



# Teach Your Computer to Code FizzBuzz

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<https://pragprog.com/book/fbmach/genetic-algorithms-and-machine-learning-for-programmers>

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Workshop materials can be found at

<http://www.fet.uwe.ac.uk/~clsimons/ACCU2019/>

JCLEC requires Java SE Development Kit, e.g. version 8

<https://www.oracle.com/technetwork/java/javase/overview/index.html>

# Workshop

Evolutionary computing

Frameworks for evolutionary computing

Java Class Library for Evolutionary Computing (JCLEC)

Optimisation problems:

- 1 - ‘OneMax’ Problem
- 2 - How to program your way out of a paper bag
- 3 - FizzBuzz

Genetic Programming, Genetic Improvement

# Slides

Evolutionary computing

Frameworks for evolutionary computing

# Coding

Optimisation problems:

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- 3 - FizzBuzz

# Slides

Genetic Programming, Genetic Improvement

# When we want to optimise:

- We might want to maximise ‘quality’, and/or
- We might want to minimise ‘cost’...

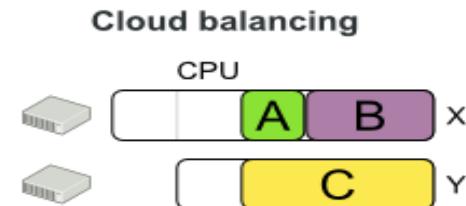
We want to find the ‘best’ (for whatever ‘best’ means for our situation)

# So what's the problem?

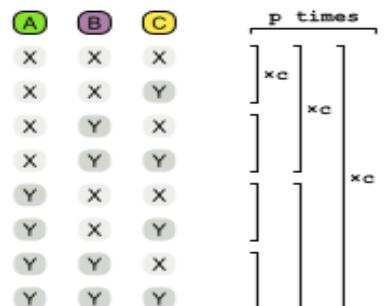
*combinatorial explosion*

## Calculate the size of the search space

Given a Solution model, how many different combinations can it represent?

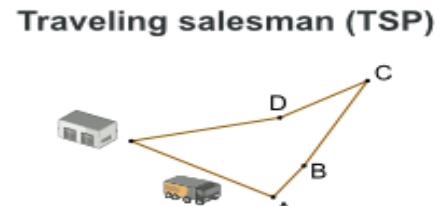


Model: Computer → Process

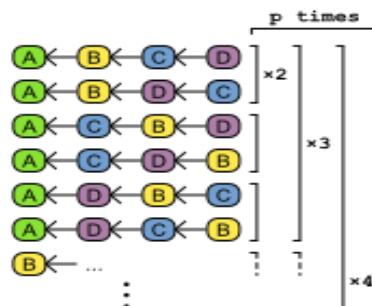


Search space:  $c^p$

# computers	# processes	search space
2	3	8
100	300	$10^{600}$
200	600	$10^{1380}$
400	1200	$10^{6967}$

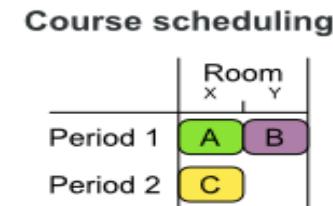


Model: linked list

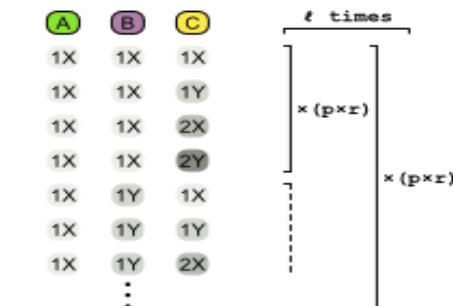


Search space:  $n!$

# customers	search space
4	24
100	$10^{157}$
1000	$10^{2567}$
10000	$10^{35659}$



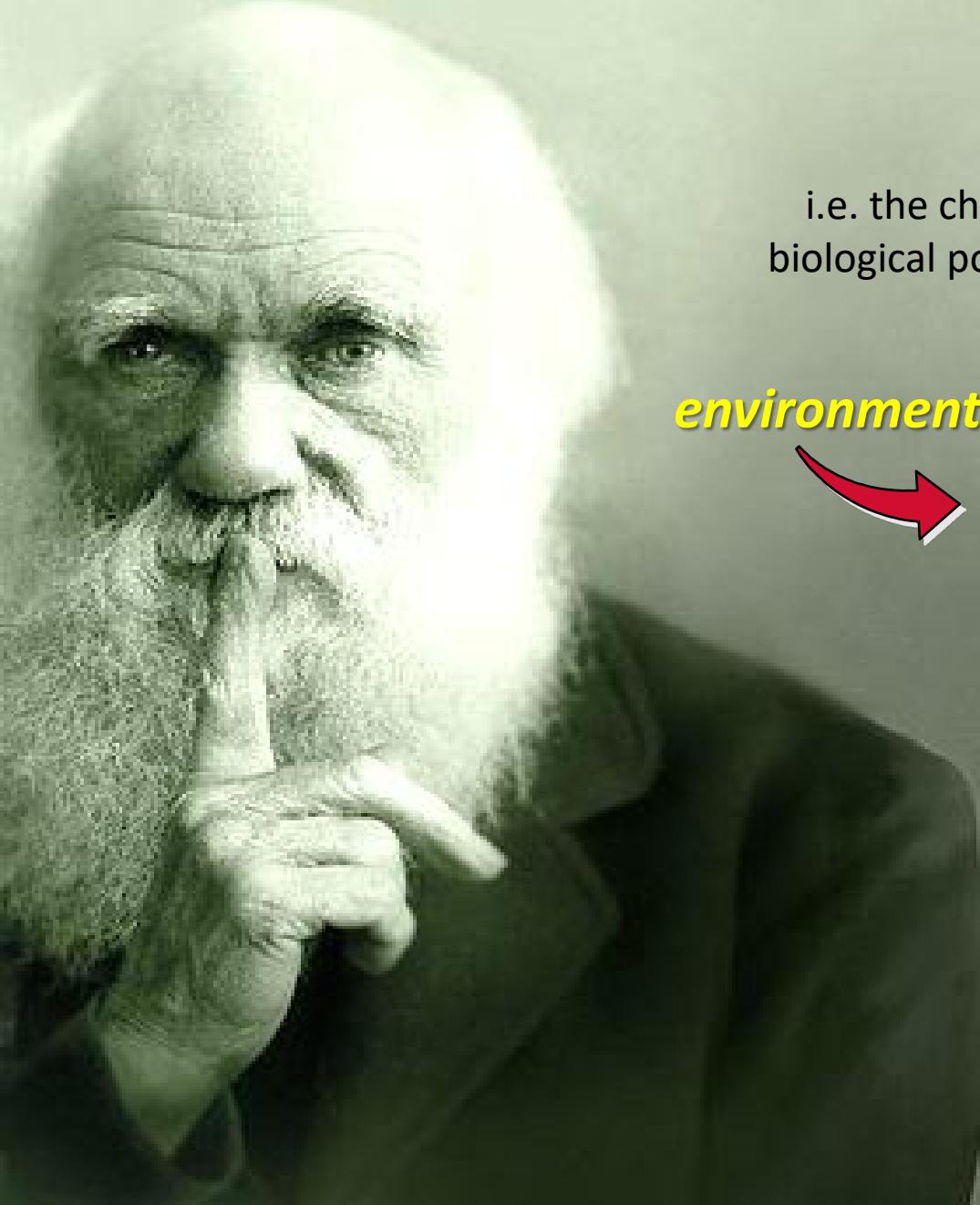
Model: Period → Room



Search space:  $(pxr)^t$

# periods	# rooms	# lectures	space
2	2	3	64
36	6	100	$10^{233}$
36	18	400	$10^{1124}$
36	36	800	$10^{2490}$

# What can we do about it?



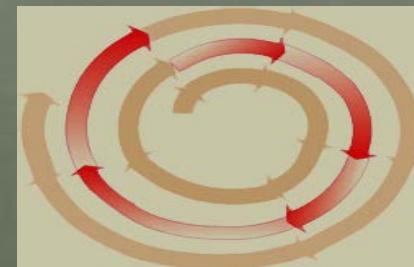
## natural evolution

i.e. the change in the inherited characteristics of biological populations over successive generations.

*environment*



Selection of fittest individuals



sexual reproduction for  
population diversity / variety

# Computational evolution

**Representation** of an “individual” solution  
e.g. arrays, trees, models, code etc. etc.

```
initialise population at random
evaluate each individual
while( not done )
    select parents
    recombine pairs of parents
    mutate new candidate individuals
    evaluate each individual
    select candidates for next generation
end while
```

# Ideas from biology (1)

Information concerning the characteristics of a solution *individual* is encoded in ‘genes’ – all the gene values of an individual is known as the *genotype*.

Typically, many individuals make up a *population*.

Individuals can become *parents* from whom *offspring* are created. The offspring help to form the new *generation*, and can themselves become parents in the next generation.

Evolutionary algorithms can run for many generations, until some *termination condition*.

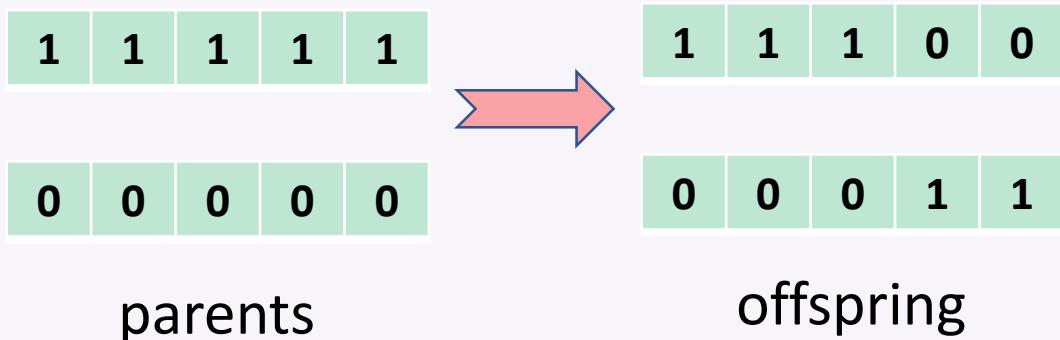
# Ideas from biology (2)

**Evaluation** of a solution **individual** gives some **fitness** value or **cost** value that is to be optimised, either **maximised** or **minimised**.

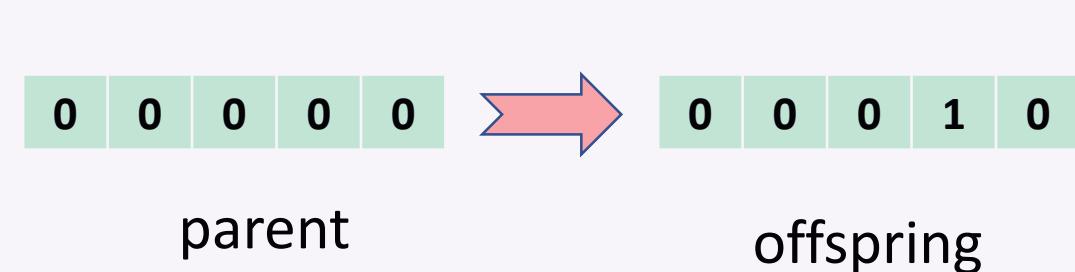
Only the fittest solution **individuals** are **selected** to breed **offspring**; individuals can enter a **tournament**, the fittest wins the right to breed.

**Diversity** in the **population** is maintained by:

**Recombination** (sexual reproduction)



**Mutation** (asexual reproduction)



# Many applications of Evolutionary Computing

Examples include many well-known optimisation problems such as

- course timetabling,
- nurse rostering,
- process scheduling,
- network routing,
- vehicle delivery scheduling,
- load balancing,
- Etc. etc.



