



# 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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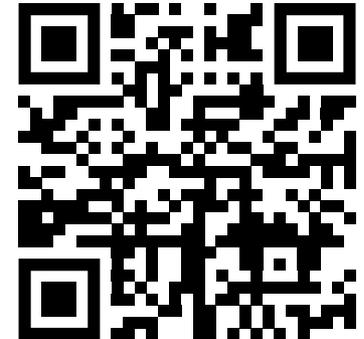
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## Abstract

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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[https://  
github.com/  
maximilian-  
gelbrecht/MCBB.jl](https://github.com/maximilian-gelbrecht/MCBB.jl)



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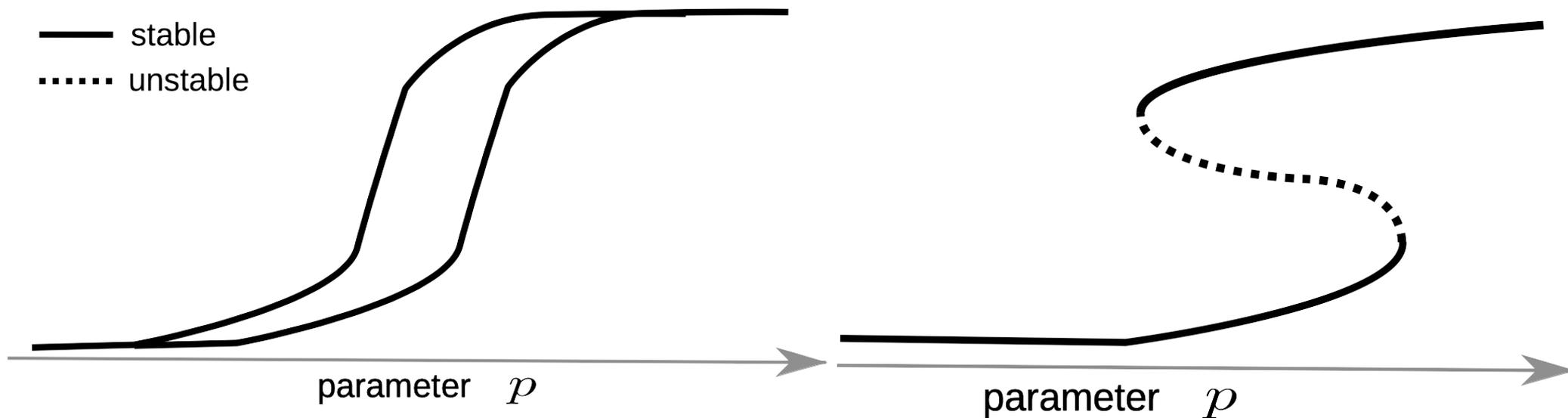
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File	Description	Time
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paper	uploaded scripts for paper	2 months ago
src	hiddenparvar now also with all IC gen options	2 months ago
test	added utility constructor for multi dim setups	3 months ago
.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



# Motivation

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- 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

# Method

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- **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  $\mathcal{U}_p$

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  $\{\mathcal{S}_i\}$   
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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  $S_{i,i_d,i_m}$



statistic  $i_m \in [1, N_m]$

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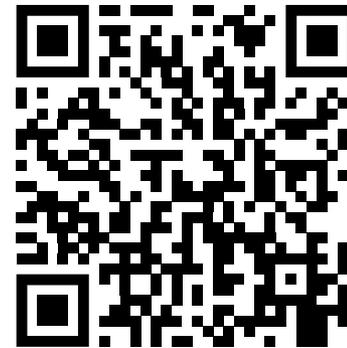
- Open Source software package MCBB.jl available
- Julia lang, easy to read and write, fast programming language
- Excellent state-of-the-art differential equations solvers (thanks to DifferentialEquations.jl)

