the joint density of X_1, \dots, X_d and $f_i(x_i)$ the marginal density of X_i for $i = 1, \dots, d$. Based on this equation, the joint density in terms of the copula density is given by

$$f(x_1, \dots, x_d) = c(u_1, \dots, u_d) \prod_{i=1}^d f_i(x_i)$$

4.3.3 Symmetric Archimedean copulas and generator functions

As Nelsen (p. 109) states, symmetric Archimedean copulas (SACs) are widely applied due to their large variety and easy construction. However, SACs only allow the same dependence strength and structure among all possible pairs of variables as [Zhang and Singh \(2019\)](#) (p.124) point out. Therefore, they are not suitable for our analysis as concluded from figure 4.3 which suggested 3 distinct correlation values. However, SACs remain an important building block for more complex copula models. Thus, this section introduces the concept of a generator function as it determines the family a SACs belongs to. Then, we specifically focus on bivariate SACs because following models are based on these.

We first give the general idea of a generator, then the representation of a copula in terms of the generator and in the end the copula families we use for our analysis.

[Nelsen \(2006\)](#) (p. 110, 111) defines a generator to be a continuous and strictly decreasing function $\phi : [0, 1] \rightarrow [0, \infty)$ such that $\phi(1) = 0$. If $\phi(0) \rightarrow \infty$, the generator is considered to be strict. The inverse $\phi^{-1} : [0, \infty) \rightarrow [0, 1]$ of such generators is strictly decreasing on $[0, \phi(0)]$. We only apply strict generators as seen towards the end of this section.

For a generator to yield a valid d -dimensional copula, [Grimaldi and Serinaldi \(2006\)](#) and [Zhang and Singh \(2019\)](#) (p. 124) mention that the inverse requires to be completely monotone which is given if it has derivatives of all orders with alternating sign

$$(-1)^k \frac{d^k \phi^{-1}(t|\theta)}{dt^k} \geq 0.$$

Now, we are in the position to formulate the general representation of a 3-dimensional SAC in terms of its generator. The relation is given by [Zhang and Singh \(2019\)](#) (p. 123) as

$$C(u_1, u_2, u_3 | \theta) = \phi^{-1}(\phi(u_1 | \theta) + \phi(u_2 | \theta) + \phi(u_3 | \theta) | \theta).$$

Equation (4.3.3) shows that SACs are uniquely defined by their generator function and a parameter vector θ which we introduce next. As mentioned by [Nelsen \(2006\)](#) (p. 110, 111, 114), the assumed functional form of the generator translates to a specific copula family. Or, vice versa, assuming a copula family implies assuming a specific generator function. The θ vector, on the other hand, influences the dependence strength within the assumed copula family as seen in [Zhang and Singh \(2019\)](#) (p. 86). This parameter vector takes on an important role in fitting a copula to observed data. That is, for an assumed copula family, this parameter vector remains to be estimated from the data. The exact approach is further discussed in section 4.3.7. For now, note that we focus on the 3 generator functions with a one-dimensional θ vector. These are specified in table 4.1.

TABLE 4.1: Generator functions of selected Archimedean copulas according to [Zhang and Singh \(2019\)](#) (p. 130) and tail dependencies according to [Zhang and Singh \(2019\)](#) (p. 132).

Copula Family	Parameter θ	Generator Function $\phi(t)$	Tail Dep.
Clayton	$\theta \in [-1, \infty) \setminus \{0\}$	$\phi(t \theta) = \frac{1}{\theta}(t^{-\theta} - 1)$	Lower
Gumbel-Hougaard	$\theta \in [1, \infty)$	$\phi(t \theta) = (-\ln t)^\theta$	Upper
Frank	$\theta \in (-\infty, \infty) \setminus \{0\}$	$\phi(t \theta) = -\ln\left(\frac{e^{-\theta t} - 1}{e^{-\theta} - 1}\right)$	None

Finally, equation (4.3.3) shows that the arguments to the SAC are exchangeable (see [Nelsen \(2006\)](#) (p. 38)). Exchangeability is a form of symmetry and implies that the copula treats all its arguments the same. Thereby, this representation displays the aforementioned restriction of SACs being able to only depict one unique dependence structure.

4.3.4 Taildependence and Rotation

After explaining what it means for a copula to be of a certain family, the following introduces the family specific concept of tail dependence. Also, we briefly explain how copulas are manipulated to extend the possible dependence structure one family captures.

Tail dependence is differentiated into upper and lower tail dependence. Their formulas are given by [Czado \(2019\)](#) (p. 34 - 35) as

$$\lambda^{\text{upper}} = \lim_{t \rightarrow 1^-} \mathbb{P}(X_2 > F_2^{-1}(t) \mid X_1 > F_1^{-1}(t)) = \lim_{t \rightarrow 1^-} \frac{1 - 2t + C(t, t|\theta)}{1 - t}$$

and

$$\lambda^{\text{lower}} = \lim_{t \rightarrow 0^+} \mathbb{P}(X_2 \leq F_2^{-1}(t) \mid X_1 \leq F_1^{-1}(t)) = \lim_{t \rightarrow 0^+} \frac{C(t, t|\theta)}{t}.$$

As seen from both equations, tail dependence is defined as conditional probability that both variables are above or below a threshold quantile. Thereby, tail dependence measures how likely it is for both random variables to jointly exhibit extreme behavior. However, upper tail dependence refers to both variables attaining large values while lower tail dependence means both variables are jointly small. Note that both, upper and lower tail dependence, depend on the copula function C and parameter θ and, thus, on the copula family. The tail dependencies implied by the copula families we consider are also listed in table 4.1 as stated in [Zhang and Singh \(2019\)](#) (p. 132).

Finally, tail dependence is a family specific property, however, a copula function may be rotated to change its native tail dependence behavior. Following [Pan et al. \(2024\)](#), this is done by modelling $u'_i = 1 - u_i$ instead of u_i itself. Every such transformation concludes in a 90 degree rotation of the copula function in the corresponding direction. This approach is based on the definition of the copula function as multivariate CDF. Because if instead of u_i the transformation u'_i is modelled, the probabilistic statement of the copula in the continuous case changes to $C(u'_1, u'_2|\theta) = \mathbb{P}(X_1 \geq x_1, X_2 \leq x_2)$ and $C(u_1, u'_2|\theta) = \mathbb{P}(X_1 \leq x_1, X_2 \geq x_2)$, respectively.

4.3.5 Fully Nested Archimedean copulas

Fully nested Archimedean copulas (FNACs) build upon SACs and partially alleviate their restrictions. Note that these are the models [Grimaldi and Serinaldi \(2006\)](#) made extensive use of.

As our analysis applies to the trivariate case, FNACs and vines in section 4.3.6 are introduced for this trivariate case only.

FNACs are built by nesting bivariate SACs

$$C(u_1, u_2, u_3 \mid \theta) = C_1(C_2(u_1, u_2 \mid \theta_2), u_3 \mid \theta_1),$$

where θ_1 and θ_2 are the parameters corresponding to copula function C_1 and C_2 and $\theta = (\theta_1, \theta_2)$ is a vector containing all parameters. Note that there are only 2 distinct parameters θ_i which is why FNACs, in the trivariate case, are only able to capture 2 distinct dependence structures. This allows 2 conclusions. First, partial exchangeability remains which means that within the bivariate nested copulas C_2 , the two arguments u_1, u_2 are interchangeable (see [Embrechts et al. \(1999\)](#) p. 375). So to a degree, symmetry prevails. And second, the nested variables u_1, u_2 have the same marginal relation with u_3 . That is, $C(a, 1, u_3|\theta) = C(1, a, u_3|\theta) = C_1(a, u_3|\theta_1)$. In essence, the statements are equivalent but the important take away is that FNACs are not able to display 3 distinct dependence structures. Thereby, they are not suitable for our analysis. As also [Grimaldi and Serinaldi \(2006\)](#) observed 3 distinct correlations, their results are questionable for the same reason.

Additionally to this restriction, [Górecki et al. \(2016\)](#) mention that FNACs require the sufficient nesting condition to be fulfilled for equation (4.3.5) to yield a valid copula. We limit our considerations to FNACs where all nested copulas are of the same family. This corresponds to what [Grimaldi and Serinaldi \(2006\)](#) used in their analysis. Then, the sufficient nesting condition is fulfilled if deeper nested variables have a stronger degree of dependence, i.e. $\theta_1 \leq \theta_2$. (see [Grimaldi and Serinaldi \(2006\)](#)).

Note that equation (4.3.5) may also be represented in terms of the generator function. Thereby, additional requirements regarding the composition of generator functions emerge, as mentioned by [Zhang and Singh \(2019\)](#) (p. 174). However, discussing these requirements is beyond the purpose of this paper. The interested reader is referred to [Zhang and Singh \(2019\)](#) (p. 174).

4.3.6 Vine Copulas

Vine copulas use the pair-copula construction (PCC) explained by [Czado \(2019\)](#) (p. 77 - 80) to characterize multivariate dependence structures. That is, PCC decomposes multivariate densities into products of (conditional) bivariate densities (see [Czado \(2019\)](#) p. 88).

There exist multiple vine copula classes, depending on the structure the PCC implies, but as we are concerned with the trivariate case, these constructions are equivalent.

We use [Czado \(2019\)](#) (p. 78, 90) and define the trivariate copula density as

$$c_{123}(u_1, u_2, u_3 | \theta) = c_{12}(u_1, u_2 | \theta_{12}) \cdot c_{23}(u_2, u_3 | \theta_{23}) \cdot c_{13|2}(u_1 | u_2, u_3 | u_2 | \theta_{13|2})$$

where $u_i | u_j = F_{ij}(x_i | x_j)$ denotes the conditional probability. Note that this copula density is based on the simplifying assumption for vines ([Czado \(2019\)](#) p. 90, [Vatter and Nagler \(2018\)](#)) which means that the conditional bivariate density $c_{13|2}$ is independent of exact x_2 values. It only depends on the conditional probabilities.

Visible from the number of parameters in the θ -vector, vines are able to capture all 3 distinct dependence structures in the trivariate case. Also, in contrast to FNACs from section 4.3.5 where we followed [Grimaldi and Serinaldi \(2006\)](#), we do not require the bivariate copulas to be of the same family. Thereby, not only the strength of dependence between all 3 variables may differ, also the dependence structure is allowed to change from pair to pair.

4.3.7 Estimation and Selection Process

In practice, we need not only to estimate the parameter vector θ , but also select the best fitting copula. Thus, the following briefly introduces the pseudo maximum likelihood (ML) approach. Also, we give a reminder on the Akaike Information Criterion (AIC) as it is our information criterion of choice to select a copula model.

