

# Graph Coloring

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## 1 Introduction

Graph coloring refers to the problem of coloring vertices of a graph in such a way that no two adjacent vertices have the same color. This is also called the vertex coloring problem. If coloring is done using at most  $m$  colors, it is called  $m$ -coloring. The first results about graph coloring deal almost exclusively with planar graphs in the form of map coloring. While trying to color a map of the counties of England, Francis Guthrie postulated the four color conjecture, noting that four colors were sufficient to color the map so that no regions sharing a common border received the same color. Guthrie's brother passed on the question to his mathematics teacher Augustus De Morgan at University College, who mentioned it in a letter to William Hamilton in 1852. Arthur Cayley raised the problem at a meeting of the London Mathematical Society in 1879.

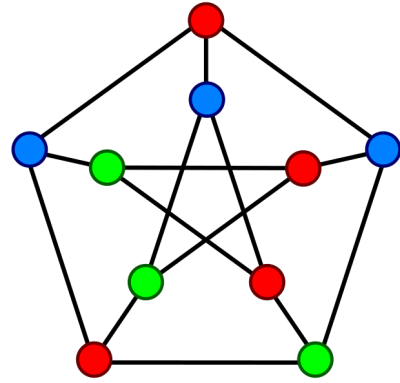


Figure 1: Petersen graph 3-coloring

More formally, we have a Graph Coloring Problem (GCP) Instance  $I = (\mathcal{G}, C)$  composed of a graph  $\mathcal{G} = (\mathcal{V}, \mathcal{E})$  and  $C \in \mathbb{N}, C > 2$ . Find a function  $f : \mathcal{V} \rightarrow \mathcal{C}$  such that  $\forall u, v \in \mathcal{V}, u \neq v, f(u) \neq f(v)$ .

## 2 Benchmark Instances

The selection of benchmark instances is an important factor in the empirical analysis of an algorithm's behaviour, and the use of inadequate benchmark sets can lead to questionable results and misleading conclusions. The criteria for benchmark selection depend significantly on the problem domain under consideration, on the hypotheses and goals of the empirical study, and on the algorithms being analysed.

Here is a list of Graph-Vertex Coloring instances used in the literature. For each instance the table gives:

- $|V|$  number of vertices

- $|E|$  number of edges
- $d\%$  graph density, corresponding to the number of edges divided by  $(|V| \text{ choose } 2)$
- $w\_LB(G)$  lower bound to the size of the maximum clique
- $w\_c(G)$  the size of the maximum clique
- $\theta(G)$  the theta number computed via semi definite programming
- $X\_f(G)$  the fractional chromatic number
- $X\_LB(G)$  lower bound to the chromatic number if computed in a way different from the previous
- $X(G)$  chromatic number
- $X\_UB(G)$  upper bound to the chromatic number, aka heuristic solution

The instances are classified according to their difficulty. We follow the SteinLib classification:

All instances for which no polynomial time algorithm is known are classified as NP.

The letter after this identifier indicates how long it takes to solve the problem using state-of-the-art soft- and hardware.

- s - seconds means less than a minute (this includes instances which can be solved in fractions of a second)
- m - minutes means less than an hour
- h - hours is less than a day
- d - days is less than a week
- w - weeks means it takes really a long time to solve this instance
- ? means the instance is not solved or the time is not known. If the chromatic number is given for this instances, then it is known by construction.

See Figure 2, Figure 3, Figure 4 and Figure 5.

1		V	E	d%	w <sub>LB</sub> (G)	w <sub>(G)</sub>	theta(G)	X <sub>f</sub> (G)	X <sub>LB</sub> (G)	X(G)	X <sub>UB</sub> (G)
2	1-FullIns_3	30	100	0,23	3	3				4	
3	1-Insertions_4	67	232	0,10	2	2				5	
4	2-FullIns_3	52	201	0,15	4	4				5	
5	2-Insertions_3	37	72	0,11	2	2				4	
6	3-Insertions_3	56	110	0,07	2	2				4	
7	anna	138	493	0,05	11	11				11	
8	ash331GPIA	662	4181	0,02	3	3				4	
9	david	87	406	0,11	11	11				11	
10	DSJC125.1	125	736	0,09	4	4				5	
11	DSJR500.1	500	3555	0,03	11	11				12	
12	fpsol2.i.1	496	11654	0,09	65	65				65	
13	fpsol2.i.2	451	8691	0,09	30	30				30	
14	fpsol2.i.3	425	8688	0,10	30	30				30	
15	games120	120	638	0,09	9	9				9	

Figure 2: NP-s instances

1		V	E	d%	w <sub>LB</sub> (G)	w(G)	theta(G)	X <sub>f</sub> (G)	X <sub>LB</sub> (G)	X(G)	X <sub>UB</sub> (G)
2	1-FullIns_4	93	593	0,14	3					5	
3	2-FullIns_4	212	1621	0,07	4					6	
4	3-FullIns_3	80	346	0,11	5					6	
5	4-FullIns_3	114	541	0,08	6					7	
6	5-FullIns_3	154	792	0,07	7					8	
7	4-Insertions_3	79	156	0,05	2					4	
8	ash608GPIA	1216	7844	0,01	3					4	
9	ash958GPIA	1916	12506	0,01	3					4	
10	le450_15a	450	8168	0,08	15					15	
11	mug100_1	100	166	0,03	3					4	
12	mug100_25	100	166	0,03	3					4	
13	qg.order40	1600	62400	0,05	40					40	
14	wap05a	905	43081	0,11	50					50	
15	myciel6	95	755	0,17	2					7	

Figure 3: NP-m instances

1		N	V	d%	w_LB(G)	w(G)	theta(G)	X_f(G)	X_LB(G)	X(G)		X_UB(G)
2	flat300_28_0	300	21695	0,48	12			28		28		28
3	r1000.5	1000	238267	0,48	234			234		234		234
4	r250.5	250	14849	0,48	65			65		65	[GM10]	
5	DSJR500.5	500	58862	0,47	122			122		122		122
6	DSJR500.1c	500	121275	0,97	83			85		85	[GM10]	
7	DSJC125.5	125	3891	0,50	10			16		17	[GM10]	
8	DSJC125.9	125	6961	0,90	34			43		44	[GM10]	
9	DSJC250.9	250	27897	0,90	43			72		72	[HSC11]	72
10	queen10_10	100	2940	0,59	10			10		11	[GM10]	
11	queen11_11	121	3960	0,55	11			11		11	[GM10]	
12	queen12_12	144	5192	0,50	12			12		12	[Vas04]	
13	queen13_13	169	6656	0,47	13			13		13	[Vas04]	
14	queen14_14	196	4186	0,22	14			14		14	[Vas04]	
15	queen15_15	225	5180	0,21	15			15		15	[Vas04]	

Figure 4: NP-h instances

1		V(G)	E(G)	d%	w_LB(G)	w(G)	theta(G)	X_f(G)	X_LB(G)	X(G)		X_UB(G)
2	le450_5a	450	5714	0,06	5			5	5	5		5
3	le450_5b	450	5734	0,06	5			5	5	5		5
4	le450_15b	450	8169	0,08	15			15	15	15		15
5	le450_15c	450	16680	0,17	15			15	15	15		15
6	le450_15d	450	16750	0,17	15			15	15	15		15
7	le450_25c	450	17343	0,17	25			25	25	25		25
8	le450_25d	450	17425	0,17	25			25	25	25		25
9	myciel7	191	2360	0,13	2			5	5	8	[MT09]	8
10	1-FullIns_5	282	3247	0,08	3			4	4	6	[MT09]	6
11	2-FullIns_4	212	1621	0,07	4			5	5	6	[MT09]	6
12	2-FullIns_5	852	12201	0,03	4			5	5	7	[MT09]	7
13	3-FullIns_4	405	3524	0,04	5			6	6	7	[MT09]	7
14	4-FullIns_4	690	6650	0,03	6			7	7	8	[MT09]	8
15	5-FullIns_4	1085	11395	0,02	7			8	8	9	[MT09]	9

Figure 5: NP-? instances

### 3 Baseline Algorithms

Previous algorithms that show good results are the Greedy Algorithm(choose to assign colors to uncolored vertexes in a particular order),DSATUR, Tabu Search(starts with invalid k colorings and attempts to reduce the number of total conflicts by finding better solutions through exploring the neighborhood via tabu search),Quantum Simulated Annealing(add the forces of attraction and repulsion as a way of visiting the neighborhood of a solution to the Simulated Annealing).

### 3.1 Particle Swarm Optimization

Particle Swarm Optimization (PSO) stands as a noteworthy paradigm in the realm of optimization algorithms, harnessing inspiration from the collective behaviors observed in nature to tackle complex problem spaces. Originating in the late 1990s through the seminal work of Eberhart and Kennedy, PSO has evolved into a widely recognized meta-heuristic, admired for its simplicity, computational efficiency, and efficacy in navigating high-dimensional solution spaces [2].

The quest for an improved version of Particle Swarm Optimization (PSO) stems from the inherent desire to enhance the algorithm’s robustness and efficiency in addressing complex optimization challenges. While the standard PSO exhibits admirable simplicity and effectiveness, its performance can be influenced by factors such as parameter settings and convergence speed. An improved iteration seeks to overcome these limitations by introducing refinements that enhance convergence rates, promote a more thorough exploration of the solution space, and mitigate premature convergence.

The framework of the conventional PSO algorithm is derived from [2]. The provided pseudocode outlines the structure of the Particle Swarm Optimization (PSO) algorithm. The algorithm initializes a population of particles with random positions and velocities. It then defines the personal best positions for each particle and initializes the global best position. The fitness function, used to evaluate the quality of solutions, is also specified. The core of the algorithm lies within a loop, where particles are iteratively evaluated, and their positions and velocities are updated based on the PSO update equations. The process continues until certain termination conditions are met.

The variant of the Particle Swarm Optimization presented in the lines above is global also called ( $g_{best}$ ). Unlike the local version ( $l_{best}$ ) in which several neighborhoods are considered, in this case, all particles are in the same neighborhood, as can be seen in fig. 1.

This variant was also used in the implementation carried out in order to execute the proposed experiment - determining the optimum values for graphs provided as inputs in terms of number of colors used for graph coloring.

An important aspect that should be mentioned is the fact that the  $l_{best}$  approach leads to greater diversity, but this approach is slower in terms of execution time required to determine the optimal value compared to the  $g_{best}$  approach.

The pseudocode in **Algorithm 1** serves as a high-level representation of the key steps and operations involved in the standard PSO algorithm.

The velocity update equation for particle  $i$  is given by:

$$v_i = wv_i + c_1r_1(pbest_i - p_i) + c_2r_2(gbest - p_i) \quad (1)$$

The position update equation for particle  $i$  is given by:

$$p_i = p_i + v_i \quad (2)$$

Here,  $w$  represents the inertia weight,  $c_1$  and  $c_2$  are the acceleration coefficients,  $r_1$  and  $r_2$  are random numbers,  $pbest_i$  is the personal best position for particle  $i$ ,  $gbest$  is the

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**Algorithm 1** Particle Swarm Optimization (PSO)

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Initialize particles with random positions and velocities  
Initialize personal best positions for each particle  
Initialize global best position  
Define fitness function for evaluation termination conditions not met each particle  
Evaluate fitness using the fitness function  
Update personal best if current position is better  
Update global best if any particle has a better personal best  
Update velocity and position using the PSO update equations

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global best position, and  $p_i$  and  $v_i$  are the current position and velocity of particle  $i$ , respectively.

## 4 SOTA

### 4.1 Graph Colouring Meets Deep Learning: Effective Graph Neural Network Models for Combinatorial Problems

The abstract of the paper [4] discusses the recent advancements in deep learning, particularly in symbolic domains, and introduces the concept of employing Graph Neural Networks (GNNs) for solving combinatorial problems. It highlights the success of GNNs in training on complex relational data, including NP-Complete problems like SAT and TSP. The study focuses on how a simple GNN architecture can effectively tackle the fundamental combinatorial problem of graph coloring. The results demonstrate high accuracy on random instances and generalization to unseen graph distributions, outperforming several baseline methods. Additionally, the study shows how vertex embeddings can be clustered to provide constructive solutions, even though the model is trained solely as a binary classifier. Overall, the findings contribute to bridging the gap in understanding the algorithms learned by GNNs and provide empirical evidence of their capability in solving hard combinatorial problems, furthering the integration of robust learning and symbolic reasoning in deep learning systems.

One approach to incorporating relational structures into neural models involves ensuring permutation invariance through neural modules with parameter sharing. This leads to the development of various models such as message-passing neural networks, recurrent relational networks, and Graph Neural Networks (GNNs). The paragraph also discusses recent work on GNNs tackling NP-Complete problems like boolean satisfiability (SAT) and introduces a new model to solve the graph coloring problem (GCP) without prior reductions. The GNN framework is employed for this purpose, leveraging its ability to handle various types of edges. The authors aim to promote the adoption and further research on GNN-like models integrating deep learning and combinatorial optimization.

#### 4.1.1 A GNN model for decision GCP

In typical GNN models, vertices and edges in the graph are assigned multidimensional representations or embeddings  $\in \mathbb{R}^d$ , which are refined based on adjacency information through message-passing iterations. The adjacency information filters valid incoming messages for each vertex or edge, aggregates these messages, and computes updates to the embeddings using a Recurrent Neural Network (RNN). The Graph Network model also allows for global graph attributes, which is suitable for the k-colorability problem. However, since each color needs its own embedding, the authors chose to model the k-colorability problem using a Graph Neural Network framework, which can handle multiple types of nodes. Given a GCO instance  $I = (\zeta, C)$  composed of a graph  $\zeta = (\vartheta, \varepsilon)$  and a number of colours  $C \in \mathbb{N} | C > 2$ , each colour is assigned to a random initial embedding over an uniform distribution  $C[i] \sim v(0, 1) | \forall c_i \in C$  and the model initially assigns the same embedding  $\in \mathbb{R}^d$  to all  $V$  vertices: this embedding is randomly initialised and then it becomes a trained parameter learned by the model.

To allow the communication between neighbouring vertices and between vertices and colours, besides the vertex-to-vertex adjacency matrix  $M_{VV} \in \{0, 1\}^{|\mathcal{V}| \times |\mathcal{V}|}$ , the model also requires a vertex to-colour adjacency matrix  $M_{VC} \in \{0, 1\}^{|\mathcal{V}| \times |\mathcal{C}|}$ , that connects each colour to all vertices since we chose to give no prior information to the model; i.e., a priori any vertex can be assigned to any colour. After this initialisation adjacent vertices and colours communicate and update their embeddings during a given number of iterations. Then the resulting vertex embeddings are fed into a MLP which computes a logit probability corresponding to the model’s prediction of the answer to the decision problem: “does the graph  $G$  accept a  $C$ -coloration?”. This procedure is summarised in Algorithm 1 (see Figure 6).

