

# Analysing Conceptual Climate Models with Monte Carlo Basin Bifurcation Analysis (MCBB)

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Joint work with Frank Hellmann

Based on M. Gelbrecht, J. Kurths, F. Hellmann: “Monte Carlo Basin Bifurcation Analysis”

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## PAPER

# Monte Carlo basin bifurcation analysis

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**Keywords:** nonlinear dynamics, complex systems, bifurcation, basin stability

## Abstract

Many high-dimensional complex systems exhibit an enormously complex landscape of possible asymptotic states. Here, we present a numerical approach geared towards analyzing such systems. It is situated between the classical analysis with macroscopic order parameters and a more thorough,

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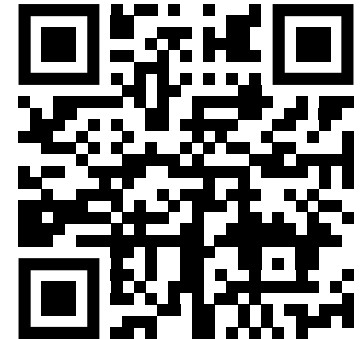
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<https://github.com/maximilian-gelbrecht/MCBB.jl>

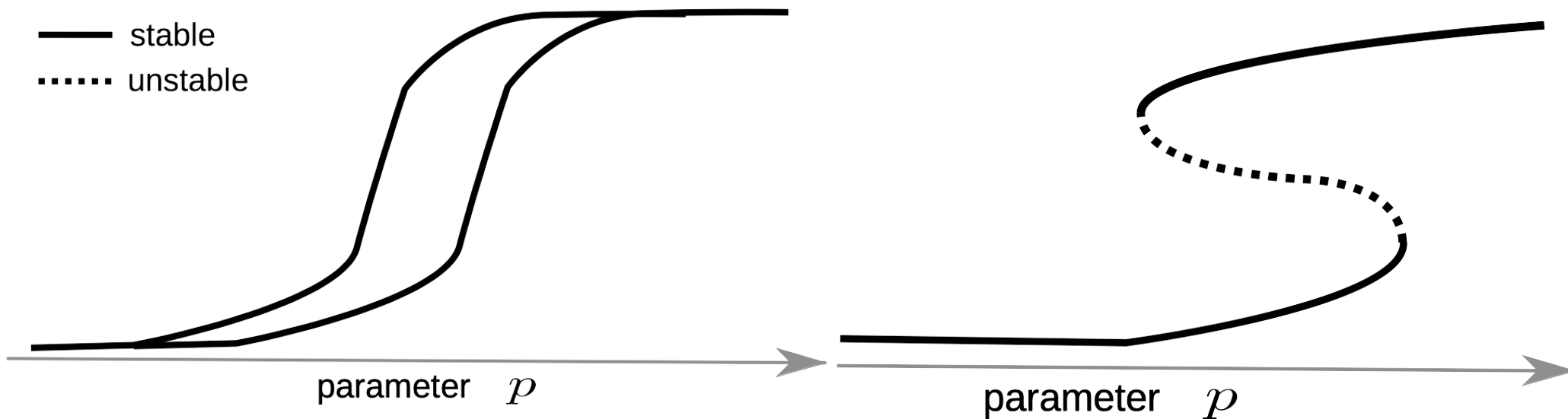


The screenshot shows the GitHub interface for the repository 'maximilian-gelbrecht / MCBB.jl'. At the top, there's a navigation bar with 'Pull requests', 'Issues', 'Marketplace', and 'Explore'. Below the repository name, there are statistics: 0 Watch, 1 Star, and 0 Fork. The main section shows the repository name 'Monte Carlo Basin Bifurcation Analysis' with an 'Edit' button. Below this, there's a summary of the repository: 503 commits, 6 branches, 0 packages, 0 releases, 1 environment, and 2 contributors. A 'View license' button is also present. The 'Branch: master' dropdown is set to 'master', and there's a 'New pull request' button. Below this, there's a table of files and folders:

File/Folder	Description	Last Commit
docs	typo in URL	23 days ago
paper	uploaded scripts for paper	2 months ago
src	hiddenparvar now also with all IC gen options	2 months ago
test	added utility constructor for multi dim setups	3 months ago
.gitignore	more docs	16 months ago

# Motivation

- **Multistability** is a universal phenomenon of complex systems
- Magnetism, human brain, gene expression networks, human perception, power grids, climate systems and many more exhibit multistable regimes
- Volume of the basin of attraction often interesting as well



