# **FIGURE College of Engineering**

#### **INTRODUCTION**

Many mathematical and practical problems require elements of a set to be arranged in a way that satisfies specific conditions. Familiar examples include Sudoku, timetabling, scheduling, and resource distribution (Russell & Norvig, 2009). Such problems, called constraint-satisfaction problems (CSPs), are solved when values from a domain are assigned to variables in a way that does not violate a set of constraints. Because some CSPs are NP-complete (Bulatov, Krokhin, & Jeavons, 2000), no polynomial-time algorithm to solve the general constraint-satisfaction problem is known. In order to reduce the time required to solve large instances, an Ant Colony Optimization (ACO) algorithm called the MAX - MIN Ant System (*MMAS*) (Stützle & Hoos, 2000) was applied to combinatorial CSPs with focus on a specific CSP (the Costas-array problem, which remains open for most sizes  $M \ge 30$ ).

#### BACKGROUND

**Definition 1.** A *constraint-satisfaction problem* (CSP) (Bulatov et al., 2000) is an ordered triple  $\langle V, D, C \rangle$ , where

- *V* is the set of *variables*
- *D* is the *domain*
- *C* is the set of *constraints*

Each constraint  $C_i$  is of the form  $\langle s_i, \rho_i \rangle$ , where  $s_i$  describes a tuple of variables  $s_i : \{1, \dots, m_i\} \rightarrow V$  and  $\rho_i$  is an  $m_i$ -ary Boolean relation on  $D, \rho_i \subseteq D^{m_i}$ 

A *solution* of the constraint-satisfaction problem  $\langle V, D, C \rangle$  is a function  $f: V \to D$  such that for all constraints  $C_i = \langle s_i, \rho_i \rangle$ ,  $f \circ s_i \in \rho_i$ 

**Definition 2.** A *Costas array* (Drakakis, 2011) of size *M* is a permutation of the integers from 1 through M such that its permutation matrix contains no equal displacement vectors between distinct pairs of distinct elements.

4 2 1 5 3 6	$1 \ 2 \ 3 \ 4 \ 6 \ 5$	$3\ 2\ 4\ 6\ 1\ 5$	$2 \ 5 \ 3 \ 4 \ 1 \ 6$
0 0 1 0 0 0	1, 0, 0, 0, 0, 0	0 0 0 0 <b>1</b> 0	0 0 0 0 <b>1</b> 0
0 1 0 0 0 0	0 $1$ $0$ $0$ $0$ $0$	0 1 0 0 0 0	$1 \ 0 \ 0 \ 0 \ 0 \ 0$
$0/0 \ 0 \ 0 \ 1 \ 0$	0  0 <b>1</b> 0  0  0	$1  0 \setminus 0  0  0  0$	0 0 1 0 0 0
$1_0 0 0 0 0$	0  0  0 <b>1</b> 0  0	0 $0$ $1$ $0$ $0$ $0$	0 0 0 <b>1</b> 0 0
$0 \ 0 \ 0 \ 1 \ 0 \ 0$	$0 \ 0 \ 0 \ 0 \ 0 \ 1$	$0 \ 0 \ 0 \ 0 \ 1$	0 1 0 0 0 0
0 0 0 0 0 1	$0 \ 0 \ 0 \ 0 \ 1 \ 0$	$0  0  0  \mathbf{\hat{1}}  0  0$	0 0 0 0 0 1
A	В	C	D

Figure 1: Three permutations (*A*, *B*, and *C*) that violate the Costas-array property and one (*D*) that satisfies it.

The constraint-satisfaction problem of searching for Costas arrays of size *M* can be formally defined as

 $\langle \{x_1,\ldots,x_M\},\{1,\ldots,M\},$  $\{ \langle i \mapsto x_i, \{t : \forall i, j (t(i) = t(j) \to i = j) \} \rangle,$  $\langle i \mapsto x_i, \{t : \forall i, j, k, l \ (i \neq j \land i \neq k \to j - i \neq l - k \lor t(j) - t(i) \neq t(l) - t(k))\} \rangle$ 

For  $M \ge 30$ , it is not in general known whether a solution to the Costas-array problem exists; the problem for a given size *M* is solved if a solution is found.

**Definition 3.** If a parallel program with *N* workers solves a problem in *T* time and the same program with one worker solves the problem in T<sub>0</sub> time, the *speedup* S and *efficiency* E of the program for that problem are

S = -

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$$E = \frac{S}{N} = \frac{T_0}{N \cdot T}$$

## Parallel MAX - MIN Ant System for combinatorial constraint-satisfaction problems

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## **DESIGN GOAL AND METHODS**

Because general CSP solvers run in exponential time, the runtime of programs solving combinatorial CSPs grows rapidly. The goal of the project was to create a parallelized and heuristic-accelerated MMAS framework that:

- Distributes work to local processors to solve the Costas-array problem
- Maintains high efficiency (E > 0.5) at high numbers of workers (up to N = 250)
- Exhibits lower average time-to-first solution with the new optimizations than without.

A detailed description of the research process can be found in the accompanying journal.

called pheromones (Fig. 2).



Figure 2: A diagram of ant-foraging behavior. Initially, ants follow random paths. Over time, better paths acquire higher concentrations of pheromone, and all ants are drawn to a single optimal path. Image taken from http://mutenet.sourceforge.net/howAnts.shtml

ACO algorithms (Dorigo & Di Caro, 1999) are based on the behavior of foraging ants, which indirectly communicate with each other by secreting and detecting chemicals

- In order to apply ACO to an optimization problem, the problem is represented as a graph connecting domain values called the *construction graph*.
- Processes called ants stochastically traverse this graph, altering the properties that govern ant behavior.

