Finding Good Compiler Optimization Sets A Case-based Reasoning Approach

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Abstract:

Case-Based Reasoning have been used for a long times to solve several problems. The first Case-Based Reasoning used to find good compiler optimization sets, for an unseen program, proposed several strategies to tune the system. However, this work did not indicate the best parametrization. In addition, it evaluated the proposed approach using only kernels. Our paper revisit this work, in order to present an detail analysis of an Case-Based Reasoning system, applied in the context of compilers. In addition, we propose new strategies to tune the system. Experiments indicate that Case-Based Reasoning is a good choice to find compiler optimization sets that outperform a well-engineered compiler optimization level. Our Case-Based Reasoning approach achieves an average performance of 4.84% and 7.59% for cBench and SPEC CPU2006, respectively. In addition, experiments also indicate that Case-Based Reasoning outperforms the approach proposed by Purini and Jain, namely Best10.

1 INTRODUCTION

Case-Based Reasoning (CBR) (Richter and Weber, 2013), an approach considered a subfield of machine learning (Mitchell, 1997; Shalev-Shwartz and Ben-David, 2014), tries to solve a new problem using a solution of an previous similar situation. It can be seen as a learning process (Aamodt and Plaza, 1994), which stores past experiences in a knowledge database, and can be updated incorporating new experiences (Jimenez et al., 2011).

Over the years, this approach have been applied to several problems, such as: estimate the project cost to web hypermedia (Mendes and Watson, 2002), estimate the Q-factor of an optical network (Jimenez et al., 2011), management of typhoon disasters (Zhou and Wang, 2014), and estimate good compiler optimization sets (Lima et al., 2013).

Compilers are programs that transform source code from one language (source language) to another (target language) (Aho et al., 2006; Srikant and Shankar, 2007; Cooper and Torczon, 2011). During this process, the compiler applies several optimizations (Muchnick, 1997), in order to improve the target code. However, some optimizations can be good to a class of programs, and bad to another. Then, the most appropriate approach is to find the best compiler optimizations to each program. The literature presents different approaches to mitigate this problem (Zhou and Lin, 2012; Lima et al., 2013; Purini and Jain, 2013). The first CBR approach in this context (Lima et al., 2013) indicates that it approach is able to infer good solutions to new problems - good compiler optimization sets to unseen programs.

In this paper we revisit the work of Lima *et al.* (Lima et al., 2013) to explore new strategies of finding good compiler optimization sets to a unseen program. Lima *et al.* (Lima et al., 2013) used dynamic features to represent analogies and a leave-one-out cross-validation approach. Our work uses dynamic features or static features to represent analogies. In addition, our work uses different training and test datasets for cross-validation.

The main contributions of this paper are:

- We describe different CBR parametrization.
- We give a new similarity metric to measure the similarity between two programs.
- We present a program characterization approach using static features.
- We present different strategies to build a collection of past experiences.
- We present a detail experimental analysis of CBR approach in the context of compilers.

This paper is organized as follows. Section 2 presents related work. Section 3 explains briefly

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CBR. Section 4 details our CBR approach, which goal is to find good compiler optimization sets. Section 5 presents the experimental setup. Section 6 details the experimental results. Finally, Section 7 presents conclusions and future work.

2 RELATED WORKS

In the context of compilers, the work of Purini and Jain finds a small set of compiler optimizations sets (COS), which cover several programs (Purini and Jain, 2013). Using iterative compilation, Zhou evaluates a random and a genetic strategy, in order to find good compiler optimizations (Zhou and Lin, 2012). Our work tries to mitigate the same problem. However, using a different approach.

Lima *et al.* proposed the first CBR approach to find good COS to a unseen program. This approach uses different strategies to select past results and measure the similarity between programs (Lima et al., 2013). The work of Lima *et al.* does not indicate which parametrization is the best. In addition, the benchmark used is composed only by kernels. It is a problem, in general we use complete applications and not kernels. Therefore, we revisit this work in order to cover these gaps.

Jimenez used a CBR approach to estimate Q-factor in optical networks. His approach obtained a successful classification in 94% of cases (Jimenez et al., 2011). Erbacher used a same approach to automatically report hostile actors in a network (Erbacher and Hutchinson, 2012). To implement a system able to generate combat strategies, Kim *et al.* (Kim et al., 2014) proposed a CBR approach that retrieves past experiences and modify them to the current situation. The main difference between these works and the our is the context where the CBR is applied.

3 CASE-BASED REASONING

CBR, a machine learning approach, can be subdivided in four processes:

- 1. Retrieve a case from a collection of (past experiences) previous cases by similarity measure.
- 2. Reuse the knowledge of an old case to solve a new case.
- 3. Revise the result of this new case, evaluating the success of the solution.
- 4. Retain the useful experience for future reuses.

Every CBR, in specially the retrieve process, needs some parameters, such as:

- **Collection guide** indicates the strategy used to build the collection of previous cases.
- **Similarity measure** measures the level of similarity between a previous case and a new one.
- **Standardization** transforms all attributes values according to a specific rule.
- **Number of analogies** indicates the number of previous cases that will be used to estimate a solution to an unseen problem.

In general, the cases that compose the collection of previous cases come from real world experiences. However, in our context, there is not a public real world collection, used by the scientific community. Therefore, we need to generate a collection of past experiences to use it as previous cases.

4 FINDING GOOD COMPILER OPTIMIZATION SETS USING A CBR APPROACH

The main goal of the CBR described in this paper is to find a COS, which is able to achieve a performance improvement over a well-engineered compiler optimization level.

The CBR approach divides the process of finding an effective COS into:

- 1. An offline phase; and
- 2. An online phase.

The offline phase collects pieces of information about a set of training programs, and downsamples the search space in order to provide a small space, which can be handled by the online phase in a easy and fast way. Therefore, the offline phase creates a collection of previous cases, which will be used to determine the knowledge that will be used to solve an unseen case, in other words, to determine a COS that should be enabled on an unseen program.

The online phase will infer from the cases provided by the offline phase, the best COS that fits the feature of unseen program as defined by its input.

