

Learning from Experience: Case Injected Genetic Algorithm Design of Combinational Logic Circuits

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Abstract. This paper presents a new approach to genetic algorithm based design. We use genetic algorithms augmented with a case-based memory of past design problem solving attempts to obtain better performance over time on sets of similar design problems. Rather than starting anew on each design, we periodically inject a genetic algorithm's population with *appropriate intermediate design solutions* to similar, previously solved problems. Experimental results on a configuration design problem; the design of a parity checker circuit, demonstrate the performance gains from our approach and show that our system learns to take less time to provide quality solutions to a new design problem as it gains experience from solving other similar design problems.

1 Introduction

Design problems seldom exist in isolation. Any useful design system must expect to confront many related problems over its lifetime and we would like such a system to improve its performance with experience. Such a learning system requires memory; a place for storing past experiences to guide future operations. The storage area may be distributed or localized, but a system without a memory is forced to start from scratch in trying to solve every given problem. Genetic algorithms (GAs) are randomized parallel search algorithms that search from a population of points [11,8]. Current genetic algorithm based machine learning systems use rules to store past experience to improve their performance over time [11,8] and [28,13,9]. However, many application areas, especially in the design domain, are more suited to a case-based storage of past experience [22,12] and [29,7].¹ We propose and describe a system that uses a case-base as a long term knowledge store in a new genetic algorithm based design system that learns from experience. Results from the design of parity checkers, applicable to problems from engineering design

¹ See also the web page at <http://www.ai-cbr.org/projects.html> for a list of case-based design projects.

and architecture, show 1) that our system, with experience, takes less time to solve new problems and produces better quality solutions and 2) a simple syntactic similarity metric can result in this performance improvement thus avoiding the indexing problem endemic to purely case-based systems.

These performance gains imply that fewer evaluations are needed to reach a specified design solution quality and that the organization deploying this system builds a knowledge base of cases.

Typically, a genetic algorithm randomly initializes its starting population so that the GA can proceed from an unbiased sample of the search space. However, as pointed out earlier, problems do not usually exist in isolation and we often confront sets of similar problems. It makes little sense to start a problem solving search attempt from scratch when previous search attempts may have yielded useful information about the search space. Instead, periodically injecting a genetic algorithm's population with *relevant* (we describe what we mean by relevant later) solutions or partial solutions to similar previously solved problems can provide information (a search bias) that reduces the time taken to find a quality solution. Our approach borrows ideas from case-based reasoning (CBR) in which old problem and solution information, stored as cases in a case-base, helps solve a new problem [25]. In our system, the data-base, or case-base, of problems and their solutions supplies the genetic problem solver with a long term memory. The system does not require a case-base to start with and can bootstrap itself by learning new cases from the genetic algorithm's attempts at solving a problem.

The case-base does what it is best at – memory organization; the genetic algorithm handles what it is best at – adaptation. The resulting combination takes advantage of both paradigms; the genetic algorithm component delivers robustness and adaptive learning while the case-based component speeds up the system.

The **C**ase **I**njected **G**enetic **A**lgo**R**ithm (CIGAR) system presented in this paper can be applied in a number of domains from computational science and engineering to operations research. We concentrate on configuration design problems, specifically the design of combinational logic circuits.

We define the CIGAR system in the next section. We then discuss related work and delineate the difference between problem and solution similarity and apply our system to two problems that highlight fundamental issues addressed by this work. The last section presents conclusions and directions for future work.

2 Case Injected Genetic Algorithms

How do we combine a genetic algorithm with case-based memory? Our first approach worked as follows. When confronted with a (new) problem, the CBR module looks in its case base for *similar problems* and their associated solutions. Note that case-based reasoning research has shown that defining

a problem similarity metric is non-trivial [16]. If the system finds any similar problems, a small number of their solutions get injected into the *initial* population of the genetic algorithm. The rest of the population is initialized randomly (to maintain diversity) and the GA searches from this combined population. This works well if we have a good measure of problem similarity.

