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Motivation

- Code size is more important for embedded systems
- Traditional optimizations are designed to minimize execution time
- Developers can tolerate longer compile times
- Interplay between optimizations depends on the program
- Take exponential time to find optimal solution

Why GA is Good?

- Exponential time and space makes it impossible to find optimal solution
- Easy evaluation function
- Objective function is discrete and nonlinear
- Can be run on a program-by-program basis (even module-by-module)

Optimization Passes



- 1. Constant propagation (cprop)
- 2. Dead Code Elimination (dead)
- 3. Empty Basic Block removal (clean)

Optimization Passes

- 4. Global Value Numbering (valnum)
- 5. Lazy Code Motion (lazy)
- 6. Partial Redundancy Elimination (partial)
- 7. Peephole Optimization (combine)
- 8. Reassociation (shape)
- 9. Register Coalescing (coalesce)
- 10. Operator Strength Reduction (strength)

Genetic algorithm

• Genetic algorithm is a **black box** optimizer



• Genetic algorithm is a search inspired by Darwinian Evolution Principle



Genetic algorithm

• Genetic algorithm is a black box optimizer



• Genetic algorithm is a search inspired by Darwinian Evolution Principle





- A solution **x** is represented by a chromosome
 - A vector with L genes
 - (In this work, gene = opt flag)
 - (In this work, chromosome = opt sequence)
 - L = 12 in this work, i.e. only conisder opt seq with size 12.



- A solution **x** is represented by a chromosome
 - A vector with L genes
- Initialize: Randomly generate N chromosomes
 - N chromosomes are called **population**.

x1='(1121210	020′
x2= '2	1212110	221′
x3= '(0212210	221′
Gene	pool: {0.	1.2}

- A solution **x** is represented by a chromosome
 - A vector with L genes
- Initialize: Randomly generate N chromosomes
 - N chromosomes are called population.
 - N=20 in this work.

x1= '01121210020'
x2= '21212110221'
x3= '00212210221'
Gene pool: {0, 1, 2}

- A solution **x** is represented by a chromosome
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- Genetic algorithm generation: Update the population
 - Evaluation-> Selection->Crossover->Mutation



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 - **Evaluation**-> Selection->Crossover->Mutation
 - The evaluated objective f(x) is called fitness.



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 - Elicitism: (μ , λ)-selection



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 - Evaluation-> Selection->Crossover->Mutation
 - 2-point crossover: Discover new candidate solution from 2 chromosomes



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- N chromosomes are called population.
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 - Discover new candidate solution from 1 chromosome.

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 - A vector with L genes
- Initialize: Randomly generate N chromosomes



- N chromosomes are called population.
- Genetic algorithm generation: Update the population
 - Evaluation-> Selection->Crossover->Mutation
 - Discover new candidate solution from 1 chromosome.
 - Flip genes with some probability.

- A solution **x** is represented by a chromosome
 - A vector with L genes
- Initialize: Randomly generate N chromosomes
- x3= '20212110120' x4= '01101210221' x2= '21212110211' Gene pool: {0, 1, 2}

- N chromosomes are called population.
- Genetic algorithm generation: Update the population
 - Iterative update until average fitness converges

Genetic algorithm: Why it works

Problem structure & GA encoding: Let's consider a simple objective function f(x) = sum(x_i), x is binary vector.

Genetic algorithm: Why it works

Problem structure & GA encoding: Let's consider a simple objective function f(x) = sum(x_i), x is binary vector.

 Segment '..1..' is a good schemata that need to be preserved to form a better candidate -> Selection + Crossover !

Experiment

- Fortran
 - fmin, rkf45, seval, solve, svd, urand, zeroin
- C code
 - adpcm, compress, fft, dfa, dhrystone, nsieve
- 1000 generations / **1 day**

	Unoptimized		GA results				
Benchmark	static	dynamic	static	% red.	dynamic	% red.	gen. found
adpcm	438	17221981	351	19.9	12290460	28.6	6
compress	1753	8402188	1318	24.8	5545480	34.0	(77,79)
dfa	1744	842382	1107	36.5	496164	41.1	806
dhrystone	760	4920264	536	29.5	3200191	35.0	(22, 920)
fft	2415	18339859	1757	27.2	14574279	20.5	2
fmin	374	2192	187	50.0	963	56.1	32
nsieve	353	761244374	202	42.8	539954218	29.1	(0, 189)
rkf45	1525	511251	745	51.1	201470	60.6	74
seval	1061	3594	288	72.9	842	76.6	39
solve	1023	2729	437	57.3	1029	62.3	(33, 58)
svd	2087	13049	972	53.4	4760	63.5	26
tomcatv	2250	1379982621	551	75.5	232833969	83.1	90
urand	204	1563	93	54.4	613	60.1	(0, 18)
zeroin	273	1815	150	45.1	809	55.4	(239, 270)

