

Figure 1: Diagram of the implemented ant colony search algorithm with visualisation of pheromones for one iteration with one ant

2. Update pheromones

For each ant the pheromones connected to visited boards are updated by a fraction of the value of cost function of the whole series of movements.

$$v_{new} = v_{old} + \frac{i}{m} cost \quad (2)$$

where v_{new} is the new value, v_{old} is the old value, i is the index of the movement in this series of movements, m is the length of the series of movements and $cost$ is the value of cost function.

3. Dissipate pheromones

Pheromone for each of the boards that has been visited at least once by any ant in this game is decreased by multiplying the value by a parameter.

$$v_{new} = v_{old} * (1 - dissipation) \quad (3)$$

where v_{new} is the new value, v_{old} is the old value and $dissipation$ is a parameter.

Pheromones can be saved to a file. The file consists of a list of pheromones, each is described with two lines:

1. String representation of a board, left to right, bottom to up, where # means no chessman, upper case letters mean white chessmen and lower case letters mean black chessmen
2. A list of moves. Each move is described by five integer values. First two are the coordinates of chessman to move, third and fourth where to move the chessman to, the fifth is a special value used for promotion (when a pawn becomes another chess piece) and the sixth a real value of pheromone describing its effectiveness.

4.1.2 Experiments

Firstly, it has to be accentuated that chess is a complex game, the number of possible boards that have to be remembered in pheromones is huge and for each ant move

another move has to be done by external Artificial Intelligence. Because of these reasons the learning phase of this metaheuristic takes a very long time. To get results appearing significantly different from completely random ones, hours have to be spent on learning. This makes it very difficult and time-consuming to experiment with parameters properly.

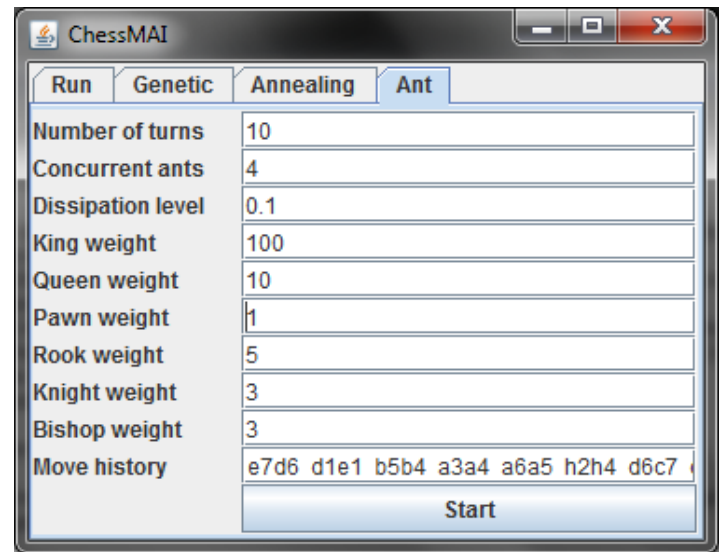


Figure 2: Ant colony dialog boxes

For the experiments the parametrization of many ant search specific variables has been put into dialog boxes in the application(Figure 2). This way user can change these values easily and create his own colonies. The user-available parameters are as follows:

- Number of turns
The number of turns for each iteration. Each iteration consists of an all concurrent ants playing one game of chance up to a win, lose, draw or the artificial end of game, when it takes too long. This parameter should be kept low if we want the learning phase to take less

time and if we want ants to have more broad "knowledge" of possible moves in the beginning of the game.

- Concurrent ants
The number of ants playing a game in each iteration.
The more of them the longer each iteration takes.
- Dissipation level
The dissipation parameter of ant search algorithm dissipation equation (Equation 3).
- Piece weight
Each piece has its own weight used in cost function (Equation ??)
- Move history
This parameter sets the starting point to start the game from any point, provided a valid chess move history. Each ant in each iteration starts its game from this point and plays up until the end of the game (including the end of maximum number of turns defined as another parameter)

The more obvious rules had to be applied to get to the point of metaheuristic being better than a random algorithm and able to win one game out of hundreds when playing against the artificial intelligence. Most importantly the weight of king should be far bigger than other figures, the number of turns small, and a move history provided that gives a possibility of winning in a few turns. Increasing the number of concurrent ants and, at the same time, the dissipation level makes the results possibly even better but by a very small margin at the cost of a longer learning phase (making it difficult to test).

4.1.3 Result

Meddling in all of these parameters proved to be insufficient in obtaining better results. After careful debugging of the application, the conclusion has been made, that the possible reason for improvement could be to increase the probability of choosing good moves instead of dwell on the bad moves. Even though the move choosing equation (Equation 1) gives more probability to good moves, it is not very significant to the sum of all the probabilities, because there are generally a lot of bad moves. One of the ideas was to sum up the logarithms of values in equation, which is often a way of dealing with those kind of issues. This however only made the values of pheromones more close to each other so the probability of choosing a good move didn't really change that much.

The really important change was observed when the Equation 1 was modified to one with a "tolerance" factor multiplied by each value except the best one. This way the probability of choosing the best move in training is big, enabling the colony to thoroughly establish how good of a situation on the board it results in. This can quite easily culminate in a fall into a local maximum of cost function, but it is not a bad situation to be in chess - at least we end up in a better situation than before. Additionally to really fall into a local maximum the moves have to contain a lot of weight gain, so it would be most probably a checkmate anyway, which is, for all intents and purposes, a global maximum. Indeed, if a winning sequence of moves has been established in this version the colony started winning very often.

4.2 Genetic algorithm

4.2.1 what?

4.2.2 gui/experiment

4.2.3 result

4.3 Simulated Annealing

4.3.1 what?

4.3.2 gui/experiment

4.3.3 result

5 Improvements

- put all the variables/modes etc in GUI - adding new engines and randomize their use between moves

6 Conclusion

It was fun / not fun.

References

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