

Parameter Control and Adaptation

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1 Parameter Control and Adaptation

```
[1]: import random
import pprint
import math
import copy
import matplotlib.pyplot as plt

# For presenting as slides
#plt.rcParams['figure.figsize'] = [12, 8]
#plt.rcParams.update({'font.size': 22})
#plt.rcParams['lines.linewidth'] = 3
```

Evolutionary algorithms have many different parameters and implementation choices. There are qualitative parameters in the design of an algorithm (e.g., what representation, what recombination operators, what selection operators, etc.) and quantitative parameters (e.g., mutation rate, crossover rate, selection bias, population size, etc.) So far, we used intuition and common default values for many of these parameters. However, parameters can have a large influence on the performance of an algorithm on a particular problem. This chapter looks at how to optimise the selection of parameter values.

1.1 Example Problem: Pairwise Testing

As an example search problem we will consider pairwise testing, a form of combinatorial interaction testing where we want to generate covering arrays of strength 2. The problem is as follows: We are testing a system with x parameters, each parameter can have a different number of parameter values. In order to properly test the system, we would need to check all possible combinations of parameter values, but the combinatorial explosion usually makes this practically impossible. Since most failures are triggered by combinations of only few parameter values, the idea of pairwise testing is to try to cover all combinations of values for any pair of 2 parameters, which greatly reduces the number of tests.

We'll use a very basic genetic algorithm to solve the problem. We'll use our wrapper class because we will be caching some values later on:

```
[2]: class L(list):
    """
    A subclass of list that can accept additional attributes.
    Should be able to be used just like a regular list.
```

```

"""
def __new__(self, *args, **kwargs):
    return super(L, self).__new__(self, args, kwargs)

def __init__(self, *args, **kwargs):
    if len(args) == 1 and hasattr(args[0], '__iter__'):
        list.__init__(self, args[0])
    else:
        list.__init__(self, args)
    self.__dict__.update(kwargs)

def __call__(self, **kwargs):
    self.__dict__.update(kwargs)
    return self

```

Let's assume we have a system with a number of parameters, each with a number of possible values:

```

[3]: parameters = { 0 : ["a", "b", "c", "d", "x"],
                    1 : ["d", "e", "f", "g", "x"],
                    2 : ["g", "h", "i", "j", "x"],
                    3 : ["g", "h", "i", "j", "x"],
                    4 : ["g", "h", "i", "j", "x"],
                    5 : ["j", "k", "l", "m", "x"]}

```

A test case is a list of values, one for each parameter of our system:

```

[4]: def get_random_test():
    num_params = len(parameters)
    test = []
    for p in range(num_params):
        test.append(random.choice(parameters[p]))
    return test

```

The aim of our optimisation problem is to produce a set of tests that covers as many as possible combinations of parameter-value pairs, at the same time we don't want too many test cases. In principle we could make this a multi-objective problem and optimise for size and coverage, but to simplify our examples we will just assume a fixed number of tests, and try to cover as many as possible pairs with that exact number of tests:

```

[5]: num_tests = 25

```

An individual of our search is a list of `num_tests` of such tests:

```

[6]: def get_random_solution():
    solution = L([])
    for test in range(num_tests):
        solution.append(get_random_test())
    return solution

```

As a simple (though maybe somewhat computationally inefficient) fitness function we count how

many distinct pairs of pairwise parameter-value combinations we are covering by putting them in a set and then counting the size. The larger the set, the more pairs we have covered:

```
[7]: def get_fitness(individual):
    pairs = set()

    for num1 in range(len(parameters) - 1):
        for num2 in range(num1 + 1, len(parameters)):
            for row in individual:
                pairs.add((num1, num2, row[num1], row[num2]))

    return len(pairs)
```

In previous chapters, the parameters were dispersed throughout the notebooks with some global variables. Since we are focusing on parameters now, we will store them explicitly in a dictionary. Let's set some default values:

```
[8]: configuration = {
    "P_xover": 0.7,
    "P_mutation": 0.07,
    "population_size": 100,
    "tournament_size": 2
}
```

We are only optimising quantitative parameters, so the operators will be the ones we've used previously, adapted only to make use of our configuration dictionary:

```
[9]: def tournament_selection(population):
    # Make sure the sample isn't larger than the population
    candidates = random.sample(population, min(len(population),
    configuration["tournament_size"]))
    winner = max(candidates, key=lambda x: get_fitness(x))

    return winner
```

We'll use single-point crossover for tests, and just need to make sure to preserve our wrapper classes:

```
[10]: def crossover(parent1, parent2):
    pos = random.randint(1, len(parent1))

    offspring1 = L(copy.deepcopy(parent1[:pos] + parent2[pos:]))
    offspring2 = L(copy.deepcopy(parent2[:pos] + parent1[pos:]))

    return offspring1, offspring2
```

In the past we dynamically set the mutation rate based on the size of individuals. To make it easier to evaluate the effects of the mutation rate parameter, we will now set it explicitly. Each parameter in our list of tests is replaced with a different parameter value with that probability. We'll pass in the probability as a parameter rather and read the value of the configuration dictionary outside the operator.

```
[11]: def mutate(individual, P_mutate):
    mutated = L(copy.deepcopy(individual))
    for num_row in range(len(mutated)):
        for num_col in range(len(parameters)):
            if random.random() < P_mutate:
                choice = parameters[num_col][:]
                choice.remove(mutated[num_row][num_col])
                mutated[num_row][num_col] = random.choice(choice)

    return mutated
```

The reproduction-related probabilities are considered during a step of our algorithm:

```
[12]: def evolution_step(population):
    new_population = []

    while len(new_population) < len(population):
        parent1 = selection(population)
        parent2 = selection(population)

        if random.random() < configuration["P_xover"]:
            offspring1, offspring2 = crossover(parent1, parent2)
        else:
            offspring1, offspring2 = parent1, parent2

        offspring1 = mutate(offspring1, configuration["P_mutation"])
        offspring2 = mutate(offspring2, configuration["P_mutation"])

        new_population.append(offspring1)
        new_population.append(offspring2)

    population.clear()
    population.extend(new_population)
```

Finally, we'll just set some parameters and global variables for our experiments, such as the list of fitness values to observe what happened, and the number of fitness evaluations as our stopping criterion:

```
[13]: max_evaluations = 10000
    selection = tournament_selection
    fitness_values = []
```

The algorithm itself should by now be well known:

```
[14]: def ga():
    population = [get_random_solution() for _ in
    ↪range(configuration["population_size"])]
    best_individual = max(population, key=lambda k: get_fitness(k))
    best_fitness = get_fitness(best_individual)
```

```

print(f"Initial population, best fitness: {best_fitness}")

iteration = 0
while iteration < (max_evaluations / configuration["population_size"]):
    fitness_values.append(best_fitness)
    iteration += 1
    evolution_step(population)
    current_best = max(population, key=lambda k: get_fitness(k))
    current_fitness = get_fitness(current_best)
    if current_fitness > best_fitness:
        print(f"Iteration {iteration}, best fitness: {best_fitness}")
        best_individual = copy.deepcopy(current_best)
        best_fitness = current_fitness

    fitness_values.append(best_fitness)
return best_individual

```

We can now run our algorithm and look at the resulting test suite:

```

[15]: fitness_values = []
      solution = ga()

      # Pretty print output
      pp = pprint.PrettyPrinter(depth=6)
      solution.sort()
      pp.pprint(solution)
      print(len(solution))

