Alloy Program Synthesizer with Genetic Algorithm (APS-GA) as part of the Program Synthesis Model Finder (PSMF) project

Installing and running procedures (version 2023-February)

Purpose

Program synthesis aims to mechanize the task of programming. We built a PSMF system to extend the synthesis competency of a general purpose imperative program synthesizer, which works on the integers' domain, accepting as specification sketches and a tiny set of user examples. The PSMF system embeds into the synthesizer some concepts of genetic algorithms and genetic programming's mutations. So far, the system has helped the synthesizer to perform better the synthesis task of programs that itself already had been able to and to discover new ones, reducing the user's intervention at different stages of the process.

For Windows Operating Systems

1. Unzip the psmf.zip file to any directory of your disk. For example to root directory C:/

Files into those folders (table below):

Folder	Files	Description							
psmf	PSMF.jar, README.TXT	README.TXT PSMF program's main file, and read me file.							
lib	AlloyAlsToXML.jar, hola-0.2.jar, jgap.jar	Mandatory libraries to execute PSMF program.							
alloyFiles	imp.als	ain file of the Alloy* synthesizer.							
impMs	semantics.als, syntax.als, wellF.als	Complementary files of the Alloy* synthesizer.							
	-	Alloy* model files (.als), test cases file (.txt), and configuration files (.cfg) have to be over here.							

- 2. The current folder or the folder ../alloyFiles/ have to be in the Path Environment Variables. Here is a how to do that in windows 10! You will able to run PSMF in *Linux* or *MacOS*, just configuring the Path Environment Variables in such Operation Systems.
- 3. Setting up precondition and post-condition on an Alloy* model file. For example open up p02max3.als into ../psmf/alloyFiles/examples/ folder.
 - 3.1. Have a note that setting up precondition and post-condition contract might be not mandatory, its purpose is to constraint the search space. In most cases, setting up the scope of the input/output variables (see examples at Section 3.2) would be enough to able the synthesizer to find out a solution. In other cases the user may be leave precondition and post-condition empty.
 - 3.2. Write down sketch over the predicate PreC (or PosC), for example:

4. Preparing Test Cases Dataset. Note that the .txt filename must have the same of the Alloy* model. For example for Max3 program synthesis, the chosen Alloy* model file was p02max3.als, so the test cases dataset file must be p02max3.txt. Then, open up the and fill it with any set o test cases you want (try to fill the file using input/output values that covers as many as possible cases for testing). Note that each line has four values separated by semicolon symbol, and they mean, respectively: expected output; first input variable value (i.e. x variable); second input variable value (i.e. y variable); and third input variable value (i.e. z variable). The content of the p02max3.txt, is shown as follows:

```
6;2;6;1
0;0;0;0
-1;-1;-1;-1
0;-1;0;-3
0;-1;-1;0
4;2;1;4
```

5. Preparing PSMF configuration file. Note that the .cfg filename can be any, for example psmf.cfg. Then, open up the file and fill it with parameters values you want to execute the task of synthesis. The file content is self explained. The content of the psmf.cfg, is shown as follows. It might be better open the psmf.cfg file on the examples folder copy and rename it to make any changes you want:

```
//Configuration file for Program Synthesis via Model Finder - PSMF system
//TO adjust System's parameters.
// A pair of slashes stands for comments.
//Inform the alloy file name (whitout extension name, for example:
 ok06max2).
InputFileName = p02max3;
//Inform the alloy file extension (including the dot symbol, for
 example: .als).
InputFileExtension = .als;
//Inform number of executions that the system will perform, for simulation
//For example: NumSysExec=5; means the user will call the system once, and
 it will perform 5 times.
NumSysExec=1;
//Distance Metric: explore all available [eAll] or read what is set into
  this configuration file [rFile]?
DistanceMetricExploreAllOrReadConfFile = rFile;
//The name of the distance metric: discrete | manhattan |
  euclidean quadratic | hellinger | chebyshev | minkowski
DistanceMetricType=discrete;
//... there are a lot of parameter, please have a look at the configuration
  file in the examples folder.
//For all signature's scope values these values are accepted:
// lower:upper bound value pair (for example 1:3 or any other zero or
 positive integer pair)
// auto (the system will define automatically)
// no
          (the system will not take account of such signature)
IntVar = 4:4; IntVal = auto; Assign = auto; SComp = auto; CondS = auto;
  While = 0:0; Add = 0:0; Sub = 0:0; Mult
                                                  = 0:0; EQ
                                                                = 0:0; NEQ
  = 0:0; LEQ
                              = auto; GTH
                = auto; GEQ
                                              = 0:3:
//end of file.
```

6. Run the system typing java command line at ../psmf folder. If you have memory issues, you may use the -Xms java parameter to allocate at least 1GB of RAM memory by typing -Xms1q:

```
java -jar -Xms1g "PSMF.jar"
```

7. Then the PSMF system will ask you three questions: sp01 just press enter if you do not want to change the default configuration folder name; sp02 just press enter if you do not want to change the default configuration file extension name; and sp03 just press enter if you do not want to change the default configuration filename.

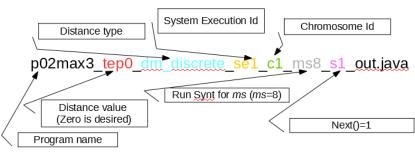
```
sp01 -> Inform the config file's folder name. Press enter for default value
  [c:/data/psmf/alloyFiles/examples/]:
sp02 -> Inform the config file's extension name. Press enter for default
  value [.cfg]:
sp03 -> Inform the config file name. Press enter for default value [psmf]:
```

8. PSMF system's outcomes. The PSMF system used write out two kind of files:

Folder	File	Description				
psmf	ibuzmaxa rebu dm discrete sel ci msk si olir.lava	program candidate synthesized source code.				
examples	p02max3_out.als	Alloy* model file.				

8.1. A set of Java program candidates synthesized source code — despite the source code, those filenames carry out the following information: (i) the input filename program, which was max3 (p02max3); (ii) the name of the distance metric (_dm_), for example _dm_discrete means discrete metric was used to calculate the fitness function; (iii) The number of executions that the system will perform within the same session (_se), for instance _se5 means the user did call the system once, and it was perform 5 times; (iv) The chromosome id (_c), for instance _c5, means

the program candidate was the chromosome number five; (v) The max sequence (_ms), for instance _ms8, means the program candidate has used the synthesizer command using the max sequence equals to eight, i.e. to run the Alloy* synthesizer directly on the



Alloy Analyzer interface the user have to execute the run Synt for 8 but... command; (vi) The subset (_s), for instance _s3, means the program candidate has used the next command equals to three, i.e. to run the Alloy* synthesizer directly on the Alloy Analyzer interface the user have to execute the next command three times after the run Synt for 8 but... command has been invoked; (vii) the calculated distance between the program candidate's output and the test cases (_tep). As less the number after _tep as better, for example _tep10 is better than _tep63. The _tep0 (Zero) is the desired situation which means there were no errors (distance=zero);

- 8.2. The Alloy* model file due the file p02max3.als does not have the run Synt for... command, this outcome file p02max3_out.als does. In other words, the PSMF system generates a text file with the Alloy* model implementation, where the run Synt for command (with all those signatures) are ready for execute directly on the Alloy Analyzer interface, if the user wants to.
- 9. Experimental benchmark results concerning 17 well-known programs using the parameters in the configuration files (*.cfg).

			p03														
Prog	Max2	Max3	Max4	Maj3	Maj5	Add2	Add3	Add4	Fib	Mult	IntS QRT	Fact	Modu	SumS QR	Medi an	GCD	Maj8