## COMP6202 Assignment2

# How Do Coevolutionary Dynamics in a Minimal Substrate Change with an Increase in the Number of Populations?

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#### Abstract

Three sets of results from a paper[1] are successfully reproduced to illustrate coevolutionary failures. The experiments are then repeated with more populations to confirm the hypothesis that increasing the number of populations from 2 to 10 solves the problem of 'focusing' or 'over specialization' within species containing multiple dimensions contributing to objective fitness.

## 1 The Original Paper

The original paper[1] explores and illustrates the following three cases pertinent to coevolutionary failure: Loss of gradient, over-specialisation and relativism. Three experiments are devised to aid this discussion, all of which make use of a minimal substrate. This makes it easier to attribute issues to coevolutionary techniques and diagnose that they are not a product of the complexity of the task

## 1.1 Experiment One: Loss of Gradient

Experiment one illustrates the concept of a 'loss of gradient' in a coevolutionary set-up. This is achieved using the coevolution of scalar values; where the scalar value being evolved is the unitation of a bit string i.e. the fitter the individual the more ones in its bit string.

A loss of gradient causes a population to drift without improvement[1]. This coevolutionary pathology can be explained by the Red Queen Effect[2]. The subjective performance of individuals in a species does not improve despite improvements in their objective fitness as the competing species improves at the same rate.

The experiment is formalised using equation 1[1]. The value returned is with respect to a random sample of individuals S from the competing population.

$$f(a,S) = \sum_{i=1}^{|S|} score(a,S_i)$$
 /eq.1

where score(a,b)=1 if a>b, 0 otherwise.

This creates a distinction between the objective and subjective fitness of an individual. Individuals are evolved based on their performance against the other population as opposed to an objective fitness metric.

## 1.2 Distinction Between Objective and Subjective Fitness

An important distinction between the objective and subjective fitness of an individual is clarified in the original paper[1]. The objective fitness is defined as the metric that we are attempting to optimize, whilst the subjective fitness is the perceived performance of the co-evolving individual. This relates to the number of ones in an individuals bit string and the result of the defined subjective fitness functions (see equations) respectively.

# 1.3 Mutation Biases and Justification for Approach

The nature of the approach (representing individuals using bit strings) results in an inherent mutation bias. A mutation bias of 0.005 is used in the original paper. This value is derived from the string length, 100, and the natural bias towards half 0s and half 1s[5]. The chosen mutation bias results in, on average, 1 bit being assigned a new random value per individual selected for reproduction for each generation.

# 1.4 Experiment Two: Over Specialization

Experiment Two introduces multiple dimensions to an individuals definition. Instead of having one dimension 100 bits long. 10 dimensions each 10 bits long are used for representation.

A single dimension determines the outcome of a match between competing individuals. For Experiment Two, this is the dimension with the largest difference between the individuals. Equation 2[1] characterizes this approach:

This set-up is used to illustrate 'over specialization' or 'focussing'. A phenomenon where high performance

$$f2(a,S) = \sum_{i=1}^{|S|} score2(a,S_i)$$
 where 
$$score2((a_x,a_y),(b_x,b_y)) = \begin{cases} score(a_x,b_x), \text{ if } (|a_x-b_x|>|a_y-b_y|) \\ score(a_y,b_y), \text{ otherwise.} \end{cases}$$

cannot simultaneously be maintained in all dimensions. Whilst one dimension is focussed on and developed to improve against the other population the other nine drift towards their neutral "50:50" composition.

### 1.5 Experiment Three: Relativism

The third experiment demonstrates intransitive superiority [2] in a coevolutionary framework. The aim is to replicate a cyclical scenario where, for three members, "a beats b beats c beats a"[1]. A slight alteration to equation 2 so the dimension with the *smallest* difference is used to determine the score between two individuals, can introduce circular dominance relations. Equation 3[1] represents this altered approach.

$$f3(a,S) = \sum_{i=1}^{|S|} score3(a,S_i)$$
 where 
$$score3((a_x,a_y),(b_x,b_y)) = \begin{cases} score(a_x,b_x), & \text{if } (|a_x-b_x| < |a_y-b_y|) \\ score(a_y,b_y), & \text{otherwise.} \end{cases}$$

and, as before, score(a,b)=1 if a>b, 0 otherwise.