```

```

Initial population, best fitness: 259
Iteration 1, best fitness: 259
Iteration 12, best fitness: 269
Iteration 56, best fitness: 271
[['a', 'd', 'g', 'i', 'i', 'l'],
 ['a', 'd', 'j', 'x', 'h', 'm'],
 ['a', 'e', 'h', 'g', 'x', 'k'],
 ['a', 'x', 'i', 'j', 'h', 'l'],
 ['b', 'd', 'h', 'j', 'x', 'x'],
 ['b', 'd', 'j', 'h', 'j', 'x'],
 ['b', 'e', 'h', 'g', 'g', 'x'],
 ['b', 'f', 'i', 'i', 'i', 'l'],
 ['b', 'g', 'x', 'h', 'x', 'x'],
 ['c', 'd', 'g', 'g', 'h', 'k'],
 ['c', 'd', 'g', 'j', 'x', 'j'],
 ['c', 'e', 'x', 'i', 'j', 'l'],
 ['c', 'f', 'h', 'j', 'g', 'x'],
 ['c', 'f', 'i', 'j', 'x', 'm'],
 ['c', 'f', 'x', 'j', 'j', 'k'],
 ['d', 'd', 'i', 'h', 'g', 'l'],

```

```

['d', 'f', 'j', 'j', 'i', 'x'],
['d', 'g', 'x', 'i', 'x', 'x'],
['d', 'x', 'x', 'h', 'h', 'j'],
['x', 'd', 'i', 'x', 'x', 'x'],
['x', 'e', 'g', 'h', 'x', 'm'],
['x', 'f', 'g', 'h', 'h', 'j'],
['x', 'g', 'j', 'i', 'g', 'l'],
['x', 'x', 'h', 'x', 'i', 'j'],
['x', 'x', 'i', 'g', 'j', 'm']]
25

```

```

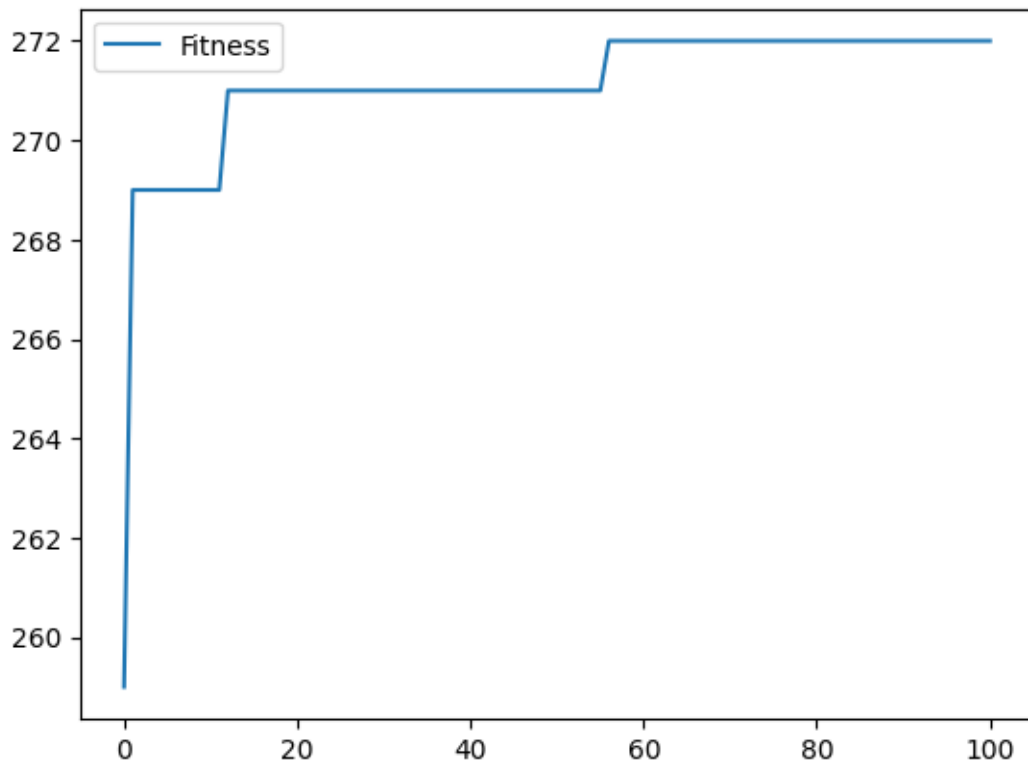
[16]: plt.plot(fitness_values, label="Fitness")
      plt.legend()

```

```

[16]: <matplotlib.legend.Legend at 0x112565c90>

```



1.2 Parameter Tuning

Tuning refers to a (systematic) process of optimising the values of the parameters of our algorithm. There are many considerations when doing so: Which parameters, and which levels? We can't look at all possible values for our parameters, so we will have to consider some relevant values. For example, let's consider the following parameters and their possible values:

```
[17]: search_parameters = {
    "population_size": [5, 37, 52, 67, 99],
    "tournament_size": [1, 2, 3, 4, 5],
    "P_mutation": [0.01, 0.34, 0.5, 0.65, 0.99],
    "P_xover": [0.01, 0.34, 0.5, 0.65, 0.99]
}
```

To find the best configuration based on these options, one way is to produce a factorial design, i.e., a combination of all possible values (very similar to the example problem we are solving with search in this chapter!) Let's define a function that produces such a factorial design:

```
[18]: def get_factorial_design(search_parameters):
    configurations = [{}]

    for parameter in search_parameters.keys():
        new_configurations = []
        for configuration in configurations:
            for option in range(len(search_parameters[parameter])):
                cp = configuration.copy()
                cp[parameter] = search_parameters[parameter][option]
                new_configurations.append(cp)

        configurations = new_configurations

    return configurations
```

For our choice of parameters and values, this is the resulting factorial design:

```
[19]: # This produces a loooong list...
    # get_factorial_design(search_parameters)
```

```
[20]: len(get_factorial_design(search_parameters))
```

```
[20]: 625
```

In order to find the best parameter setting, we would have to run the search on some example problems for each of these configurations. That would take very long, so one option is to produce a fractional factorial design where we omit or restrict certain combinations. Let's simply reduce the value choices to 2 per parameter:

```
[21]: search_parameters = {
    "population_size": [20, 100],
    "tournament_size": [1, 3],
    "P_mutation": [0.01, 0.5],
    "P_xover": [0.0, 0.7]
}
```

```
[22]: len(get_factorial_design(search_parameters))
```

[22]: 16

In order to evaluate how well each of the configurations performs, we need to define what *performance* actually means. A simple measurement of performance is given by the fitness value of the resulting solution. However, since the algorithm is randomised, we need to apply it repeatedly. Furthermore, we need to choose which problem instances to evaluate the algorithm on. To keep things simple, we will only consider the specific problem listed earlier, and we'll measure the performance as the average of the best fitness values for a number of repetitions:

```
[23]: from IPython.utils import io
def get_performance(config):
    global max_evaluations
    results = []
    repetitions = 4 # Should be more, but to accelerate the notebook...
    old_evaluations = max_evaluations
    max_evaluations = 3000 # Should be more, but to accelerate the notebook...
    for i in range(repetitions):
        fitness_values = []
        configuration.update(config)
        with io.capture_output() as captured:
            result = ga()
            fitness = get_fitness(result)
        results.append(fitness)
    max_evaluations = old_evaluations
    return sum(results)/len(results)
```

Now we just have to consider each configuration of our factorial design and measure the performance. The best configuration is the one with the best performance:

```
[24]: best_config = None
best_performance = 0
#max_evaluations = 10000 # Limit number of fitness evaluations so we don't have
↳to wait too long
for config in get_factorial_design(search_parameters):
    print(f"Current configuration: {config}")
    performance = get_performance(config)
    print(f"Performance: {performance}")
    if performance > best_performance:
        best_config = config
        best_performance = performance

print(f"Chosen best configuration: {best_config}")
configuration.update(best_config)
```

```
Current configuration: {'population_size': 20, 'tournament_size': 1,
'P_mutation': 0.01, 'P_xover': 0.0}
```

```
Performance: 261.25
```

```
Current configuration: {'population_size': 20, 'tournament_size': 1,
'P_mutation': 0.01, 'P_xover': 0.7}
```