To avoid assumptions on the marginal distributions, we estimate the parameter vector θ using the pseudo-likelihood proposed by [Genest et al. \(1995\)](#)

$$\hat{\theta} = \operatorname{argmax}_{\theta} l(\theta) = \operatorname{argmax}_{\theta} \sum_{k=1}^n \log[c(u_{1k}, u_{2k}, u_{3k} | \theta)],$$

where u_{ik} denotes the marginal empirical distribution function scaled by $\frac{n}{n+1}$. These transformed variables are referred to as pseudo-observations. Depending on the copula model, the definition of the copula density follows either from equation (4.3.5) or (4.3.6). Thus, the exact estimation process slightly differs between copula models.

The AIC is given by [Fahrmeir et al. \(2013\)](#) (p. 164) as

$$AIC = -2l(\hat{\theta}) + 2(|M| + 1)$$

where $l(\hat{\theta})$ represents the log-likelihood of the copula model fit and $|M|$ the number of parameters included in the model. We select that copula model with the smallest AIC.

4.3.8 Identifying univariate margins

While the estimation process described in section 4.3.7 utilizes the empirical distribution function, an empirical function has undesired properties when re-transforming copula data. That is, during the estimation process, the empirical distribution functions ensure we do not affect the copula model fit by

missspecifying the marginal distributions. However, after fitting the copula, whenever the inverse of the empirical distribution is applied, it bins any continuous data because it is a step-wise function. This, of course, limits the power of our copula analysis. Thus, we decided to fit a Generalized Extreme Value (GEV) distribution to the marginal distribution only for re-transforming any results from the fitted copula models. The distribution function for the GEV family is given by [Coles \(2001\)](#) (p. 47)

$$G(z) = \exp \left\{ - \left[1 + \xi \left(\frac{z - \mu}{\sigma} \right) \right]^{-1/\xi} \right\},$$

defined for $\{z | 1 + \xi \left(\frac{z - \mu}{\sigma} \right) > 0\}$ where $-\infty < \mu < \infty$ denotes the location parameter, $\sigma > 0$ the scale parameter and $-\infty < \xi < \infty$ the shape parameter. In practice, these parameters are usually unknown, but [Coles \(2001\)](#) (p. 50) describes how these parameters are estimated from data using the ML approach.

Since GEV distributions are not at essence for our work, we refer the interested reader to [Coles \(2001\)](#) (chapter 3) for a more detailed consideration.

4.3.9 Kendall's τ

According to [Kendall and Gibbons \(1990\)](#) (p. 6), τ is a measure of association between two random variables that distinguishes concordant and discordant pairs. Concordance means that the two variables move in the same direction while discordance means moving in opposite directions.

[Zhang and Singh \(2019\)](#) (p. 86) and [Nelsen \(2006\)](#) (p. 159, 161 - 164) show that Kendall's τ is directly connected to the generator function and, thus, a function in the parameter θ .

$$\tau(t) = 4 \int_0^1 \frac{\phi(t|\theta)}{\phi'(t|\theta)} dt + 1$$

where $\phi'(t|\theta)$ denotes the derivative of the generator function. Note that this relation is positive which is seen in [Zhang and Singh \(2019\)](#) (p. 134). That is, if the parameter of copula increases, the strength of dependence increases. Vice versa, if correlation increases, θ increases, too. Also, note that this implies that estimating a θ implicitly estimates a τ value. This is a relation we will use during our simulation.

Empirically, there are multiple versions of Kendall's τ depending on the data structure. Since this paper focuses on continuous variables only, the formula given by [Kendall and Gibbons \(1990\)](#) (p. 5) is applicable

$$t = \frac{P - Q}{\frac{1}{2}n(n - 1)}.$$

P denotes the number of concordant and Q the number of discordant pairs in the data.

4.3.10 Software

The whole analysis is implemented using the programming language R. We used the `hydroEvents` package by [Wasko and Guo \(2025\)](#). This method relies on the BFI as explained in the data section using a default BFI of 0.5. Additionally, the `eventBaseflow` function calculates the BFI at each time step and extracts discrete flood events when the BFI falls below the specified threshold for a user-specified minimum duration. For copulas, we relied on the `copula` package by [Hofert et al. \(2025\)](#) and the `pobs` function to apply the empirical distribution function as described in section 4.3.7. FNACs are dealt with using the HAC by [Okhrin and Ristig \(2014\)](#), especially the function `estimate.copula` for FNAC fitting which implements the ML approach from section 4.3.7. This required some additional code to select the best fitting copula according to the AIC. For vine copulas, `VineCopula` package by [Nagler et al. \(2024\)](#) was consulted. Especially the function `RVineCopSelect` which implicitly fits a selection of copulas and also selects the one with the smallest AIC. Finally, GEV distribution were fitted using the function `fevd` from the `extRemes` package by [Gilleland and Katz \(2016\)](#) which uses MLE to determine the parameters mentioned in 4.3.8.

4.4 Simulation

To examine the incapability of FNACs to capture 3 distinct dependence structures in a trivariate setting, we ran a simulation. The true underlying model builds a vine copula models. This section is limited to the most crucial finding which is highly relevant for the interpretation of our results in the following section.

We set up the simulation by drawing 27000 random samples of size 15, 30, 50, 1000. The sample size of 50 represents our real world conditions while 1000 observations aim to examine large sample behavior. The two smaller samples sizes are interesting because we earlier decided to remove stations due to their small sample size.

For each drawn sample, the underlying vine copula model has 2 wheels to tweak: First, each copula density of the underlying vine copula model is allowed to be of one of the copula families listed in table 4.1 leading to a possible total of $3^3 = 27$ copula family combinations. Second, we allowed for 4 different correlation values, 2 of which correspond to the observed average correlation values in the Isar and the Danube. The remaining 2 aimed to examine general behavior of FNACs. That is, we added a Low-Medium-High (LMH) correlation structure with correlation values of 0.1, 0.5, 0.85 and a Low-High-High (LHH) structure using 0.1, 0.8, 0.8. Thereby, we had 27000 data points per sample size which are split among all 27 possible copula family combinations and 4 possible correlation structures. This leads to roughly $\frac{27000}{4 \cdot 27} = 250$ data points per setup. It is not exactly 250 because we used a uniform draw to select copula families and correlation structure as it drastically simplified implementation.

The one result we want to focus on is described in figure 4.5.

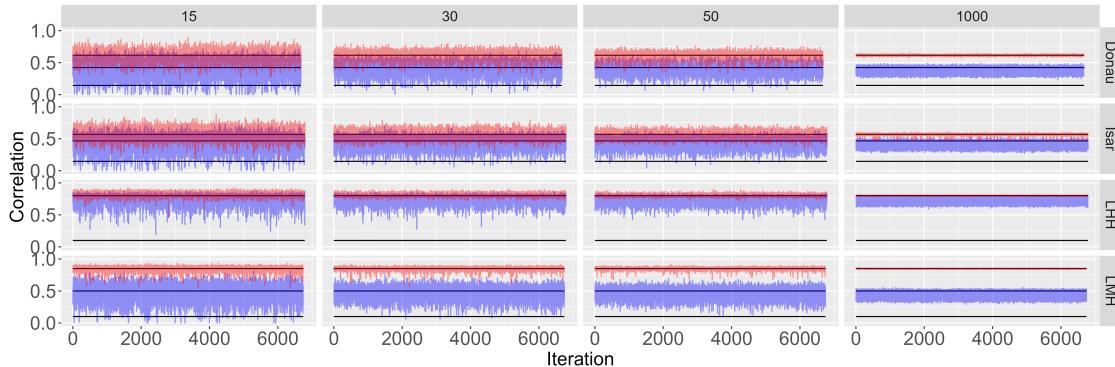


FIGURE 4.5: The simulation results by sample size and correlation structure.

This figure uses multiple subplots displaying trace plots of the τ values estimated by the FNAC model. Kendall's τ estimates are displayed on the y-axis while the index of the iteration in which this model was fitted is on x-axis. Note that this plot does not differ by copula family combination leading to roughly $\frac{27000}{4} = 6750$ iterations each. The black lines in each subplot refers to the true underlying correlation values. The name of the corresponding correlation structure is to the right hand side of the plot. Additionally, the figure is divided by the sample size on which the estimated FNAC model is based on. From a sample size of 15 on the left up to a size of 1000 on the right. Color-wise, the red line refers to the $\hat{\tau}$ of the nested FNAC and the blue line to the outer estimate. Due to the sufficient nesting condition (see section 4.3.5), the inner $\hat{\tau}$ in each iteration is always larger than the outer.

The first observation is the decrease in variance of the estimators as the sample size increases. This holds for all possible correlation structures and was expected. Now, focus only on the column of sample size 1000. Especially in the Low-Medium-High subplot, we observe that the inner τ estimate moves around the most upper black line. Thereby, the nested copula in a FNAC correctly captures the largest correlation value. The blue line, however, moves around the second largest black line and its volatility remains comparably large. We conclude that, first, the τ estimate of the outer copula of a FNAC model varies within a set interval for large sample sizes. This is similar to what Grimaldi and Serinaldi (2006) found in their work when they

examined how SACs perform if the true underlying model is a FNAC. Second, and highly relevant for our analysis, the $\hat{\tau}$ based on the outer copula of a FNAC tends towards the second largest correlation value in the (simulated) data. This implies that, due to the same copula margins mentioned in section 4.3.5, FNACs systematically overestimate the weakest dependence strength. This is an important result because it explains not only the comparably bad performance of FNACs during our application, but also their bias in the results.

4.5 Application

Due to our finding during the simulation, we focus on presenting the fitted vine copulas models. Only for the model comparison, we jointly consider FNACs and vines to discuss the effect of the bias in FNACs and its practical meaning.

4.5.1 Goodness of Fit

The following discusses the goodness of fit for the vine copula model based on the Munich station using figure 4.6.

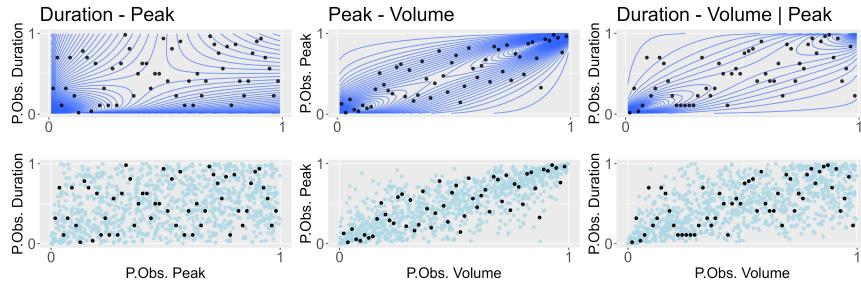


FIGURE 4.6: The contour lines and synthetic data plots for the Munich station.

This figure consists of 6 subplots. Each column of subplots refers to a pair of variables specified in the column header. The variable named first is displayed on the y-axis. Note that all subplots are on copula level. Thus, all axes range from 0 to 1. The top row of subplots shows the contour lines for the fitted copula density and the bottom row a synthetic random sample from the copula model. The black points in every subplot depict the pseudo observations of the corresponding variables.

