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UNIVERSITY OF ZAGREB
FACULTY OF SCIENCE
DEPARTMENT OF MATHEMATICS

Luka Jovanović

**PHYSICS-INSPIRED DYNAMICAL
SYSTEMS FOR OPTIMIZATION**

Master's thesis

Advisors:
assoc. prof. dr. sc. Ilya Gogić,
dr. sc. Daniel Ebler

Zagreb, December 2023

SVEUČILIŠTE U ZAGREBU
PRIRODOSLOVNO–MATEMATIČKI FAKULTET
MATEMATIČKI ODSJEK

Luka Jovanović

**Dinamički sustavi za optimizaciju
inspirirani fizikom**

Diplomski rad

Voditelji rada:
dr. sc. Ilya Gogić, izv. prof.;
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Zagreb, prosinac 2023.

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*To my parents and grandparents, to my brothers, to my friends, and to my dear one –
for the support and love they shared with me throughout my studies.*

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Introduction

The quadratic unconstrained binary optimization (QUBO) task is a minimization task defined by

$$\min_{s_1, \dots, s_n \in \{0,1\}} \sum_{i,j=1}^n J_{ij} s_i s_j + \sum_{i=1}^n h_i s_i + c$$

where the coefficients $J_{ij}, h_i, c \in \mathbb{R}$ are given. This simple combinatorial optimization task has many applications in the sense that it can be used to model various real-world problems. These real-world problems include combinatorial optimization problems which arise in finance [1], cluster analysis [2], economic analysis [3], computer-aided layout design [4], integrated chip design [5], physics [5, 6], and many more [7]. Although simple to state, these combinatorial optimization problems are in general very hard to solve if the number of variables is large. One obvious algorithm for solving such a problem would be to try every possible combination $s_i \in \{0, 1\}$, evaluate it, and then take the minimum result found. However, the problem with this algorithm is that the number of possible combinations which need to be evaluated is 2^n which grows exponentially with respect to the number of variables. For example, solving a problem with $n = 100$ variables on a computer which can perform 10^8 multiply/add operations per second would require more than $2^{100} * 100^2 / 10^8 > 10^{26}$ seconds which is more time than the predicted current age of the Universe (according to [8]).

The question which now emerges is whether there exists a significantly faster algorithm which would find the exact solution of QUBO? The short answer to this question is — *probably not*. In fact, QUBO is an archetype of a NP-hard combinatorial optimization problem [7, 9], meaning that there is no polynomial-time algorithm which would solve it, unless $P=NP$ which is suspected to be false [10]. Furthermore, many NP-complete and NP-hard problems, including all of Karp's 21 NP-complete problems, can be reduced to a QUBO problem efficiently [11].

Although being NP-hard, there is a vast number of algorithms for solving QUBO problem exactly. Each of these algorithms exploits some properties of QUBO and provides a smarter way to reach the exact solution than trying all combinations. Nevertheless, their complexity is still non-polynomial and thus they work efficiently for up to a few hundred

variables at most. Many of these algorithms, developed before 2014, can be found in a survey [7].

On the other hand, there is a multitude of heuristic and metaheuristic algorithms for solving QUBO approximately but quickly. These algorithms usually do not have a guarantee of achieving certain accuracy, but their performance is rather based on empirical evidence. These algorithms include variations and adaptations of tabu search [12], simulated annealing [13, 14], genetic algorithms [15], Hopfield neural networks [16], and many others [7]. There is also one notable metaheuristic algorithm called Breakout Local Search (BLS) [17] which provides high quality approximate solutions to the MAX-CUT problem, which is equivalent to QUBO 1.2.6.

Along these classical algorithms designed for running on conventional computers, significant effort was put into building Ising machines – a special type of hardware devices designed for sampling the exact, or high-quality approximate solutions of the QUBO problem [18]. One particular type of Ising machines are quantum computers capable of performing *adiabatic quantum computing* [19]. The working principle is as follows. First, the combinatorial optimization problem is encoded into a Hamiltonian H_P in such a way that the ground state of this Hamiltonian encodes the optimum solution of the combinatorial problem. In order to find the ground state of H_P , one first prepares the quantum state to be the ground state of some initial Hamiltonian H_0 whose ground state is easy to find and construct. Then, the system is *slowly* evolved in time from H_0 to H_P according to $H(t) = (1 - \frac{t}{T})H_0 + \frac{t}{T}H_P$. According to the adiabatic theorem of quantum mechanics [20], if the system is evolved slowly enough, the state will remain the ground state at each time instance. Thus, at the final time instance T , the ground state of the Hamiltonian $H(T) = H_P$ will be prepared. After measurement, it provides the solution of the original combinatorial optimization problem. There is, however, a debate whether this technique would be useful in practice because, in order to satisfy the assumptions of the adiabatic theorem of quantum mechanics, one often finds that the required time for evolution depends exponentially on the problem size [11]. Besides this, one would first need a quantum computer with a large enough number of qubits in order to outperform the existing classical computer architectures. The current state of the art quantum annealers have around 5000 qubits (by D-Wave [21, 22]). However, the topological embedding of these qubits limits the number of variables that can be encoded into the annealer to around tens or hundreds of variables. Thus, existing quantum devices are not able to handle moderate and large instances of problems.

Some other physical devices which have been proposed as Ising machines include a network of coupled optical parametric oscillators [23], electronic oscillators [24], and quantum-mechanical oscillators [25]. Some of these machines have been built and demonstrated good performance on problems of size up to 100,000 variables [26].

Recently, a new paradigm for heuristic approaches has been proposed in [27] and [25]. Instead of building a physical hardware for an Ising machine, one can simulate its be-

behaviour on standard hardware devices such as CPU, GPU, and FPGA. In order to do that, one first writes a set of differential equations which approximately describe the behaviour of an Ising machine. For example, instead of evolving a quantum Hamiltonian on a quantum device, one can define the corresponding classical dynamical system by approximating the expected value of the annihilation operator a , present in the quantum Hamiltonian H , by a complex number $x + iy$ where $x, y \in \mathbb{R}$. This way, one obtains the equations of motion for x and y .

The derived set of differential equations represents a dynamical system which is then further simplified such that it can be efficiently simulated with standard numerical algorithms on a classical computer. This is how Coherent Ising Machine (CIM) [28], Simulated Bifurcation (SB) [29], Simulated Coherent Ising Machine (SimCIM) [30], Ballistic Simulated Bifurcation (bSB) [31], Discrete Simulated Bifurcation (dSB) [31], and some other related algorithms were born. These algorithms are easily parallelized on GPUs or similar hardware devices which enables them to quickly provide high-quality solutions for large instances of QUBO and other combinatorial problems.

The goal of this master's thesis is presenting these physics-inspired dynamical systems, and analyzing them from a mathematical perspective in order to understand their mechanisms for generating high-quality solutions of combinatorial optimization problems. To understand how a continuous dynamical system can provide a solution of the combinatorial optimization problem, consider a state vector which evolves in time $\mathbf{x} : [0, \infty) \rightarrow \mathbb{R}^n$. Taking the sign of each component at some time instance t , $\sigma_i := \text{sign } x_i(t)$ provides a candidate solution $(\sigma_1, \dots, \sigma_n)$ for the QUBO task. The evolution of a continuous dynamical system is determined by its vector field and initial conditions. The vector field is defined at each point in space and determines the velocity $\dot{\mathbf{x}}$ of the system if it passes through that point. The initial conditions $\mathbf{x}_0 \in \mathbb{R}^n$ determine the starting position of the dynamical system $\mathbf{x}(0) = \mathbf{x}_0$. The goal is obtaining a dynamical system which *attracts* various trajectories towards such areas of space that provide high-quality approximate solutions of the QUBO problem, according to the mapping mentioned above $\sigma_i := \text{sign } x_i(t)$.

In chapter 1, a QUBO task is studied. Several other combinatorial optimization problems are presented including MAX-CUT and Traveling Salesman Problem, and their embedding into a QUBO task is given.

In chapter 2, a mathematical theory for dynamical systems and certain other topics are briefly presented, with the main purpose of providing a good understanding of topics in chapter 3.

Chapter 3 is the core chapter as it presents and analyzes these aforementioned physics-inspired dynamical systems and their corresponding numerical simulations which together are used for approximately solving a QUBO problem. These algorithms include Coherent Ising Machine (CIM), Simulated Bifurcation (SB), Simulated Coherent Ising Machine (SimCIM), and Ballistic Simulated Bifurcation (bSB). In order to provide a bigger pic-

ture and the mutual relation between these algorithms, they may be presented in slightly different form than in the original articles.

All of these algorithms have efficient implementations which are run on classical computers or even parallelized architectures such as GPUs and FPGAs. They can be used for fast sampling of high-quality QUBO solutions. The results obtained by these algorithms are presented in the final chapter 4. These results cover the performance of algorithms on a benchmark dataset GSet along with methods used for fine-tuning parameters. Finally, a *dropout* technique is proposed for improving the performance of these algorithms.

Contributions

This thesis focuses on analyzing physics-inspired dynamical systems purely from a formal mathematical perspective. Thus, some arguments mentioned in the original articles are refined here and formalized as much as possible. In order to do that, it was first necessary to extract the relevant existing theorems from the theory of dynamical systems and other fields. Some arguments for explaining the mechanism of these algorithms are only applicable to autonomous dynamical systems. Since dynamical systems of interest are nonautonomous, the author proposes a theorem for connecting nonautonomous system whose vector field changes slowly in time to the corresponding autonomous systems with the vector field frozen in time. This is the theorem 2.1.9.

As proposed in the original articles, the dynamical systems discussed in this thesis are solved approximately by simulating them numerically. In this thesis, the exact solutions of SimCIM and bSB algorithms are derived which is, as far as the author is aware, not yet reported in the literature, but could potentially be useful for further research.

Although momentum has been introduced for SimCIM algorithm, it seems not to be used for CIM. It was observed that adding momentum to CIM improved the solution quality for the QUBO task. During fine-tuning, the momentum had the option of being 0 (original CIM), or some other larger values (for example 0.8, 0.9, 1.0). It indeed turned out to always pick some value larger than 0. In chapter 4, results which compare the performance of CIM with and without momentum have also been provided.

Furthermore, it is shown in chapter 3 that CIM with momentum is in a certain sense a generalization of the SB algorithm. Similarly, SimCIM with momentum is in some sense a generalization of the bSB algorithm, which is even able to exactly reproduce the behaviour of bSB.

Most of these algorithms have already been benchmarked on GSet. However, the thesis provides another independent source of these results. This can be understood as comparing all of these algorithms in a consistent and unified way.

Finally, a new technique for adding a *meaningful noise* was proposed in chapter 4, which we call *dropout*. It works by randomly and temporarily (throughout one iteration of the algorithm) dropping out some vertices – meaning that all of the connections of such vertices are temporarily deleted. This technique was tested on GSet instances and seems to improve the solution quality obtained by CIM, SimCIM, and bSB algorithms. How exactly does this technique enable these algorithms to find better solutions is yet to be researched.

Chapter 1

Quadratic Unconstrained Binary Optimization

The quadratic unconstrained binary optimization (QUBO) problem is a combinatorial optimization problem defined by

$$\min_{s_1, \dots, s_n \in \{0,1\}} \sum_{i,j=1}^n J_{ij} s_i s_j + \sum_{i=1}^n h_i s_i + c \quad (1.1)$$

where coefficients $J_{ij}, h_i, c \in \mathbb{R}$ are given. In other words, the task is to minimize a given quadratic polynomial in n variables over the discrete domain $\{0, 1\}^n$.

We are usually interested not only in finding a minimum but finding an argument which minimizes the function. Also, since finding such an optimum is computationally very hard for large instances, we will be interested in finding as low value as possible and the corresponding argument. The domain $\{-1, +1\}^n$ is called the *search space* while the elements of this set are called *candidate solutions* or *feasible solutions* or simply *solutions*. An *optimum solution* is then the minimizer of the function over the set of feasible solutions - i.e. the best possible solution.

Since $h_i s_i = h_i s_i^2$ for $s_i \in \{0, 1\}$, all terms of degree one can be transformed into quadratic terms. The constant c does not play any role in the minimization task. That being said, an equivalent formulation of the QUBO task is minimizing a homogeneous quadratic polynomial in n variables over the domain $\{0, 1\}^n$,

$$\min_{s_1, \dots, s_n \in \{0,1\}} \sum_{i,j=1}^n J_{ij} s_i s_j \quad (1.2)$$

Coefficients J_{ij} can always be taken symmetrically i.e. such that $J_{ij} = J_{ji}$. Indeed, if they are not symmetric, taking coefficients $\frac{1}{2}(J_{ij} + J_{ji})$ in place of both J_{ij} and J_{ji} will make them symmetric.

As stated in the Introduction part, an obvious brute-force algorithm for solving a QUBO problem is to evaluate the polynomial at all possible candidate solutions $s_i \in \{0, 1\}$ and take the minimum among them. However, the problem with this algorithm is that there are 2^n combinations in total, so the search space grows exponentially with respect to the number of variables. Depending on the computer's performance, this will work efficiently for a number of variables of order 10^1 . However, already for $n = 100$ this becomes too many operations to perform on a computer within any reasonable amount of time.

The key point here is that, since QUBO is a NP-hard problem [7], there is no polynomial-time algorithm which would solve it exactly (unless $P=NP$ which is suspected to be false [10]). Thus, there is probably no algorithm for solving QUBO exactly which would have significantly lower computational complexity than exponential one.

Nevertheless, there are many algorithms which solve QUBO exactly by leveraging various properties of the problem and using different techniques. Many of these methods are listed in [7]. However, all of these methods work effectively for number of vertices up to few hundreds at most.

For larger instances of QUBO, various heuristic and metaheuristic algorithms have been developed which provide some relatively high-quality approximate solution. A lot of such algorithms can be found in [7] as well. However, there is one special kind of heuristic algorithms recently proposed. These algorithms are based on physics-inspired dynamical systems. These algorithms are the core of this thesis and are presented in chapter 3.

It will be shown in 1.2 that QUBO problem is in fact equivalent to a specific graph problem called MAX-CUT. There are some approximation algorithms in the literature for solving MAX-CUT which are thus directly applicable to solving QUBO.

The question which emerges is whether there exists a polynomial-time algorithm which would provide arbitrarily good approximate solution. In fact, QUBO problem is APX-hard, which under the assumption $P \neq NP$ implies that there is no polynomial-time approximation scheme (PTAS) for it [32, 33, 34]. In other words, no polynomial-time algorithm can guarantee to provide a solution which is as close to the optimal solution as we would require in advance (if $P \neq NP$).

To address the question of how close to the optimal solution can some algorithm get with a guarantee, there is an article by Goemans and Williamson [35] proposing a randomized algorithm for solving MAX-CUT problem (see section 1.2) based on semidefinite programming which always provides solutions with expected value at least 0.87856 times the optimal solution. MAX-CUT is equivalent to QUBO 1.2.6 in such a way that there exists a mapping from candidate solutions of one problem to candidate solutions of the other problem, which preserves value obtained by these candidate solutions. Thus, algorithm which provides solutions with expected value at least 0.87856 times the optimal solution of the MAX-CUT problem could also be used to provide solutions of the QUBO problem with the same accuracy.

If the unique games conjecture [36] is true, this is the best possible approximation ratio which can be guaranteed for MAX-CUT (see [34, 36], subsection 1.2 and remark 1.2.6), and thus for QUBO as well.

Equivalent forms

Let $Q'(s_1, \dots, s_n) = \sum_{i,j=1}^n J'_{ij} s_i s_j + \sum_{i=1}^n h'_i s_i + c'$ be a quadratic polynomial with the domain $\{0, 1\}^n$ as in 1.1. By taking a linear change of variables given by $\sigma_i = 2s_i - 1$, a new quadratic polynomial $Q(\sigma_1, \dots, \sigma_n)$ is given over the domain $\{-1, +1\}^n$, satisfying $Q'(s_1, \dots, s_n) = Q(\sigma_1, \dots, \sigma_n)$. Thus, the QUBO task 1.1 has an equivalent form

$$\min_{\sigma_1, \dots, \sigma_n \in \{-1, +1\}} \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i + c \quad (1.3)$$

for some new coefficients J_{ij}, h_i, c .

Although equivalent, each of these formulations has its own benefit. As it will be seen in the following sections, some other combinatorial optimization problems can be embedded into QUBO task. Sometimes it will be easier to think of an embedding in 1.1 form and sometimes in 1.3 form. Form 1.3 might be more suitable for analysis because the domain $\{-1, +1\}^n$ consists of elements with equal norm. When solving QUBO task on a computer, the algorithm will often be implemented either as Q over the domain $\{-1, +1\}^n$ or as Q' over the domain $\{0, 1\}^n$. It is thus useful to have an explicit relation between coefficients in Q and Q' . That relation is given by

$$\begin{array}{ll} Q: \{-1, +1\}^n \rightarrow \mathbb{R} & Q': \{0, 1\}^n \rightarrow \mathbb{R} \\ J_{ij} = \frac{1}{4} J'_{ij} & J'_{ij} = 4J_{ij} \\ h_i = \frac{1}{4} \sum_{j=1}^n (J'_{ij} + J'_{ji}) + \frac{1}{2} h'_i & h'_i = -2 \sum_{j=1}^n (J_{ij} + J_{ji}) + 2h_i \\ c = \frac{1}{4} \sum_{i,j=1}^n J'_{ij} + \frac{1}{2} \sum_{i=1}^n h'_i + c' & c'_i = \sum_{i,j=1}^n J_{ij} - \sum_{i=1}^n h_i + c \end{array} \quad (1.4)$$

Another equivalent form emerges if minimization task is replaced with maximization. Indeed,

$$\max_{\sigma_1, \dots, \sigma_n \in \{-1, +1\}} Q(\sigma_1, \dots, \sigma_n) = - \min_{\sigma_1, \dots, \sigma_n \in \{-1, +1\}} -Q(\sigma_1, \dots, \sigma_n) \quad (1.5)$$

and $-Q$ is still a quadratic polynomial.

Homogenizing

Let us assume that a quadratic polynomial

$$Q(\sigma_1, \dots, \sigma_n) = \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i + c$$

is given over the domain $\{-1, +1\}^n$. For the minimization task, the constant term c is irrelevant so here we assume without loss of generality that $c = 0$. By introducing one auxiliary variable $\sigma_{n+1} \in \{-1, +1\}$ we define a polynomial

$$P(\sigma_1, \dots, \sigma_n, \sigma_{n+1}) := \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i \sigma_{n+1}$$

which is homogeneous of degree 2. For each configuration $(\sigma_1, \dots, \sigma_n) \in \{-1, +1\}^n$, there are two corresponding configurations in $\{-1, +1\}^{n+1}$.

Those are $(\sigma_1, \dots, \sigma_n, +1)$ and $(-\sigma_1, \dots, -\sigma_n, -1)$. It follows that ($c = 0$ by assumption)

$$Q(\sigma_1, \dots, \sigma_n) = P(\sigma_1, \dots, \sigma_n, +1) = P(-\sigma_1, \dots, -\sigma_n, -1) \quad (1.6)$$

Thus, finding a minimum of a (nonhomogeneous) quadratic polynomial with n variables over the domain $\{-1, +1\}^n$ is equivalent to finding a minimum of the corresponding homogeneous quadratic polynomial with $n+1$ variables over the domain $\{-1, +1\}^{n+1}$. Because of (1.6), not only the minima are in correspondence, but all values are. Although increasing the number of variables by 1 does not play any important role in a sense of computational complexity, the performance of some algorithms might be affected because the *extra* variable σ_{n+1} is (possibly) coupled to all other variables. If some algorithm leverages the coupling structure of the original problem (for example sparsity of J), the new coupling might lose this structure after introducing σ_{n+1} (sparsity might be compromised because σ_{n+1} could be coupled to all other variables), thus causing the algorithm to drop in performance. Adding this extra variable could also often cause issues with convergence of physics-inspired dynamical systems and corresponding algorithms presented in chapter 3.

Nevertheless, we will not be concerned with the effect of adding this *extra* variable in the rest of the thesis, but rather take this theoretical result as a justification for solving and analyzing mostly the homogeneous case

$$\min_{\sigma_1, \dots, \sigma_n \in \{-1, +1\}} \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j \quad (1.7)$$

Matrix-Vector notation

Given a quadratic polynomial with real coefficients

$$Q(\sigma_1, \dots, \sigma_n) = \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i + c$$

it's evaluation can be written in matrix-vector notation as follows. Put all coefficients J_{ij} into the matrix $J \in M_n(\mathbb{R})$, coefficients h_i into the vector $\mathbf{h} \in \mathbb{R}^n$ and arguments σ_i into the vector $\boldsymbol{\sigma} \in \{-1, +1\}^n$. Then we write

$$\sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i + c = Q(\sigma_1, \dots, \sigma_n) = Q(\boldsymbol{\sigma}) = \boldsymbol{\sigma}^T J \boldsymbol{\sigma} + \boldsymbol{\sigma}^T \mathbf{h} + c$$

and so the minimization task is rewritten as

$$\min_{\boldsymbol{\sigma} \in \{-1, +1\}^n} \boldsymbol{\sigma}^T J \boldsymbol{\sigma} + \boldsymbol{\sigma}^T \mathbf{h} + c \quad (1.8)$$

This transition between vectors, matrices and their coefficients, will be used in what follows without further noticing.

1.1 Analysis of the problem

Bounds with coefficients

Define the following matrix and vector norms

$$\begin{aligned} \|J\|_1 &:= \sum_{i,j=1}^n |J_{ij}| & \|\mathbf{h}\|_1 &:= \sum_{i=1}^n |h_i| \\ \|J\|_\infty &:= \max_{i,j=1,\dots,n} |J_{ij}| & \|\mathbf{h}\|_\infty &:= \max_{i=1,\dots,n} |h_i| \end{aligned}$$

Then we have the following bounds for the value in QUBO

$$\left| \sum_{i,j=1}^n J_{ij} \sigma_i \sigma_j + \sum_{i=1}^n h_i \sigma_i + c \right| \leq \|J\|_1 + \|\mathbf{h}\|_1 + |c| \leq n^2 \|J\|_\infty + n \|\mathbf{h}\|_\infty + |c| \quad (1.9)$$

Bounds with the largest and the smallest eigenvalue

Let $\lambda_1, \dots, \lambda_n$ be the eigenvalues of the symmetric coupling matrix J sorted in descending ascending order, meaning that $\lambda_1 \leq \lambda_2 \leq \dots \leq \lambda_n$. Let v_1, \dots, v_n be the orthonormal basis

which diagonalizes the coupling matrix J , corresponding to eigenvalues $\lambda_1, \dots, \lambda_n$. Then we have the following:

$$\begin{aligned}\boldsymbol{\sigma} &= \sum_i \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma} \\ \boldsymbol{\sigma}^T J \boldsymbol{\sigma} &= \left(\sum_i \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma} \right)^T J \left(\sum_i \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma} \right) = \sum_{i,j} \boldsymbol{\sigma}^T \mathbf{v}_i \mathbf{v}_i^T J \mathbf{v}_j \mathbf{v}_j^T \boldsymbol{\sigma} = \sum_i \lambda_i \boldsymbol{\sigma}^T \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma}\end{aligned}$$

Thus,

$$\begin{aligned}\boldsymbol{\sigma}^T J \boldsymbol{\sigma} &\leq \max\{\lambda_1, \dots, \lambda_n\} \sum_i \boldsymbol{\sigma}^T \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma} = \max\{\lambda_1, \dots, \lambda_n\} \boldsymbol{\sigma}^T \boldsymbol{\sigma} \\ \boldsymbol{\sigma}^T J \boldsymbol{\sigma} &\geq \min\{\lambda_1, \dots, \lambda_n\} \sum_i \boldsymbol{\sigma}^T \mathbf{v}_i \mathbf{v}_i^T \boldsymbol{\sigma} = \min\{\lambda_1, \dots, \lambda_n\} \boldsymbol{\sigma}^T \boldsymbol{\sigma}\end{aligned}$$

Since for all vectors $\boldsymbol{\sigma} \in \{-1, +1\}^n$ the quantity $\boldsymbol{\sigma}^T \boldsymbol{\sigma}$ is equals n , we obtain the bounds

$$n \cdot \min\{\lambda_1, \dots, \lambda_n\} \leq \boldsymbol{\sigma}^T J \boldsymbol{\sigma} \leq n \cdot \max\{\lambda_1, \dots, \lambda_n\} \quad (1.10)$$

Degree of synchronization

Let $Q(\mathbf{x}) = -\mathbf{x}^T J \mathbf{x}$ and consider the following minimization QUBO task

$$\min_{\boldsymbol{\sigma} \in \{-1, +1\}^n} Q(\boldsymbol{\sigma}) \quad (1.11)$$

We have that $\lambda_{\min}(-J) = -\lambda_{\max}(J)$ with eigenvectors corresponding to each other. According to the previous subsection, if the eigenvector \mathbf{v}_{\max} corresponding to the largest eigenvalue $\lambda_{\max}(J)$ consists only of components with magnitude 1 i.e. $\mathbf{v}_{\max} \in \{-1, +1\}^n$, then the solution of the QUBO problem (1.11) is exactly the vector \mathbf{v}_{\max} . Intuitively this should also hold in case that \mathbf{v}_{\max} is very close to some point from the feasible set $\{-1, +1\}^n$ (ignoring the scale and considering all vectors to be normalized). This is formally justified in the following result from [37].

Definition 1.1.1. *Let \mathbf{x} be an arbitrary vector in R^n . The feasible solution for QUBO task corresponding to this vector is considered to be $\boldsymbol{\sigma} = \text{sign}(\mathbf{x})$. We define the degree of synchronization of \mathbf{x} to be*

$$\alpha^2(\mathbf{x}) = \left(\frac{\mathbf{x}^T \boldsymbol{\sigma}}{\|\mathbf{x}\| \|\boldsymbol{\sigma}\|} \right)^2 \quad (1.12)$$

where $\boldsymbol{\sigma} = \text{sign}(\mathbf{x})$.

Theorem 1.1.2. Consider the minimization QUBO task (1.11). Denote with Q_0 the optimum (minimum) solution for this minimization task. Denote with Q_1 the second smallest value if it exists, otherwise let $Q_1 = Q_0$. Denote with $\Delta Q = Q_1 - Q_0$. Let $\lambda_{\max} = \lambda_1 \geq \dots \geq \lambda_n = \lambda_{\min}$ be all eigenvalues of J sorted in descending order. Assume that the largest eigenvalue's multiplicity is 1 (i.e. $\lambda_1 \neq \lambda_2$). Denote with \mathbf{v}_{\max} a normalized eigenvector corresponding to λ_{\max} .

If for the degree of synchronization of vector \mathbf{v}_{\max} it holds

$$\alpha^2(\mathbf{v}_{\max}) \geq 1 - \frac{\Delta Q}{n(\lambda_{\max} - \lambda_{\min})} \quad (1.13)$$

then the corresponding solution $\sigma = \text{sign}(\mathbf{v}_{\max})$ is the optimum solution of QUBO task (1.11).

Proof. If $\Delta Q = 0$ we have that all feasible solutions are minima so there is nothing to prove. Thus, let us assume that $\Delta Q > 0$.