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1: procedure GNN-GCP( $\mathcal{G} = (\mathcal{V}, \mathcal{E}), C$ )
2:
3:   // Compute binary adjacency matrix from vertex to vertex
4:    $\mathbf{M}_{\mathcal{V}\mathcal{V}}[i, j] \leftarrow 1$  iff  $(\exists e \in \mathcal{E} | e = (v_i, v_j)) \mid \forall v_i \in \mathcal{V}, v_j \in \mathcal{V}$ 
5:
6:   // Compute binary adjacency matrix from vertices to colours
7:    $\mathbf{M}_{\mathcal{V}\mathcal{C}}[i, j] \leftarrow 1 \forall v_i \in \mathcal{V}, c_j \in \mathcal{C}$ 
8:
9:   // Compute initial vertex embeddings
10:   $\overset{(1)}{\mathbf{V}}[i] \sim \mathcal{N}(0, 1) \mid \forall v_i \in \mathcal{V}$ 
11:
12:  // Compute initial colour embeddings
13:   $\overset{(1)}{\mathbf{C}}[i] \sim \mathcal{U}(0, 1) \mid \forall c_i \in \mathcal{C}$ 
14:
15:  // Run  $t_{max}$  message-passing iterations
16:  for  $t = 1 \dots t_{max}$  do
17:    // Refine each vertex embedding with messages received
    from its neighbours and candidate colours
18:     $\overset{(t+1)}{\mathbf{V}}_h, \overset{(t+1)}{\mathbf{V}} \leftarrow V_u(\overset{(t)}{\mathbf{V}}_h, \mathbf{M}_{\mathcal{V}\mathcal{V}} \times \overset{(t)}{\mathbf{V}}, \mathbf{M}_{\mathcal{V}\mathcal{C}} \times \underset{msg}{C}(\overset{(t)}{\mathbf{C}}))$ 
19:    // Refine each colour embedding with messages received
    from all vertices
20:     $\overset{(t+1)}{\mathbf{C}}_h, \overset{(t+1)}{\mathbf{C}} \leftarrow C_u(\overset{(t)}{\mathbf{C}}_h, \mathbf{M}_{\mathcal{V}\mathcal{C}}^T \times \underset{msg}{V}(\overset{(t)}{\mathbf{V}}))$ 
21:    // Translate vertex embeddings into logit probabilities
22:     $V_{logits} \leftarrow V_{vote} \left( \overset{(t_{max})}{\mathbf{V}} \right)$ 
23:    // Average logits and translate to probability (the operator
     $\langle \rangle$  indicates arithmetic mean)
24:    prediction  $\leftarrow \text{sigmoid}(\langle \mathbf{V}_{logits} \rangle)$ 

```

Figure 6: Algorithm 1 Graph Neural Network Model for GCP



The GNN-based algorithm updates vertex and colour embeddings, along with their respective hidden states, according to the following equations:

$$V^{(t+1)}, V_h^{(t+1)} \leftarrow \vartheta_u(V_h^{(t)}, M_{\vartheta\vartheta} \times (V^{(t)}), M_{\vartheta C} \times C_{msg}(C^{(t)})) \quad (1)$$

$$C^{(t+1)}, C_h^{(t+1)} \leftarrow C_u(C_h^{(t)}, M_{\vartheta C} \times V_{msg}(V^{(t)})) \quad (2)$$

In this case, the GNN model needs to learn the message functions (implemented via MLPs)  $C_{msg} : \mathbb{R}^d \rightarrow \mathbb{R}^d$ , which will translate colours embeddings into messages that are intelligible to a vertex update function, and  $V_{msg} : \mathbb{R}^d \rightarrow \mathbb{R}^d$ , responsible for translating vertices embeddings into messages. Also, it learns a function (RNN) responsible for updating vertices  $V_u : \mathbb{R}^d \rightarrow \mathbb{R}^d$  given its hidden state and received messages and another RNN to do the analogous procedure to the colors  $V_c : \mathbb{R}^d \rightarrow \mathbb{R}^d$ .

#### 4.1.2 Training Metodology

With this embedding as a graph problem as well as the given number of colors  $\mathcal{C}$ , we then let the RNN simulate the message passing (passing of the information from nodes adjacent to each other). This network will work as a binary classifier in order to predict the chromatic number  $\chi$  of the GCP. In order to train this network, we can use randomly generated instances  $I$  with  $\chi(I) = \mathcal{C}$  and create an adversarial instance  $I'$  that has the chromatic number  $\chi(I') = \mathcal{C} + 1$ .

The MLPs responsible for message computing are three-layered (64,64,64) with ReLU nonlinearities as the activations for all layers except for the linear activation on the output layer. The RNN cells are basic LSTM cells with layer normalisation and ReLU activation.

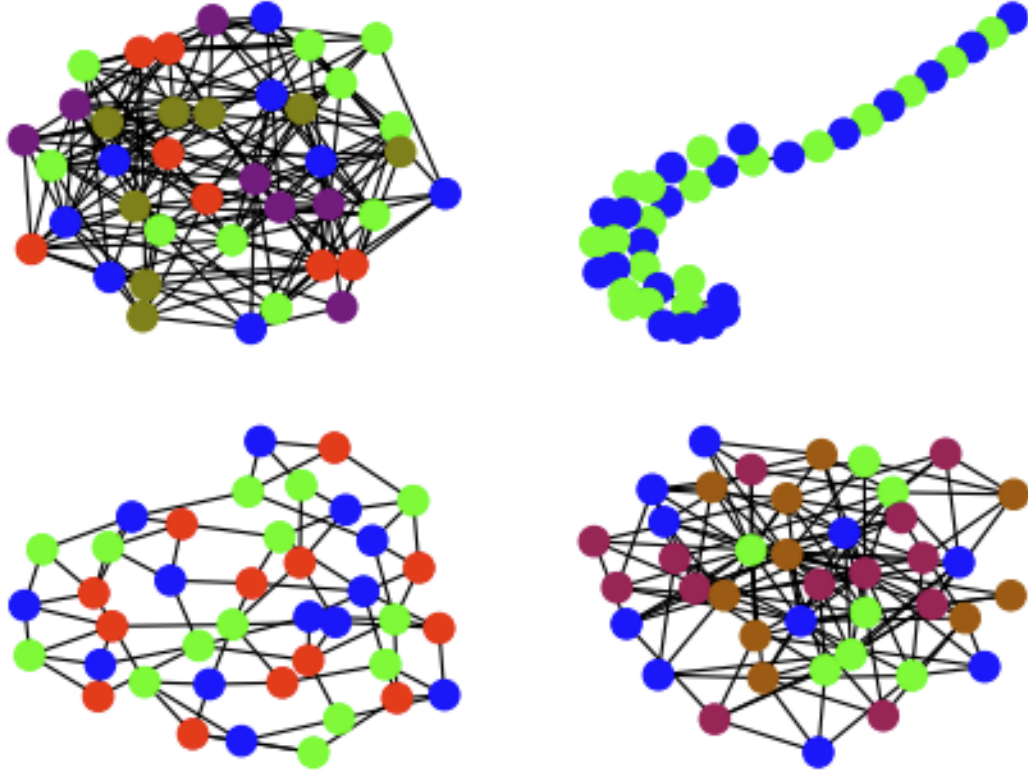


Figure 7: Pictorial representation of random training instances (top left) and some of the structured test instances (clockwise: power-law tree, power-law cluster, small-world). All instances are coloured with a number of colours equal to their chromatic number.

#### 4.1.3 Experimental results and analyses

The authors stopped the training procedure when the model achieved 82% accuracy and 0.35 Binary Cross Entropy loss averaged over 128 batches containing 16 instances at the end of 5300 epochs. In Figure 8 you can see a comparison between GNN and other algorithms. Figure 9 and Figure 10 present more results of the GNN in comparison with the others.

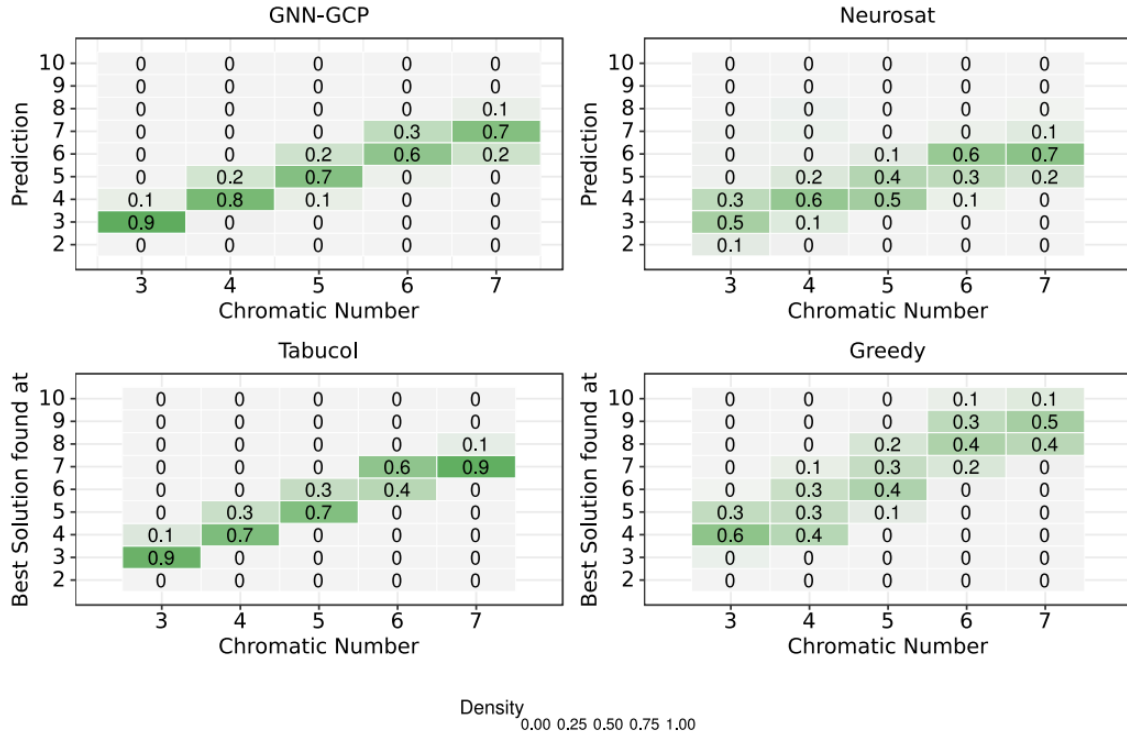


Figure 8: Prediction distributions over 4096 unseen test instances, with similar features to those seen in training, for GNN model, Neurosat, Tabucol and the Greedy algorithm. Note that the darker the main diagonal, the better the predictions.

Instance	Size	$\chi$	GNN	Computed $\chi$	
				Tabucol	Greedy
queen5_5	25	5	6	<b>5</b>	8
queen6_6	36	7	<b>7</b>	8	11
myciel5	47	6	5	<b>6</b>	<b>6</b>
queen7_7	49	7	8	8	10
queen8_8	64	9	8	10	13
1-Insertions_4	67	4	<b>4</b>	5	5
huck	74	11	8	<b>11</b>	<b>11</b>
jean	80	10	7	<b>10</b>	<b>10</b>
queen9_9	81	10	9	11	16
david	87	11	9	<b>11</b>	12
mug88_1	88	4	3	<b>4</b>	<b>4</b>
myciel6	95	7	<b>7</b>	<b>7</b>	<b>7</b>
queen8_12	96	12	10	<b>12</b>	15
games120	120	9	6	<b>9</b>	<b>9</b>
queen11_11	121	11	12	NA	17
anna	138	11	<b>11</b>	<b>11</b>	12
2-Insertions_4	149	4	<b>4</b>	5	5
queen13_13	169	13	14	NA	21
myciel7	191	8	NA	<b>8</b>	<b>8</b>
homer	561	13	14	<b>13</b>	15

Figure 9: The chromatic number produced by the GNN model and two heuristics on some instances of the COLOR02/03/04 dataset. As the GNN model faces unseen graph sizes and larger chromatic numbers it tends to underestimate its answers.

Distribution	GNN Accuracy [%]	Tabucol Accuracy [%]	Greedy Accuracy [%]
Power Law Tree	100.0	93.0	100.0
Small-world	90.0	77.0	9.0
Holme and Kim	54.1	76.4	100.0

Figure 10: Strict accuracy of the GNN model and the two algorithms considering three random graph distributions.

## 4.2 Acceleration Based Particle Swarm Optimization for Graph Coloring Problem

Particle swarm optimization is a simple and powerful technique employed to solve the graph coloring problem. Its main drawback is its tactlessness of being trapped in the local optimum. Therefore, to overcome this, an efficient Acceleration based Particle Swarm Optimization (APSO) was introduced. Empirical study of the proposed APSO algorithm is performed on the second DIMACS challenge benchmarks. The APSO results are compared with the standard PSO algorithm and experimental results validates the superiority of the proposed APSO.

### 4.2.1 Description of the concept

Particle swarm optimization is a swarm intelligence approach which is inspired by the intelligence, experience-sharing, social behavior of bird flocking and fish schooling. In 1995 Kennedy and Eberhart developed PSO to solve continuous-valued space but later on it was modified for binary/discrete optimization problems. PSO is a searching technique in which a collection of particles find the global minimum that move through the search space. In PSO, a group of a particle's position and particle velocity is initialized randomly. Each particle position represents a possible solution and particle velocity represents the rate of changes of the next position with respect to the current position.

In PSO, there are local and global extremes. The fitness value is computed. If this fitness value has better-quality than the best fitness value then the current fitness value is set as new local best. Now the particle with the best fitness value of all particles is chosen as the global best. Finally for each particle velocity and particle position is calculated and updated using equations (1) and (2) respectively. It has been used across a wide range of applications, such as image and video analysis, design and restructuring of electricity networks, control, antenna design, electronics and electromagnetic. The advantages of particle swarm optimization are there is no centralized controller in the system. Hence, failure of any particle does not affect the search process. PSO is a simple, easy and only few parameters needed to be adjusted.

$$Vid(t + 1) = wVid(t) + c_1R_1(Pid(t) - Xid(t)) + c_2R_2(Pgd(t) - Xid(t)) \quad (3)$$

$$Xid(t + 1) = Xid(t) + Vid(t + 1) \quad (4)$$

where,

Vid - Velocity of the  $i^{th}$  particle of dimension d

Pid - Best previous position of the  $i^{th}$  Particle of dimension d

Pgd - Best position of the neighbours of dimension d

Xid - Current position of the  $i^{th}$  particle of dimension d

$R_1, R_2$  - Random function in the range [0, 1]

w - Inertia weight that forces the particle to move in the same direction of the previous iteration

$c_1$  &  $c_2$  - Acceleration constants. The first coefficient “ $c_1$ ” controls the impact of the cognitive component on the particle trajectory and the second coefficient, “ $c_2$ ” controls the impact of the social component. While  $c_1$  component is responsible to maintain the diversity,  $c_2$  component is responsible to ensure convergence.

If  $c_1 \ll c_2$ , then particles may result into premature convergence as particles are attracted more towards the global best position then to their personal best positions. If,  $c_2 \ll c_1$  then particles may result in slow convergence or may not converge at all because each particle is more attracted to its personal best positions then to global best. Since the PSO relies on a combination of both personal and social knowledge of the given search space, so the coefficients  $c_1$  and  $c_2$  are chosen such that  $c_1 \cong c_2$ . For too large values of  $c_1$  and  $c_2$ , particle velocities accelerate too fast, leading in swarm divergence. On the other hand for too small values of  $c_1$  and  $c_2$ , swarm convergence time increases as particles move too slowly. In order to find the optimum solution efficiently proper control on global exploration and local exploration is crucial. Generally high diversity is necessary during the early part of the search to allow the use of the full range of the search space on the other hand, during the latter part of the search, when the algorithm is converging to the optimal solution, fine tuning of the solution is important to find the global optima efficiently. Therefore a proper control of these two components is very important to find the optimum solution accurately and efficiently. Particles draw their strength from their cooperative nature, and are most effective when  $c_1$  and  $c_2$  are adaptive to the particle value to facilitate exploitation and exploration of the search area. Hence to enhance the performance of PSO, Acceleration based PSO(APSO) was introduced.

#### 4.2.2 Acceleration Based Particle Swarm Optimization (APSO)

In this research work [1], an Acceleration based Particle Swarm Optimization (APSO) algorithm is developed to solve premature convergence problem of the standard PSO. In APSO, the acceleration coefficients are selected based on the fitness value which will increase the accuracy of the results. The selection of acceleration coefficient values in APSO algorithm is described as follows:

$$nc_1 = c_1 \times (1 - \lambda) \quad (5)$$

$$nc_2 = c_2 \times (1 + \lambda) \quad (6)$$

In eq. (3) and (4), the  $\lambda$  value is computed based on the fitness value and is calculated as:

$$\lambda = \frac{\chi(1 + \varphi(f_{max} - f_{min})^\omega - (f_{avg}))}{\delta(f_{max} - f_{min})^\omega - f_{avg}^\omega} \quad (7)$$

$$\lambda = \left( \frac{f_{max} - f_{min}}{f_{avg}} \right)^\omega \quad (8)$$

Where,

$\chi$  - Alteration probability

$\omega, \varphi$  - Coefficient factors

$F_{max}, F_{min}, F_{avg}$  - Maximum, minimum and average fitness of the particles

By exploiting eq. (3) and (4), the acceleration coefficients values, the velocity formula which is given in eq. (1) is updated by eq. (7). However the position update equation remains the same:

$$Vid(t+1) = wVid(t) + nc_1R_1(Pid(t) - Xid(t)) + nc_2R_2(Pgd(t) - Xid(t)) \quad (9)$$

#### 4.2.3 Experiments and Results

The values of parameters in the APSO algorithm for the graph coloring problem are presented in Table 1. These values are selected based on some preliminary trials. To validate the performance of the proposed APSO algorithm, an extensive experiments on some benchmarks are conducted. Benchmark graphs are derived from the well known DIMACS challenge benchmarks illustrated in Table 2. These instances cover a variety of types and sizes of graphs. For each instance in table 2, 10 independent runs of the algorithm were carried out. Table 3 shows the experimental results. Column 2 records the graph name, column 3 records the expected chromatic number. Column 4 is reported for standard PSO which is divided in 2 sub columns in which first column represent the total steps and other column represent PSO result. Column 5 is recorded for APSO which is divided into two parts in which one part represent the total steps and second part represent the APSO result. We run APSO independently on every graph and found that in case of Mycielski graphs such as myciel3.col, myciel4.col, myciel5.col, APSO found the optimal solution in each run. Then Stanford GraphBase (SGB) such as huck.col, jean.col, david.col. games120.col, anna.col is tested independently and APSO found an optimal solution in each run. For miles graphs was found that for miles250.col APSO is able to produce an optimal solution in each run while miles1000.col produces an optimal solution in 9 runs. On queen graph i.e on queen 5\_5.col an optimal solution with the predefined parameter was not obtained. Hence. the parameter value of population size was updated to 50 and maximum step size to 500 then the optimal solution was obtained using APSO.

#### 4.2.4 PSO VS APSO comparison

Standard PSO was compared with the proposed APSO algorithm. Experimental results show that the APSO algorithm for graph coloring problem is feasible and robust. The algorithm was able to scale across the different graphs and produce optimum solutions in each case. APSO is capable of finding an optimal solution for all the graphs with a very high probability. APSO finds an optimal solution in each run for all graphs except miles1000.col and queen 5\_5.col in 9 and 7 runs respectively while standard PSO was unable to find an optimal solution in 100 steps. The standard PSO algorithm failed to produce the optimum solution for all graphs except the two graphs named myciel3.col and myciel4.col. From the simulation result, it has been observe that APSO is competitive with the standard PSO algorithm.

Table 1: Parameter setting

Parameter	Population size	Maximum step size	Inertia weight	Maximum velocity
Value	100	100	0.9	10

Table 2: Benchmark Graphs

S No	Graph Name	Vertices	Edges
1	myciel3.col	11	20
2	myciel4.col	23	71
3	myciel5.col	47	236
4	huck.col	74	301
5	jean.col	80	254
6	david.col	87	406
7	games120.col	120	638
8	miles250.col	128	387
9	miles1000.col	128	3216
10	anna.col	138	493
11	queen5_5.col	25	160



Table 3: Experimental Results

S No	Graph	Expect. $\chi(G)$	Steps PSO	PSO res	Steps APSO	APSO res
1	myciel3.col	4	14	4	1	4
2	myciel4.col	5	59	5	1	5
3	myciel5.col	6	100	-	1	6
4	huck.col	11	100	-	1	11
5	jean.col	10	100	-	1	10
6	david.col	11	100	-	1	11
7	games120.col	9	100	-	1	9
8	miles250.col	8	100	-	1	8
9	miles1000.col	42	200	-	2	42
10	anna.col	11	100	-	1	11
11	queen5_5.col	5	100	-	32	5

#### 4.2.5 Results analysis

In this paper was proposed an Acceleration based Particle Swarm Optimization method (APSO) to solve the graph coloring problem. With a view to achieve a highly accurate optimal outcome, the defects inherent in the PSO technique such as premature convergence and loss of diversity. These have to be properly tackled and surmounted by initiating effective alteration or augmentation in the procedure of PSO. Therefore, to attain a further precise outcome and to steer clear of the PSO defects, rather than fixing the value of acceleration coefficient, an Acceleration base Particle swarm optimization (APSO) technique in which acceleration coefficient values are updated on the basis of evaluation function was introduced. The algorithm was tested on a set of ten DIMACS benchmark test while limiting the number of usable colors to the expected optimal coloring number. Compared with the standard PSO, was found that APSO succeeded in solving the sample data set and even outperformed the standard PSO algorithm in terms of the minimum number of colors and minimum number of steps. APSO finds an optimal solution in one step for all graphs, while standard PSO is unable to find an optimal solution in 100 steps. The standard PSO algorithm failed to produce the optimum solution for all graphs except the two graphs named myciel3.col and myciel4.col. Hence computational results show that APSO is feasible and competitive with the standard PSO algorithm.

### 4.3 Hybrid Evolutionary Algorithm in a Duet

To introduce a variation on a hybrid evolutionary algorithm [5], we use a population of 2 individuals ( $p_1$  and  $p_2$ ) initialized at random and we follow first try to extract to children from the parents by applying a Greedy Partition Crossover(GPX) after which we apply a Tabu Search from the 2 children obtained.

We also have 2 other candidate solutions (similar to elite solutions), elite1 and elite2, in order to reintroduce some diversity to the duet. Indeed, after a given number of generations, the two individuals of the population become increasingly similar within the search-space. To maintain the population diversity, the idea is to replace one of the two candidates solutions by a solution previously encountered by the algorithm. We define one cycle as a number of Itercycle generations. Solution elite1 is the best solution found during the current cycle and solution elite2 the best solution found during the previous cycle. At the end of each cycle, the elite2 solution replaces one of the population individuals. Figure 12 presents the graphic view of algorithm 2.

This elitist mechanism provides relevant behaviors to the algorithm as it can be observed in the results. Indeed, elite solutions have the best fitness value of each cycle. It is clearly interesting in terms of intensification. Moreover, when the elite solution is reintroduced, it is generally different enough from the other individuals to be relevant in terms of diversification.

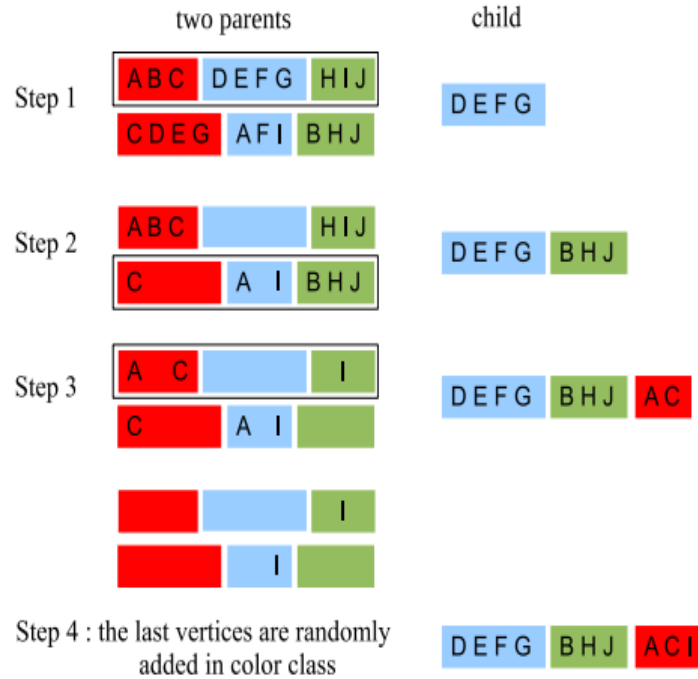


Figure 11: Greedy Partition Crossover

---

**Algorithm 2:** *HEAD* - second version of *HEAD* with two extra elite solutions

---

**Input:**  $k$ , the number of colors;  $Iter_{TC}$ , the number of *TabuCol* iterations;  $Iter_{cycle} = 10$ , the number of generations into one cycle.

**Output:** the best  $k$ -coloring found:  $best$