4.1 The Offline Phase

The offline phase builds a collection of previous cases storing for training programs several success cases. It means that the downsampling technique should be guided to retain good cases, besides pruning the

Algorithm 1	: Offlin	ne Phase.
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search space. The Algorithm 1 describes briefly the offline phase.

Algorithm 1 indicates that the feature for each training program is normalized. It is performed by the offline phase, due to different programs have different features, for example the number of hardware instructions executed, runtime, and others. In addition, each feature should be collected compiling and running the training program without any compiler optimization enabled. It guarantees that the optimizations will not influence the program behavior.

4.2 The Online Phase

The online phase performs the CBR, in order to find an effective COS, which should be enabled on the unseen (test) program. The Algorithm 2 describes briefly this phase.

For the test program, the online phase collects its features and compares them with the features of each training program, with the help of a similarity model that ranks the training programs. This rank indicates what training program is the most similar to test program. In summary, the online phase selects from the most similar training programs previous cases, evaluates these cases and returns the best one.

Algorithm 2: Online Phase.
Input: C // Collection
P // Test program
B // Baseline
N // Number of analogies (cases)
S // Similarity
Output: The best case
$cases \leftarrow []$
$best_case \leftarrow B$
// Compile the program P using the baseline,
// and get the number of hardware
// instructions executed
$best_perf \leftarrow getPerformance(B,P)$
// Get the features of the program P,
// and normalize them
$info \leftarrow \text{getNormalizedInformation}(P)$
// Build a collection with only success cases
// (cases that improved the performance // of training programs)
$C' \leftarrow filterCollection(C)$
// Rank the training programs
// based on similarity model S
$rank \leftarrow getRank(C', info, S)$
// Get the potential previous cases
$cases \leftarrow getAnalogies(C', L, rank)$
// Evaluate the potential cases
for each $case \in cases$ do
// Compile the program P using the
// case, and get the number
// of hardware instructions executed
$perf \leftarrow getPerformance(case, P)$
if <i>perf</i> < <i>best_perf</i> then
$best_perf \leftarrow perf$
$best_case \leftarrow case$
return <i>best_case</i> , <i>best_perf</i>

4.3 Parametrization

The parametrization of our CBR is as follows.

Collection Guide. Several strategies can be used to build a collection of previous cases. In our work, we use iterative and metaheuristic algorithms to perform this task. The iterative algorithm generates the collection by a uniform random sampling of the optimization space. While, the metaheuristic algorithms use a sophisticated way to build a collection. The metaheuristic are genetic algorithm with rank selection, genetic algorithm with tournament selector, and simulated annealing. In addition, we build a collections.

- **Similarity Model.** In order to measure the similarity between two programs, the CBR system can be tunned to use a strategy chosen from:
 - **Cosine.** In this model, the similarity between P_i and P_j is defined as:

$$sim(F_i, F_j) = rac{\sum\limits_{w=1}^{m} F_{iw} \times F_{jw}}{\sqrt{\sum\limits_{w=1}^{m} (F_{iw})^2} \times \sqrt{\sum\limits_{w=1}^{m} (F_{jw})^2}}$$

Jaccard. In this model, the similarity between F_i and F_j is defined as:

$$sim(F_i, F_j) = \frac{1}{m} \sum_{w=1}^{m} \frac{\min(F_{iw}, F_{jw})}{\max(F_{iw}, F_{jw})}$$

Euclidean. In this model, the similarity between F_i and F_j is defined as:

$$sim(F_i, F_j) = \frac{1}{\sqrt{\sum_{w=1}^m (F_{iw} - F_j)}}$$

In both models, *m* is the quantity of features. The first two similarity models were proposed by Lima *et al.* (Lima et al., 2013). In our work, we use these two models and a model based on Euclidean distance. As mentioned previously by Lima *et al.*, these models are based on similarity coefficients used to compare statistical sampling, and indicate that large difference between two feature vectors should return a low similarity value.

- Feature. A similarity model measures the similarity between two programs, based on their features. Our CBR is able to use dynamic or static features to describe the program behavior. Dynamic features are composed by hardware performance counters (Mucci et al., 1999), which describe the program behavior during its execution. Static features are composed by compiler statistics, which describe the program behavior during its compilation. Table 1 presents the dynamic features used in our work, and Table 2 presents the static features. In addition, these tables indicate the most important feature (*). Our CBR system can use all features, only the most important features, or a weighted strategy. In the weighted strategy the feature vectors are weighted to reflect the relative importance of each feature. In our system, the weight of an important feature has value 2, while the other has value 1.
- **Standardization.** The system should divide each feature for the most important one (**). It transforms all attributes values in order to standardize the features of training and test programs.
- Number of Analogies. The best strategy is to choose several analogies to increase the probability of

Table 1: Dynamic Feature.

Туре	Performance Counter	er
	PAPI_L2_DCR	PAPI_L3_DCR
	PAPI_L2_TCA*	PAPLL3_TCA
	PAPI_L2_DCW	PAPI_L3_DCW
	PAPI_L1_ICM	PAPI_L2_STM
	PAPI_L1_DCM	PAPI_L3_TCM
	PAPI_L2_TCM	PAPI_L3_TCR
Cache	PAPI_L2_TCR	PAPLL3_DCA
Cache	PAPI_L2_DCA	PAPI_L3_TCW
	PAPI_L2_TCW	PAPI_L2_ICR
	PAPI_L2_DCH*	PAPI_L1_STM
	PAPI_L1_TCM	PAPI_L1_LDM
	PAPI_L2_DCM	PAPI_L2_ICA
	PAPI_L3_ICR	PAPI_L2_ICM
	PAPI_L3_ICA	PAPI_L2_ICH*
	PAPI_BR_PRC	PAPI_BR_UCN
D 1	PAPI_BR_NTK	PAPI_BR_INS*
Branch	PAPI_BR_MSP	PAPI_BR_TKN
	PAPI_BR_CN	
SIMD	PAPI_VEC_SP*	PAPI_VEC_DP*
	PAPI_FDV_INS	PAPI_FP_INS
Floating Point	PAPL FP_OPS	PAPLSP_OPS
-	PAPL_DP_OPS	
TLB	PAPI_TLB_DM*	PAPI_TLB_IM
<i>a</i> .1	PAPLREF_CYC	PAPLTOT_CYC*
Cycles	PAPI_STL_ICY	PAPI_STL_ICY
Instructions	PAPL TOT INS **	

Table 2: Static Feature.