$\alpha^2(\mathbf{v}_{\max}) = 1$ is equivalent to the fact that σ is proportional to \mathbf{v}_{\max} . In this case, σ reaches the bound value of (1.10) so σ must be the optimum solution and the proof is done.

Consider the opposite case where σ is not proportional to \mathbf{v}_{\max} . Since $\lambda_i < \lambda_{\max}, \forall i > 1$ we have that

$$Q_0 > \min_{\|\mathbf{x}\|^2=n} Q(\mathbf{x}) = -n\lambda_{\max} \quad (1.14)$$

which implies

$$Q_1 > -n\lambda_{\max} + \Delta Q \quad (1.15)$$

Using the fact that $J = \sum_i \lambda_i \mathbf{v}_i \mathbf{v}_i^T$ we have that

$$\begin{aligned} Q(\sigma) &= -\sigma^T J \sigma = -\sigma^T \left(\sum_i (\lambda_i - \lambda_{\min}) \mathbf{v}_i \mathbf{v}_i^T \right) \sigma - \sum_i \lambda_{\min} \sigma^T \mathbf{v}_i \mathbf{v}_i^T \sigma \\ &= -\sum_i (\lambda_i - \lambda_{\min}) (\mathbf{v}_i^T \sigma)^2 - n\lambda_{\min} \\ &\leq -(\lambda_{\max} - \lambda_{\min}) (\mathbf{v}_{\max}^T \sigma)^2 - n\lambda_{\min} \end{aligned} \quad (1.16)$$

where for obtaining the last inequality we have dropped all but one summation terms. Since $n\alpha^2(\mathbf{v}_{\max}) = (\mathbf{v}_{\max}^T \sigma)^2$, by assumption of the theorem we have that

$$(\mathbf{v}_{\max}^T \sigma)^2 \geq n - \frac{\Delta Q}{\lambda_{\max} - \lambda_{\min}} \quad (1.17)$$

Combining these we get

$$Q(\sigma) \leq \Delta Q - n(\lambda_{\max} - \lambda_{\min}) - n\lambda_{\min} = \Delta Q - n\lambda_{\max} < Q_1 \quad (1.18)$$

Thus, $Q(\sigma) = Q_0$ i.e. σ is the optimum solution of the QUBO task. \square

1-opt solution

Let $Q(\sigma_1, \dots, \sigma_n) = \sum_{i,j=1}^n J_{ij}\sigma_i\sigma_j + \sum_{i=1}^n h_i\sigma_i + C$.

Definition 1.1.3. For the maximization QUBO task

$$\max_{\sigma \in \{-1, +1\}^n} Q(\sigma), \quad (1.19)$$

vector $\sigma \in \{-1, +1\}^n$ is called a 1-opt solution if for every $i = 1, \dots, n$ we have that

$$\Delta_i Q(\sigma_1, \dots, \sigma_n) \leq 0 \quad (1.20)$$

where

$$\Delta_i Q(\sigma_1, \dots, \sigma_n) := Q(\sigma_1, \dots, \sigma_{i-1}, -\sigma_i, \sigma_{i+1}, \dots, \sigma_n) - Q(\sigma_1, \dots, \sigma_{i-1}, \sigma_i, \sigma_{i+1}, \dots, \sigma_n) \quad (1.21)$$

That is, for a 1-opt solution, switching the value of a single variable cannot produce a better solution.

By replacing \leq with \geq in equation (1.21) we get the definition of a 1-opt solution for the minimization QUBO task.

In a similar way we can define a k -opt solution for integer $k > 1$.

Writing out further the formula for $\Delta_i Q$ and using the fact that $J_{ij} = J_{ji}$, $J_{ii} = 0$, we get

$$\Delta_i Q(\sigma_1, \dots, \sigma_n) = -2 \sum_{j=1}^n (J_{ij} + J_{ji})\sigma_i\sigma_j - 2h_i\sigma_i = -4 \sum_{j=1}^n J_{ij}\sigma_i\sigma_j - 2h_i\sigma_i \quad (1.22)$$

1.2 Applications

Many combinatorial optimization problems can be embedded into a QUBO problem. Those include all of Karp's 21 NP-complete problems. [11] This embedding refers to the fact that the original combinatorial optimization problem can be reformulated into a QUBO task in such a way that the solution of QUBO task encodes the solution of the original optimization problem. In the following subsections, some examples of these embeddings are provided. Namely, those are MAX-CUT problem, Number partitioning problem, and Traveling salesman problem. Many more examples can be found in [11].

There is one notable difference among these embeddings of different combinatorial optimization problems. As an example, let us consider the Traveling salesman problem (TSP). In order to embed this problem into the QUBO formulation, we need to introduce some *ancillary* variables. So, for a TSP problem with n variables (cities), we would need

to solve the QUBO task with n^2 variables. Nevertheless, the number of variables required for embedding all of Karp's 21 NP-complete problems is at most cubic with respect to the number of variables in the original problem [11]. Additionally to ancillary variables, we will need to impose some *constraints* on those variables i.e. we will have to solve a *constrained* quadratic binary optimization task. These constraints can often be embedded into the QUBO task as well, but this might significantly affect the algorithm's performance and the quality of the solution.

Handling constraints

In this section we will describe a way of handling constraints by introducing *penalty* functions. Consider a minimization task

$$\min_{\mathbf{s} \in A} Q(s_1, \dots, s_n) \quad (1.23)$$

where $A \subset \{0, 1\}^n$ is some allowable set over which we want to minimize a quadratic polynomial Q . Denote with A^c its complement. Let us suppose that $P(s_1, \dots, s_n)$ is a quadratic polynomial which is precisely equal zero on the set A , while it is strictly greater than zero outside of A . Now, let us define

$$Q'(\mathbf{s}) := Q(\mathbf{s}) + C \cdot P(\mathbf{s}) \quad (1.24)$$

for some $C > 0$.

First, we have not changed the values obtained when evaluating solutions from the allowable set. That is, $Q'(\mathbf{s}) = Q(\mathbf{s}), \forall \mathbf{s} \in A$.

Second, by choosing $C > 0$ to be large enough, every configuration which is not allowed evaluates Q' to something which is too large to be the minimum of Q . That is, $\forall \mathbf{s} \in A^c, Q'(\mathbf{s}) > \min_{\mathbf{s} \in A} Q(\mathbf{s})$

Third, Q' is still a quadratic polynomial.

This means that solving an unconstrained (QUBO) problem

$$\min_{\mathbf{s} \in \{0, 1\}^n} Q'(s_1, \dots, s_n) \quad (1.25)$$

provides us with a solution of the constrained problem (1.23).

Example 1.2.1. Suppose we want to minimize some quadratic polynomial $Q(s_1, \dots, s_n)$ over the discrete set $\{0, 1\}^n$, but we are only interested in balanced configurations, i.e. configurations (s_1, \dots, s_n) which contain equal amount of 0's as 1's (assume that n is even).

For the penalty function

$$P(s_1, \dots, s_n) := \left(\frac{n}{2} - \sum_{i=1}^n s_i \right)^2 \quad (1.26)$$

we have that $P(s_1, \dots, s_n) \geq 0$ for all configurations, but $P(s_1, \dots, s_n) = 0$ if and only if the configuration is balanced. Furthermore, P is a quadratic polynomial.

By selecting $C > 0$ appropriately large, and defining $Q' := Q + C \cdot P$, we have that Q' is a quadratic polynomial whose minimum over the whole set $\{0, 1\}^n$ corresponds to the minimum of Q over the set of all possible balanced configurations.

Similar procedures can be done for handling other types of constraints. If we have multiple constraints which need to be satisfied at the same time, we can just sum the penalties which correspond to each constraint separately.

Although theoretically perfectly valid, handling constraints with penalty functions can impose problems in practice because algorithms for solving the QUBO often do not necessarily provide *the minimum*, but rather an *approximate* solution. Choosing a constant $C > 0$ (from above example) which is too large might force the algorithm to produce solutions which satisfy the constraints, but are completely unaware of the actual value of Q which we are trying to minimize. On the other hand, making the value $C > 0$ too small might allow the algorithm to produce solutions which violate constraints. Therefore, we would definitely need to fine tune the value C . However, for approximate solvers this sometimes just does not work well (for example if many constraints need to be satisfied at the same time).

It is very problem-specific and also algorithm-specific to determine how to efficiently embed the constraints – whether there are some alternative penalty functions which might perform better, and whether there are some completely different embeddings which could perform better.

MAX-CUT

This section is based on [34].

MAX-CUT is a combinatorial optimization problem on graphs which can be directly modeled with QUBO task. Consider an undirected weighted graph $G = (V, E)$ with no self-loops and no multiple edges between vertices. V denotes the set of vertices, E the set of edges. Each edge $e \in E$ is a triplet of the form $e = (a, b, J_{ab})$ where vertices $a, b \in V$ are connected with an edge with weight J_{ab} . If the number of vertices is $n = |V|$ then all weights can be stored in a symmetric adjacency matrix $J \in M_n(\mathbb{R})$. The fact that vertices $a, b \in V$ are connected with an edge having weight w is represented in adjacency matrix by $J_{ab} = J_{ba} = w$. Throughout this section, only graphs of this type will be considered.

In order to describe the MAX-CUT problem we need a precise definition of what the *graph cut* means.

Definition 1.2.2. Given a graph $G = (V, E)$ with adjacency matrix J , a (graph) cut is a bipartition $\mathcal{P} = \{A, B\}$ of the vertices (meaning that $A, B \subset V; A \cap B = \emptyset; A \cup B = V$).

The value of a certain graph cut $\mathcal{P} = \{A, B\}$ is a number

$$\sum_{a \in A, b \in B} J_{ab} \quad (1.27)$$

A cut is maximum if no other cut produces greater value.

The MAX-CUT problem is then straightforward

Problem 1.2.3 (MAX-CUT problem). *Given a graph G , determine its maximum cut.*

MAX-CUT problem is a NP-hard problem.

There is a similar version with *yes-no* solution

Problem 1.2.4 (MAX-CUT problem, binary). *Given a graph G and a value k , determine whether or not there exists a cut with value at least k in G .*

This version is a NP-complete problem and is on the Karp's list of 21 NP-complete problems.

Remark 1.2.5. *A dual definition of "MIN-CUT" of a graph could be provided and so the problem of finding a MIN-CUT would be equivalent to the problem of finding a MAX-CUT by just taking the opposite sign of each edge weight. However, in the literature a MIN-CUT is usually considered only for weighted graphs whose weights are strictly positive and with the requirement that the cut is not trivial i.e. $A, B \neq \emptyset$ (otherwise a trivial cut $A = V, B = \emptyset$ would always be the solution). Under these restrictions, a MIN-CUT problem becomes P instead of NP-hard. It is in fact a dual problem to the max-flow problem i.e. finding the maximum flow from source to sink (source and sink are newly added vertices) which is as well solvable in polynomial time. Here, only graphs with arbitrary real weights are considered, so in order to stay aligned with nomenclature in the literature, only MAX-CUT problem will be considered, as defined in 1.2.2.*

Let us formulate the MAX-CUT problem as a QUBO task. Suppose a graph $G = (V, E)$ is given, with adjacency matrix J and the set of vertices being $V = \{1, \dots, n\}$. Assume that the graph is weighted, has no self-loops, and is undirected. For a given cut $\mathcal{P} = \{A, B\}$, define a vector $\sigma = (\sigma_1, \dots, \sigma_n) \in \{-1, +1\}^n$ such that $\sigma_i = -1$ if $i \in A$, and $\sigma_i = +1$ if $i \in B$. It is clear that this mapping is a one-to-one correspondence between all possible cuts and all vectors $\{-1, +1\}^n$. Furthermore, define

$$Q(\sigma) := \frac{1}{4} \sum_{i,j=1}^n J_{ij}(1 - \sigma_i \sigma_j) \quad (1.28)$$

which is a quadratic polynomial over the domain $\{-1, +1\}^n$. For a particular cut $\mathcal{P} = \{A, B\}$, plugging in σ corresponding to \mathcal{P} , the value of $Q(\sigma)$ becomes precisely the value of that

cut. To see this, note that $(1 - \sigma_i \sigma_j)$ is equals to 2 in case that i and j belong to different parts of the partition (i.e. $i \in A, j \in B$ or $i \in B, j \in A$) while it is 0 otherwise.

The conclusion is that finding a MAX-CUT of the graph can be embedded into solving a QUBO task given by

$$\max_{\sigma \in \{-1, +1\}^n} Q(\sigma)$$

with Q defined by 1.28.

Note that up to a constant term (which is irrelevant for maximization task), Q is a homogeneous polynomial. The number of variables that we required to embed a MAX-CUT problem into a QUBO task is $O(n)$, where n is the number of vertices in the graph.

On the other hand, consider an arbitrary QUBO problem given by

$$\max_{\sigma \in \{-1, +1\}^n} \sigma^T J \sigma + \sigma^T \mathbf{h} + c \quad (1.29)$$

As discussed in subsection 1, this is equivalent to solving the QUBO task with a corresponding homogeneous polynomial with $n + 1$ variables. In this correspondence, all candidate solution values are preserved. Thus, we may assume that $\mathbf{h} = \mathbf{0}$ and $c = 0$ without loss of generality. Let us define an undirected graph $G = (V, E)$ with adjacency matrix given by $-4J$. Up to a constant term, all the cuts of this graph and their values correspond to the evaluation of the given quadratic polynomial, as seen in 1.28. Thus, solving a QUBO problem can be embedded into a MAX-CUT problem.

These two statements are summarized in the following remark.

Remark 1.2.6. *The QUBO problem and the MAX-CUT problem are equivalent in a sense that given an arbitrary instance of one of these problems with n variables, one can formulate an instance of the other problem with at most $n + 1 = O(n)$ variables in such a way that there is a mutual correspondence between candidate solutions as well as between values produced by these candidate solutions.*

Thus, statements and techniques for solving MAX-CUT can almost always be applied to QUBO, and vice versa.

Number Partitioning

Number partitioning is the following combinatorial problem.

Problem 1.2.7 (Number Partitioning). *A sequence of n real numbers a_1, \dots, a_n is given. Determine whether or not there exists a bipartition $\mathcal{P} = \{A, B\}$ of those numbers ($A, B \subset \{1, \dots, n\}; A \cup B = \{1, \dots, n\}; A \cap B = \emptyset$) such that $\sum_{i \in A} a_i = \sum_{i \in B} a_i$. If such a bipartition exists, determine it.*

For example, can we divide a set of assets with values a_1, \dots, a_n fairly between two people?

This problem is known to be NP-complete [11].

To formulate this problem in terms of a QUBO task, let us define the following quadratic polynomial with n variables over the domain $\{-1, +1\}^n$.

$$Q(\sigma_1, \dots, \sigma_n) := \left(\sum_{i=1}^n a_i \sigma_i \right)^2 \quad (1.30)$$

There is a one-to-one correspondence between each bipartition $\mathcal{P} = \{A, B\}$ and the domain $\{-1, +1\}^n$ given by the rule: $\sigma_i = -1$ if $i \in A$, $\sigma_i = +1$ if $i \in B$. It is clear that a solution of the number partitioning problem exists if and only if there exists a configuration $(\sigma_1, \dots, \sigma_n) \in \{-1, +1\}^n$ that evaluates to $Q(\sigma_1, \dots, \sigma_n) = 0$. Additionally, those solutions are in correspondence with minimizers of Q by the described rule. Evaluation $Q(\sigma_1, \dots, \sigma_n) = 0$ means that the minimum of Q has been found and it is equals to 0. Thus, minimizing Q is equivalent to solving the Number partitioning problem.

If there does not exist a solution to the Number partitioning problem, then one might want to find a bipartition which is the *closest* possible to the fair partition. Minimizing Q again solves this problem.

The polynomial Q is generally not homogeneous. The number of variables required for encoding the number partitioning problem into a QUBO task is linear with respect to the problem size, i.e. $\mathcal{O}(n)$, where n is the number of assets that are being partitioned.

Traveling Salesman Problem

There are many variants of the traveling salesman problem. We will present the following version and it's QUBO formulation. Many other variants can be formulated as QUBO task in a similar fashion.

Problem 1.2.8 (Traveling Salesman Problem (TSP)). *A weighted directed complete graph $G = (V, E)$ is given. Edges of the form $(a, b, W_{ab}) \in E$ represent that there is an edge from a to b with weight W_{ab} . All weights are positive. The fact that G is directed here refers to the possibility that $W_{ab} \neq W_{ba}$. Let us assume that vertices are numbered $V = (1, \dots, n)$. A tour v_1, \dots, v_n is a permutation of vertices V representing the order for visiting each vertex, that is $v_1, v_2, \dots, v_n, v_1$ because one wants to return to the starting vertex.*

The task is to find a tour which minimizes the sum of traversed edges i.e. the sum

$$\sum_{i=1}^{n-1} W_{v_i, v_{i+1}} + W_{v_n, v_1}$$

which will often be referred to as tour length.

One example of the traveling salesman problem is: how to plan the route for a delivery vehicle given that it needs to visit each of the specified locations once, while minimizing the total path length and thus minimizing the transport costs.

Let us present a QUBO formulation of the TSP. Since the route length does not depend on the starting vertex, we may choose vertex $a = 1$ as the starting one. Define $(n - 1)^2$ variables $s_{i,u} \in \{0, 1\}$ where $i \in \{2, \dots, n\}$ represents the step and $u \in V \setminus \{a\}$ represents the vertex to be visited in i -th step. $s_{i,u} = 1$ means that we want to visit vertex u at step i , while $s_{i,u} = 0$ otherwise. Since we want a permutation of vertices, we want exactly one $s_{i,u} = 1$ per turn i , and all others to be $s_{i,u} = 0$. The same should hold per each vertex $u \in V \setminus \{a\}$. Let us thus define the following quadratic polynomial

$$Q_A(\mathbf{s}) := \sum_{i=2}^n (1 - \sum_{u \in V \setminus \{a\}} s_{i,u})^2 + \sum_{u \in V \setminus \{a\}} (1 - \sum_{i=2}^n s_{i,u})^2 \quad (1.31)$$

We have that $Q_A(\mathbf{s}) = 0$ if and only if \mathbf{s} represents a permutation of vertices as described above, and $Q_A(\mathbf{s}) \geq 1$ otherwise.

Now define the following quadratic polynomial

$$Q_B(\mathbf{s}) := \sum_{u \in V \setminus \{a\}} W_{a,u} s_{2,u} + \sum_{i=3}^n \sum_{u \in V \setminus \{a\}} \sum_{v \in V \setminus \{a\}} W_{u,v} s_{i-1,u} s_{i,v} + \sum_{u \in V \setminus \{a\}} W_{u,a} s_{n,u} \quad (1.32)$$

We have that $Q_B \geq 0$. Considering only those configurations \mathbf{s} that already represent a valid permutation as described above, the value $Q_B(\mathbf{s})$ is precisely the length of the tour corresponding to the given permutation. Now define

$$Q(\mathbf{s}) := C \cdot Q_A(\mathbf{s}) + Q_B(\mathbf{s}) \quad (1.33)$$

for properly chosen constant $C > 0$. It should be chosen in such a way that any configuration \mathbf{s} which violates the above requirements for producing a valid permutation, evaluates to suboptimal $Q(\mathbf{s})$. For example, for any \mathbf{s} corresponding to a valid permutation of vertices, $Q_B(\mathbf{s}) \leq n \cdot \max_{u,v \in V} W_{u,v}$. Choosing any $C > n \cdot \max_{u,v \in V} W_{u,v}$ as a constant will be sufficient. Indeed, for any \mathbf{s} which does not correspond to a valid permutation, the value $Q(\mathbf{s})$ will be higher than every possible tour length, so any valid route encoding \mathbf{s}' will provide $Q(\mathbf{s}') < Q(\mathbf{s})$. Minimizing Q thus simplifies to minimizing Q_B over the set of all valid permutations. This is exactly minimizing the tour length from the traveling salesman problem.

Since Q is a quadratic polynomial over the domain $\{0, 1\}^{(n-1)^2}$, we have showed that solving a traveling salesman problem can be embedded into solving a minimization QUBO task. The number of variables required for embedding a traveling salesman problem into a QUBO task is $\mathcal{O}(n^2)$ where n is the number of vertices in the graph.

Chapter 2

Preliminary theory

This chapter is a brief recapitulation of certain well-known topics in mathematics. Its first purpose is to give a mathematical background for dynamical systems, along with a few examples which will be essential for understanding the behaviour of dynamical systems in chapter 3. Its second purpose is to serve as a reference for some arguments used in chapter 3.

In terms of notation, a time derivative is denoted by a dot, $\dot{x} = \frac{dx}{dt}$. Column vectors in \mathbb{R}^n are denoted by bold letters, while their components are denoted with regular letters and

a subscript denoting their index of component $\mathbf{x} = \begin{bmatrix} x_1 \\ \vdots \\ x_n \end{bmatrix} = (x_1, \dots, x_n)^T$. The superscript

T denotes a transpose. For a vector function $\mathbf{f} : \Omega \subset \mathbb{R}^n \rightarrow \mathbb{R}^n$, its *matrix derivative* or *differential* is considered to be a matrix of its partial derivatives arranged as

$$\frac{\partial \mathbf{f}}{\partial \mathbf{x}} = \begin{bmatrix} \frac{\partial f_1}{\partial x_1} & \cdots & \frac{\partial f_1}{\partial x_n} \\ \vdots & \ddots & \vdots \\ \frac{\partial f_n}{\partial x_1} & \cdots & \frac{\partial f_n}{\partial x_n} \end{bmatrix}$$

So, for a scalar function $E : \mathbb{R}^n \rightarrow \mathbb{R}$, $\frac{\partial E}{\partial \mathbf{x}}$ is a row vector. Its *second differential* (also called *Hessian matrix*) is considered to be a matrix

$$\frac{\partial^2 E}{\partial \mathbf{x}^2} = \begin{bmatrix} \frac{\partial^2 E}{\partial x_1 \partial x_1} & \cdots & \frac{\partial^2 E}{\partial x_1 \partial x_n} \\ \vdots & \ddots & \vdots \\ \frac{\partial^2 E}{\partial x_n \partial x_1} & \cdots & \frac{\partial^2 E}{\partial x_n \partial x_n} \end{bmatrix}$$

Throughout the thesis, the second differential will always be taken of C^2 functions, so the matrix $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ will always be symmetric due to the Schwarz's theorem.

Matrix derivations of higher order and shape could be defined as well, but will not be used in the thesis.

2.1 Dynamical systems

In this section, the results and concepts from the theory of dynamical systems are briefly given. All systems discussed are continuous dynamical systems meaning that the time is continuous and they are governed by ordinary differential equations.

Let $I \subset \mathbb{R}$, $U \subset \mathbb{R}^n$ be open sets, and $\mathbf{f} \in C^1(I \times U; \mathbb{R}^n)$. A (*continuous*) *dynamical system* is an ordinary differential equation of the form

$$\dot{\mathbf{x}}(t) = \mathbf{f}(t, \mathbf{x}) \quad (2.1)$$

where \mathbf{f} is a given vector field. When *initial condition* $\mathbf{x}(0) = \mathbf{x}_0$ is specified, we call such system

$$\begin{cases} \dot{\mathbf{x}}(t) = \mathbf{f}(t, \mathbf{x}) \\ \mathbf{x}(0) = \mathbf{x}_0 \end{cases} \quad (2.2)$$

an *initial value problem*.

The following two theorems are fundamental for dynamical systems in a sense that given a vector field, and the initial condition, the trajectory of the dynamical system exists and is uniquely determined. Moreover, this solution can be extended in time until it either reaches the domain boundary or blows up to infinity.

Theorem 2.1.1 (Existence-Uniqueness). [38] *Let $t_0 \in I \subset \mathbb{R}$, $\mathbf{x}_0 \in U \subset \mathbb{R}^n$ be open sets, and $\mathbf{f} \in C^1(I \times U; \mathbb{R}^n)$. Then there exist open subsets $t_0 \in I_0 \subset I$, $\mathbf{x}_0 \in U_0 \subset U$ and a function $\phi : I_0 \times I_0 \times U_0 \rightarrow \mathbb{R}^n$ such that for each $(t_1, \mathbf{x}_1) \in I_0 \times U_0$, the function $t \mapsto \phi(t, t_1, \mathbf{x}_1)$ is the unique solution defined on I_0 of the initial value problem (2.2).*

Theorem 2.1.2 (Extension). [38] *Let $t_0 \in I \subset \mathbb{R}$, $\mathbf{x}_0 \in U \subset \mathbb{R}^n$ be open sets, and $\mathbf{f} \in C^1(I \times U; \mathbb{R}^n)$. Let $\langle \alpha, \beta \rangle$ be the maximal open interval where the solution \mathbf{x} of dynamical system (2.2) exists, with $-\infty \leq \alpha \leq \beta \leq \infty$. Then either $\|\mathbf{x}(t)\|$ approaches ∞ or $\mathbf{x}(t)$ approaches the boundary of U as $t \rightarrow \beta$.*

When \mathbf{f} does not depend on the time variable t , we call such dynamical system an *autonomous dynamical system*. Otherwise, we call it a *nonautonomous dynamical system*.

When $\mathbf{f}(t, \mathbf{x}) = A\mathbf{x}$ for some constant matrix $A \in M_n(\mathbb{R})$, we say that the dynamical system is *linear*. Otherwise we call it a *nonlinear dynamical system*.

Remark 2.1.3. *Nonautonomous dynamical systems are often referred to as a generalization of autonomous ones, so if we do not care if \mathbf{f} depends on t or not, we may just say a nonautonomous dynamical system.*

Analogously, nonlinear dynamical systems are a generalization of linear ones.

Consider a nonautonomous dynamical system

$$\dot{\mathbf{x}}(t) = \mathbf{f}(t, \mathbf{x}) \quad (2.3)$$

Definition 2.1.4. An equilibrium point of a nonautonomous dynamical system is a point $\mathbf{x}_0 \in \Omega$ which is also a trajectory of the system, meaning that $\mathbf{x}(t) := \mathbf{x}_0$ is the solution of (2.3).

The definition of equilibrium point is equivalent to the fact that $\mathbf{f}(t, \mathbf{x}_0) = \mathbf{0}, \forall t$. When studying a dynamical system, it is often useful to look for its equilibrium points. The character of these points can often tell us about the qualitative behavior of the dynamical system in the neighborhood of these points. The main result about this is the Hartman-Grobman theorem for nonlinear autonomous dynamical systems.

The Hartman-Grobman theorem

This section is based on [38]. Consider a nonlinear autonomous dynamical system

$$\dot{\mathbf{x}}(t) = \mathbf{f}(\mathbf{x}) \quad (2.4)$$

where $\mathbf{f} \in C^1(U; \mathbb{R}^n), U \subset \mathbb{R}^n$.

Definition 2.1.5. For each $\mathbf{x}_0 \in U$ let \mathbf{x} be the solution of 2.4 with initial condition $\mathbf{x}(0) = \mathbf{x}_0$. Let us define a one-parameter family of functions ϕ_t called the flow of dynamical system 2.4 as

$$\phi_t(\mathbf{x}_0) = \mathbf{x}(t)$$

for every t in which \mathbf{x} is defined.