The 2006 NASA ST5 spacecraft antenna. This complicated shape was found by an evolutionary computer design program to create the best radiation pattern.

[https://en.wikipedia.org/wiki/Evolved\\_antenna](https://en.wikipedia.org/wiki/Evolved_antenna)

# Workshop

Evolutionary computing

**Frameworks for evolutionary computing**

Java Class Library for Evolutionary Computing (JCLEC)

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- 2 - How to program your way out of a paper bag
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Genetic Programming, Genetic Improvement

# Frameworks for Evolutionary Computing

Characteristics include:

- mechanisms for the integration of problem-specific knowledge, such as problem constraints and fitness function(s);
- components to configure and monitor the execution, allowing the user to set execution parameters and visualise intermediate results;
- designed with *best practices* and *design patterns* in mind.

Language	Framework	version	Date
C++	Evolutionary Computation Framework (ECF)	1.4.2	2017
	Evolving Objects (EO)	1.3.1	2012
	jMetalCpp	1.7	2016
	Mallba	2.0	2009
	Open Beagle	3.0.3	2007
	OptFrame	2.2	2017
	PaGMO	2.5	2017
Java	ParadiseO	2.0.1	2012
	Java-based Evolutionary Computation Research System (ECJ)	24.0, 25.0	2017
	Evolutionary Algorithms Workbench (EvA)	2.2.0	2015
	Java Class Library for Evolutionary Computation (JCLEC)	4.0	2014
	jMetal	5.3	2017
	Multi-Objective Evolutionary Algorithm (MOEA) Framework	2.12	2017
	Opt4J	3.1.4	2015
C#	GeneticSharp (A C# Genetic Algorithm Library)	On-going	2017
	HeuristicLab (A Paradigm-Independent and Extensible Environment for Heuristic Optimization)	3.3.14	2016
Python	Distributed Evolutionary Algorithms in Python (DEAP)	1.1.0	2017
	jMetalPy	On-going	2017
	Pyevolve	0.6rc_1	2015
	PyGMO	On-going	2017
Matlab	Pyvolution	1.1	2012
	Genetic and Evolutionary Algorithm Toolbox for Matlab (GEATbx)	3.8	2017
	Global Optimisation Toolbox	R2017b	2017
	Matlab Platform for Evolutionary Multi-objective Optimisation (PlatEMO)	1.3	2017

Many frameworks available!

For further details, see Overload 142

<https://accu.org/index.php/journals/c380/>

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# Time to look at an example of a evolutionary computing framework

The screenshot shows the homepage of the JCLEC website. At the top left is the JCLEC logo, which includes a circular design with the text "Class Library for Evolutionary Computation - JCLEC". The main title "JCLEC - Java Class Library for Evolutionary Computation" is centered above a detailed description of the software. Below the description is a bulleted list of features. On the left side, there's a "MAIN MENU" with links to Home, Features, Documentation, Examples, Download, VisualJCLEC, and EC Software Links. There's also a "LATEST NEWS" section with a link to a tutorial and information about the classification module.

**JCLEC - Java Class Library for Evolutionary Computation**

JCLEC is a software system for Evolutionary Computation (EC) research, developed in the Java programming language. It provides a high-level software framework to do any kind of Evolutionary Algorithm (EA), providing support for genetic algorithms (binary, integer and real encoding), genetic programming (Koza's style, strongly typed, and grammar based) and evolutionary programming.

JCLEC architecture follows strong principles of object oriented programming, where abstractions are represented by loosely coupled objects and where it is common and easy to reuse code. JCLEC provide an EC environment with the following main features:

- **Genericity.** Any kind of EC algorithm can be executed using JCLEC, as some minimum requirements are fulfilled. The only necessary condition is to have a population of individuals to which a sequence of evolving operations is iteratively applied. So far, JCLEC supports most mainstream EC flavors such genetic programming, bit string, integer-valued vector and real-valued vector genetic algorithms, and evolution strategy. It also supports advanced EC techniques such as multi-objective optimization. The user can make use of any of these specialized frameworks, or even modify them to create his/her own customized evolutionary algorithm.
- **User friendly.** Considerable efforts were made to build JCLEC in a pretty and pleasant form. JCLEC provides diverse mechanisms that offer a user friendly programming interface. It follows a high-level programming style, which allows the rapid prototyping of applications.
- **Portability.** The JCLEC system has been coded in the Java programming language that ensures its portability between all platforms that implement a Java Virtual Machine.

<http://jlec.sourceforge.net>

# Time to download the framework <http://jclec.sourceforge.net>

The screenshot shows the JCLEC project page on SourceForge. On the left, there's a sidebar with a logo, a main menu (Home, Features, Documentation, Examples, Download, VisualJCLEC, EC Software Links), and a latest news section. A red arrow points from the 'Download' link in the main menu to the 'Downloads' section of the page. The 'Downloads' section lists various ways to obtain the source code, including SourceForge, SVN, GIT, CVS, Eclipse, and NetBeans. It also provides instructions for starting testing and mentions the JCLEC wiki, tutorial, and classification tutorial. A large red arrow points from the 'Download' link in the sidebar to the 'Download Latest Version' button on the right. The main content area has tabs for Summary, Files, Reviews, Support, Mailing Lists, Bugs, and News. The 'Files' tab is active, showing a list of files. A green button labeled 'Download Latest Version jclec4-base.zip (3.7 MB)' is highlighted with a red arrow. Below it is a blue 'Get Updates' button. The file list table has columns for Name, Modified, and Size. It contains three items: 'jclec4-base.zip' (3.7 MB, modified 2014-07-03), 'jclec4-classification.zip' (6.5 MB, modified 2014-07-03), and 'jclec4-tutorial.zip' (1.6 MB, modified 2013-04-18). The total size is 11.8 MB. At the bottom, it says 'Totals: 3 Items'. To the left of the file list, there are three annotations with arrows: 'We need this FIRST' pointing to the 'jclec4-base.zip' row, 'But we don't need this' pointing to the 'jclec4-classification.zip' row, and 'And we need this SECOND' pointing to the 'jclec4-tutorial.zip' row.

We need this **FIRST** ➔

But we don't need this ➔

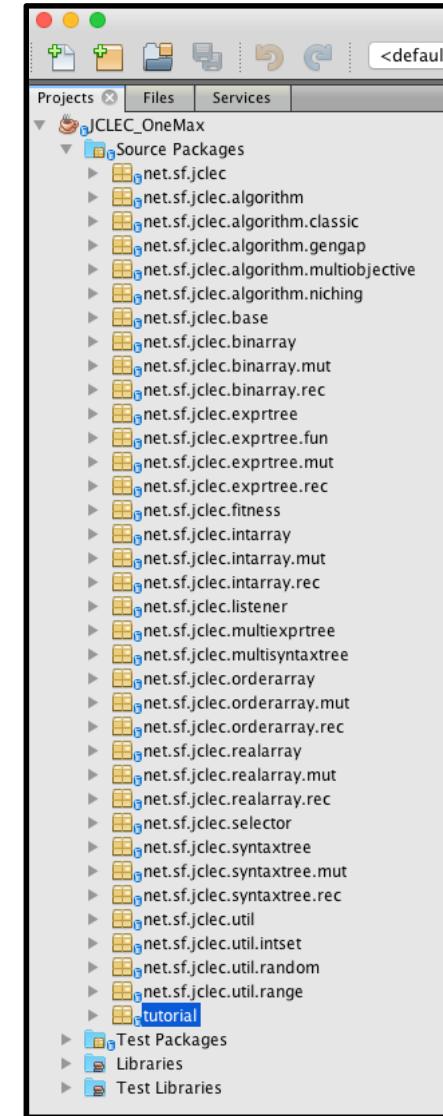
And we need this **SECOND** ➔

Name	Modified	Size
jclec4-base.zip	2014-07-03	3.7 MB
jclec4-classification.zip	2014-07-03	6.5 MB
jclec4-tutorial.zip	2013-04-18	1.6 MB