## 2 Reimplementing Results

### 2.1 Experimental Set Up

The experiments described in the previous section have been re-implemented, the parameters used can be found in table 1. The run function found in **AP-PENDIX** defines the general approach for each experiment. *Length* (of each dimension), *dimensions*, eq (equation used to calculate subjective fitness) and

generations (number of generations to run the simulation) are the main variables between experiments (the sample size was altered on one occasion).

Parameter	Value
Evolutionary Algorithm	Generational
Selection Scheme	Roulette Wheel
Sample Size (S)	Variable
Representation	Bit string
Population Size	25
Dimensions	Variable
Dimension Length	Variable
Subjective Fitness Equation	Variable

Table 1: Parameters for reimplementing results

A Generational approach is taken; to produce a new generation fitness proportionate selection (roulette wheel selection) is used. The number of times the 'wheel' is spun is equal to the size of the population, each time the selected individual is passed through a mutate function, where each bit has a probability equal to the mutation bias of being assigned a new random value.

# 2.2 Reimplemented Results with Discussion

All of the reimplemented results were run for double the number of generations, to corroborate the expected behavior and in some cases showcase downward trends and separation between the populations multiple times (ruling out anomalous characteristics). This decision was substantiated by the fact that running the simulations was not very computationally expensive.

#### 2.2.1 Experiment One: Loss of Gradient

The parameters shown in **Table 2** are used to replicate the results for the first part of Experiment One from the original paper. **Figure 1** shows the original and reimplemented results. Both sets of results show the same trend: The populations reach the optimal

solution after approximately 400 generations, with a variation in the subjective fitness of one populations equal to around one minus the subjective fitness of the other. Due to the performance improvement of opponents at the same rate, the average subjective fitness of the population plateaus once the optimal objective fitness is reached. This is problematic in scenarios where an objective fitness of an individual is not measurable. However, the optimal solution could be inferred when the average subjective fitness has converged.

Parameter	Value
Sample Size (S)	15
Population Size	25
Dimensions	1
Dimension Length	100
Subjective Fitness Equation	eq1

Table 2: Parameters for reimplementing experiment one

The parameter values used to replicate the second part of Experiment One are the same as those shown in Table 2, save the sample size, S, which is now 1. Now each member competes with a randomly selected member of the other population. Comparing the results shown in **Figure 2** it is clear that in both set there are multiple downward and upward trends, with periods of complete separation between the populations. These downward trends coincide with a period of polarisation in the subjective fitnesses of the populations for both graphs. Showing that all the individuals in one population beat all the individuals in the other.

#### 2.2.2 Experiment Two: Over Specialization

Table 3 provides the parameters used to replicate the results for Experiment Two. Figure 3 compares the reimplemented results with original set for this experiment and shows that the experiment was successfully reimplemented. Both results show the same

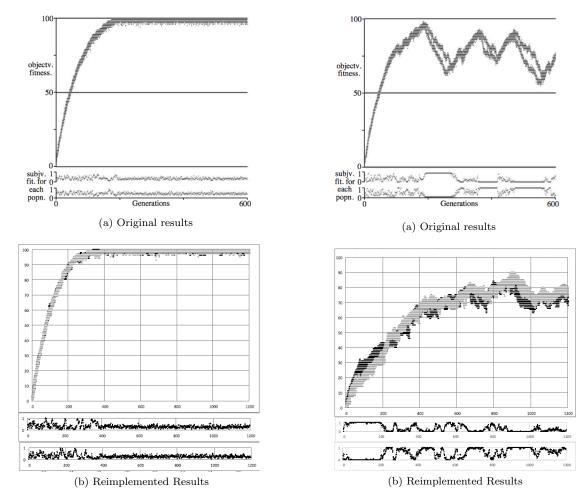


Figure 1: Reimplemented results for experiment one (original on top)