Static operation

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Benchmark	GA sequence	reduced sequence
adpcm	tonosdnzscno	osnzc
compress	ncsnzosvndvs	nczvnds
dfa	rztonvncodvs	rztcodvs
dhrystone	covolcnocdvc	covlcodvc
fft	ovocdvscnnos	ovcdvs
fmin	tdcotcscnnvo	dcsvo
nsieve	snzdvvcdsvss	nzdvcs
rkf45	ovndtdcvsndd	ovndtcvsd
seval	rnldoncvconv	ldovco
solve	dondvnsdsdcv	ovndcv
svd	odvdvdsnnnss	odvs
tomcatv	dntvvccocvdv	tvcodv
urand	cnottcdtvooc	nodvc
zeroin	rsdosvncnsss	rsosvcs

gene	optimization	
с	cprop	
d	dead	
1	partial	
n	clean	
0	combine	
r	shape	
S	coalesce	
\mathbf{t}	strength	
v	valnum	
Z	lazy	

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compress	ncsnzosvndvs	nczvnds
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dhrystone	covolcnocdvc	covlcodvc
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fmin	tdcotcscnnvo	dcsvo
nsieve	snzdvvcdsvss	nzdvcs
rkf45	ovndtdcvsndd	ovndtcvsd
seval	rnldoncvconv	ldovco
solve	dondvnsdsdcv	ovndcv
svd	odvdvdsnnnss	odvs
tomcatv	dntvvccocvdv	tvcodv
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Z	lazy	

duplicated?

Benchmark	GA sequence	reduced sequence
adpcm	tonosdnzscno	osnzc
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dfa	rztonvncodvs	rztcodvs
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fft	ovocdvscnnos	ovcdvs
fmin	tdcotcscnnvo	dcsvo
nsieve	snzdvycdsvss	nzdvcs
rkf45	ovndtdcvsndd	ovndtcvsd
seval	rnldoncvconv	ldovco
solve	dondvnsdsdcv	ovndcv
svd	odvdvdsnnnss	odvs
tomcatv	dntvvccocvdv	tvcodv
urand	cnottcdtvooc	nodvc
zeroin	rsdosvncnsss	rsosvcs

gene	optimization
с	cprop
d	dead
1	partial
n	clean
0	combine
r	shape
s	coalesce
t	strength
v	valnum
z	lazy

rnldoncvconv

) .				
Benchmark	GA sequence	reduced sequence		gene	optimization		↓
adpcm	tonosdnzscno	osnzc		с	cprop		roldonguconu
compress	ncsnzosvndvs	nczvnds		d	dead		
dfa	rztonuncodus	rztcodys		1	partial		
uiu N	1200nviicouvs	IZCCOUVS		n	clean	ſ	•
dhrystone	covolcnocdvc	covlcodvc		0	combine		nldoncycony
fft	ovocdvscnnos	ovcdvs		r	shape		rldonevconv
fmin	tdcotcscnnvo	dcsvo		\mathbf{S}	coalesce		nuoneveonv
ngiovo	enzduucdevee	nzduce		t	strength		rndoncvconv
INSTEVE	SHZUVVCUSVSS	IIZUVCS		v	valnum	12	
rkf45	ovndtdcvsndd	ovndtcvsd		Z	lazy	12	
seval	rnldoncvconv	ldovco					
solve	dondvnsdsdcv	ovndcv					• valdo pavany
svd	odvdvdsnnnss	odvs					
tomcatv	dntvvccocvdv	tvcodv					
urand	cnottcdtvooc	nodvc					rnidoncvcon
zeroin	rsdosvncnsss	rsosvcs					