```

1  $p_1, p_2, elite_1, elite_2, best \leftarrow \text{init}()$  /* initialize with
   random  $k$ -colorings */
2  $generation, cycle \leftarrow 0$ 
3 do
4    $c_1 \leftarrow GPX(p_1, p_2)$ 
5    $c_2 \leftarrow GPX(p_2, p_1)$ 
6    $p_1 \leftarrow \text{TabuCol}(c_1, Iter_{TC})$ 
7    $p_2 \leftarrow \text{TabuCol}(c_2, Iter_{TC})$ 
8    $elite_1 \leftarrow \text{saveBest}(p_1, p_2, elite_1)$  /* best  $k$ -coloring of
   the current cycle */
9    $best \leftarrow \text{saveBest}(elite_1, best)$ 
10  if  $generation \% Iter_{cycle} = 0$  then
11     $p_1 \leftarrow elite_2$  /* best  $k$ -coloring of the
   previous cycle */
12     $elite_2 \leftarrow elite_1$ 
13     $elite_1 \leftarrow \text{init}()$ 
14     $cycle++$ 
15     $generation++$ 
16 while  $nbConflicts(best) > 0$  and  $p_1 \neq p_2$ 

```

---

Figure 12: HEAD Algorithm

## 4.4 Results for HEAD

Test instances are selected among the most studied graphs since the 1990s, which are known to be very difficult (the second DIMACS challenge of 1992-1993)

Graphs	HEAD	LS	Hybrid algorithm					
		1987/2008 <i>TabuCol</i> [22, 18]	1999 <i>HEA</i> [14]	2008 <i>AmaCol</i> [27]	2010 <i>MACOL</i> [28]	2011 <i>EXTRACOL</i> [29]	2012 <i>IE<sup>2</sup>COL</i> [26]	2012 <i>QA-col</i> [21]
dsjc250.5	<b>28</b>	<b>28</b>	<b>28</b>	<b>28</b>	<b>28</b>	-	-	<b>28</b>
dsjc500.1	<b>12</b>	13	-	<b>12</b>	<b>12</b>	-	-	-
dsjc500.5	<b>47</b>	49	48	48	48	-	-	<b>47</b>
dsjc500.9	<b>126</b>	127	-	<b>126</b>	<b>126</b>	-	-	<b>126</b>
dsjc1000.1	<b>20</b>	-	<b>20</b>	<b>20</b>	<b>20</b>	<b>20</b>	<b>20</b>	<b>20</b>
dsjc1000.5	<b>82</b>	89	83	84	83	83	83	<b>82</b>
dsjc1000.9	<b>222</b>	227	224	224	223	<b>222</b>	<b>222</b>	<b>222</b>
r250.5	<b>65</b>	-	-	-	<b>65</b>	-	-	<b>65</b>
r1000.1c	<b>98</b>	-	-	-	<b>98</b>	101	<b>98</b>	<b>98</b>
r1000.5	245	-	-	-	245	249	245	<b>234</b>
dsjr500.1c	<b>85</b>	<b>85</b>	-	86	<b>85</b>	-	-	<b>85</b>
le450_25c	<b>25</b>	26	26	26	<b>25</b>	-	-	<b>25</b>
le450_25d	<b>25</b>	26	-	26	<b>25</b>	-	-	<b>25</b>
flat300_28_0	31	31	31	31	<b>29</b>	-	-	31
flat1000_50_0	<b>50</b>	<b>50</b>	-	<b>50</b>	<b>50</b>	<b>50</b>	<b>50</b>	-
flat1000_60_0	<b>60</b>	<b>60</b>	-	<b>60</b>	<b>60</b>	<b>60</b>	<b>60</b>	-
flat1000_76_0	<b>81</b>	88	83	84	82	82	<b>81</b>	<b>81</b>
C2000.5	146	-	-	-	148	146	<b>145</b>	<b>145</b>
C4000.5	266	-	-	-	272	260	<b>259</b>	<b>259</b>

Table 1: Best coloring found

Figure 13: Results of the HEAD

## 5 Our approaches

### 5.1 GNN for chromatic number estimation

This approach consists of a graph neural network that will output an approximation of the chromatic number of a graph given as input.

#### 5.1.1 Graph Neural Networks introduction

Recently, deep learning on graphs has emerged to one of the hottest research fields in the deep learning community. Graph Neural Networks (GNNs) aim to generalize classical deep learning concepts to irregular structured data (in contrast to images or texts) and to enable neural networks to reason about objects and their relations.

This is done by following a simple neural message passing scheme, where node features  $x_v^{(l)}$  of all nodes  $v \in \vartheta$  in a graph  $G = (\vartheta, \varepsilon)$  are iteratively updated by aggregating localized information from their neighbors  $N(\nu)$ :

$$x_v^{(l+1)} = f_{\theta}^{(l+1)}(x_v^{(l)}, \{x_w^{(l)} : w \in N(\nu)\})$$

### 5.1.2 PyTorch Geometric (PyG) library

PyTorch Geometric is an extension library to the popular deep learning framework PyTorch, and consists of various methods and utilities to ease the implementation of Graph Neural Networks. Each graph in PyTorch Geometric is represented by a single Data object, which holds all the information to describe its graph representation. This data object holds 4 attributes:

1. The `edge_index` property holds the information about the graph connectivity, i.e., a tuple of source and destination node indices for each edge.
2. Node features as `x` (each of the 34 nodes is assigned a 34-dim feature vector)
3. Node labels as `y` (each node is assigned to exactly one class)
4. There also exists an additional attribute called `train_mask`, which describes for which nodes we already know their community assignments. In total, we are only aware of the ground-truth labels of 4 nodes (one for each community), and the task is to infer the community assignment for the remaining nodes.

### 5.1.3 Creating the GNN architecture

The GCN layer is defined as  $x_v^{(l+1)} = W^{(l+1)} \sum_{w \in N(\nu) \cup \{\nu\}} \frac{1}{C_{w,\nu}} x_w^{(l)}$  where  $W^{(l+1)}$  denotes a trainable weight matrix of shape `[num_output_features, num_input_features]` and  $C_{w,\nu}$  refers to a fixed normalization coefficient for each edge. PyG implements this layer via `GCNConv`, which can be executed by passing in the node feature representation `x` and the COO graph connectivity representation `edge_index`.

We chose as an architecture with 3 Graph Convolutional Layers along with a global aggregation layer as well as a linear layer for regression. The reason behind this is to see how well a simple network like this performs with a small number of parameters. We also employed dropout for the linear layers and convolutional layers since we intend to have a high number of epochs to further help the aggregation of information and learning.