Figure 2: Reimplemented results for experiment one (original on top)

leading to a focus on improving that dimension, sub-

sequently the improvements in the original dimension

trends in objective and subjective fitness. The objective performance never reaches 100 for an individual and the subjective performances continue to show an inversely proportional relationship.

are lost and the competing population is improving at the same rate in the new dimension - so the cycle continues.

t

2.2.3 Experiment Three: Relativism

Table 4 details the parameters used to replicate the

In both cases the trends are explained by the fact that top performance across all dimensions cannot be maintained due to "focussing" within a single dimension. Here we can see the principles of the Red Queen Effect applied to dimensions within an individual. Although the performance across one dimension is improving, the performance across that dimension is improving at the same rate in the other population. Once maximal performance in this dimension is achieved, a change in performance in another dimension can cause a population to improve subjectively,

Table 4 details the parameters used to replicate the results for Experiment Three. Figure 4 provides a comparison between the original and reimplemented results for this experiment. Both sets of results show that the objective fitness is being driven down, below the natural bias towards 50. This is more apparent in the reimplemented version. A reduction in the objective fitness in a dimension for an individual can lead

Parameter	Value
Sample Size (S)	15
Population Size	25
Dimensions	10
Dimension Length	10
Subjective Fitness Equation	eq2

Table 3: Parameters for reimplementing experiment two

Parameter	Value
Sample Size (S)	15
Population Size	25
Dimensions	10
Dimension Length	10
Subjective Fitness Equation	eq3

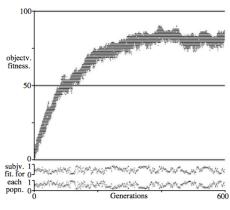
Table 4: Parameters for reimplementing experiment two

to improvements in subjective performance, explaining this behavior. Reducing objective performance can increase the difference in that dimension between two competing individuals and change the applicable dimension when calculating the subjective fitness.

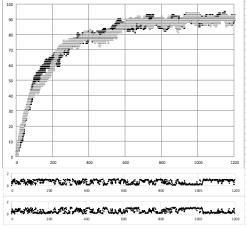
## 3 Coevolutionary Behavior With More Populations - Does this Fix Focussing?

The experiments from the original paper are extended to cover a coevolutionary set-up where more than 2 populations are used in the aim to fix the problem of over specialization illustrated in Experiment Two. The author hypothesizes that increasing the number of populations will lead to generalization in the case of Experiment Two. Drawing the sample, S, randomly from multiple populations (excluding the population that the individual is a part of) could lead to a situation where 'forgetting' objective improvement in a single dimension is detrimental to the individual's subjective performance.

No discernible improvement in the issues described in Experiment's One and Three is expected with an



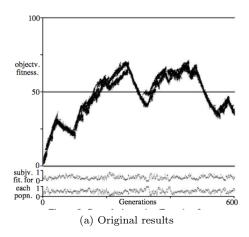




(b) Reimplemented Results

Figure 3: Reimplemented results for experiment two (original on top)

increase in the number of populations. For Experiment One (Figure 2), it is likely that there will still be periods of polarisation where some populations are dominant and others are not, leading to a downward drift towards the neutral performance position owing to a lack of selective pressure. The nature of the game defined for Experiment Three, means that intransitive superiority cannot be solved simply by increasing the number of populations. Objective performance levels can still be driven down to produce a gain in subjective performance.



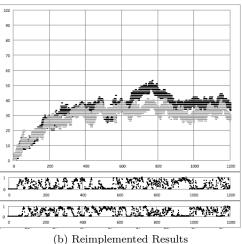


Figure 4: Reimplemented results for experiment three (original on top)

#### 3.1 Justification

A similar coevolutionary hypothesis is proposed by Zhal et al.[4], where multiple populations are used for multiple objectives, so the fitness in a population is defined by a single objective. However, this extension does not assign each population a specialist 'dimension' or 'objective', but aims to produce a generalist across all populations; so improvements learnt in one dimension are not forgotten, when improvements are made in another.