	Static Data
Binary Instructions	Number of Add insts
Binary instructions	Number of Sub insts
	Number of Store insts*
	Number of Load insts*
Memory Instructions	Number of memory instructions*
	Number of GetElementPtr insts
	Number of Alloca insts
Terminator Instructions	Number of Ret insts
Terminator Instructions	Number of Br insts
	Number of ICmp insts
Other Instructions	Number of PHI insts
Ould hisudeholis	Number of machine instrs printed*
	Number of Call insts
Function	Number of non-external functions
Basic block	Number of basic blocks
Floating Point Instructions	Number of floating point instructions*
Total Instructions	Number of instructions (of all types)* **

choosing a good one. Our system is able to evaluate several number of cases. However, the most similar training program can not be able to provide the required number of analogies. If it is the case, the second most similar program will provide it, and so on.

5 EXPERIMENTAL SETUP

In our experiments, we will evaluate different configurations of our CBR system. The main configuration of the experimental environment is given by:

- Hardware. We used a machine with a Intel processor Core I7-3779, 8 MB of cache, and 8 GB of RAM.
- **Operating System.** The operating system was Ubuntu 14.04, with kernel 3.13.0-37-generic.
- **Compiler.** We adopted LLVM 3.5 (Lattner and Adve, 2004; LLVM Team, 2014) as compiler in-frastructure.
- **Baseline.** The baseline is the LLVM's highest compiler optimization level, -O3. The baseline indicates the threshold that our system should overcome.
- **Optimizations.** The optimizations used to compose a case are present in Table 3. We use only the optimizations used by the highest compiler optimization level -O3.

Optimizations						
-inline	-prune-eh -scalar-evolution					
-argpromotion	-inline-cost	-indvars				
-gvn	-functionattrs	-loop-idiom				
-slp-vectorizer	-sroa	-loop-deletion				
-globaldce	-domtree	-loop-unroll				
-constmerge	-early-cse	-memdep				
-targetlibinfo	-lazy-value-info	-memcpyopt				
-no-aa	-jump-threading	-sccp				
-tbaa	-loop-unswitch	-dse				
-basicaa	-tailcallelim	-adce				
-notti	-reassociate	-barrier				
-globalopt	-loops	-branch-prob				
-ipsccp	-loop-simplify	-block-freq				
-deadargelim	-lcssa	-loop-vectorize				
-instcombine	-loop-rotate	-strip-dead-prototypes				
-simplifycfg	-licm	-verify				
-basiccg	-correlated-propag	ation				

Table 3: LLVM's optimizations used by -O3.

Cases. The process of creating a case is guided by the criteria:

- Every optimization appears only once in a case;
- Every optimization can appear in any position;
- Every optimization should address the compilation infrastructure rules; and
- All cases have the same length.

The first criterion indicates that the offline phase does not explore the use of one optimization several times. Although, this occurs in LLVM's -O3 optimization level. Second can be viewed as an

organization of the case (or COS). In the collection, every case is represented as a sequence of optimizations. It means that there is a predefined order to apply each specific optimization. The third indicates that a new case can not violate the safety of the infrastructure. By the fourth criterion, the offline phase tries to give to every case the same characteristic.

- **Collection Guide.** The creation of the collections was guided as follows.
 - **Random.** This iterative algorithm generates in a random way 500 cases.
 - Genetic Algorithm with Rank Selector. The
 - parameters chosen in this strategy were: chromosome_size $\leftarrow 40$ (the number of optimizations in a case), population $\leftarrow 60$, generation $\leftarrow 100$, mutation_rate $\leftarrow 0.02$, and crossover_rate $\leftarrow 0.9$. The algorithm will finish whether the standard deviation of the current fitness score is less than 0.01 or the best fitness score does not change in three consecutive generations. In addition, this strategy uses elitism. It means that the best solution of the generation N - 1 is kept in generation N.
 - Genetic Algorithm with Tournament Selector. It is similar to the previous strategy, but instead of using a rank selector it uses a tournament selector.
 - **Simulated Annealing.** In this strategy, the initial temperature is the half of hardware instructions executed, the perturbation function only changes one random optimization in a random position, the acceptance probability is given by the Equation 1, and the temperature is adjusting multiplying it by the constant α , which value is 0.95. The stop criteria is 500 iterations.

$$P(N) = e^{-\frac{\Delta(C) - \Delta(N)}{T}} \tag{1}$$

- **All.** This strategy only merges the previous collections.
- **Training Programs.** For the generation of the collections of previous cases, we used 61 microkernel programs take from LLVM's test-suite. All the programs are single-file, and have short running times. Table 4 shows the microkernels.
- **Test Programs.** We used for evaluating our CBR approach the benchmarks, cBench (cBench, 2014) with dataset 1, and SPEC CPU2006 (Henning, 2006) with training dataset.
- Validation. The results is based on the arithmetic average of five executions, excluding the best

	Microkernel pro	grams	
ackermann	hash	perlin	
ary3	heapsort	perm	
bubblesort	himenobmtxpa	pi	
chomp	huffbench	puzzle	
dry	intmm	puzzle-stanford	
dt	lists	queens	
fannkuch	lowercase	queens-mcgill	
fbench	lpbench	quicksort	
ffbench	mandel-2	random	
fib2	mandel	realmm	
fldry	matrix	recursive	
flops-1	methcall	reedsolomon	
flops-2	misr	richards_benchmark	
flops-3	n-body	salsa20	
flops-4	nestedloop	sieve	
flops-5	nsieve-bits	spectral-norm	
flops-6	objinst	streat	
flops-7	oourafft	towers	
flops-8	oscar	treesort	
flops	partialsums	whetstone	
fp-convert			

Table 4: Microkernels.

and the worst values. In the experiments, the machine workload was minimum as possible, in other words, each instance was executed sequential. In addition, the machine did not have external interference, and the runtime variance was less than 0.01.

5.1 An Overview of the Collections

To analyze the improvement obtained by each strategy of creating the collections, we summaries the collections in Table 5. In this table, the column #P means the number of programs with good cases.