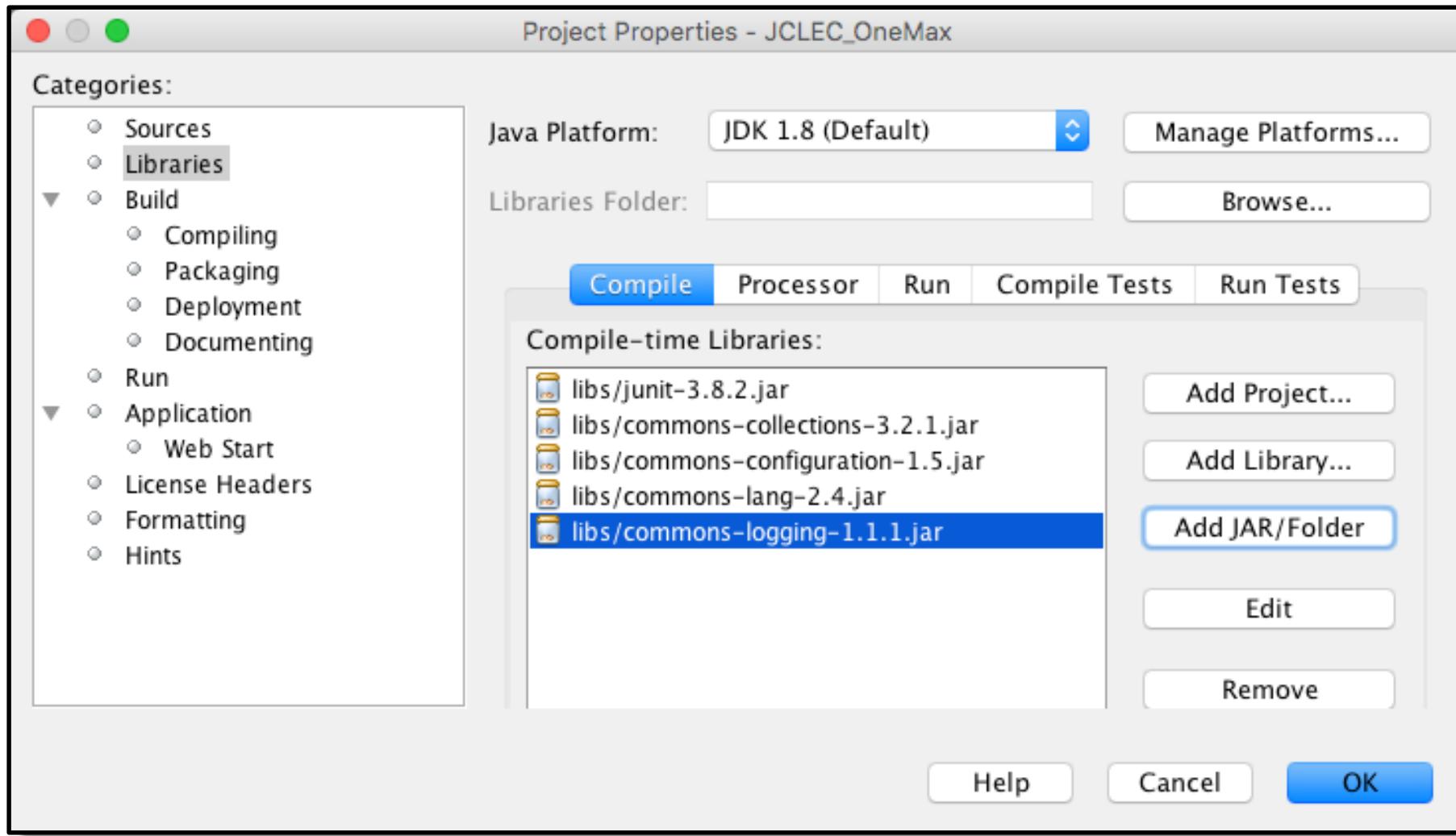
Totals: 3 Items 11.8 MB

You can copy an Eclipse or NetBeans project, or

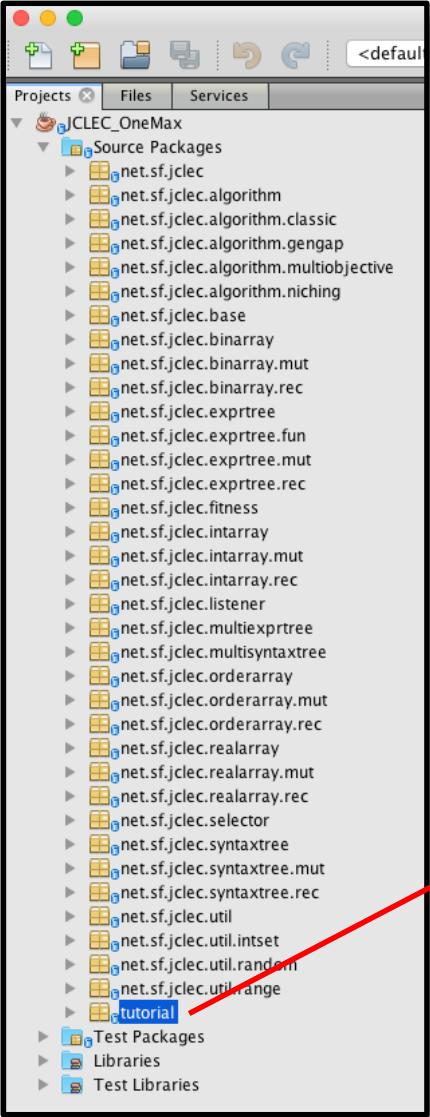
From [jclec4\\_base.zip](#), copy extracted JCLEC source files to an IDE of your choosing, e.g.



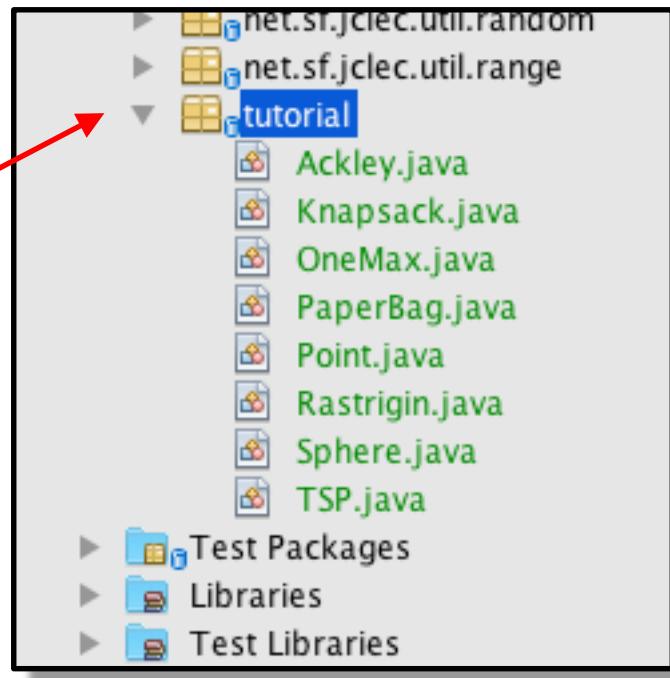
From **jclec4\_base.zip**, extract and let the IDE know about the required libraries, e.g.



## From **jclec4-tutorial.zip**,



place tutorial source files  
in a package called ‘tutorial’

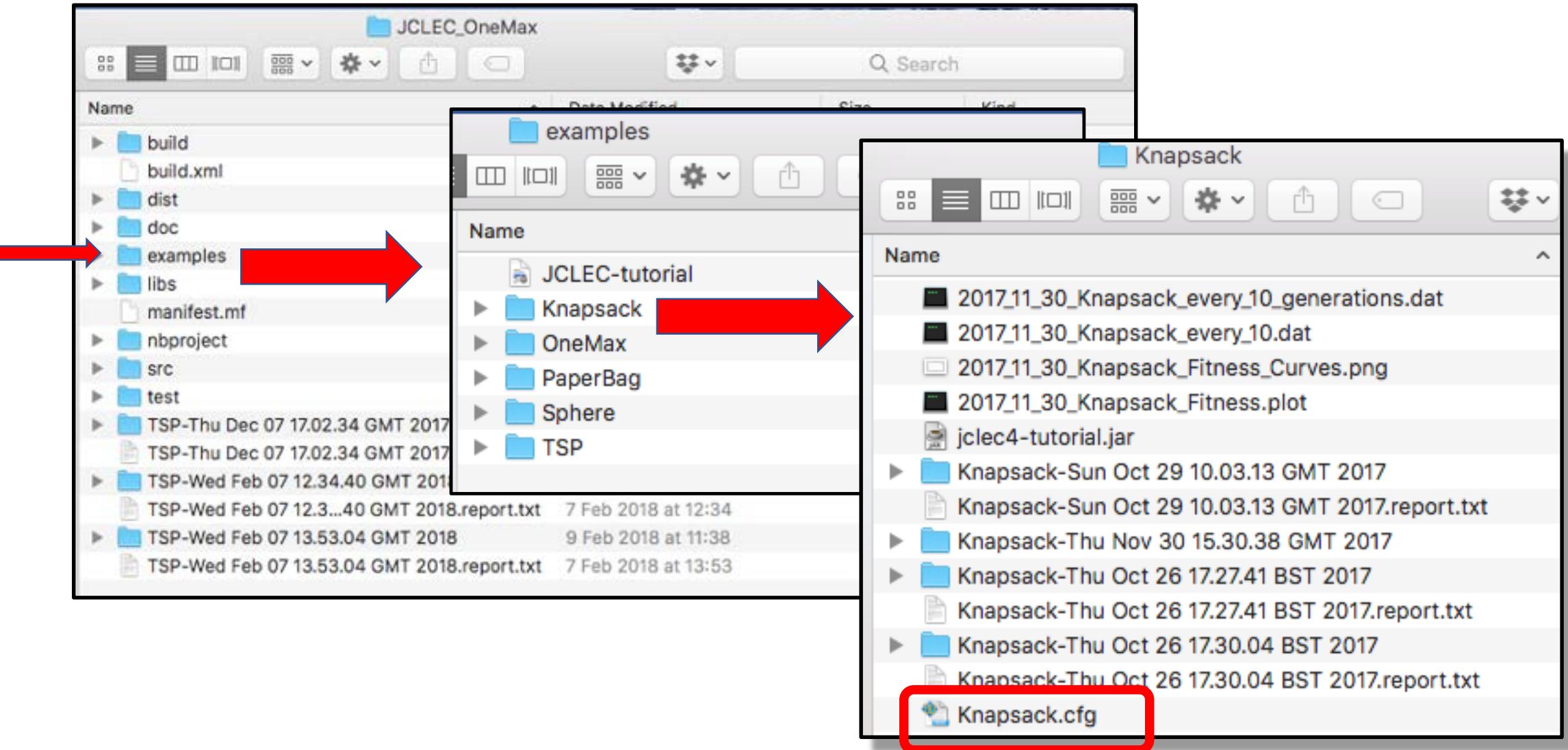


Each JCLEC project has an XML-based configuration file to specify:

1. Evolutionary algorithm components used, and
2. parameter set up

XML configuration files have a .cfg suffix

Also from [jlec4-tutorial.zip](#), copy tutorial example configuration files to a folder called 'examples', e.g.



# Note

- For the Travelling Salesman Problem (TSP) example, you may also need `orderarray` package from [\*\*jlec4-tutorial.zip\*\*](#)

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Genetic Programming, Genetic Improvement

The "hello world" of evolutionary computing!

The OneMax problem consists of maximising the number of ones in a bitstring.

Let's take a length of 100 bits for the bitstring.

e.g. <http://tracer.lcc.uma.es/problems/onemax/onemax.html>

Yes, I know, we can do this in our heads : ) but it's a good example of getting going with the framework...

# Algorithm design and parameter set up – let's apply some patterns...

## **Representation**

- how to encode a candidate solution?

a binary array

**Initialisation:** random

**Evolution:** simple generational with elitism (SGE)

## **Fitness**

- how to evaluate the fitness of a candidate solution?

count the number of 1s

... and suggested parameters

**Population size:** 100 individuals

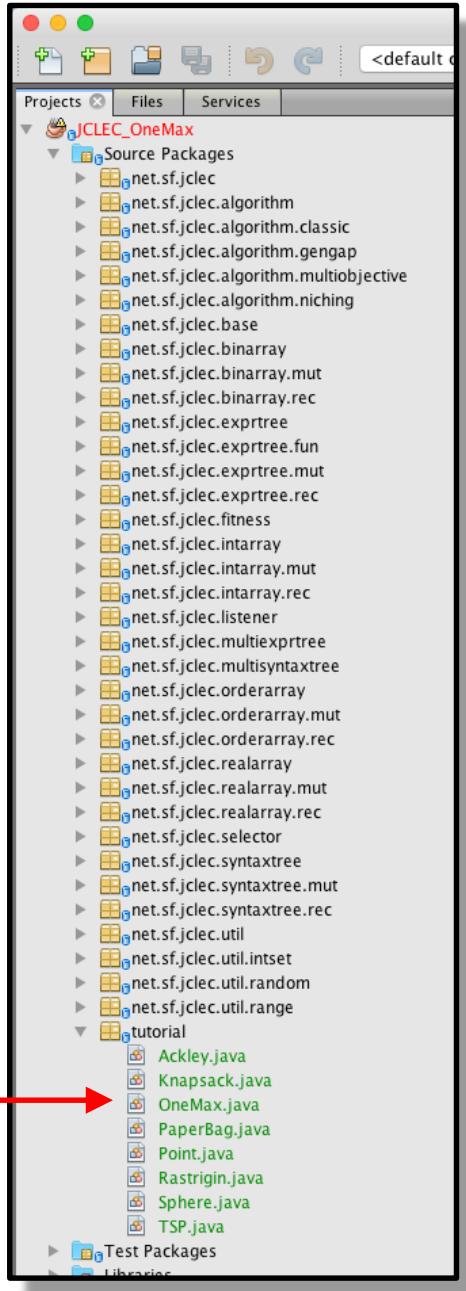
**Stop Criterion:** 50 generations

## **Diversity**

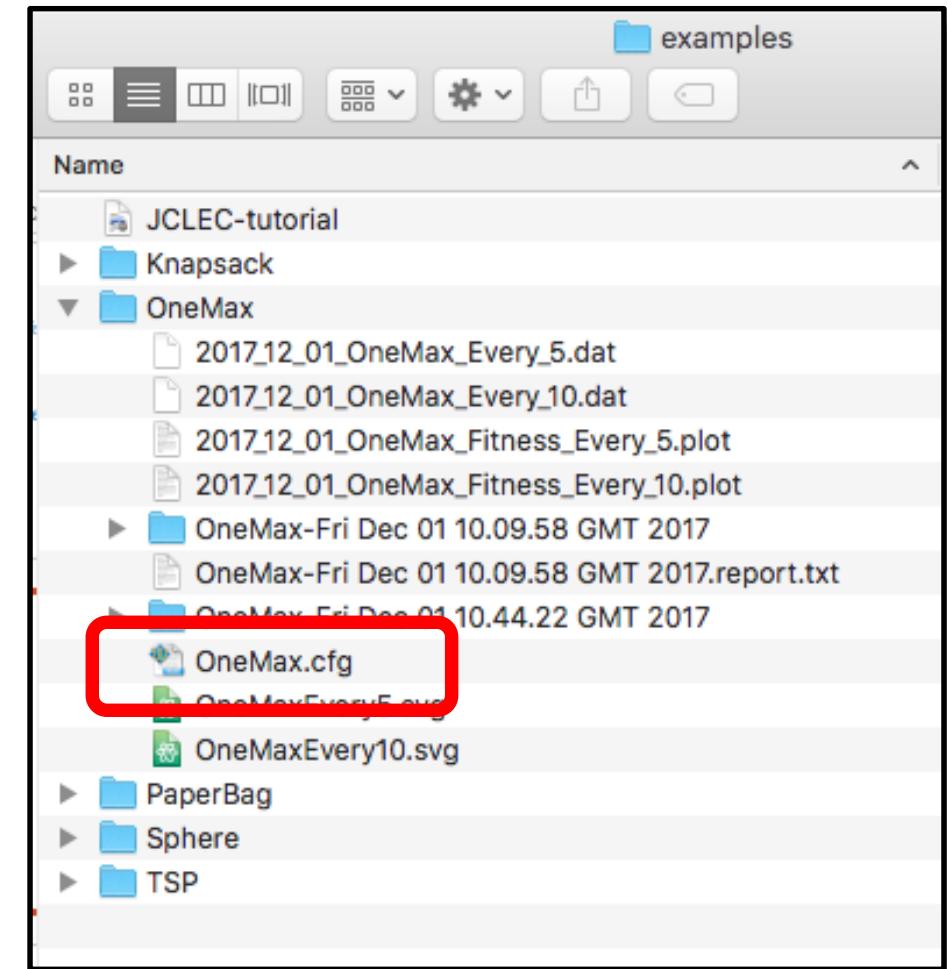
- how to make offspring different to parents?