#### Experimental Set Up 3.2

The exact same parameters described in section 2.2

difference in the experimental set-up is that 10 populations are used instead of 2. When calculating the subjective fitness of an individual, the sample, S, consists of members randomly drawn from the remaining 9 populations. This has two consequences: 1) The random sample contains a greater genetic diversity 2) The selective pressure is drawn from a larger pool of individuals, whilst maintaining the independence of reproduction and selection between populations.

#### 3.3 Results

Figure 5 shows the coevolutionary dynamics for 10 competing populations. These results will be compared with **Figure 2**, hence, the parameters used are the same as those given in Table  ${\bf 2}$  apart from the sample size, S, which equals 1.

Figure 6b shows the results for Experiment Two with multiple populations. The objective fitness took longer than 1200 generations to converge, hence, in Figure 6a the original experiment is repeated for the same number of generations to provide a fair comparison. The parameters given in **Table 3** are used to produce both figures.

The extended results for Experiment Three are illustrated in **Figure 7** for 1 and 10 populations. The same parameters in **Table 4** are used to produce this figure.

#### 3.4 Discussion

Figure 5 shows an interesting dynamic. Unlike the case illustrated in Figure 2, the maximal objective fitness is reached by some populations. However, once this occurs, there is a general drift towards the natural bias, as expected. Figure 6 shows the subjective fitnesses for the 10 populations. As in Figure 2 some polarisation is observable. This is accompanied by the downward trend of some populations towards are used to produce the extended results. The only the natural bias. A particular matter of intrigue is

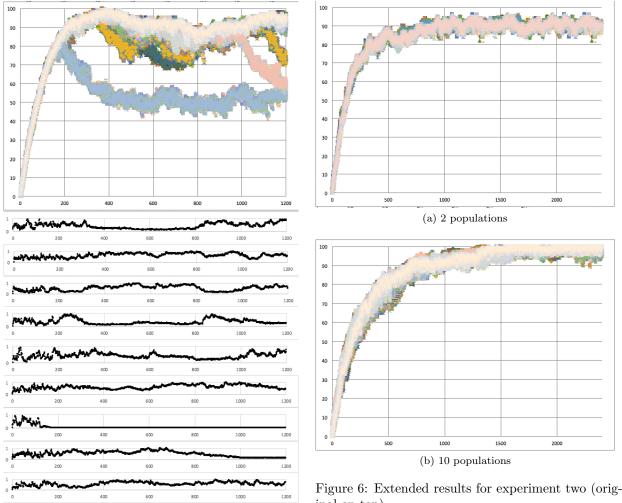


Figure 5: Extended Results for experiment one

the complete domination over population 7. A clear

divergence at around 200 generations can be seen in Figure 5, corroborated by the polarization of the subjective fitness in Figure 2. This reveals that if a population does not improve at fast enough rate objectively, it can be 'left behind' by the other populations which continue to improve and not drift to the same level as they play against other competing populations that are improving objectively. In the case of two populations domination by one population will be resolved as both start to drift towards the natural

Comparing the results in Figure 6, it is apparent

bias.

inal on top)

that introducing more populations to the coevolutionary game has lead to the optimal fitness across all dimensions being found. The solution took a greater number of generations to converge, but this is expected, as it takes longer to maintain optimal performance across all dimensions.

Figure 7 illustrates that, as predicted, there is not much improvement in the objective fitness of the populations. Intransitive superiority still exists between multiple populations. The populations continue to repeatedly travel the same part of strategy space.