Collection	#P	#Sets	Good Cases (%)	Worst Perf. (%)	Best Perf. (%)
Random	55	30500	19.57	-99.99	58.33
Genetic Algorithm Rank	57	27973	39.42	-99.99	51.87
Genetic Algorithm Tournament	56	27543	56.77	-99.99	51.87
Simulated Annealing	45	30500	7.42	-99.99	54.22
All	57	116455	30,54	-99.99	58.33

Table 5: Summary.

Most cases generated by simulated annealing do not overcome the baseline. However, this strategy was able to find some cases that achieve a good improvement over the baseline. It indicate this strategy has difficulty to escape from some bad improvement, which is probably due to the disturbing function (the function that chooses the neighbor). Besides, the initial solution influences the final result, which could be the source to several cases achieve bad improvements.

The overview of the other strategies shows a different scenario. The genetic algorithms generated a better distribution. They have more success cases, if we take the average as criteria to evaluate the collection quality. However, if we take the maximum improvement obtained for each program, simulated annealing generates better results than genetic algorithms, and random strategy.

These differences can be justified by the characteristics of each implementation. Simulated annealing just change one optimization in each new case, while the others metaheuristics try more changes in each new case.

The number of cases is a small portion of the optimization space. It indicates that the strategies used to generate the collections was able to downsample the search space in order to provide a small space with good cases.

and the worst values. In the experiments, the 6 EXPERIMENTAL RESULTS

The goal of our CBR approach is to find a COS that outperforms the well-engineered compiler optimization level -O3, in terms of hardware instructions executed.

Our experiments was conducted in a way to answer the following questions:

- What is the best parametrization of the CBR approach applied in the context of compilers?
- What is the best characterization of programs?
- What is the improvement obtained by the CBR approach in real applications?
- What is the best strategy to create a collection of previous cases?

Tables 6 and 7 show the results obtained by each CBR configuration. In these tables, API stands for average percentage improvement, APIE stands for average percentage improvement excluding the programs showing no improvement, and NPI stands for the number of programs achieving improvement.

Strategies that use static features do not generate different results. It means that the use of these configurations always ranks the training programs in the same way. It explains the use of only one entry for static features.

6.1 Overview

Collection Guide. Analyzing the different strategies to construct the collections shows that the random

	cBe	ench		1	Analogy	,	3	Analogies	s	5	Analogie	S	
Base		Similarity		API	APIE	NPI	API	APIE	NPI	API	APIE	NPI	
		4 11	Cosine	-1.88	5.91	12	-1.12	6.58	14	0.71	7.15	19	
		All (AL)	Jaccard	-2.86	5.55	13	-1.61	5.78	16	-1.21	5.8	17	
		· · · ·	Euclidean	-1.18	6.27	13	-0.62	6.45	15	0.7	6.7	19	
Genetic	Dynamic	Most	Cosine	-5.51	3.58	13	-3.41	4.28	14	-1.47	6.32	16	
Algorithm	Feature	Informative	Jaccard	-1.89	4.81	13	-0.56	5.28	17	0.18	5.79	18	