Starting with bottom row, a good model fit implies that the light blue points capture the structure of the pseudo observations. This seems to be the case for all variable pairs as not only the shape of the black points is nicely reflected in the synthetic data points, but also the strength of dependence matches. This is validated when comparing the empirical correlation values with the correlation implied by the fitted models as seen in table 4.2. The largest absolute difference between the correlation values is just 0.02.

TABLE 4.2: Descriptives on the fitted copula model for Munich station.

Pair	Copula Family	Empirical τ	Fitted τ	$ \Delta\tau $
Duration – Peak	Clayton	0.15	0.16	0.01
Peak – Volume	Gumbel-Hougaard	0.49	0.51	0.02
Duration – Volume	180° Rotated Gumbel-Hougaard	0.60	0.59	0.01

Now, consider the top row of figure 4.6 in combination with the corresponding copula families mentioned

in table 4.2. The Clayton and 180° rotated Gumbel-Hougaard copula imply lower tail dependence for the relation between duration and peak as well as duration and volume, respectively. Thereby, if the flood peak is small, the duration tends to be short, too. Also, the volume of a flood is more likely to be small given a short flood duration. However, due to its small strength of dependence, the tail dependence for duration and peak is rather small. This is not only suggested by the contour lines, but follows from the low τ value which implies that parameter θ is small, too, as mentioned in section 4.3.9. This in turn affects the tail dependence as it is a function in θ , as discussed in section 4.3.4. Finally, a Gumbel copula is fitted to the peak and volume pair implying upper tail dependence. Thus, given a large peak, the flood volume tends to be large, too.

4.5.2 Fitted Tail Dependencies

Because tail dependence is an important concept from a hydrological point of view, we extend the tail dependence analysis from the previous section to all considered stations.

Contemplate figure 4.7 for a visual assessment of the tail dependence structure.

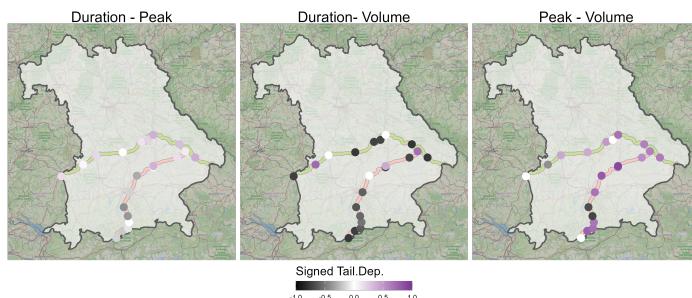


FIGURE 4.7: The signed tail dependencies of the three variable combinations.

This plot shows 3 different subplots, one for each variable pair. The subplots are based on figure 4.1 and display all stations colored by their tail dependence value. Because we only allowed for copula families with either upper, lower or no tail dependence, we visualize the signed tail dependence. That is, upper tail dependence values remain unchanged, but lower tail dependencies are multiplied by -1 . Thereby, the blacker a station is colored in, the stronger the lower tail dependence and the more purple the stronger the upper tail dependence. A totally white point on the map refers to no tail dependence. In these cases, a Frank copula has been fitted to the variable pair.

In general, we observe lighter colors for the duration-peak pairs which is due to the rather small correlation values. This follows from the reasoning in the previous section. Duration and volume exhibit the darkest colors suggesting that if the flood duration is short, the flood volume tends to be small, too, for the majority of the considered stations. This is especially the case at the Isar stations. But, the relationship begins to shift towards a positive dependence approximately at the middle of the river, likely due to the influence of tributaries that contribute additional water as the river progresses downstream. In contrast, the Danube displays mixed signals for the duration-volume pair, indicating a more complex or variable relationship, possibly reflecting the diverse hydrological influences across its larger catchment area.

For the peak-volume pair, both rivers show a positive tail dependence, indicating that higher peak flows are generally associated with larger flood volumes. This is consistent with expected hydrological patterns where significant peak events often appear with substantial volumes of water. The duration-peak pair exhibits the weakest tail dependence, suggesting that the flood duration does not strongly influence the peak discharge. This weak correlation is indicative of the more dynamic nature of peak discharges, which may be influenced by short-term, localized weather events rather than the overall duration of the flood.

Across all three pairs, the Isar consistently demonstrates more extreme tail dependence trends compared to the Danube. This difference is likely due to the more extreme flood characteristics in the Isar, which are influenced by factors such as snowmelt, topography, and high spring rainfall events (Parajka et al. (2010)).

4.5.3 Event Probability

The following examines the difference between a univariate and the multivariate approach for flood characterization and thereby addresses one of our research questions. To answer it, the following focuses on the flood event in Munich 2024 which had a peak of $462\text{m}^3/\text{s}$, took 20 days and had a volume of 406m^3 .

Figure 4.8 displays the marginal fit of a GEV distribution to the peak data for the Munich station.

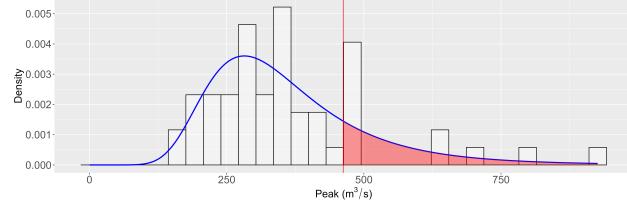


FIGURE 4.8: The GEV fit for the Munich station. The red area delineates floods with a peak equal or higher to the flood event of 2024.

The x-axis of this plot denotes the peak values, the y-axis the density of the fitted GEV model. The histogram displays the original data to which the GEV distribution was fitted. The resulting smooth distribution is marked by the blue line. The red vertical line marks the peak value of $462\text{m}^3/\text{s}$ and the shaded area visualizes the probability to observe a peak at least as large. Based on the GEV fit, the probability is calculated to be 19%. Thereby, this approach assigns the a return period of such a peak to be $\frac{1}{0.19} \approx 5$ years. Thus, characterizing the whole flood event only by its peak, the return period of the whole flood in Munich of 2024 according to the univariate model is 5 years.

The trivariate copula model allows to also consider volume and duration values to characterize a flood event. Figure 4.9 is based on figure 4.6 and visualizes how the multivariate model determines the probability for a flood event to be at least as severe. Also, this plot helps to understand the quite stark differences in the probabilities.

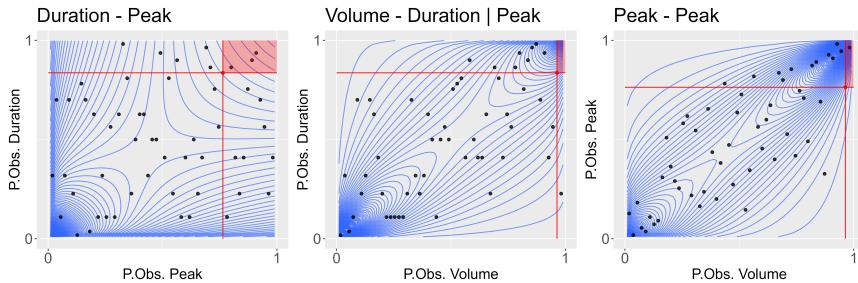


FIGURE 4.9: The copula fit for the Munich station. The red area delineates floods with a peak, volume and duration equal or higher to the flood event of 2024.

Note that, as before, this figure displays the data in terms of their pseudo observations. The red dot in each subplot marks the event in 2024 and the red lines correspond to the univariate univariate observed pseudo values. Thereby, a flood event is at least as severe as the flood in Munich if it lies within the shaded area. Thus, the probability is obtained by integrating the joint copula density over the cube made up by the shaded area. According to our model, this adds up to be 2.7% which corresponds to a return period of merely $\frac{1}{0.027} \approx 37$ years. Thereby, the return period for such an extreme flood is $\frac{37}{5} \approx 7$ times longer if the flood is characterized not only by its peak, but also by its volume and duration value. The reason for this drastic difference is seen in figure 4.9. While the pseudo observation of the peak value of the event is around 0.76, the volume is at 0.96. That is, the volume during this flood was exceptionally large which decreases the probability of such an event to occur. Visually, this is seen from the shaded area being very slim. In contrast, the univariate consideration of peak values only is not capable to account for this.

4.5.4 Model Comparison

Finally, this section examines the effect of different peak values onto the other characteristics of a flood. Mainly, we are interested in the average discharge value during a flood event as it is a measure of the average energy the system has to deal with.

First, we fit station specific GEV distributions onto the observed peak values and determine peaks for the return periods of 2, 5, 10, 20 and 50 years. We choose the upper bound of 50 years to validate the model predictions using the available data. Conditional on these peaks, the most likely combination of duration and volume are determined as well as the average discharge values. Figure 4.10 is based on figure 4.4 and displays our results.

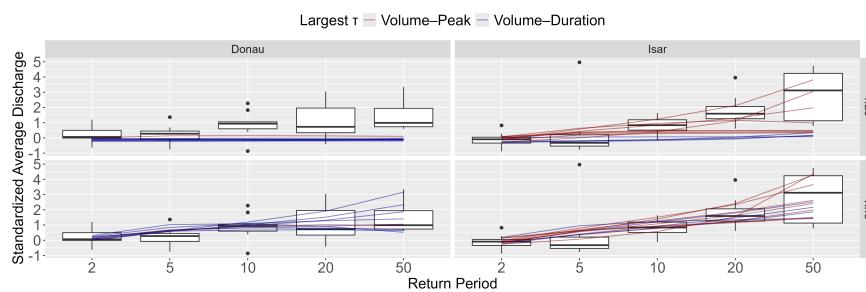


FIGURE 4.10: The model comparison by river between FNACs and vines.

The rows of the figure refer to the model structure applied to predict the average discharges. Additionally, we colored each model prediction by which variable pair had the highest correlation. First, consider the vine model fits in the bottom row. The models correctly capture the trend suggested by the boxplots independent of which pair of variables has the highest correlation. For a return period of 50, model predictions and boxplots increase in variance. This is reasonable because the manner in which an event is extreme depends on the station. Thereby, a joint behavior in boxplot and model prediction suggests a good fit. For FNACs, consider their performance within the stations of the Isar first. There is a visible difference in model performance depending on which variable pair has a larger correlation value. If volume-peak have a larger correlation value, the model captures the underlying data structure more reliable than for volume and duration being stronger correlated pair. For the Danube, FNAC models barely move at all failing to capture any trend in the data whatsoever. To quantify the visual analysis, consider the mean absolute difference of the model prediction to the median of the data: For FNACs, the error for the Danube is 0.70 and the error for the Isar 0.90 . Vine outperformed FNACs with an error of 0.35 for the Danube and 0.53 for the Isar. However, take these with a grain of salt because especially for a return period of 50 years, we deal with an extreme event. And it is reasonable to expect extreme events to be highly station specific. Thereby, a model perfectly predicting the median of the average discharge data is not necessarily desirable.