Performance: 267.0
 Current configuration: {'population_size': 20, 'tournament_size': 1, 'P_mutation': 0.5, 'P_xover': 0.0}
 Performance: 266.0
 Current configuration: {'population_size': 20, 'tournament_size': 1, 'P_mutation': 0.5, 'P_xover': 0.7}
 Performance: 268.25
 Current configuration: {'population_size': 20, 'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.0}
 Performance: 300.25
 Current configuration: {'population_size': 20, 'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.7}
 Performance: 305.25
 Current configuration: {'population_size': 20, 'tournament_size': 3, 'P_mutation': 0.5, 'P_xover': 0.0}
 Performance: 267.25
 Current configuration: {'population_size': 20, 'tournament_size': 3, 'P_mutation': 0.5, 'P_xover': 0.7}
 Performance: 267.75
 Current configuration: {'population_size': 100, 'tournament_size': 1, 'P_mutation': 0.01, 'P_xover': 0.0}
 Performance: 263.75
 Current configuration: {'population_size': 100, 'tournament_size': 1, 'P_mutation': 0.01, 'P_xover': 0.7}
 Performance: 266.0
 Current configuration: {'population_size': 100, 'tournament_size': 1, 'P_mutation': 0.5, 'P_xover': 0.0}
 Performance: 268.0
 Current configuration: {'population_size': 100, 'tournament_size': 1, 'P_mutation': 0.5, 'P_xover': 0.7}
 Performance: 268.25
 Current configuration: {'population_size': 100, 'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.0}
 Performance: 291.25
 Current configuration: {'population_size': 100, 'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.7}
 Performance: 291.25
 Current configuration: {'population_size': 100, 'tournament_size': 3, 'P_mutation': 0.5, 'P_xover': 0.0}
 Performance: 268.0
 Current configuration: {'population_size': 100, 'tournament_size': 3, 'P_mutation': 0.5, 'P_xover': 0.7}
 Performance: 267.0
 Chosen best configuration: {'population_size': 20, 'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.7}

Finding vectors of parameter values is a complex optimisation task. Our performance function defines a *utility landscape* – an abstract landscape where the locations are the parameter vectors

of an EA and the height reflects the utility (performance). Thus, technically we can treat our tuning problem as a search problem. To do so, we'll define range limits for the parameters we are optimising:

```
[25]: search_parameter_ranges = {
    "population_size": [1, 100],
    "tournament_size": [1, 5],
    "P_mutation": [0.0, 0.2],
    "P_xover": [0.0, 1.0]
}
```

In order to apply meta-heuristic search to the problem of optimising the parameters of a meta-heuristic search algorithm, we need to define a representation, which is a parameter vector. We'll define a helper function that gives us a random value for a chosen parameter, using the range limits defined above:

```
[26]: def get_random_parameter(parameter_name):
    value = random.uniform(*search_parameter_ranges[parameter_name])
    if type(1) == type(search_parameter_ranges[parameter_name][0]):
        value = int(value)
    return value
```

An individual of our parameter-search is a dictionary of parameter values:

```
[27]: def get_random_configuration():
    config = {}
    for param in search_parameter_ranges.keys():
        config[param] = get_random_parameter(param)
    return config
```

```
[28]: get_random_configuration()
```

```
[28]: {'population_size': 36,
    'tournament_size': 2,
    'P_mutation': 0.08087349436294278,
    'P_xover': 0.5500358931778431}
```

A very basic mutation of parameter dictionaries is to replace individual parameters with different values.

```
[29]: def mutate_configuration(config):
    copy = config.copy()
    P_mutate = 1/len(copy)
    while copy == config:
        for key in copy.keys():
            if random.random() < P_mutate:
                copy[key] = get_random_parameter(key)
    return copy
```

Technically, we can use any search algorithm for this optimisation problem. However, measuring fitness requires running our `get_performance` function, which is expensive. Thus, some algorithms are better suited than others. One popular algorithm in practice is *differential evolution*, which we haven't covered yet, so to keep things simple we'll just use a (1+1)EA:

```
[30]: def oneplusone():
    current = best_config #get_random_configuration()
    fitness = get_performance(current)
    iteration = 0
    print(f"Iteration {iteration}: Fitness {fitness} - {current}")

    while iteration < max_steps:
        iteration += 1
        candidate = mutate_configuration(current)
        candidate_fitness = get_performance(candidate)
        print(f"Iteration {iteration}: Fitness {candidate_fitness} -
↪{candidate}")
        if candidate_fitness >= fitness:
            current = candidate
            fitness = candidate_fitness
            fitness_values.append(fitness)

    return current
```

```
[31]: max_steps = 20
tuned_opo = oneplusone()
tuned_opo
```

```
Iteration 0: Fitness 305.0 - {'population_size': 20, 'tournament_size': 3,
'P_mutation': 0.01, 'P_xover': 0.7}
Iteration 1: Fitness 316.25 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.01, 'P_xover': 0.7}
Iteration 2: Fitness 271.5 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.17412544485762843, 'P_xover': 0.7}
Iteration 3: Fitness 278.25 - {'population_size': 90, 'tournament_size': 4,
'P_mutation': 0.04170512760424956, 'P_xover': 0.7}
Iteration 4: Fitness 273.0 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.13774178729337586, 'P_xover': 0.7654648240337393}
Iteration 5: Fitness 309.5 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.01, 'P_xover': 0.23601978075990593}
Iteration 6: Fitness 304.0 - {'population_size': 20, 'tournament_size': 3,
'P_mutation': 0.01, 'P_xover': 0.7}
Iteration 7: Fitness 317.25 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.006448523301229781, 'P_xover': 0.7}
Iteration 8: Fitness 315.25 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.006448523301229781, 'P_xover': 0.6192003088149942}
Iteration 9: Fitness 262.25 - {'population_size': 9, 'tournament_size': 1,
'P_mutation': 0.006448523301229781, 'P_xover': 0.7}
```

```

Iteration 10: Fitness 269.25 - {'population_size': 53, 'tournament_size': 2,
'P_mutation': 0.16556268128001964, 'P_xover': 0.7}
Iteration 11: Fitness 270.25 - {'population_size': 20, 'tournament_size': 2,
'P_mutation': 0.12785415444421147, 'P_xover': 0.7}
Iteration 12: Fitness 315.75 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.006448523301229781, 'P_xover': 0.08618137683907989}
Iteration 13: Fitness 275.5 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.07902040893925338, 'P_xover': 0.7}
Iteration 14: Fitness 284.5 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.03659142926130086, 'P_xover': 0.16171271553605926}
Iteration 15: Fitness 279.0 - {'population_size': 57, 'tournament_size': 4,
'P_mutation': 0.050206000624864404, 'P_xover': 0.7}
Iteration 16: Fitness 290.0 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.025670218865608677, 'P_xover': 0.10277573846720711}
Iteration 17: Fitness 273.25 - {'population_size': 97, 'tournament_size': 4,
'P_mutation': 0.1256462864496599, 'P_xover': 0.03842289311892222}
Iteration 18: Fitness 317.0 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.006448523301229781, 'P_xover': 0.42918200505162385}
Iteration 19: Fitness 281.5 - {'population_size': 20, 'tournament_size': 4,
'P_mutation': 0.04911416429863549, 'P_xover': 0.7}
Iteration 20: Fitness 268.0 - {'population_size': 20, 'tournament_size': 1,
'P_mutation': 0.1615797844364074, 'P_xover': 0.7}

```

```

[31]: {'population_size': 20,
      'tournament_size': 4,
      'P_mutation': 0.006448523301229781,
      'P_xover': 0.7}

```

To reduce the enormous costs of fitness evaluations in this search, the *response surface methodology* systematically samples the utility space, creates a regression model that predicts the utility for parameter values, and then uses this regression model as a surrogate for querying the performance.