	Benchmark	GA sequence	reduced sequence		gene	optimization
1	adpcm	tonosdnzscno	osnzc]	с	cprop
	compress	ncsnzosvndvs	nczynds		d	dead
	J.f		unt as due		1	partial
	dia	rztonyncoavs	rztcodvs		n	clean
	dhrystone	covol <mark>c</mark> nocdvc	covlcodvc		0	combine
	fft	ovocdvscnnos	ovcdvs		r	shape
	fmin	tdcotcscnnvo	dcsvo		s	coalesce
	nsieve	snzdvycdsvss	nzdvcs		t	strength
	mbiove	51124V CubVbb	inzavos		v	valnum
	rkf45	ovndtdcvsndd	ovndtcvsd		z	lazy
	seval	rnldoncvconv	ldovco			
	solve	dondvnsdsdcv	ovndcv			
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	urand	cnottcdtvooc	nodvc			
	zeroin	rsdosvncnsss	rsosvcs			

	↓ rnldoncvconv ↓
	nldoncvconv rldoncvconv rndoncvconv
12	· · r nldoncvcnv rnldoncvcov rnldoncvcon

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urand	cnottcdtvooc	nodvc
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optimization	
cprop	
dead	rnidoncvconv
partial	
clean	¥
combine	pldopovcopy
shape	Πασπενεστιν
coalesce	rldoncvconv
strength	rndoncvconv
valnum	
lazy	•
	-
	rnldoncvcnv
	rnldoncvcov
	rnldoncvcon

...

Benchmark	GA sequence	reduced sequence	ge	ene	optimization		
adpcm	tonosdnzscno	osnzc	с		cprop		roldongugonu
compress	ncsnzosvndvs	nczvnds	d		dead		
dfa	rztonyncodys	rztcodys	1		partial		
uiu	1200m/mcouvb	12000000	n		clean		· · · · · · · · · · · · · · · · · · ·
dhrystone	covolcnocdvc	covlcodvc	0		combine		nldoncycony
fft	ovocdvscnnos	ovcdvs	r		shape		
fmin	tdcotcscnnvo	dcsvo	s		coalesce		ridoncvconv
			t		strength		rndoncvconv
nsieve	snzdvycdsvss	nzdvcs	v		valnum		
rkf45	ovndtdcvsndd	ovndtcvsd	z		lazy		•
seval	rnldoncvconv	ldovco	~				•
solve	dondvnsdsdcv	ovndcv					•
svd	odvdvdsnnnss	odvs		$\overline{}$	<		rnldonevenv
tomcaty	dntwyccocydy	tycody			\sim		rnldoncvcov
comcatv	antiviccocvav	tvecav					rnldoncycon
urand cnottcdtvooc		nodvc				<	rindoneveon
zeroin	rsdosvncnsss	rsosvcs					
							<u> </u>
							•••

GA against fixed sequence

- System default that produces low *dynamic* operation counts
 - rvzcodtvzcod
- Observations of optimization sequences returned by the GA
 nodvcodvs

GA against fixed sequence

	rvzcodtvz	cod	nodvcodvs	
Benchmark	operations	%	operations	%
adpcm	362	3.0	356	1.4
compress	1412	6.7	1325	0.5
dfa	1168	5.2	1145	3.3
dhrystone	574	6.6	544	1.5
fft	1973	10.9	1757	0.0
fmin	203	7.9	198	5.6
nsieve	227	11.0	202	0.0
rkf45	832	10.5	751	0.8
seval	313	8.0	297	3.0
solve	609	28.2	438	0.2
svd	1641	40.8	973	0.1
tomcatv	770	28.4	565	2.5
urand	93	0.0	93	0.0
zeroin	158	5.1	154	2.6

	rvzcodtv	zcod	nodvcodvs	
Benchmark	operations	%	operations	%
adpcm	11965360	-2.7	12440360	1.2
compress	7494982	26.0	5548268	0.1
dfa	546396	9.2	511349	3.0
dhrystone	3390200	5.6	3270193	2.1
fft	15088485	3.4	14574279	0.0
fmin	955	-0.8	947	-1.7
nsieve	570608938	5.4	554450986	2.6
rkf45	214146	5.9	215620	6.6
seval	809	-4.1	985	14.5
solve	1099	6.4	1033	0.4
svd	4731	-0.6	4758	-0.0
tomcatv	186355888	-24.9	239463612	2.8
urand	631	2.9	613	0.0
zeroin	829	2.4	811	0.2

GA against fixed sequence

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Conclusion

- Program-specific optimization sequence
- Reduction of code size up to 40%

Discussion

- Compare to known optimization flag O2, O3
- Convergence too slow
- Different fitness function
 - Execution time, power consumption