The reason for the underperformance of FNACs is their incapability of capturing 3 unique dependence structures which leads to 2 distinct effects. First and mainly, due to the shared bivariate marginal copula, the performance of FNAC models is highly impacted by which variable pair is higher correlated. Because by the sufficient nesting condition, the stronger correlated variable pair is required to build the nested copula. If duration-volume is the higher correlated pair and, thus, peak in the outer copula, the dependence between duration-peak and volume-peak is forced to be identical. Thereby, conditioning on a peak value has the identical effect on both nested variables. However, if peak is inside the nested copula, duration-peak and volume-peak remain a distinct relation structure to peak. Thereby, both models are wrong, but one of them remains more flexible.

The second effect is independent of the nesting structure. As result of the simulation we observed a systematic overestimation of the lower correlation value. Because in our data, the smallest correlation value is always between duration-peak, this strength of dependence is systematically overestimated. This implies that an increase in peak affect duration more than it should. And because all observed correlation values between the two variable are positive, the duration of a flood for given an increasing peak is always overestimated. This implies that the average discharge predicted by FNACs is systematically too low because the volume is

divided by a duration that is too long. In conclusion, even if some of the FNAC models look as if they captured the underlying structure, they really did not. Thereby, this application nicely depicts the shortcomings of FNACs and the reason why they are not suitable if their assumptions are violated.

4.6 Conclusion

The aim of this work was to investigate the potential of copula-based multivariate models for flood risk assessment in Bavaria, focusing on the joint behaviour of flood peak, volume and duration. Three main research questions were addressed: the applicability of copula models to Bavarian flood data, the comparative performance of NACs and vine copulas, and the impact of using multivariate rather than univariate approaches to estimate return periods. Our results confirm that copula theory provides a robust framework for modelling the dependence structure between multiple hydrological variables. By applying both vine and FNACs to 21 stations, we demonstrated that multivariate models are capable of capturing complex, asymmetric dependencies that are not accounted for in traditional univariate analyses. A key finding is the spatial variability in tail dependence structures across the study area. The Isar, in particular, exhibited more pronounced and often more extreme dependencies—especially between volume and duration, suggesting stronger hydrological controls related to topography, snowmelt, and spring precipitation patterns. In contrast, the Danube showed more mixed signals, underscoring the necessity for flexible dependence structures that can adapt to regional hydrological conditions. When comparing the model structures, vine copulas consistently outperformed FNACs which suffered from structural biases. This limitation was particularly evident when analysing conditional average discharge values, where FNACs systematically underestimated flood severity. Finally, the comparison of univariate and multivariate return period estimations underscores the need of a multivariate perspective. For the 2024 Munich flood event, the return period increased from 5 to 37 years when considering the joint severity of peak, volume, and duration. This highlights the risk of underestimating extreme events when relying solely on univariate peak discharge values. In conclusion, the integration of flexible multivariate copula models, especially vine copulas, represents a significant advancement in flood risk analysis. Their ability to capture spatial and structural dependencies can greatly improve the reliability of flood risk assessments and help create more resilient flood management strategies in Bavaria.



5

Introduction

Author:

Supervisor:

5.1 Intro About the Seminar Topic

5.2 Outline of the Booklet



6

Co-occurrence of extremes

Author: Chongjun

Supervisor: Henri Funk

6.1 Abstract

This report investigates the spatial dependence of extreme co-occurrence globally by employing hypothesis testing. The analysis aims to assess whether extreme events, such as temperature and precipitation extremes, exhibit spatial dependencies across the Earth's land surface. Using observational data from the Climatic Research Unit (CRU), the Berkeley Earth (BE) dataset, and the Global Precipitation Climatology Centre (GPCC), we explore both positive and negative spatial dependencies between extreme temperature and precipitation events. The study quantifies the intensity of spatial dependence by introducing the Probability Multiplication Factor (PMF), which provides a numerical representation of the strength of the spatial correlation between extreme events. Furthermore, we examine how spatial dependence varies with distance and latitude, offering insights into the geographic and climatological factors that influence the co-occurrence of extreme events. The findings demonstrate significant patterns of both positive and negative spatial dependencies, highlighting the complex relationships between extreme weather events and their spatial distribution across different regions.

6.2 Introduction

Climate extremes, such as heatwaves, heavy precipitation events, droughts, and tropical cyclones, pose severe threats to both natural and human systems. Their intensification in recent decades has led to escalating economic losses, ecosystem disruptions, and public health crises. These extremes are not random or isolated occurrences but exhibit distinct statistical dependencies. While extreme events can cluster over time (temporal dependence), they are also spatially dependent, meaning that extreme conditions in one region are often correlated with those in neighboring or even distant areas due to large-scale climate dynamics. Understanding this spatial dependence is crucial for improving risk assessments, predicting multi-region impacts, and formulating climate adaptation strategies.

Although temporal dependence plays a role in the persistence of climate extremes, this report focuses exclusively on their spatial dependence—the extent to which extreme climate conditions co-occur across different locations. Spatial dependence arises due to shared meteorological drivers, such as atmospheric circulation patterns, teleconnections, and geographic influences. For instance, heatwaves are often linked across multiple regions by high-pressure blocking systems, while extreme precipitation events can span large areas due to stalled weather fronts. Recognizing and quantifying these spatial relationships is critical for accurately modeling extreme event risks and preparing for simultaneous disasters that affect multiple regions.

One striking example of spatial dependence in climate extremes is the 2010 Eastern Russian heatwave,

which was not confined to Eastern Russia alone but also influenced temperatures across Eastern Europe and Central Asia. The event was driven by a persistent atmospheric blocking pattern, which sustained extreme temperatures over an extensive spatial scale.

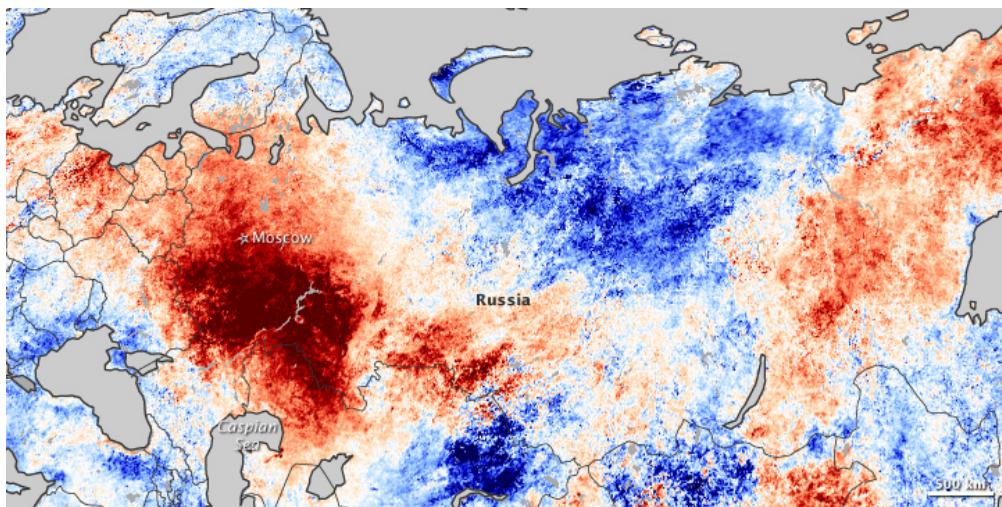


FIGURE 6.1: Source: [<https://earthobservatory.nasa.gov/images/45069/heatwave-in-russia>]. Heat waves in Eastern Russia, Eastern Europe and Central Asia in 2010.

Another example is tropical cyclones in the Atlantic, where clusters of hurricanes can develop due to favorable oceanic and atmospheric conditions, causing sequential or simultaneous destruction across multiple coastal regions. Similarly, compound flooding events, where storm surges coincide with extreme rainfall across interconnected watersheds, demonstrate the importance of spatially correlated extremes in shaping disaster impacts.

The study of spatial dependence in climate extremes has gained increasing attention as traditional models often assume independent occurrences, leading to the underestimation of multi-region risks. Modern statistical approaches, such as extreme value theory (EVT), copula models, and geostatistical methods, provide powerful tools for quantifying spatial dependence and improving predictions of concurrent extreme events. Additionally, the role of climate teleconnections, such as the El Niño-Southern Oscillation (ENSO) and the North Atlantic Oscillation (NAO), further highlights how distant regions can experience correlated extremes.

Despite advancements in understanding spatial dependencies, significant challenges remain. Non-stationarity in climate extremes—driven by global warming, land-use changes, and shifting atmospheric patterns—complicates the identification of stable spatial dependence structures. Furthermore, as climate extremes become more frequent and intense, their spatial footprints may expand, necessitating continuous refinement of statistical models. Addressing these challenges is crucial for enhancing resilience against climate risks and improving predictive capabilities in disaster-prone regions.

This report will discuss methodologies for quantifying the spatial dependence of climate extremes and examine key characteristics of this dependence on a global scale. By deepening our understanding of how extreme climate events co-occur across space, we can better identify large-scale patterns, improve predictive models, and refine risk assessments to support climate adaptation strategies.

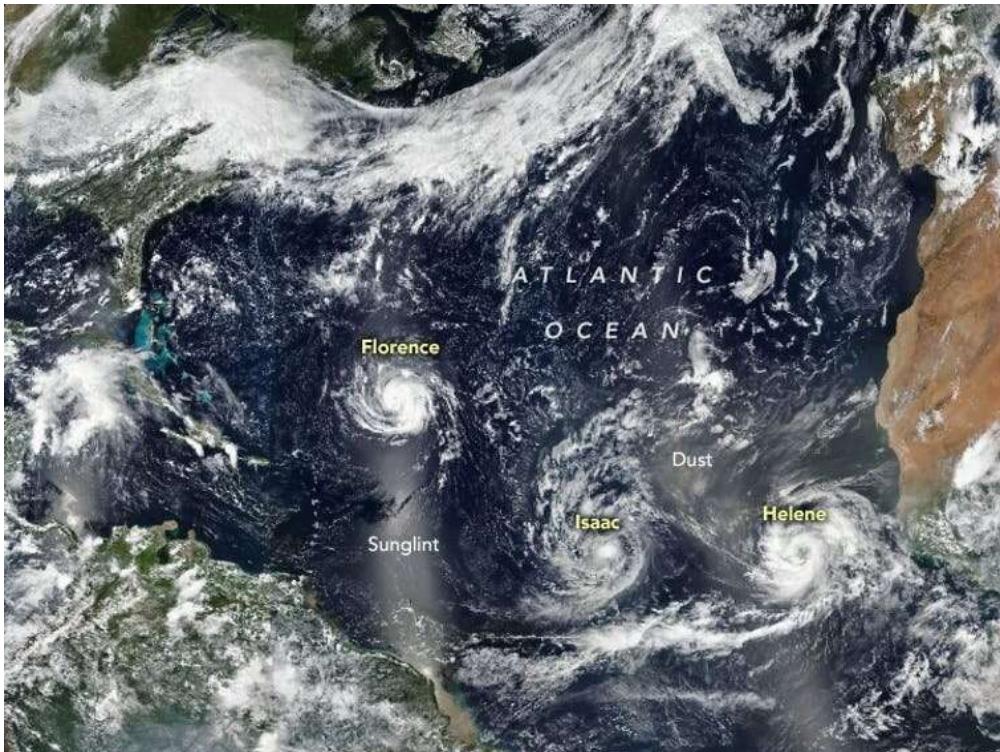


FIGURE 6.2: Source: [<https://phys.org/news/2019-03-big-storm-clusters-hurricane-hotspots.html>]. Hurricanes occur simultaneous in the Atlantic in 2019.