Rank	(DF)	(MI)	Euclidean	-0.4	4.57	14	0.09	4.83	16	1.5	5.35	20	
Selection		Weight	Cosine Jaccard	-2.13 -2.86	5.94 5.55	12 13	-1.02	6.49 5.78	14 16	0.79	7.07 6.11	19 17	
(GR)		(WE)	Euclidean	-2.80	6.29	13	-0.46	6.07	16	0.78	6.39	20	
-	St	tatic	Cosine	-7.59	5.45	8	-6.8	5.15	10	-6.71	5.15	10	
		ature	Jaccard	-1.88	5.84	16	0.17	5.77	18	0.19	5.79	18	
	(!	SF)	Euclidean	-4.28	6.06	12	-3.71	6.03	13	-3.41	5.61	14	
			Cosine	-1.87	4.4	15	-0.41	4.97	18	0.1	5.52	18	
		All	Jaccard	-1.52	4.97	16	0.14	5.83	18	0.48	5.95	19	
		(AL)	Euclidean	-1.87	4.4	15	-0.41	4.97	18	0.1	5.52	18	
Genetic	Dynamic	Most	Cosine	-4.38	4.31	13	-2.69	5.87	14	-2.28	6.06	15	
Algorithm	Feature	Informative	Jaccard	-3.77	4.39	13	-1.23	6.13	17	-0.82	5.92	19	
Tournament	(DF)	(MI)	Euclidean	-2.8	3.5	15	-1.5	3.67	18	-1.06	4.08	18	
Selection		Weight	Cosine	-1.61	4.03	16	-0.45	4.45	19	0.05	4.92	19	
(GT)		(WE)	Jaccard	-1.85	4.97	16	-0.19	5.83	18	0.15	5.95	19	
		tatic	Euclidean Cosine	-0.57 -5.84	4.12 4.02	15 11	0.61	4.54	18 - 12	1.26 -5.43	4.98 4.17	19 12	
SCIE		ature	Jaccard	-2.66	6.02	13	-2.21	5.63	12	-1.82	5.63	12	
		SF)	Euclidean	-3.31	3.67	13	-2.16	5.11	10	-2.11	5.13	10	
	(.		Cosine	-4.97	3.88	12	-2.04	5.78	17	-1.72	5.87	18	
		All	Jaccard	-2.97	6.29	12	0.79	7.11	17	2.35	6.64	20	
		(AL)	Euclidean	-5.89	4.39	10	-3.43	6.94	13	-2.8	6.58	16	
	Dynamic	Most	Cosine	-8.73	4.94	8	-1.65	6.25	13	0.74	6.39	17	
S' 1.4.1	Feature		Informative	Jaccard	-4.19	5.46	13	-0.4	6.38	18	0.32	6.37	19
Simulated Annealing	(DF)	(MI)	Euclidean	-6.12	1.69	11	-3.57	3.85	17	-2.02	6.2	18	
(SA)		Weight	Cosine	-5.68	4.13	11	-2.75	6.07	16	-2.42	6.46	16	
(SII)		(WE)	Jaccard	-3.1	6.69	15	0.52	7.27	17	1.87	6.93	18	
		. ,	Euclidean	-6.36	4.39	10	-3.83	6.6	14	-3.31	6.94	15	
		tatic	Cosine	-8.88 -4.4	4.73 3.66	9 9	-1.49	5.56 4.08	15 17	-0.93 -1.25	5.37 4.72	18 18	
		ature SF)	Jaccard Euclidean	-4.4	4.54	12	-2.01	4.08 6.04	17	0.04	5.92	18	
	()	51)										1	
		All	Cosine Jaccard	-3.16 -3.25	5.23 7.11	11 10	1.21 0.05	6.28 6.62	20 16	1.92 1.14	6.88 6.59	21 19	
		(AL)	Euclidean	-2.95	4.84	10	1.05	6.03	20	1.14	6.73	21	
	Dynamic	Most	Cosine	-2.01	5.41	11	1.03	6.43	18	2.14	6.39	21	
	Feature	Informative	Jaccard	-0.98	6.6	13	2.02	6.52	19	2.58	6.84	20	
Random	(DF)	(MI)	Euclidean	-3.06	3.24	12	2.02	5.18	22	3.05	6.24	23	
(RA)		Weight	Cosine	-2.49	5.23	11	1.14	6.18	20	1.83	6.78	21	
		(WE)	Jaccard	-3.42	6.9	10	-0.04	6.35	16	1.05	6.37	19	
	~		Euclidean	-1.72	5.12	12	2.2	6.41	19	2.92	7.02	20	
		tatic	Cosine	-3.34	6.71	11	0.5	6.47	18	1.46	7.27	19	
		ature SE)	Jaccard Fuclidean	0.27	7.78 4.98	15	3.0	7.28 5.39	20 21	3.95	7.61	22	
	(;	SF)	Euclidean	-5.81		13	0.21			1.28	6.44	21	
		All	Cosine	0.45	3.86	16	2.43	5.52	20	3.16	5.97	20	
	Dynamic	(AL)	Jaccard Euclidean	-3.53 0.56	5.35 3.8	16 15	0.38 2.67	5.65 5.06	20 20	0.61 3.17	5.65 5.27	20 21	
		Most	Cosine	-7.38	2.53	8	-2.17	7.36	12	-1.12	7.31	15	
	Feature	Informative	Jaccard	-4.69	4.99	11	-0.77	5.79	12	-0.67	5.9	19	
All	(DF)	(MI)	Euclidean	-3.88	3.95	9	-1.24	5.43	15	1.29	4.75	21	
(AL)			Cosine	-1.46	4.1	15	0.6	5.36	19	1.36	5.59	20	
		Weight (WE)	Jaccard	-3.53	5.35	16	0.38	5.65	20	0.61	5.65	20	
			Euclidean	0.65	3.81	15	2.78	5.07	20	3.33	4.86	23	
		tatic	Cosine	-6.82	3.01	10	-5.36	3.66	12	-5.06	3.89	13	
		ature SF)	Jaccard Euclidean	-3.21 -6.96	3.61 3.37	12 10	-1.24 -1.83	5.12 5.12	18 14	-1.03 -1.54	4.86 5.22	19 15	