## 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

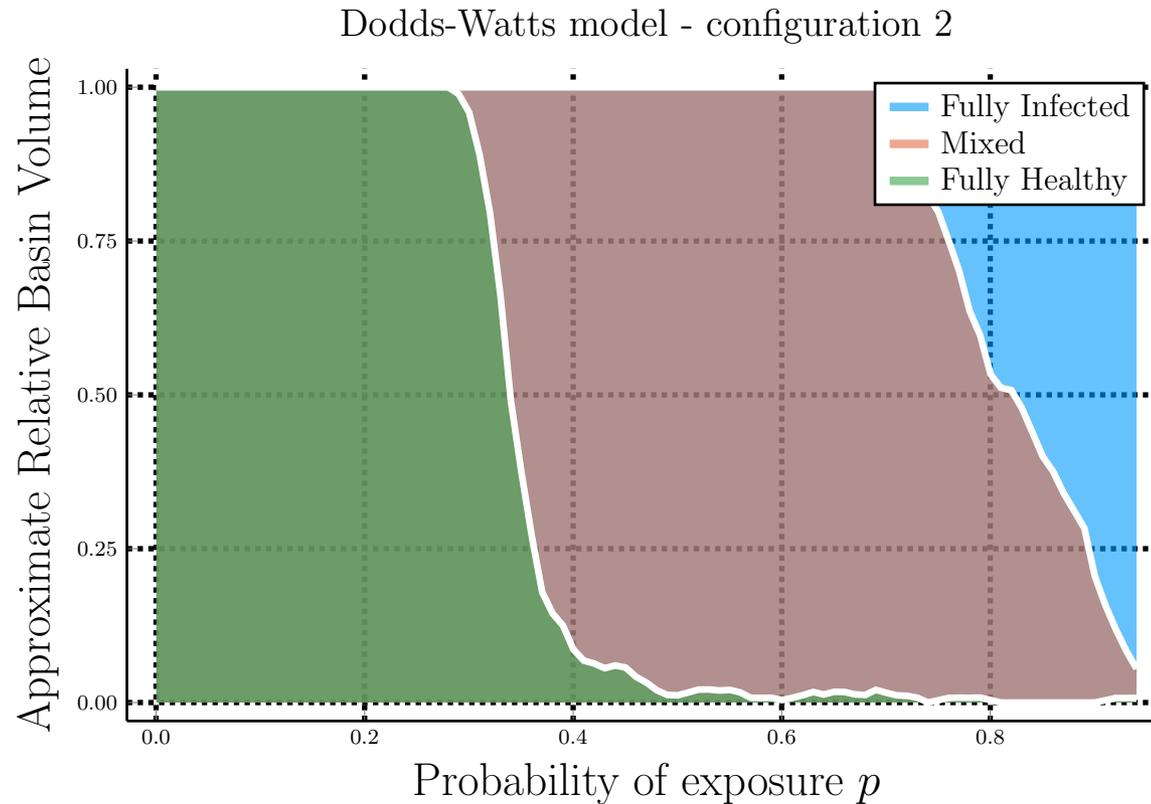
# Application: Dodds Watts model

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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**nfected 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 “wiggle” to the EBM to invoke more stable states than the regular cold / warm state
- add noise -> SDE