6.3 Tail Dependence

Tail dependence refers to the relationship between the extreme values of two or more variables, specifically whether extreme values of one variable tend to coincide with extreme values of another variable. Unlike traditional correlation $r(G_i, G_j)$, which measures the linear relationship between variables, tail dependence focuses on the co-occurrence of extreme events, such as the occurrence of severe climate conditions in different regions simultaneously. The objective of this research is not to analyze correlation between regions but to explore how extreme values in one region might depend on extreme values in another region, even if there is no linear correlation.

Statistically, tail dependence is a measure used to quantify the likelihood of simultaneous extreme values in the tails of distributions. There are various methods to measure tail dependence, with the most commonly used being the upper and lower tail dependence coefficients. These coefficients describe the probability of one variable exceeding a threshold given that the other variable has already exceeded a similar threshold, capturing the essence of tail events and their joint occurrences. Tail dependence is crucial in risk management, particularly in fields such as finance and climate science, where extreme events (e.g., market crashes or heatwaves) can have substantial consequences.

It is important to note that while correlation can provide insights into the likelihood of tail dependence, a high correlation between two grid cells (G_i and G_j) suggests a higher probability of their extreme values occurring together. However, a high correlation does not necessarily imply that extreme values will co-occur, nor does the absence of correlation rule out the possibility of tail dependence. In other words, tail dependence can manifest even when two variables are not correlated, and vice versa.

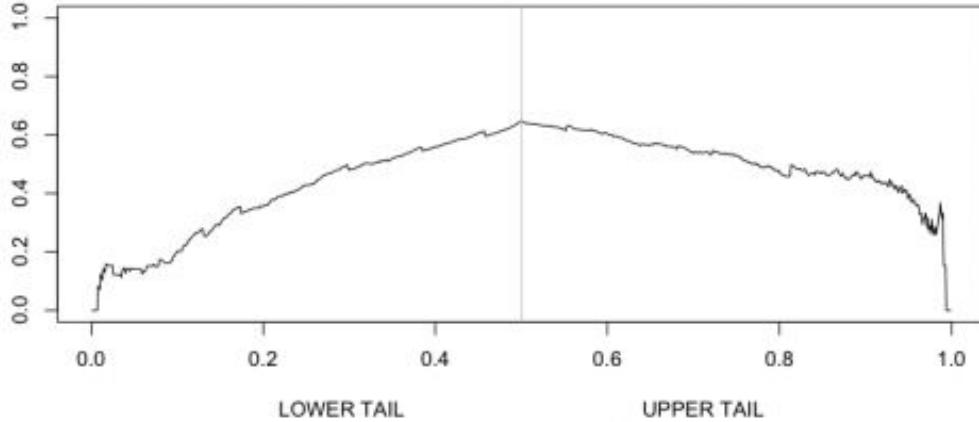


FIGURE 6.3: Source: [https://freakonometrics.hypotheses.org/2435]. Tail dependence of 2 variables.

6.4 Hypothesis Test

We conducted a hypothesis test to assess the spatial dependence of climate extremes. The null hypothesis, H_0 , is that climate extremes occur independently at two grid cells, G_i and G_j , where $i \neq j$, at a significance level of $\alpha = 0.05$. We aim to identify spatial dependence by rejecting H_0 . The identification of extremes in a grid cell is defined by whether the value exceeds the 95th percentile or falls below the 5th percentile of the distribution. Therefore, the probability that a single grid cell, G_i , is extreme is $P(G_i \text{ is extreme}) = 0.1$ (10%).

Next, we consider the joint probability of two grid cells, G_i and G_j , being extreme. Specifically, the probability that both G_i and G_j are extreme is calculated based on two cases: both cells exceeding the 95th percentile, or both cells falling below the 5th percentile. This defines positive concurrence of extremes. For negative concurrence, we consider the case where G_i is greater than the 95th percentile and G_j is less than the 5th percentile, or vice versa. In mathematical terms:

- Positive concurrence: $P(G_i > 0.95 \text{ and } G_j > 0.95) \text{ or } P(G_i < 0.05 \text{ and } G_j < 0.05)$
- Negative concurrence: $P(G_i > 0.95 \text{ and } G_j < 0.05) \text{ or } P(G_i < 0.05 \text{ and } G_j > 0.95)$

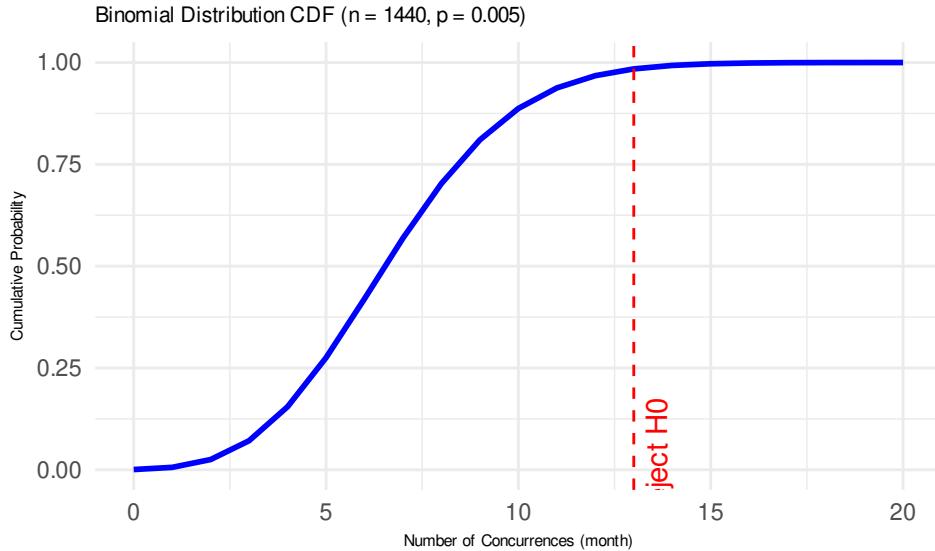
Under the assumption of no spatial dependence, the probability of either positive or negative concurrence, p_0 , is computed as:

$$p_0 = 2 \times 0.0025 = 0.005$$

This is the probability of concurrent extremes occurring at both grid cells, regardless of whether the concurrence is positive or negative, under the null hypothesis of no spatial dependence.

Next, we consider a time frame of 120 years, or 1440 months. The expected number of months during which both grid cells experience concurrent extremes is $E(X) = n \times p_0 = 1440 \times 0.005 = 7.2$ months. To test for spatial dependence, we model the number of concurrent extreme months using a binomial distribution:

$$X \sim \text{Binomial}(n = 1440, p = 0.005)$$



Using the binomial distribution's cumulative probability, we reject H_0 if the number of months with concurrent extremes, x , is greater than or equal to 13. That is, we reject H_0 at the 0.05 significance level when $x \geq 13$. This corresponds to the condition where:

$$P(X \geq 13) \leq 0.05$$

Thus, if the number of months with concurrent extremes exceeds 13 in the 120-year period, we conclude that there is spatial dependence between the two grid cells. The corresponding probability threshold for rejecting H_0 is when $p_0 \geq 0.009$.

6.5 Probability Multiple Factor

In order to assess the strength of spatial dependence, the study introduces the Probability Multiplication Factor (PMF), which quantifies the degree of extreme concurrence between two grid cells. The PMF is defined as the ratio of the actual probability of extreme concurrence to the baseline probability of extreme concurrence under the null hypothesis, denoted as p_0 . Mathematically, this can be expressed as:

$$\text{PMF} = \frac{P(\text{extreme concurrence})}{p_0}$$

The threshold for the PMF is calculated by dividing $p_0 = 0.005$ by the baseline probability of extreme concurrence $p_0 = 0.005$, yielding a threshold value of:

$$\frac{0.009}{0.005} = 1.8$$

When the PMF exceeds or equals this threshold, i.e., when:

$$\text{PMF} \geq 1.8$$

it indicates the presence of spatially dependent extremes. Furthermore, the larger the PMF, the stronger the spatial dependence between the extreme events in the two grid cells.

6.6 Results

The study utilizes temperature and precipitation products from the Climatic Research Unit (CRU), the temperature product from Berkeley Earth (BE), and the precipitation product from the Global Precipitation Climatology Centre (GPCC). These datasets provide time series data spanning a 120-year period (1901–2020), enabling a comprehensive analysis of extreme event co-occurrence and spatial dependence.

The study divides the globe into $2^\circ \times 2^\circ$ grid cells, selecting only those with a land area greater than 30% for analysis. Consequently, the research primarily focuses on spatial dependence over land, excluding oceanic regions.

Analysis of temperature and precipitation extremes reveals distinct patterns of spatial dependence. Temperature extremes predominantly exhibit positive spatial co-occurrence, with a mean Probability Multiplication Factor (PMF) of 3.25 across 58% of grid cell pairs. In contrast, negative dependence is detected in only 15% of grid cell pairs. Precipitation extremes display both positive and negative spatial dependencies, observed in 27% and 25% of grid cell pairs, respectively, with mean PMF values of 2.64 and 2.40.

The strength of positive concurrence is particularly pronounced in geographically proximate regions and diminishes with increasing distance. Specifically, for both temperature (Figure 4. A) and precipitation (Figure 4. B) extremes, PMF remains consistently higher when the distance between grid cells is less than 5000 km. Beyond this threshold, PMF exhibits only a marginal decline as distance increases, suggesting a limited attenuation of spatial dependence at larger scales. However, in cases of extreme distances, a PMF exceeding 1.8 alone is not sufficient to confirm spatial dependence, as atmospheric circulation patterns, including heat advection and precipitation transport, can facilitate the transfer of extreme conditions between distant grid cells within a monthly timeframe. Nevertheless, the fact that PMF does not significantly decrease beyond 5000 km indicates that distance is not the primary determinant of spatial dependence. This implies the presence of additional underlying factors contributing to the strong spatial correlation observed between geographically distant regions.

The negative concurrence strength of temperature extremes (Figure 4. C) is also pronounced in neighboring regions. In contrast, no distinct spatial pattern is observed for the negative concurrence of precipitation extremes (Figure 4. D).

The study further identified that extreme events tend to co-occur more strongly in the tropics and high-latitude regions of both hemispheres. In the analysis, solid lines represent the probability mass function (PMF) of regions with positive concurrence, while dashed lines indicate regions with negative concurrence. For both temperature and precipitation extremes, the mean PMF for negative concurrence remains consistently lower than that for positive concurrence, suggesting that positively dependent extremes are more prominent.