As an alternative to a full factorial design, the response surface methodology is often performed using a *central composite design*. For the central composite design we obtain five levels for each parameter: {0, 1, -1, , - }.

The design consists of three matrices: - A factorial design with two levels for each factor (1, -1) - A set of center points (0) - A set of axial points, configurations identical to the centre points except for one factor that is varied with values below and above the median (, -).

(The choice of 1, -1, , - is actually more complicated, but out of scope for this notebook)

For example, assume we have derived the following levels (- , -1, 0, 1,) for our factors:

```

[32]: search_parameters = {
      "population_size": [5, 37, 52, 67, 99],
      "tournament_size": [1, 2, 3, 4, 5],
      "P_mutation": [0.005, 0.01, 0.05, 0.1, 0.2],
      "P_xover": [0.01, 0.34, 0.5, 0.65, 0.99]
    }

```

We will define helper functions that produce the three component matrices:

```
[33]: def get_factorial_for(search_parameters, options = [1, 3]):
        configurations = [{}]
```

```
        for parameter in search_parameters.keys():
            new_configurations = []
            for configuration in configurations:
                for option in options:
                    cp = configuration.copy()
                    cp[parameter] = search_parameters[parameter][option]
                    new_configurations.append(cp)

            configurations = new_configurations

        return configurations
```

```
[34]: get_factorial_for(search_parameters)
```

```
[34]: [{'population_size': 37,
        'tournament_size': 2,
        'P_mutation': 0.01,
        'P_xover': 0.34},
        {'population_size': 37,
        'tournament_size': 2,
        'P_mutation': 0.01,
        'P_xover': 0.65},
        {'population_size': 37,
        'tournament_size': 2,
        'P_mutation': 0.1,
        'P_xover': 0.34},
        {'population_size': 37,
        'tournament_size': 2,
        'P_mutation': 0.1,
        'P_xover': 0.65},
        {'population_size': 37,
        'tournament_size': 4,
        'P_mutation': 0.01,
        'P_xover': 0.34},
        {'population_size': 37,
        'tournament_size': 4,
        'P_mutation': 0.01,
        'P_xover': 0.65},
        {'population_size': 37,
        'tournament_size': 4,
        'P_mutation': 0.1,
        'P_xover': 0.34},
        {'population_size': 37,
        'tournament_size': 4,
        'P_mutation': 0.1,
        'P_xover': 0.34}]
```

```
{'population_size': 37,
 'tournament_size': 4,
 'P_mutation': 0.1,
 'P_xover': 0.65},
{'population_size': 67,
 'tournament_size': 2,
 'P_mutation': 0.01,
 'P_xover': 0.34},
{'population_size': 67,
 'tournament_size': 2,
 'P_mutation': 0.01,
 'P_xover': 0.65},
{'population_size': 67,
 'tournament_size': 2,
 'P_mutation': 0.1,
 'P_xover': 0.34},
{'population_size': 67,
 'tournament_size': 2,
 'P_mutation': 0.1,
 'P_xover': 0.65},
{'population_size': 67,
 'tournament_size': 4,
 'P_mutation': 0.01,
 'P_xover': 0.34},
{'population_size': 67,
 'tournament_size': 4,
 'P_mutation': 0.01,
 'P_xover': 0.65},
{'population_size': 67,
 'tournament_size': 4,
 'P_mutation': 0.1,
 'P_xover': 0.34},
{'population_size': 67,
 'tournament_size': 4,
 'P_mutation': 0.1,
 'P_xover': 0.65}]
```

```
[35]: def get_center_point(parameters, center = 2):
        configuration = {}
        for parameter in parameters.keys():
            configuration[parameter] = parameters[parameter][center]
        return configuration
```

```
[36]: get_center_point(search_parameters)
```

```
[36]: {'population_size': 52,
        'tournament_size': 3,
```

```
'P_mutation': 0.05,  
'P_xover': 0.5}
```

```
[37]: def get_axial_points(parameters, axial_points = [0, 4]):  
    configurations = []  
  
    for parameter in parameters.keys():  
        for axial_point in axial_points:  
            configuration = get_center_point(parameters)  
            configuration[parameter] = parameters[parameter][axial_point]  
            configurations.append(configuration)  
    return configurations
```

```
[38]: get_axial_points(search_parameters)
```

```
[38]: [{ 'population_size': 5,  
        'tournament_size': 3,  
        'P_mutation': 0.05,  
        'P_xover': 0.5},  
       { 'population_size': 99,  
        'tournament_size': 3,  
        'P_mutation': 0.05,  
        'P_xover': 0.5},  
       { 'population_size': 52,  
        'tournament_size': 1,  
        'P_mutation': 0.05,  
        'P_xover': 0.5},  
       { 'population_size': 52,  
        'tournament_size': 5,  
        'P_mutation': 0.05,  
        'P_xover': 0.5},  
       { 'population_size': 52,  
        'tournament_size': 3,  
        'P_mutation': 0.005,  
        'P_xover': 0.5},  
       { 'population_size': 52,  
        'tournament_size': 3,  
        'P_mutation': 0.2,  
        'P_xover': 0.5},  
       { 'population_size': 52,  
        'tournament_size': 3,  
        'P_mutation': 0.05,  
        'P_xover': 0.01},  
       { 'population_size': 52,  
        'tournament_size': 3,  
        'P_mutation': 0.05,  
        'P_xover': 0.99}]
```

The full central composite design now consists of all three of these:

```
[39]: configurations = get_factorial_for(search_parameters) + \
      [ get_center_point(search_parameters) ] + \
      get_axial_points(search_parameters)
```

```
[40]: len(configurations)
```

```
[40]: 25
```

Now we need some patience while we determine the performance for each of these configurations:

```
[41]: responses = []
      max_evaluations = 10000
      for config in configurations:
          r = get_performance(config)
          config["performance"] = r
          print(config)
          responses.append(r)
```

```
{'population_size': 37, 'tournament_size': 2, 'P_mutation': 0.01, 'P_xover':
0.34, 'performance': 292.0}
{'population_size': 37, 'tournament_size': 2, 'P_mutation': 0.01, 'P_xover':
0.65, 'performance': 293.0}
{'population_size': 37, 'tournament_size': 2, 'P_mutation': 0.1, 'P_xover':
0.34, 'performance': 271.5}
{'population_size': 37, 'tournament_size': 2, 'P_mutation': 0.1, 'P_xover':
0.65, 'performance': 270.25}
{'population_size': 37, 'tournament_size': 4, 'P_mutation': 0.01, 'P_xover':
0.34, 'performance': 308.5}
{'population_size': 37, 'tournament_size': 4, 'P_mutation': 0.01, 'P_xover':
0.65, 'performance': 308.25}
{'population_size': 37, 'tournament_size': 4, 'P_mutation': 0.1, 'P_xover':
0.34, 'performance': 274.25}
{'population_size': 37, 'tournament_size': 4, 'P_mutation': 0.1, 'P_xover':
0.65, 'performance': 274.0}
{'population_size': 67, 'tournament_size': 2, 'P_mutation': 0.01, 'P_xover':
0.34, 'performance': 287.25}
{'population_size': 67, 'tournament_size': 2, 'P_mutation': 0.01, 'P_xover':
0.65, 'performance': 289.0}
{'population_size': 67, 'tournament_size': 2, 'P_mutation': 0.1, 'P_xover':
0.34, 'performance': 271.0}
{'population_size': 67, 'tournament_size': 2, 'P_mutation': 0.1, 'P_xover':
0.65, 'performance': 271.5}
{'population_size': 67, 'tournament_size': 4, 'P_mutation': 0.01, 'P_xover':
0.34, 'performance': 302.5}
{'population_size': 67, 'tournament_size': 4, 'P_mutation': 0.01, 'P_xover':
0.65, 'performance': 306.25}
{'population_size': 67, 'tournament_size': 4, 'P_mutation': 0.1, 'P_xover':
```



```

0.34, 'performance': 272.75}
{'population_size': 67, 'tournament_size': 4, 'P_mutation': 0.1, 'P_xover':
0.65, 'performance': 272.5}
{'population_size': 52, 'tournament_size': 3, 'P_mutation': 0.05, 'P_xover':
0.5, 'performance': 277.75}
{'population_size': 5, 'tournament_size': 3, 'P_mutation': 0.05, 'P_xover': 0.5,
'performance': 277.5}
{'population_size': 99, 'tournament_size': 3, 'P_mutation': 0.05, 'P_xover':
0.5, 'performance': 274.0}
{'population_size': 52, 'tournament_size': 1, 'P_mutation': 0.05, 'P_xover':
0.5, 'performance': 268.75}
{'population_size': 52, 'tournament_size': 5, 'P_mutation': 0.05, 'P_xover':
0.5, 'performance': 280.5}
{'population_size': 52, 'tournament_size': 3, 'P_mutation': 0.005, 'P_xover':
0.5, 'performance': 306.0}
{'population_size': 52, 'tournament_size': 3, 'P_mutation': 0.2, 'P_xover': 0.5,
'performance': 270.5}
{'population_size': 52, 'tournament_size': 3, 'P_mutation': 0.05, 'P_xover':
0.01, 'performance': 276.75}
{'population_size': 52, 'tournament_size': 3, 'P_mutation': 0.05, 'P_xover':
0.99, 'performance': 278.75}