crossover and mutation

**Parent selection:** tournament of 2 individuals

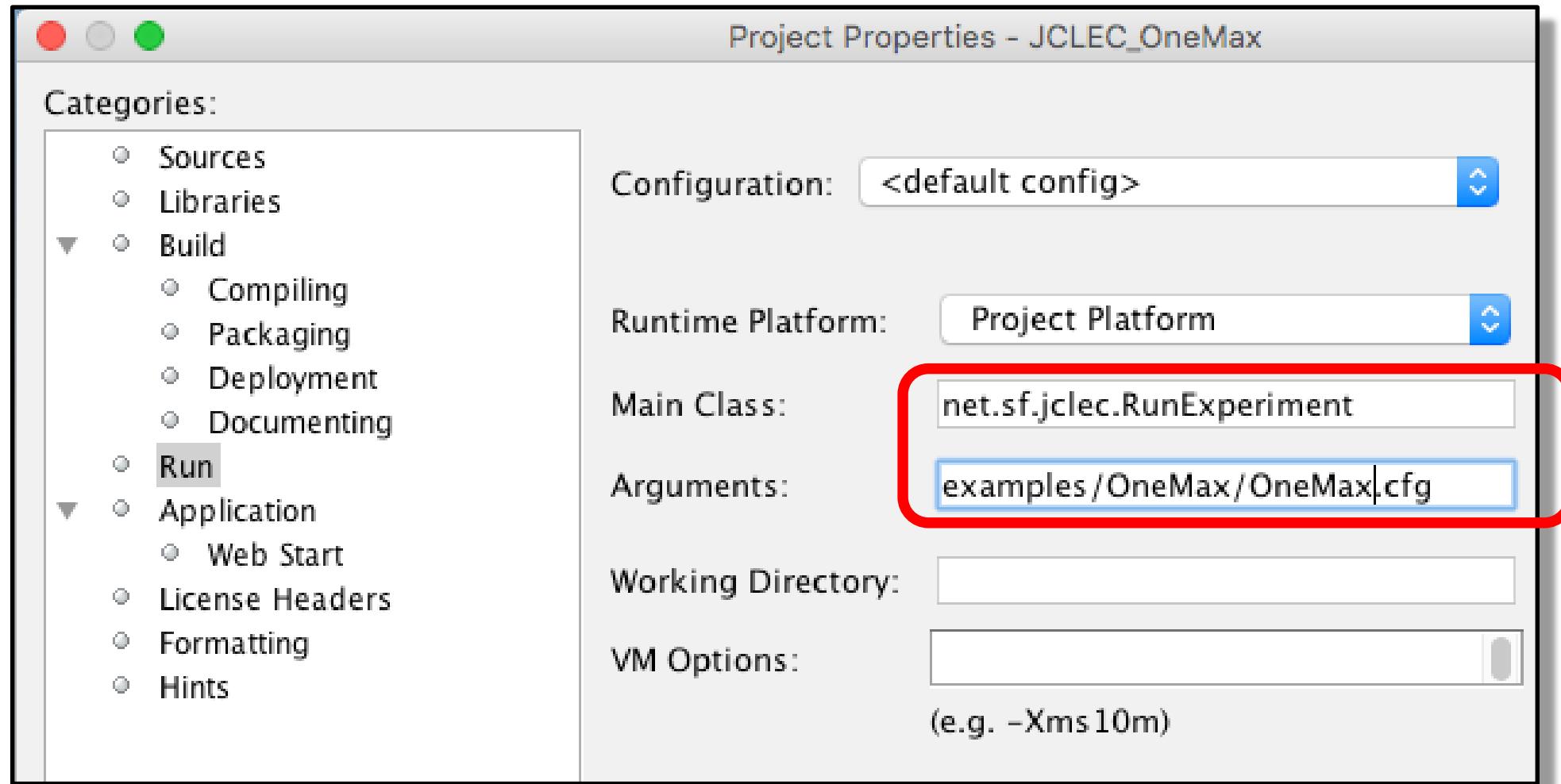


And make a new folder “OneMax” in “examples” for the configuration file – “OneMax.cfg”



Make a new file  
“OneMax.java”

Don't forget to invoke the executable with "OneMax.cfg" as an argument



# Enjoy the programming!

Here's one way to solve the OneMax optimisation problem...

Here's one possible configuration:

```
<experiment>
  <process algorithm-type="net.sf.jclec.algorithm.classic.SGE">
    <rand-gen-factory type="net.sf.jclec.util.random.RanecuFactory" seed="987328938" />
    <population-size>100</population-size>
    <max-of-generations>50</max-of-generations>
    <species type="net.sf.jclec.binarray.BinArrayIndividualSpecies" genotype-length="100" />
    <evaluator type="tutorial.OneMax" />
    <provider type="net.sf.jclec.binarray.BinArrayCreator" />
    <parents-selector type="net.sf.jclec.selector.TournamentSelector">
      <tournament-size>2</tournament-size>
    </parents-selector>
    <recombinator type="net.sf.jclec.binarray.rec.UniformCrossover" rec-prob="0.9" />
    <mutator type="net.sf.jclec.binarray.mut.OneLocusMutator" mut-prob="0.2" />
    <listener type="net.sf.jclec.listener.PopulationReporter">
      <report-frequency>5</report-frequency>
      <report-on-file>true</report-on-file>
      <save-complete-population>true</save-complete-population>
      <report-title>"OneMax-</report-title>
    </listener>
  </process>
</experiment>
```

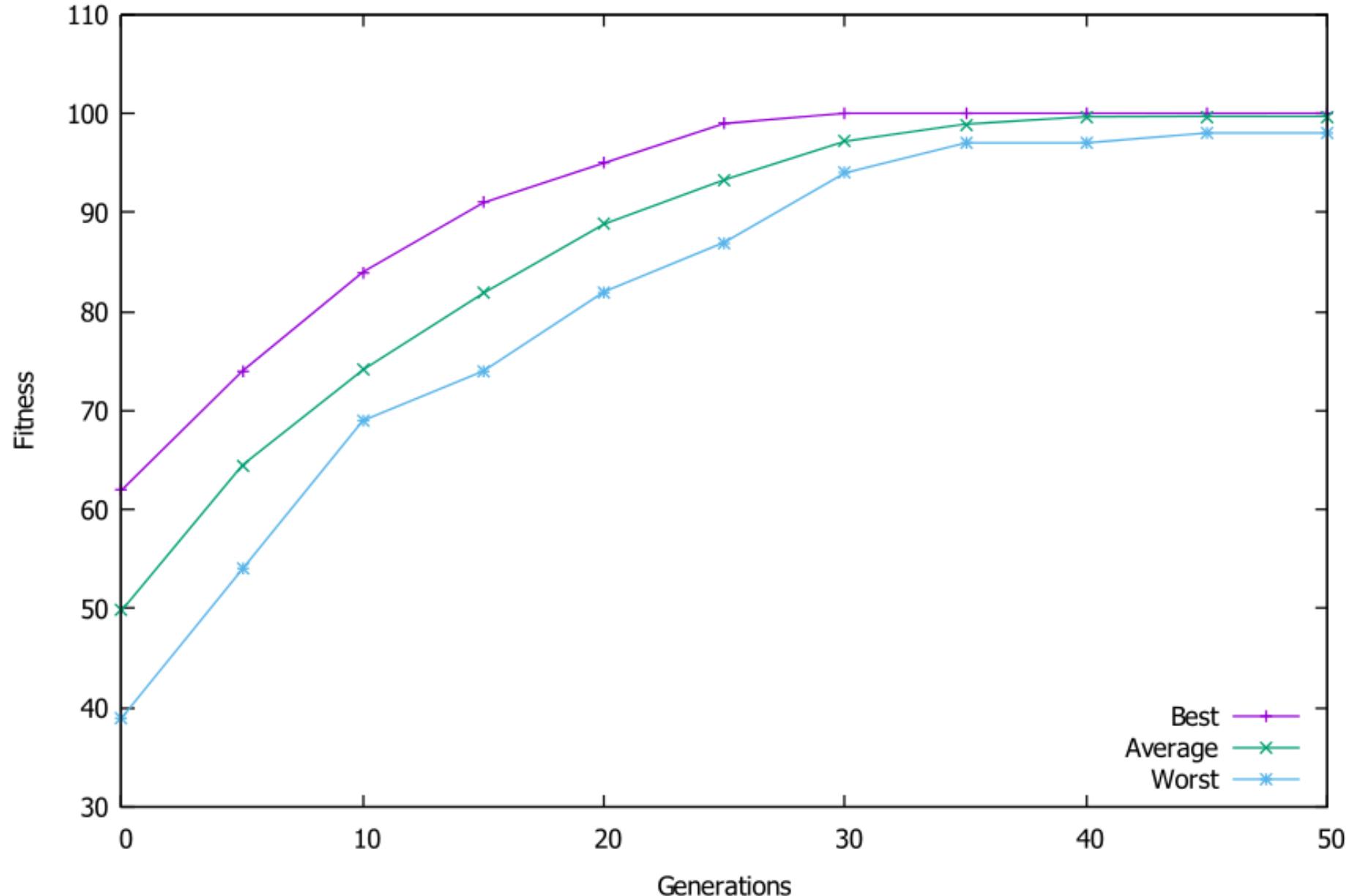
## One way of solving the fitness evaluation:

```
@Override  
protected void evaluate( IIndividual ind ) {  
    // Individual genotype  
    byte[ ] genotype = ( (BinArrayIndividual)ind ).getGenotype( );  
    int bitCount = 0;  
  
    for( int i = 0; i < genotype.length; i++ ) {  
        if( genotype[ i ] == 1 ) {  
            bitCount++;  
        }  
    }  
    ind.setFitness( new SimpleValueFitness( bitCount ) );  
}
```