The impact of long-term warming trends can be evaluated by comparing the red and blue lines in the figures. In the tropics, the strong positive dependence of temperature extremes (Figure 5.E) is largely due to warming trends, as indicated by PMF values reaching up to 6 when the trend is included. However, after removing the trend, the PMF drops to 4.5, emphasizing the substantial influence of warming on the observed concurrence of temperature extremes. In contrast, negative dependence in temperature extremes is more commonly found in the northern high-latitude regions. The historical simulations from CMIP6 models show a closer match with observed data when trends are included, whereas the historical natural (hist nat) simulations better align with observations when trends are removed.

For precipitation extremes (Figure 5.F), the spatial pattern of dependence is similar to that of temperature extremes. However, in this case, long-term warming trends have only a minor effect on PMF, suggesting that other factors play a more significant role in shaping the spatial dependence of precipitation extremes. Another key difference from temperature extremes is that the results from historical simulations and historical natural

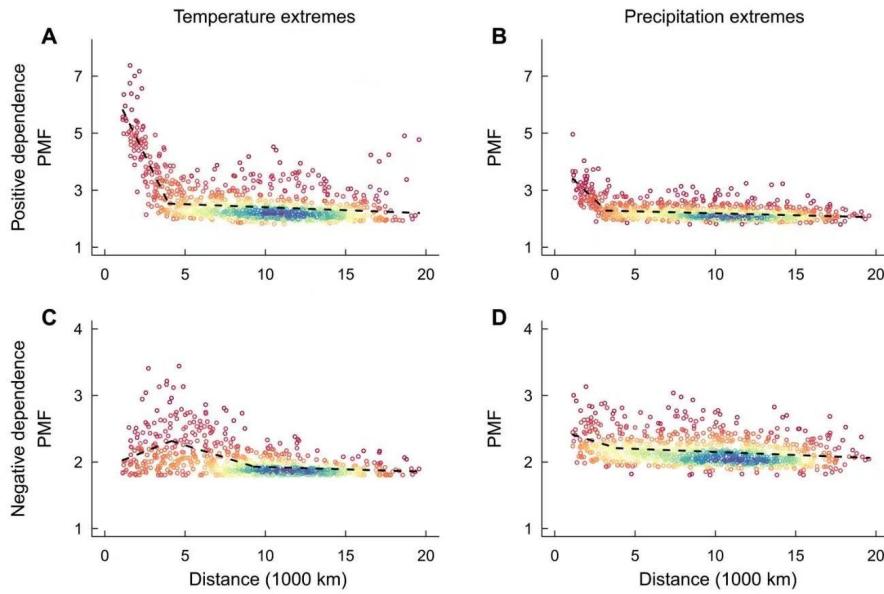


FIGURE 6.4: Source: [https://www.science.org/doi/10.1126/sciadv.abo1638]. The distribution of PMF values across different grid cell pairs at varying distances. Each scatter point on the plot represents the PMF of a grid cell pair, while the color indicates the density of point distribution. Blue signifies the highest density, indicating more concentrated pairs, whereas red represents the lowest density, corresponding to more dispersed pairs. The data used is observed data.

(hist nat) simulations are nearly identical. This suggests that the warming trend does not substantially alter the spatial dependence of precipitation extremes, unlike its effect on temperature extremes.

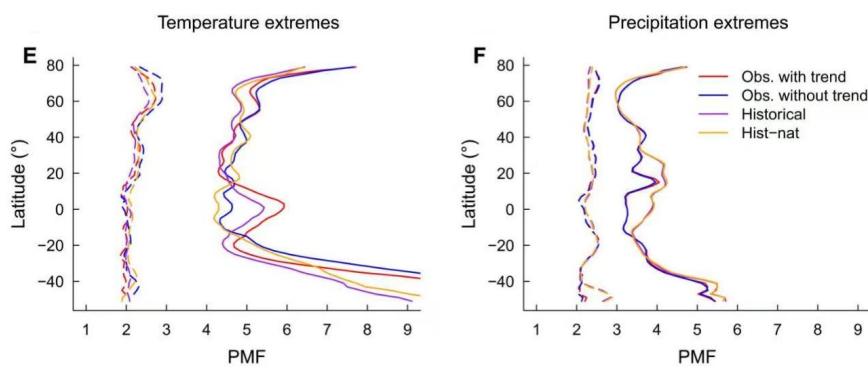


FIGURE 6.5: Source: [https://www.science.org/doi/10.1126/sciadv.abo1638]. The distribution of PMF values across different grid cell pairs at varying latitudes. Solid lines represent the PMF of positively concurrent regions, dashed lines indicate negatively concurrent regions. Four scenarios: observation with/without trend, historical simulated data and hist-nat simulated data (natural forcing only).

6.7 Sensitivity Test

This study also investigated how spatial dependence responds to different thresholds used to define extreme events. The hypothesis test was conducted with a threshold of 0.05/0.95, while the sensitivity analysis explored a range from 0.01/0.99 to 0.1/0.9. The 0.01/0.99 threshold represents the strictest criterion, meaning only the most extreme events are considered, while the 0.1/0.9 threshold is the least strict, allowing more events to be included. The findings suggest that as the threshold becomes stricter, fewer grid cell pairs exhibit extreme concurrence, leading to a smaller number of identified regions. At the same time, the mean probability mass function (PMF) increases, indicating that spatial dependence is stronger under stricter threshold conditions.

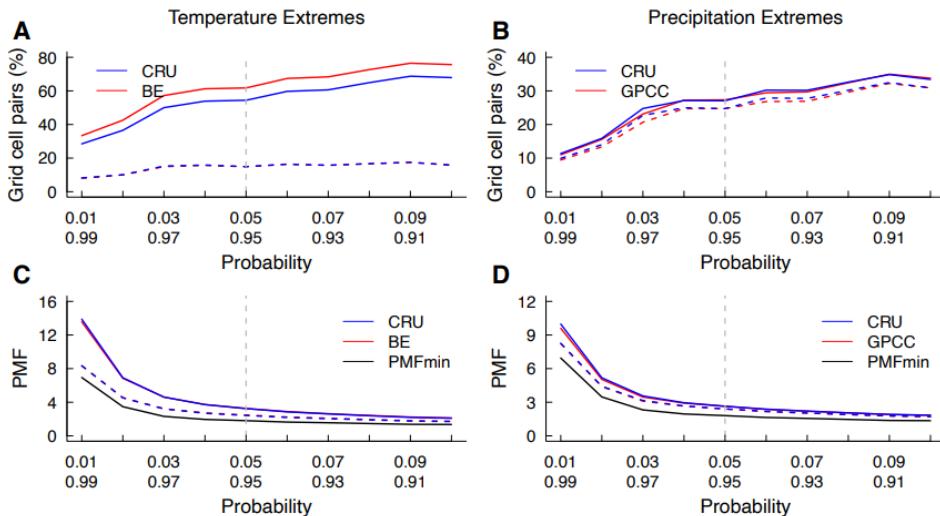


FIGURE 6.6: Source: [https://www.science.org/doi/10.1126/sciadv.abo1638]. The proportion of land grid cell pairs where climate extremes exhibit significant positive (solid lines) or negative (dashed lines) dependence, assessed using probability thresholds ranging from 0.9/0.1 to 0.99/0.01, based on observational data from 1901 to 2020. (C,D) Corresponding results for (A,B), but depicting the mean probability multiplication factor (PMF) for positively (solid lines) and negatively (dashed lines) dependent climate extremes.

Author: Yijie Chen

Supervisor: Helmut Küchenhoff

7.1 Introduction

Public health is at increasing danger from the effects of climate change—particularly heat-related effects on health and diseases which increase with rising temperatures. To identify the complex relationships connecting climate variables and human health is the key to effective intervention and prevention efforts. In recent years, both statistical and machine learning methods have been applied to quantify this relationship and forecast risks in future, Shang says.

This study examines the applications of a variety of methods to examine risks for climate-sensitive diseases that are influenced by temperature fluctuations. We apply some statistical methods, including splines and Generalized Additive Models (GAMs), as well as machine learning algorithms such as Gradient Boosting and XGBoost. These approaches enable us to model the nonlinear and intricately entwined relationships between different climate indicators, health outcomes, and socio-economic factors.

To illustrate the practical use of these methods, we present two case studies based on the 2024 Europe Report of the Lancet Countdown on Health and Climate Change ([van Daalen et al., 2024](#)), with a regional focus on Europe. The first study investigates heat-related mortality, analyzing how temperature impacts human deaths, as previously demonstrated in a large-scale observational study ([Gasparrini et al., 2015](#)). The second study examines the transmission risk of West Nile Virus (WNV), using predictive models informed by a combined dataset of climatic and socio-economic variables, with the modeling approach based on the XGBoost framework ([Chen and Guestrin, 2016](#)).

7.2 Background

7.2.1 Climate and Public Health

Climate change is increasingly recognized as a major global driver of public health risks. Among climate-sensitive outcomes, two of the most pressing are heat-related mortality and the transmission of vector-borne diseases, such as West Nile Virus (WNV). Rising temperatures and more frequent heatwaves have been linked to excess mortality, particularly among vulnerable populations such as the elderly and individuals with pre-existing health conditions. At the same time, changes in temperature and precipitation patterns have affected the habitat and seasonality of disease-carrying vectors like mosquitoes, contributing to the spread of WNV in Europe. These public health threats have become a central focus in environmental epidemiology and climate-health research. For example, the 2024 Lancet Countdown report highlights the growing health burden of both heat exposure and infectious disease risk in the context of climate change ([van Daalen et al., 2024](#)).

7.2.2 Related Literature

In addition to model flexibility and performance, interpretability plays a vital role in selecting suitable methods for public health applications. While traditional linear models often fall short in capturing the complex temporal dynamics of environmental exposure, the use of splines provides an elegant solution. Hansen et al. systematically outlined the theoretical foundation for spline-based modeling, emphasizing their ability to flexibly represent smooth functions while controlling for overfitting. This makes splines particularly suitable for applications where relationships between variables change gradually over time or vary across geographic regions. ([Hansen et al., 2006](#))

Building on these developments, distributed lag non-linear models (DLNM) have emerged as a powerful approach for simultaneously modeling exposure-response and lag-response functions, especially in the context of climate-related health outcomes such as heat-related mortality. These models enable researchers to estimate both immediate and delayed effects of temperature, as demonstrated by Gasparrini et al. in a large-scale, multi-country analysis ([Gasparrini et al., 2015](#)).

Altogether, these methodological advancements—GAMs for flexible modeling, splines for smooth function estimation, DLNMs for time-varying effects, and XGBoost for classification in high-dimensional, imbalanced data—form a comprehensive analytical toolkit. They allow for a nuanced understanding of how climate variability translates into health risks, providing the basis for the two empirical applications explored in this report.

7.3 3 Theoretical framework

This section outlines the theoretical foundation of the statistical and machine learning methods used in this study. The aim is to model complex relationships and non-linear dependencies among health outcomes, climate indicators, as well as socio-economic variables.