Table 6: Results obtained by cBench.

[SPEC C	PU2006		1	Analogy		3	Analogie	s	5	Analogie	s
Base	Similarity			API	APIE	NPI	API	APIE	NPI	API	APIE	NPI
		<u>,</u>	Cosine	-3.56	6.61	7	-2.46	7.09	7	-1.88	6.73	8
		All	Jaccard	-6.4	5.06	6	-4.96	7.56	6	-4.33	6.92	7
		(AL)	Euclidean	-3.12	5.48	8	-1.9	6.14	8	-1.29	6.07	9
Genetic	Dynamic	Most	Cosine	-5.55	6.62	7	-1.34	7.15	8	-0.63	6.96	9
Algorithm	Feature	Informative	Jaccard	-8.67	5.81	6	-4.8	5.55	8	-0.41	5.09	9
Rank	(DF)	(MI)	Euclidean Cosine	-6.35 -3.93	5.34 7.18	7 6	-5.15 -2.82	5.1 7.74	8 6	-4.25 -2.24	5.03 7.24	9 7
Selection		Weight	Jaccard	-5.95	5.06	6	-2.82	7.56	6	-4.33	6.92	7
(GR)		(WE)	Euclidean	-3.42	5.44	8	-2.21	6.1	8	-1.6	6.04	9
	S	tatic	Cosine	-12.03	2.97	2	-8.39	2.42	3	-7.97	2.97	4
		ature	Jaccard	-5.58	5.09	4	-4.92	5.15	4	-4.58	4.32	5
	(SF)	Euclidean	-11.71	2.58	3	-8.08	2.27	4	-7.81	3.18	4
		All	Cosine	-3.4	4.86	7	-2.9	5.4	7	-2.74	5.79	7
		(AL)	Jaccard	-1.63	5.99	7	-1.11	6.11	8	-0.56	6.61	8
	Dunamia	Most	Euclidean Cosine	-4.57 -8.34	4.86	7	-4.12 -4.91	5.4 4.25	7	-3.96 -2.87	5.79 4.64	7
Genetic	Dynamic Feature	Informative	Jaccard	-8.54	7.45	5	-4.91	6.24	7	-2.87	6.9	7
Algorithm	(DF)	(MI)	Euclidean	-5.15	4.49	8	-1.21	6.99	9	-0.82	6.9	10
Tournament			Cosine	-4.87	4.68	8	-1.81	5.32	9	-1.65	5.63	9
Selection (GT)		Weight (WE)	Jaccard	-2.35	5.99	7	-1.84	6.11	8	-1.3	6.61	8
			Euclidean	-5.31	5.07	8	-2.37	5.4	9	-2.21	5.72	9
		tatic ANL	Cosine	-8.65	3.45	6	-8.12	3.25	9	-7.71	3.25	7—
		ature SF)	Jaccard Euclidean	-3.26 -8.6	3.14 3.5	7	-2.72 -8.09	3.14	8 7	-2.26 -7.68	3.16	8
	(31)		-					-			-
		All	Cosine Jaccard	-3.12 -8.13	9.07 6.12	3 7	-1.32	8.37 7.89	6 7	-1.05	7.18 8.07	7
		(AL)	Euclidean	-3.12	9.07	3	-1.28	7.26	7	-0.94	6.53	8
	Dynamic	Most	Cosine	-3.93	8.1	6	-2.3	9.35	7	-1.63	9.7	7
Simulated	Feature	Informative	Jaccard	-4.97	5.55	7	-1.26	7.45	8	-0.07	7.57	8
Annealing	(DF)	(MI)	Euclidean	-6.15	11.38	2	-3.47	8.87	5	-2.65	8.78	6
(SA)		Weight	Cosine	-3.4	9.07	3	-1.67	9.04	5	-1.16	6.87	7
()		(WE)	Jaccard	-8.65	4.7	7	-2.45	6.41	7	-2.03	6.86	7
	S	tatic	Euclidean Cosine	-2.75 -10.04	7.94 5.49	5 5	-1.51 0.33	8.14 5.08	6 8	-1.03 0.67	7.38 6.01	7
		ature	Jaccard	-7.09	3.21	5	-1.93	3.82	5	-0.9	5.18	6
		SF)	Euclidean	-10.42	5.49	5	-0.13	5.08	8	0.35	6.01	7
			Cosine	-5.69	6.64	4	0.21	6.45	9	0.57	6.61	9
		All	Jaccard	-6.53	4.95	5	-1.39	5.78	7	-0.69	6.19	8
		(AL)	Euclidean	-4.38	5.33	5	0.47	5.86	10	0.87	6.09	10
	Dynamic	Most	Cosine	-4.94	8.79	6	0.51	9.13	9	1.23	9.61	9
	Feature	Informative	Jaccard	-5.07	5.94	6	-1.51	7.39	7	-0.97	7.32	8
Random (RA)	(DF)	(MI)	Euclidean Cosine	-6.92 -6.02	7.41 6.31	3	-2.63 -2.17	7.02	7 8	-1.74 -1.53	7.15	8
(KA)		Weight	Jaccard	-6.53	4.95	5	-1.39	5.78	8 7	-0.69	6.19	8
		(WE)	Euclidean	-5.86	6.7	5	-2.37	6.19	8	-0.64	5.84	10
	S	tatic	Cosine	-3.29	3.9	6	0.85	3.84	9	2.08	5.14	10
		ature	Jaccard	-4.83	3.1	7	0.24	2.88	10	1.66	4.33	11
	()	SF)	Euclidean	-3.42	4.01	6	0.57	3.82	9	1.79	4.73	10
		All	Cosine	-5.51	5.44	6	-3.07	6.44	7	-2.9	6.91	7
		(AL)	Jaccard	-4.11	4.46	6	-3.86	4.99	6	-3.06	5.78	7
	Dranssis		Euclidean	-5.77	5.14	6	-3.1	6.46	7	-2.9	7.01	7
	Dynamic Feature	Most Informative	Cosine Jaccard	-6.58 -6.95	10.56 4.8	5 6	-4.28 -4.66	9.04 5.62	7 6	-1.69 -1.49	9.41 6.24	7 7
All	(DF)	(MI)	Euclidean	-0.93	5.19	6	-4.60	5.92	8	-1.49	6.15	8
(AL)	(21)		Cosine	-5.68	5.62	6	-3.16	6.59	7	-2.98	7.06	7
()		Weight	Jaccard	-4.11	4.46	6	-3.86	4.99	6	-3.06	5.78	7
		(WE)	Euclidean	-5.76	5.09	6	-3.12	6.42	7	-2.92	6.97	7
		tatic	Cosine	-10.31	5.17	4	-6.76	5.55	4	-6.54	5.93	4
		ature	Jaccard	-3.63	4.52	5	-1.74	4.36	6	-1.57	4.36	6
	(SF)	Euclidean	-10.26	5.26	4	-6.73	5.65	4	-6.51	6.03	4

Table 7: Results obtained by SPEC CPU2006.

strategy got the best results in general. The use of random collection reached the best improvements. This strategy is the best for cBench in API, APIE, and NPI. The improvements of the simulated annealing are qualitative improvements, i. e., the cases in this collection reaches good improvements, but they cover a small number of programs. It can be seen in the results obtained by SPEC CPU2006. We also must highlight that the random approach in SPEC CPU2006 achieves the best API and NPI. The use of the collection with all cases not always obtained the best improvement. It occurs due to the potential previous cases is selected based on training programs, and not based on test programs. When the system uses the collection that store all cases, it can chooses a different case to validate the same test program. Note that a good case for a training program, not always is best for a test program.

- Similarities. The similarity models have different performance. In cBench, the Jaccard similarity model reached the best results. This model achieved the best improvements, and covered the most programs. In SPEC CPU2006, we have an scenario that Euclidean distance improved the most programs in general (NPI). However, the best percentage improvement for all programs (API) was obtained by Jaccard, and the best percentage improvement excluding the programs showing no improvement (APIE) was obtained by Cosine.
- Analogies. Increasing the number of analogies increases the API, APIE and NPI. This increase happens because when we choose more optimization sets to evaluate, we increase the probability of chosing a good case. In general, 3 analogies increases the performance up to 15%, and the coverage up to 61%, respectively. While, using 5 analogies increases the performance up to 4%, and the coverage up to 71%. Both, comparing with a configuration that uses only 1 analogy. This give us an idea that if we have two similar programs P and Q, and the optimization set S that is good for P, there is a high probability of S be good for Q. Otherwise, if S is a bad solution for P it also has a high probability of being a bad solution for Q.
- **Test Programs.** Observing our two benchmarks, SPEC CPU2006 can not be covered by our past examples. cBench reached best results evaluating this criteria. It indicates that cBench is more similar to microkernels than SPEC CPU2006.
- Feature. The most informative dynamic features obtained the best results, specially in SPEC

CPU2006. For cBench, not only dynamic features are required to cover all programs, we need some static features too.

