

# Group 2 Project Proposal

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**Abstract:** Group 2 proposes to study the convergence properties of simulated annealing, genetic algorithms, and ant-colony approaches towards finding optimal solutions for various travelling salesman problems. Code will be tested to find convergence rate, wall-clock time, and for solution space exploration. For future studies, we will consider looking at parallel and/or GPU solutions to accelerate algorithms.

## 0.1 Introduction

## 0.2 Simulated Annealing

Simulated annealing is a stochastic approach towards minimizing an energy function.

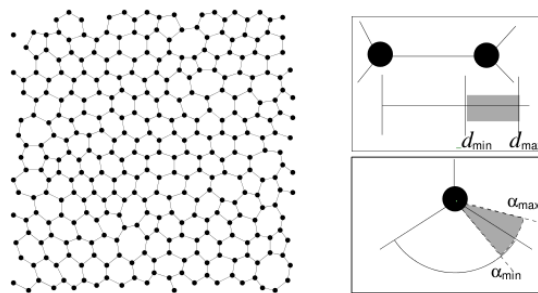
## 0.3 Genetic Algorithms

Unlike simulated annealing, genetic algorithms rely on a population approach to

## 0.4 Ant-Colony Approaches

## 0.5 Go with the Winner Approaches

Simulated annealing probabilistically converges to the optimal solution at the rate of  $O(\frac{1}{n})$  where  $n$  is the number of independent runs. Using heuristic schemes such as the “Go with the Winner” [1], hereby referred to as GWW, one can improve this convergence



**Fig. 1.** left: an example of an amorphous structure with vertex degree 3 in two dimensions; right: the bond length and bond angle constraints

rate to  $O(\log \frac{1}{n})$  and have recently found their way into molecular modeling applications [2]. In general terms, the GWW algorithm mirrors that of simulated annealing; however, each independent run is periodically reassessed to determine if one is converging upon a losing solution. Runs with no chance for success are immediately discarded while runs with the best chance of finding an optimal solution are replicated; thus, GWW generates an ensemble of independent runs filled with “winners”.

## 0.6 Analysis

To assess the performance of our optimization schemes, we would

## References

- [1] D. Aldous and Vazirani U. Go with the winners algorithms. *Proceedings of the 35th IEEE Symposium on Foundations of Computer Science*, pages 492–501, 1994.
- [2] Marcus Peinado and Thomas Lengauer. ‘go with the winners’ generators with applications to molecular modeling. *RANDOM*, pages 135–149, 1997.