7.3.1 3.1 Splines

Splines are flexible mathematical tools that model smooth nonlinear relationships across defined intervals. They are especially useful in health-related data modeling, such as temperature effects, due to their ability to control smoothness and avoid overfitting. ([Hansen et al., 2006](#))

7.4 Generalized Additive Models (GAM)

Generalized Additive Models (GAMs) extend the classical Generalized Linear Models (GLMs) by incorporating smooth, non-parametric functions of predictor variables. This flexibility allows GAMs to capture complex, nonlinear relationships between predictors and the response variable, without requiring a strict parametric form. They are particularly valuable in environmental and public health contexts, where such relationships often evolve over time or vary across spatial or demographic dimensions.

The concept of GAMs was first formally introduced by Hastie and Tibshirani ([Hastie and Tibshirani, 1986](#)), who proposed a statistical framework that combines the interpretability of linear models with the adaptability of smoothing splines and other basis functions. In their foundational work, they demonstrated how GAMs could be used to model structured nonlinearity while maintaining additive interpretability. Since then, GAMs

have become a widely used tool in epidemiology, climate-health modeling, and beyond, especially when the assumption of linearity is inappropriate.

7.4.1 Gradient Boosting

Gradient Boosting sequentially adds weak learners to build predictive models. With each iteration, the algorithm fits a new learner to the errors remaining from what ensembles have been made so far—honing its overall model. This policy helps attenuate both the bias and variance of predictions, typically leading to improved prediction performance on structured data sets and the like.

7.4.2 XGBoost

XGBoost (Extreme Gradient Boosting) is a scalable and efficient tree boosting algorithm that performs second-order Taylor approximation and incorporates regularization to prevent overfitting. It is particularly effective for high-dimensional, structured data in health research. ([Chen and Guestrin, 2016](#))

7.4.2.1 3.4.1 Modeling High-Dimensional Climate-Health Data

1. Climate Variables: Temperature, humidity and precipitation, which have a direct effect on mortality and disease transmission.
2. Bioclimatic Variables: Long-term climate trends relevant to vector-borne diseases.
3. Sociodemographic Data: For example population density, degree of urbanization, or economic metrics, revealing vulnerability and exposure from context.

7.4.2.2 Greedy Feature Selection

XGBoost employs a greedy division method, using information gain to optimize decision trees: Highly efficient in reducing forecast error and identifying the most relevant variables.

7.4.3 Lag Effects and CrossBasis Function

To account for the potential delayed effects of heat exposure on mortality, we introduced a lag structure in the modeling framework. Specifically, we used a cross-basis function, which allows for simultaneous modeling of both the exposure-response relationship and the lag-response relationship.

The temperature-mortality association was modeled using a natural cubic spline with internal knots placed at the 10th, 50th, and 90th percentiles of each region's weekly temperature distribution. The lag dimension was constructed using discrete lags at 0, 2, and 3 weeks, capturing both immediate and short-term delayed responses.

This cross-basis setup enhances the flexibility of the model in capturing the non-linear and cumulative nature of heat-related mortality. While the most substantial effects were typically observed in the immediate period (0–3 days), delayed impacts spanning up to several weeks were also evident, particularly in regions with sustained heat stress. The cross-basis approach is particularly valuable for identifying at-risk populations and informing time-sensitive public health interventions.

7.5 Application

7.5.1 Heat-related mortality

7.5.1.1 Data Description and Modeling Approach

Based on data from 1 January 2015 to 30 November 2022, this chapter is a systematic investigation into fatal ailments due to high temperature that took in 823 localities across 35 European countries. There were more than 45 million ciphered deaths in the study, among which over 22 million were female deaths and more than 21 million were of men. To capture the temperature–mortality relation, this study melds climate data, mortality data and demographic material into a coherent whole within the setting of statistics. Counted mortality data on a daily basis were taken from national sources, with population figures from Euro-stat. The temperature figures were derived from the ERA5-Land reanalysis data set which provided high temporal and spatial resolution suitable for impact studies in health. Merging those three data sets produced a pan-European mortality record that is standardized to population. To model temperature effects on mortality, a quasi-Poisson regression model was used. This approach can accommodate over-dispersion in count data and is therefore well suited to modelling mortality. In order to control for seasonality and long-term trends, a natural cubic spline of time with eight degrees of freedom per year was included. This adjustment ensures that the estimated temperature effects are not confounded by broader temporal fluctuations which are not climate related. Furthermore, as heat exposure may not have immediate mortality effects the model takes lag structures into account. Lag effects were estimated using a cross-basis function from a Distributed Lag Non-Linear Model (DLNM), which offers a joint modeling of exposure–response and lag–response relationship. A natural cubic spline is used to model the temperature effect, with internal knots placed at the 10th, 50th and 90th percentiles of the local weekly temperature distribution. The lag dimension was integrated to integer lags of 0, 2 and 3 weeks, allowing both immediate and later physiological effects of heat exposure to be captured. This provides a means for looking at the cumulative effects of heat across various time horizons. To quantify the burden of heat-attributable mortality, heat weeks were defined as those weeks in which the average daily temperature surpassed a region’s minimum mortality temperature. Then the heat-linked deaths were calculated as extra deaths per 100 thousand residents. These estimates were made separately for two reference periods: 2003–2012 and 2013 to 2022. Where data gaps are possible, imputation methods were used to ensure continuity of trends. In order to estimate the uncertainty around empirical mortality risk, Monte Carlo simulations were employed. This technique yields multiple draws from the distribution of model parameters, producing probabilistic confidence intervals for the estimated rates of heat-attributable mortality.

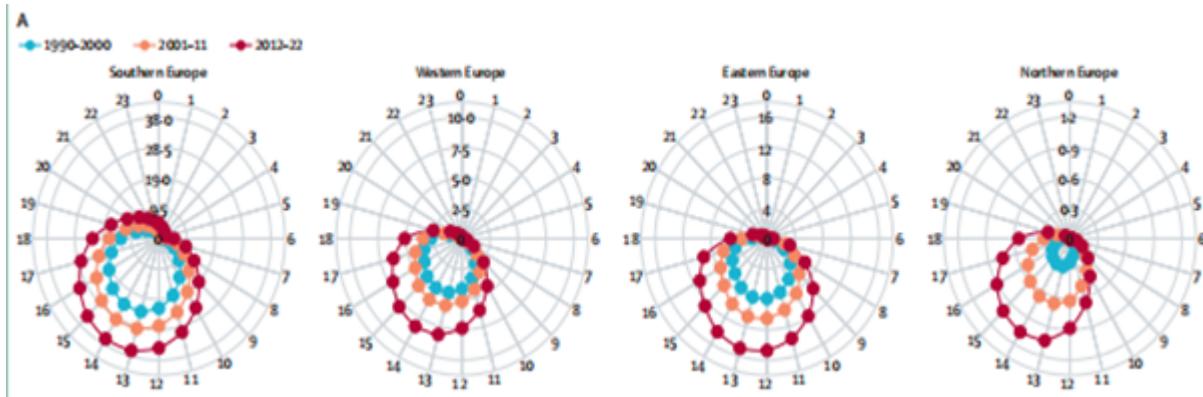


FIGURE 7.1: Radar plots illustrating the 24-hour distribution of heat stress risk across Southern, Western, Eastern, and Northern Europe during three decades: 1990–2000, 2001–2011, and 2012–2022. Adapted from Van Daalen, et al. (2024).

7.5.1.2 Result

Based on data from 1 January 2015 to 30 November 2022, this chapter is a systematic investigation into fatal ailments due to high temperature that took in 823 localities across 35 European countries. There were more than 45 million ciphered deaths in the study, among which over 22 million were female deaths and more than 21 million were of men.

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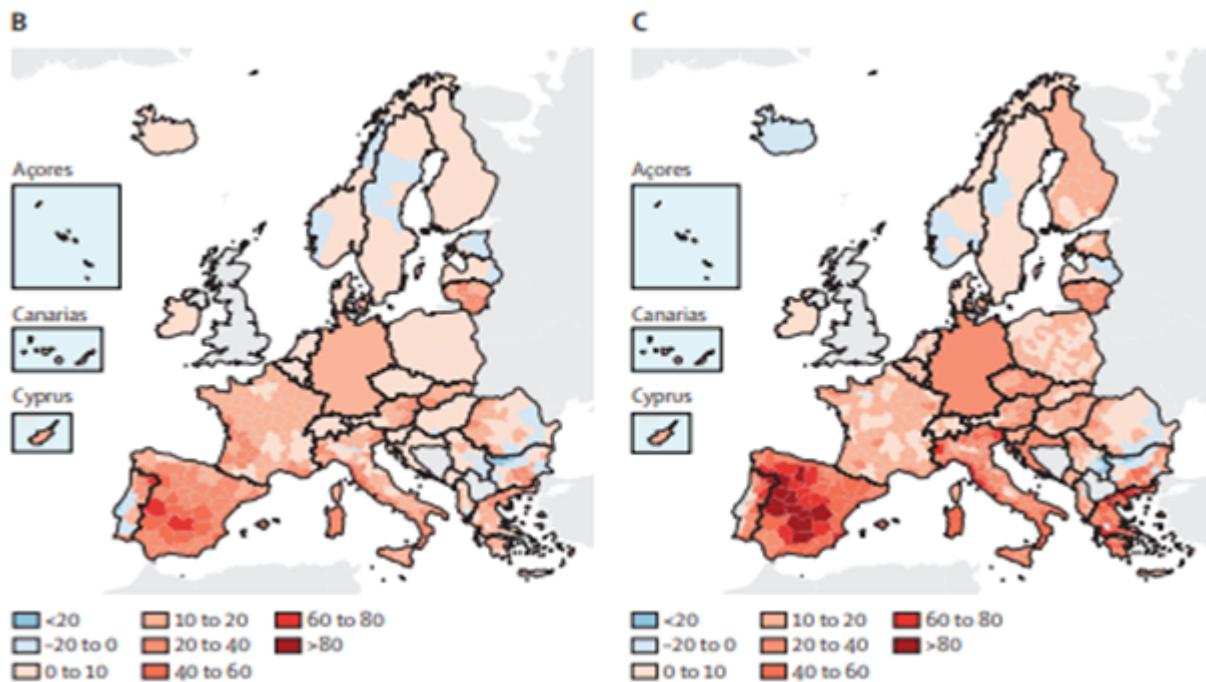


FIGURE 7.2: Heat-attributable mortality rates per 100,000 population across Europe, comparing two periods: 2003–2012 and 2013–2022 Adapted from Van Daalen, et al. (2024).