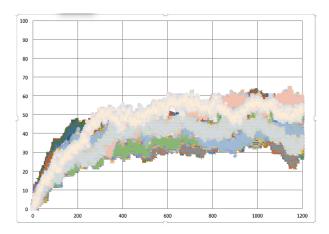


Figure 7: Extended Results for experiment three

### 4 Conclusion

The results from the original paper were successfully reimplemented and discussed. The parameters used for each experiment are clearly provided as well as the general approach taken. An extension to this paper was then proposed: Does increasing the number of populations in a coevolutionary set-up fix focusing? The experiments were repeated with 10 populations and compared against the results for two populations. As predicted, repeating Experiments One and Three illustrated the same issues of a loss of gradient and over specialization for more than two populations. Repeating Experiment Two, showed that increasing the number of populations does indeed solve the problem of over specialization or focussing in a coevolutionary set-up. Forgetting improvements in one dimension lead to poorer subjective performance as an individual could compete against a population containing specialists in that dimension in any generation.

### References

[1] Watson, Richard A. and Pollack, Jordan B., Spector, Lee(ed.) (2001) Coevolutionary Dynamics in a Minimal Substrate Proceedings of the Genetic and Evolutionary Computation Conference (GECCO 2001), pp. 702-709.

- [2] Cliff, D, & Miller, GF, 1995, "Tracking the Red Queen: Measurements of adaptive progress in coevolutionary simulations", Third European conference on Artificial Life, pp 200-218, Springer-Verlag, LNCS 929.
- [3] Ficici, SG. & Pollack, JB., 1998, "Challenges in Coevolutionary Learning: Arms-Race Dynamics, Open- Endedness, and Mediocre Stable States." Proceedings of the Sixth International Conference on Artificial Life. Adami, et al, eds. Cambridge: MIT, Press.
- [4] Zhi-Hui Zhan, Jingjing Li, Jiannong Cao, Jun Zhang, H. Chung and Yu-Hui Shi, "Multiple Populations for Multiple Objectives: A Coevolutionary Technique for Solving Multiobjective Optimization Problems", IEEE Transactions on Cybernetics, vol. 43, no. 2, pp. 445-463, 2013.
- [5] Fisher, RA, 1930, The genetical theory of natural selection, Clarendon press, oxford.