- For high-dimensional systems a traditional bifurcation analysis is often challenging
- Often one is also interested in classes of asymptotic states instead of every single possible asymptotic state
- **Aim:**
  - Fill gap between thorough bifurcation analysis and macroscopic order parameters
  - Learn classes of similar attractors that collectively have the largest basin of attraction
  - Understand how their *basin volumes change as a function of the parameters*
  - Get insights into the dynamics of these classes of asymptotic states
  - Apply it to climate dynamics

- **Idea:**
  - Combine a sampling based approach with a clustering analysis
  - Don't compare the high-dimensional trajectory tails with each other directly but with per-dimension measures

# Algorithm

## Given:

system $\dot{\mathbf{x}} = F(\mathbf{x}, t; \mathbf{p})$ or $\mathbf{x}_{n+1} = F(\mathbf{x}_n, \mathbf{x}_{n-1}, \dots; \mathbf{p})$ with system dimension $N_d$	A set of $N_m$ statistics $\{\mathcal{S}_i\}$ on the components $\mathbb{R}^{N_t} \rightarrow \mathbb{R}$ (e.g. mean and variance)	Distribution $\mathcal{U}_{IC}$ of the initial conditions and parameters $\mathcal{U}_p$
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sample  $N$  initial conditions and  $N$  parameters from  $\mathcal{U}_{IC}$  and  $\mathcal{U}_p$

for every sample  $i \in [1, N]$

    solve system for a long trajectory, only save the tail  $\mathbf{x}(t; p)$

    for every system dimension  $i_d \in [1, N_d]$  and for every statistic  $i_m \in [1, N_m]$

        compute matrix of statistics  $S_{i,i_d,i_m} = \mathcal{S}_i(x_{i_d})$

**Obtained:**  $N$   $(N_d \times N_s)$ -matrices  $\mathbf{S}_i$

compute  $(N \times N)$  distance matrix  $\mathbf{D}$  of all  $\mathbf{S}_i$  to each other

Density-based clustering of  $\mathbf{D}$  (e.g. DBSCAN)

analyse cluster memberships and measures for each cluster dependent on  $\mathbf{p}$

# Algorithm

## Given:

system $\dot{\mathbf{x}} = F(\mathbf{x}, t; \mathbf{p})$ or $\mathbf{x}_{n+1} = F(\mathbf{x}_n, \mathbf{x}_{n-1}, \dots; \mathbf{p})$ with system dimension $N_d$	A set of $N_m$ statistics $\{\mathcal{S}_i\}$ on the components $\mathbb{R}^{N_d} \rightarrow \mathbb{R}$ (e.g. mean and variance)	Distribution $\mathcal{U}_{IC}$ of the initial conditions and parameters $\mathcal{U}_p$
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for every sample  $i \in [1, N]$

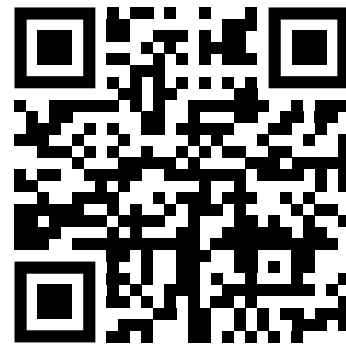
**more detailed information in the paper**

**<https://doi.org/10.1088/1367-2630/ab7a05>**

solve system for a long trajectory, only save

for every system dimension  $i_d \in [1, N_d]$

compute matrix of statistics  $S_{i,i_d,i_m}$



**Obtained:**  $N$   $(N_d \times N_s)$ -matrices  $S_i$

compute  $(N \times N)$  distance matrix  $\mathbf{D}$  of all  $S_i$  to each other

Density-based clustering of  $\mathbf{D}$  (e.g. DBSCAN)

analyse cluster memberships and measures for each cluster dependent on  $\mathbf{p}$

# Software Implementation: MCBB.jl

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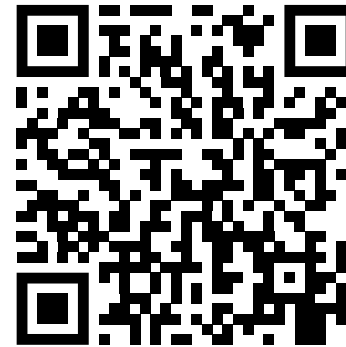
- Open Source software package MCBB.jl available
- Julia lang, easy to read and write, fast programming language
- Excellent state-of-the-art differential equations solvers (thanks to DifferentialEquations.jl)

## GitHub repository



<https://github.com/maximilian-gelbrecht/MCBB.jl>

## Documentation



<https://maximilian-gelbrecht.github.io/MCBB.jl/dev/>



# Applications

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- MCBB is a modular, flexible method suitable for many different kinds of mid- to high-dimensional complex systems
- Maps, ODEs, ...
- **Examples**
  - **Dodds-Watts model of social and biological contagion**
  - Kuramoto network
  - Stuart-Landau oscillator network
- conceptual climate models (work in progress)
- **modified Lorenz 96 model** (here)