 $\mathcal{M}\mathcal{M}$  AS was applied to the problem of Costas arrays as follows:

- Paths through the construction graph represent permutations.
- The *cost* of a path is the number of Costas-array-property violations.
- The *heuristic value*  $\eta_{si}$  associated with adding *i* to permutation *s* is inversely proportional to the cost incurred.
- The *pheromone value*  $\tau_{si}$  is based on the learned favorability of following *s* with *i*.
- The probability that an ant will add *i* to permutation *s* is based on a function of  $\eta_{si}$  and  $\tau_{si}$ , the *ant-routing table*  $A_{si}$ .



Figure 4: Path representation of permutations for pheromone association. (A) Pheromone is applied to all five unique subsequences of the permutation (1,3,2,4); clockwise from top left (+1,-2), (+1), (-2), (-2,+1,-2), (-2,+1). (B) All suffixes of the permutation (4, 2, 1, 3) are considered when computing the ant-routing table  $A_{s3}$  at s = (4, 2, 1); from left to right (+2, +1, -2), (+1, -2), (-2).

• The queen applies pheromone to all subsequences of the best paths found (Fig. 4A).

• The amount of pheromone applied to a path's subsequences is inversely proportional to that path's cost.

The construction graph supports parallel querying of pheromone values to compute the ant-routing table  $\mathcal{A}$ .

• As a new optimization, the construction graph uses the longest suffix encountered when determining pheromone values (Fig. 4B).

used:

The design process started with the implementation of a singlethreaded *MMAS* solver for Costas arrays. Over the course of several months, this solver was parallelized and new optimizations, such as map-based pheromone storage, were developed. Two languages were

• C++ for the *MMAS* framework

• Java for the visualizer program

No libraries outside the C++17 STL and Java standard were used.

The ant system was tested on a machine with 256 identical Xeon Phi Knight's Landing processors.

A discussion of the techniques used to apply MMAS to the Costasarray problem and the new optimizations added follows.

### **FRAMEWORK DESIGN**



Figure 3: A high-level schematic of the parallel *MMAS* framework. Purple arrows indicate the flow of information through the system; green ones indicate the flow of control.

The *MMAS* framework includes two types of processes that run in parallel and indirectly communicate with each other through two shared data structures (Fig. 3).

Some number *N* of parallel worker ants repeatedly traverse the construction graph, constructing paths that represent possible solutions. • Worker ants construct paths stochastically, favoring paths with high heuristic and pheromone values.

• The best (lowest-cost) paths are inserted into a shared list.

• As a new optimization, ants constructing a path the cost of which exceeds the cost of the best path found by a *quality threshold*  $\vartheta$ abandon the path and immediately start a new one.

A single queen process periodically updates pheromone values.

• The queen operates based on ant time periods called epochs; over one epoch, each ant constructs a set number of paths.

Simplified visualizations of the evolution of the pheromone graph over the course of  $\mathcal{MMAS}$  Costas-array searches show that the final solution found by the ants utilizes high-pheromone components of non-Costas array permutations (Fig. 5).



Figure 5: Visualizations of the 2D pheromone graph for an  $M = 14 \mathcal{MMAS}$  Costas-array search. From left to right, the state of the graph at time t = 5, t = 15, t = 30, t = 45, t = 60, t = 75, and t = 90 (in ant epochs). The color of a cell in the *i*<sup>th</sup> row and *j*<sup>th</sup> column represents the learned favorability of following *i* with *j* in a Costas-array candidate, with darker cells indicating higher pheromone concentration and lighter cells representing lower pheromone concentration.



Figure 6: Graphs of  $\mathcal{MMAS}$  performance. (A) A graph of median E (as a ratio of times-to-firstsolution) with respect to N and M (n = 100). Error bars represent the first and third quartiles. (B) A graph of decrease in average time-to-first-solution after new optimizations were added.

• For  $N \ge 200$ ,  $E \approx 0.6$  for all  $M \ge 14$ , implying that an efficiency of ~0.6 is likely to extend for greater worker counts and problem sizes.

Fig. 6B indicates that the new optimizations generally improve  $\mathcal{M}\mathcal{M}AS$  time-to-first-solution.

• New optimizations are most effective at larger problem sizes and processor counts; for N = 250 and  $M \ge 14$ , the new optimizations cause a  $\sim 30\%$  runtime decrease.

The performance data collected indicate that the *MMAS* framework developed satisfies the project goals:

• *MM* AS maintains an approximately constant high efficiency across several larger processor counts and problem sizes, indicating that *E* is likely to remain above 0.5 even for larger *M* and N.

• The optimizations decrease average *MMAS* time-to-first solution by a significant margin (~30%).

• Application of distributed computing techniques to pool the computational power of multiple machines without shared memory.

• Application of *MMAS* to unsolved problem instances (namely, the search for M = 32 Costas arrays).



#### VISUALIZATIONS

Fig. 6A indicates that, for  $M \ge 14$ ,  $E \ge 0.5$  for all values of N tested.

• For *N* < 200, *E* varies with *M* and decreases as *N* increases.

• For N > 1,  $\mathcal{MMAS}$  with the new optimizations is consistently faster than  $\mathcal{MMAS}$  without (exception: N = 150, M = 13).

#### DISCUSSION

Future goals for the *MMAS* program include:

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