Proposition 2.1.6. The flow ϕ_t of the dynamical system 2.4 has the following properties

a) $\phi_{t+s} = \phi_t \circ \phi_s$

b) $\phi_0(\mathbf{x}) = \mathbf{x}$

whenever both sides of the equation are defined.

Let us assume that the dynamical system 2.4 has an equilibrium point at \mathbf{x}_0 .

The behavior of a nonlinear autonomous dynamical system near an equilibrium point \mathbf{x}_0 is qualitatively the same as the dynamics of the corresponding *linearization* i.e.

$$\dot{\mathbf{x}}(t) = A\mathbf{x} \quad (2.5)$$

where $A = \frac{\partial \mathbf{f}}{\partial \mathbf{x}}(\mathbf{x}_0)$. That is formally stated in the Hartman-Grobman theorem.

Theorem 2.1.7 (The Hartman-Grobman Theorem [38]). Let $\mathbf{f} \in C^1(\mathbb{R}^n; \mathbb{R}^n)$ be a smooth vector field, and let ϕ_t be the flow of the nonlinear system (2.4). Suppose that \mathbf{x}_0 is a hyperbolic equilibrium point, meaning that $\mathbf{f}(\mathbf{x}_0) = \mathbf{0}$ and none of the eigenvalues of $A = \frac{\partial \mathbf{f}}{\partial \mathbf{x}}(\mathbf{x}_0)$ have zero real part. Let ψ_t be the flow of the corresponding linearization (2.5). Then there exist open sets $\mathbf{x}_0 \in U \subset \mathbb{R}^n, \mathbf{0} \in V \subset \mathbb{R}^n$ and a homeomorphism $H : U \rightarrow V$ such that $H(\phi_t(\mathbf{x})) = \psi_t(H(\mathbf{x}))$ whenever $\mathbf{x} \in U$ and both sides of the equation are defined.

A connection between autonomous and nonautonomous systems

In chapter 3, we will be dealing with nonautonomous dynamical systems. However, the vector field \mathbf{f} will *slowly* vary in time, meaning that $\frac{\partial \mathbf{f}}{\partial t}$ has relatively small norm. In this case, if we want to analyze the local (in time) behavior of the dynamical system, it might be tempting to analyze the behaviour of the dynamical system which has its vector field *frozen in time*, i.e. the dynamical system

$$\dot{\mathbf{x}} = \mathbf{f}(t_0, \mathbf{x}) \quad (2.6)$$

where t_0 is some time instance at which the vector field is frozen.

For analyzing the global (in time) behavior of the system, this strategy is inadequate and even misleading, as seen from the following example.

Example 2.1.8. Consider a one-dimensional nonautonomous dynamical system

$$\dot{x}(t) = t - x(t) \quad (2.7)$$

If we were to analyze this system by observing the system which has its vector field frozen in time, that is

$$\dot{x}(t) = t_0 - x(t) \quad (2.8)$$

for some $t_0 \in \mathbb{R}$, we would see that it has equilibrium point at $x = t_0$. The linearization of such system is

$$\dot{x}(t) = -x(t) \quad (2.9)$$

which by Hartman-Grobman theorem (or by solving this directly) would imply that it is an attractive equilibrium. It might be tempting to conclude that the system follows the trajectory given by instantaneous equilibrium points, that is the trajectory $x_{ISP}(t) = t$.

However, the exact solutions of this equation are $x(t) = Ce^{-t} + t - 1$ for some $C \in \mathbb{R}$. It is clear that all of these solutions converge to the solution $x_{DHT}(t) = t - 1$ as $t \rightarrow \infty$.

First of all, $x_{ISP}(t)$ is not even a solution of the above differential equation. Second, and even more important, the system globally behaves in such a way that it converges towards the trajectory $x_{DHT}(t)$, and not $x_{ISP}(t)$.

In [39] they develop the concept of a *distinguished hyperbolic trajectory* which in the above example is precisely $x_{DHT}(t)$, contrary to the *instantaneous stagnation points* which are $x_{ISP}(t)$ in this example.

However, the global (in time) behavior of the dynamical system will not be of great interest in chapter 3, but rather a local one. In this case, analyzing the system as if its vector field was frozen in time makes sense. This concept is formalized in the following.

Let $\mathbf{f} \in C^1(\mathbb{R} \times \mathbb{R}^n; \mathbb{R}^n)$ be a given vector field, $\mathbf{x}_0 \in \mathbb{R}^n$, $t_0 < t_1$. Consider the following dynamical system

$$\begin{cases} \dot{\mathbf{x}}(t) = \mathbf{f}(t, \mathbf{x}(t)) \\ \mathbf{x}(t_0) = \mathbf{x}_0 \end{cases} \quad (2.10)$$

and the corresponding dynamical system with vector field frozen at time instance t_0 , with the same initial conditions.

$$\begin{cases} \dot{\mathbf{y}}(t) = \mathbf{f}(t_0, \mathbf{y}(t)) \\ \mathbf{y}(t_0) = \mathbf{x}_0 \end{cases} \quad (2.11)$$

For vector $\mathbf{v} \in \mathbb{R}^n$ we will denote the norm $\|\mathbf{v}\|_\infty = \max_{i=1, \dots, n} |v_i|$, and for a vector function $\mathbf{g} : \Omega \subset \mathbb{R}^m \rightarrow \mathbb{R}^n$ we will denote the norm

$$\|\mathbf{g}\|_{L^\infty(\Omega)} := \max_{i=1, \dots, n} \|g_i\|_{L^\infty(\Omega)} \quad (2.12)$$

Let us formally define the successive approximations as

$$\begin{aligned} \mathbf{x}_0(t) &:= \mathbf{x}_0 \\ \mathbf{x}_{k+1}(t) &:= \mathbf{x}_0 + \int_{t_0}^t f(\tau, \mathbf{x}_k(\tau)) d\tau \\ \mathbf{y}_0(t) &:= \mathbf{x}_0 \\ \mathbf{y}_{k+1}(t) &:= \mathbf{x}_0 + \int_{t_0}^t f(t_0, \mathbf{y}_k(\tau)) d\tau \end{aligned} \quad (2.13)$$

Theorem 2.1.9 (Connection between autonomous and nonautonomous dynamical systems). *Let $\mathbf{f} \in C^1(\mathbb{R} \times \mathbb{R}^n; \mathbb{R}^n)$, $\mathbf{x}_0 \in \mathbb{R}^n$, $t_0 < t_1$. Assume that all successive approximations (2.13) for dynamical systems (2.10) and (2.11) are well defined and continuous on $[t_0, t_1]$ and that they uniformly converge to solutions*

$$\mathbf{x}_k \xrightarrow{L^\infty([t_0, t_1])} \mathbf{x} \quad (2.14)$$

$$\mathbf{y}_k \xrightarrow{L^\infty([t_0, t_1])} \mathbf{y} \quad (2.15)$$

Assume that $D \subset \mathbb{R}^n$ is a convex compact set which contains both solutions and all successive approximations $\mathbf{x}([t_0, t_1])$, $\mathbf{x}_k([t_0, t_1])$, $\mathbf{y}([t_0, t_1])$, $\mathbf{y}_k([t_0, t_1]) \subset D$, $\forall k$. Let $L > 0$ be a Lipschitz constant in second variable for \mathbf{f} in a sense that

$$\|\mathbf{f}(t, \mathbf{x}) - \mathbf{f}(t, \mathbf{y})\|_\infty \leq L \|\mathbf{x} - \mathbf{y}\|_\infty, \quad \forall t \in [t_0, t_1], \forall \mathbf{x}, \mathbf{y} \in D \quad (2.16)$$

Let

$$M := \left\| \frac{\partial \mathbf{f}}{\partial t} \right\|_{L^\infty([t_0, t_1] \times D)} \quad (2.17)$$

Then,

$$\|\mathbf{x} - \mathbf{y}\|_{L^\infty([t_0, t_1])} \leq \frac{M}{L^2} e^{L(t-t_0)} - \frac{M}{L}(t-t_0) - \frac{M}{L^2}, \quad \forall t \in [t_0, t_1] \quad (2.18)$$

Proof. First, let's show that \mathbf{f} is Lipschitz continuous in second variable with some constant $L > 0$ in a sense of (2.16). By following the proof of proposition 2.3.4 and applying it to each component of \mathbf{f} , we get that f_1, \dots, f_n are Lipschitz continuous in second variable. From here, (2.16) follows easily.

By inductive argument we prove a bound for $\|\mathbf{x}_k(t) - \mathbf{y}_k(t)\|_\infty$. For $k = 0$ we have $\|\mathbf{x}_0(t) - \mathbf{y}_0(t)\|_\infty = 0$. Assume that for some k it holds

$$\|\mathbf{x}_k(t) - \mathbf{y}_k(t)\|_\infty \leq M \sum_{i=2}^{k+1} \frac{L^{i-2}}{i!} (t-t_0)^i, \quad \forall t \in [t_0, t_1] \quad (2.19)$$

Then,

$$\|\mathbf{x}_{k+1}(t) - \mathbf{y}_{k+1}(t)\|_\infty \leq \int_{t_0}^t \|\mathbf{f}(\tau, \mathbf{x}_k(\tau)) - \mathbf{f}(\tau, \mathbf{y}_k(\tau))\|_\infty d\tau \quad (2.20)$$

$$= \int_{t_0}^t \|\mathbf{f}(\tau, \mathbf{x}_k(\tau)) - \mathbf{f}(\tau, \mathbf{y}_k(\tau)) + \mathbf{f}(\tau, \mathbf{y}_k(\tau)) - \mathbf{f}(\tau, \mathbf{y}_k(\tau))\|_\infty d\tau \quad (2.21)$$

$$\leq \int_{t_0}^t L \|\mathbf{x}_k(\tau) - \mathbf{y}_k(\tau)\|_\infty + M(\tau - t_0) d\tau \quad (2.22)$$

$$\leq \int_{t_0}^t LM \sum_{i=2}^{k+1} \frac{L^{i-2}}{i!} (\tau - t_0)^i + M(\tau - t_0) d\tau \quad (2.23)$$

$$= LM \sum_{i=2}^{k+1} \frac{L^{i-2}}{i!} \int_{t_0}^t (\tau - t_0)^i d\tau + \int_{t_0}^t M(\tau - t_0) d\tau \quad (2.24)$$

$$= M \sum_{i=2}^{k+1} \frac{L^{i-1}}{(i+1)!} (t-t_0)^{i+1} + \frac{M}{2} (t-t_0)^2 \quad (2.25)$$

$$= M \sum_{i=2}^{k+2} \frac{L^{i-2}}{i!} (t-t_0)^i \quad (2.26)$$

for arbitrary $t \in [t_0, t_1]$. In order to get (2.22), we used triangle inequality, mean value theorem, Lipschitz condition (2.16) and bound for the partial derivative of \mathbf{f} by t (2.17).

Now, by induction we have that (2.19) holds for each k . Thus, for all k we have

$$\|\mathbf{x}_k - \mathbf{y}_k\|_{L^\infty([t_0, t_1])} \leq M \sum_{i=2}^{k+1} \frac{L^{i-2}}{i!} (t - t_0)^i \quad (2.27)$$

$$\leq M \sum_{i=2}^{\infty} \frac{L^{i-2}}{i!} (t - t_0)^i \quad (2.28)$$

$$= \frac{M}{L^2} \sum_{i=0}^{\infty} \frac{L^i}{i!} (t - t_0)^i - \frac{M}{L} (t - t_0) - \frac{M}{L^2} \quad (2.29)$$

$$= \frac{M}{L^2} e^{L(t-t_0)} - \frac{M}{L} (t - t_0) - \frac{M}{L^2}, \quad \forall t \in [t_0, t_1] \quad (2.30)$$

Since

$$\mathbf{x}_k \xrightarrow{L^\infty([t_0, t_1])} \mathbf{x}, \quad k \rightarrow \infty \quad (2.31)$$

$$\mathbf{y}_k \xrightarrow{L^\infty([t_0, t_1])} \mathbf{y}, \quad k \rightarrow \infty \quad (2.32)$$

we get the bound

$$\|\mathbf{x} - \mathbf{y}\|_{L^\infty([t_0, t_1])} \leq \frac{M}{L^2} e^{L(t-t_0)} - \frac{M}{L} (t - t_0) - \frac{M}{L^2}, \quad \forall t \in [t_0, t_1] \quad (2.33)$$

□

Gradient dynamical systems

A special type of dynamical systems are *gradient dynamical systems*. The name is derived by the fact that the vector field is negative gradient of some landscape function. More formally, let $\Omega \subset \mathbb{R}^n$, $I \subset \mathbb{R}$ be open sets and $E \in C^2(I \times \Omega; \mathbb{R})$. The dynamical system

$$\dot{\mathbf{x}}(t) = -\frac{\partial E^T}{\partial \mathbf{x}}(t, \mathbf{x}) \quad (2.34)$$

is a *gradient dynamical system* with corresponding *landscape* function E .

Generally, the function E can vary over time. However, let us analyze an example where E is time-independent, i.e. $E = E(\mathbf{x})$.

Example 2.1.10. Let $A \in M_n(\mathbb{R})$ be a real symmetric matrix and $E(\mathbf{x}) = \frac{1}{2} \mathbf{x}^T A \mathbf{x}$. Consider a gradient dynamical system with landscape function E .

$$\begin{aligned} \dot{\mathbf{x}}(t) &= -\frac{\partial E^T}{\partial \mathbf{x}}(\mathbf{x}(t)) \\ &= -A \mathbf{x} \end{aligned} \quad (2.35)$$

Since A is symmetric, it is orthogonally diagonalizable, so let $U^T A U = \Lambda = \text{diag}(\lambda_1, \dots, \lambda_n)$ be its orthogonal diagonalization.

Define a change of variables $\bar{\mathbf{x}} := U^T \mathbf{x}$. The equation (2.35) becomes

$$\begin{aligned}\dot{\bar{\mathbf{x}}} &= -U^T A U \bar{\mathbf{x}} \\ &= -\Lambda \bar{\mathbf{x}}\end{aligned}\tag{2.36}$$

which is a system of decoupled ordinary differential equations in each component. Written out component-wise, the system is

$$\dot{\bar{x}}_i = -\lambda_i \bar{x}_i; \quad i = 1, \dots, n\tag{2.37}$$

The solution for each component is thus

$$\bar{x}_i(t) = \bar{x}_i(0)e^{-\lambda_i t}; \quad i = 1, \dots, n\tag{2.38}$$

where $\bar{\mathbf{x}}(0)$ is the initial condition. The solution in the original reference frame is then $\mathbf{x}(t) = U \bar{\mathbf{x}}(t)$.

This change of reference provides us not only with the exact solution, but also with a qualitative understanding of solution's behavior. For each component we have different behavior depending on the sign of the corresponding eigenvalue. For $\lambda_i > 0$ the solution component rapidly collapses towards 0. For $\lambda_i = 0$, the solution component is stationary. For $\lambda_i < 0$ the solution component rapidly expands to $\pm\infty$.

This gives us the qualitative understanding of the behavior of $\mathbf{x}(t)$. When viewed in reference frame which diagonalizes A , the solution's components either collapse, stay stationary, or expand, all with respect to the sign of the corresponding eigenvalue.

Hamiltonian dynamical systems

Let $I \subset \mathbb{R}$, $\Omega \subset \mathbb{R}^{2n}$ be open sets and $H \in C^2(I \times \Omega)$ where $H = H(t, \mathbf{x}, \mathbf{y})$, $\mathbf{x}, \mathbf{y} \in \mathbb{R}^n$. A system of the form

$$\begin{aligned}\dot{\mathbf{x}} &= \frac{\partial H^T}{\partial \mathbf{y}}(t, \mathbf{x}, \mathbf{y}) \\ \dot{\mathbf{y}} &= -\frac{\partial H^T}{\partial \mathbf{x}}(t, \mathbf{x}, \mathbf{y})\end{aligned}\tag{2.39}$$

is called a Hamiltonian dynamical system.

Function H is called a *Hamiltonian function* or *total energy* of the system.

Hamiltonian H can take various forms.

The motion of the particles in space can be modeled by taking H to be of the following form

$$H(t, \mathbf{x}, \mathbf{y}) = \sum_{i=1}^n \frac{m_i}{2} y_i^2 + E(t, \mathbf{x}) \quad (2.40)$$

Each particle has certain number of degrees of freedom (for example 1D, 2D or 3D motion). Each degree of freedom has its corresponding *position* x_i and *momentum* y_i . The quantity $\sum_{i=1}^n \frac{m_i}{2} y_i^2$ is called the *kinetic energy* of the system, while $E(t, \mathbf{x})$ is a function called *potential energy* of the system. Kinetic energy depends only on constants m_i which represent the *mass* of each particle, and momenta y_i . On the other hand, the potential energy depends only on the position vector \mathbf{x} and the time instance in case that the potential energy changes over time.

Example 2.1.11. Let $A \in M_n(\mathbb{R})$ be a real symmetric matrix and $E(\mathbf{x}) = \frac{1}{2} \mathbf{x}^T A \mathbf{x}$. Consider an autonomous Hamiltonian dynamical system corresponding to the motion of particles, each with mass m , in time-independent potential $E(\mathbf{x})$

$$\begin{aligned} H(\mathbf{x}, \mathbf{y}) &= \sum_{i=1}^n \frac{m}{2} y_i^2 + \frac{1}{2} \mathbf{x}^T A \mathbf{x} \\ \dot{\mathbf{x}} &= \frac{\partial H}{\partial \mathbf{y}} \\ &= m \mathbf{y} \\ \dot{\mathbf{y}} &= -\frac{\partial H}{\partial \mathbf{x}} \\ &= -A \mathbf{x} \end{aligned} \quad (2.41)$$

Since A is symmetric, it is orthogonally diagonalizable, so let $U^T A U = \Lambda = \text{diag}(\lambda_1, \dots, \lambda_n)$ be its orthogonal diagonalization.

Define a change of variables $\bar{\mathbf{x}} := U^T \mathbf{x}$, $\bar{\mathbf{y}} := U^T \mathbf{y}$. The dynamical system (2.41) becomes

$$\begin{aligned} \dot{\bar{\mathbf{x}}} &= m \bar{\mathbf{y}} \\ \dot{\bar{\mathbf{y}}} &= -U^T A U \bar{\mathbf{x}} = -\Lambda \bar{\mathbf{x}} \end{aligned} \quad (2.42)$$

or written out component-wise

$$\begin{aligned} \dot{\bar{x}}_i &= m \bar{y}_i; & i = 1, \dots, n \\ \dot{\bar{y}}_i &= -\lambda_i \bar{x}_i; & i = 1, \dots, n \end{aligned} \quad (2.43)$$

By taking the derivative of $\dot{\bar{x}}_i$ we obtain an equivalent equation

$$\ddot{\bar{x}}_i = -m \lambda_i \bar{x}_i; \quad i = 1, \dots, n \quad (2.44)$$

This is a system of decoupled linear equations so for each component we have the solution depending on the sign of λ_i .

If $\lambda_i > 0$, then

$$\bar{x}_i(t) = \frac{1}{\sqrt{m\lambda_i}} \dot{\bar{x}}_i(0) \sin(\sqrt{m\lambda_i}t) + \bar{x}_i(0) \cos(\sqrt{m\lambda_i}t) \quad (2.45)$$

which means that this component periodically oscillates around 0.

If $\lambda_i = 0$, then

$$\bar{x}_i(t) = \dot{\bar{x}}_i(0)t + \bar{x}_i(0) \quad (2.46)$$

which means that this component linearly expands.

If $\lambda_i < 0$,

$$\bar{x}_i(t) = \left(\frac{\bar{x}_i(0)}{2} + \frac{\dot{\bar{x}}_i(0)}{2\sqrt{-m\lambda_i}} \right) e^{\sqrt{-m\lambda_i}t} + \left(\frac{\bar{x}_i(0)}{2} - \frac{\dot{\bar{x}}_i(0)}{2\sqrt{-m\lambda_i}} \right) e^{-\sqrt{-m\lambda_i}t} \quad (2.47)$$

which means that this component exponentially expands.

By changing the reference frame back to the original $\mathbf{x} = U\bar{\mathbf{x}}$, we can obtain the exact solution of the dynamical system. However, the qualitative behavior of this dynamical system can be understood by analyzing each solution component of the system in reference frame which diagonalizes A . Each component in this reference frame either oscillates around zero, linearly expands, or exponentially expands, with respect to the sign of the corresponding eigenvalue.

2.2 Numerical simulations

In this section, two numerical methods for solving ordinary differential equations are presented. Those are Euler method and symplectic Euler method.

Euler method is a numerical method which can be used to approximately solve any initial-value problem and thus simulate any dynamical system. Although being very simple to state and implement, its weakness is being less precise than some other numerical methods such as various other Runge-Kutta methods.

Symplectic Euler method is a modification of the Euler method used for simulating special kind of ordinary differential equations, namely Hamiltonian dynamical systems. Although the error bound is of the same order as for the Euler method, symplectic Euler method conserves the energy (for time-independent Hamiltonian) much better than the Euler method. Thus, symplectic Euler method is more appropriate for simulating Hamiltonian dynamics.

Many other numerical methods for solving various ordinary differential equations can be found in [40] and [41], each with certain advantages and disadvantages. However, only these two methods will be used in chapter 3.

Euler method

This section is based on [40]. Consider a given initial value problem

$$\begin{aligned}\dot{\mathbf{x}}(t) &= \mathbf{f}(t, \mathbf{x}) \\ \mathbf{x}(t_0) &= \mathbf{x}_0\end{aligned}\tag{2.48}$$

The Euler method for approximately solving this initial value problem is given by the iterative formula

$$\begin{aligned}\mathbf{x}^{(0)} &= \mathbf{x}_0 \\ \mathbf{x}^{(k+1)} &= \mathbf{x}^{(k)} + \mathbf{f}(t_k, \mathbf{x}^{(k)}) \cdot \Delta_t, \quad k = 0, \dots, N_{\text{iter}} - 1\end{aligned}\tag{2.49}$$

where $\Delta_t = \frac{T}{N_{\text{iter}}}$, $t_k = k\Delta_t$, and T is the time interval on which we need to approximate the solution. For each k , $\mathbf{x}^{(k)}$ is the approximate value for $\mathbf{x}(t_k)$.

Euler method is a first-order method, meaning that the bound for error between the approximate and exact solutions is proportional to Δ_t , and is not proportional to any higher power of Δ_t . Details can be found in [40].

Symplectic Euler method

This section is based on [42]. Symplectic Euler method (sometimes also called semi-implicit Euler method) can be used for approximately solving the initial value problem of the form

$$\begin{aligned}\dot{\mathbf{x}} &= \mathbf{f}(t, \mathbf{y}) \\ \dot{\mathbf{y}} &= \mathbf{g}(t, \mathbf{x}) \\ \mathbf{x}(t_0) &= \mathbf{x}_0 \\ \mathbf{y}(t_0) &= \mathbf{y}_0\end{aligned}\tag{2.50}$$

This system typically arises in Hamiltonian dynamics if the Hamiltonian has separable variables

$$H(t, \mathbf{x}, \mathbf{y}) = E(t, \mathbf{x}) + T(t, \mathbf{y})$$

Often E is the potential energy of the system, while T is the kinetic energy.

Symplectic Euler method for approximately solving this system is given by the iterative formula

$$\begin{aligned}\mathbf{x}^{(0)} &= \mathbf{x}_0 \\ \mathbf{y}^{(0)} &= \mathbf{y}_0 \\ \mathbf{x}^{(k+1)} &= \mathbf{x}^{(k)} + \mathbf{f}(t_k, \mathbf{y}^{(k)}) \cdot \Delta_t \\ \mathbf{y}^{(k+1)} &= \mathbf{y}^{(k)} + \mathbf{g}(t_k, \mathbf{x}^{(k+1)}) \cdot \Delta_t, \quad k = 0, \dots, N_{\text{iter}} - 1\end{aligned}\tag{2.51}$$

where $\Delta_t = \frac{T}{N_{\text{iter}}}$, $t_k = k\Delta_t$, and T is the time interval on which we need to approximate the solution. For each k , $\mathbf{x}^{(k)}$ and $\mathbf{y}^{(k)}$ are respectively approximate values for $\mathbf{x}(t_k)$ and $\mathbf{y}(t_k)$. The difference between original Euler method is that $\mathbf{x}^{(k+1)}$ is used for calculating $\mathbf{y}^{(k+1)}$, instead of $\mathbf{x}^{(k)}$.

A second variant of this method is given by *reversing the order of calculations*, i.e.

$$\begin{aligned} \mathbf{x}^{(0)} &= \mathbf{x}_0 \\ \mathbf{y}^{(0)} &= \mathbf{y}_0 \\ \mathbf{y}^{(k+1)} &= \mathbf{y}^{(k)} + \mathbf{g}(t_k, \mathbf{x}^{(k)}) \cdot \Delta_t \\ \mathbf{x}^{(k+1)} &= \mathbf{x}^{(k)} + \mathbf{f}(t_k, \mathbf{y}^{(k+1)}) \cdot \Delta_t, \quad k = 0, \dots, N_{\text{iter}} - 1 \end{aligned} \tag{2.52}$$

Although symplectic Euler method is also a first-order method, i.e. the error is proportional to Δ_t , it is usually a more appropriate method for simulating Hamiltonian dynamical systems compared to the original Euler method. This is due to the fact that symplectic Euler method conserves energy (for time-independent Hamiltonians) better than Euler method. In fact, Euler method often persistently increases the energy, making it less accurate. Details about this method can be found in [42] and [41].

2.3 Other

This section serves only as a reference for some arguments used in the rest of the thesis.

Theorem 2.3.1 (Weyl's inequality [43]). *Let $M = N + R$, N , and R be $n \times n$ Hermitian matrices, with their respective eigenvalues μ_i, η_i, ρ_i ordered in descending order as follows:*

$$\begin{aligned} M : \quad & \mu_1 \geq \dots \geq \mu_n, \\ N : \quad & \eta_1 \geq \dots \geq \eta_n, \\ R : \quad & \rho_1 \geq \dots \geq \rho_n \end{aligned}$$

Then the following inequalities hold:

$$\eta_i + \rho_n \leq \mu_i \leq \eta_i + \rho_1, \quad i = 1, \dots, n$$

Proposition 2.3.2. *Let $\Omega \subset \mathbb{R}^n$ be an open, convex and connected set. Let $E \in C^2(\Omega)$ such that the second differential $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ is positive semidefinite, i.e. all of its eigenvalues are greater than 0, at each point of Ω .*

Then E is a convex function.

Definition 2.3.3. [44] *Consider a second-order differential equation*

$$\ddot{x}(t) = tx(t) \tag{2.53}$$

Two linearly independent solutions of this equation are called Airy functions and are denoted by $A(t), B(t)$.