6.2 CBR and Best10

In order to compare the performance of our CBR approach, we implemented the Best10 algorithm proposed by Purini and Jain (Purini and Jain, 2013). The Best10 algorithm finds the best 10 compiler optimization sets that cover several programs. To find these sets, it is necessary to downsample the compiler search space. It is done extracting from each training program, the best case from each collection of previous cases. In our experiments this new collection has 183 cases. After excluding the redundancies, the Best10 algorithm reduces the sample space in 10 cases. The work of Purini and Jain details this algorithm (Purini and Jain, 2013).

Tables 8 and 9 show the best results for each benchmark, using CBR with 5 analogies (the best configuration). In addition, these tables show the results obtained by Best10 algorithm.

The best results obtained by each program indicates that our CBR apprach is able to outperform the well-engineered compiler optimization level O3, and Best10 in several programs. CBR outperforms Best10 in 21 programs of cBench, and 15 programs of SPEC CPU2006.

The results show several configurations reaches the best improvements, mainly for SPEC CPU2006 and CRC 32^1 . In addition, these improvements are better than that obtained by Best10.

The results also indicate that CBR approach is better when using with cBench than SPEC CPU2006. In cBench, only for 6.45% of programs our CBR approach did not find a good previous case. This percentage increases in SPEC CPU2006 (26.32%). It indicates that our approach needs to be improved, in order to achieve better performance in complex benchmarks and cover more programs.

The improvement obtained by the CBR approach is better than that obtained by Best10, in cBench and SPEC CPU2006. In fact, Best10 does not outperform CBR. It indicates that the best choice is to analyze

¹For CRC32 the configurations that reach the best improvement are AL.DF.MI.E, AL.SF.AL.J, AL.SF.AL.C, AL.SF.AL.E, SA.DF.AL.J, SA.DF.MI.J, SA.DF.WEJ, SA.DF.AL.C, SA.DF.MI.C, SA.DF.WE.C, SA.DF.AL.E, SA.DF.MI.E, SA.DF.WE.E, SA.SF.AL.J, SA.SF.AL.C, SA.SF.AL.E, RA.DF.MI.J, RA.DF.MI.C, RA.DF.MI.E, RA.SF.AL.J, RA.SF.AL.C, RA.SF.AL.E, GR.DF.MI.C, GR.SF.AL.J, GR.SF.AL.J, GT.DF.WE.L, GT.DF.AL.J, GT.DF.MI.J, GT.DF.WE.J, GT.DF.AL.C, GT.DF.WE.C, GT.DF.AL.E, GT.DF.MI.E, GT.DF.WE.E, GT.SF.AL.J, GT.SF.AL.C, and GT.SF.AL.E.

Configurations

Program

Program	CBR		Best10
Flogram	Configurations	Imp. (%)	(%)
bitcount	AL.DF.AL.C, AL.DF.WE.C AL.DF.AL.E, AL.DF.MI.E AL.DF.WE.E	1.276	-45.748
qsort1	GR.DF.MI.C	10.958	6.709
susan_c	RA.SF.AL.J	31.561	27.042
susan_e	SA.DF.MI.C	4.279	1.862
susan_s	SA.DF.AL.J, SA.DF.WE.J SA.DF.AL.C, SA.DF.WE.C SA.DF.AL.E, SA.DF.MI.E SA.DF.WE.E	1.119	0.784
bzip2d	SA.DF.MI.J	20.796	39.972
bzip2e	RA.DF.MI.J	5.996	4.394
jpeg_c	RA.DF.AL.J, RA.DF.MI.J RA.DF.WE.J	11.436	5.686
jpeg_d	RA.SF.AL.C, RA.SF.AL.E	8.086	4.894
lame	AL.SF.AL.J	7.554	8.748
mad	GR.SF.AL.J	2.038	1.303
tiff2bw	RA.DF.MI.E	16.784	7.449
tiff2rgba	SA.DF.MI.E	14.194	7.485
tiffdither	SA.DF.AL.J, SA.DF.WE.J SA.DF.MI.C	-0.048	0.121
tiffmedian	RA.DF.MI.J, RA.DF.AL.C RA.DF.WE.C, RA.DF.AL.E RA.DF.WE.E	16.972	23.049
dijkstra	GR.DF.AL.J, GR.DF.WE.J	1.586	0.415
patricia	GT.DF.MI.E	0.321	0.562
ghostscript	SA.DF.MI.J	0.191	0.318
rsynth	RA.DF.MI.C	0.482	0.376
stringsearch1	RA.DF.MI.J, RA.DF.MI.E	-2.432	-18.242
	GR.DF.MI.E, GT.DF.MI.J		
blowfish_d blowfish_e	GR.DF.MI.E, GT.DF.AL.J GT.DF.MI.J GT.DF.WE.J	4.074 4.053	4.124 3.943
pgp_d	RA.SF.AL.J, RA.SF.AL.C	4.159	1.902
pgp_e	SA.SF.AL.E	1.216	0.587
rijndael_d	SA.DF.AL.C, SA.DF.WE.C SA.DF.AL.E, SA.DF.MI.E SA.DF.WE.E	13.125	0.003
rijndael_e	SA.DF.AL.C, SA.DF.WE.C SA.DF.AL.E, SA.DF.MI.E SA.DF.WE.E	14.659	-2.211
sha	SA.DF.AL.J, SA.DF.WE.J	6.969	7.605
CRC32	*1	2.126	2.126
adpcm_c	AL.DF.AL.C, GT.DF.AL.C GT.DF.WE.C, GT.DF.AL.E GT.DF.WE.E	13.336	5.831
adpcm_d	RA.SF.AL.J	16.96	11.183
gsm	SA.DF.AL.C, SA.DF.WE	1.609	1.967
API		7.593	3.656
APIE	1	8.202	6.412
APIE		0.202	0.112

Table 8: The Best Results obtained by cBench.

Table 9: Best Results obtained by SPEC CPU2006.