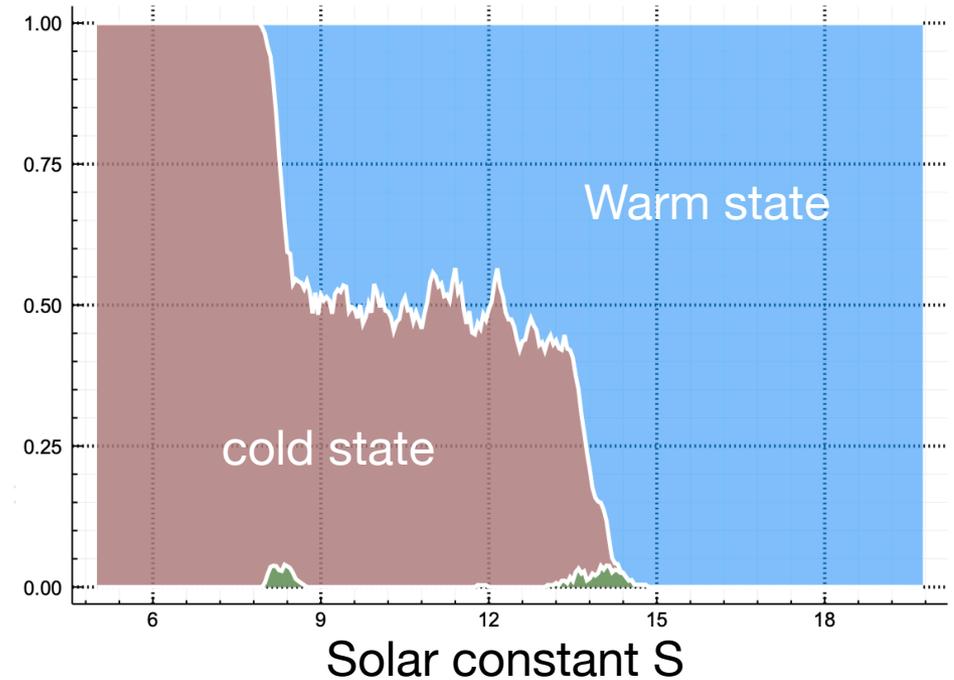
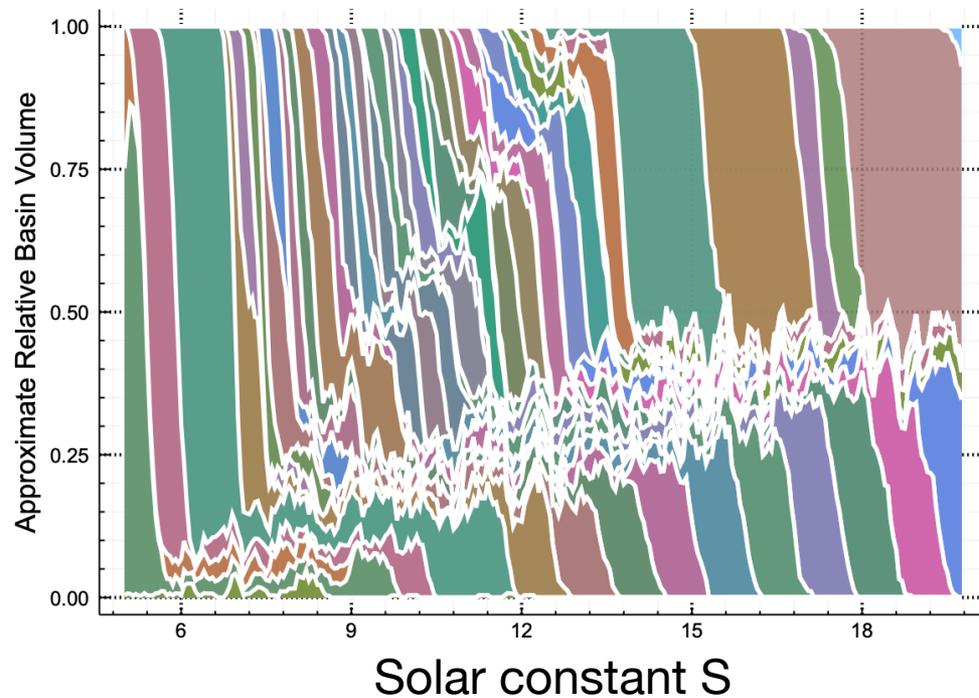
$$dX_k = \underbrace{(-X_{k-2}X_{k-1} + X_{k+1}X_{k+1} - X_k + F)}_{\text{Lorenz 96}} dt + \underbrace{\sigma_X dW}_{\text{Noise}}$$

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

# Application: Lorenz 96

$$\sigma_{EBM} = 0$$

$\sigma_{EBM}$  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)