However, if we assume that similar solutions must have come from similar problems, CIGAR can operate on the basis of solution similarity and avoid the issue of defining problem similarity metrics. In this scenario, shown in Figure 1, we **periodically inject** a small number of solutions *similar to the current best member (candidate solution)* of the GA population into the *current* population, replacing the worst members. We call this the “closest to the best” strategy, one of several possible injection strategies. The GA continues searching with this combined population.

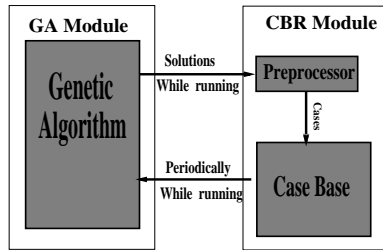


Fig. 1. Conceptual view of CIGAR.

What are cases and how do we save them to the case-base? During GA search, whenever the fitness of the best individual in the population increases, the new best individual is stored in the case-base. A case is a member of the population (a candidate solution) together with ancillary information including fitness, the generation this case was generated, and its parents [20]. Reusing old solutions has been a traditional performance improvement procedure. Our work differs in that 1) we attack a set of tasks, 2) store and reuse **intermediate** candidate solutions, and 3) do not depend on the existence of a problem similarity metric avoiding indexing problems common to case-based systems.

What happens if our similarity measure is noisy and/or leads to unsuitable retrieved solutions? By definition, unsuitable solutions will have low fitness and will quickly be eliminated from the GA’s population. CIGAR may suffer from a slight performance hit but will not break or fail – the genetic search component will continue making progress towards a solution. CIGAR is robust. As noted earlier, we can choose schemes other than injecting the closest to the best; furthest from the worst and probabilistic versions of both have proven effective.

3 Related Work

One early attempt at reuse can be found in Ackley’s work with SIGH [1]. Ackley periodically restarts a search in an attempt to avoid local optima and increase the quality of solutions. Eshelman’s CHC algorithm, a genetic algorithm with elitist selection and cataclysmic mutation, also restarts search when the population diversity drops below a threshold [6]. Other related work includes Koza’s automatically defined functions [15] and Schoenauer’s constraint satisfaction method [26]. More recently, Ramsey and Grefenstette come closest to our approach and use previously stored solutions to initialize a genetic algorithm’s initial population and thus increase a genetic algorithm’s performance in an anytime learning environment that changes with time [24,10]. Automatic injection of the best solutions to previously encountered problems biases the search toward relevant areas of the search space and results in the reported consistent performance improvements. Sheppard and Salzburg combined memory-based learning with genetic algorithms in finding better plans for the pursuer-evader game within a reinforcement learning framework [27]. Their experiments indicate that the combined approach performed better than either approach alone. To our knowledge, the earliest work in combining genetic algorithms and case-based reasoning was done by Louis, McGraw, and Wyckoff who used case-based reasoning principles to explain solutions found by genetic algorithm search [20]. Preliminary work in this area solved pairs of similar problems with a clear performance advantage for the combined system [17]. Work that uses case-based reasoning in extracting design patterns, short term memory in genetic programming, Robert Reynold’s cultural algorithms, and circuit design that may be relevant are [23,14] and [2,30].

These approaches only attack a single problem not a related *class* of problems. Moreover, these approaches do not relate problem/solution similarity to the quality of injected solutions and performance – a fundamental result of this paper.

Work on multitask learning suggests, as we do, that it is easier to learn many tasks at once, rather than to learn these same tasks separately [4,5] and [3]. Multitask learning can be applied to clusters of related tasks in parallel or in sequence and provides an inductive bias that often leads to better generalization performance on the tasks. While MTL addresses generalization, we lean towards improving search performance – in design domains, a mapping from function to structure – for sequential (design) tasks. Parallel genetic algorithms using the island model, provide one possible avenue towards information exchange on related tasks in parallel for genetic based machine learning systems.

Lifelong learning seeks to address the problem of knowledge transfer between related tasks in the context of learning control algorithms for robotics [31,32]. The authors believe that knowledge transfer is essential for scaling robot learning algorithms to more realistic and complex domains. Lifelong learning

differs from our approach in that they are interested in an incremental learning situation seeking to build behavioral complexity with experience. Work by Louis provides strong support for the CIGAR approach to control problems in robotics thus indirectly and independently providing more evidence for the utility of lifelong learning [19].