```

class GNNRegression3(torch.nn.Module):
    def __init__(self, device: torch.device,
                  no_hidden_units: int, layer_aggregation: str,
                  global_layer_aggregation: str,
                  linear_layer_dropout: float, conv_layer_dropout: float):
        super(GNNRegression3, self).__init__()

        self.device = device

        self.conv1 = GraphConv(1, no_hidden_units, aggr=layer_aggregation)
        self.batch_norm1 = BatchNorm(no_hidden_units)

        self.conv2 = GraphConv(no_hidden_units, no_hidden_units, aggr=layer_aggregation)
        self.batch_norm2 = BatchNorm(no_hidden_units)

        self.conv3 = GraphConv(no_hidden_units, no_hidden_units, aggr=layer_aggregation)
        self.batch_norm3 = BatchNorm(no_hidden_units)

        self.regression_layer = Linear(no_hidden_units, 1)

        self.global_layer_aggregation = global_layer_aggregation
        self.linear_layer_dropout = linear_layer_dropout
        self.conv_layer_dropout = conv_layer_dropout

        self.to(device)

```

Figure 14: GNN architecture

```

def forward(self, x, edge_index, batch):
    x = self.conv1(x, edge_index)
    x = self.batch_norm1(x)
    x = F.dropout(x, p=self.conv_layer_dropout, training=self.training)
    x = x.relu()

    x = self.conv2(x, edge_index)
    x = self.batch_norm2(x)
    x = F.dropout(x, p=self.conv_layer_dropout, training=self.training)
    x = x.relu()

    x = self.conv3(x, edge_index)
    x = self.batch_norm3(x)
    x = F.dropout(x, p=self.conv_layer_dropout, training=self.training)

    if self.global_layer_aggregation == "add":
        x = global_add_pool(x, batch)
    elif self.global_layer_aggregation == "max":
        x = global_max_pool(x, batch)
    else:
        # Assume mean
        x = global_mean_pool(x, batch)

    x = F.dropout(x, p=self.linear_layer_dropout, training=self.training)
    x = self.regression_layer(x)