```

The next step is to create a regression model that predicts the performance of any given combination of parameter values. We'll cut this short and simply throw our data at a multilayer perceptron regression model, and doing this properly and checking the fit of the model are left as an exercise.

```

[42]: import numpy as np
import pandas as pd
from sklearn.neural_network import MLPRegressor

```

```

[43]: model = MLPRegressor(solver="lbfgs", max_iter=1000)

```

```

[44]: df = pd.DataFrame(configurations)
y = df['performance'].values
x = df[['population_size', 'tournament_size', 'P_mutation', 'P_xover']].values
# This will need some adjusting...
model.fit(x, y)

```

```

[44]: MLPRegressor(max_iter=1000, solver='lbfgs')

```

We can now take any configuration and query the model. For example, let's query the configuration we determined best with our simplified factorial design initially:

```

[45]: model.predict([list(best_config.values())])

```

```

[45]: array([335.82585836])

```

We'll use a (1+1)EA again to do the search, but instead of querying the fitness function, we query the model:

```
[46]: def surrogate_oneplusone():
    current = best_config #get_random_configuration()
    fitness = model.predict([list(current.values())])[0]
    iteration = 0
    print(f"Iteration {iteration}: Fitness {fitness} - {current}")

    while iteration < max_steps:
        iteration += 1
        candidate = mutate_configuration(current)
        candidate_fitness = model.predict([list(candidate.values())])[0]
        if candidate_fitness >= fitness:
            if candidate_fitness > fitness:
                print(f"Iteration {iteration}: Fitness {candidate_fitness} - {candidate}")
                current = candidate
                fitness = candidate_fitness
            fitness_values.append(fitness)

    return current
```

Since querying the model is very cheap compared to measuring the actual performance, we can run many more iterations of the search algorithm.

```
[47]: max_steps = 1000
rsm_config = surrogate_oneplusone()
```

```
Iteration 0: Fitness 335.8258583555003 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.7}
Iteration 15: Fitness 337.72768919123956 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.7887918298287592}
Iteration 21: Fitness 341.16426931993766 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.01, 'P_xover': 0.9492373501379531}
Iteration 25: Fitness 342.24718147424284 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.0043827291783720135, 'P_xover':
0.9492373501379531}
Iteration 42: Fitness 342.37054209518993 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.0043827291783720135, 'P_xover':
0.9549967557890329}
Iteration 95: Fitness 343.05556033562635 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.0043827291783720135, 'P_xover':
0.9869785818754603}
Iteration 228: Fitness 343.0693309114329 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.004311298588763424, 'P_xover':
0.9869785818754603}
Iteration 433: Fitness 343.2821416845018 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.004311298588763424, 'P_xover':
0.996914196333372}
```

```
Iteration 513: Fitness 343.59030364491974 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.0027128040278620036, 'P_xover':
0.996914196333372}
Iteration 535: Fitness 344.07462109700106 - {'population_size': 20,
'tournament_size': 3, 'P_mutation': 0.00020055750706422516, 'P_xover':
0.996914196333372}
```

```
[48]: rsm_config
```

```
[48]: {'population_size': 20,
'tournament_size': 3,
'P_mutation': 0.00020055750706422516,
'P_xover': 0.996914196333372}
```

Is the optimised configuration really better? Let's find out by comparing against our initial default values.

```
[49]: configuration['P_xover'] = 0.7
configuration['P_mutation'] = 0.07
configuration['population_size'] = 100
configuration['tournament_size'] = 2

fitness_values = []
ga()
# Stretch this by the population size to make plots comparable
default_fitness = [item for item in fitness_values for i in
↳range(configuration['population_size'])]

configuration.update(tuned_opo)
fitness_values = []
ga()
# Stretch this by the population size to make plots comparable
opo_tuned_fitness = [item for item in fitness_values for i in
↳range(configuration['population_size'])]

configuration.update(rsm_config)
fitness_values = []
ga()
# Stretch this by the population size to make plots comparable
rsm_tuned_fitness = [item for item in fitness_values for i in
↳range(configuration['population_size'])]
```

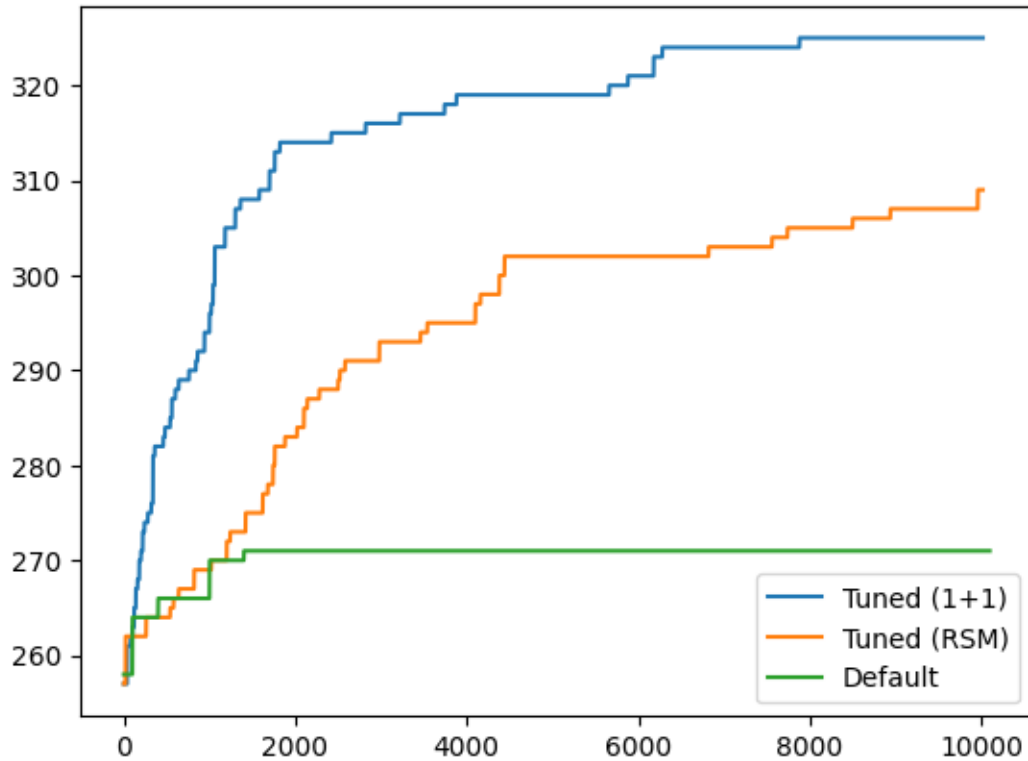
```
Initial population, best fitness: 258
Iteration 1, best fitness: 258
Iteration 4, best fitness: 264
Iteration 10, best fitness: 266
Iteration 14, best fitness: 270
```