To quantify the burden of heat-attributable mortality, heat weeks were defined as those weeks in which the average daily temperature surpassed a region's minimum mortality temperature. Then the heat-linked deaths were calculated as extra deaths per 100 thousand residents. These estimates were made separately

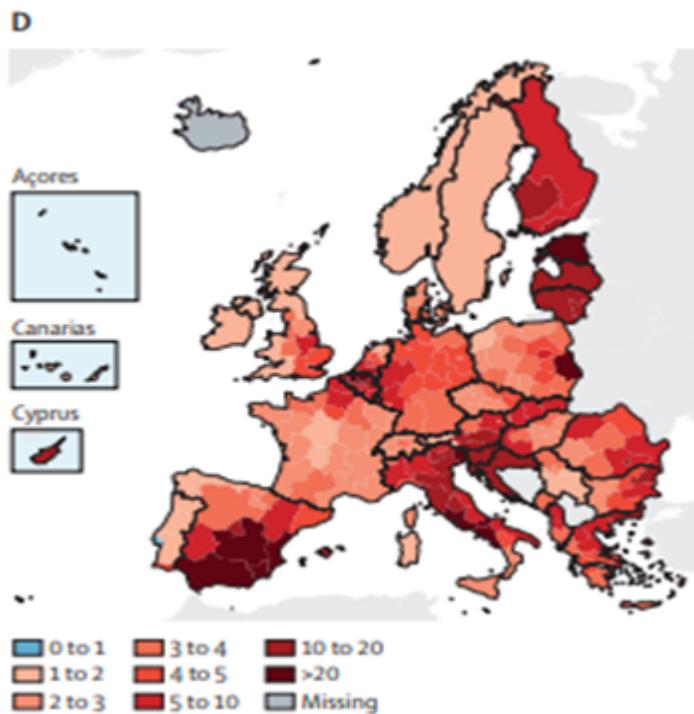


FIGURE 7.3: Heat-attributable mortality rates per 100,000 population across Europe, comparing two periods: 2003–2012 and 2013–2022 Adapted from Van Daalen, et al. (2024).

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7.6 West Nile Virus

7.6.1 Data and Method

To estimate the probability of West Nile Virus (WNV) transmission, our study integrates three main categories of input: climate data, WNV infection records, and socioeconomic factors. Besides temperature and rainfall – two factors with significant impacts on mosquito ecology – another 19 bioclimatic variables are taken from the ERA5-Land reanalysis dataset. This gives information at a resolution of 0.1° if needed, sufficient to capture truly fine-grained geographic detail. Among the 19 bioclimatic variables are ones that pick out long-term climatic trends. It may be possible in future work to use these to model the spread of the WNV more accurately both in space and time. The Y variable is human WNV infection distribution-data obtained from the European Centre for Disease Prevention and Control (ECDC). The dataset covers 800 NUTS3-level administrative regions across Europe. Despite the territorial coverage, however, the data is extremely unbalanced: Only 4% of the regions report WNV positive detections while 96% do not.

This affects the results of predictive models and special handling is necessary. The inclusion of socioeconomic information on the part of the human host is surely essential. Prompting these questions whether the environment of man, including land use zoning, urban development intensity (especially new urban area density),

and overall economic situation (measured not just financially but from all angles: there are indicators that try to take in income, expenditure, investing practices) plays a role in West Nile Virus transmission. The influences for these socio-economic factors on someone living in habitations within 1 kilometer radius will in general differ over very short distances greater than 2 kilometers out from any given point of residence. The XGBoost method was used for the WNV risk model, a gradient boosting approach particularly well-suited to highly dimensional, imbalanced and symbolically labeled datasets. The data set was split temporally to evaluate model performance over different period.

Thus, it was used training data for 2010 to 2019 in order to allow the model to learn from recent infection trends. For testing data from 1950–2009 and 2020–2022 is used to reflect the model's performance both backwards in time, and forward. This approach has the merit of ensuring that the model is reliable in discerning historical patterns and in forecasting future risks. Various strategies are employed by the models to tackle the serious class imbalance.

For one, it uses custom loss functions and class weighting to ensure that the minority class – regions with WNV infections – does not get swamped by the majority class. Hyperparameter tuning is performed to optimize the way decision trees are structured, enhancing the model's ability to find rare events. Furthermore, the scale position weight parameter in XGBoost is adjusted to enhance sensitivity to WNV-positive cases. Together, all these strategies help the model to better differentiate high-risk regions – even in cases where positive cases are rare.

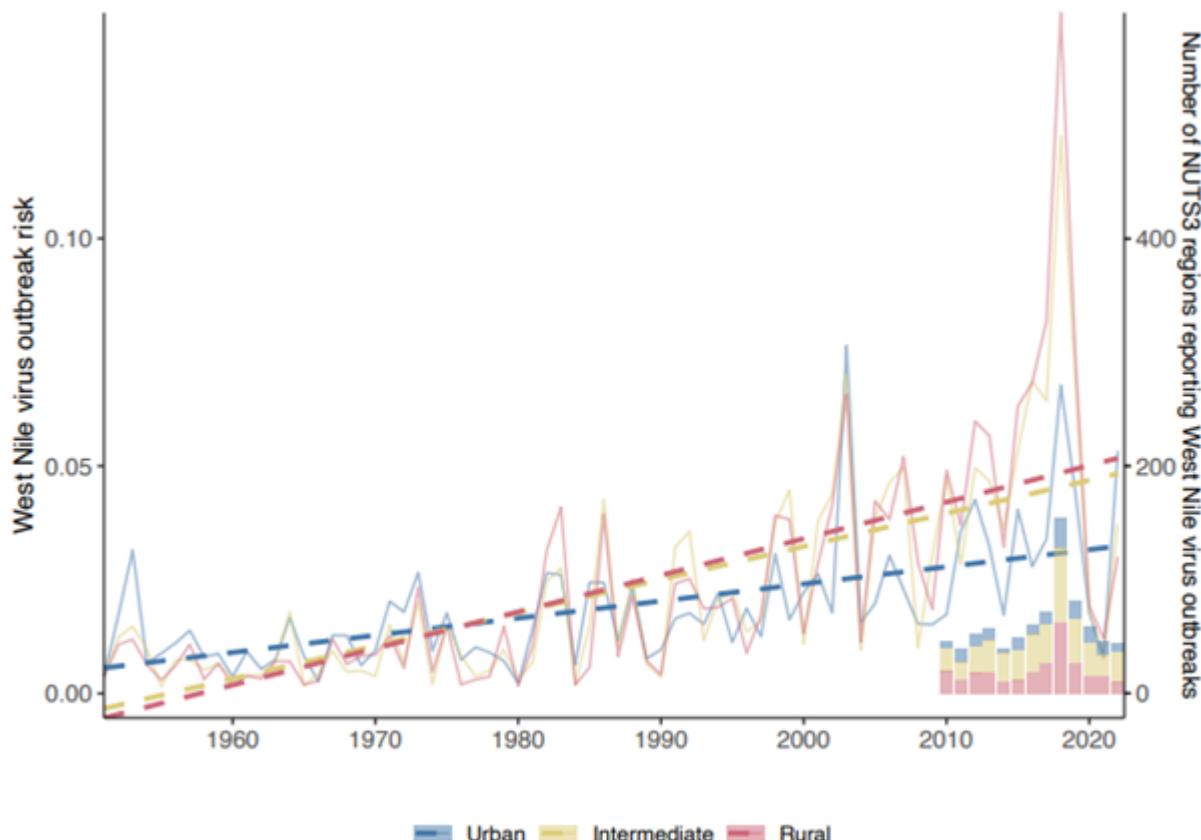


FIGURE 7.4: Predicted West Nile Virus (WNV) outbreak risk from 1950 to 2022 across urban, intermediate, and rural areas. Adapted from Van Daalen, et al. (2024).

7.6.2 Results

The model outputs confirm that WNV outbreak risk is increasing across Europe, with rural regions being most affected. This finding aligns with previous observations that rural and intermediate areas are more prone to outbreaks due to favorable environmental conditions for mosquito breeding and virus transmission. The model effectively captures historical outbreak trends, providing evidence for its reliability in forecasting future WNV risk. Climate change and related environmental factors appear to play a critical role in shaping these risks.

Results further demonstrate a consistent rise in WNV risk across the decades from 1950 to 2022. The sharp increase in predicted risk after the year 2000 supports the hypothesis that rising temperatures and altered precipitation patterns have contributed to an expanded range of the virus. Rural and semi-urban regions show particularly elevated risks, suggesting that these areas may require targeted surveillance and intervention.

Importantly, model predictions closely align with observed data, strengthening the credibility of the XGBoost model. By successfully reproducing known outbreak distributions, the model not only validates its approach but also highlights the strong connection between climatic variables and WNV transmission. The combination of temperature, precipitation, and bioclimatic indicators, along with socioeconomic context, emerges as a powerful framework for understanding viral spread.

Beyond modeling accuracy, the findings have important implications for public health planning. They highlight the need for enhanced disease surveillance and the development of early-warning systems in high-risk regions. Furthermore, climate-adaptive intervention strategies should be prioritized to address the growing influence of environmental change on vector-borne diseases. By linking long-term climate trends to outbreak risk, the study underscores the urgency of proactive and climate-informed public health measures

7.7 Conclusion

This study has explored the intersection of climate change and public health by examining two critical dimensions of climate-related health risks in Europe: heat-related mortality and the spread of West Nile Virus (WNV). Drawing on a combination of statistical modeling and machine learning techniques, the analysis aimed to capture the nonlinear, delayed, and regionally varying nature of these impacts.

In the first part of the study, we discussed heat-related mortality, drawing on findings from previous literature that analyzed temperature–mortality relationships using quasi-Poisson regression combined with distributed lag non-linear modeling (DLNM). This approach, as shown in the work of Gasparini et al. ([Gasparini et al., 2015](#)), allows for the estimation of both immediate and delayed mortality effects of high temperatures. These findings highlight the increased health burden associated with heat, particularly in Southern and Western Europe, and among vulnerable subgroups such as the elderly and women. The analysis illustrates how mortality patterns can evolve over the course of a heatwave, underscoring the need for timely public health interventions.

In the second part of the study, we assessed the spatial risk of WNV transmission using XGBoost, a tree-based gradient boosting algorithm well-suited for high-dimensional and imbalanced data. By incorporating climate, bioclimatic, and socioeconomic variables, the model effectively identified regions at elevated risk and aligned well with historical outbreak trends. The results show a rising trend in WNV risk, especially after 2000, with rural and semi-urban areas emerging as key hotspots. These findings reinforce the importance of integrating climate data into infectious disease monitoring and forecasting systems.

Together, these two case studies illustrate the value of combining traditional epidemiological approaches with modern machine learning tools to study climate-sensitive health outcomes. They also underscore the growing urgency of public health adaptation in response to climate variability. As global temperatures continue to rise, the capacity to detect, predict, and respond to climate-related health threats will become increasingly vital.

Future research should build upon this work by incorporating additional climate-sensitive diseases, refining spatial and temporal resolutions, and extending the analysis to non-European contexts. Moreover, the integration of early warning systems with public health infrastructure will be essential for mitigating the adverse effects of climate change on human health. By advancing our understanding of climate-health interactions, this study contributes to the broader goal of building more resilient and adaptive public health systems in the face of global environmental change.



8

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The authors of this work take full responsibilities for its content.

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³<http://www.en.uni-muenchen.de/index.html>



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