Proposition 2.3.4. *Let $D \subset \mathbb{R}^n$ be a compact and convex set. Let $f \in C^1(D)$. Then, f is Lipschitz continuous.*

Proof. Choose arbitrary $\mathbf{x}, \mathbf{y} \in D$ and consider the line between them parametrized by $\gamma : [0, 1] \rightarrow D, \gamma(t) = (1 - t)\mathbf{x} + t\mathbf{y}$. Since D is convex, γ is indeed contained in D . Define $g : [0, 1] \rightarrow \mathbb{R}, g = f \circ \gamma$. Then, by the mean value theorem there exists $\xi \in \langle 0, 1 \rangle$ such that

$$\frac{\partial f}{\partial \mathbf{x}}(\gamma(\xi)) \cdot (\mathbf{y} - \mathbf{x}) = g'(\xi) = g(1) - g(0) = f(\mathbf{y}) - f(\mathbf{x})$$

Thus,

$$|f(\mathbf{y}) - f(\mathbf{x})| \leq [\text{Cauchy-Schwartz}] \leq \left\| \frac{\partial f}{\partial \mathbf{x}}(\gamma(\xi)) \right\| \cdot \|\mathbf{y} - \mathbf{x}\| \leq L \cdot \|\mathbf{y} - \mathbf{x}\|$$

where

$$L = \max_{\mathbf{x} \in D} \left\| \frac{\partial f}{\partial \mathbf{x}}(\mathbf{x}) \right\| < \infty$$

Thus, f is Lipschitz with a Lipschitz constant L . □

Chapter 3

Algorithms

This chapter presents and analyses physics-inspired dynamical systems and their corresponding numerical simulations, which together comprise algorithms for heuristically solving QUBO problem. These algorithms are namely Coherent Ising Machine (CIM) 3.2, Simulated Bifurcation (SB) 3.4, Simulated Coherent Ising Machine (SimCIM) 3.8, and Ballistic Simulated Bifurcation (bSB) 3.9. There are other similar variants of these physics-inspired algorithms, such as Discrete Simulated Bifurcation (dSB) [31] and heated versions of simulated bifurcation [45], but those are not covered in the thesis.

These physics-inspired algorithms were proposed as adaptations of quantum algorithms (algorithms designed for running on a quantum computer) for solving QUBO and other combinatorial optimization problems. These adaptations are done by approximating the expected value of annihilation operator a , present in the quantum Hamiltonian H , by a complex number $x + iy$ where $x, y \in \mathbb{R}$. What is left after this *dequantization* are the equations of motion for variables x, y . This is where physics-inspired dynamical systems for solving combinatorial optimization problems came from — the details can be found in [27], [29], and [25]. Modifying these dynamical systems further enabled to design algorithm which are easier to simulate, while preserving, and even improving the solution quality [31, 30].

Although motivated by quantum systems, these dynamical systems are completely classical, often even deterministic. Thus, in what follows, those systems will be analyzed from a mathematical perspective of dynamical systems.

First, in order to understand how a dynamical system generates candidate solutions of QUBO problem, consider a state vector which evolves in time $\mathbf{x} : [0, \infty) \rightarrow \mathbb{R}^n$. Taking the sign of each component at some time instance t , $\sigma_i := \text{sign } x_i(t)$ provides a candidate solution for the QUBO task $(\sigma_1, \dots, \sigma_n)$. The goal is to obtain a dynamical system which *attracts* various trajectories towards such areas of space that provide high-quality approximate solutions of the QUBO problem, according to the mapping mentioned above

$\sigma_i := \text{sign } x_i(t)$.

Finally, CIM and SB were chronologically developed earlier, and they are respectively a gradient and a Hamiltonian system over the same energy/landscape function E . Motivated by CIM and SB, and by modifying the function E , new algorithms were proposed. This is how bSB and SimCIM were born. They are also respectively a gradient and a Hamiltonian systems over this new energy/landscape function.

3.1 Introduction to CIM and SB

Let us define the following energy functional

$$E(\mathbf{x}; \alpha, \beta, \mu) = \frac{\alpha}{4} \sum_{i=1}^n (x_i^2 - \mu)^2 - \frac{\beta}{2} \sum_{i,j=1}^n J_{ij} x_i x_j \quad (3.1)$$

and the corresponding gradient vector field

$$\begin{aligned} \mathbf{f}(\mathbf{x}; \alpha, \beta, \mu) &= -\frac{\partial E}{\partial \mathbf{x}}^T(\mathbf{x}; \alpha, \beta, \mu) \\ &= -\alpha[x_1^3, \dots, x_n^3]^T + \alpha\mu\mathbf{x} + \beta J\mathbf{x} \end{aligned} \quad (3.2)$$

both of which will be used in sections 3.2 and 3.4. E and \mathbf{f} are functions of position \mathbf{x} and parameters α, β, μ denoted after the semicolon. It is essential to analyze this energy functional E with respect to parameters because it defines the vector field \mathbf{f} which will govern the dynamical systems in sections 3.2 and 3.4. This energy function will vary in time itself because the parameters will vary in time.

The given coupling matrix J is symmetric (if not, we may always symmetrize it by taking coefficients $\frac{1}{2}(J_{ij} + J_{ji})$ instead of J_{ij} and J_{ji}). $\alpha(t) > 0$, $\beta(t) > 0$ and $\mu(t) \in \mathbb{R}$ are parameters which (might) vary over time and thus produce different landscape over which the point \mathbf{x} moves. However, both in CIM and SB algorithm, parameters α and β are constant in time. The crucial part is that $\mu(t)$ is time-dependent which causes the bifurcations of the system. When talking about CIM and SB, a *bifurcation* refers to the emergence of new or vanishing of some existing local minima of energy functional E . The second part, $-\frac{\beta(t)}{2} \sum_{i,j} J_{ij} x_i x_j$, corresponds to the value of the QUBO task and is the actual function that we want to minimize but over the set $\{-1, +1\}^n$. The first part, $\frac{\alpha(t)}{4} \sum_i (x_i^2 - \mu(t))^2$, thus might be viewed as a penalty term. It penalizes x_i 's whose magnitude is different than $\sqrt{\mu}$ (whenever $\mu \geq 0$). More about penalty terms can be found in [46].

When μ is small enough, the only local and global minimum of E is the origin. As μ increases, new local minima occur. For large enough μ , E will have 2^n local minima corresponding to (existing in the close vicinity of) the points $\{-\sqrt{\mu}, +\sqrt{\mu}\}^n$. The value of

the energy functional E at such a point $\mathbf{x} = \sqrt{\mu}\boldsymbol{\sigma}$, $\boldsymbol{\sigma} \in \{-1, +1\}^n$ is given by

$$E(\sqrt{\mu}\boldsymbol{\sigma}; \alpha, \beta, \mu) = -\frac{\beta\mu}{2} \sum_{i,j} J_{ij}\sigma_i\sigma_j$$

which is proportional to the function that we are minimizing over the discrete set.

Increasing the parameter μ from the starting value μ_0 to the final value μ_1 causes new bifurcation of the system. As already stated, if we evolve the system long enough, i.e. until μ crosses certain large enough value, E will have 2^n local minima, each corresponding to one feasible solution from $\{-1, +1\}^n$. This is the statement of the next theorem.

Theorem 3.1.1. *Let $D := \max_i \sum_{j=1}^n |\beta J_{ij}|$ For $\mu > \frac{1}{2\alpha}(1 + 3\sqrt{3})D$, there exist 2^n local minima of function E , which are all in a one-to-one correspondence with the feasible set $\{-1, +1\}^n$. This correspondence is given by taking the sign of each component — for a local minimum \mathbf{x} , the corresponding element of the feasible set is $\boldsymbol{\sigma} = \text{sign } \mathbf{x}$.*

Proof. The proof can be found in [37]. □

Let us calculate the second differential of E .

$$\frac{\partial^2 E}{\partial \mathbf{x}^2} = 3\alpha \text{diag}(x_1^2, \dots, x_n^2) - \alpha\mu I - \beta J \quad (3.3)$$

Proposition 3.1.2. *The origin is a stationary point of E , meaning that $\frac{\partial E}{\partial \mathbf{x}}(\mathbf{0}) = \mathbf{0}^T$, for all $\mu \in \mathbb{R}$. For*

$$\mu < -\frac{\beta}{\alpha} \lambda_{\max}(J) \quad (3.4)$$

the origin is a local minimum of E . For

$$\mu > -\frac{\beta}{\alpha} \lambda_{\max}(J) \quad (3.5)$$

the origin is not a local minimum of E .

Proof. Use formula (3.2) to see that the origin is stationary. The second differential of E at the origin is

$$\frac{\partial^2 E}{\partial \mathbf{x}^2}(\mathbf{0}) = -\alpha\mu I - \beta J$$

If (3.4) holds, then $\lambda_{\min}(\frac{\partial^2 E}{\partial \mathbf{x}^2}(\mathbf{0})) = -\alpha\mu - \beta\lambda_{\max}(J) > \alpha\frac{\beta}{\alpha}\lambda_{\max}(J) - \beta\lambda_{\max}(J) = 0$ which means that all eigenvalues of $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ are positive which implies that the origin is a local minimum of E . Analogous argument shows that when (3.5) holds, the origin is not a local minimum because at least one eigenvalue is negative. □

For

$$\mu \leq -\frac{\beta}{\alpha} \lambda_{\max}(J) \quad (3.6)$$

we can actually analyze the second differential of E to see that

$$\begin{aligned} \lambda_{\min} \left(\frac{\partial^2 E}{\partial \mathbf{x}^2} \right) &\geq [\text{Weyl's inequality (2.3.1)}] \\ &\geq \lambda_{\min}(3\alpha \text{diag}(x_1^2, \dots, x_n^2)) + \lambda_{\min}(-\alpha\mu I - \beta J) \\ &= 3\alpha \min_{i=1, \dots, n} x_i^2 - \alpha\mu - \beta \lambda_{\max}(J) \\ &\geq -\alpha\mu - \beta \lambda_{\max}(J) \\ &\geq \alpha \frac{\beta}{\alpha} \lambda_{\max}(J) - \beta \lambda_{\max}(J) = 0 \end{aligned} \quad (3.7)$$

So, when condition (3.6) is satisfied, we know from 2.3.2 and 3.1.2 that E is a convex function with its global minimum centered at the origin.

The condition

$$\mu = -\frac{\beta}{\alpha} \lambda_{\max}(J) \quad (3.8)$$

is thus called the *first bifurcation point*.

It thus makes sense to increase μ , for example linearly, from certain starting value μ_0 to certain ending value μ_1 , while taking

$$\mu_0 \geq -\frac{\beta}{\alpha} \lambda_{\max}(J) \quad (3.9)$$

because this is the point when bifurcations in landscape function E start to emerge. Selecting the concrete value of μ_0 and μ_1 should be based on empirical evidence i.e. by experimenting which parameter provides the best result.

Example 3.1.3. *In this example we will observe what happens for the case $n = 2$. Since the coupling matrix is symmetric and with zero diagonal elements, there are, up to scale, two possibilities for the coupling matrix, and those are given by $J = \begin{bmatrix} 0 & \epsilon \\ \epsilon & 0 \end{bmatrix}$, $\epsilon \in \{-1, +1\}$.*

Consider the QUBO task given by

$$\max_{\sigma_1, \sigma_2 \in \{-1, +1\}} \sum_{i,j=1}^2 J_{ij} \sigma_i \sigma_j \quad (3.10)$$

and define the corresponding energy landscape as in (3.1)

$$E = \frac{\alpha}{4} \sum_{i=1}^2 (x_i^2 - \mu)^2 - \beta \epsilon x_1 x_2 \quad (3.11)$$

Diagonalization of J provides us with eigenvectors $\begin{bmatrix} 1 \\ -1 \end{bmatrix}$, $\begin{bmatrix} 1 \\ 1 \end{bmatrix}$ corresponding respectively to eigenvalues $-\epsilon$, $+\epsilon$. First, we will determine stationary points of E for some fixed μ .

$$\begin{aligned} \alpha x_1(\mu - x_1^2) + \beta \epsilon x_2 &= 0 \\ \alpha x_2(\mu - x_2^2) + \beta \epsilon x_1 &= 0 \end{aligned} \quad (3.12)$$

which is, by summing and subtracting these equations, equivalent to

$$\begin{aligned} \alpha \mu(x_1 + x_2) - \alpha(x_1 + x_2)(x_1^2 - x_1x_2 + x_2^2) + \beta \epsilon(x_1 + x_2) &= 0 \\ \alpha \mu(x_1 - x_2) - \alpha(x_1 - x_2)(x_1^2 + x_1x_2 + x_2^2) - \beta \epsilon(x_1 - x_2) &= 0 \end{aligned} \quad (3.13)$$

Case 1: $x_1 + x_2 = 0$, $x_1 - x_2 = 0$

This is the trivial case $x_1 = x_2 = 0$ which is the stationary point for all μ .

Case 2: $x_1 + x_2 = 0$, $x_1 - x_2 \neq 0$

From first equation we have $x_1 = -x_2$ so plugging it into the above we get

$$\begin{aligned} 2\alpha \mu x_1 - 2\alpha x_1^3 - 2\beta \epsilon x_1 &= 0 \\ \Leftrightarrow x_1^2 = \mu - \frac{\beta \epsilon}{\alpha} \end{aligned} \quad (3.14)$$

From here we conclude that

$$-x_2 = x_1 = \pm \sqrt{\mu - \frac{\beta \epsilon}{\alpha}}, \quad \text{for } \mu \geq \frac{\beta \epsilon}{\alpha} \quad (3.15)$$

Case 3: $x_1 + x_2 \neq 0$, $x_1 - x_2 = 0$

From second equation we have $x_1 = x_2$ so plugging it into the above we get

$$\begin{aligned} 2\alpha \mu x_1 - 2\alpha x_1^3 + 2\beta \epsilon x_1 &= 0 \\ \Leftrightarrow x_1^2 = \mu + \frac{\beta \epsilon}{\alpha} \end{aligned} \quad (3.16)$$

From here we conclude that

$$x_2 = x_1 = \pm \sqrt{\mu + \frac{\beta \epsilon}{\alpha}}, \quad \text{for } \mu \geq -\frac{\beta \epsilon}{\alpha} \quad (3.17)$$

Case 4: $x_1 + x_2 \neq 0$, $x_1 - x_2 \neq 0$

This provides

$$\begin{aligned} \alpha \mu - \alpha(x_1^2 - x_1x_2 + x_2^2) + \beta \epsilon &= 0 \\ \alpha \mu - \alpha(x_1^2 + x_1x_2 + x_2^2) - \beta \epsilon &= 0 \end{aligned} \quad (3.18)$$

$$\Leftrightarrow \begin{aligned} x_1^2 - x_1x_2 + x_2^2 &= \mu + \frac{\beta\epsilon}{\alpha} \\ x_1^2 + x_1x_2 + x_2^2 &= \mu - \frac{\beta\epsilon}{\alpha} \end{aligned} \quad (3.19)$$

$$\Leftrightarrow [\text{by introducing } y_1 = x_1 + x_2, y_2 = x_1 - x_2]$$

$$\Leftrightarrow \begin{aligned} y_1^2 + 3y_2^2 &= 4\left(\mu + \frac{\beta\epsilon}{\alpha}\right) \\ 3y_1^2 + y_2^2 &= 4\left(\mu - \frac{\beta\epsilon}{\alpha}\right) \end{aligned} \quad (3.20)$$

which is a full-rank linear system with quadratic indeterminates whose solution is then

$$\begin{aligned} y_1^2 &= \frac{1}{2}\left[-\left(\mu + \frac{\beta\epsilon}{\alpha}\right) + 3\left(\mu - \frac{\beta\epsilon}{\alpha}\right)\right] = \mu - 2\frac{\beta\epsilon}{\alpha} \\ y_2^2 &= \frac{1}{2}\left[3\left(\mu + \frac{\beta\epsilon}{\alpha}\right) - \left(\mu - \frac{\beta\epsilon}{\alpha}\right)\right] = \mu + 2\frac{\beta\epsilon}{\alpha} \end{aligned} \quad (3.21)$$

leading to 4 solutions

$$\begin{aligned} x_1 &= \frac{1}{2}\left[(-1)^k \sqrt{\mu - \frac{2\beta\epsilon}{\alpha}} + (-1)^l \sqrt{\mu + \frac{2\beta\epsilon}{\alpha}}\right] \\ x_2 &= \frac{1}{2}\left[(-1)^k \sqrt{\mu - \frac{2\beta\epsilon}{\alpha}} - (-1)^l \sqrt{\mu + \frac{2\beta\epsilon}{\alpha}}\right] \end{aligned}, \quad k, l \in \{0, 1\} \quad (3.22)$$

which holds whenever $\mu \geq |\epsilon| \frac{2\beta}{\alpha}$.

First two non-trivial cases are local minima, which can be seen by taking the second partial derivatives of E :

$$-\frac{\partial^2 E}{\partial \mathbf{x}^2} = \begin{bmatrix} \alpha\mu - 3\alpha x_1^2 & \beta\epsilon \\ \beta\epsilon & \alpha\mu - 3\alpha x_2^2 \end{bmatrix} \quad (3.23)$$

If $-\frac{\partial^2 E}{\partial \mathbf{x}^2}$ is negative-definite (all of its eigenvalues are less than zero), then we have a local minimum. Calculating the eigenvalues is straightforward for this 2×2 symmetric matrix, and plugging in from first two cases the fact that $x_1^2 = x_2^2$ we obtain that E has local minimum when

$$\beta|\epsilon| < \alpha(3x_1^2 - \mu) \quad (3.24)$$

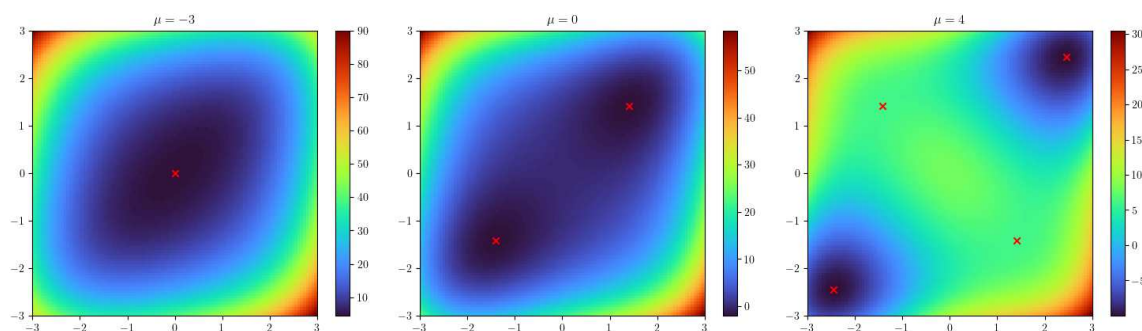
In table 3.1 we systematically write all the cases and conclude that those local minima that correspond to optimum solution of the minimization problem (3.10) bifurcate earlier than other local minima.

In figure 3.1 we can see how these local minima start to emerge as μ increases.

	$\epsilon = +1$	$\epsilon = -1$
$x_1 = -x_2$ (case (3.15))	$\mu > \frac{2\beta}{\alpha}$	$\mu > -\frac{\beta}{\alpha}$
$x_1 = x_2$ (case (3.17))	$\mu > -\frac{\beta}{\alpha}$	$\mu > \frac{2\beta}{\alpha}$
solution for $\max_{\sigma_1, \sigma_2 \in \{-1, +1\}} 2\epsilon\sigma_1\sigma_2$	$\sigma_1 = \sigma_2$	$\sigma_1 = -\sigma_2$

Table 3.1: Behaviour of E on a system with two variables

We see that those local minima that correspond to the optimum solution (highlighted in gray) bifurcate earlier than suboptimal local minima.

Figure 3.1: Visualization of energy landscape E for CIM and SB

Overview: This is a visual representation of energy landscape E for a system with two variables and coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. As the parameter μ increases, it causes the qualitative appearance of the landscape function E to change — new local minima emerge, and they correspond to candidate solutions of QUBO. Notice that local minima corresponding to optimum solutions of QUBO bifurcate earlier than suboptimal ones.

Details: The values of parameter μ are denoted on the top of each subplot. Red crosses denote local minima of E . $\alpha = 1, \beta = 2$.

3.2 Coherent Ising machine (CIM)

This section is based on [28] and [37].

Dynamics

CIM is a gradient dynamical system with time-dependent landscape function E defined by equation 3.1. The vector field \mathbf{f} describing this dynamical system is thus given by the negative gradient of E , as defined by 3.2

$$\begin{aligned}\mathbf{f}(\mathbf{x}; \alpha, \beta, \mu) &= -\frac{\partial E}{\partial \mathbf{x}}(\mathbf{x}; \alpha, \beta, \mu) \\ &= -\alpha[x_1^3, \dots, x_n^3]^T + \alpha\mu\mathbf{x} + \beta J\mathbf{x}\end{aligned}\quad (3.25)$$

Parameters are written after the semicolon and those are α, β, μ . Time dependence is achieved by varying parameters in time. In this algorithm only parameter μ is monotonically increased from certain initial value μ_0 to certain final value μ_1 . Parameters $\alpha, \beta > 0$ are kept constant.

The dynamical system describing CIM is thus given by

$$\dot{\mathbf{x}}(t) = \mathbf{f}(\mathbf{x}(t); \alpha, \beta, \mu(t)) \quad (3.26)$$

Written out component-wise without denoting the dependence on time

$$\dot{x}_i = f_i = \alpha x_i(\mu - x_i^2) + \beta \sum_j J_{ij} x_j \quad ; \quad i = 1, \dots, n \quad (3.27)$$

Algorithm

CIM algorithm is a simulation of the described dynamical system on a computer using Euler method.

Algorithm 1 CIM

- 1: Initialize vector \mathbf{x} randomly around $\mathbf{0}$
 - 2: **for** k in range(0, N_{iter}) **do**
 - 3: $x_i \leftarrow x_i + \left[\alpha x_i(\mu(t_k) - x_i^2) + \beta \sum_j J_{ij} x_j \right] \cdot \Delta_t, \quad i = 1, \dots, n$
 - 4: **end for**
-

In figure 3.2 we observe that as local minima bifurcate away from the origin, the system follows some of these local minima. Local minima corresponding to $(+, +)$ and $(-, -)$ are lower than those corresponding to $(+, -)$ and $(-, +)$ so it is more likely that the system

will converge towards $(+, +)$ or $(-, -)$ that the other two minima. Also, local minima corresponding to $(+, +)$, $(-, -)$ bifurcate earlier, i.e. for smaller values of μ , than the other two local minima. That's why in the middle subplot the system will surely converge to the optimal solution, since other local minima have not yet even emerged.

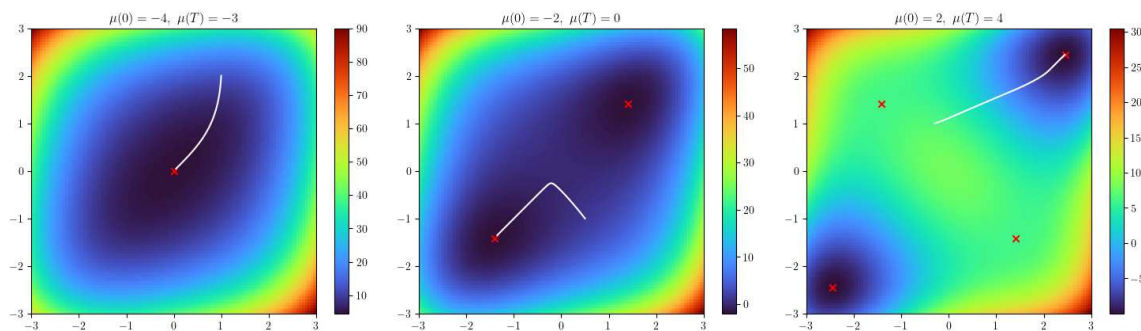


Figure 3.2: CIM algorithm

Overview: This is a visual representation of CIM algorithm on a system with two variables with coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. In the left subplot, the parameter μ did not cross the first bifurcation point, so the dynamical system collapses towards the origin (starting far away from it). As the time passes, μ increases, causing a change in the qualitative appearance of landscape E . In the middle subplot, the origin splits into two local minima (red crosses) which correspond to the solution of QUBO problem. In the right subplot, two new local minima emerge, corresponding to suboptimal solutions of QUBO. However, they are not as *attractive* for the dynamical system as the optimal solutions. In both cases, the dynamical system ends up converging towards a local minimum corresponding to an optimal solution of QUBO.

Details: The white line represents the trajectory obtained by running the CIM algorithm. Red crosses denote local minima of E . Each subplot corresponds to a new run of the algorithm. The parameters are $\alpha = 1, \beta = 2$. The energy landscape E is plotted only for the final value of μ , that is $E(\cdot; \alpha, \beta, \mu(T))$ is plotted on each of the subplots. In each subplot, μ increases linearly from starting to ending value. Starting and ending values are respectively denoted as $\mu(0)$ and $\mu(T)$ on top of each subplot. Number of iterations is 400, while $\Delta_t = 0.01$. Initial positions are $\mathbf{x}(0) = (1, 2)$ for the left subplot, $\mathbf{x}(0) = (0.5, -1.0)$ for the middle subplot, and $\mathbf{x}(0) = (-0.3, 1.0)$ for the right subplot.

3.3 Gradient descent and momentum

Gradient descent algorithms are used extensively for various continuous optimization tasks, such as for training neural networks in machine learning models. These algorithms could be understood as Euler method applied to the gradient system where the gradient is taken of the function which is being optimized. Suppose that we want to find a (local) minimum of a function $E : \mathbb{R}^n \rightarrow \mathbb{R}$. A gradient descent algorithm is then

$$\begin{aligned} \mathbf{x}^{(0)} &= \mathbf{x}_0 \\ \mathbf{x}^{(k+1)} &= \mathbf{x}^{(k)} - \frac{\partial E}{\partial \mathbf{x}}(\mathbf{x}^{(k)}) \cdot \Delta_t, \quad k = 0, \dots, N_{\text{iter}} - 1 \end{aligned} \quad (3.28)$$

where Δ_t is a parameter called *learning rate*, N_{iter} is the number of iterations, and \mathbf{x}_0 is some initial position which is often randomly chosen, ideally in a close vicinity of the target (local) minimum.

On top of this, there is a technique which is often used for improving the convergence rate of such optimization algorithms, and it is called the *momentum* [47]. The momentum acts in such a way that it accumulates the previous displacements and takes those into account when performing the next update. The algorithm is the following

$$\begin{aligned} \mathbf{x}^{(0)} &= \mathbf{x}_0 \\ \mathbf{x}^{(1)} &= \mathbf{x}^{(0)} - \frac{\partial E}{\partial \mathbf{x}}(\mathbf{x}^{(0)}) \cdot \Delta_t \\ \mathbf{x}^{(k+1)} &= \mathbf{x}^{(k)} - \frac{\partial E}{\partial \mathbf{x}}(\mathbf{x}^{(k)}) \cdot \Delta_t + \gamma (\mathbf{x}^{(k)} - \mathbf{x}^{(k-1)}), \quad k = 1, \dots, N_{\text{iter}} - 1 \end{aligned} \quad (3.29)$$

where all parameters are as earlier, and $\gamma \in [0, 1]$ is the momentum parameter. Adding the momentum has been shown to increase the convergence rate significantly [47]. According to [47], adding momentum is actually equivalent to the numerical simulation of a physical system in which a Newtonian particle moves through a viscous medium under the influence of a conservative force field. This interpretation is of special interest to us because we are already dealing with continuous dynamical systems.

Now let's return to the CIM algorithm.

Adding momentum to CIM

Since CIM is essentially a gradient descent algorithm, it might make sense to apply the momentum to it for improving its performance.