What do we mean by problem similarity? What is the difference between problem similarity and solution similarity? The next section provides concrete answers to these questions in the context of combinational logic design.

4 Indexing and similarity

Indexing, or how we define similarity, is a basic issue in all the systems described above. Previous work has dealt with the simpler case where a similarity metric exists in the problem space [18,24]. Simply put, when we know that two *problems* are similar, the system can use information from attempting one problem to improve performance on the other. However, a problem similarity metric is not easy to come by and remains a major issue for case-based reasoning systems. In this paper, we study the more realistic case where we do *not* need a similarity measure on the problem space. Since genetic algorithms solutions are encoded as binary or real strings, purely syntactic similarity measures lead to surprisingly good performance – a needed property for applications to poorly-understood systems. Solution similarity measures completely avoid CBR’s indexing problems.

4.1 Problem Similarity

Combinational circuit design is an example of a configuration design problem. The general problem can be stated as: Given a function and a target technology to work within, design an artifact that performs the function subject to constraints. For parity checkers, the function is parity checking and the target technology is combinational logic gates such as boolean AND and OR gates. A genetic algorithm can be used to design combinational circuits as described in [21]. A five-bit parity checker is one of $2^{2^5} = 2^{32} = 4294967296$ different five-input one-output combinational circuits (boolean functions). If we are trying to solve this class of problems, one way of indexing, defining similarity of problems, can be: Concatenate the output bit for all possible 5-bit input combinations counting from 0 through 31 in binary. This results in a binary string of output bits, S_o , of length 32. Strings that are one bit different from S_o define a set of boolean functions, as do strings that are two bits different and so on. This way of naming boolean functions provides a simple distance metric and indexing mechanism for the combinational circuit design problems that we consider in this paper. That is, we have a metric for measuring *problem similarity*. Problems are constructed from the parity problem by randomly changing bits in the output bit string. The fitness of a

candidate circuit is the number of correct output bits. Thus 5-bit problems would have a maximum fitness of $2^5 = 32$. As a simple example, consider the 3-bit parity checker as shown in Table 4.1. The first column in Table 4.1 shows all possible inputs for a 3-bit parity checker, the second column shows the correct output for each input. We construct a 3-input, 1-output problem that is similar to the 3-bit parity checker by randomly choosing and flipping one of the output bits of the 3-bit parity checker as shown by the boxed 0 in the table. The correct output bits for the new problem (1 bit different from the 3-bit parity checker) are shown in the third column. This new problem is distance 1 away from the parity problem in terms of our problem similarity metric.

Table 1. Outputs of the 3-bit parity checker (a 3-0) and a 3-1 problem.

Inputs	3-bit parity problem	3-1 problem
000	0	0
001	1	0
010	1	1
011	0	0
100	1	1
101	0	0
110	0	0
111	1	1

4.2 Representation

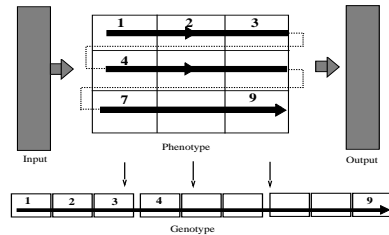


Fig. 2. Mapping from 1D chromosome to 2D circuit.

Since a solution similarity metric depends on how solutions/circuits are encoded, we describe our representation. An individual in the GA's population encoded as a bit string maps to a two-dimensional circuit as shown in figures 2 and 3. We use four (4) bits to encode all $2^{2^1} = 16$ possible two-input, one-output gates. An additional bit specifies the location of the second

input. A gate $S_{i,j}$ gets its first input from $S_{i,j-1}$ and its second, from one of $S_{i+1,j-1}$ or $S_{i-1,j-1}$ as shown in figure 3. If the gate is in the first or last

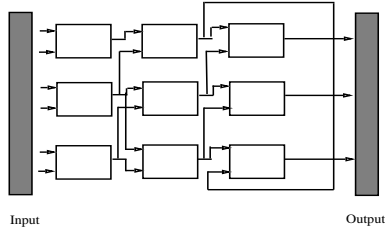


Fig. 3. A gate in a 2D template gets its second input from one of two gates in the previous column.

rows, the row number for the second input is calculated modulo the number of rows. The gates in the first column, $S_{i,0}$ receive the input to the circuit.