    return x

```

Figure 15: GNN architecture

#### 5.1.4 Mini-batching of graphs

A good idea is to batch the graphs before inputting them into a Graph Neural Network to guarantee full GPU utilization. In the image or language domain, this procedure is typically achieved by rescaling or padding each example into a set of equally-sized shapes, and examples are then grouped in an additional dimension. The length of this dimension is then equal to the number of examples grouped in a mini-batch and is typically referred to as the `batch_size`.

However, for GNNs the two approaches described above are either not feasible or may result in a lot of unnecessary memory consumption. Therefore, PyTorch Geometric opts for another approach to achieve parallelization across a number of examples. Here, adjacency matrices are stacked in a diagonal fashion (creating a giant graph that holds multiple isolated subgraphs), and node and target features are simply concatenated in

the node dimension:

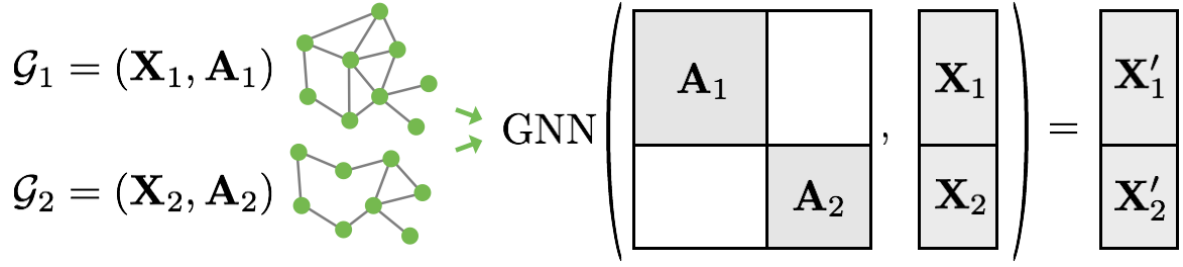


Figure 16: Graph batch

This procedure has some crucial advantages over other batching procedures:

1. GNN operators that rely on a message passing scheme do not need to be modified since messages are not exchanged between two nodes that belong to different graphs.
2. There is no computational or memory overhead since adjacency matrices are saved in a sparse fashion holding only non-zero entries, i.e., the edges.

PyTorch Geometric automatically takes care of batching multiple graphs into a single giant graph with the help of the `torch_geometric.data.DataLoader` class.

### 5.1.5 Training a Graph Neural Network (GNN)

Training a GNN for graph classification usually follows a simple recipe:

1. Embed each node by performing multiple rounds of message passing
2. Aggregate node embeddings into a unified graph embedding (readout layer)
3. Train a final classifier on the graph embedding

There exists multiple readout layers in literature, but the most common one is to simply take the average of node embeddings:

$$x_G = \frac{1}{|V|} \sum_{v \in V} x_v^{(L)}$$

PyTorch Geometric provides this functionality via `torch_geometric.nn.global_mean_pool`, which takes in the node embeddings of all nodes in the mini-batch and the assignment vector batch to compute a graph embedding of size `[batch_size, hidden_channels]` for each graph in the batch.



### 5.1.6 Training instances

It is very important to have a big balanced training set, otherwise, if most of our instances have the chromatic number 4 and some instances with chromatic number 2, 3 and 10, the neural network will tend to predict numbers close to 4 for graphs with chromatic number of 7,8, 12 for example. It would be ideal to have training instances for all the possible chromatic numbers the graphs given as input can have. To find the chromatic number of a random generated instance we used ORTools to described each graph coloring problem as a constraint satisfaction problem after which we find the chromatic number and the coloring associated using binary search.

Unfortunately, because of the complexity of generating even instances with a chromatic number of 6 taking too much, we instead generated we generated random graph instances with 20 - 60 nodes and 7.5% - 25%. Due to this as well as observing that the models tend to overfit quickly to big datasets, we chose to generate 4 datasets with 10000 instances (in the ranges approximately specified above) and another dataset with 124 hard instances found on the internet. They are:

- D1: Randomly generated
- D2: Randomly generated by adding a constant number of edges until a minimum chromatic number was found in the range  $[3, 6]$ . This set is also balanced
- D3: Randomly generated upon which we form a clique and add a number of random edges. This enforces the chromatic number to be at least the clique size. This dataset is also balanced and is in the range  $[3, 8]$
- D4: Randomly generated from 100k instances chosen at random and is in the range  $[2, 6]$
- D5: Instances that are considered 'harder' with a chromatic number in the range  $[3, 15]$ . They are only about 124 instances

For a visual representation of the distributions of each dataset:

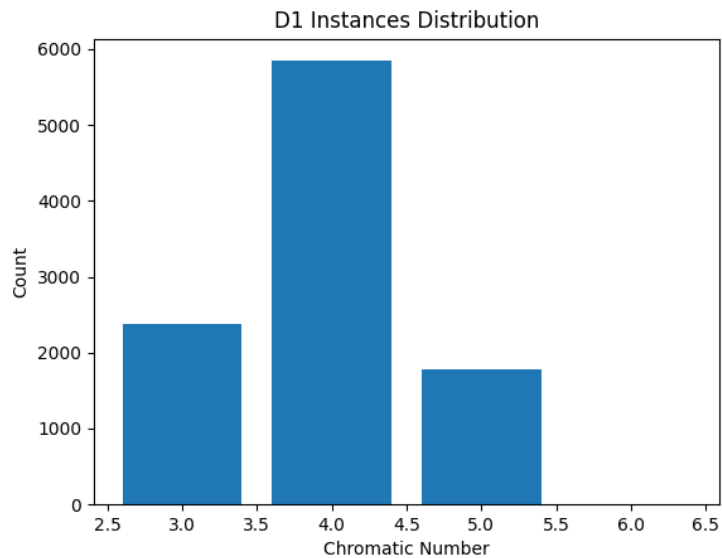


Figure 17: D1 Instance Distribution

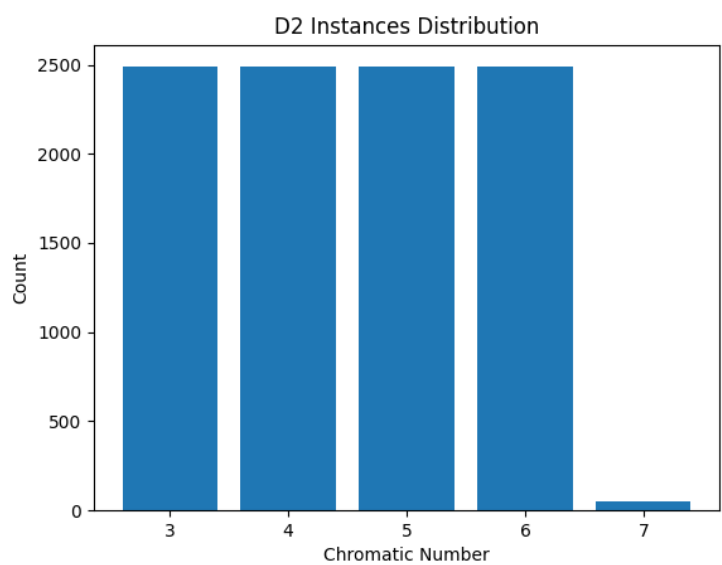


Figure 18: D2 Instance Distribution

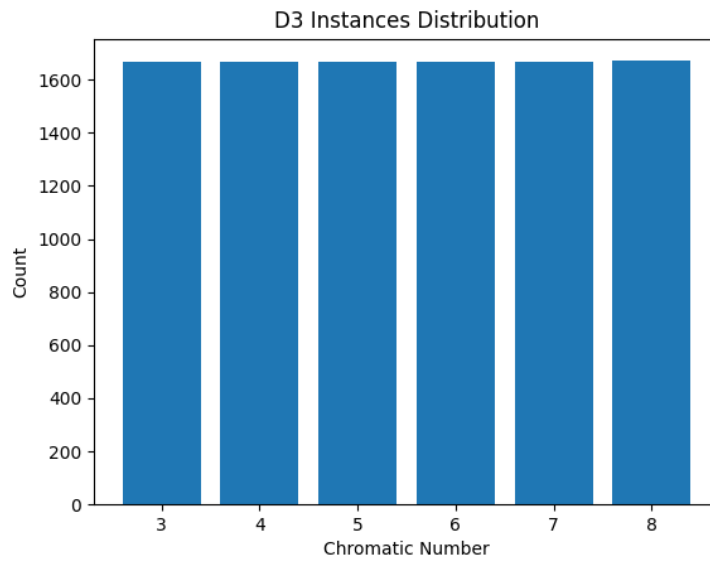


Figure 19: D3 Instance Distribution

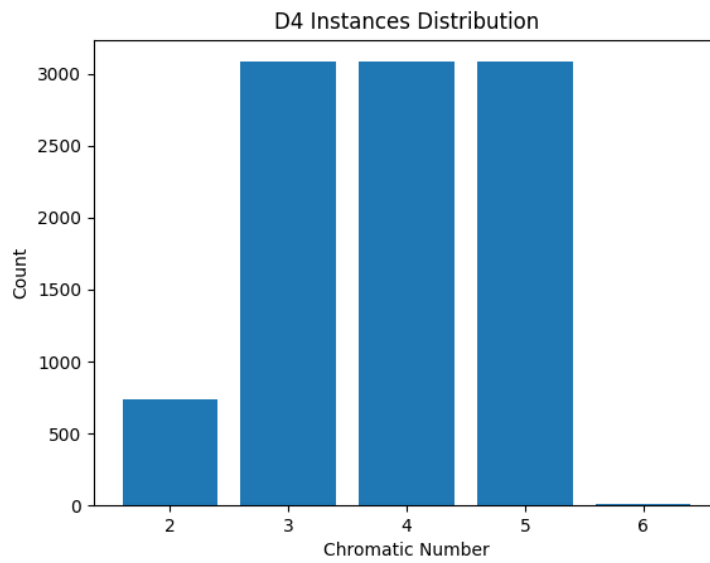


Figure 20: D4 Instance Distribution

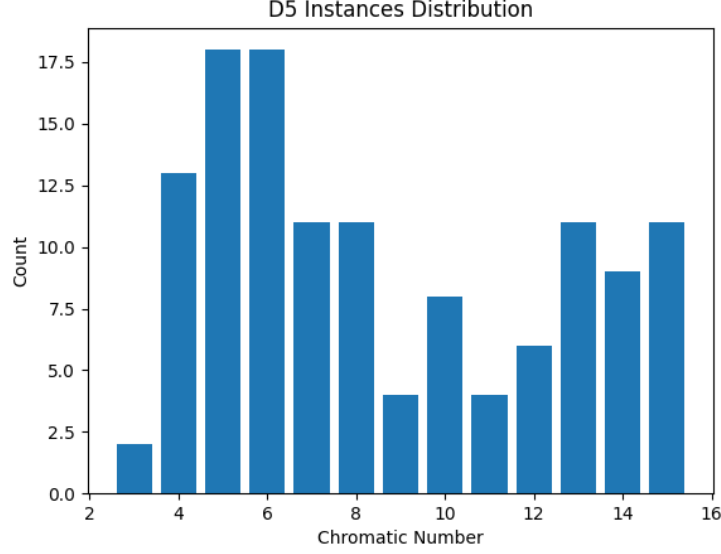


Figure 21: D5 Instance Distribution

For the validation dataset we chose to use the D5 dataset MAE loss since this better tests whether our model is generalizing than choosing a validation dataset from the training dataset. Thus, we will save the model that has the best loss for the D5 dataset as the final model. For other parameters, we used learning rate of  $1e-3$  with Adam. We chose 3 layers with a dropout of 0.5 for each layer and in order to ensure fairness we adjusted the parameters such that each model had approximately the same number of parameters (difference arising from LSTM Aggregation).