Adding the momentum to algorithm 1, we get the following: First, choose the momentum parameter $0 \leq \gamma \leq 1$. Momenta will be stored in vector \mathbf{y} . We could also initialize the momentum vector randomly.

Algorithm 2 CIM with momentum

-
- 1: Initialize vectors \mathbf{x}, \mathbf{y} randomly around $\mathbf{0}$
 - 2: **for** k in range($0, N_{\text{iter}}$) **do**
 - 3: $y_i \leftarrow \gamma \cdot y_i + \alpha x_i(\mu(t_k) - x_i^2) + \beta \sum_j J_{ij} x_j, \quad i = 1, \dots, n$
 - 4: $x_i \leftarrow x_i + y_i \cdot \Delta t, \quad i = 1, \dots, n$
 - 5: **end for**
-

For momentum parameter $\gamma = 0$, CIM with momentum (algorithm 2) becomes precisely CIM (algorithm 1).

Remark 3.3.1. *Throughout the rest of the thesis, 'CIM' and 'CIM with momentum' will often be used interchangeably while referring to CIM with momentum, unless specified otherwise.*

3.4 Simulated Bifurcation (SB)

This section is based on [29].

Dynamics

Simulated Bifurcation (SB) is a Hamiltonian dynamical system with time-dependent Hamiltonian given by

$$H(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu) = \frac{m}{2} \sum_{i=1}^n y_i^2 + E(\mathbf{x}; \alpha, \beta, \mu) \quad (3.30)$$

where E is the potential function given by 3.1. The first part of the summation corresponds to kinetic energy depending only on momenta \mathbf{y} while the second part corresponds to potential energy depending only on positions \mathbf{x} . Parameters $\alpha, \beta, m > 0$ will be kept constant over time, while the parameter $\mu(t) \in \mathbb{R}$ will monotonically increase from certain initial to certain final value. The system of differential equations governing the dynamics for this Hamiltonian system is thus given by

$$\begin{aligned} \dot{\mathbf{x}}(t) &= \frac{\partial H^T}{\partial \mathbf{y}} (\mathbf{x}(t), \mathbf{y}(t); \alpha, \beta, m, \mu(t)) = m\mathbf{y}(t) \\ \dot{\mathbf{y}}(t) &= -\frac{\partial H^T}{\partial \mathbf{x}} (\mathbf{x}(t), \mathbf{y}(t); \alpha, \beta, m, \mu(t)) = \mathbf{f}(\mathbf{x}(t); \alpha, \beta, \mu(t)) \end{aligned} \quad (3.31)$$

where \mathbf{f} is the negative gradient of E , defined by equation (3.2) i.e.

$$\begin{aligned}\mathbf{f}(\mathbf{x}; \alpha, \beta, \mu) &= -\frac{\partial E^T}{\partial \mathbf{x}}(\mathbf{x}; \alpha, \beta, \mu) \\ &= -\alpha[x_1^3, \dots, x_n^3]^T + \alpha\mu\mathbf{x} + \beta J\mathbf{x}\end{aligned}\quad (3.32)$$

Written out further component-wise and omitting the time variable we get

$$\begin{aligned}\dot{x}_i &= \frac{\partial H}{\partial y_i} = my_i \\ \dot{y}_i &= -\frac{\partial H}{\partial x_i} = \alpha x_i(\mu - x_i^2) + \beta \sum_j J_{ij}x_j\end{aligned}\quad (3.33)$$

Algorithm

SB algorithm is a numerical simulation of the above dynamical system on a computer. The simulation is performed by symplectic Euler method rather than standard Euler method because it is more stable for simulating Hamiltonian systems. The update step with standard symplectic Euler method would thus be

$$\begin{aligned}x_i^{(k+1)} &= x_i^{(k)} + my_i^{(k)} \cdot \Delta_t \\ y_i^{(k+1)} &= y_i^{(k)} + [\alpha x_i^{(k+1)} \cdot (\mu(t_k) - (x_i^{(k+1)})^2) + \beta \sum_j J_{ij}x_j^{(k+1)}] \cdot \Delta_t\end{aligned}\quad (3.34)$$

where $t_k = \Delta_t k$ is the time discretization, having Δ_t fixed.

Additionally, as the authors proposed in the article [29], a modified symplectic Euler method provides even better results in terms of solution quality and computation speed. Since the computation of $\sum_j J_{ij}x_j^{(k+1)}$ is computationally the most expensive part, the Hamiltonian is split into $M + 1$ parts

$$\begin{aligned}H(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu) &= M \frac{H_1(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu)}{M} + H_2(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu) \\ H_1(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu) &:= \frac{m}{2} \sum_i y_i^2 + \frac{\alpha}{4} \sum_i (x_i^2 - \mu)^2 \\ H_2(\mathbf{x}, \mathbf{y}; \alpha, \beta, m, \mu) &:= -\frac{\beta}{2} \sum_{i,j} J_{ij}x_i x_j\end{aligned}\quad (3.35)$$

M is some positive integer.

Applying arguments about symplectic maps, a modified explicit symplectic Euler method for Simulated Bifurcation is obtained, which is called the SB algorithm.

Algorithm 3 SB

```

1: Initialize vectors  $\mathbf{x}, \mathbf{y}$  randomly around 0
2: for  $k$  in range(0,  $N_{\text{iter}}$ ) do
3:   for  $l$  in range(0,  $M$ ) do
4:      $x_i \leftarrow x_i + my_i \delta_t, \quad i = 1, \dots, n$ 
5:      $y_i \leftarrow y_i + [\alpha x_i \cdot (\mu(t_k) - x_i^2)] \cdot \delta_t, \quad i = 1, \dots, n$ 
6:   end for
7:    $y_i \leftarrow y_i + \beta \sum_{j=1}^n J_{ij} x_j \cdot \Delta_t, \quad i = 1, \dots, n$ 
8: end for

```

Δ_t is fixed discretized time interval and δ_t is its smaller refinement given by $\delta_t = \frac{\Delta_t}{M}$. Updates with respect to the first part of the Hamiltonian are refined by iterating $m = 0, \dots, M - 1$ sub-updates with smaller time step δ_t .

In figure 3.3 we see a visualization of SB algorithm on a system with two variables.

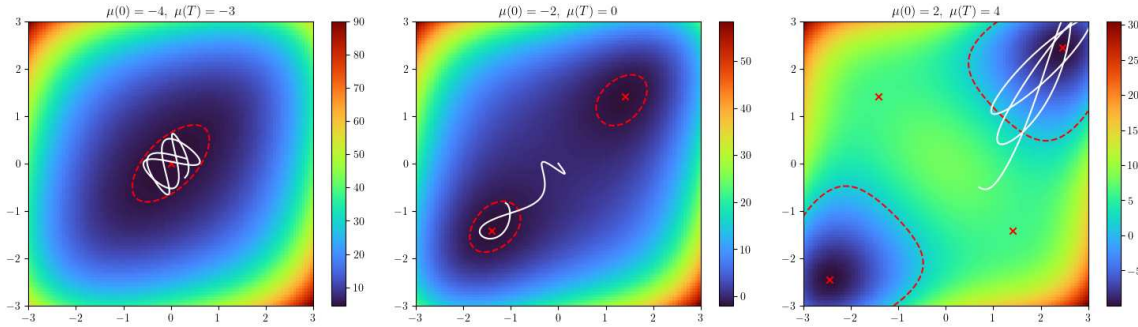


Figure 3.3: SB algorithm

Overview: This is a visual representation of SB algorithm on a system with two variables with coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. In the left subplot, the parameter μ did not cross the first bifurcation point, so the dynamical system revolves around the origin. As the time passes, μ increases, causing a change in the qualitative appearance of landscape E . In the middle subplot, the origin splits into two local minima (red crosses) which correspond to the solution of QUBO problem. In the right subplot, two new local minima emerge, corresponding to suboptimal solutions of QUBO. However, they are not as *attractive* for the dynamical system as the optimal solutions. In both cases, the dynamical system ends up circulating through the area (red dashed line) corresponding to an optimal solution of QUBO.

Details: The white line represents the trajectory obtained by running the SB algorithm. Red crosses denote local minima of E . The red dotted line represents the boundary for the feasible region of the Hamiltonian system, that is the level set of E at value equals to the total energy of the system at that time instance. The parameters are $\alpha = 1, \beta = 2$. Each subplot corresponds to a new run of the algorithm. The energy landscape E is plotted only for the final value of μ , that is $E(\cdot; \alpha, \beta, \mu(T))$ is plotted on each of the subplots. In each subplot, μ increases linearly from the starting to the ending value. Starting and ending values are respectively denoted as $\mu(0)$ and $\mu(T)$ on top of each subplot. Initial positions are $\mathbf{x} = (0, 0)$ in each subplot. For the leftmost subplot, initial positions are $\mathbf{x} = [0.3, -0.3]$, initial momenta are $\mathbf{y} = [0.5, 0.5]$, $\Delta_t = 0.1$, and the number of iterations is 100. For the middle subplot, initial positions are $\mathbf{x} = [0, 0]$, initial momenta are $\mathbf{y} = [0.3, -0.4]$, $\Delta_t = 0.05$, and the number of iterations is 150. For the rightmost subplot, initial positions are $\mathbf{x} = [0, 0]$, initial momenta are $\mathbf{y} = [0.3, -0.2]$, $\Delta_t = 0.05$, and the number of iterations is 150.

3.5 Mechanism of CIM and SB

Let us explain the mechanism of CIM and SB. Although CIM and SB are nonautonomous dynamical systems; in order to analyze them locally in time, it is sufficient to consider them as autonomous systems with their vector fields frozen at some time instance of interest. Indeed, using the connection theorem 2.1.9 and the fact that these vector fields change relatively *slowly* in time, we have that the error made by such approximation will be relatively small, especially for short time intervals.

The behaviour of these two systems around the first bifurcation point could be understood in the light of examples 2.1.11 and 2.1.10 which provide solutions of linear gradient and Hamiltonian systems. Indeed, by the Hartman-Grobman theorem 2.1.7, we have that the qualitative behaviour of CIM and SB around the origin is equivalent to the dynamics of the corresponding linearized systems. Thus, we need to analyze $\frac{\partial f}{\partial x} = -3\alpha \text{diag}(x_1^2, \dots, x_n^2) + \alpha\mu I + \beta J$ which at the origin is $\frac{\partial f}{\partial x}(\mathbf{0}) = \alpha\mu I + \beta J$. To understand this, we need to observe what happens in the reference frame which diagonalizes $\frac{\partial f}{\partial x}(\mathbf{0})$. Before the first bifurcation point, all eigenvalues of $\frac{\partial f}{\partial x}(\mathbf{0})$ are smaller than 0, so CIM will collapse towards the origin, while SB will revolve around it. As the time passes, and μ increases, some of the largest eigenvalues of $\frac{\partial f}{\partial x}(\mathbf{0})$ will cross 0 and become positive. Both the CIM and SB will start to rapidly expand towards these directions, while staying bounded or even contract in all other directions. Since these direction correspond to eigenvectors of J , both CIM and SB provide good approximate solutions of the QUBO problem at early stages of the algorithm.

After the system expands far enough from the origin, and as μ continues to increase, the behaviour around the origin stops being relevant. Instead, new local minima of E emerge and the system starts either converging towards them (CIM) or revolving around them (SB). Under the assumption that local minima corresponding to better QUBO solutions emerge earlier (for smaller μ) than other local minima, or that these local minima will have higher *attractivity*, we expect that the system will be able to converge towards these local minima and thus provide high-quality QUBO solutions.

As discussed in [37] for CIM, those variables which bifurcate from the origin at early stages *usually* do not return to the origin nor change their sign anymore. This potentially allows us to consider only the system reduced to those variables which did not bifurcate yet, while keeping others frozen.

Analyzing the bifurcation pattern (the relative order of emergence of local minima) would be essential for understanding the behavior of CIM. This would enable us to predict the performance of CIM and SB algorithms — provide us a way to tell when will these algorithms provide good solutions of QUBO problem and when it will fail to do so. However, it is not clear how to systematically treat and analyze this bifurcation pattern or develop theory about it. Analyzing this further is out of the scope of this thesis.

3.6 Comparing CIM and SB

Bifurcation pattern

In the figure 3.4 we compare the bifurcation behaviour of a single-variable system and the evolution of the dynamics through time.

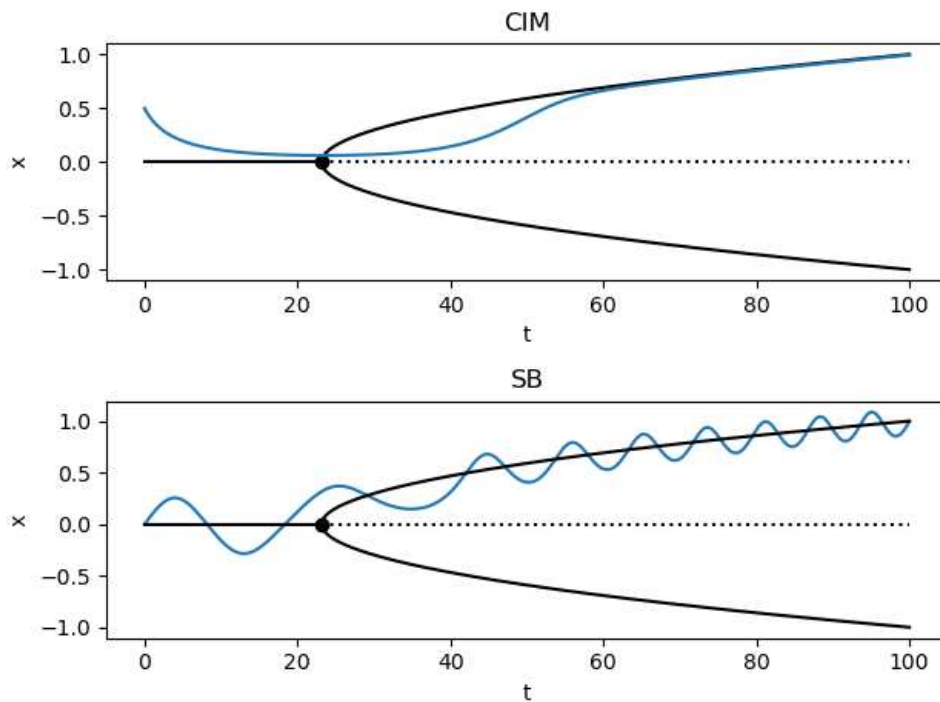


Figure 3.4: Comparison of CIM and SB with one variable

Overview: This is a visual representation of CIM and SB algorithms performing on only one variable. As the time evolves, and μ crosses the first bifurcation point (black dot), the landscape function E changes its qualitative appearance and thus forces the dynamical system to bifurcate towards one of the two stable branches (solid black lines).

Details: Solid black lines represent the positions of local minima of landscape function E .

Dashed line represents the position of an unstable stationary point. Black dot is the bifurcation point. Blue lines represent the evolution of CIM and SB algorithms. These are locally a good approximation of the exact solutions of CIM and SB dynamical systems. The following parameters have been used. $J = [0]$, $\alpha = 0.5$, $\beta = 0.5$, $\mu_0 = -0.3$, $\mu_1 = 1.0$, $\Delta_t = 0.1$. The number of iterations is 1000. Initial conditions for CIM are $x_1(0) = 0.5$, while for SB are $x_1(0) = 0.0$, $y_1(0) = 0.1$. Momentum is not used for this CIM simulation.

Relation between CIM and SB

Although the momentum parameter is usually kept smaller than 1, it is very interesting what happens when $\gamma = 1$. In this case, CIM with momentum (algorithm 2) potentially becomes a simulation of SB dynamical system. That is, we can choose hyperparameters in CIM with momentum in such a way that the update step becomes a symplectic Euler method applied to SB dynamics.

Let us distinguish between CIM's and SB's parameters with a superscript 'CIM' and 'SB'.

For given parameters for SB algorithm $\alpha^{\text{SB}}, \mu_0^{\text{SB}}, \mu_1^{\text{SB}}, \beta^{\text{SB}}, \Delta_t^{\text{SB}}, m^{\text{SB}}$, and assuming that $M^{\text{SB}} = 1$, the SB algorithm could be written like this (notice that the order of equations for x_i and y_i is flipped but this remains a correct numerical simulation of the dynamics as described in section 2.2)

Algorithm 4 variant of SB with $M = 1$

- 1: Initialize vectors \mathbf{x}, \mathbf{y} randomly around 0
 - 2: **for** k in range(0, N_{iter}) **do**
 - 3: $y_i \leftarrow y_i + \left[\alpha^{\text{SB}} x_i \cdot (\mu^{\text{SB}}(t_k) - x_i^2) + \beta^{\text{SB}} \sum_{j=1}^n J_{ij} x_j \right] \cdot \Delta_t^{\text{SB}}, \quad i = 1, \dots, n$
 - 4: $x_i \leftarrow x_i + m^{\text{SB}} y_i \Delta_t^{\text{SB}}, \quad i = 1, \dots, n$
 - 5: **end for**
-

Define the parameters for CIM as follows

$$\begin{aligned} \gamma^{\text{CIM}} &= 1 & \alpha^{\text{CIM}} &= \alpha^{\text{SB}} \Delta_t^{\text{SB}} \\ \beta^{\text{CIM}} &= \beta^{\text{SB}} \Delta_t^{\text{SB}} & \mu^{\text{CIM}} &= \mu^{\text{SB}} \\ \Delta_t^{\text{CIM}} &= m^{\text{SB}} \Delta_t^{\text{SB}} \end{aligned}$$

Plugging in the above parameters into CIM with momentum (algorithm 2), we obtain

Algorithm 5 CIM with momentum mimicking SB

- 1: Initialize vectors \mathbf{x}, \mathbf{y} randomly around $\mathbf{0}$
 - 2: **for** k in range(0, N_{iter}) **do**
 - 3: $y_i \leftarrow y_i + \alpha^{\text{SB}} \Delta_t^{\text{SB}} x_i (\mu^{\text{SB}}(t_k) - x_i^2) + \beta^{\text{SB}} \Delta_t^{\text{SB}} \sum_{j=1}^n J_{ij} x_j, \quad i = 1, \dots, n$
 - 4: $x_i \leftarrow x_i + y_i \cdot m^{\text{SB}} \Delta_t^{\text{SB}}, \quad i = 1, \dots, n$
 - 5: **end for**
-

which is precisely the above variant of SB, algorithm 4. Although these implementations of CIM and SB are not completely equivalent, CIM with momentum has the ability to simulate the dynamics of SB if the parameters are selected appropriately (especially $\gamma = 1$). In this context, CIM with momentum is in some sense a generalization of SB algorithm.

3.7 Introduction to SimCIM and bSB

Let us define the following energy functional

$$E(\mathbf{x}; \alpha, \beta) = \frac{\alpha}{2} \sum_i x_i^2 - \frac{\beta}{2} \sum_{i,j} J_{ij} x_i x_j, \quad \text{for } \mathbf{x} \in [-1, +1]^n \quad (3.36)$$

$$E(\mathbf{x}; \alpha, \beta) = +\infty, \quad \text{otherwise} \quad (3.37)$$

which will be used both in sections 3.8 and 3.9.

Let us define the corresponding vector field \mathbf{f} as a negative gradient of the *regular* part of E (3.36)

$$\begin{aligned} \mathbf{f}(\mathbf{x}; \alpha, \beta) &:= -\frac{\partial E}{\partial \mathbf{x}}^T(\mathbf{x}; \alpha, \beta) \\ &= -\alpha \mathbf{x} + \beta J \mathbf{x} \end{aligned} \quad (3.38)$$

This energy functional is a function of position \mathbf{x} and parameters α, β denoted after the semicolon.

Remark 3.7.1. *Sometimes when we choose and fix some concrete values for α and β , we will pretend that E is only a function of position \mathbf{x} and thus write $E(\mathbf{x}) \equiv E(\mathbf{x}; \alpha, \beta)$.*

For SimCIM and bSB we will only be interested in the dynamics inside the region $[-1, +1]^n$, and we actually want to prevent the system from going outside of this region. That's why we formally define that $E(\mathbf{x}) = +\infty$ outside of $[-1, +1]^n$. This potentially complicates the mathematical formalism but it will not be an issue for the analysis because we will only be interested in the local behaviour in time in the regular region. When some components hit the wall, these components will be (temporarily) frozen so all the others will evolve in their (restricted) regular region. That is why we further define the vector field \mathbf{g} which handles these boundary conditions, and will thus be suitable for defining both SimCIM and bSB dynamical systems.

$$\mathbf{g}(\mathbf{x}; \alpha, \beta) := A(\mathbf{x}; \alpha, \beta) \cdot \mathbf{f}(\mathbf{x}; \alpha, \beta) \quad (3.39)$$

where the diagonal matrix A regulates which components are allowed to evolve and which should be paused because they hit the boundary and additionally their vector field components point out of the regular region. That is, we have $A(\mathbf{x}; \alpha, \beta) = \text{diag}(a_1, \dots, a_n)$; $a_i = 1$ if $x_i \in \langle -1, +1 \rangle$ or $(x_i \in \{-1, +1\} \ \& \ x_i f_i < 0)$, $a_i = 0$ otherwise.

It is essential to analyze the energy functional E with respect to parameters because it defines vector fields \mathbf{f} and \mathbf{g} ; and thus governs the SimCIM and bSB dynamical systems in sections 3.8 and 3.9. This energy function will vary in time itself because the parameters

will vary in time. Concretely, $\beta > 0$ will be constant over time, while it is crucial that $\alpha \equiv \alpha(t)$ will decrease monotonically from some initial value α_0 to some final value α_1 . This decreasing regime varies among algorithms. In bSB, $\alpha(t)$ changes linearly, i.e. $\alpha(t) = \alpha_0 - \frac{\alpha_0 - \alpha_1}{T}t$. In the original SimCIM algorithm, $\alpha(t)$ is a sigmoid function given by the hyperbolic tangent law $\alpha(t) = -(\alpha_1 - \alpha_0) \tanh(A(t - \frac{T}{2})) + \alpha_1$, where $A > 0$ is some positive constant, for example $A = 3$. Although different regimes for $\alpha(t)$ produce different solutions and some regimes might be better than the others, we cannot predict which regime would be the best. The choice of the regime is not carved into the stone and it even seems arbitrary to a certain degree. Nevertheless, what is important is that $\alpha(t)$ continuously, monotonically, and relatively *slowly* decreases in time starting from α_0 and ending with α_1 . Thus, we will usually make α linearly decrease from α_0 to α_1 , as it is originally proposed for bSB.

Let us explain the behaviour of energy functional E for arbitrary $\alpha \in \mathbb{R}$. Since J is symmetric, it is thus orthogonally diagonalizable. Let $U^T J U = \Lambda = \text{diag}(\lambda_1, \dots, \lambda_n)$ be the diagonalization of J such that eigenvalues of J are sorted in descending way $\lambda_{\max} = \lambda_1 \geq \dots \geq \lambda_n = \lambda_{\min}$. Denote the change of variables $\bar{\mathbf{x}} := U^T \mathbf{x}$. We have that

$$E(\mathbf{x}) = \frac{\alpha}{2} \mathbf{x}^T \mathbf{x} - \frac{\beta}{2} \mathbf{x}^T J \mathbf{x} = \frac{\alpha}{2} \bar{\mathbf{x}}^T \bar{\mathbf{x}} - \frac{\beta}{2} \bar{\mathbf{x}}^T \Lambda \bar{\mathbf{x}} = \bar{\mathbf{x}}^T \left(\frac{\alpha}{2} I - \frac{\beta}{2} \Lambda \right) \bar{\mathbf{x}} \quad (3.40)$$

Here the nice thing is that $(\frac{\alpha}{2} I - \frac{\beta}{2} \Lambda)$ is a diagonal matrix so E represented in these transformed coordinates is more comprehensible.

Let us further calculate the second differential of E .

$$\frac{\partial^2 E}{\partial \mathbf{x}^2} = \alpha I - \beta J \quad (3.41)$$

The eigenvalues of $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ are $\alpha - \beta \lambda_1, \dots, \alpha - \beta \lambda_n$ with corresponding eigenvectors being the same as J 's. The sign of these eigenvalues will locally determine the qualitative behavior of the dynamical systems which follow in the next sections.

Lemma 3.7.2. *For $\alpha > \beta \lambda_{\max}$, E is convex with the global minimum at the origin $\mathbf{x}_0 = \mathbf{0}$.*

Proof. On one hand, $E(\mathbf{0}) = 0$. Write E in the form $E(\mathbf{x}) = \frac{\alpha}{2} \mathbf{x}^T \mathbf{x} - \frac{\beta}{2} \mathbf{x}^T J \mathbf{x}$ so because $\mathbf{x}^T J \mathbf{x} \leq \lambda_{\max} \mathbf{x}^T \mathbf{x}$, we have

$$E(\mathbf{x}) \geq \frac{\alpha}{2} \mathbf{x}^T \mathbf{x} - \frac{\beta}{2} \lambda_{\max} \mathbf{x}^T \mathbf{x} = \frac{1}{2} (\alpha - \beta \lambda_{\max}) \mathbf{x}^T \mathbf{x} \quad (3.42)$$

So, for $\mathbf{x} \neq \mathbf{0}$, this implies $E(\mathbf{x}) > 0$ i.e. the origin is the global minimum.

Taking the minimum eigenvalue of $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ provides us with

$$\lambda_{\min} \left(\frac{\partial^2 E}{\partial \mathbf{x}^2} \right) = \alpha - \beta \lambda_{\max}(J) > 0$$

so all the eigenvalues of the second differential are positive, meaning that E is convex (proposition 2.3.2). \square

One more question which should be addressed are the starting and ending values α_0 and α_1 . There is some degree of freedom in choosing them, but nothing interesting happens when $\alpha > \beta\lambda_{\max}$ as E is convex and centered at the origin. For $\alpha \leq \beta\lambda_{\max}$, E starts to change because some of its eigenvalues start being negative.

The condition

$$\alpha = \beta\lambda_{\max}(J) \quad (3.43)$$

is thus called the *first bifurcation point*.

Thus, both for the gradient and for the Hamiltonian system with landscape E , it makes sense starting the dynamics with

$$\alpha_0 \leq \beta\lambda_{\max}(J) \quad (3.44)$$

Selecting the concrete values of α_0 and α_1 should be determined by empirical evidence i.e. by experimenting which parameter provides the best result.