} paper

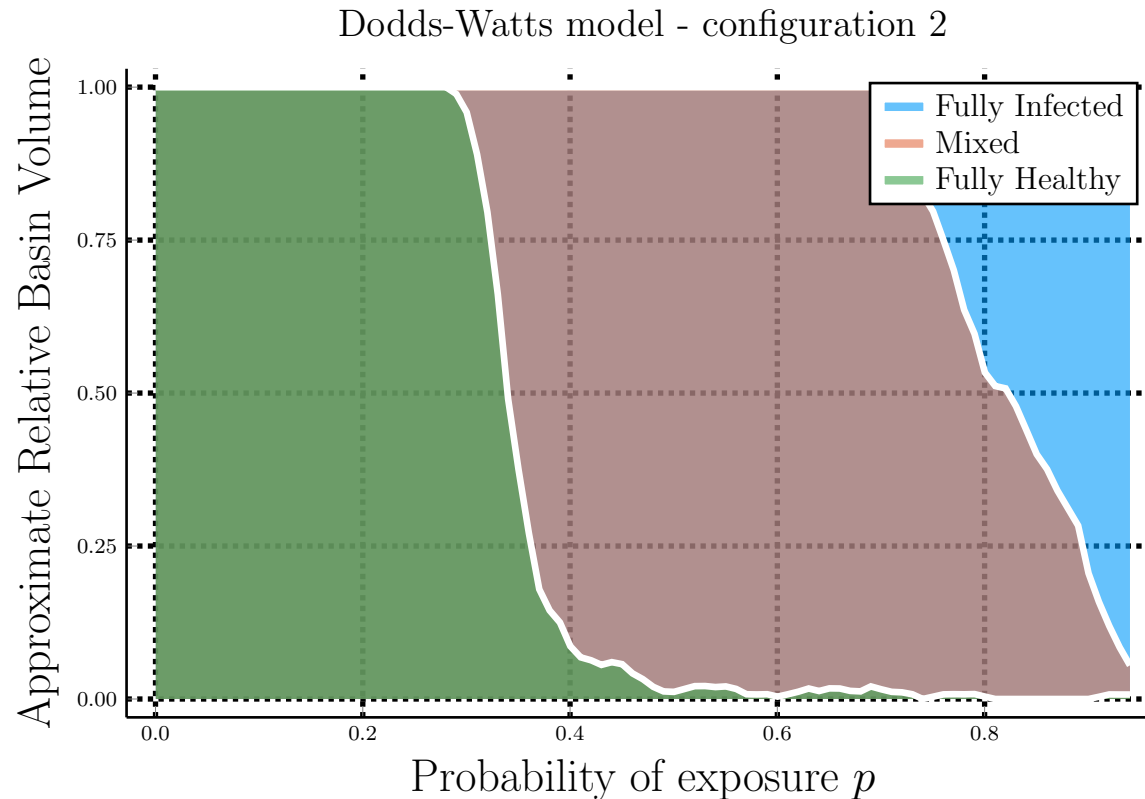
# Application: Dodds Watts model

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- Dodds-Watts model of social and biological contagion
- Generalisation of SI(RS) models from epidemiology
- Population with  $N$  Individuals that can be either **S**usceptible, **I**nfectious or **R**ecovered
- Dodds-Watts model introduces a dosage memory into these models (all details see Dodds, Watts [arXiv:1705.10783](https://arxiv.org/abs/1705.10783) )

# Application: Dodds Watts model - MCBB results

- Area in the plot corresponds to the basin volume



- Additional tools to identify the dynamics of the individual classes of the asymptotic states in the paper / library
- Here, coexistence of states where the population is fully healthy (green), only some individuals are infected (red) and fully infected (blue)

# Application: Lorenz 96

- 1-Layer Lorenz 96 model coupled to a simple EBM
- add an additional “wobble” to the EBM to invoke more stable states than the regular cold / warm state
- add noise -> SDE