## **Appendix**

	Code 33	" " "
	34	= 1 if $=$ 1, $=$ 1.
1	fromfuture import division	otherwise
2	import random, string 35	0000000 II II II
3	from random import randrange, sample $_{36}$	$\mathbf{if} \ \ \mathbf{a} [0]. \ \mathbf{count} (1) > \mathbf{b} [0]. \ \mathbf{count}$
4	import time	(1):
5	import csv 37	return 1
6	from copy import deepcopy 38	else:
7	39	return 0
8	# LENGTH = 10 40	
9	$POP\_SIZE = 25$ 41	$def exp2\_score(a,b)$ :
10	$SAMPLE\_SIZE = 15$ 42	"""
11	# DIMENSIONS = 10 43	use_exp1_score(Ax,Bx)_where_x_
12	$MUTATION\_BIAS = 0.005$	is the dimension with the largest
13		difference_in_obj_fitness
14	def create_population(length, size, 44	
	dimensions):	# find dimension with largest
15	11 11 11	differnce
16	Creates_a_population_of_size_46	$largest\_diff = -1$
	size 'Each_member_consists_of_a_47	index = 0
	string_of_length 48	for i, dim in enumerate(a):
17	'length' bits. All bits are 49	diff = abs(dim.count)
	initialised_to_0.	(1) - b[i]. count (1)
18	5555555 " " " "	)
19	pop = [[0] * length for i in 50]	if diff > largest diff
	range(dimensions)   for j in	· ·
	range(size)] 51	$_{ m largest}$ diff $=$
20	return pop	diff
21	52	index = i
22	def obj_fitness(member): 53	# use that dimension to
23	"""	determine the score
24	Evaluates the objective 54	if a [index]. count $(1) > b$ [index
	fitness_of_a_member_by_counting_the	count(1):
	_number_of_1s_in_the_bitstring. 55	return 1
25	56	else:
26	$\#$ consider using coutner <b>if</b> $_{57}$	return 0
	this is too slow 58	return 0
27	fitness = 0    59	def exp3_score(a,b):
28	for dim in member: 60	$\frac{\text{def exps_score}(a,b)}{\  \  \ }.$
29	fitness = fitness + 61	use evel score (Av Rv) where v
	$\dim . \operatorname{count}(1)$	is_the_dimension_with_the_smallest_
30	return fitness	difference_in_obj_fitness
31	62	difference in Jobj Inthess
32	def exp1_score(a,b):	

```
63
           # find dimension with smallest
                                                                          member2 [0])
                                                              if eq == 2:
                differnce
                                          88
64
            smallest diff = abs(a[0].coun89
                                                                       fitness =
               (1) - b[0]. count(1)
                                                                           fitness +
            index = 0
                                                                          exp3 score(
65
            for i, dim in enumerate(a):
66
                                                                          member,
67
                    diff = abs(dim.count
                                                                          member2[0]
                        (1) - b[i].count(190
                                                              \# fitness = fitness +
                                                                  exp1 score (member,
                        )
68
                    if diff <
                                                                  member2[0]
                                                      return fitness
                        smallest diff:
                                          91
69
                             smallest diff92
                                = diff
                                          93
                            index = i
70
                                          94
                                              def exp2 subj fitness pop(popfit1,
                                                 popfit2, eq):
71
           # use that dimension to
               determine the score
                                          95
72
            if a[index].count(1) > b[inde 26]
                                             Evaluates the fitnesses of
                                                 each_member_in_an_entire_population
               [. count (1):
73
                    return 1
                                                 , using samples from
74
            else:
                                          97 ____another_population_and_the_'
                                                 exp2 subj fitness'_function.
75
                    return 0
76
                                             Returns the fitnesses as a
                                                 list.
77
   def exp2_subj_fitness(member, sample,
                                              ____"""
       eq):
                                          99
            11 11 11
78
                                         100
                                                      fitnesses = []
79
   Evaluates the subjective
                                                      for member in popfit1:
                                         101
       fitness_of_a_member_uisng_the_sum102
                                                               fitnesses.append(
                                                                  exp2 subj fitness (
       of_exp1 score_function
80
                                                                  member [0], random.
   against an input sample.
   sample (popfit2,
81
82
            fitness = 0
                                                                  SAMPLE SIZE), eq))
83
            for member2 in sample:
                                         103
                                                      # update popfit1 with new subj
                    if eq = 0:
                                                           fitnesses
84
                             fitness =
                                         104
                                                      # popfit1 = zip(list(zip(*
85
                                                          popfit1)), fitnesses)
                                fitness +
                                exp1 scorle05
                                                      return fitnesses
                                member, 106
                                member 2 [01]0)7