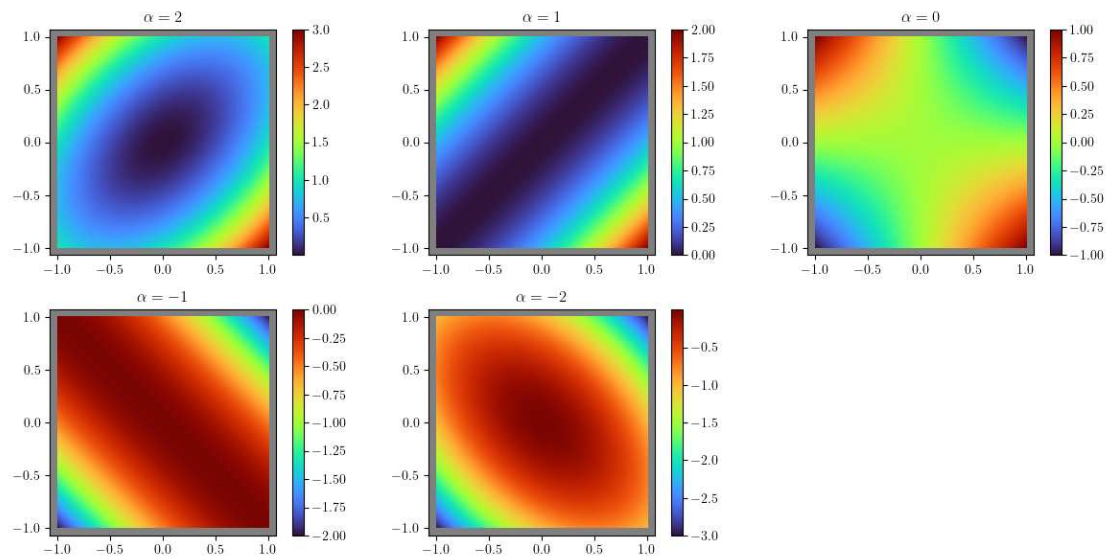


Figure 3.5: Visualization of energy landscape E for SimCIM and bSB

Overview: This is a visual representation of energy landscape E for a system with two variables and a coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. The landscape undergoes a qualitative change as α decreases.

Details: $\beta = 1$. Values of α are denoted on top of each subplot. $\alpha = 1$ and $\alpha = -1$ are critical values where some eigenvalues of D^2E change their sign, thus qualitatively changing the landscape. The gray border represents the wall beyond which the dynamics cannot occur.

3.8 Simulated Coherent Ising machine (SimCIM)

This section is based on [30].

Dynamics

SimCIM is a gradient dynamical system with time-dependent landscape function $E(\mathbf{x}; \alpha, \beta)$ given by equation (3.36). Parameter $\beta > 0$ is kept constant through time but it is essential that parameter $\alpha(t)$ is monotonically decreased in time as it will be responsible for generating bifurcations. The system of differential equations governing the dynamics for this gradient system is thus given by

$$\dot{\mathbf{x}}(t) = -\frac{\partial E^T}{\partial \mathbf{x}}(\mathbf{x}(t), \alpha(t), \beta) \quad (3.45)$$

$$= \mathbf{g}(\mathbf{x}(t); \alpha(t), \beta) \quad (3.46)$$

where \mathbf{g} is the vector field defined by equation (3.39). It is precisely equal to the negative gradient of the landscape E in the interior region. On the boundary region, it is equal zero for those components which point out of the allowed region $[-1, +1]^n$, so the system will always stay inside the allowed region.

Written out component-wise without denoting the dependence on time we get

$$\begin{cases} \dot{x}_i = -\alpha x_i + \beta \sum_j J_{ij} x_j, & \text{when } x_i \in \langle -1, +1 \rangle \\ \dot{x}_i = 0, & \text{or } (x_i \in \{-1, +1\} \ \& \ x_i f_i < 0) \\ & \text{otherwise} \end{cases} \quad (3.47)$$

Algorithm

SimCIM algorithm is a numerical simulation of the above dynamical system by Euler method.

Algorithm 6 SimCIM

- 1: Initialize vector $\mathbf{x}^{(0)}$ randomly around $\mathbf{0}$
 - 2: **for** k in range(0, N_{iter}) **do**
 - 3: $x_i^{(k+1)} \leftarrow x_i^{(k)} + \left(-\alpha(t_k)x_i^{(k)} + \beta \sum_{j=1}^n J_{ij}x_j^{(k)} \right) \Delta_t, \quad i = 1, \dots, n$
 - 4: $x_i^{(k+1)} \leftarrow \phi(x_i^{(k+1)}), \quad i = 1, \dots, n$
 - 5: **end for**
-

In algorithm 6, ϕ is the clamping function defined by

$$\phi(x) = \begin{cases} -1, & \text{for } x < -1 \\ x, & \text{for } -1 \leq x \leq 1 \\ 1, & \text{for } x > 1 \end{cases} \quad (3.48)$$

In figure 3.6 we observe that until α crosses the first bifurcation point ($\alpha = 1$), the system collapses towards the origin. As α crosses the first bifurcation point, the landscape function E changes such that the directions corresponding to $(+, +)$ and $(-, -)$ become directions where E decreases, while directions $(+, -)$ and $(-, +)$ stay directions where E increases. That's why in the top-right and bottom-left subplots we have that the system evolves straight to the point $(-1, -1)$. In the bottom-middle figure the system also evolves towards the $(-, -)$ configuration, but it hits the wall $y = -1$ in the meantime and after that continues evolving only in x component, and finally ends up in the state $(-1, -1)$.

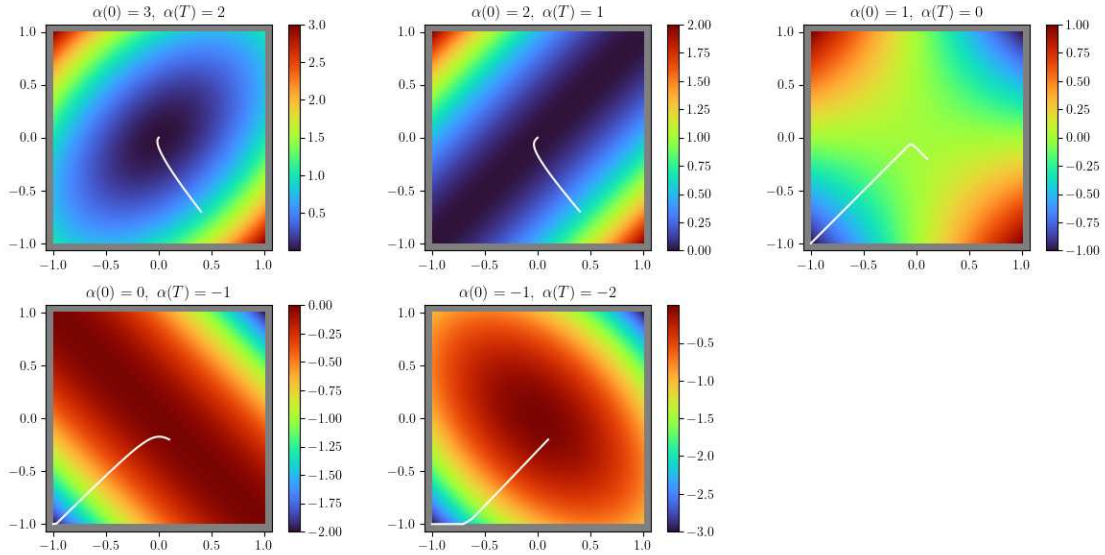


Figure 3.6: SimCIM algorithm

Overview: This is a visual representation of SimCIM algorithm on a system with two variables with a coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. In the top-left and top-middle subplots, the parameter α did not cross the first bifurcation point so the dynamical system collapses towards the origin (starting far away from it). In all other subplots, the dynamical system starts close to the origin, but converges to QUBO optimum solutions.

Details: The gray border represents the wall beyond which the dynamics cannot occur. The white line represents the trajectory obtained by running the SimCIM algorithm. Each subplot corresponds to a new run of the algorithm. The energy landscape E is plotted only for the final value of α , that is $E(\cdot; \alpha(T), \beta)$ is plotted on each of the subplots. $\beta = 1$. In each subplot, α decreases linearly from the starting value (denoted as $\alpha(0)$) to the ending value (denoted as $\alpha(T)$). The number of iterations is 200, and the step size is $\Delta_t = 0.05$.

Initial positions are $\mathbf{x}(0) = (0.4, -0.7)$ for upper left and upper middle subplot, while $\mathbf{x}(0) = (0.1, -0.2)$ for other plots.

Adding momentum

Since SimCIM is essentially a gradient descent algorithm, it makes sense to add momentum to it in order to make the convergence faster, similarly as it was done for CIM in section 3.3 (based on section 3.3 and [47]). Momentum is actually proposed in the SimCIM article [30] as well.

First we choose the momentum parameter $0 \leq \gamma \leq 1$. Momenta will be stored in vector \mathbf{y} .

Algorithm 7 SimCIM with momentum

- 1: Initialize vectors $\mathbf{x}^{(0)}, \mathbf{y}^{(0)}$ randomly around $\mathbf{0}$
 - 2: **for** k in range(0, N_{iter}) **do**
 - 3: $y_i^{(k+1)} \leftarrow \gamma y_i^{(k)} + \left(-\alpha(t_k)x_i^{(k)} + \beta \sum_{j=1}^n J_{ij}x_j^{(k)} \right), \quad i = 1, \dots, n$
 - 4: $x_i^{(k+1)} \leftarrow x_i^{(k)} + y_i^{(k+1)}\Delta t, \quad i = 1, \dots, n$
 - 5: $y_i^{(k+1)} \leftarrow 0, \quad \text{if } x_i^{(k+1)} \notin [-1, +1], \quad i = 1, \dots, n$
 - 6: $x_i^{(k+1)} \leftarrow \phi(x_i^{(k+1)}), \quad i = 1, \dots, n$
 - 7: **end for**
-

For momentum parameter $\gamma = 0$, SimCIM with momentum (algorithm 7) becomes precisely SimCIM (algorithm 6).

Remark 3.8.1. *'SimCIM' and 'SimCIM with momentum' will often be used interchangeably, while referring to SimCIM with momentum, unless specified otherwise.*

Exact solution

In this section, an exact solution of SimCIM dynamical system is derived. This solution holds until the system hits the wall. After it hits the wall, a similar approach for deriving the exact solution could be used but considering only those variables that did not hit the wall, while fixing others.

Let us revise the dynamical system describing SimCIM (3.47) for the interior region i.e. until none of the components hit the wall

$$\dot{\mathbf{x}} = -\alpha\mathbf{x} + \beta J\mathbf{x} \quad (3.49)$$

Let $U^T J U = \Lambda$ be an orthogonal diagonalization of J and denote the change of variables $\bar{\mathbf{x}} = U^T \mathbf{x}$. As described in the previous subsection, the system of coupled equations (3.49) is brought down to

$$\dot{\bar{x}}_i = -\alpha\bar{x}_i + \beta\lambda_i\bar{x}_i, \quad i = 1, \dots, n \quad (3.50)$$

which is a system of uncoupled equations of the same form, so it is sufficient to consider only single one of them. So we drop the index i in what follows. Suppose that α has the linear form $\alpha(t) = \alpha_0 - \frac{\alpha_0 - \alpha_1}{T}t$. We are thus interested in solving equation of the form

$$\dot{\bar{x}}(t) = (\beta\lambda - \alpha_0)\bar{x}(t) + \frac{\alpha_0 - \alpha_1}{T}t\bar{x}(t) \quad (3.51)$$

But this is a linear equation of the first order, so it has the solution

$$\bar{x}(t) = \bar{x}(0) \exp\left(\frac{\alpha_0 - \alpha_1}{2T}t^2 + (\beta\lambda - \alpha_0)t\right) \quad (3.52)$$

Since α is decreasing, i.e. $\alpha_0 > \alpha_1$, we see from equation (3.52) that the solution will eventually start rapidly increasing because of the t^2 term. For $\alpha_0 - \beta\lambda \leq 0$ the solution will immediately start expanding, while for $\alpha_0 - \beta\lambda > 0$ the solution will first collapse towards 0 and then at some point in time start expanding.

Thus, SimCIM dynamical system has an exact component-wise solution given by (3.52) in the reference frame which diagonalizes J . Changing the reference frame back to $\mathbf{x} = U\bar{\mathbf{x}}$ we obtain the exact solution of SimCIM until it hits the wall.

3.9 Ballistic Simulated Bifurcation (bSB)

This section is based on [31].

Dynamics

Ballistic Simulated Bifurcation (bSB) is a Hamiltonian dynamical system with time-dependent Hamiltonian given by

$$H(\mathbf{x}, \mathbf{y}; \alpha, \beta, m) = \frac{m}{2} \sum_i y_i^2 + E(\mathbf{x}; \alpha, \beta) \quad (3.53)$$

where E is the potential function given by equation (3.36). The first part of the summation corresponds to the kinetic energy – depending only on momenta \mathbf{y} , while the second part corresponds to the potential energy – depending only on positions \mathbf{x} . Parameters $m, \beta > 0$ are constant in time, while parameter α decreases monotonically (usually linearly) from certain starting value α_0 to certain ending value α_1 . The system of differential equations governing the dynamics for this Hamiltonian system is thus given by

$$\begin{aligned} \dot{\mathbf{x}}(t) &= \frac{\partial H^T}{\partial \mathbf{y}} (\mathbf{x}(t), \mathbf{y}(t); \alpha(t), \beta, m) \\ &= m\mathbf{y} \\ \dot{\mathbf{y}}(t) &= -\frac{\partial H^T}{\partial \mathbf{x}} (\mathbf{x}(t), \mathbf{y}(t); \alpha(t), \beta) \\ &= -\frac{\partial E^T}{\partial \mathbf{x}} (\mathbf{x}(t), \alpha(t), \beta, m) \\ &= \mathbf{g}(\mathbf{x}, \alpha(t), \beta) \end{aligned} \quad (3.54)$$

and it holds when the system is in the interior region. \mathbf{g} is the vector field defined by equation (3.39). It is precisely equal to the negative gradient of the energy landscape E in the interior region. On the boundary region, it is equal zero for those components which point out of the allowed region $[-1, +1]^n$.

Furthermore, in order to model the boundary of $[-1, +1]^n$ as a perfectly inelastic wall, we do not only want to set the acceleration g_i to zero, but we also need to set the velocity y_i to zero itself when certain component x_i hits the boundary and the acceleration points out of the allowed region. This will prevent the system from going out of the allowed region. This way, the system loses kinetic energy as components hit the wall.

Written out further component-wise and omitting the time variable we get the system of equations describing bSB completely

$$\begin{cases} \dot{x}_i = my_i \\ \dot{y}_i = -\alpha x_i + \beta \sum_j J_{ij} x_j, \\ y_i = 0, \end{cases} \quad \begin{array}{l} \text{when } x_i \in \langle -1, +1 \rangle \\ \text{or } (x_i \in \{-1, +1\} \ \& \ x_i f_i < 0) \\ \text{otherwise} \end{array} \quad (3.55)$$

Algorithm

Ballistic Simulated Bifurcation algorithm is a numerical simulation of the above dynamical system. The simulation is performed by symplectic Euler method rather than standard Euler method because it is more stable for simulating Hamiltonian systems.

Algorithm 8 bSB

- 1: Initialize vectors $\mathbf{x}^{(0)}, \mathbf{y}^{(0)}$ randomly around $\mathbf{0}$
 - 2: **for** k in $\text{range}(0, N_{\text{iter}})$ **do**
 - 3: $y_i^{(k+1)} \leftarrow y_i^{(k)} + (-\alpha(t_k)x_i^{(k)} + \beta \sum_{j=1}^n J_{ij}x_j^{(k)}) \Delta_t, \quad i = 1, \dots, n$
 - 4: $x_i^{(k+1)} \leftarrow x_i^{(k)} + my_i^{(k+1)} \Delta_t, \quad i = 1, \dots, n$
 - 5: $y_i^{(k+1)} \leftarrow 0, \quad \text{if } x_i^{(k+1)} \notin [-1, +1], \quad i = 1, \dots, n$
 - 6: $x_i^{(k+1)} \leftarrow \phi(x_i^{(k+1)}), \quad i = 1, \dots, n$
 - 7: **end for**
-

In algorithm 8, ϕ is the clamping function defined by (3.48). Figure 3.7 shows the behaviour of bSB algorithm.

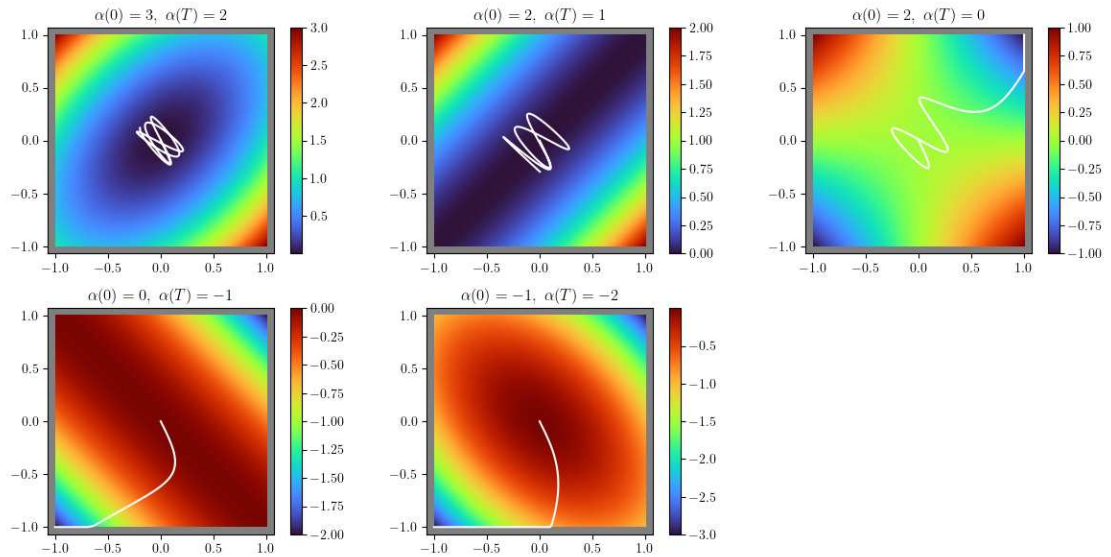


Figure 3.7: bSB algorithm

Overview: This is a visual representation of bSB algorithm on a system with two variables with coupling term $J_{12} = J_{21} = 1$. Optimum solutions of the corresponding QUBO problem are $(+, +)$ and $(-, -)$. In the top-left and top-middle subplots, the parameter α did not cross the first bifurcation point so the dynamical system circulates around the origin. In the top-right subplot, and in the bottom subplots, the dynamical system converges to QUBO optimum solutions.

Details: The gray border represents the wall beyond which the dynamics cannot occur. The white line is the trajectory obtained by running the bSB algorithm. Each subplot corresponds to a new run of the algorithm. The energy landscape E is plotted only for the final value of α , that is $E(\cdot; \alpha(T), \beta)$ is plotted on each of the subplots. $\beta = 1$. In each subplot, α decreases linearly from the starting value (denoted $\alpha(0)$) to the ending value (denoted $\alpha(T)$). The number of iterations is 250, and the step size is $\Delta_t = 0.05$. Initial positions are $\mathbf{x}(0) = (0.0, 0.0)$ and initial momenta are $\mathbf{y}(0) = (0.2, -0.4)$.

Exact solution

In this section, an exact solution of the bSB dynamical system is provided. This solution holds until the system hits the wall. After it hits the wall, a similar approach for deriving the exact solution could be used but considering only those variables that did not hit the wall, while fixing others.

Let us revise the dynamical system describing bSB in the interior region i.e. until none of the components hit the wall

$$\begin{aligned}\dot{\mathbf{x}} &= m\mathbf{y} \\ \dot{\mathbf{y}} &= -\alpha\mathbf{x} + \beta J\mathbf{x}\end{aligned}\tag{3.56}$$

Let $U^T J U = \Lambda$ be an orthogonal diagonalization of J and denote the change of variables $\bar{\mathbf{x}} = U^T \mathbf{x}$. By taking the second derivative, and changing the reference frame to $\bar{\mathbf{x}}$, the system of coupled equations (3.56) is brought down to

$$\ddot{\bar{x}}_i = m(\beta\lambda_i - \alpha)\bar{x}_i, \quad i = 1, \dots, n\tag{3.57}$$

which is a system of uncoupled equations of the same form, so it is sufficient to consider only single one of them. So we drop the index i in what follows. α is a linear function so suppose it has the form $\alpha(t) = \alpha_0 - \frac{\alpha_0 - \alpha_1}{T}t$. Now, we will shift and scale the solution to reduce the problem to Airy equation (2.53). Define

$$x'(t) := \bar{x} \left(\sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}} t + \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right)\tag{3.58}$$

Now calculate

$$\begin{aligned}\ddot{x}'(t) &= \sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}}^2 \ddot{\bar{x}} \left(\sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}} t + \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \\ &= \sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}}^2 m \left(\beta\lambda - \alpha_0 - \frac{\alpha_1 - \alpha_0}{T} \left(\sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}} t + \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \right) \cdot \dots \\ &\quad \cdot \bar{x} \left(\sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}} t + \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \\ &= -\frac{\alpha_1 - \alpha_0}{T} \frac{T}{m(\alpha_0 - \alpha_1)} m t x'(t) + \dots \\ &\quad + \sqrt[3]{\frac{T}{m(\alpha_0 - \alpha_1)}}^2 m \left(\beta\lambda - \alpha_0 - \frac{\alpha_1 - \alpha_0}{T} \cdot \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) x'(t) \\ &= t x'(t)\end{aligned}\tag{3.59}$$

so x' solves the Airy equation (2.53).

Conversely, let $x'(t) = C_1A(t) + C_2B(t)$ be any solution of Airy equation (2.53) ($C_1, C_2 \in \mathbb{R}$ are arbitrary constants, while A, B are linearly independent solutions of Airy equation). By taking

$$\bar{x}(t) = x' \left(\sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}} \left(t - \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \right) \quad (3.60)$$

we get

$$\begin{aligned} \ddot{\bar{x}}(t) &= \sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}}^2 \ddot{x}' \left(\sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}} \left(t - \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \right) \\ &= \sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}}^2 \cdot \dots \\ &\quad \cdot \left(\sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}} \left(t - \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \right) \cdot x' \left(\sqrt[3]{\frac{m(\alpha_0 - \alpha_1)}{T}} \left(t - \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \right) \quad (3.61) \\ &= \frac{m(\alpha_0 - \alpha_1)}{T} \left(t - \frac{(\beta\lambda - \alpha_0)T}{\alpha_1 - \alpha_0} \right) \bar{x}(t) \\ &= m \left(-\frac{(\alpha_1 - \alpha_0)}{T} t + \beta\lambda - \alpha_0 \right) \bar{x}(t) \\ &= m(\beta\lambda - \alpha(t)) \bar{x}(t) \end{aligned}$$

which is exactly (3.57).

Thus, bSB dynamical system has an exact solution given by (3.60) in the reference frame which diagonalizes J . Choosing constants C_1, C_2 such that initial conditions are satisfied, and changing the reference frame back to $\mathbf{x} = U\bar{\mathbf{x}}$ we obtain an exact solution of bSB until it hits the wall.

3.10 Mechanism of SimCIM and bSB

Let us explain the mechanism of SimCIM and bSB. Although SimCIM and bSB are nonautonomous dynamical systems, by the connection theorem 2.1.9, in order to analyze such system locally (in time), it is sufficient to consider it as an autonomous system with frozen vector field at some time instance of interest. The mechanism of SimCIM and bSB can now be understood in the light of examples 2.1.11 and 2.1.10 which provide solutions of linear gradient and Hamiltonian systems. By changing the reference frame to the one that diagonalizes the coupling matrix J , one obtains a comprehensible behaviour of the dynamical system. As the parameter α decreases through time, the qualitative appearance of E changes. That is, some of the largest eigenvalues of $\frac{\partial^2 E}{\partial \mathbf{x}^2} = -\frac{\partial \mathbf{f}}{\partial \mathbf{x}}$ cross zero and become negative. Until that first bifurcation point, SimCIM dynamical system will collapse towards the origin, and bSB will revolve around the origin. After the first bifurcation point, the system will start to rapidly expand towards eigenvectors corresponding to smallest (now negative) eigenvalues of $\frac{\partial^2 E}{\partial \mathbf{x}^2}$ i.e. largest eigenvalues of J . This way, the system provides good approximate solutions of QUBO at early stages of the algorithm. This behaviour holds until some components hit the wall. After hitting the wall, this argument does not hold anymore in the reference frame which diagonalizes J . However, a similar argument can be applied, but one needs to consider only those components which have not hit the wall yet and diagonalize such submatrix.

As discussed in [37] for CIM, those variables which bifurcate from the origin at early stages *usually* do not return to the origin nor change their sign anymore. This seems consistent with the clipping mechanism for SimCIM and bSB, and could be potentially used for analyzing the system which is reduced to only those variables which did not hit the wall yet.

There is one more mechanism applied both to SimCIM and bSB and it is regarding the final state of the system. The moment when $\alpha(t) = 0$ has a significant role because at that point, if the system has bifurcated enough, it will provide a 1-opt solution. This will be discussed in the following two sections. In terms of parameter settings, it would make sense to stop the dynamics at that point and thus make $\alpha_1 = 0$. However, sometimes it is found that making α_1 slightly smaller than 0 provides better solution. It does not hurt making α_1 smaller than 0 as long as we track the best possible solution found, so we still have a chance of capturing that 1-opt solution at $\alpha(t) = 0$, but in case that required conditions are not met at $\alpha(t) = 0$, we still have a chance of finding better solution for $\alpha(t) < 0$. Making α_1 too small however usually is not good because we want $\alpha(t)$ to change slowly in time, in order for the system to be able to evolve properly.

Let us now explain why SimCIM and bSB have a tendency of generating 1-opt solutions for QUBO.

Final state

Let us observe what happens at (usually the final) time $\alpha(t) = 0$. Assume that the system has bifurcated completely in a sense that $\forall i, x_i \in \{-1, +1\}$. Assume also that the component i has stopped in a sense that $g_i = 0$. For SimCIM this means that the component i has stopped, while for bSB it means that the component i has zero acceleration (which combined with the fact that the boundary was hit means as well that this component has stopped). This implies $x_i f_i \geq 0$. The solution provided by SimCIM and bSB is, as usual, $\sigma_i := \text{sign } x_i$ which in this case provides $\sigma_i = x_i$. We have, since $\alpha(t) = 0$,

$$\beta \sum_{j=1}^n J_{ij} \sigma_i \sigma_j = x_i \beta \sum_{j=1}^n J_{ij} x_j = x_i f_i \geq 0 \quad (3.62)$$

Flipping the i -th binary variable provides a change in QUBO functional (see (1.22))

$$\begin{aligned} \Delta_i Q &= Q(\sigma_1, \dots, \sigma_{i-1}, -\sigma_i, \sigma_{i+1}, \dots, \sigma_n) - Q(\sigma_1, \dots, \sigma_{i-1}, \sigma_i, \sigma_{i+1}, \dots, \sigma_n) \\ &= -4 \sum_{j=1}^n J_{ij} \sigma_i \sigma_j \end{aligned} \quad (3.63)$$

This, combined with (3.62), provides

$$\Delta_i Q \leq 0 \quad (3.64)$$

so for the maximization task which we are interested in here, flipping the value of σ_i cannot improve the solution. If (3.62) holds for each i , i.e. the system has stopped completely, then the solution obtained by SimCIM or bSB algorithm is, by definition 1.1.3, a 1-opt solution. That is why SimCIM and bSB have a tendency to provide QUBO solutions which are 1-opt.

3.11 Relation between SimCIM and bSB

The relation between SimCIM and bSB is completely analogous to the relation between CIM and SB.

For momentum parameter $\gamma = 0$, SimCIM with momentum (algorithm 7) becomes precisely SimCIM (algorithm 6).