### 5.1.7 Experiments and results

For each dataset (from D1 to D5) we will train 4 models, all with 50 epochs and dropout of 0.5, all with 3 layers of graph convolutions:

- GraphConv-uses learnable graph convolutions with add aggregation
- GraphConv\_LSTM-uses LSTM as the aggregation
- SAGEConv
- SAGEConv\_LSTM-uses LSTM for the aggregation

For each dataset we will show the training loss of each model.

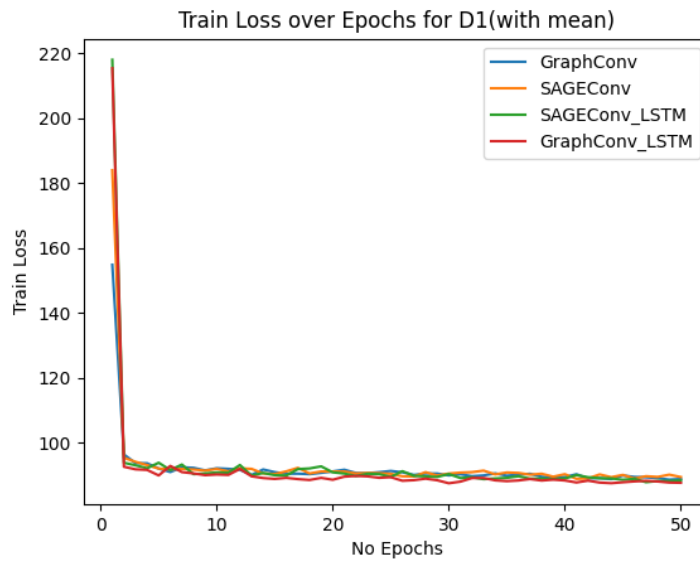


Figure 22: D1 Train Loss

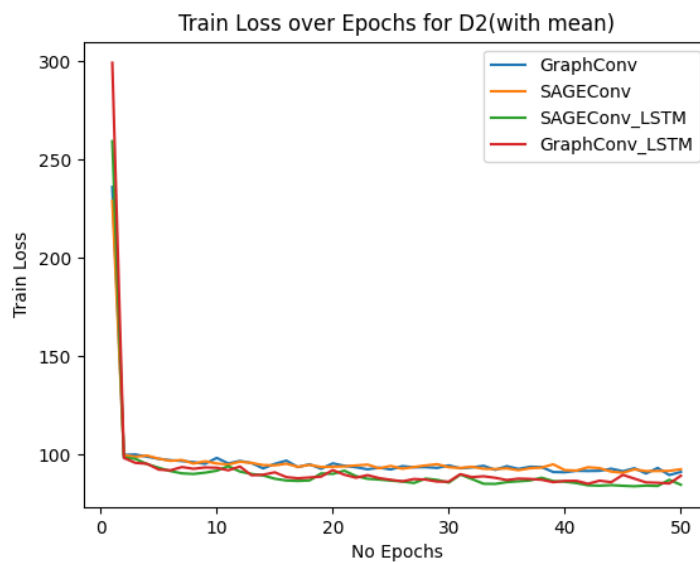


Figure 23: D2 Train Loss

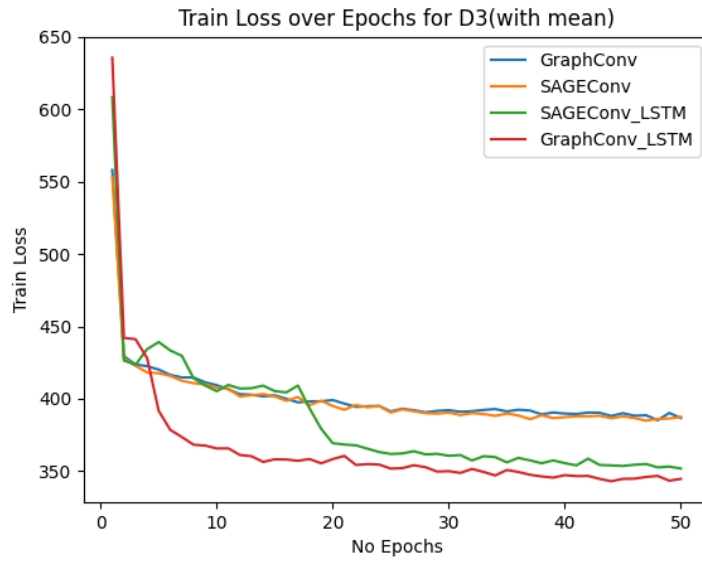


Figure 24: D3 Train Loss

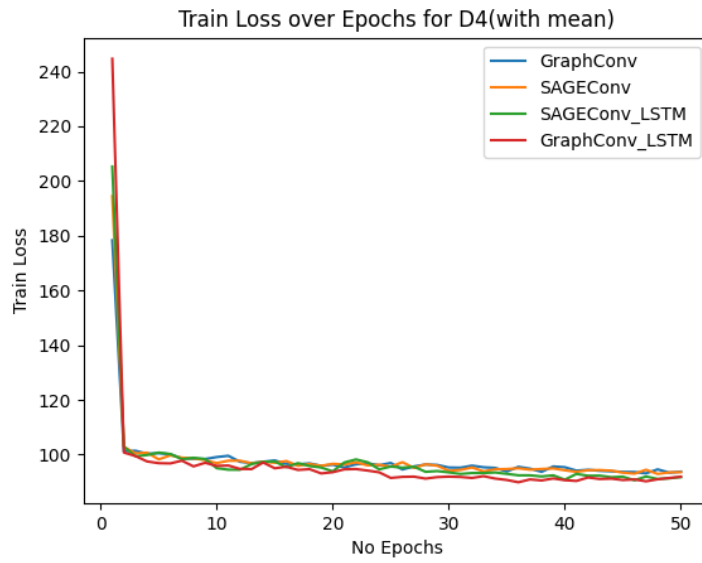


Figure 25: D4 Train Loss

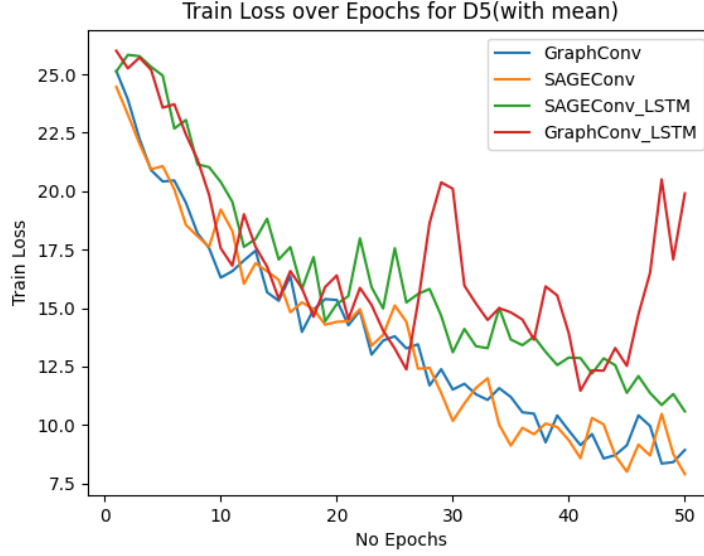


Figure 26: D5 Train Loss

From these plots, we can see how quickly each model improves its training loss for the D1, D2 and D4 datasets, but struggles a lot more with loss for the D3 dataset where we generated instances using Cliques, meaning that it struggles to find the clique to use as a factor for the coloring. For the D5 dataset it is better. We can also see a lot of spikes which is probably due to dropout.

For the validation loss of each model for each dataset:

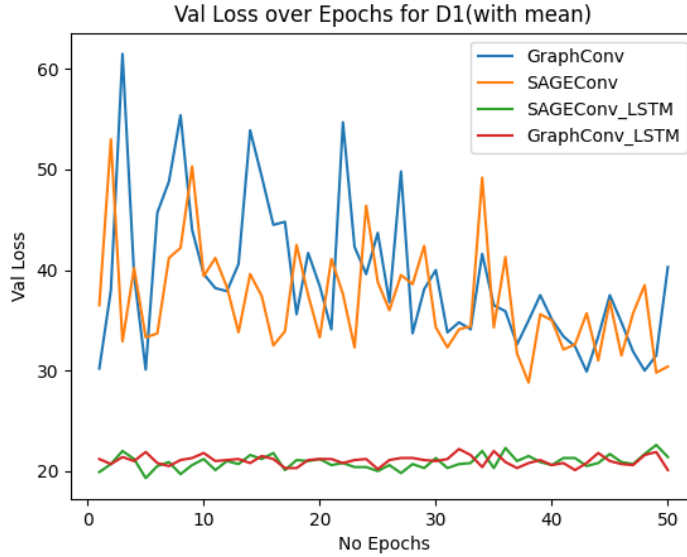


Figure 27: D1 Val Loss

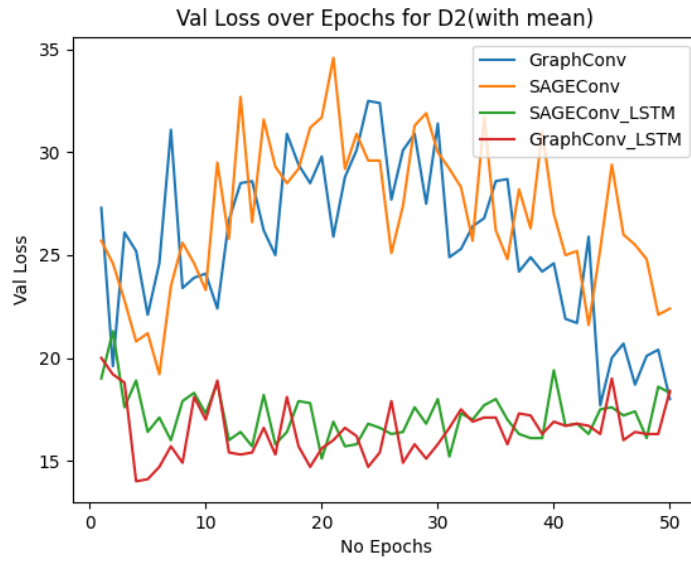


Figure 28: D2 Val Loss

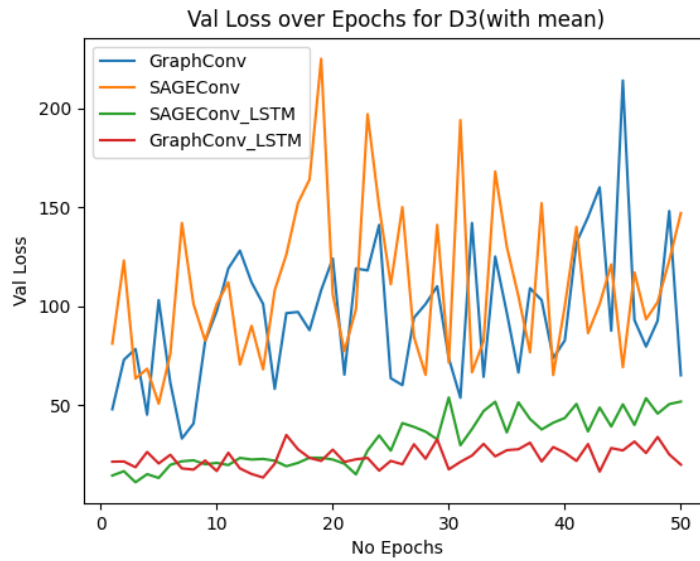


Figure 29: D3 Val Loss



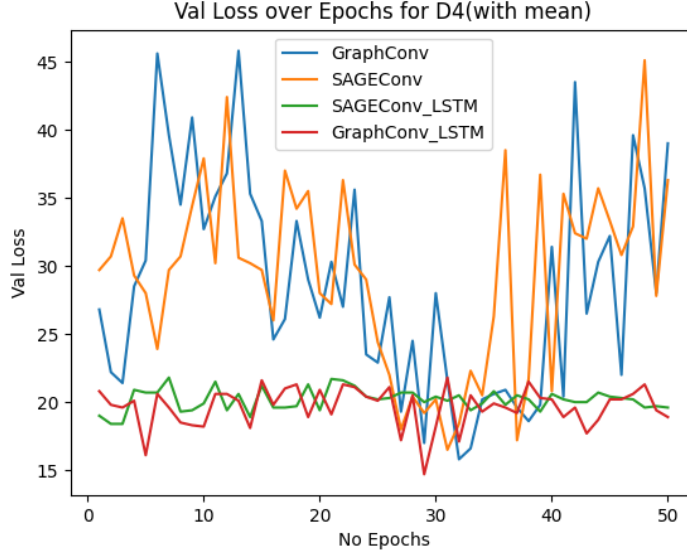


Figure 30: D4 Val Loss

Interestingly, here we can see a clear difference between models with LSTM and without, since those without even start at a lower validation loss and drop or remain at the same level, whereas the other 2 clearly learn and then start to overfit.

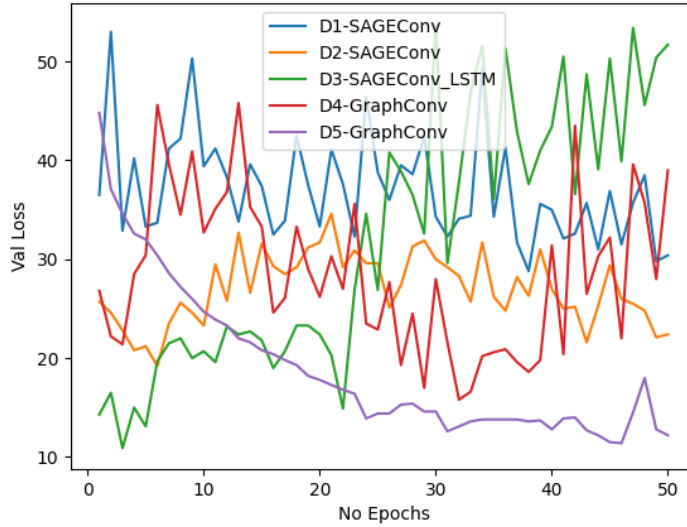


Figure 31: D1 Val Loss

Comparing the best model (having the lowest MAE error for test instances) from each dataset for the validation loss (this excludes the models D5). Interestingly, there is only one model with LSTM on D3, which makes sense seeing how the plots for D3 show LSTM having a far better validation loss. However, this also shows that the D3 dataset is seen as the hardest and we can also see how some models are the best in some datasets and worse in others.

Instances	Size	$\chi$	Computed $\chi$			
			GraphConv(D2)	SAGEConv(D2)	GraphConv_LSTM(D3)	SAGEConv_LSTM(D3)
queen5.5	25	5	6	6	7	10
queen6.6	36	7	6	6	6	9
myciel5	47	6	5	<b>6</b>	12	8
queen7.7	49	7	6	<b>7</b>	6	9
queen8.8	64	9	7	7	5	<b>9</b>
huck	74	11	4	5	14	7
jean	80	10	4	4	12	6
queen9.9	81	10	8	9	6	9
david	87	11	5	5	14	7
myciel6	95	7	6	<b>7</b>	13	9
queen8_12	96	12	9	10	8	10
games120	120	9	6	6	7	8
anna	138	11	3	3	14	7
queen13.13	169	13	14	16	8	8
myciel7	191	8	7	<b>8</b>	14	9
homer	561	13	3	4	14	6
MAE Loss	-	-	52.8194	49.5995	53.4215	44.6211

Table 4: Chromatic number by GNN obtained by the best models via MAE Loss, where each model trained on datasets D1, D2, D3 or D4 were considered

From this table, we chose the models that had lowest validation loss from each of the dataset. We can see how the LSTM models are very good for D3 (which is the Clique+Random Edges instances) and the other 2 are good at the randomly generated instances by only adding edges. Both datasets are balanced. From this we can see based in the MAE Loss that the best model is SAGEConv\_LSTM, followed by SAGE.

Instances	Size	$\chi$	Computed $\chi$			
			GraphConv(D5)	SAGEConv(D5)	GraphConv_LSTM(D5)	SAGEConv_LSTM(D5)
queen5.5	25	5	7	7	<b>5</b>	<b>5</b>
queen6.6	36	7	8	8	4	5
myciel5	47	6	7	7	<b>6</b>	<b>6</b>
queen7.7	49	7	9	8	4	5
queen8.8	64	9	<b>9</b>	8	4	5
huck	74	11	7	7	6	10
jean	80	10	7	6	6	8
queen9.9	81	10	9	8	3	5
david	87	11	9	8	6	<b>11</b>
myciel6	95	7	10	9	5	10
queen8_12	96	12	9	8	3	5
games120	120	9	8	8	5	6
anna	138	11	8	7	6	10
queen13.13	169	13	9	7	3	5
myciel7	191	8	14	11	5	11
homer	561	13	7	6	6	11
MAE Loss	-	-	42.8294	47.2741	72.4097	46.4171

Table 5: Chromatic number by GNN obtained for the best models, where each model was on the D5 dataset

From both tables, we would get that the best model to generalize is GraphConv trained on the D5 dataset (hard instances) followed closely by SAGEConv\_LSTM trained on D3.

We also tried fine tuning based on a validation dataset for minimizing the validation loss but this lead to overfitting quickly(in the first few epochs) resulting in models that always print values in the range [3, 5].

## 5.2 Improved PSO Algorithm (PSO-AWDV)

In this study [1], the authors propose an enhanced version of Particle Swarm Optimization (PSO) named Adaptive Weighted Delay Velocity PSO (PSO-AWDV). The research introduces a novel approach by incorporating weighted delay velocity into a new PSO variant termed PSO with Weighted Delay Velocity (PSO-WDV). Following this, the authors develop the Adaptive PSO-AWDV algorithm, designed to dynamically update the velocity inertia weight. This adaptive approach leverages a unique estimation method that evaluates the evolutionary state of the particle swarm. Through comprehensive testing on well-known benchmark functions, the authors demonstrate that their proposed Adaptive PSO-AWDV algorithm outperforms several established PSO variants and intelligent optimization algorithms documented in the literature. The new update equations for velocity and position are:

$$v_i = wv_i + (1 - w)v_{i-1} + c_1r_1(pbest_i - p_i) + c_2r_2(gbest - p_i) \quad (10)$$

$$p_i = p_i + v_i \quad (11)$$

Here, the only difference from standard PSO version is the term  $v_{i-1}$ , multiplied by  $(1 - w)$  which is the delayed velocity. The new adaptive scheme is developed according to the estimation of the evolutionary state of the particle swarm, thereby a new PSO with adaptive weighted delay velocity (PSO-AWDV) is presented on the basis of the updating functions:

$$c_1 = (c_{1i} - c_{1f}) \times \frac{k_{max} - k}{k_{max}} + c_{1f} \quad (12)$$

$$c_2 = (c_{2i} - c_{2f}) \times \frac{k_{max} - k}{k_{max}} + c_{2f} \quad (13)$$

where  $c_{1i}$  and  $c_{1f}$  indicate, respectively, the initial and final values of the acceleration factor  $c_1$ ; likewise, the initial and final values of the acceleration factor  $c_2$  are represented by  $c_{2i}$  and  $c_{2f}$ , respectively;  $k$  and  $k_{max}$  denote, respectively, the current and maximal iterations in the optimization. In particular, the inertia weight of velocity is adaptively regulated in accordance with the evolutionary state in the optimization, and it is calculated as follows:

$$w = 1 - \frac{a}{1 + e^{b \cdot E(k)}} \quad (14)$$

where  $a$  and  $b$  are two parameters that can be designed to adjust the search performance of PSO;  $E(k)$  is the estimation value (EV) of the evolutionary state at the  $k$ -th iteration and can be computed as follows:

$$E(k) = \frac{|f_{max}(k)| - |f_{min}(k)|}{|f_{max}(k)|} \quad (15)$$

where  $f_{max}$  and  $f_{min}$  stand for, respectively, the maximal and minimal fitness values of the particles at the  $k$ -th iteration. In sum, the PSO-AWDV algorithm pseudocode is illustrated in **Algorithm 2**.

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**Algorithm 2** PSO-AWDV Algorithm

---

**Initialize** the parameters of the PSO-AWDV algorithm, including maximum iteration  $k_{max}$ , parameters of the inertia weight  $a$  and  $b$ , parameters of the acceleration factor  $c_{1i}$ ,  $c_{1f}$ ,  $c_{2i}$ , and  $c_{2f}$ ;

**repeat**

    Evaluate the fitness of every particle of the swarm and estimate the evolutionary state  $E(k)$  of the particle swarm at the current iteration according to Equation (8);

    Compare the fitness value of each particle with its  $pbest$  and  $gbest$  of the swarm, and select the corresponding better particle to replace the original one;

    Compute the acceleration factors and inertia weight according to Equations (5) - (7);

    Update the velocity and position vectors of the particles at the current iteration according to Equations (3) and (4);

**until** Maximum iteration

---

The particle swarm optimization (PSO) algorithm was adapted for the graph coloring problem by modifying the position and velocity update rules. A particle's position represents the colors assigned to the nodes in the graph. To ensure the correctness of a solution, the PSO was modified such that every position of a particle is a list with the length equal to number of nodes containing random integer between 0 and the number of colors - 1 (representing the encoding of the colors). The velocities were clipped between a specific range to prevent them from becoming too large or too small. The positions were rounded and converted to integers in each iteration to ensure that they represented valid color assignments.

The number of iterations was set to 2500 for small graph coloring instances and 5000 for larger graph coloring instances. This number of iterations was chosen based on empirical testing, which showed that it was sufficient to find good solutions for most graphs.

In addition to the modifications mentioned above, the PSO algorithm was also implemented with a number of other parameters, including the the initial and final values for acceleration factors

$$c_{1i} = c_{2i} = 1.8$$

$$c_{1f} = c_{2f} = 1.2$$

and the specific factors used for the evolutionary state calculation

$$a = 0.8, b = 0.5$$

The configuration of the parameters was chosen to minimize the number of adjacent nodes having the same color.

The fitness function consists of the count of the violated coloring restrictions. It should be minimized and a perfect graph coloring should have a fitness function equal to 0.

### 5.3 Grid Search Optimization Technique applied on PSO

Despite the reduced complexity compared to other optimization techniques, Grid Search is still one of the most powerful strategies for identifying the optimal solution in case of various optimization problems, but not only.

It is well known that, Grid Search is a hyperparameter tuning technique used in optimization processes in order to find the best set of hyperparameters for a solution designed for a given problem.

Hyperparameters are variables which are set by the user before execution, and hence they are not learned during running process. In this case, different hyperparameters considered are: number of iterations, number of particles and threshold values for cognitive, and social coefficients.

Grid Search works following a simple principle through exhaustive exploration of a predefined grid of possible values for each hyperparameter.