Initial population, best fitness: 257
Iteration 2, best fitness: 257
Iteration 4, best fitness: 261
Iteration 5, best fitness: 262
Iteration 6, best fitness: 263
Iteration 7, best fitness: 265
Iteration 8, best fitness: 267
Iteration 9, best fitness: 268
Iteration 10, best fitness: 270
Iteration 11, best fitness: 271
Iteration 12, best fitness: 273
Iteration 14, best fitness: 274
Iteration 16, best fitness: 275
Iteration 17, best fitness: 276
Iteration 18, best fitness: 281
Iteration 23, best fitness: 282
Iteration 24, best fitness: 283
Iteration 27, best fitness: 284
Iteration 28, best fitness: 285
Iteration 30, best fitness: 287
Iteration 32, best fitness: 288
Iteration 38, best fitness: 289
Iteration 42, best fitness: 290
Iteration 43, best fitness: 291
Iteration 47, best fitness: 292
Iteration 50, best fitness: 294
Iteration 51, best fitness: 296
Iteration 52, best fitness: 297
Iteration 53, best fitness: 299
Iteration 59, best fitness: 303
Iteration 65, best fitness: 305
Iteration 68, best fitness: 307
Iteration 79, best fitness: 308
Iteration 85, best fitness: 309
Iteration 88, best fitness: 311
Iteration 91, best fitness: 313
Iteration 121, best fitness: 314
Iteration 141, best fitness: 315
Iteration 161, best fitness: 316
Iteration 187, best fitness: 317
Iteration 194, best fitness: 318
Iteration 283, best fitness: 319
Iteration 294, best fitness: 320
Iteration 309, best fitness: 321
Iteration 314, best fitness: 323
Iteration 394, best fitness: 324
Initial population, best fitness: 257
Iteration 1, best fitness: 257

```
Iteration 13, best fitness: 262
Iteration 27, best fitness: 264
Iteration 29, best fitness: 265
Iteration 32, best fitness: 266
Iteration 41, best fitness: 267
Iteration 51, best fitness: 269
Iteration 60, best fitness: 270
Iteration 62, best fitness: 272
Iteration 71, best fitness: 273
Iteration 81, best fitness: 275
Iteration 84, best fitness: 277
Iteration 87, best fitness: 278
Iteration 88, best fitness: 280
Iteration 94, best fitness: 282
Iteration 101, best fitness: 283
Iteration 105, best fitness: 284
Iteration 107, best fitness: 286
Iteration 114, best fitness: 287
Iteration 125, best fitness: 288
Iteration 126, best fitness: 289
Iteration 129, best fitness: 290
Iteration 149, best fitness: 291
Iteration 173, best fitness: 293
Iteration 177, best fitness: 294
Iteration 205, best fitness: 295
Iteration 208, best fitness: 297
Iteration 219, best fitness: 298
Iteration 222, best fitness: 300
Iteration 341, best fitness: 302
Iteration 378, best fitness: 303
Iteration 387, best fitness: 304
Iteration 425, best fitness: 305
Iteration 447, best fitness: 306
Iteration 498, best fitness: 307
```

```
[50]: plt.plot(opo_tuned_fitness, label="Tuned (1+1)")
      plt.plot(rsm_tuned_fitness, label="Tuned (RSM)")
      plt.plot(default_fitness, label="Default")
      plt.legend()
```

```
[50]: <matplotlib.legend.Legend at 0x1685d73d0>
```



1.3 Parameter Control

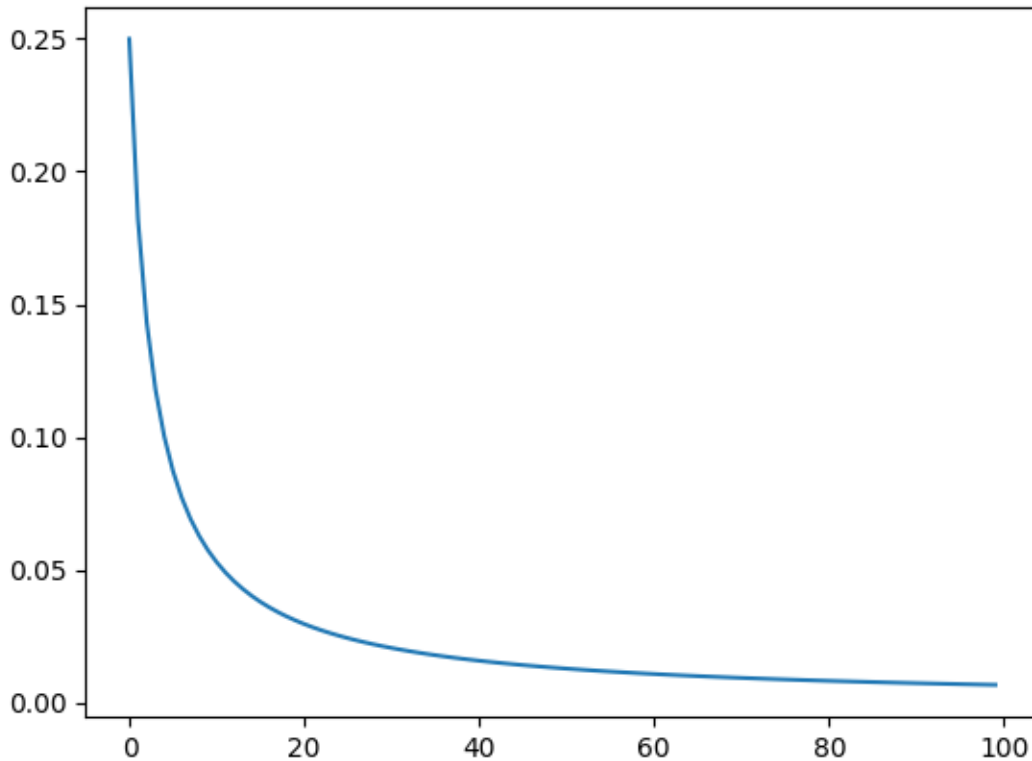
There's actually something fundamentally missing in the tuning experiments: We did all our tuning on the exact same problem. That's obviously not what we would do in practice: In order to find good parameters we performed search on that problem; once we have found a good set of parameters we know that it is a good set of parameters because we have found a good solution already, so we don't need to run the search again on that problem. What is missing in our experiments is that we would in practice of course do the tuning on several problems, and then hope that these parameters generalise to other, new problems.