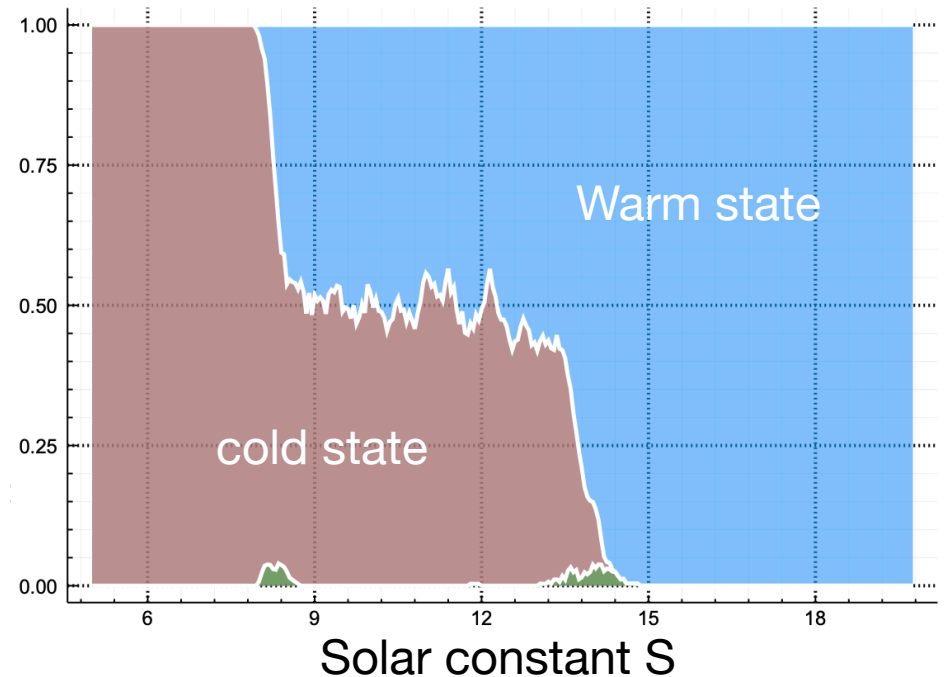
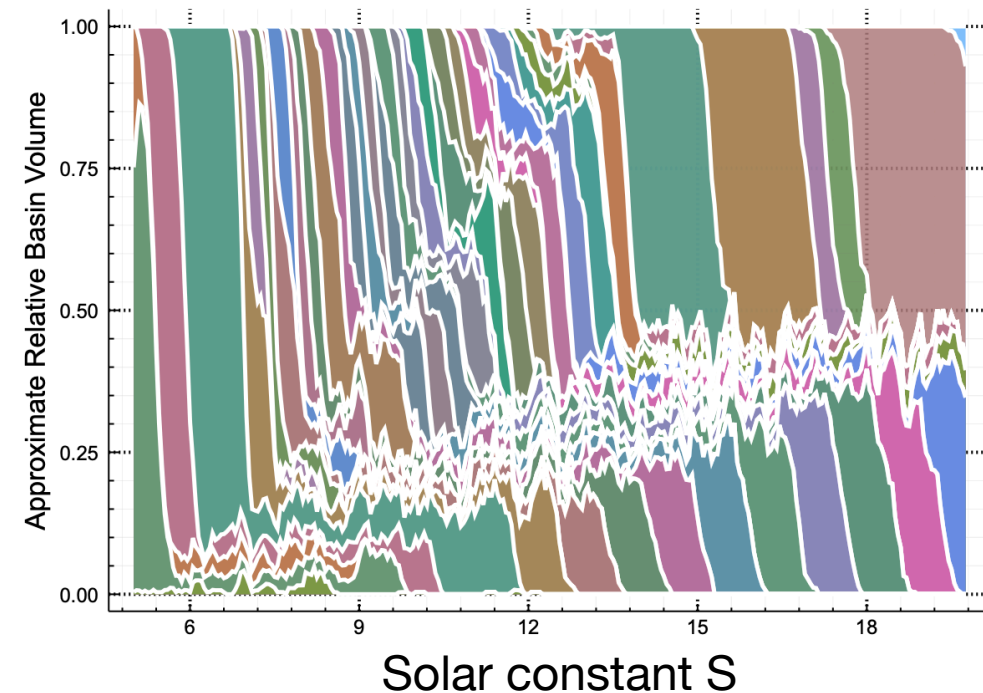
$$dX_k = \underbrace{(-X_{k-2}X_{k-1} + X_{k+1}X_{k+1} - X_k + F)}_{\text{Lorenz 96}} dt + \underbrace{\sigma_X dW}_{\text{Noise}}$$

$$dF = (EBM(\mathbf{X}, F) + A \cdot \underbrace{\sin(\omega(F - F_0))}_{\text{Additional Wiggle}}) dt + \underbrace{\sigma_{EBM} dW}_{\text{Noise}}$$

# Application: Lorenz 96

$$\sigma_{EBM} = 0$$

$$\sigma_{EBM} \text{ large}$$



- Sinus wiggle introduces many additional stable states
- For large noise amplitudes only the “deepest” states in the EBM are relevant and the sinus-wiggle is not important anymore
- Further analysis with MCBB possible (and also experiments with two parameters)