# Demonstration



OneMax example fitness curves



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**Optimisation problems:**

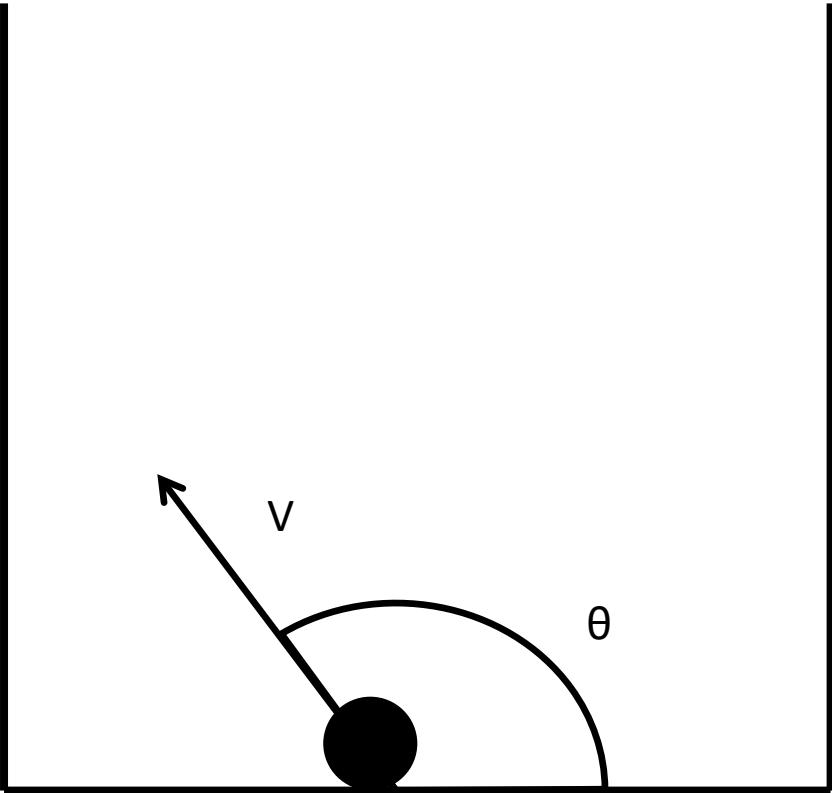
1 - 'OneMax' Problem

**2 - How to program your way out of a paper bag**

3 - FizzBuzz

Genetic Programming, Genetic Improvement

Let's suppose there's a canon in a paper bag



Let's also suppose:  
width of bag is 10.0,  
height of bag is 5.0.

- *Overload*, 21(118):7–9, December 2013
  - <http://accu.org/index.php/journals/1821>

Given a bag with bottom left corner at  $(k, 0)$ , of width  $w$ , and height  $h$ , assuming the projectile is smaller than the bag, the cannon is a point of no size, and given the acceleration due to gravity,  $g$ , after time  $t$  the projectile will be at point  $(x, y)$  where

$$x = k + \frac{1}{2}w + vt \cos \theta$$

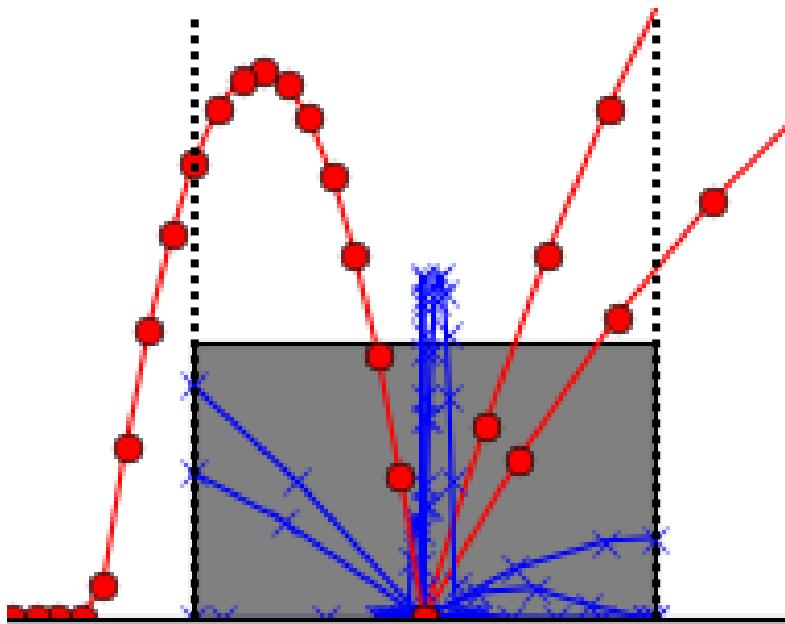
$$y = vt \sin \theta - \frac{1}{2}gt^2$$

$x$  is the horizontal displacement and  $y$  the vertical displacement. The projectile will just escape when  $y \geq h$  and  $x < k$  or  $x > k + w$ .

$g$  will be taken as 9.81 m/s<sup>2</sup>. For simplicity, the code will assume  $k$  is zero.

# Fitness evaluation

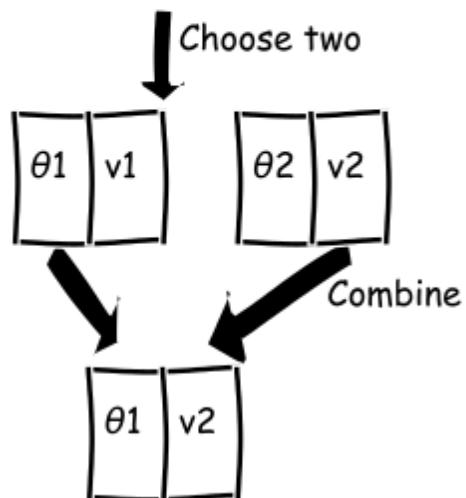
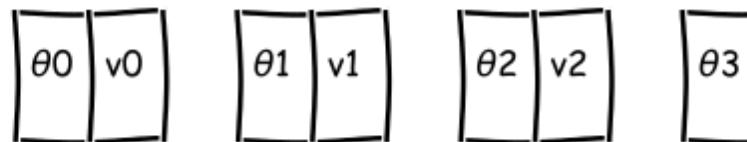
- Launching at random, something either ends in or out of the bag
- But some fail cases are less bad than others



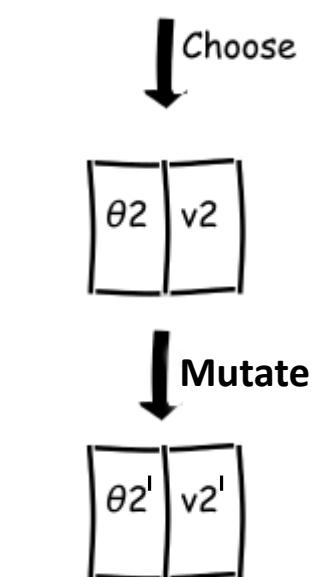
- 3 escape
- 2 on left get “close”
- Could “close” mean height  
(at edge of bag)?
- Fitness = height

# Diversity Preservation

## Recombination



## Mutation



# Algorithm design and parameter set up – let's again apply some patterns...

## **Representation**

- how to encode a candidate solution in the population?

?

## **Fitness**

- how to evaluate the fitness of a candidate solution?

?

## **Diversity**

- how to make offspring different to parents?

?

**Initialisation:** random?

**Evolution:** simple generational with elitism (SGE)?

... and suggested parameters

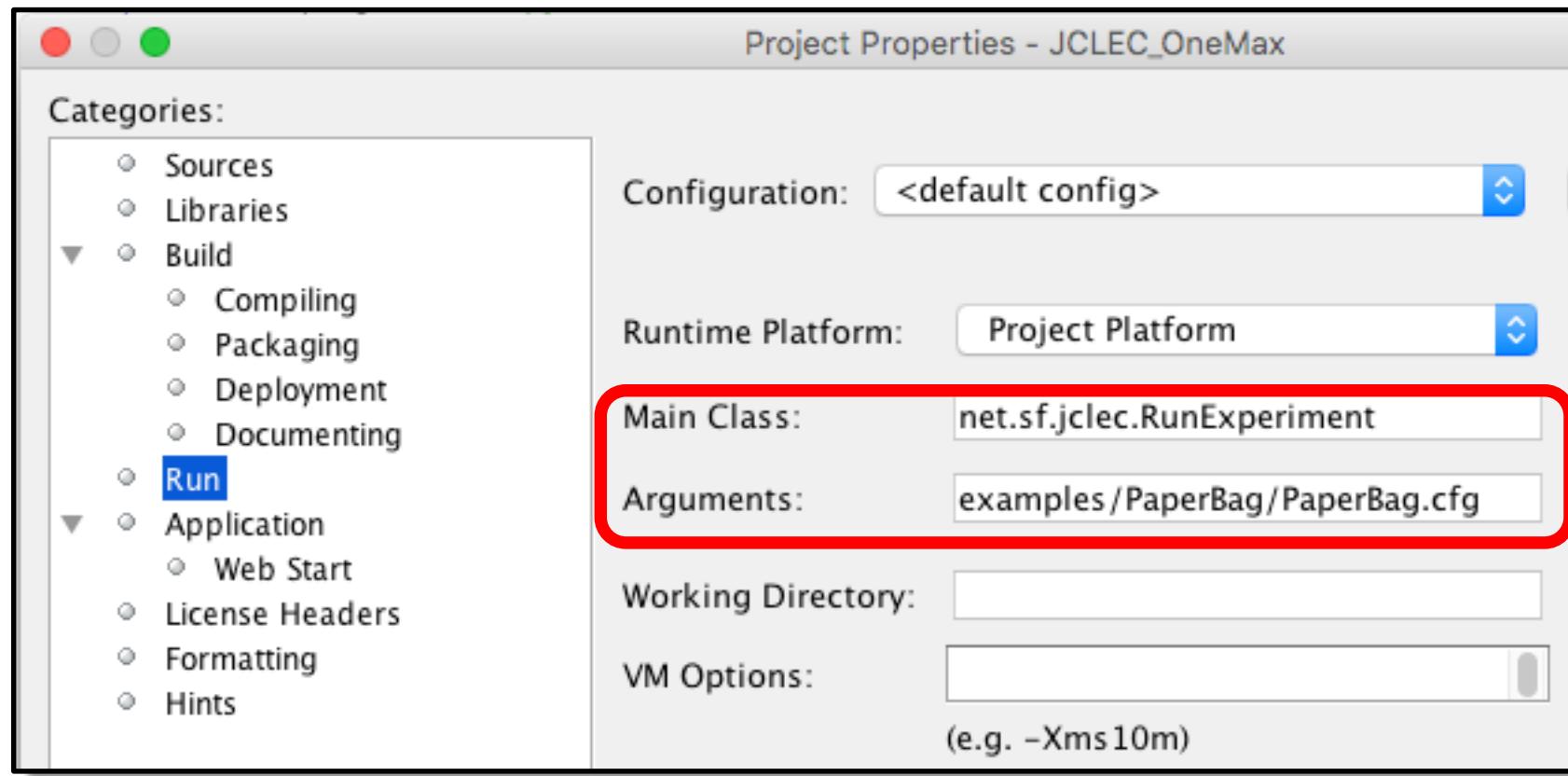
**Population size:** 12 individuals

**Stop Criterion:** 20 generations

**Parent selection:** tournament of 2 individuals

```
<experiment>
  <process algorithm-type="net.sf.jclec.algorithm.classic.SGE">
    <population-size>12</population-size>
    <max-of-generations>20</max-of-generations>
    <rand-gen-factory type="net.sf.jclec.util.random.RanecuFactory" seed="124321453"/>
    <species type="net.sf.jclec.realarray.RealArrayIndividualSpecies">
      <genotype-schema>
        <locus type="net.sf.jclec.util.range.Interval" left="0.0" right="20.0"
               closure="closed-closed"/>
        <locus type="net.sf.jclec.util.range.Interval" left="0.0" right="180.0"
               closure="closed-closed"/>
      </genotype-schema>
    </species>
    <evaluator type="tutorial.PaperBag" />
    <provider type="net.sf.jclec.realarray.RealArrayCreator" />
    <parents-selector type="net.sf.jclec.selector.TournamentSelector"
                      tournament-size="2" />
    <mutator type="net.sf.jclec.realarray.mut.NonUniformMutator" mut-prob="0.15" />
    <recombinator type="net.sf.jclec.realarray.rec.BLXAlphaCrossover" rec-prob="0.9"
                  alpha="0.3" />
    <listener ... </listener>
  </process>
</experiment>
```

Don't forget to invoke the executable with "PaperBag.cfg" as an argument



# Enjoy the programming!

Hint – think about converting the angle theta to radians before applying `sin()` and `cos()`

Here's one way to solve the 'Out of a Paper Bag' optimisation problem...