CBR

Best10

(%)

Imp. (%)

perlbench 12.926 10.138 AL.DF.MI.C, SA.DF.MI.C RA.DF.MI.C 6.882 7.280 bzip2 GR.DF.MI.C 13.43 -2.929 gcc RA.SF.AL.C, RA.SF.AL.E 10.128 mcf 9.307 GT.DF.AL.J, GT.DF.WE.J 3.49 -1.568 milc GT.DF.WE.C. GT.DF.MI.E GT.DF.WE.E 5.582 4.928 namd AL.DF.MI.J, SA.DF.MI.J RA.DF.AL.J, RA.DF.WE.J gobmk -0.176 -2.006 GR.DF.AL.J, GR.DF.MI.J dealII -9.654 -4.262 GR.DF.WE.J -0.174 AL.DF.MI.C 0.376 soplex GR.DF.AL.E, GR.DF.MI.E povray 0.485 0.679 GR.DF.WE.E SA.DF.MI.E, RA.DF.MI.E 6.57 3 064 hmmer SA.DF.MI.C, RA.DF.MI.C 6.75 5.495 sjeng SA.DF.AL.J, SA.DF.MI.J SA.DF.AL.C, SA.DF.MI.C SA.DF.WE.C, SA.DF.AL.E SA.DF.MI.E, SA.DF.WE.E libquantum 19.126 19.287 RA.DF.MI.J, RA.DF.AL.C RA.DF.MI.C, RA.DF.WE.C RA.DF.AL.E, RA.DF.MI.E RA.DF.WE.E h264ref GT.DF.MI.C 1.825 -1.352 SA.SF.AL.C, SA.SF.AL.E 0.587 0.557 lbm RA.SF.AL.C, RA.SF.AL.E AL.SF.AL.J, GR.SF.AL.J -1 303 -1 427 omnetpp GT.SF.AL.C, GT.SF.AL.E 3.062 -6.417 astar GT.DF.AL.J, GT.DF.MI.J sphinx3 13.99 12.811 GT.DF.WE.J AL.DF.AL.C, AL.DF.MI.C AL.DF.WE.C, AL.DF.AL.E AL.DF.MI.E. AL.DF.WE.E -1.719 -3.017 xalancbmk GR.DF.AL.C, GR.DF.MI.C GR.DF.WE.C, GR.DF.AL.E GR.DF.MI.E, GR.DF.WE.E API 4.840 2.897 APIE 7.499 6.706 NPI 14 11

the characteristic of a specific program, and not try to cover several programs.

6.3 Coverage

The results shown in the previous section presented a difficult, in using a CBR approach to find the best previous case that outperforms the well-engineered compiler level -O3, namely: it is necessary to try several configurations. This is a problem, due to the high response time. However, it is possible to use few configurations and obtain good results.

Using only three different configurations, it is possible to reach good results, without significant loss of performance. It is shown in Tables 10.

These results show a trade-off between performance and response time. It means that the best performance requires a high response time.

In cBench, it is necessary to tune several parameters to reach good results. However, there is a loss of performance up to 22.10%. For cBench, it is difficulty to reduce the variety of parameters. cBench needs at least three different collections of previous cases, and to use dynamic and static features.

In SPEC CPU2006, the system need to be tunned only in the similarity model. For this benchmark, the

cBench					
	Description		Number		
Restriction	of the best	APIE	of		
	Group		groups		
	RA.SF.AL.J				
No Restriction	SA.SF.AL.E	6.389	6		
	AL.DF.MI.E				
Only two Collections	-	-	-		
Only one Collection	-	-	-		
5	SPEC CPU2006				
	Description		Number		
Restriction	of the best	APIE	of		
	Group		groups		
	RA.DF.MI.C				
No Restriction	GR.DF.MI.E	6.678	76		
	GR.DF.MI.C				
	RA.DF.MI.C	_			
Only two Collections	GR.DF.MI.E	6.678	49		
	GR.DF.MI.C				
	GR.DF.MI.E				
Only one Collection	GR.DF.MI.C	5.978	1		
	GR.DF.MI.J	<u> </u>			

Table 10: Coverage Summary.

lost of improvement ranges from 10.95% to 20.28%. In this benchmark, the reduced number of collections is an excellent result, because this reduces the time spent in the offline phase.

7 CONCLUSIONS AND FUTURE WORK

- **Case-based Reasoning Approach.** In this paper we revisited the work of Lima *et al.*, in oder to explore new strategies of finding good compiler optimization sets. The strategy is to create an exploratory space that will be used by the Casebased Reasoning approach to predict the compiler optimization set, which should be enabled on an unseen program.
- **Results.** Our work indicate that if the main goal is to find the best configuration that achieves the best results, an interesting way it to use a random strategy to build a collection of previous cases, static features and Jaccard similarity model. On the other hand, if the main goal is to cover more programs, the use of dynamic features will be more useful, and metaheuristics. The use of metaheuristics improves the covering range.

In our results, the random strategy was a good choice to create a collection. Although, it was not expected. It does not mean we do not have to use metaheuristics to create collections. The results indicate that the collection created by the genetic algorithm with rank selector cover the maximum programs of the SPEC CPU2006 benchmark.

It is possible to find few configurations that achieves a good performance. It reduces the time spend in offline and online phases. However, there is a trade-off between response time and performance. Reducing the response time, in general, decreases the average percentage improvement.

The CBR approach obtained good improvements. It obtained an average percentage improvement of 4.84% for SPEC CPU2006, where some programs achieved a improvement up to 10%. For cBench, the CBR obtained an average of 7.499%, where some improvements up to 15%. Besides, our CBR approach outperforms the approach proposed by Purini and Jain, namely Best10.

Critical Discussion. This work shows that it is difficult to achieve performance based on only one configuration. Although, it is possible to achieve a performance better than state-of-the-art algorithms, it is necessary a high response time. Besides, it is difficult to achieve a good performance for complex programs.

We should note that finding the best compiler optimization set for a specific program, as defined by its input, is a problem without solution. Therefore, the metric that we use is the best improvement achieves by the best algorithm, presented in the literature (the state-of-the-art).

The deficiency of all compiler optimizations orchestration strategies is to handle programs (training and test) as a black box. Programs are composed by several different blocks. It indicates that each block will probably be best optimized by a specific compiler optimization set. It has to be investigated by new projects, in order to improve the state-of-the-art.

Future Work. We plan to propose new strategies to characterize programs, new strategies to create collections of previous cases, and new strategies to select previous cases. In addition, we are interested in proposing a CBR approach that is able to find different previous cases for different parts of the program.

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