Although the momentum parameter is usually kept smaller than 1, it is very interesting what happens when $\gamma = 1$. In this case, SimCIM with momentum (algorithm 7) becomes precisely bSB algorithm (algorithm 8). That is, we can choose hyperparameters in SimCIM with momentum in such a way that the update step becomes exactly the same as the update step of bSB algorithm.

Let us distinguish between SimCIM's and bSB's parameters with a superscript "SimCIM" and "bSB".

For given parameters for bSB algorithm, $\alpha_0^{\text{bSB}}, \alpha_1^{\text{bSB}}, \beta^{\text{bSB}}, \Delta_t^{\text{bSB}}, m^{\text{bSB}}$ let us choose a special set of parameters for SimCIM defined with

$$\begin{aligned}
\gamma^{\text{SimCIM}} &= 1 \\
\alpha_0^{\text{SimCIM}} &= \alpha_0^{\text{bSB}} \cdot \Delta_t^{\text{bSB}} \\
\alpha_1^{\text{SimCIM}} &= \alpha_1^{\text{bSB}} \cdot \Delta_t^{\text{bSB}} \\
\beta^{\text{SimCIM}} &= \beta^{\text{bSB}} \cdot \Delta_t^{\text{bSB}} \\
\Delta_t^{\text{SimCIM}} &= m^{\text{bSB}} \Delta_t^{\text{bSB}}
\end{aligned} \tag{3.65}$$

Algorithm 9 bSB, revisited

- 1: Initialize vectors $\mathbf{x}^{(0)}, \mathbf{y}^{(0)}$ randomly around $\mathbf{0}$
 - 2: **for** k in $\text{range}(0, N_{\text{iter}})$ **do**
 - 3: $y_i^{(k+1)} \leftarrow y_i^{(k)} + \left(-\alpha^{\text{bSB}}(t_k)x_i^{(k)} + \beta^{\text{bSB}} \sum_{j=1}^n J_{ij}x_j^{(k)} \right) \Delta_t^{\text{bSB}}, \quad i = 1, \dots, n$
 - 4: $x_i^{(k+1)} \leftarrow x_i^{(k)} + m^{\text{bSB}} y_i^{(k+1)} \Delta_t^{\text{bSB}}, \quad i = 1, \dots, n$
 - 5: $y_i^{(k+1)} \leftarrow 0, \quad \text{if } x_i^{(k+1)} \notin [-1, +1], \quad i = 1, \dots, n$
 - 6: $x_i^{(k+1)} \leftarrow \phi(x_i^{(k+1)}), \quad i = 1, \dots, n$
 - 7: **end for**
-

Plugging in the above parameters into SimCIM with momentum (algorithm 7) we get

Algorithm 10 SimCIM with momentum mimicking bSB

- 1: Initialize vectors $\mathbf{x}^{(0)}, \mathbf{y}^{(0)}$ randomly around $\mathbf{0}$
 - 2: **for** k in $\text{range}(0, N_{\text{iter}})$ **do**
 - 3: $y_i^{(k+1)} \leftarrow y_i^{(k)} + \left(-\alpha^{\text{bSB}}(t_k) \Delta_t^{\text{bSB}} x_i^{(k)} + \beta^{\text{bSB}} \Delta_t^{\text{bSB}} \sum_{j=1}^n J_{ij} x_j^{(k)} \right), \quad i = 1, \dots, n$
 - 4: $x_i^{(k+1)} \leftarrow x_i^{(k)} + y_i^{(k+1)} m^{\text{bSB}} \Delta_t^{\text{bSB}}, \quad i = 1, \dots, n$
 - 5: $y_i^{(k+1)} \leftarrow 0, \quad \text{if } x_i^{(k+1)} \notin [-1, +1], \quad i = 1, \dots, n$
 - 6: $x_i^{(k+1)} \leftarrow \phi(x_i^{(k+1)}), \quad i = 1, \dots, n$
 - 7: **end for**
-

This shows that SimCIM with momentum is able to exactly reproduce bSB steps if parameters are chosen accordingly. In this context, SimCIM is a generalization of bSB algorithm.

3.12 First bifurcation point

As it was discussed earlier in this chapter, we want to choose parameters such that at the beginning of the dynamics we are already at the first bifurcation point or have even crossed it. Although the best parameter setting should be determined experimentally, by observing empirical evidence for which parameter provides the best solution, we expect that it will be something around the first bifurcation point.

Let us revise the first bifurcation points.

For CIM and SB this is equation 3.8 i.e.

$$\mu = -\frac{\beta}{\alpha}\lambda_{\max}(J)$$

For SimCIM and bSB this is equation 3.43 i.e.

$$\alpha = \beta\lambda_{\max}(J)$$

We see that both of these depend on the largest eigenvalue of the coupling matrix J . So, how do we determine the first bifurcation point exactly?

The first approach would be to use some standard library with numerical algorithms and use it to numerically approximate the largest eigenvalue of J . However, for real applications this might be undesirable (for example if the matrix is very large this will be time consuming).

The workaround, proposed in [29], is to *guess* the approximation of the largest eigenvalue. As stated in [29], according to Wigner's semicircle law in random matrix theory, λ_{\max} is approximately given by

$$\lambda_{\max} \approx 2\sqrt{n}\sigma \quad (3.66)$$

where n is the number of variables and σ is the standard deviation of the nondiagonal elements of J , which can be easily calculated from J .

3.13 Overview

In table 3.2 we summarize and systematically show all four dynamical systems discussed in this thesis – CIM, SB, SimCIM, and bSB.

Table 3.2: Overview of physics-inspired dynamical systems for combinatorial optimization

Dynamical system	Landscape & Vector field	Allowed region	Type	Equations of motion
CIM	$E(\mathbf{x}; \alpha, \beta, \mu)$ $= \frac{\alpha}{4} \sum_i (x_i^2 - \mu)^2 - \frac{\beta}{2} \sum_{i,j} J_{ij} x_i x_j$ $f_i(\mathbf{x}; \alpha, \beta, \mu) = \alpha x_i (\mu - x_i^2) + \beta \sum_j J_{ij} x_j$	$\mathbf{x} \in \mathbb{R}^n$	gradient	$\dot{x}_i = \alpha x_i (\mu - x_i^2) + \beta \sum_j J_{ij} x_j$
SB			Hamiltonian	$\begin{cases} \dot{x}_i = my_i \\ \dot{y}_i = \alpha x_i (\mu - x_i^2) + \beta \sum_j J_{ij} x_j \end{cases}$
SimCIM	$E(\mathbf{x}; \alpha, \beta) = \frac{\alpha}{2} \sum_i x_i^2 - \frac{\beta}{2} \sum_{i,j} J_{ij} x_i x_j$ $f_i(\mathbf{x}; \alpha, \beta) = -\alpha x_i + \beta \sum_j J_{ij} x_j$	$\mathbf{x} \in [-1, +1]^n$	gradient	$\begin{cases} \dot{x}_i = -\alpha x_i + \beta \sum_j J_{ij} x_j, & \text{when } x_i \in \langle -1, +1 \rangle \\ \dot{x}_i = 0, & \text{or } (x_i \in \{-1, +1\} \ \& \ x_i f_i < 0) \\ & \text{otherwise} \end{cases}$
bSB			Hamiltonian	$\begin{cases} \dot{x}_i = my_i \\ \dot{y}_i = -\alpha x_i + \beta \sum_j J_{ij} x_j, & \text{when } x_i \in \langle -1, +1 \rangle \\ y_i = 0, & \text{or } (x_i \in \{-1, +1\} \ \& \ x_i f_i < 0) \\ & \text{otherwise} \end{cases}$

Vector field \mathbf{f} is equal to the negative gradient of landscape function E , i.e. $\mathbf{f} = (f_1, \dots, f_n) = \frac{\partial E}{\partial \mathbf{x}}^T$. For gradient systems, vector field \mathbf{f} governs the dynamics. For Hamiltonian systems, the Hamiltonian is defined as $H = \frac{m}{2} \sum_i y_i^2 + E(\mathbf{x}; \alpha, \beta, \mu)$, where the first part of the summation corresponds to the kinetic energy, and the second part to the potential energy. The equations of motion for Hamiltonian systems are then $\dot{x}_i = \frac{\partial H}{\partial y_i} = my_i$, $\dot{y}_i = -\frac{\partial H}{\partial x_i} = f_i$. For SimCIM and bSB the dynamics is constrained inside the allowed region $[-1, +1]^n$ so the boundary of this region is modeled as a wall. When some component i hits the wall, we set the velocity for that component to 0 – meaning that for the gradient system (SimCIM) we set $\dot{x}_i = 0$, and for the Hamiltonian system (bSB) we set $y_i = 0$.

Chapter 4

Experiments

Physics-inspired dynamical systems presented in the previous chapter can be efficiently simulated on conventional hardware devices such as a CPU. Furthermore, since they almost completely comprise of a vast amount of simple add/multiply operations, they can be simulated even more efficiently on parallelized hardware devices such as GPUs or FPGAs. This enables them to easily scale to large instances of QUBO, MAX-CUT, and other problems of interest.

This chapter focuses on presenting the results of various experiments, explaining them, and discussing them further, all in order to investigate the performance of algorithms presented in chapter 3. These simulations include Coherent Ising Machine (CIM), Simulated Bifurcation (SB), Simulated Coherent Ising Machine (SimCIM), and Ballistic Simulated Bifurcation (bSB) algorithms. Besides these algorithms, some results obtained by classical algorithms are provided for comparison.

Furthermore, the performance of the original CIM algorithm is compared to the version of the algorithm where momentum is applied. Finally, a new technique called *dropout* is presented. It acts as adding noise in a *meaningful* way to the above algorithms, often providing improvement in solution quality.

4.1 GSet Dataset

GSet is a benchmark dataset publicly available on the link [48]. It consists of many weighted graph instances with different graph topologies. On some instances, the edges' weight is always 1, while on other instances, the edges' weight can be either +1 or negative -1.

Since the MAX-CUT problem is equivalent to QUBO problem 1.2.6, this dataset in fact serves as a QUBO benchmark. The best known cut value for each graph instance is provided according to the article [17], and will be referred to as 'max known cut' in the

following tables. These solutions are, in fact, also obtained by their heuristic algorithm called Breakout Local Search (BLS).

The experiments will be run on first 21 instances of GSet, G1-G21. Every instance G1-G21 has 800 nodes. The number of edges is denoted in the 'edge' column. Weights can be either always +1 or combined +1 and -1 which is denoted in the column 'weights'.

4.2 Method

This section explains details regarding experiment setup and execution.

Each experiment consists first of parameter fine-tuning and then testing. For each algorithm, and for each GSet instance this process is started from scratch.

In order to fine-tune the parameters, some predetermined set of candidates is chosen for each parameter. This set of candidates is chosen based on manual experimenting, and determining which range for each parameter provides sensible behaviour in terms of stability and solution quality. For example, in order to fine-tune the SimCIM or bSB algorithm, we need to choose starting coefficient $\alpha_0 \in \mathbb{R}$. If we choose $\beta = 1$, it makes sense to choose α_0 to be at the first bifurcation point $\beta\lambda_{\max}(J)$, or slightly less. So, for example, the candidates for α_0 could be chosen to be $\beta\lambda_{\max}(J) \cdot \eta$, $\eta \in \{1.0, 0.9, 0.8, 0.5, 0.1\}$.

Note that the momentum was used both for CIM and SimCIM algorithms, with the possibility for the momentum parameter γ to be fine-tuned, including set to 0 (which for CIM is the original algorithm without momentum). For example, for SimCIM the set of possible momentum parameters used was $[0, 0.8, 0.9, 0.99, 1.0]$. Both for SimCIM and CIM it turned out that the algorithm consistently provided better results with momentum strictly larger than 0. In section 4.4 we will provide an explicit comparison of the performance of CIM with and without momentum.

After choosing the possible candidates for each parameter, the fine-tuning process begins. For every parameter combination, selected in a grid-search fashion, the experiment is run 5 times using different initial conditions. The result from each run is taken and the mean value of these results is recorded. A parameter combination — which turned out to be stable (in a sense that it succeeded in all 5 tries), and with the highest mean value obtained — is chosen for testing, and will be referred to as *fine-tuned parameters*.

Next, a proper testing is performed. Using these fine-tuned parameters, the experiment is run 50 times with various initial conditions. The result from each run is taken and the minimum, mean, and maximum values are recorded, and presented in the following tables in section 4.3 under the columns 'min', 'mean', and 'max'.

In order to remove the possible advantage of certain parameter combinations just because of luck, i.e. having better initial conditions, each parameter combination receives the same initial conditions. In fact, 5 initial conditions are generated at the beginning, and their copy is provided to each parameter combination at the beginning of their run.

In order to remove the effect of approximating λ_{\max} by some proposed ways in chapter 3, for all of these experiments the value of λ_{\max} is calculated in advance using standard numerical algorithms for obtaining spectral decomposition of a symmetric matrix.

4.3 Benchmarking

The following tables represent results obtained by each algorithm on instances G1-G21 of GSet. For each algorithm, for each graph instance, first the parameters are fine-tuned as described in the previous section. Then, these fine-tuned parameters are selected and a new experiment is performed with 50 tries. In each try, different initial conditions (initial positions and initial momenta) are chosen. After performing these 50 tries, the minimum, mean and maximum cut values found by the algorithm are denoted in the corresponding columns 'min', 'mean', 'max'.

The following figures represent a single run of the experiment of each algorithm on GSet instance G18. This shows how the algorithm performs through time (or actually number of iterations). On x -axis, the number of iterations is plotted. On the *Cut value* subplot, the value of the cut is plotted obtained by taking $\text{sign}(\mathbf{x})$ as the partition of the graph. The red dotted line in this plot represents the maximum known cut from the table. For G18 it is equals to 992. On *Landscape value E* and *Potential energy E* subplots, the value of the function E (defined in chapter 3) is plotted at each iteration step $E(\mathbf{x}(t); \cdot(t))$. Saturation graph represents how many variables have absolute value larger than 0.98. It shows how many variables have moved from the origin. For SimCIM and bSB it actually represents how many variables have (probably) stopped their motion. The subplots denoted with α or μ represent the value of corresponding parameters changing through time. The subplot denoted with *Spins norm* is the norm of vector \mathbf{x} . The red dashed line represents the norm of feasible set $\{-1, +1\}^n$.

For SimCIM and CIM, the coupling matrices are normalized (not a necessary step) in a sense that $J/\|J\|$ is taken in place of J . That's why the α and E might be scaled differently than expected. The purpose of these plots is to see the overall behavior — the trend of each function.

All of these plots could be used to analyze the performance of each algorithm and potentially improve algorithms' bottlenecks.

instance	edges	weights	max known cut	min	mean	max
G1	19176	1	11624	11525	11561.2	11594
G2	19176	1	11620	11531	11565.9	11590
G3	19176	1	11622	11525	11566.8	11606
G4	19176	1	11646	11567	11608.3	11627
G5	19176	1	11631	11566	11594	11602
G6	19176	+1,-1	2178	2093	2142.7	2147
G7	19176	+1,-1	2006	1921	1955	1976
G8	19176	+1,-1	2005	1916	1950.3	1973
G9	19176	+1,-1	2054	1939	1990.9	2032
G10	19176	+1,-1	2000	1918	1947.4	1982
G11	1600	+1,-1	564	544	550.3	554
G12	1600	+1,-1	556	534	543.1	550
G13	1600	+1,-1	582	558	567.5	574
G14	4694	1	3064	2966	2986.2	3010
G15	4661	1	3050	2955	2975.4	2985
G16	4672	1	3052	2959	2974.2	2984
G17	4667	1	3047	2958	2975.4	2988
G18	4694	+1,-1	992	898	924.7	945
G19	4661	+1,-1	906	815	853.1	854
G20	4672	+1,-1	941	852	889.5	891
G21	4667	+1,-1	931	814	850.6	887

Table 4.1: CIM results on GSet

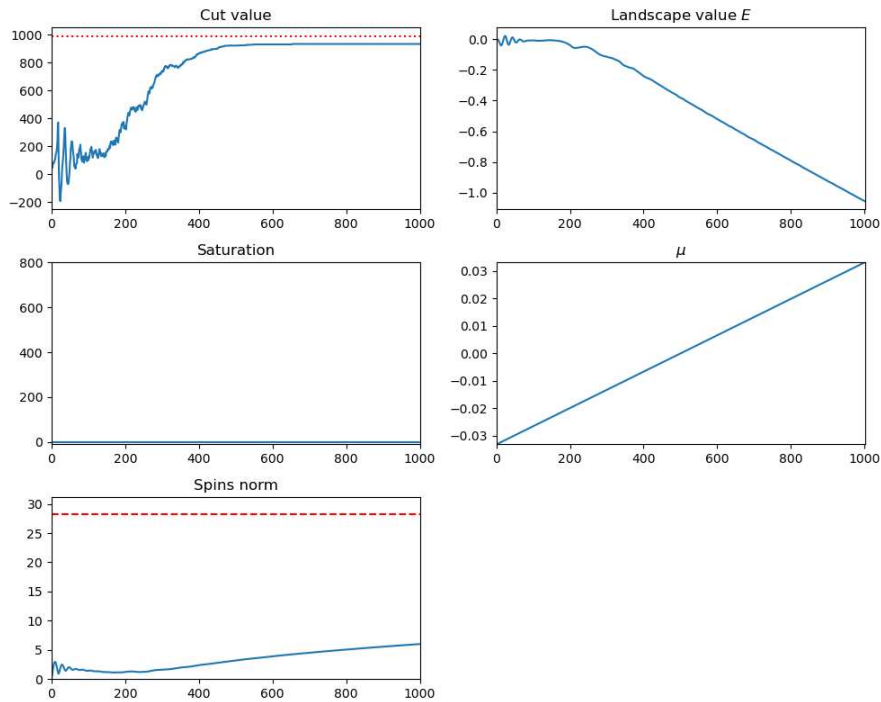


Figure 4.1: Single run of CIM algorithm on GSet instance G18

instance	edges	weights	max known cut	min	mean	max
G1	19176	1	11624	11550	11582.2	11615
G2	19176	1	11620	11569	11580	11599
G3	19176	1	11622	11563	11589.8	11617
G4	19176	1	11646	11587	11614.5	11632
G5	19176	1	11631	11595	11607.2	11618
G6	19176	+1,-1	2178	2149	2158.7	2168
G7	19176	+1,-1	2006	1960	1978.1	1998
G8	19176	+1,-1	2005	1936	1977.8	1990
G9	19176	+1,-1	2054	2000	2020.7	2038
G10	19176	+1,-1	2000	1944	1968.6	1986
G11	1600	+1,-1	564	548	553.6	558
G12	1600	+1,-1	556	542	549.8	554
G13	1600	+1,-1	582	558	571.2	578
G14	4694	1	3064	3004	3016.5	3033
G15	4661	1	3050	2980	2996	3011
G16	4672	1	3052	2980	3000.7	3018
G17	4667	1	3047	2981	2993.7	3007
G18	4694	+1,-1	992	914	946.4	972
G19	4661	+1,-1	906	473	818	871
G20	4672	+1,-1	941	874	899.5	917
G21	4667	+1,-1	931	890	893.1	899

Table 4.2: SB results on GSet

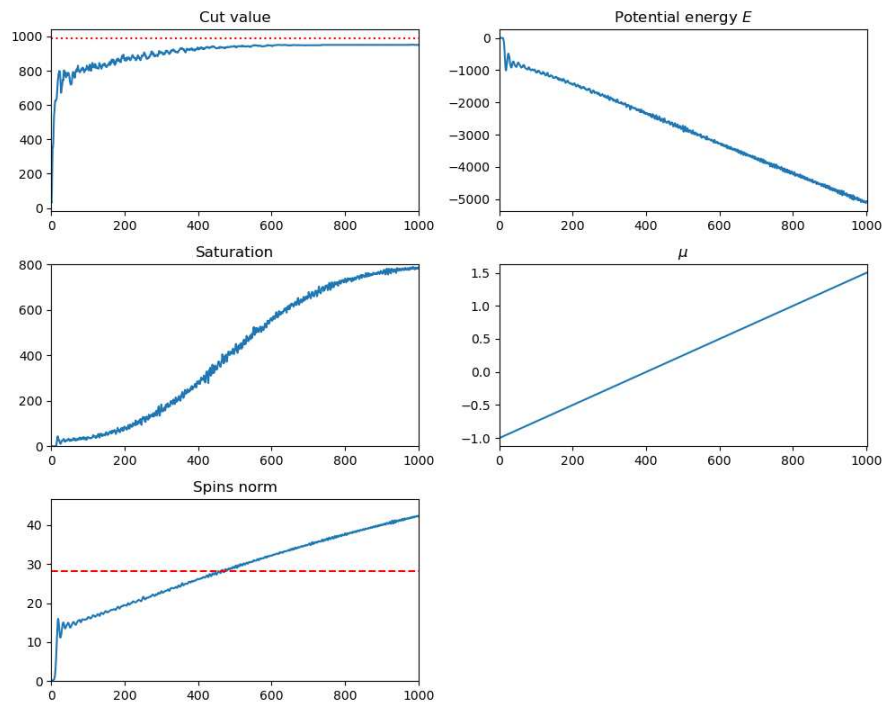


Figure 4.2: Single run of SB algorithm on GSet instance G18

instance	edges	weights	max known cut	min	mean	max
G1	19176	1	11624	11566	11608.8	11623
G2	19176	1	11620	11580	11595	11609
G3	19176	1	11622	11568	11618.7	11622
G4	19176	1	11646	11617	11636.8	11642
G5	19176	1	11631	11599	11624.2	11625
G6	19176	+1,-1	2178	2139	2172.6	2174
G7	19176	+1,-1	2006	1975	1986.9	2002
G8	19176	+1,-1	2005	1969	1986.6	2005
G9	19176	+1,-1	2054	2012	2035.9	2046
G10	19176	+1,-1	2000	1952	1975.6	1992
G11	1600	+1,-1	564	544	549.6	558
G12	1600	+1,-1	556	536	544.5	550
G13	1600	+1,-1	582	558	569.2	576
G14	4694	1	3064	3031	3033.6	3034
G15	4661	1	3050	3022	3022	3022
G16	4672	1	3052	3026	3027.3	3029
G17	4667	1	3047	3021	3021	3021
G18	4694	+1,-1	992	964	964.6	965
G19	4661	+1,-1	906	869	878.3	880
G20	4672	+1,-1	941	926	926.1	931
G21	4667	+1,-1	931	907	913.6	917

Table 4.3: SimCIM results on GSet

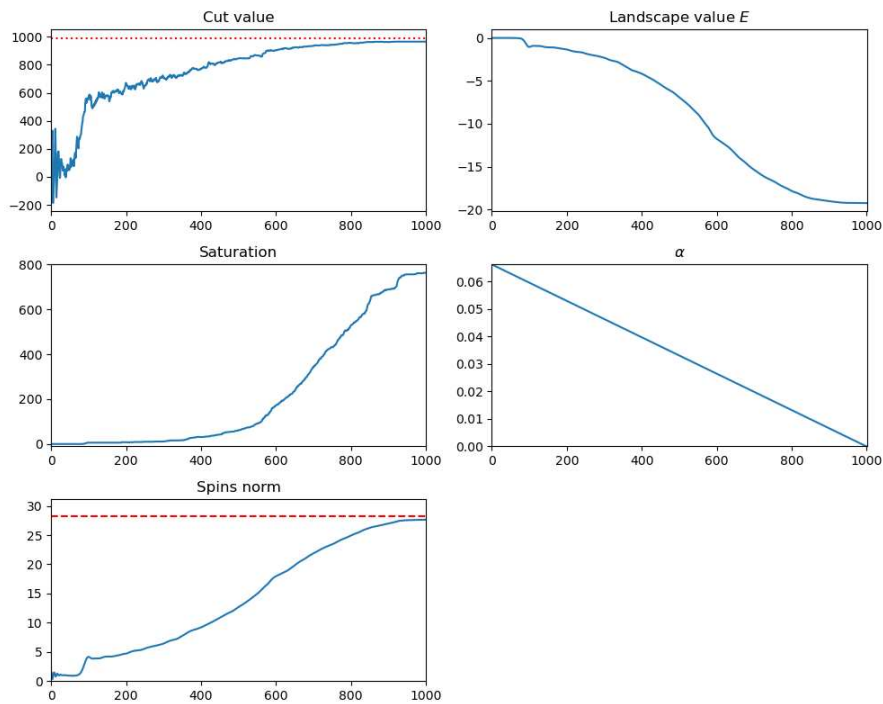


Figure 4.3: Single run of SimCIM algorithm on GSet instance G18

instance	edges	weights	max known cut	min	mean	max
G1	19176	1	11624	11582	11608.5	11623
G2	19176	1	11620	11569	11595.6	11612
G3	19176	1	11622	11573	11619.1	11622
G4	19176	1	11646	11623	11637.6	11638
G5	19176	1	11631	11625	11625	11625
G6	19176	+1,-1	2178	2143	2171.3	2174
G7	19176	+1,-1	2006	1958	1985.2	1998
G8	19176	+1,-1	2005	1959	1985.7	2004
G9	19176	+1,-1	2054	2012	2031.9	2046
G10	19176	+1,-1	2000	1969	1982.9	1991
G11	1600	+1,-1	564	546	552.7	556
G12	1600	+1,-1	556	538	545.5	550
G13	1600	+1,-1	582	562	569.1	578
G14	4694	1	3064	3020	3028.6	3038
G15	4661	1	3050	3011	3021.3	3036
G16	4672	1	3052	3016	3025.3	3038
G17	4667	1	3047	3003	3017	3033
G18	4694	+1,-1	992	953	963.7	976
G19	4661	+1,-1	906	855	874.5	879
G20	4672	+1,-1	941	881	912.5	930
G21	4667	+1,-1	931	875	906.2	917

Table 4.4: bSB results on GSet

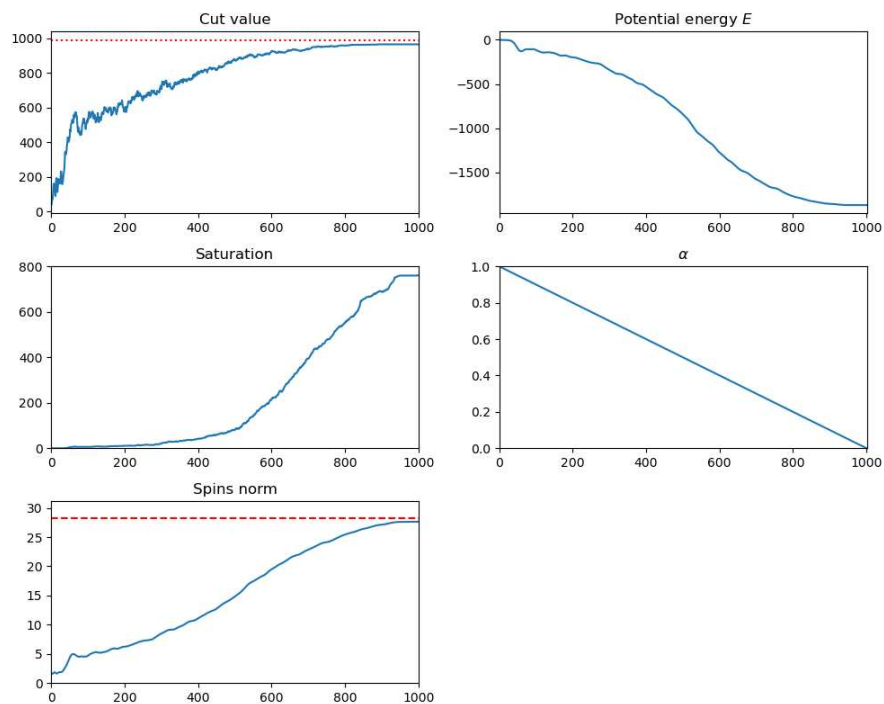


Figure 4.4: Single run of bSB algorithm on GSet instance G18

instance	edges	weights	max known cut	CIM max	SB max	SimCIM max	bSB max	Eigen cut	Random cuts (1k)
G1	19176	1	11624	11594	11615	11623	11623	11265	9785
G2	19176	1	11620	11590	11599	11609	11612	11248	9819
G3	19176	1	11622	11606	11617	11622	11622	11264	9794
G4	19176	1	11646	11627	11632	11642	11638	11290	9806
G5	19176	1	11631	11602	11618	11625	11625	11326	9771
G6	19176	+1,-1	2178	2147	2168	2174	2174	1905	329
G7	19176	+1,-1	2006	1976	1998	2002	1998	1659	151
G8	19176	+1,-1	2005	1973	1990	2005	2004	1640	135
G9	19176	+1,-1	2054	2032	2038	2046	2046	1776	176
G10	19176	+1,-1	2000	1982	1986	1992	1991	1596	178
G11	1600	+1,-1	564	554	558	558	556	466	94
G12	1600	+1,-1	556	550	554	550	550	470	62
G13	1600	+1,-1	582	574	578	576	578	508	86
G14	4694	1	3064	3010	3033	3034	3038	2635	2462
G15	4661	1	3050	2985	3011	3022	3036	2550	2443
G16	4672	1	3052	2984	3018	3029	3038	2644	2442
G17	4667	1	3047	2988	3007	3021	3033	2493	2435
G18	4694	+1,-1	992	945	972	965	976	590	154
G19	4661	+1,-1	906	854	871	880	879	514	55
G20	4672	+1,-1	941	891	917	931	930	600	84
G21	4667	+1,-1	931	887	899	917	917	564	77

Table 4.5: Performance comparison of various algorithms on GSet

Table 4.5 compares performance of physics-inspired algorithms presented in this thesis. For all physics-inspired algorithms (CIM, SB, SimCIM, bSB), the maximum value is shown, which is found among 50 tries with various initial conditions with fine-tuned parameters as described earlier. Column 'max known cut', as earlier, represents the best known solution on particular instance, according to [17], which is also the solution found by BLS heuristic. The column 'Eigen cut' represents the cut value of the partition obtained by taking the sign of each component of the eigenvector corresponding to smallest eigenvalue of the adjacency matrix (this is an obvious heuristic based on 1.1.2). The column 'Random cuts (1k)' represents the maximum cut found among 1000 random cuts. These last two columns are here as a reference for the lower bound which we definitely want to beat.