Let's start with a 'Point' class (with public x & y attributes for convenience):

```
public class Point {  
    public double x;  
    public double y;  
  
    public Point( ) {  
        x = 0.0;  
        y = 0.0;  
    }  
}
```

```
public class PaperBag extends AbstractEvaluator {  
  
    protected boolean maximize = true;  
  
    private Comparator<IFitness> COMPARATOR;  
  
    /* list of x,y points of the projectile trajectory */  
    private List< Point > pointsList;  
  
    private DecimalFormat df; // for debugging  
  
    private static final double GRAVITY = 9.81; // gravity i.e. 9.81 m/sec2  
  
    private static final double WIDTH = 10.0; // width of the paper bag  
  
    private static final double HEIGHT = 5.0; // height of the paper bag  
  
    private static final double STEP = 0.1; // the "time step"  
  
    // ...
```

Continued....

## One way of solving the fitness evaluation:

```
protected void evaluate( IIndividual ind ) {  
    // Individual genotype  
    double[ ] genotype = ((RealArrayIndividual)ind).getGenotype( );  
    double velocity = genotype[ 0 ];  
    double theta = genotype[ 1 ];  
  
    pointsList = new ArrayList< >( ); // clear out the list of points  
  
    // calculate the points of the parabolic trajectory  
    for( double time = 0.0; time < END; time += STEP ){  
        Point p = getPointAtTime( time, velocity, theta );  
        pointsList.add( p );  
    }  
  
    // ...
```

Continued....

```
double fitness = 0.0;

// calculate fitness value from the parabolic trajectory points
for( Point p : pointsList ) {
    if( p.x <= 0.0 || p.x >= WIDTH ) {
        fitness = p.y;
        break;
    }
}

ind.setFitness( new SimpleValueFitness( fitness ) );
}
```

## The getPointAtTime( ) method:

```
private Point getPointAtTime( final double time, final double velocity,
    final double theta )      {
    double inRadians = Math.toRadians( theta ); // convert to radians

    double xTemp = 0.5 * WIDTH;
    double xIncrement = velocity * time * Math.cos( inRadians );
    xTemp += xIncrement;

    double yTemp = velocity * time * Math.sin( inRadians );
    double yIncrement = 0.5 * GRAVITY * ( time * time );
    yTemp -= yIncrement;
    // can't have a negative y value - this is the ground!
    if( yTemp < 0.0 ) yTemp = 0.0;

    Point p = new Point( );
    p.x = xTemp; p.y = yTemp;
    return p;
}
```

# Demonstration

JCLEC\_OneMax - NetBeans IDE 8.1

<default config> | Search (%+)

Output - JCLEC\_OneMax (run) | X

```
Median individual: net.sf.jclec.realarray.RealArrayIndividual@12a00e1genotype={13.598527962322732,76.20177283896926}, fitness=net.sf.jclec.fitness.SimpleValueFitness@20fa25c1[value=8.57295523277901]
Average fitness = 8.57295523277901
Fitness variance = 0.0

Generation 15 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@3581c5f3[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2530c12[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@3581c5f3[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2530c12[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@3581c5f3[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2530c12[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

Generation 16 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@73c6c3b2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@64a294a6[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@73c6c3b2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@64a294a6[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@73c6c3b2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@64a294a6[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

Generation 17 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@7e0b37bc[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@6ae40994[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@7e0b37bc[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@6ae40994[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@7e0b37bc[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@6ae40994[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

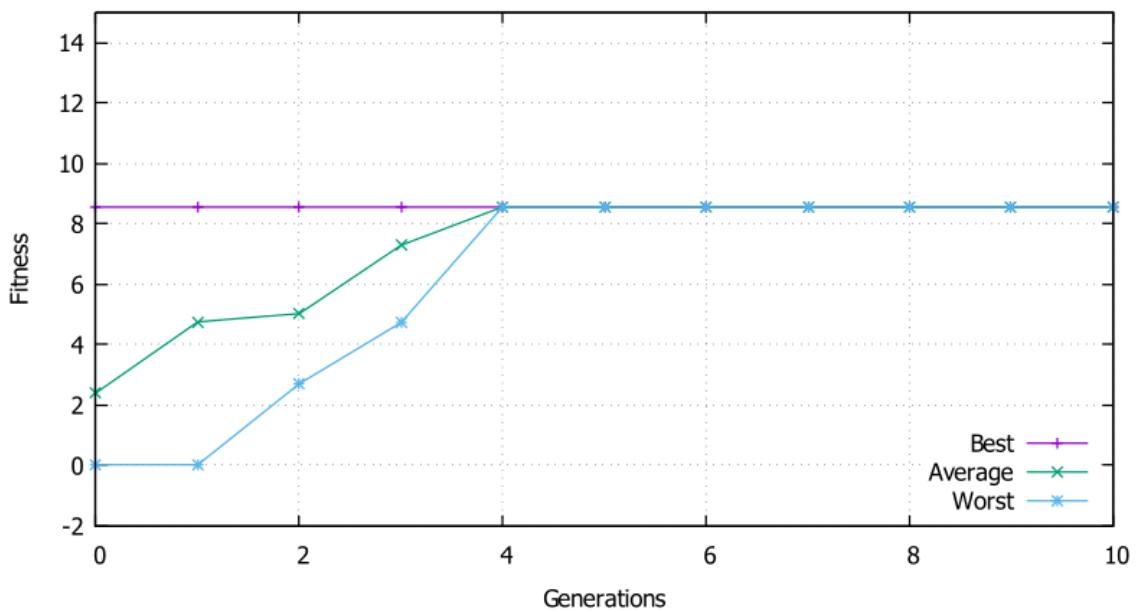
Generation 18 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@1a93a7ca[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2b05039f[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@1a93a7ca[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2b05039f[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@1a93a7ca[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@2b05039f[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

Generation 19 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@61e717c2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@4dcbadb4[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@61e717c2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@4dcbadb4[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@61e717c2[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@4dcbadb4[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

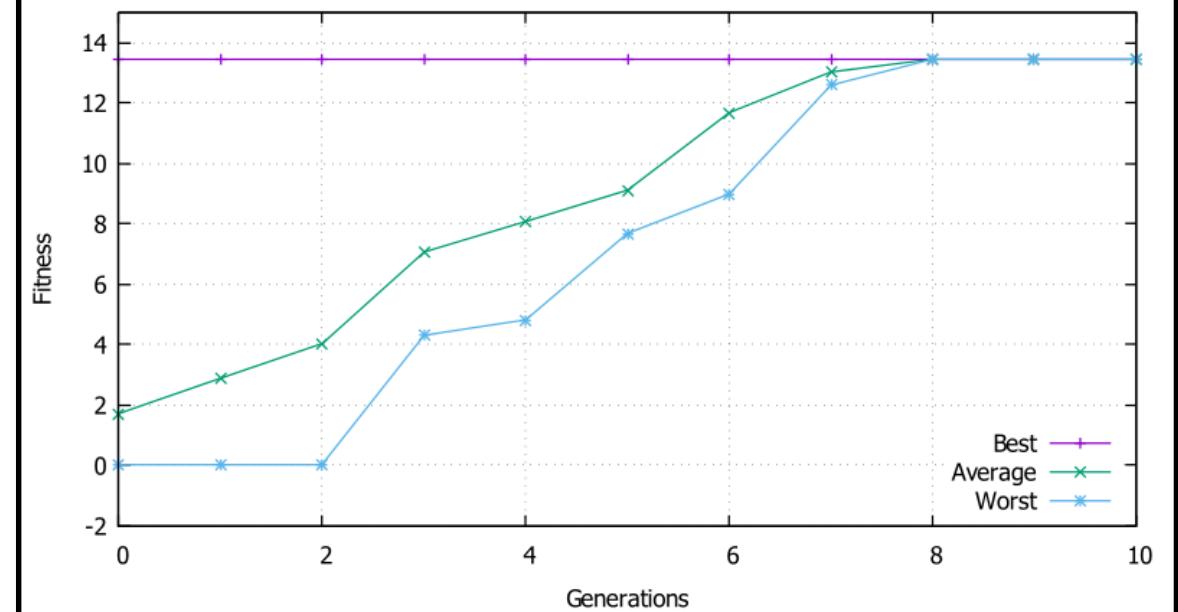
Generation 20 Report
Best individual: net.sf.jclec.realarray.RealArrayIndividual@4e515669[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@1b9e1916[value=8.57295523277901]]
Worst individual: net.sf.jclec.realarray.RealArrayIndividual@4e515669[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@1b9e1916[value=8.57295523277901]]
Median individual: net.sf.jclec.realarray.RealArrayIndividual@4e515669[genotype={13.598527962322732,76.20177283896926},fitness=net.sf.jclec.fitness.SimpleValueFitness@1b9e1916[value=8.57295523277901]]
Average fitness = 8.57295523277901
Fitness variance = 0.0

Algorithm finished
Job finished
BUILD SUCCESSFUL (total time: 0 seconds)
```

PaperBag fitness curves, population size 12



PaperBag fitness curves, population size 100



# Workshop

Evolutionary computing

Frameworks for evolutionary computing

Java Class Library for Evolutionary Computing (JCLEC)

**Optimisation problems:**

- 1 - 'OneMax' Problem
- 2 - How to program your way out of a paper bag
- 3 - FizzBuzz**

Genetic Programming, Genetic Improvement

# Rules of FizzBuzz

A typical game of FizzBuzz involves counting through a sequence of numbers starting at one, but multiples of three are substituted with ‘Fizz’, multiples of five are substituted with ‘Buzz’, and multiples of fifteen are substituted with ‘FizzBuzz’.

# Let's start by evolving a sequence of 100 integers...

- First define the actual sequence in the configure method in a FizzBuzz class
  - Here are some constants that might be useful...

```
private List< Integer > sequence; // to evolve against

private static final int SEQUENCE_SIZE = 100;

private static final int FIZZ = 3;
private static final int BUZZ = 5;
private static final int FIZZ_BUZZ = 15;

private static final Integer FIZZ_IDENTIFIER = 101;
private static final Integer BUZZ_IDENTIFIER = 102;
private static final Integer FIZZ_BUZZ_IDENTIFIER = 103;
```

# Enjoy the programming!

Here's one way to solve the 'FizzBuzz as a sequence' problem...

```
public void configure(Configuration settings)  {
    sequence = new ArrayList< >( );
    for( int i = 1; i <= SEQUENCE_SIZE; i++ )  {
        sequence.add( new Integer( i ) );
    }
    for( int i = 1; i < sequence.size( ); i++ )  {
        int number = sequence.get( i );
        if( number % FIZZ_BUZZ == 0 )  {
            sequence.set( i, new Integer( FIZZ_BUZZ_IDENTIFIER ) );
        }
        else if( number % BUZZ == 0 )  {
            sequence.set( i, new Integer( BUZZ_IDENTIFIER ) );
        }
        else if( number % FIZZ == 0 )  {
            sequence.set( i, new Integer( FIZZ_IDENTIFIER ) );
        }
    }
}
```

## Here's the fizzbuzz.cfg file (1 of 2):

```
<experiment>
  <process algorithm-type="net.sf.jclec.algorithm.classic.SGE">
    <rand-gen-factory type="net.sf.jclec.util.random.RanecuFactory">
      seed="123456789" />
    <population-size>100</population-size>
    <max-of-generations>3000</max-of-generations>

    <provider type="net.sf.jclec.intarray.IntArrayCreator" />
      <species type="net.sf.jclec.intarray.IntArrayIndividualSpecies">
        genotype-length="100">
        <genotype-schema>
          <locus type="net.sf.jclec.util.intset.Interval">
            left="1" right="103" closure="closed-closed" />
          <locus type="net.sf.jclec.util.intset.Interval">
            left="1" right="103" closure="closed-closed" />
          ...
          ... Etc. for each integer in the array of 100 integers ...
        </genotype-schema>
      </species>
```

Continued...