A further limitation of parameter tuning is that it tries to select a good set of parameter values, but these parameters remain constant throughout the search. However, different phases of the search may benefit from different parameter values. For example, during initial phases of the search we may want to put more focus on exploration, thus using higher mutation rates than in later phases (like we saw with the MIO algorithm). Thus, a basic approach to control parameters is to make the parameter values dependent on the phase of the search:

```
[51]: def calculate_probability(size, iteration, max_iterations):
      return math.pow(4 + (size - 2)/(max_iterations - 1) * iteration, -1)

[52]: mutation_values = [calculate_probability(num_tests*len(parameters), x, 100) for
      ↪ x in range(100)]
      plt.plot(mutation_values)
```

[52]: [<matplotlib.lines.Line2D at 0x168713400>]



We'll now create a version of the genetic algorithm which doesn't use the mutation rate from our configuration dictionary, but derives it using our probability function:

```
[53]: def controlled_evolution_step(population, P_mutation):
    new_population = []

    while len(new_population) < len(population):
        parent1 = selection(population)
        parent2 = selection(population)

        if random.random() < configuration["P_xover"]:
            offspring1, offspring2 = crossover(parent1, parent2)
        else:
            offspring1, offspring2 = parent1, parent2

        offspring1 = mutate(offspring1, P_mutation)
        offspring2 = mutate(offspring2, P_mutation)

        new_population.append(offspring1)
        new_population.append(offspring2)
```

```

population.clear()
population.extend(new_population)

```

```

[54]: mutation_values = []
def controlled_ga():
    population = [get_random_solution() for _ in
↳range(configuration["population_size"])]
    best_individual = max(population, key=lambda k: get_fitness(k))
    best_fitness = get_fitness(best_individual)
    print(f"Initial population, best fitness: {best_fitness}")

    iteration = 0
    max_iterations = (max_evaluations / configuration["population_size"])
    while iteration < max_iterations:
        fitness_values.append(best_fitness)
        P_mutation = calculate_probability(num_tests * len(parameters),
↳iteration, max_iterations)
        mutation_values.append(P_mutation)
        iteration += 1
        controlled_evolution_step(population, P_mutation)
        current_best = max(population, key=lambda k: get_fitness(k))
        current_fitness = get_fitness(current_best)
        if current_fitness > best_fitness:
            print(f"Iteration {iteration}, best fitness: {best_fitness}")
            best_individual = copy.deepcopy(current_best)
            best_fitness = current_fitness

    fitness_values.append(best_fitness)
    return best_individual

```

As point of reference, we will compare against our initial default values again:

```

[55]: fitness_values = []
mutation_values = []
max_evaluations = 10000

configuration['P_xover'] = 0.7
configuration['P_mutation'] = 0.07
configuration['population_size'] = 100
configuration['tournament_size'] = 2

controlled_ga()
controlled_fitness = fitness_values

fitness_values = []
ga()

```



```

Initial population, best fitness: 259
Iteration 3, best fitness: 259
Iteration 6, best fitness: 263
Iteration 9, best fitness: 264
Iteration 11, best fitness: 268
Iteration 17, best fitness: 271
Iteration 28, best fitness: 272
Iteration 34, best fitness: 273
Iteration 35, best fitness: 274
Iteration 42, best fitness: 275
Iteration 44, best fitness: 276
Iteration 47, best fitness: 278
Iteration 52, best fitness: 280
Iteration 61, best fitness: 281
Iteration 64, best fitness: 282
Iteration 68, best fitness: 283
Iteration 73, best fitness: 285
Iteration 75, best fitness: 286
Iteration 79, best fitness: 287
Iteration 82, best fitness: 288
Iteration 88, best fitness: 290
Iteration 89, best fitness: 291
Iteration 90, best fitness: 292
Iteration 93, best fitness: 294
Iteration 95, best fitness: 295
Iteration 96, best fitness: 296
Initial population, best fitness: 264
Iteration 3, best fitness: 264
Iteration 21, best fitness: 268
Iteration 28, best fitness: 269

```

```

[55]: [['d', 'g', 'x', 'g', 'h', 'k'],
       ['d', 'x', 'h', 'x', 'x', 'm'],
       ['b', 'f', 'i', 'j', 'j', 'l'],
       ['a', 'g', 'x', 'x', 'h', 'l'],
       ['c', 'g', 'x', 'h', 'h', 'x'],
       ['x', 'd', 'i', 'g', 'x', 'k'],
       ['x', 'e', 'i', 'g', 'j', 'l'],
       ['a', 'f', 'i', 'j', 'g', 'm'],
       ['b', 'g', 'g', 'i', 'j', 'j'],
       ['b', 'd', 'h', 'j', 'h', 'j'],
       ['d', 'e', 'i', 'j', 'h', 'm'],
       ['c', 'x', 'h', 'g', 'i', 'l'],
       ['d', 'd', 'i', 'g', 'g', 'j'],
       ['c', 'e', 'x', 'j', 'x', 'x'],
       ['a', 'f', 'x', 'i', 'h', 'm'],
       ['x', 'x', 'j', 'i', 'h', 'm'],

```

```

['b', 'f', 'j', 'x', 'x', 'k'],
['d', 'x', 'g', 'x', 'j', 'j'],
['a', 'd', 'j', 'x', 'x', 'l'],
['b', 'd', 'x', 'h', 'j', 'l'],
['a', 'e', 'i', 'i', 'g', 'x'],
['x', 'g', 'g', 'h', 'g', 'x'],
['d', 'f', 'g', 'g', 'x', 'x'],
['c', 'e', 'i', 'i', 'g', 'k'],
['b', 'e', 'j', 'h', 'i', 'm']]

```

```

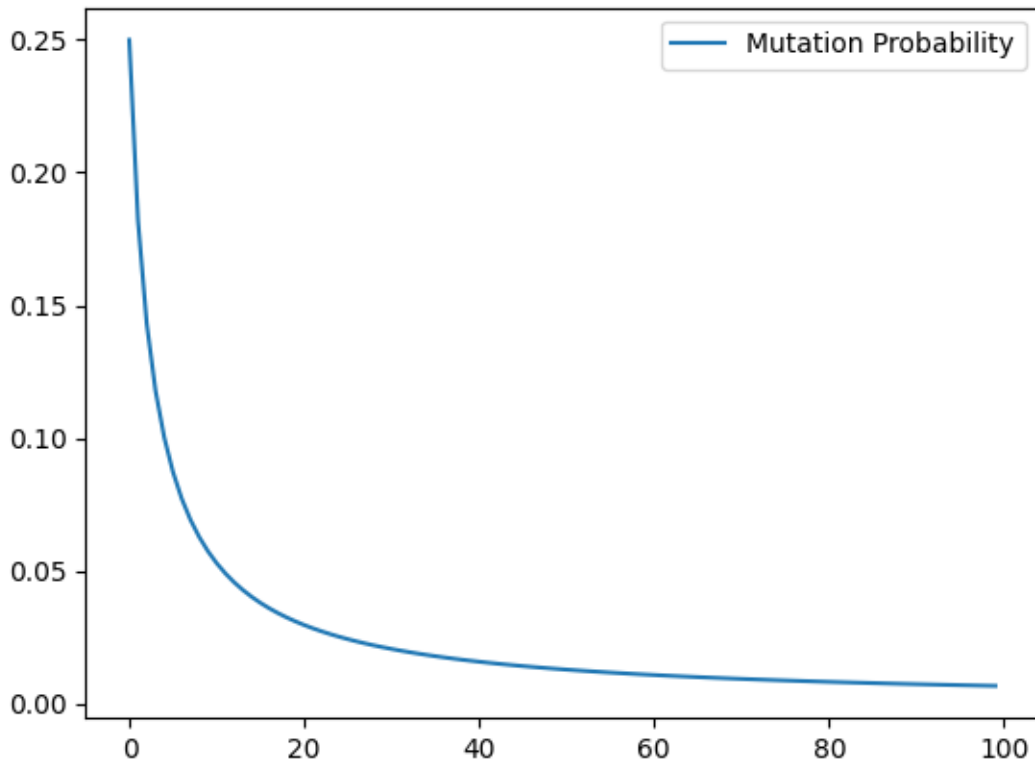
[56]: plt.plot(mutation_values, label="Mutation Probability")
      plt.legend()