                                              def exp4 subj fitness pop(popfit index
86
                    if eq = 1:
                                                 , popfit, eq):
                                                      11 11 11
87
                             fitness =
                                         108
                                fitness +109
                                             ____Evaluates_the_fitnesses_of_
                                exp2 score(
                                                 each_member_in_an_entire_population
                                                 , \_using\_samples\_from
                                member,
```

```
if random.
110 ____all_other_populations_and_the35
        'exp2 subj fitness'_function.
                                                                              random() <
111
    Returns the fitnesses as a
                                                                              mutation rate
        list.
                                                                              :
    112
                                            136
                                                                                   #
113
             fitnesses = []
                                                                                       member
             other pops = []
114
                                                                                       [ i ]
             for i, pop in enumerate (popfit
115
                                                                                       =
                 ):
                                                                                       !
                      if i == popfit index:
116
                                                                                       member
                              continue
                                                                                       [ i ]
117
                      for member in pop:
                                                                                   dim[i]
118
                                           137
119
                              other pops.
                                                                                        =
                                                                                       0
                                  append (
                                                                                       i f
                                  member)
120
             for member in popfit [
                                                                                       random
                 popfit_index]:
121
                      fitnesses.append(
                                                                                       random
                         exp2 subj fitness (
                                                                                       ()
                         member [0], random.
                                                                                       <
                         sample (other pops,
                                                                                       0.5
                         SAMPLE SIZE), eq))
122
             return fitnesses
                                                                                       else
123
                                                                                        1
124
    def mutate member (member original,
                                                         return member
                                           138
        mutation rate):
                                            139
             11 11 11
125
                                            140
                                                def make new gen (pop fit,
126
                                                    mutation rate):
    ____Mutates_a_member_in_a_
        popoulation_with_probability_'
                                           141
                                                         new pop fit = []
        mutation rate'_that_a_bit_is
                                            142
                                                         for n in range (0, len (pop fit))
127
    ____assigned_a_new_random_value_(
                                                                 # select member for
        or_is_flipped).
                                            143
    ____"""
128
                                                                     mutation
129
             member = []
                                            144
                                                                  total = 0
                                                                  wheel = []
130
             member = deepcopy((
                                            145
                 member original))
                                                                  # create wheel
                                            146
             for dim in member:
                                                                  for member in pop fit:
131
                                            147
                      for i, bit in enumeral 48
                                                                          total = total
132
                         (dim):
                                                                              + member [1]
133
                              \# \operatorname{test} =
                                                                               + 1
                                  random. 149
                                                                          wheel.append(
                                  random()
                                                                              total)
134
                              # print test150
                                                                 # pick from wheel
```