4.3 Solution Similarity

The solution similarity metric needs to measure the distance between encoded circuits (solutions). We are assuming that similar circuits have similar function. Since we use a binary string to encode combinational circuits, the hamming distance between these bit strings suffices as our similarity metric on this problem. We show that this simple similarity metric works well for these parity checker design problems. Real number strings could use Euclidean distance.

4.4 Configuration Design

Using the parity problem as a basis, we randomly generate 50 problems that are similar in terms of our problem similarity metric by randomly flipping 12% of the output bits of the 6-bit parity checker's output bit string.

Figure 4 shows the distribution of problems in terms of problem similarity. Most problems' output strings are between 6 and 7 bits different, no two problems are the same, and the similarities are normally distributed.

In this paper, CIGAR uses a population of size 30 run for a maximum of 30 generations to solve 6-bit combinational circuit design problems (similar to parity checkers). This results in a 150 length chromosome (6 rows, 5 columns, 5 bits per gate) with a corresponding search space of 2^{150} . The probability of crossover is 0.95 and the probability of mutation is 0.05. We inject three cases (10% of population size) into the population every five generations. Previous work explains these parameter values [17]. All plots are averages over ten (10) runs. Figure 5(left) compares performance in terms of time taken to find a solution for CIGAR and a randomly initialized genetic algorithm. The figure

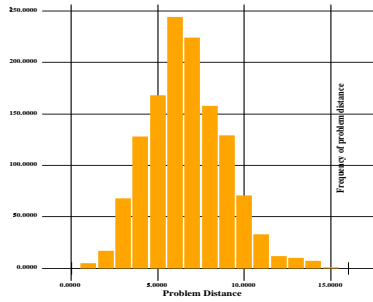


Fig. 4. Frequency distribution of problem similarities

plots time taken to find the best solution on the vertical axis and the number of problems solved on the horizontal axis. The Randomly Initialized Genetic Algorithm (RIGA) uses the same set of genetic algorithm parameters. We ran both ten times with different random seeds.

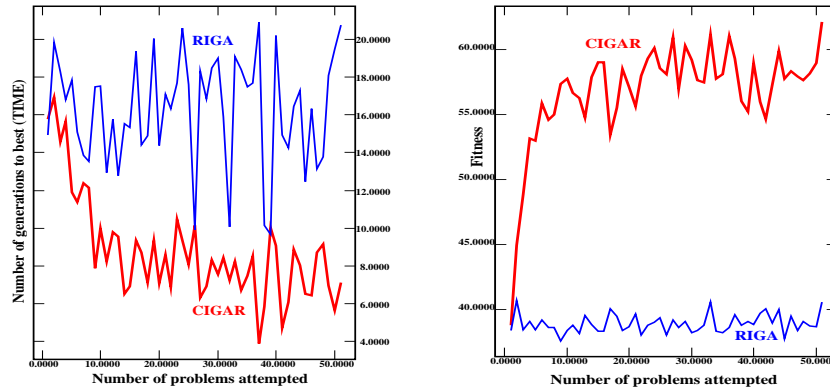


Fig. 5. Left: Convergence speed. As more problems are attempted, CIGAR reduces the time to convergence. Right: Solution quality. As more problems are solved CIGAR produces better solutions

Figure 5 (right) plots the fitness of the best solution found on the vertical axis and number of problems solved on the horizontal axis. The figures clearly show that CIGAR takes less time to produce better quality solutions as it gains experience from attempting more problems. Figure 6 plots the distribution of solutions' similarities produced by CIGAR and RIGA. CIGAR solutions tend to be more similar indicating that CIGAR is storing and reusing solutions and partial solutions to previously solved problems.

Figure 7 (right) plots the solution quality as a function of experience for the following injection strategies.