4.4 Momentum

In table 4.6 we compare the performance of the original CIM (algorithm 1) and its version with momentum (algorithm 2). The method is the same as described in section 4.2, except that either we choose the momentum to be precisely 0 (denoted by ' $m = 0$ ' in the table) or we choose it from the set $\{0.5, 0.7, 0.8, 0.9, 1.0\}$ (denoted by ' $m > 0$ ' in the table). For each of these momentum choices, a parameter fine-tuning procedure is performed as described in 4.2. After the parameter fine-tuning, a full experiment was run, consisting of 50 tries with different initial conditions, and obtained cut values were recorded. Mean and maximum of these recorded values is taken, for each momentum parameter. The maximum among these 5 mean values is in the column ' $m > 0$ mean'. The maximum among these 5 max values is in the column ' $m > 0$ max'. Columns ' $m = 0$ mean' and ' $m = 0$ max' represent the corresponding values when momentum parameter is set to 0. "Diff" column represents the difference $(m > 0) - (m = 0)$, so its positive value implies that adding momentum increases the performance.

In order to remove the possible advantage of momentum results due to larger number of total tries (because 5 different momentum parameters were tried), initial conditions are taken exactly the same as for the experiment without momentum. Thus, 50 random initial conditions were sampled at the beginning, and their copy was provided to each experiment. The same was done during fine-tuning, but with 5 random initial conditions.

It can be concluded that introducing momentum to CIM algorithm significantly improves its performance.

instance	edges	weights	max known cut	$m = 0$ mean	$m > 0$ mean	mean diff	$m = 0$ max	$m > 0$ max	max diff
G11	1600	+1,-1	564	531.8	550.3	18.5	542	554	12
G12	1600	+1,-1	556	526	543.1	17	536	550	14
G13	1600	+1,-1	582	551	567.5	16.5	562	574	12
G14	4694	1	3064	2940.1	2986.2	46.1	2962	3010	48
G15	4661	1	3050	2932	2975.4	43.5	2948	2985	37
G18	4694	+1,-1	992	877.9	924.7	46.8	906	945	39
G19	4661	+1,-1	906	811.8	853.1	41.3	838	854	16
G20	4672	+1,-1	941	849.5	889.5	40.1	873	891	18

Table 4.6: Momentum results for CIM

4.5 Dropout

This section describes a potential way of improving the performance of physics-inspired algorithms, presented earlier in this chapter, by adding a special type of noise. This procedure is discussed here in terms of graphs and MAX-CUT problem instead of variables and QUBO (which is equivalent 1.2.6). There are certainly many ways to add noise, including adding just white noise to the dynamical system, making it a stochastic dynamical system. One possible way of adding noise in a sensible way is to perturb the graph somehow in each step of the algorithm, for example by removing some vertices or edges from the graph temporarily. This way, intuitively, the dynamical system should on average, at each step, follow the same path, while opening the possibility to escape local minima or even make some components bifurcate earlier by removing some of the opposite forces acting on a single variable (often called *frustration*). Indeed, random removal of vertices often seems to improve the performance on GSet instances. This is done in the following way. At each iteration of the algorithm (CIM, SB, SimCIM, bSB), instead of working with the full adjacency matrix, one randomly and independently picks with probability p for each vertex whether it will stay or it will be temporarily removed. If vertex i is removed, then all edges connected to it are temporarily removed, until completing this iteration. Everything else remains the same. There is a visual example of adding dropout in figure 4.5.

Let us observe what can happen to vertex i . Each of the algorithms has the term $\sum_j J_{ij}x_j$ inside the update. If vertex i is not removed, then the summation $\sum_j J_{ij}x_j$ will not go through all vertices, but rather through all vertices which are not removed. If vertex i is removed itself, then the summation $\sum_j J_{ij}x_j$ will disappear. The rest of the update step for this vertex can either be performed, or not, depending on the implementation.

This procedure is similar to adding dropout to machine learning models during training [49]. This procedure will thus also be referred to as adding *dropout*.

For small dropout probabilities ($p \leq 0.1$), the algorithm often obtains an increase in performance.

In table 4.7, we see how adding dropout affects the performance of particular algorithm on GSet instances. The experiments were performed in a similar fashion as described in section 4.2. Dropout probabilities were selected from $p \in \{1\%, 5\%, 10\%, 20\%\}$. For each of these dropout probability, a parameter fine-tuning procedure is performed as described in 4.2. After parameter fine-tuning, a full experiment was run consisting of 50 tries with different initial conditions, and the obtained cut values are recorder. Mean and maximum of these recorded values is taken, for each dropout probability. The maximum among these 4 mean values is represented in the table under the column ' $p > 0$ mean'. The maximum among these 4 max values is represented in the table under the column ' $p > 0$ max'. Columns 'mean diff' and 'max diff' represent the difference between 'mean' and 'max' columns in this table and the corresponding columns in tables 4.1, 4.2, 4.3, and 4.4, where

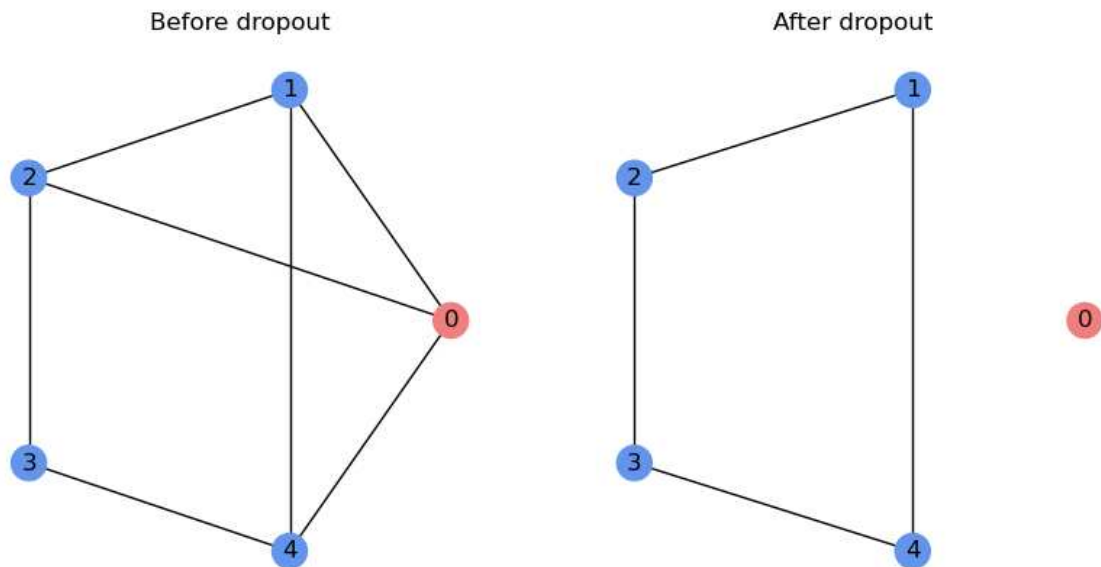


Figure 4.5: Dropout procedure

On the left we see a graph without applied dropout. Only the vertex 0 is chosen to be dropped out. The effect of dropout is that all edges connected to this vertex are removed, and it can be seen on the right.

there is no dropout — denoted as $p = 0$. Thus, the 'diff' columns represent the difference $(p > 0) - (p = 0)$ for mean and max columns. So, if this difference is positive, it means that applying dropout increased the performance. Otherwise, adding dropout did not impact the performance, or worsened it.

In order to remove the possible advantage of dropout results due to larger number of total tries (because 4 different dropout probabilities were tried), initial conditions are taken exactly the same as for the experiment with no dropout. Thus, 50 random initial conditions were sampled already for experiment with no dropout, and exactly these initial conditions were used as 50 tries of each experiment with added dropout. The same was done during fine-tuning, but with 5 random initial conditions.

General info		CIM				SB				SimCIM				bSB			
instance	max known cut	$p > 0$ mean	mean diff	$p > 0$ max	max diff	$p > 0$ mean	mean diff	$p > 0$ max	max diff	$p > 0$ mean	mean diff	$p > 0$ max	max diff	$p > 0$ mean	mean diff	$p > 0$ max	max diff
G1	11624	11571.8	10.6	11615	21	11546.1	-36.1	11586	-29	11605.9	-2.9	11624	1	11608.3	-0.2	11624	1
G2	11620	11567.1	1.2	11596	6	11552.2	-27.8	11577	-22	11600.5	5.5	11616	7	11601.1	5.5	11615	3
G3	11622	11563.8	-3	11594	-12	11551	-38.7	11575	-42	11614.4	-4.3	11622	0	11612.6	-6.5	11622	0
G4	11646	11611.2	2.8	11628	1	11593.7	-20.8	11618	-14	11634.6	-2.2	11646	4	11637.2	-0.4	11646	8
G5	11631	11592.7	-1.3	11610	8	11582.9	-24.3	11608	-10	11622.1	-2.1	11627	2	11622.4	-2.6	11631	6
G6	2178	2141.5	-1.2	2155	8	2125	-33.7	2157	-11	2169.6	-3.1	2175	1	2172.5	1.2	2177	3
G7	2006	1957.6	2.6	1977	1	1947.5	-30.6	1969	-29	1987.8	0.9	2005	3	1990.7	5.5	2006	8
G8	2005	1949.1	-1.3	1976	3	1946.8	-31	1973	-17	1989.5	2.8	2005	0	1992	6.2	2005	1
G9	2054	2006.6	15.7	2026	-6	1997	-23.7	2024	-14	2038.2	2.2	2048	2	2039.2	7.3	2051	5
G10	2000	1954	6.6	1980	-2	1948.2	-20.4	1975	-11	1979.4	3.8	1995	3	1983.1	0.1	1996	5
G11	564	549.6	-0.6	558	4	549.6	-4	558	0	551.4	1.8	558	0	551.6	-1.1	556	0
G12	556	543.4	0.3	552	2	542.8	-7	554	0	545.5	1	552	2	545.8	0.3	552	2
G13	582	566.7	-0.8	576	2	567.5	-3.7	576	-2	570.1	0.9	576	0	569.2	0.1	576	-2
G14	3064	2990.3	4.1	3008	-2	2995.9	-20.6	3016	-17	3033.8	0.2	3048	14	3034.9	6.3	3049	11
G15	3050	2979.7	4.3	2992	7	2984.4	-11.6	2997	-14	3021.9	-0.1	3034	12	3021.1	-0.2	3037	1
G16	3052	2978.5	4.3	2999	15	2987.7	-13	2999	-19	3027.9	0.6	3042	13	3022.7	-2.5	3040	2
G17	3047	2978.4	3.1	2999	11	2985.3	-8.5	3003	-4	3019.7	-1.3	3038	17	3018.7	1.7	3035	2
G18	992	935.7	11.1	946	1	937.9	-8.5	953	-19	965.1	0.5	982	17	967.7	4	986	10
G19	906	854.2	1.1	860	6	849.9	31.8	864	-7	876.8	-1.6	892	12	877.8	3.3	899	20
G20	941	889.2	-0.4	902	11	889.5	-10	908	-9	927.7	1.6	932	1	919.3	6.8	938	8
G21	931	878.1	27.5	896	9	883.9	-9.1	904	5	909.7	-3.9	917	0	909.8	3.6	917	0

Table 4.7: Dropout results

The results show that for CIM, SimCIM, and bSB algorithms, adding some amount of dropout often increases the performance by some small extent. It rarely decreases the performance, and most importantly, sometimes increases the performance significantly.

To further investigate whether adding dropout makes any significant difference, the following was done. GSet instance G18 was picked, and bSB algorithm was run 10.000 times with various parameters, picked randomly from some sensible search space, and with various random initial conditions, but with no dropout. The best cut value found was 982. On the other hand, doing the same experiment with 1.000 tries, and while adding dropout (this time with decreasing schedule of dropout probability), a cut with value 989 was found, which is very close to the maximum known cut 992. Thus, it can be concluded that likely it is the dropout which improved the performance of bSB algorithm on this instance.

Besides these improvements, it can be seen that adding dropout to SB algorithm makes the opposite effect — it reduces the performance significantly. It is not clear why only the SB algorithm suffers a performance loss, while CIM, SimCIM, and bSB obtain an improvement. It was observed, however, that fine-tuned parameters for CIM often had lower rates of momentum when $p > 0$ compared to the case when $p = 0$. This is consistent with the results, since SB can be seen as having fixed momentum equal to 1 (see section 3.6).

While adding dropout by removing vertices increases the performance, it has been observed that adding dropout in such a way as to remove random edges (each edge sampled independently), does not seem to improve the performance.

A step towards analyzing the dropout

In order to analyze the mechanism of dropout, it would be nice to have some small instance of a problem where dropout makes a significant improvement. Such an instance is provided in this section.

Consider a graph G with the following adjacency matrix

$$\begin{bmatrix} 0 & 10 & 10 & -1 & -10 & 1 & -10 \\ 10 & 0 & -10 & -1 & -1 & -1 & -100 \\ 10 & -10 & 0 & -1 & 1 & 10 & 0 \\ -1 & -1 & -1 & 0 & 0 & -1 & 10 \\ -10 & -1 & 1 & 0 & 0 & -100 & 0 \\ 1 & -1 & 10 & -1 & -100 & 0 & -1 \\ -10 & -100 & 0 & 10 & 0 & -1 & 0 \end{bmatrix}$$

MAX-CUT value for this graph is 26, and it is obtained for the cut (1, 0, 0, 1, 1, 1, 0). Running 100 tries of SimCIM algorithm on this graph with various sensible parameters and initial conditions (but with momentum 0 and taking the last state of the system instead

of the best – in order to remove unnecessary variance), we obtain the following distribution of cut values.

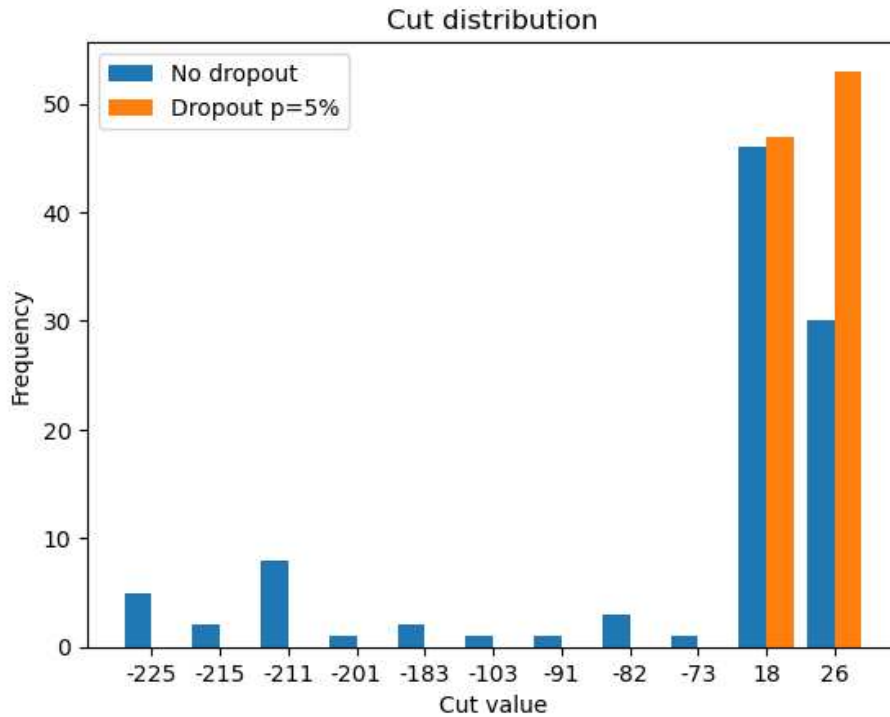


Figure 4.6: Cut distribution obtained by SimCIM on the above example

Adding dropout to this example significantly shifts the distribution towards higher values of cuts. In fact, this serves as one more evidence that adding dropout might increase the performance of the algorithm.

The question about *how* and *why* does adding dropout actually contribute to the performance gain — is still unanswered. This question is out of the scope of this thesis, and would require further testing and analysis.

Conclusion

Many combinatorial optimization problems are NP-hard, meaning that there is no (known) efficient (polynomial time complexity) algorithm for solving them exactly. At the beginning of this master's thesis, in chapter 1, a few of these problems are presented. The most attention throughout the thesis is devoted to two of them (which are in fact equivalent):

- Quadratic unconstrained binary optimization (QUBO) problem - A problem of minimizing (or maximizing) a multivariable quadratic functional over the discrete domain $\{-1, +1\}^n$
- MAX-CUT problem - Given a graph, divide its vertices into two parts such that the total edge weight between these parts (cut value) is maximized.

Since there is no known efficient algorithm for solving NP-hard problems exactly, various heuristic approaches are being developed to solve these problems approximately.

A relatively new paradigm for such heuristic algorithms are physics-inspired algorithms, including Coherent Ising Machine (CIM), Simulated Bifurcation (SB), and their variants (bSB and SimCIM). These algorithms are the central topic of this master's thesis. They are essentially numerical simulations of specific dynamical systems which encode candidate solutions of the QUBO problem. If the dynamical system has the trajectory $\mathbf{x} : [0, T] \rightarrow \mathbb{R}^n$, then this encoding is achieved by taking the sign of each variable at some time instance, i.e. taking $(\text{sign } x_1, \dots, \text{sign } x_n)$ as a candidate combinatorial solution. These dynamical systems are designed in a way that they attract the system towards those areas of space which correspond to high-quality solutions of QUBO problem. Although there is no theoretical guarantee for the quality of obtained solutions, it has been empirically found that these solutions indeed tend to be of high quality.

That these algorithms are physics-inspired means that they have been derived by observing certain physical systems capable of solving the QUBO problem (these physical systems are called *Ising machines* in the literature [18]), and then writing a set of coupled differential equations which approximately describe the physical system. The above systems essentially turn out to be gradient (CIM and SimCIM) and Hamiltonian (SB and bSB) dynamical systems over some time-dependent energy landscape function E . Study-

ing their behaviour thus boils down to analyzing the energy landscape E . Throughout the thesis, these dynamical systems are analyzed from a purely mathematical perspective. This way, one is able to clearly understand their mechanisms for generating high-quality solutions, rather than just assume their similarity to the original physical systems.

Numerically simulating these dynamical systems on a computer provides us with heuristic algorithms for solving the QUBO problem. A major advantage of these algorithms is that they can be parallelized efficiently. Since computationally the most expensive part of these algorithms are matrix-vector multiplications, these can be effectively handled by a GPU. Implementing these algorithms is relatively straightforward using some framework and library containing numerical algorithms (for example PyTorch). Then, it does not take much more work to run them on a GPU or other parallelized architecture.

In the final part of the thesis, physics-inspired algorithms are benchmarked on the GSet dataset. Along with that, some new techniques are proposed for improving the performance of these existing algorithms.

The first technique is proposing momentum to CIM algorithm. Similarly as it was proposed for SimCIM and other gradient-based systems, adding momentum to CIM algorithm improves its performance. Adding momentum to gradient systems (CIM and SimCIM) makes them some sort of generalization of the corresponding Hamiltonian systems (SB and bSB). In fact, by setting the momentum parameter equal to 1, CIM and SimCIM are able to reproduce the behaviour of SB and bSB algorithms.

The second proposed technique is dropout. This technique effectively adds some meaningful noise to the dynamical systems. Although the exact mechanism of dropout needs to be explained yet, there is empirical evidence found both on GSet dataset, and some other examples, which show that adding dropout to CIM, SimCIM, and bSB might significantly improve their performance (does not hold for SB). This technique is named according to a similar procedure used when training machine learning models.

Some interesting questions and open problems for further research include the following.

- What other dynamical systems are capable of approximately solving QUBO and MAX-CUT? For example, there are variants of SB algorithm (not studied in this thesis) called discrete simulated bifurcation (dSB) [31], and heated simulated bifurcation [45]. Except new variants of these algorithms, there might exist completely new paradigms. These dynamical system are physics-inspired and turn out to use mechanisms described in sections 3.5 and 3.10. Maybe some other mechanisms for providing high-quality combinatorial solutions could be used as a starting point, then transformed into a dynamical system which exploits such mechanisms.
- Are there any dynamical systems which would be capable of directly solving some other combinatorial optimization problems, without the need of embedding them

into QUBO and imposing constraints on it? For example, a variant of SB algorithm was proposed in [50] for binary optimization of polynomials with order higher than quadratic.

- Is there some efficient way of handling constraints, without losing the performance?
- How and why does adding dropout to CIM, SimCIM, and bSB improve their performance?
- Is there some other way to add noise which would improve the performance even more? For example, using *randomly modulated driving signals* was proposed for CIM already in [28]. Adding noise might improve the performance but there are various ways for adding it and it requires further tuning.
- How to choose parameters efficiently for various instances of problems? How to schedule the parameters (μ , α in sections 3.1 and 3.7) responsible for changing the landscape E in time? This problem commonly also arises when training neural networks. Choosing the right set of hyperparameters is often an issue and it is not clear how to do it in a good systematic way.
- Are there any theoretical guarantees for the solution quality obtained by these algorithms?

Hopefully some of these and related questions will be answered soon, providing us with better techniques for solving NP-hard combinatorial optimization problems.

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Summary

This thesis presents, analyzes, and finally proposes some improvements to physics-inspired dynamical systems – a novel paradigm of heuristic algorithms used for approximately solving NP-hard combinatorial optimization problems.

The thesis begins with a brief introduction to NP-hard combinatorial problems, with a particular emphasis on the quadratic unconstrained binary optimization problem (QUBO) and the MAX-CUT problem. Then, a brief overview is provided for the theory of dynamical systems and other mathematical concepts which are important for analyzing the following systems. At the core of this master's thesis are physics-inspired dynamical systems for solving combinatorial optimization problems. Numerical simulations of these dynamical systems provide heuristic algorithms for approximately solving some NP-hard problems, including the QUBO and the MAX-CUT problem. The algorithms that are covered in this thesis are namely the Coherent Ising Machine (CIM), Simulated Bifurcation (SB), Simulated Coherent Ising Machine (SimCIM), and Ballistic Simulated Bifurcation (bSB). The mechanism of these heuristic algorithms is analyzed, and they are further benchmarked on the GSet dataset. Finally, a few new techniques are proposed for improving the performance of these algorithms.

Sažetak

Ovaj diplomski rad izlaže, analizira te naposljetku predlaže unaprjeđenja dinamičkih sustava inspiriranih fizikom – novu vrstu heurističkih algoritama koji aproksimativno rješavaju NP-teške kombinatorne probleme.

Diplomski rad započinje sažetim uvodom u NP-teške kombinatorne probleme, s posebnim naglaskom na problem neograničene binarne optimizacije kvadratnog funkcionala (eng. *Quadratic Unconstrained Binary Optimization* – QUBO) te MAX-CUT problema. Zatim se izlažu temeljni rezultati teorije dinamičkih sustava i ostalih matematičkih koncepata koji će se koristiti u ostatku rada kako bi se analizirali sustavi koji slijede.

Glavni dio ovog rada su dinamički sustavi inspirirani fizikom koji rješavaju kombinatorne probleme. Numeričke simulacije ovih dinamičkih sustava predstavljaju heurističke algoritme za aproksimativno rješavanje NP-teških problema, uključujući QUBO i MAX-CUT. Algoritmi koji su obrađeni u ovom radu su *Coherent Ising Machine* (CIM), *Simulated Bifurcation* (SB), *Simulated Coherent Ising Machine* (SimCIM) i *Ballistic Simulated Bifurcation* (bSB). Analiziran je mehanizam ovih algoritama te je provedeno testiranje njihovih performansi na skupu podataka *GSet*. Na kraju rada predloženo je nekoliko tehnika koje unaprjeđuju performanse navedenih algoritama.

Životopis

Rođen sam 10. srpnja 1999. u Zagrebu. Školovanje sam započeo u Osnovnoj školi Josipa Račića, nakon čega sam 2014. godine upisao XV. gimnaziju u Zagrebu. Tijekom srednje i osnovne škole sudjelovao sam na brojnim informatičkim natjecanjima. Nakon završenog srednjoškolskog obrazovanja 2018. godine upisao sam preddiplomski studij Matematike na Prirodoslovno-matematičkom fakultetu u Zagrebu. Nakon stjecanja akademskog naziva *sveučilišni prvostupnik* 2021. godine upisujem diplomski studij Teorijska matematika na istom fakultetu.