List of hyperparameter values can be found below:

- Number of iterations = [2500, 5000, 7500]
- Number of particles = [80, 100, 200]
- Social and cognitive components  $c1_i$  and  $c1_f$  = [1.5, 1.5], [1.8, 1.8]
- Social and cognitive components  $c2_i$  and  $c2_f$  = [1.5, 1.5], [1.8, 1.8]

The execution and evaluation of the results are done by modeling each combination of parameters in the grid. Finally, using the graphic representation, the optimal or sub-optimal configurations can be easily selected, taking into account the metric that constitutes the objective function, namely, the minimum number of edges of identical colors.

Grid Search is an expensive optimization method from a computational point of view, a fact supported by all measurements performed considering the large number of possible values of the considered parameters. Moreover, the dimensionality of the graph for which the experiments are carried out must also be taken into account.

As already mentioned, the big disadvantage of Grid Search is the very high requirements in terms of hardware resources. Considering the relatively low processing capacities, a medium-sized instance was chosen for the measurements. The obtained results vary depending on the way of combining the values considered in the grid and thus the impact of each parameter in the context of the proposed solution is highlighted.

Although the computational costs of Grid Search are very high, it guarantees the identification of the best combination within the specified grid. Thus, a wise choice for the variation intervals of the parameters could very easily lead to the identification of a sub-optimal or even optimal solution in context of the entire space of parameter variation.

Bellow, the performed experiments will be presented, highlighting the impact of each parameter on the obtained solutions.

## 5.4 Grid Search Optimization - Experiments

### 5.4.1 mulsol.i.3.col.col

Optimal chromatic number: 11

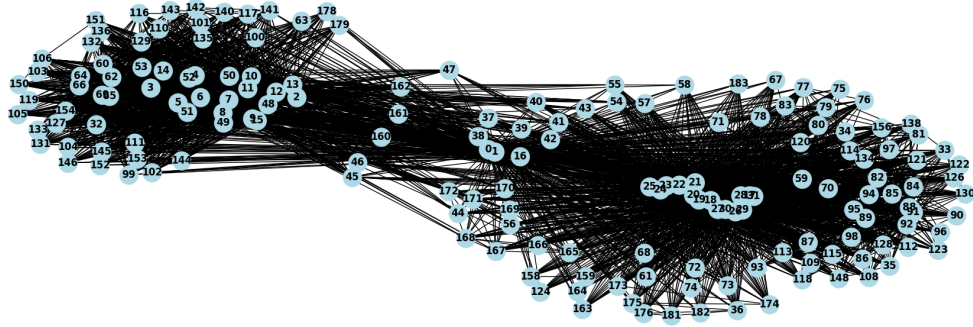


Figure 32: mulsol.i.3.col graph

#### 2500 ITERATIONS

- 1) 80 particles in swarm
- 2) 100 particles in swarm
- 3) 200 particles in swarm

#### 5000 ITERATIONS

- 1) 80 particles in swarm
- 2) 100 particles in swarm
- 3) 200 particles in swarm

#### 7500 ITERATIONS

- 1) 80 particles in swarm

It should be noted that in addition to the number of iterations, the values of the cognitive and social coefficients are particularly important. From the analysis of the graphs, it is easy to see that the increased values of these coefficients lead to a more drastic reduction in the number of collisions from the first iterations. For low values of these parameters, it is observed that the decrease is smoother and thus a greater number of iterations would be necessary to achieve a reasonable result.

The conclusion that remains from the graphs presented is that the number is particles within the swarm does not play a very important role as long as the value is reasonable (e.g. 100). An increase in the the number of particles does not imply a significant improvement of the obtained solution.

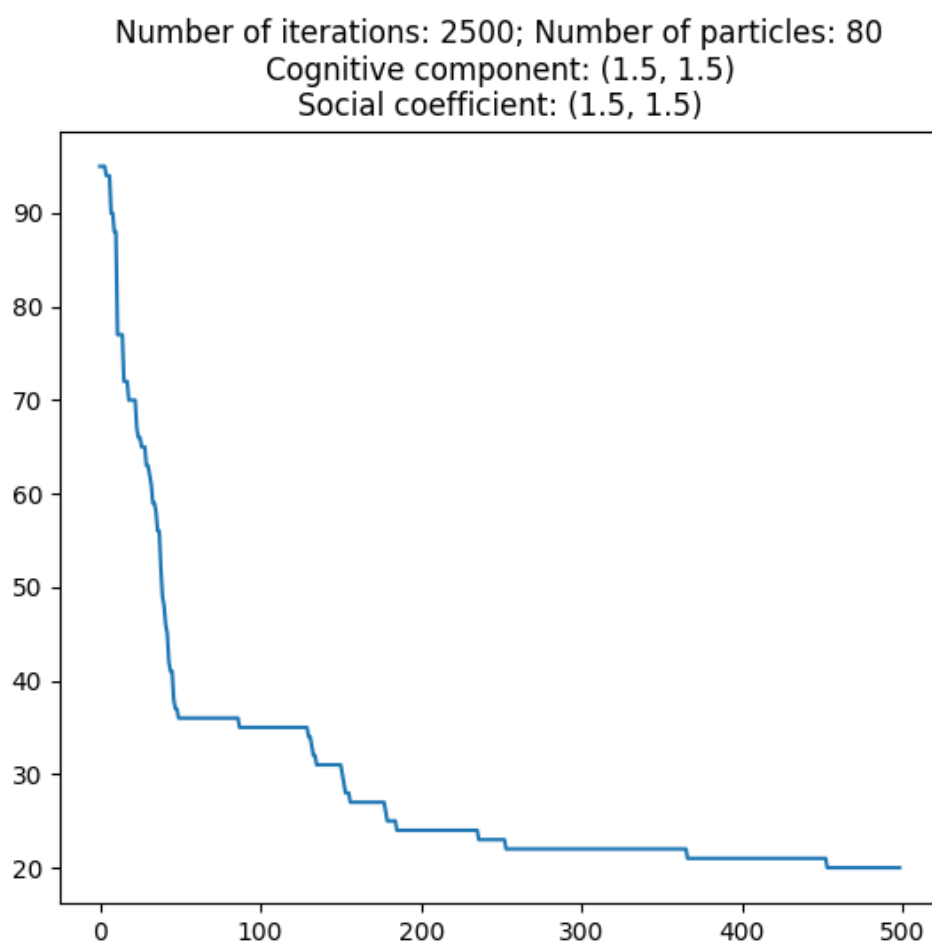


Figure 33: Evolution for mulsol.i.3.col graph

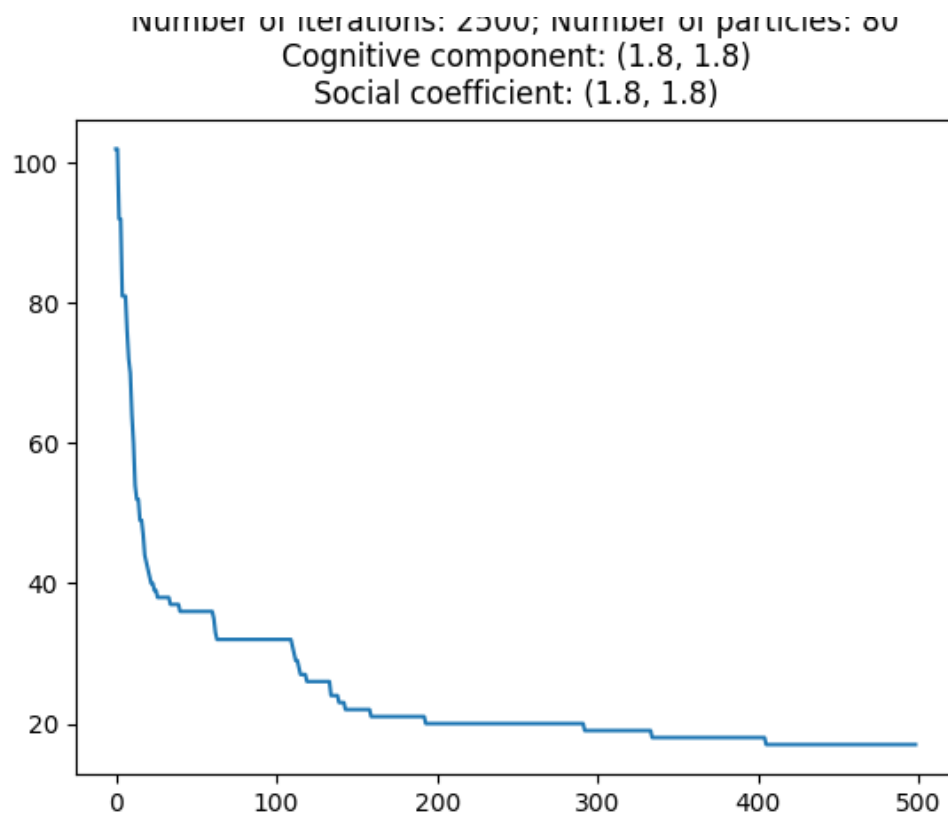


Figure 34: Evolution for mulsol.i.3.col graph



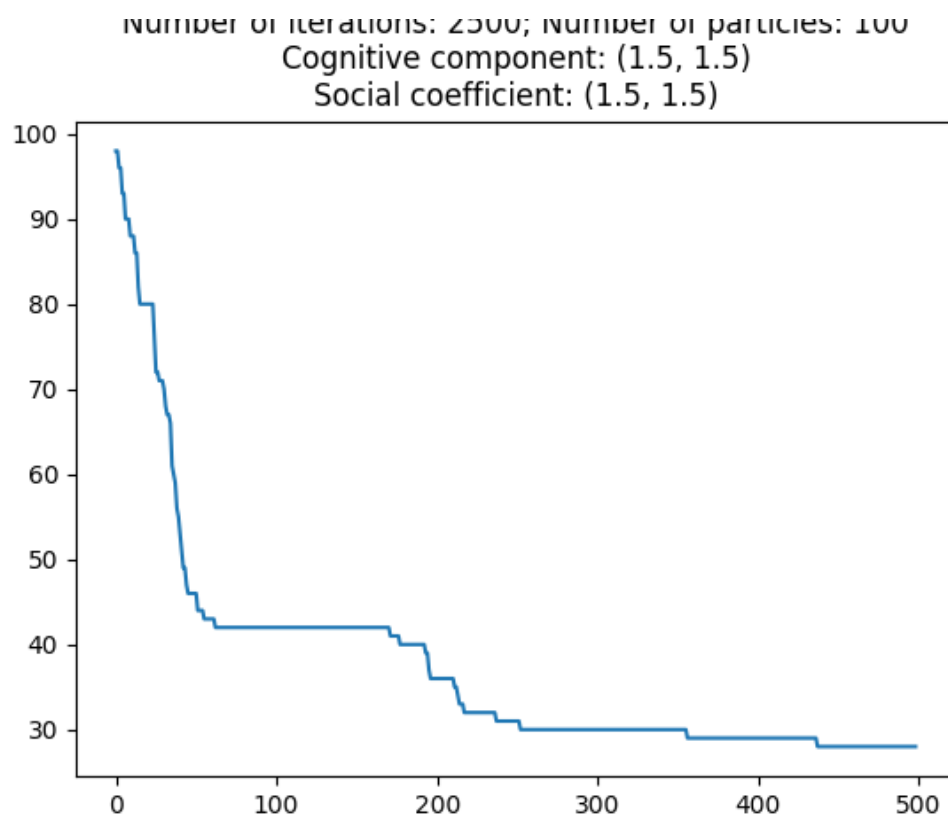


Figure 35: Evolution for mulsol.i.3.col graph

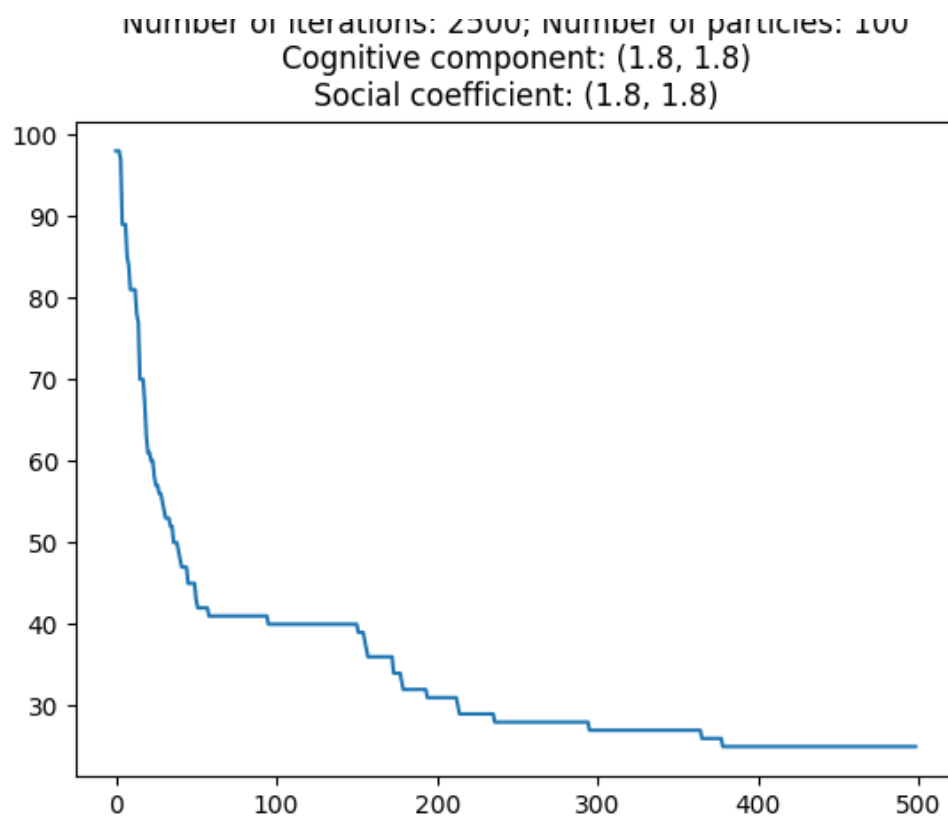


Figure 36: Evolution for mulsol.i.3.col graph

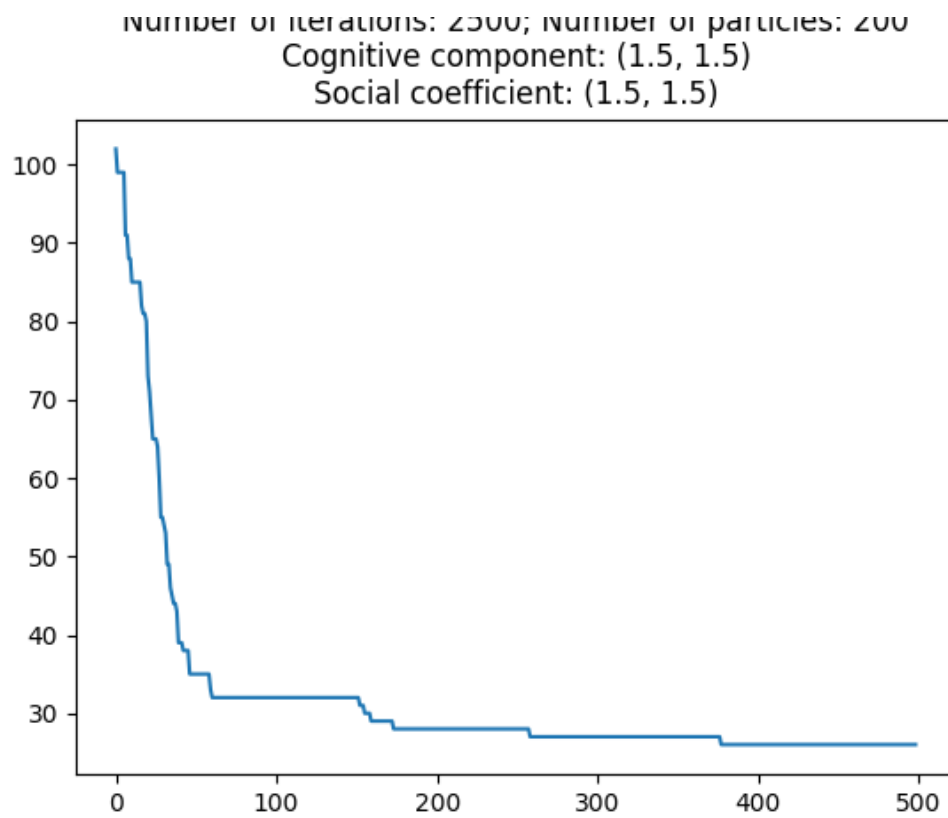


Figure 37: Evolution for mulsol.i.3.col graph

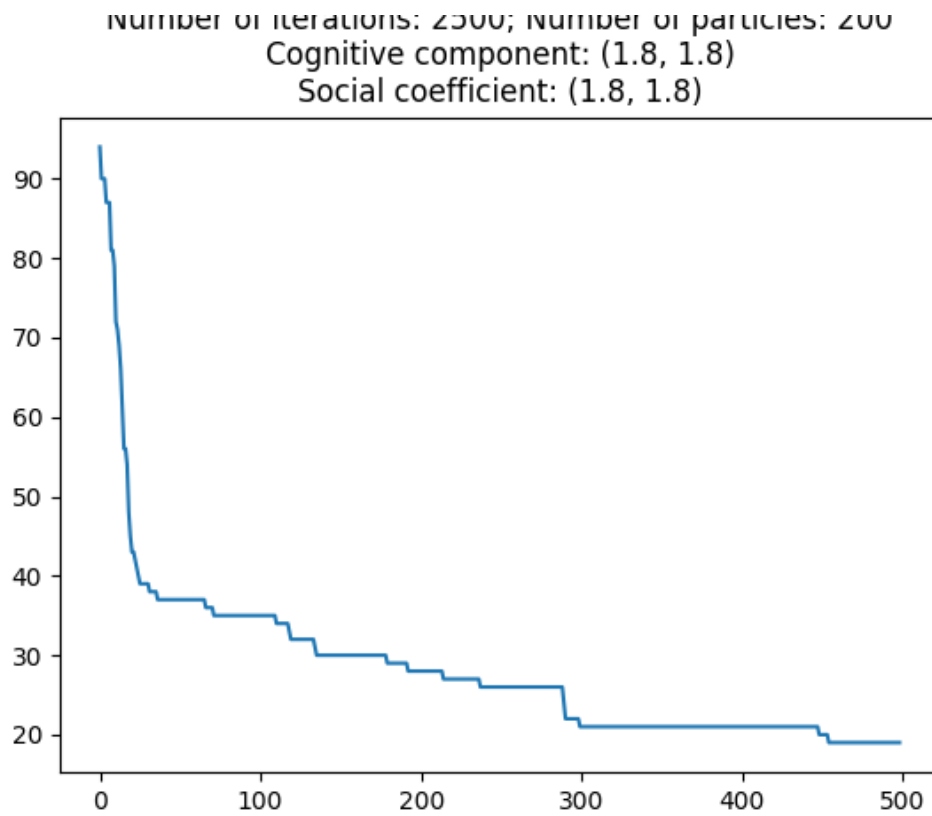


Figure 38: Evolution for mulsol.i.3.col graph

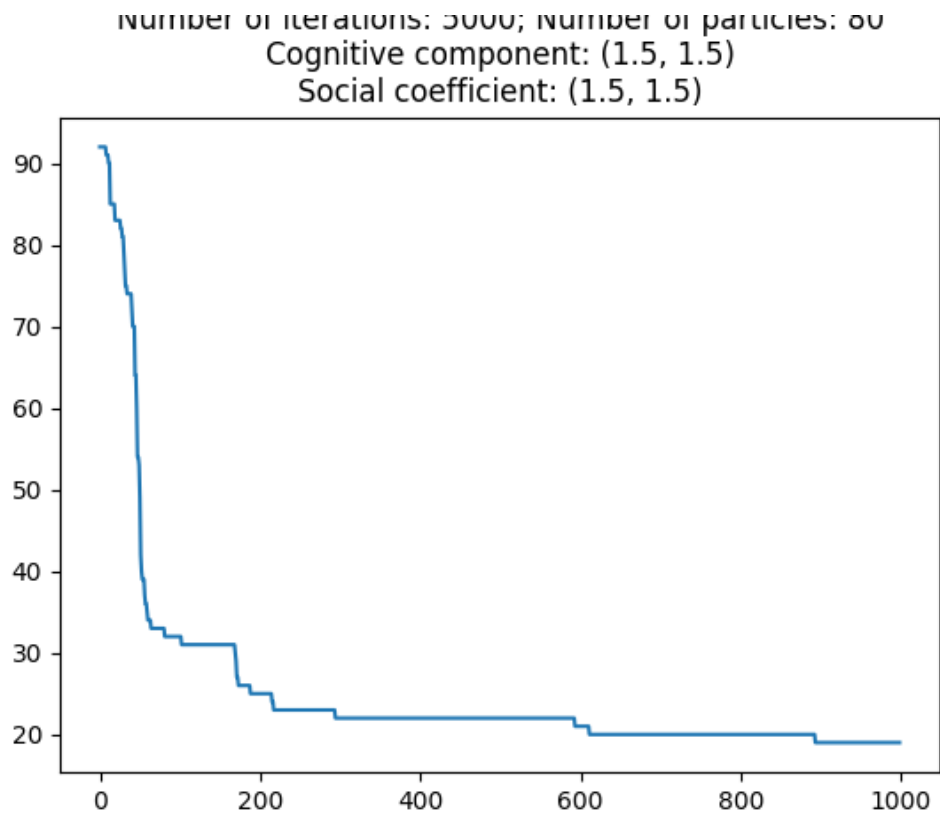


Figure 39: Evolution for mulsol.i.3.col graph

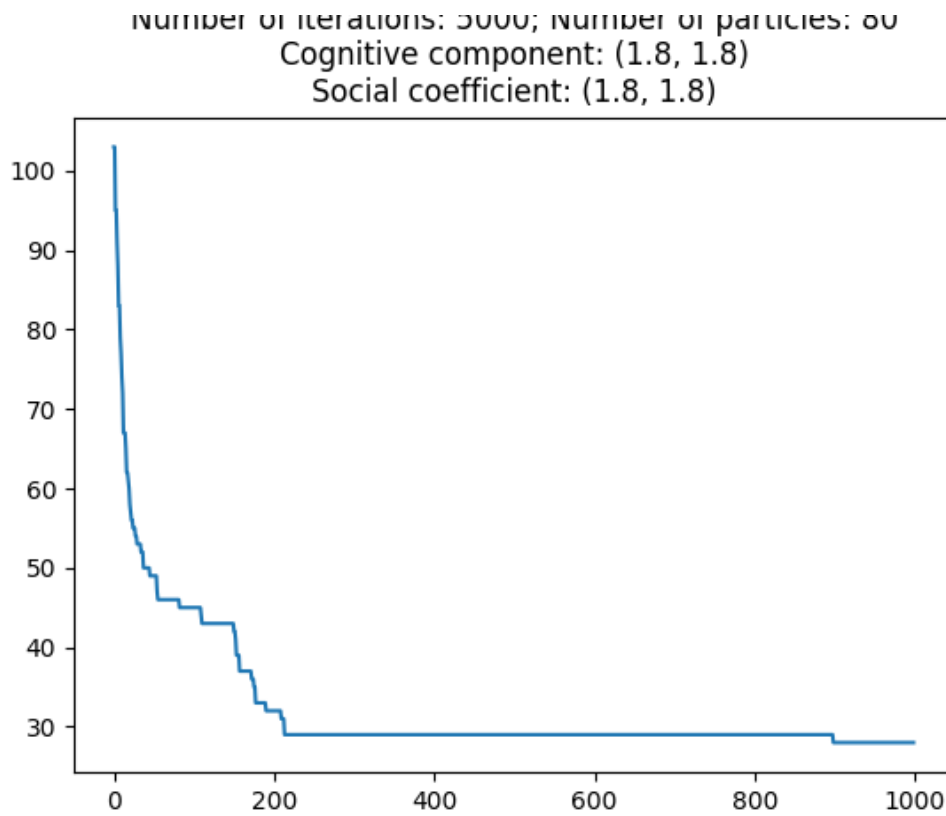


Figure 40: Evolution for mulsol.i.3.col graph

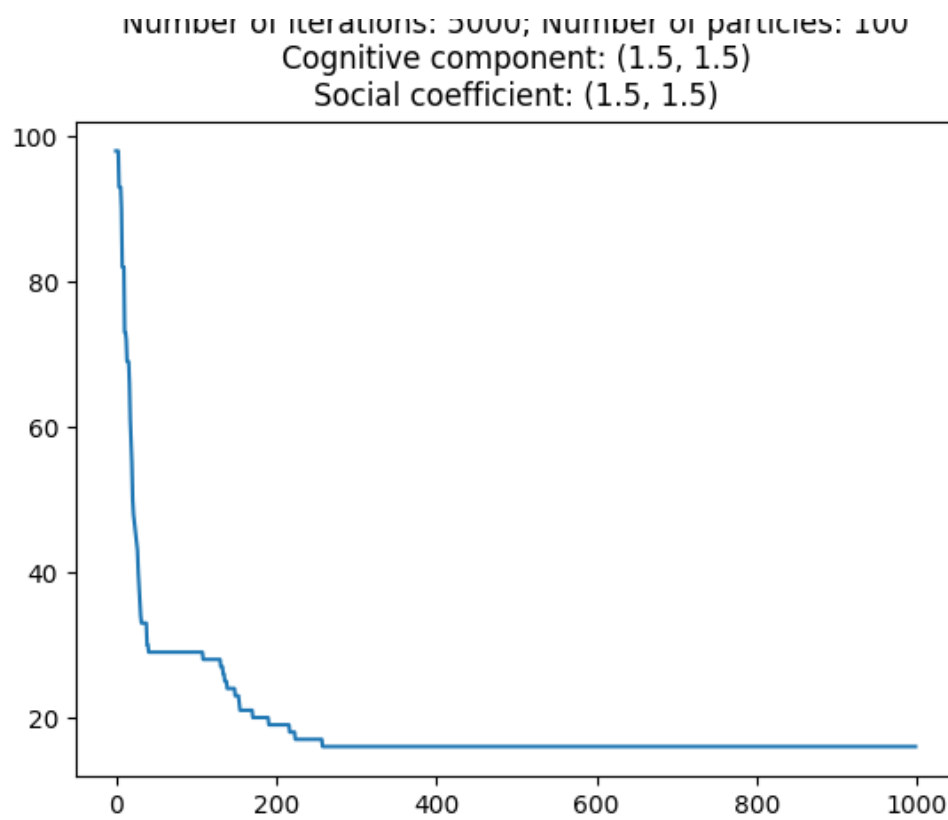


Figure 41: Evolution for mulsol.i.3.col graph

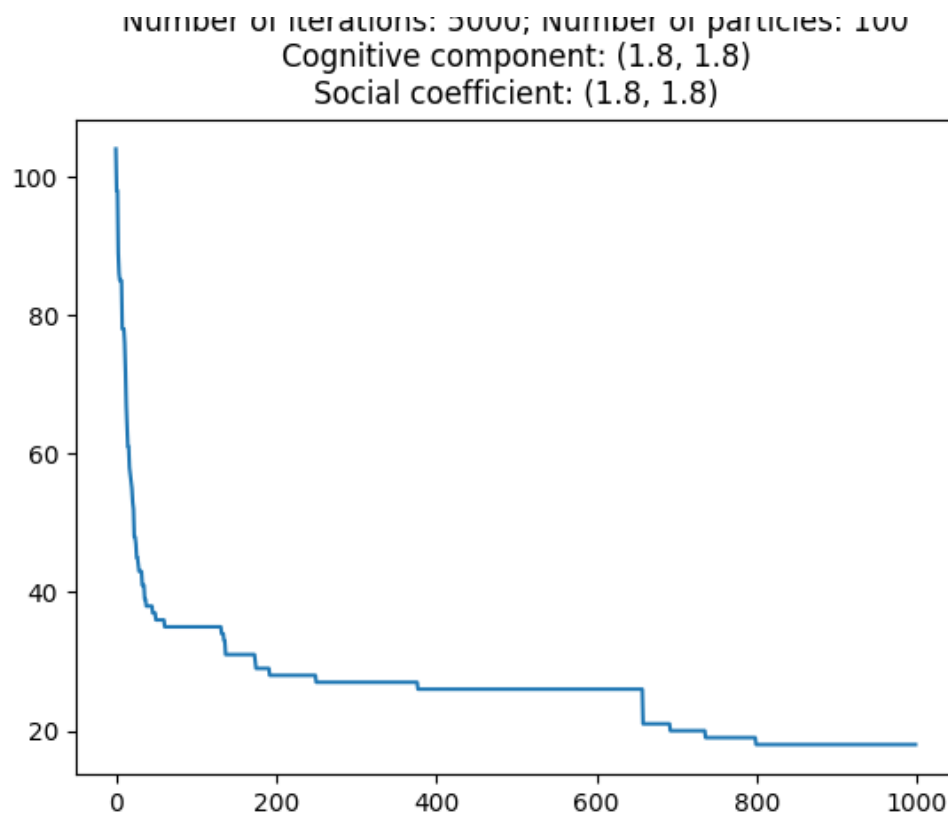


Figure 42: Evolution for mulsol.i.3.col graph



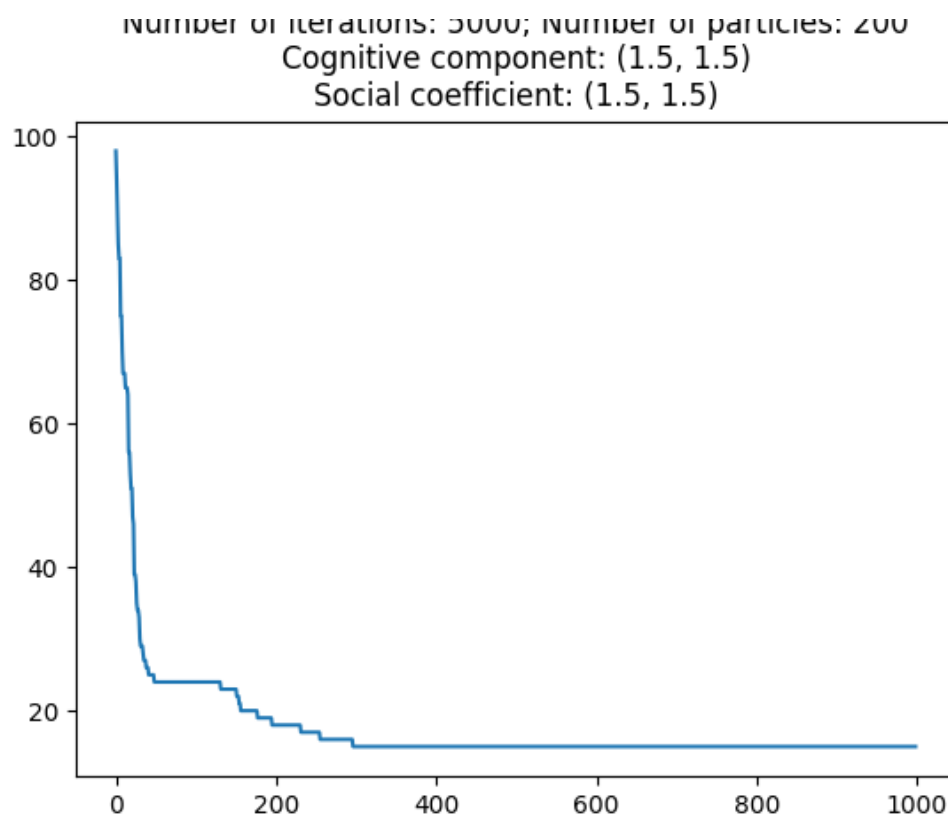


Figure 43: Evolution for mulsol.i.3.col graph

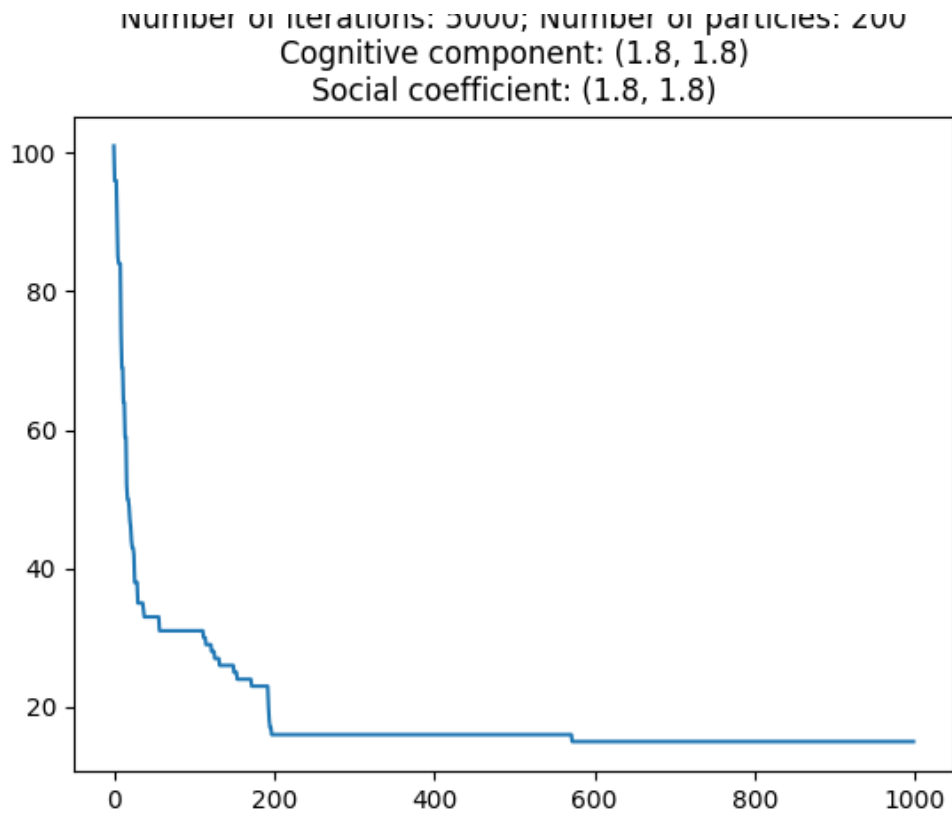


Figure 44: Evolution for mulsol.i.3.col graph

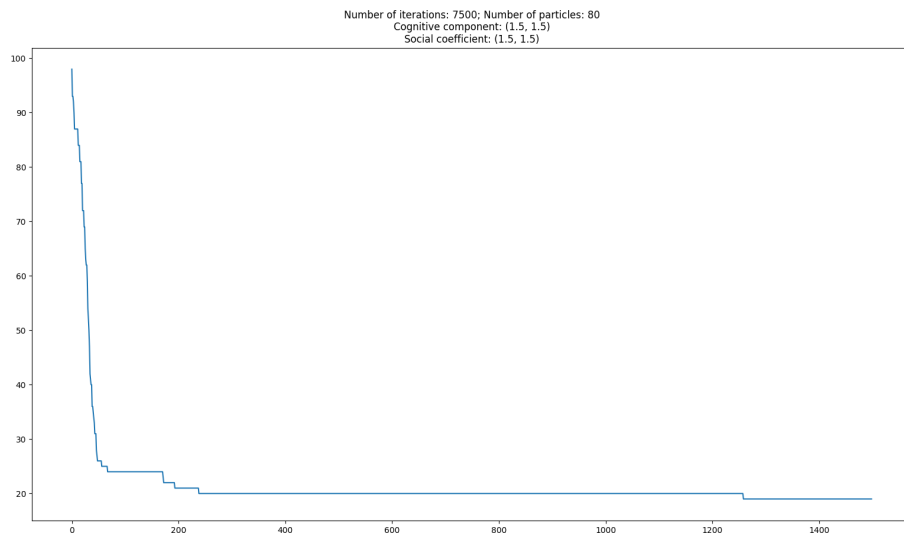


Figure 45: Evolution for mulsol.i.3.col graph

### 5.4.2 fpsol2.i.1.col

Optimal chromatic number: 65

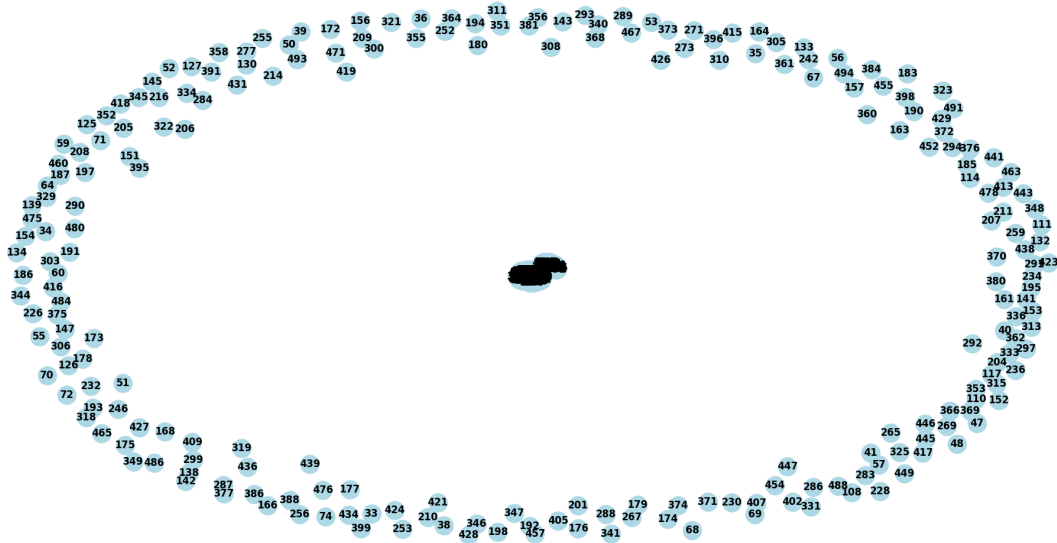


Figure 46: fpsol2.i.1.col graph

This experiment highlights an important contrast. First of all, it was one of the most expensive experiments in terms of time, taking into account that the necessary execution time was several hours.

It is obvious that the number of conflicts decreases with the increase in the number of performed iterations, a conclusion that applies to a large part of the situations in which PSO is involved in order to achieve an optimization process.

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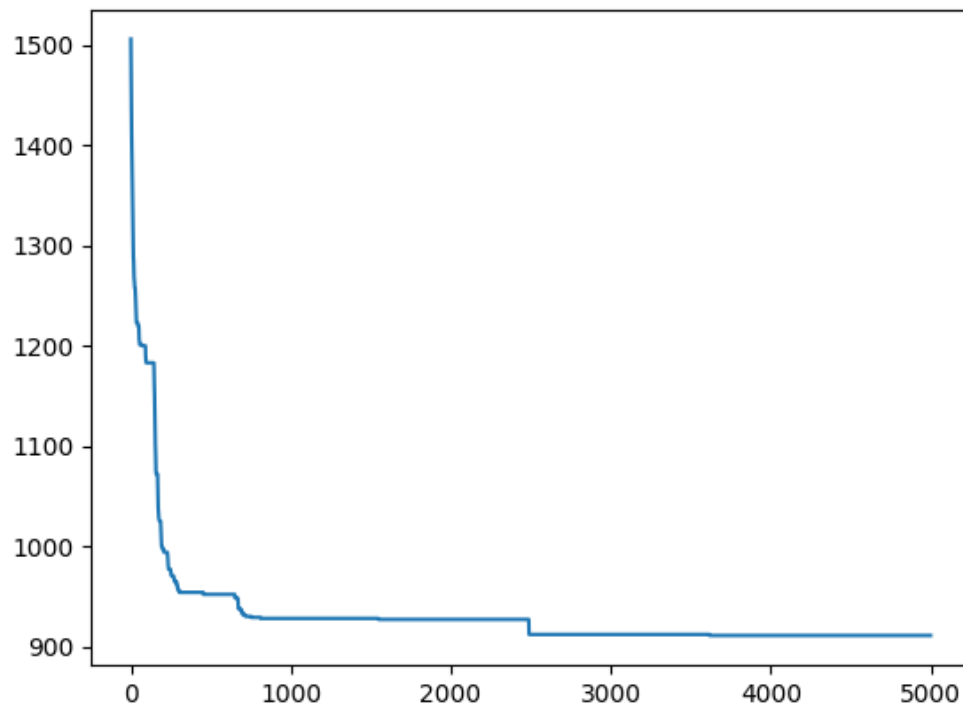


Figure 47: fpsol2.i.1.col result after 5000 iterations

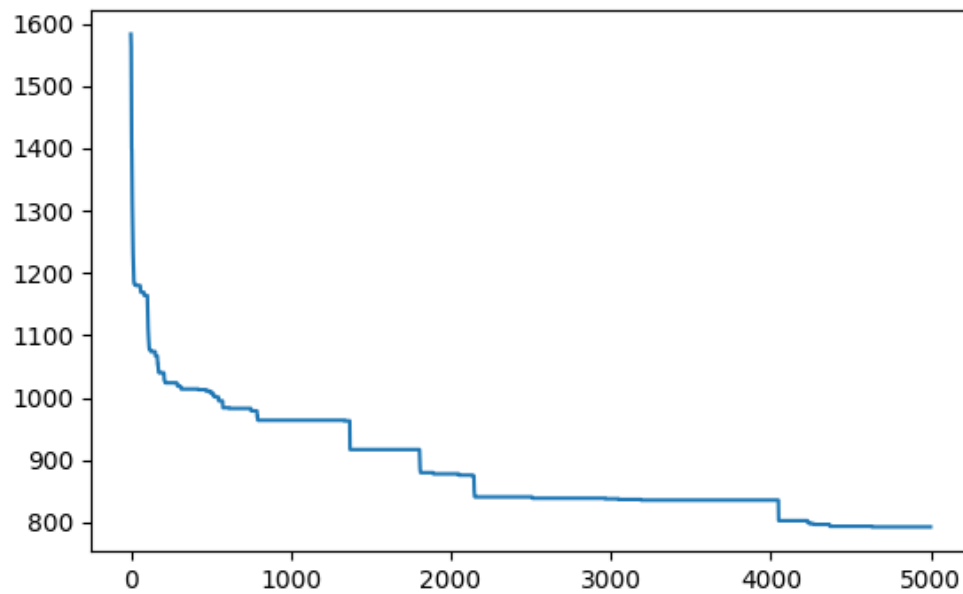


Figure 48: fpsol2.i.1.col result after 25000 iterations

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