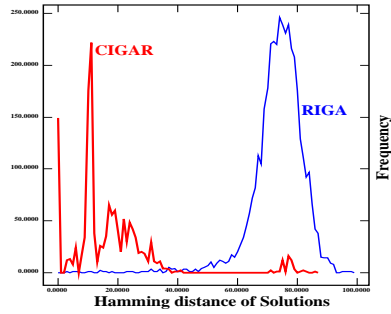


Fig. 6. Solution distribution

1. Closest to the best
2. Probabilistic closest to the best
3. Furthest from the worst
4. Probabilistic furthest from the worst
5. Randomly choosing individuals from the case-base to inject
6. Creating and injecting randomly generated individuals

In all cases, we replace the worst individuals in the population with the individuals chosen (or created) by the injection strategy. The figure shows that the probabilistic strategies and the strategy of randomly injecting individuals from the case-base result in the largest increase in quality with experience. As expected, creating and injecting random individuals does not lead to better quality with experience. Both deterministic strategies show intermediate gains in quality. Figure 7 (left) shows a similar trend for the time to solution measure of performance. We did not label the graph with injection strategies because we are highlighting the difference between deterministic and probabilistic strategies and because we have not completed our investigation of injection strategies.

5 Discussion and Conclusions

CIGAR makes the assumption that similar problems have similar solutions. According to the schema theorem, genetic algorithms process syntactic string similarities and we have shown that storing and injecting syntactically similar solutions leads to increased performance with experience. CIGAR thus combines the strengths of GAs and CBR and avoids the difficult problem of measuring similarities among problems.

Our sample results indicate that CIGAR learns to increase performance at related tasks as it gains experience. We show that the simple syntactic similarity measure of hamming distance works well on combinational circuit design problems. It is important to note that we need to save (and re-use) to the case-base, not just the best or final solution to the problem at hand, but

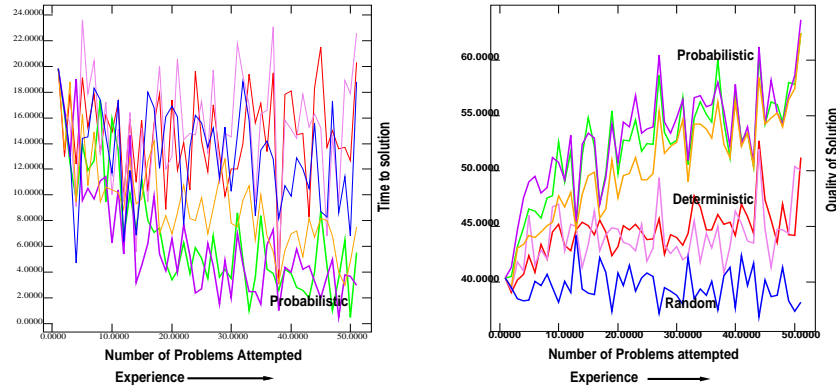


Fig. 7. Comparing Injection Strategies. Left: Average time to find the best design versus number of problems attempted. Right: Quality of design solution versus number of problems attempted

also intermediate solutions that the GA generates while working on a problem. We also find that the more similar the problems the larger the difference in performance between CIGAR and RIGA and the more individuals that can be injected without loss of performance. However, injecting too many individuals often leads to premature convergence. Since we do not usually know problem similarity injecting between 5% – 15% of the population size strikes a good balance.

Our experiments also indicate that we can use one of several ways to choose individuals to be injected with little difference in CIGAR performance. We can inject the closest (individuals in the case-base) to the best individual in the population, the farthest from the worst, and probabilistic versions of both. The probabilistic versions seem to result in greater performance improvements and we are currently working confirming these initial impressions.

We are applying CIGAR to problems in object identification, spectroscopic analysis of dense plasmas, and real-time targeting and re-targeting. Because these problems are computationally intensive or require real-time responses, we will be building and investigating a parallel CIGAR implementation, different selection schemes, and different case-base implementations. On the theoretical side we are modeling and quantifying our qualitatively derived parameter values for CIGAR.

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