151	pick = random.random()	selection to mutate
	* total	a new parent
152	for i, wheel_val in 171	$pop1\_fit =$
	${\tt enumerate (wheel):}$	$make\_new\_gen($
153	${f if}$ wheel_val	$\mathtt{pop1\_fit}$ ,
	>= pick:	MUTATION_BIAS)
154	inde <b></b> ≵72	$\mathtt{pop2\_fit} \ =$
	= i	${\rm make\_new\_gen}($
155	break	$\mathtt{pop2\_fit}$ ,
156	${ m new\_pop\_fit}$ . ${ m append}$ ( (	MUTATION_BIAS)
	$mutate\_member( 173$	# calculate new
	$pop_fit[index][0]$ ,	objective fitnesses
	$\operatorname{mutation\_rate})\;,\;\;0))$	$\mathbf{for}$ new
157	return new_pop_fit	populations
158	174	$obj\_fitnesses\_next \ = \\$
159	def run(length, dimensions, eq,	[]
	generations, filename): 175	for mem in pop1_fit:
160	$obj\_fitnesses = [[0 for i in 176]]$	$obj\_fitnesses\_next$
	$range(0,POP\_SIZE*2)]]$	$.\mathrm{append}($
161	# initialise populations and	$obj\_fitness$
	subj fitnesses	$(\operatorname{mem}\left[0\right]))$
162	pop1 = create_population( 177	$\mathbf{for}$ mem in $pop2$ _fit:
	length, POP_SIZE, 178	$obj\_fitnesses\_next$
	$\dim$ ensions)	$.\mathrm{append}($
163	${ m pop2}  =  { m create\_population}  ($	$obj\_fitness$
	length, POP_SIZE,	(mem [0])
	dimensions) 179	$\operatorname{obj}$ _ fitnesses . append (
164	$subj\_fitnesses1 = [[0 for i in$	$obj\_fitnesses\_next)$
	range (0, POP_SIZE)]] 180	# calc new subjective
165	$\mathrm{subj\_fitnesses2} = [[0 \ \mathbf{for} \ \mathrm{i} \ \mathrm{in}]]$	fitnesses for each
	range (0, POP_SIZE)]]	member in both
166	# create a list of tuples	populations
	containing members with 181	fitnesses1 =
	their subjective fitnesses	${\tt exp2\_subj\_fitness\_pop}$
167	$pop1\_fit = zip(pop1,$	(pop1_fit,pop2_fit,
	$\operatorname{subj\_fitnesses1}[0])$	$\mathrm{eq}$ )
168	$pop2_fit = zip(pop2,$ 182	$pop1_fit = zip(list($
	${\tt subj\_fitnesses2}[0])$	zip(*pop1_fit)[0]),
169	${f for}$ i in range (0, generations):	fitnesses1)
170	# make a new 183	fitnesses2 =
	generation - use	${\tt exp2\_subj\_fitness\_pop}$
	$\mathtt{fit}\mathtt{n}\mathtt{e}\mathtt{s}\mathtt{s}$	$(pop2\_fit, pop1\_fit,$
	proportionate	eq)

```
184
                      pop2 fit = zip(list(
                                                                      members with their
                          zip(*pop2 fit)[0]),
                                                                      subjective
                           fitnesses2)
                                                                      fitnesses
185
                      subj fitnesses1.appe2065
                                                                  pop fits[i] = zip(
                          (fitnesses1)
                                                                      create population (
                                                                      length, POP SIZE,
186
                      subj fitnesses2.append
                          (fitnesses2)
                                                                      dimensions),
             with open(filename + ".csv", "
187
                                                                      subj fitnesses
                 a") as fp:
                                                                      [0][0])
188
                      wr = csv.writer(fp, 206)
                          dialect='excel') 207
                                                          for i in range (0, generations):
                                                                   obj fitnesses next =
189
                      wr.writerows(
                                            208
                          obj fitnesses)
                                                                      \prod
190
             with open(filename + "
                                                                  for k, pop fit in
                                            209
                 subj fitnesses " + ".csv",
                                                                      enumerate (pop fits)
                 "a") as fp:
191
                      wr = csv.writer(fp, 210)
                                                                           # make a new
                          dialect='excel')
                                                                               generation
192
                      wr.writerows(
                                                                               – use
                          subj fitnesses1)
                                                                               fitness
                      wr.writerow("\n")
193
                                                                               proportionate
194
                      wr.writerows(
                                                                                selection
                          subj_fitnesses2)
                                                                               to mutate a
195
             return [obj fitnesses,
                                                                                new parent
                 subj fitnesses1,
                                                                           pop fits[k] =
                                            211
                 subj fitnesses2, pop1 fit,
                                                                               make new gen
                 pop2 fit]
                                                                               (pop fit,
196
                                                                               MUTATION BIAS
197
    def run2 (length, dimensions, eq, pops,
                                                                               )
         generations, filename):
                                                                           # calculate
                                            212
198
             obj fitnesses = [[0 \text{ for } i \text{ in }]]
                                                                               new
                 range(0,POP SIZE*pops)]]
                                                                               objective
199
             # initialise populations and
                                                                               fitnesses
                 subj fitnesses
                                                                               for new
             pop_fits = [0 for i in range
200
                                                                               populations
                 (0, pops)]
                                                                           for mem in
                                            213
             subj fitnesses = []
201
                                                                               pop fits [k
             for i in range (0, pops):
202
                                                                               ]:
203
                      subj fitnesses.appen@114
                                                                                    obj fitnesses next
                          ([[0 for j in range
                          (0,POP_SIZE)]])
                                                                                        append
204
                      # create a list of
```

obj fitness

tuples containing

```
(229)
                                                                           wr.writerow('
                                                                               new\_pop\ ')
                                           mem
                                           [238]0)
                                                         return [obj_fitnesses ,
                                           )
                                                             subj fitnesses]
215
                      obj fitnesses.append(
                          obj_fitnesses_next)
216
                      # calc new subjective
                          fitnesses for each
                         member in both
                          populations
                      for j in range (0, pops
217
                          ):
218
                               fitnesses =
                                  exp4\_subj\_fitness\_pop
                                   (j,
                                  pop fits,
                                  eq)
219
                               subj fitnesses
                                   [j].append(
                                   fitnesses)
220
                               pop_fits[j] =
                                  zip(list(
                                  zip(*
                                   pop_fits[j
                                   ])[0]),
                                   fitnesses)
221
222
             with open(filename + ".csv", "
                 a") as fp:
223
                      wr = csv.writer(fp,
                          dialect='excel')
224
                      wr.writerows(
                          obj fitnesses)
225
             with open(filename + "
                 _subj_fitnesses" + ".csv",
                 "a") as fp:
226
                      wr = csv.writer(fp,
                          dialect='excel')
227
                      for fitness in
                          subj fitnesses:
228
                               wr.writerows(
                                   fitness)
```