## Here's the fizzbuzz.cfg file (2 of 2):

```
<evaluator type="tutorial.FizzBuzz"> </evaluator>

<parents-selector type="net.sf.jclec.selector.TournamentSelector">
    <tournament-size>2</tournament-size>
</parents-selector>

<mutator type="net.sf.jclec.intarray.mut.OneLocusMutator" mut-prob="0.2" />
<recombinator type="net.sf.jclec.intarray.rec.OnePointCrossover" rec-prob="0.9"/>

<listener type="net.sf.jclec.listener.PopulationReporter">
    <report-frequency>10</report-frequency>
    <report-on-file>false</report-on-file>
    <save-complete-population>false</save-complete-population>
    <report-title>FizzBuzz-</report-title>
</listener>
</process>
</experiment>
```

## The evaluation( ) method is almost trivial – something like ‘OneMax’?

```
protected void evaluate( IIndividual ind ) {
    int [ ] genotype = ( ( IntArrayIndividual ) ind ).getGenotype( );
    int matchCount = 0;

    for( int i = 0; i < genotype.length; i++ ) {
        if( genotype[ i ] == sequence.get( i ) ) {
            matchCount++;
        }
    }

    ind.setFitness( new SimpleValueFitness( matchCount ) );
}
```

# Demonstration

# But there's a problem with this approach....

- How do you write the fitness function without writing code to generate the correct values?
- What properties do we need?
- Listen to @KevlinHenney

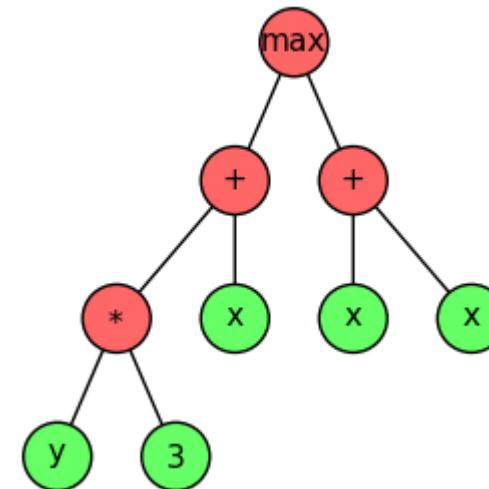
*every result is 'Fizz', 'Buzz', 'FizzBuzz' or a decimal string,  
every decimal result corresponds to its ordinal position,  
every third result contains 'Fizz',  
every fifth result contains 'Buzz',  
every fifteenth result is 'FizzBuzz',  
the ordinal position of every 'Fizz' result is divisible by 3,  
the ordinal position of every 'Buzz' result is divisible by 5,  
the ordinal position of every 'FizzBuzz' result is divisible by 15*



```
actual = [fizzbuzz(n) for n in range(1, 101)]
truths = [
    all(a in {'Fizz', 'Buzz', 'FizzBuzz'} or a.isdecimal() for a in actual),
    all(int(a) == n for n, a in enumerate(actual, 1) if a.isdecimal()),
    all('Fizz' in a for a in actual[2::3]),
    all('Buzz' in a for a in actual[4::5]),
    all(a == 'FizzBuzz' for a in actual[14::15]),
    all(n % 3 == 0 for n, a in enumerate(actual, 1) if a == 'Fizz'),
    all(n % 5 == 0 for n, a in enumerate(actual, 1) if a == 'Buzz'),
    all(n % 15 == 0 for n, a in enumerate(actual, 1) if a == 'FizzBuzz')
]
assert all(truths)
```

# Change the array to a tree

- Previous genotypes were “list/array”
- What if we use trees?
- What do trees make?
- Code



But JCLEC doesn't specifically offer AST representations... and that looked a bit like Python...

## Distributed Evolutionary Algorithms in Python (DEAP)

<https://deap.readthedocs.io/en/master/>

DEAP documentation



DISTRIBUTED  
EVOLUTIONARY  
ALGORITHMS IN  
PYTHON

DEAP is a novel evolutionary computation framework for rapid prototyping and testing of ideas. It seeks to make algorithms explicit and data structures transparent. It works in perfect harmony with parallelisation mechanism such as multiprocessing and SCOOP. The following documentation presents the key concepts and many features to build your own evolutions.

**First steps:**

- [Overview \(Start Here!\)](#)
- [Installation](#)
- [Porting Guide](#)

**Basic tutorials:**

- [Part 1: creating types](#)
- [Part 2: operators and algorithms](#)
- [Part 3: logging statistics](#)
- [Part 4: using multiple processors](#)

**Advanced tutorials:**

- [Genetic Programming](#)
- [Checkpointing](#)
- [Constraint Handling](#)
- [Benchmarking Against the Bests \(BBOB\)](#)
- [Inheriting from Numpy](#)

**Examples**

**Library Reference**

**Release Highlights**

**Contributing**

**About DEAP**

**Getting Help**

Having trouble? We'd like to help!

- Search for information in the archives of the [deap-users mailing list](#), or post a question.
- Report bugs with DEAP in our [issue tracker](#).

# So let's evolve a tree

<https://deap.readthedocs.io/en/master/tutorials/advanced/gp.html>

- You add a primitive set,

```
pset = PrimitiveSet("main", 2)
```

- Add function and number operators:

```
pset.addPrimitive(max, 2)
pset.addPrimitive(operator.add, 2)
pset.addPrimitive(operator.neg, 1)
```

- Or values:

```
pset.addTerminal(3)
```

- Then generate trees:

```
expr = genFull(pset, min_=1, max_=3)
tree = PrimitiveTree(expr)
```

# For Fizz Buzz

```
def if_then_else(x,y,z):
    if x:
        return y
    else:
        return z

def mod3(x):
    return operator.mod(x, 3) == 0

def mod5(x):
    return operator.mod(x, 5) == 0

def mod15(x):
    return operator.mod(x, 15) == 0

def both(x, y):
    return x and y

def either(x, y):
    return x or y
```

```
pset = gp.PrimitiveSet( "MAIN" , 1)
pset.addPrimitive(operator.add, 2)
pset.addPrimitive(operator.sub, 2)
pset.addPrimitive(operator.mul, 2)

pset.addPrimitive(both, 2)
pset.addPrimitive(either, 2)

pset.addPrimitive(operator.mod, 2)

pset.addPrimitive(if_then_else, 3)
pset.addPrimitive(mod3, 1)
pset.addPrimitive(mod5, 1)
pset.addPrimitive(mod15, 1)

pset.addTerminal( "Buzz" )
pset.addTerminal( "Fizz" )
pset.addTerminal( "FizzBuzz" )
```

Can get it to find numbers, e.g.

```
pset.addEphemeralConstant( "rand101" , lambda: random.randint(-1,1))
```

But don't have a working example yet

# Evolve...

```
creator.create("FitnessMax", base.Fitness, weights=(1.0, ))
creator.create("Individual", gp.PrimitiveTree,
              fitness=creator.FitnessMax)

toolbox = base.Toolbox()                                #or genFull or genGrow
                                                       # genHalfAndHalf does grow 50% or
toolbox.register("expr", gp.genHalfAndHalf,             # time, Full 50%
                pset=pset, min_=1, max_=2)
toolbox.register("individual", tools.initIterate,
                creator.Individual, toolbox.expr)
toolbox.register("population", tools.initRepeat, list,
                toolbox.individual)
toolbox.register("compile", gp.compile, pset=pset)
```

# Fitness...

```
def fizz_buzz(func, points):
    passed = 0
    def safe_run(func, x):
        try:
            return func(x)
        except:
            return -1
    results = [safe_run(func, x) for x in points]
    if every_result_is_Fizz_Buzz_FizzBuzz_or_a_decimal(results):
        passed += 1
    if every_decimal_result_corresponds_to_its_ordinal_position(results):
        passed += 1

    if every_third_result_contains_Fizz(results):
        passed += 1
    if every_fifth_result_contains_Buzz(results):
        passed += 1

    if every_fifteenth_result_contains_FizzBuzz(results):
        passed += 1
    if the_ordinal_position_of_every_Fizz_result_is_divisible_by_3(results):
        passed += 1
    if the_ordinal_position_of_every_Buzz_result_is_divisible_by_5(results):
        passed += 1
    if the_ordinal_position_of_every_FizzBuzz_result_is_divisible_by_15(results):
        passed += 1
    return passed
```

```

def register(fn):      #e.g. our fizz buzz tests fitness function

    def eval(individual, points): # This is our custom evaluation function
        # Transform the tree expression in a callable function
        func = toolbox.compile(expr=individual)
        # Evaluate the result, somehow
        return fn(func, points),

    toolbox.register("evaluate", eval, points=range(101)) # and we register it here
    toolbox.register("select", tools.selTournament, tournsize=3)
    toolbox.register("mate", gp.cxOnePoint)
    toolbox.register("expr_mut", gp.genFull, min_=0, max_=2)
    toolbox.register("mutate", gp.mutUniform, expr=toolbox.expr_mut, pset=pset)

    toolbox.decorate("mate", gp.staticLimit(key=operator.attrgetter("height"), max_value=17))
    toolbox.decorate("mutate", gp.staticLimit(key=operator.attrgetter("height"), max_value=17))

def main():
    random.seed(318)
    register(fizz_buzz)
    pop = toolbox.population(n=4000)
    hof = tools.HallOfFame(1) #The best

    pCrossover = 0.75
    pMutation = 0.5
    nGen = 75
    pop, log = algorithms.eaSimple(pop, toolbox, pCrossover, pMutation, nGen, halloffame=hof)
#can add other stats
    return pop, log, hof

```

# Ta-da!

```
['FizzBuzz', 1, 2, 'Fizz', 4, 'Buzz', 'Fizz', 7, 8, 'Fizz', 'Buzz', 11, 'Fizz', 13, 14, 'FizzBuzz', 16, 17, 'Fizz', 19, 'Buzz', 'Fizz', 22, 23, 'Fizz', 'Buzz', 26, 'Fizz', 28, 29, 'FizzBuzz', 31, 32, 'Fizz', 34, 'Buzz', 'Fizz', 37, 38, 'Fizz', 'Buzz', 41, 'Fizz', 43, 44, 'FizzBuzz', 46, 47, 'Fizz', 49, 'Buzz', 'Fizz', 52, 53, 'Fizz', 'Buzz', 56, 'Fizz', 58, 59, 'FizzBuzz', 61, 62, 'Fizz', 64, 'Buzz', 'Fizz', 67, 68, 'Fizz', 'Buzz', 71, 'Fizz', 73, 74, 'FizzBuzz', 76, 77, 'Fizz', 79, 'Buzz', 'Fizz', 82, 83, 'Fizz', 'Buzz', 86, 'Fizz', 88, 89, 'FizzBuzz', 91, 92, 'Fizz', 94, 'Buzz', 'Fizz', 97, 98, 'Fizz', 'Buzz']
```

# The Hof

```
if_then_else(mod15(if_then_else(if_then_else(mul(x, 'FizzBuzz'), 'Fizz',
'Buzz'), x, if_then_else('Buzz', 'FizzBuzz', mod3(x)))), 'FizzBuzz',
if_then_else(both(if_then_else(if_then_else(mod15(x),
either('FizzBuzz', 'FizzBuzz'), 'FizzBuzz'), if_then_else('FizzBuzz',
mod15(mod5(x)), 'Buzz'), 'Buzz'), if_then_else('Fizz', 'Buzz',
if_then_else('FizzBuzz', if_then_else(if_then_else('Buzz',
if_then_else(if_then_else(mod3(x), x, 'FizzBuzz'), if_then_else(x, x,
either('Buzz', 'Buzz')), x), 'Fizz'), 'Fizz', x),
if_then_else(either(if_then_else(x, x, mod3(x)), 'FizzBuzz'), 'Fizz',
'Fizz')))), if_then_else(mod15(x), either('FizzBuzz', either('Buzz', x)),
if_then_else(mod3(x), 'Fizz', x)), 'Buzz'))
```

# The Hof

```
if_then_else(mod15(if_then_else(if_then_else(mod15(x), 'FizzBuzz'), 'Fizz',  
'Buzz'), x, if_then_else(mod15(if_then_else(if_then_else(mod15(x), 'FizzBuzz'),  
if_then_else(if_then_else(if_then_else(if_then_else(if_then_else(if_then_else(mod15(mod5(x)),  
either('FizzBuzz', 'Buzz'), 'Buzz',  
mod15(mod5(x)), 'Buzz',  
if_then_else('FizzBuzz', 'Buzz',  
if_then_else(if_then_else(if_then_else(if_then_else(if_then_else(if_then_else(x, x,  
either('Buzz', 'Buzz',  
if_then_else(either(if_then_else(x, x, mod3(x)), 'FizzBuzz'), 'Fizz',  
'Fizz')))), if_then_else(mod15(x), either('FizzBuzz', either('Buzz', x)),  
if_then_else(mod3(x), 'Fizz', x)), 'Buzz'))
```



# Workshop

Evolutionary computing

Frameworks for evolutionary computing

Java Class Library for Evolutionary Computing (JCLEC)

**Optimisation problems:**

- 1 - ‘OneMax’ Problem
- 2 - How to program your way out of a paper bag
- 3 - FizzBuzz

**Genetic Programming, Genetic Improvement**

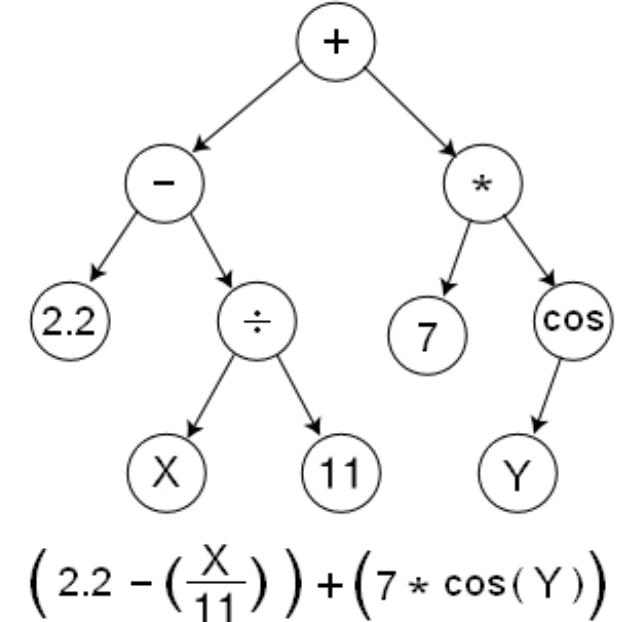
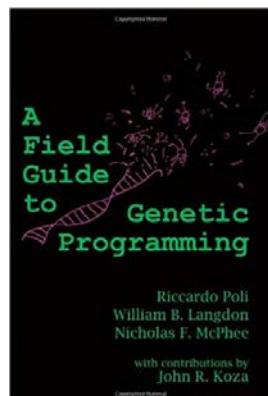
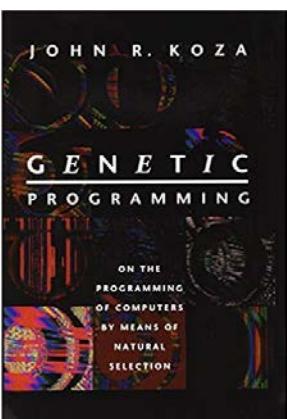
# Genetic Programming (GP) - evolution of a tree structure

Evolves the ‘innards’ (white box) of a function or expression

Each tree node is an *operator* or *variable*,  
or a *terminal node*.

Used widely to evolve functions for

- Curve fitting
- Circuit board design
- Data modelling
- Symbolic regression
- Feature selection
- Classification



[https://en.wikipedia.org/wiki/Genetic\\_programming#/media/File:Genetic\\_Program\\_Tree.png](https://en.wikipedia.org/wiki/Genetic_programming#/media/File:Genetic_Program_Tree.png)

# Could we use GP to evolve source code e.g. for FizzBuzz?

Well, using AST representations, as we saw above, yes!

**BUT** GP doesn't always scale well (e.g. due to 'code bloat' with crossover) and  
*code doesn't look like what a human programmer would produce!!!*

So in practice, maybe better to start with an existing body of code  
and use '**Genetic Improvement**' (GI) e.g.

IEEE TRANSACTIONS ON EVOLUTIONARY COMPUTATION, VOL. 22, NO. 3, JUNE 2018 415

## Genetic Improvement of Software: A Comprehensive Survey

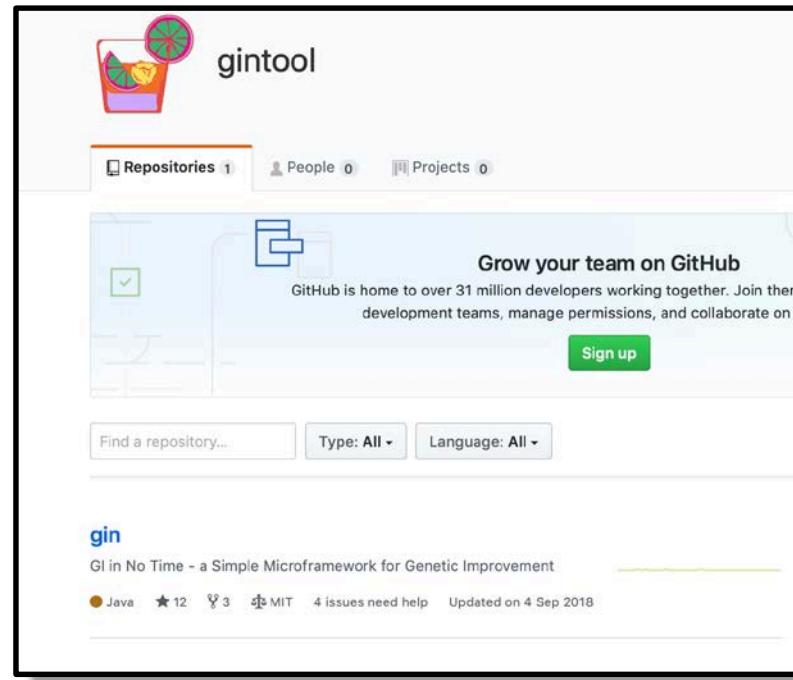
Justyna Petke, Saemundur O. Haraldsson, Mark Harman, *Member, IEEE*, William B. Langdon, David R. White, and John R. Woodward

**Abstract**—Genetic improvement (GI) uses automated search to find improved versions of existing software. We present a comprehensive survey of this nascent field of research with a focus on the core papers in the area published between 1995 and 2015. We identified core publications including empirical studies, 96% of which use evolutionary algorithms (genetic programming in particular). Although we can trace the foundations of GI back to the origins of computer science itself, our analysis reveals a significant upsurge in activity since 2012. GI has resulted in dramatic performance improvements for a diverse set of properties such as execution time, energy and memory consumption, as well as results for fixing and extending existing system functionality. Moreover, we present examples of research work that lies on the boundary between GI and other areas, such as program transformation, approximate computing, and software repair, with the intention of encouraging further exchange of ideas between researchers in these fields.

**Index Terms**—Genetic improvement (GI), survey.

Recent work on GI has received notable awards, demonstrating its acceptance and success within the wider software engineering and evolutionary computation communities. For example, work on GI for software repair and specialization won four "Humies" [1]–[5], awarded for human-competitive results produced by genetic and evolutionary computation [6]. Several papers on GI also won distinguished paper awards [1], [5] and technical challenges [7]. GI has also been the subject of attention from the broadcast media, as well as popular developer magazines, websites, and blogs [8]–[11], demonstrating its influence and reach beyond the research community to the wider developer community and the public at large.

This survey of 3132 distinct titles found, resulted in the identification of 66 core GI papers.<sup>1</sup> However, GI research draws on and potentially influences many other areas in soft-



The screenshot shows the GitHub profile for the repository 'gintool'. At the top, there's a logo of a cocktail glass with a lime wedge. Below it, the repository name 'gintool' is displayed. Underneath, there are three tabs: 'Repositories 1', 'People 0', and 'Projects 0'. A prominent green 'Sign up' button is located on the right. The main area features a large image of a computer monitor displaying a software interface with various windows and icons. Below this, there's a search bar with the placeholder 'Find a repository...', and dropdown menus for 'Type: All' and 'Language: All'. Further down, another repository card is visible for 'gin'.

gin  
GI in No Time - a Simple Microframework for Genetic Improvement

Java ★ 12 3 MIT 4 issues need help Updated on 4 Sep 2018

*For further information:*

The Pragmatic Programmers

# Genetic Algorithms and Machine Learning for Programmers

Create AI Models  
and Evolve Solutions

Frances Buontempo  
*edited by Tammy Coron*

A silhouette of a person's head and shoulders is overlaid with a grid pattern. A hand is shown holding a molecular model composed of red and blue spheres connected by black lines.

This article has been accepted for publication in a future issue of this journal, but has not been fully edited. Content may change prior to final publication. Citation information: DOI 10.1109/TSE.2018.2803055, IEEE Transactions on Software Engineering

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## A Systematic Review of Interaction in Search-Based Software Engineering

Aurora Ramírez, José Raúl Romero, *Member, IEEE*, and Christopher L. Simons

**Abstract**—Search-Based Software Engineering (SBSE) has been successfully applied to automate a wide range of software development activities. Nevertheless, in those software engineering problems where human evaluation and preference are crucial, such insights have proved difficult to characterize in search, and solutions might not look natural when that is the expectation. In an attempt to address this, an increasing number of researchers have reported the incorporation of the ‘human-in-the-loop’ during search and interactive SBSE has attracted significant attention recently. However, reported results are fragmented over different development phases, and a great variety of novel interactive approaches and algorithmic techniques have emerged. To better integrate these results, we have performed a systematic literature review of interactive SBSE. From a total of 669 papers, 26 primary studies were identified. To enable their analysis, we formulated a classification scheme focused on four crucial aspects of interactive search, i.e. the problem formulation, search technique, interactive approach, and the empirical framework. Our intention is that the classification scheme affords a methodological approach for interactive SBSE. Lastly, as well as providing a detailed cross analysis, we identify and discuss some open issues and potential future trends for the research community.

**Index Terms**—Search-Based Software Engineering, Interaction, Systematic Literature Review, Optimization

### 1 INTRODUCTION

The design and development of complex, large-scale software systems can be non-trivial and challenging for the software engineer to perform. In an attempt to

base problems in the software development lifecycle, this may also be challenging [6] (chapter 9).

In addition, results of automated search approaches

# Thanks!

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