```

```

[56]: <matplotlib.legend.Legend at 0x168710e80>

```



```

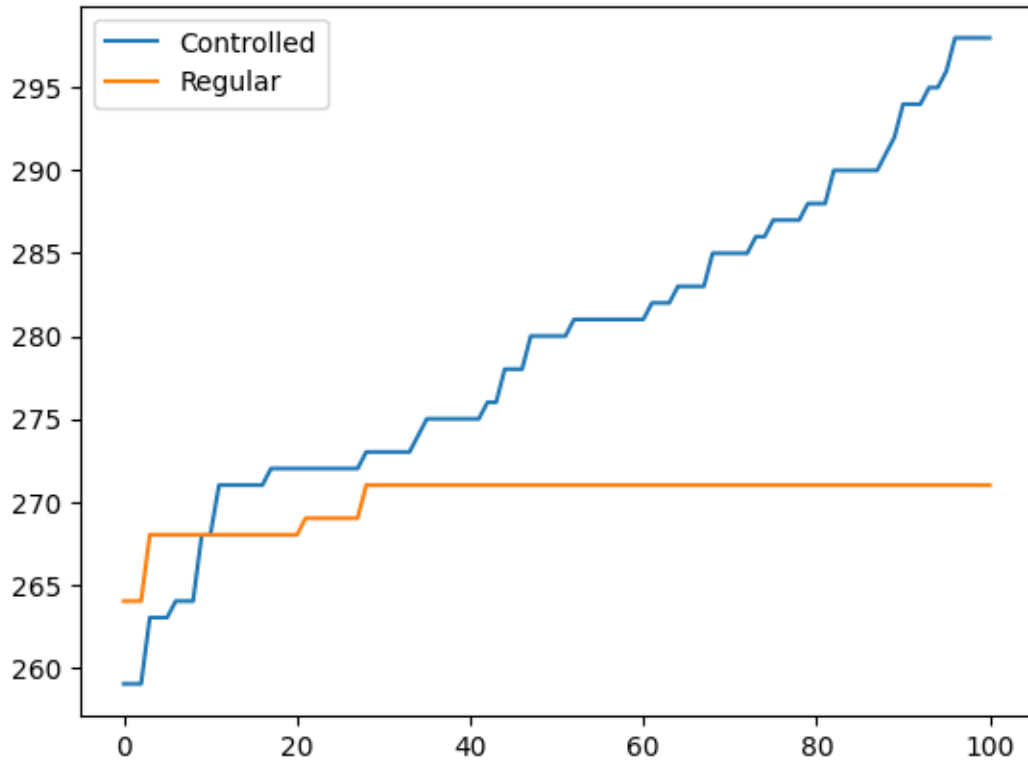
[57]: plt.plot(controlled_fitness, label="Controlled")
      plt.plot(fitness_values, label="Regular")
      plt.legend()

```

```

[57]: <matplotlib.legend.Legend at 0x168710bb0>

```



1.4 Adaptive Evolutionary Algorithms

The function that we used to control the mutation rate depends only on the phase of the search, but does not consider how the search is performing at any particular point in time. The idea of *adaptive* parameter control is to adjust parameters based on how well the search is going. As a basic example, if mutation leads to more fitness improvement than crossover, we may want to increase the probability of doing mutation, and reduce the probability of doing crossover, and vice versa. In order to do this, we need to keep track of the fitness values before and after applying these operators:

```
[58]: def adaptive_crossover(parent1, parent2):
    fitness_pre = get_fitness(parent1) + get_fitness(parent2)
    offspring1, offspring2 = crossover(parent1, parent2)
    fitness_post = get_fitness(offspring1) + get_fitness(offspring2)
    xover_performance.append(fitness_post - fitness_pre)
    return offspring1, offspring2
```

```
[59]: def adaptive_mutate(individual, P_mutate):
    fitness_pre = get_fitness(individual)
    mutated = mutate(individual, P_mutate)
    fitness_post = get_fitness(mutated)
    mutation_performance.append(fitness_post - fitness_pre)
```

```
return mutated
```

Note that, technically, we are introducing additional fitness evaluations. However, to some degree these could be avoided (though not all) using some proper caching of fitness values.

```
[60]: mutation_performance = []
      xover_performance = []

      adaptive_values = []
      xover_values = []
      mutation_values = []
```

The individual evolution step is the same as usual, we just need to make sure it uses the correct parameters:

```
[61]: def adaptive_evolution_step(population, mutation_probability,
      ↪xover_probability):
      new_population = []

      while len(new_population) < len(population):
          parent1 = selection(population)
          parent2 = selection(population)

          if random.random() < xover_probability:
              offspring1, offspring2 = adaptive_crossover(parent1, parent2)
          else:
              offspring1, offspring2 = parent1, parent2

          offspring1 = adaptive_mutate(offspring1, mutation_probability)
          offspring2 = adaptive_mutate(offspring2, mutation_probability)

          new_population.append(offspring1)
          new_population.append(offspring2)

      population.clear()
      population.extend(new_population)
```

After each step we need to re-evaluate the performance of the operators, and decide whether and how to update the parameters.

```
[62]: def adaptive_ga():
      population = [get_random_solution() for _ in
      ↪range(configuration["population_size"])]
      best_individual = max(population, key=lambda k: get_fitness(k))
      best_fitness = get_fitness(best_individual)
      print(f"Initial population, best fitness: {best_fitness}")

      iteration = 0
```

```

max_iterations = (max_evaluations / configuration["population_size"])
xover_probability = 0.7
mutation_probability = 0.07
delta = 0.01

while iteration < max_iterations:
    mutation_values.append(mutation_probability)
    xover_values.append(xover_probability)
    fitness_values.append(best_fitness)
    iteration += 1
    adaptive_evolution_step(population, mutation_probability,
↪xover_probability)

    current_best = max(population, key=lambda k: get_fitness(k))
    current_fitness = get_fitness(current_best)
    if current_fitness > best_fitness:
        print(f"Iteration {iteration}, best fitness: {best_fitness}")
        best_individual = copy.deepcopy(current_best)
        best_fitness = current_fitness

    max_mutation = max(mutation_performance) if mutation_performance else 0
    max_xover = max(xover_performance) if xover_performance else 0

    if max_mutation < max_xover:
        xover_probability = min(xover_probability + delta, 1.0)
        mutation_probability = max(mutation_probability - delta, 0.01)
    elif max_xover < max_mutation:
        xover_probability = max(xover_probability - delta, 0.0)
        mutation_probability = min(mutation_probability + delta, 1.0)

    mutation_performance.clear()
    xover_performance.clear()

    fitness_values.append(best_fitness)
    return best_individual

```

We'll compare how this performs compared to our initial default configuration again.

```

[63]: fitness_values = []
      num_mutations = []

      configuration['P_xover'] = 0.7
      configuration['P_mutation'] = 0.07
      configuration['population_size'] = 100
      configuration['tournament_size'] = 2

      mutation_values = []

```

```

xover_values = []
adaptive_ga()
adaptive_fitness = fitness_values

fitness_values = []
ga()

```

```

Initial population, best fitness: 261
Iteration 1, best fitness: 261
Iteration 14, best fitness: 267
Iteration 29, best fitness: 270
Iteration 37, best fitness: 272
Iteration 43, best fitness: 273
Iteration 44, best fitness: 274
Iteration 45, best fitness: 276
Iteration 46, best fitness: 282
Iteration 60, best fitness: 284
Iteration 61, best fitness: 286
Iteration 67, best fitness: 287
Iteration 82, best fitness: 290
Iteration 84, best fitness: 291
Iteration 91, best fitness: 292
Iteration 92, best fitness: 294
Initial population, best fitness: 254
Iteration 1, best fitness: 254
Iteration 2, best fitness: 262
Iteration 4, best fitness: 264

```

```

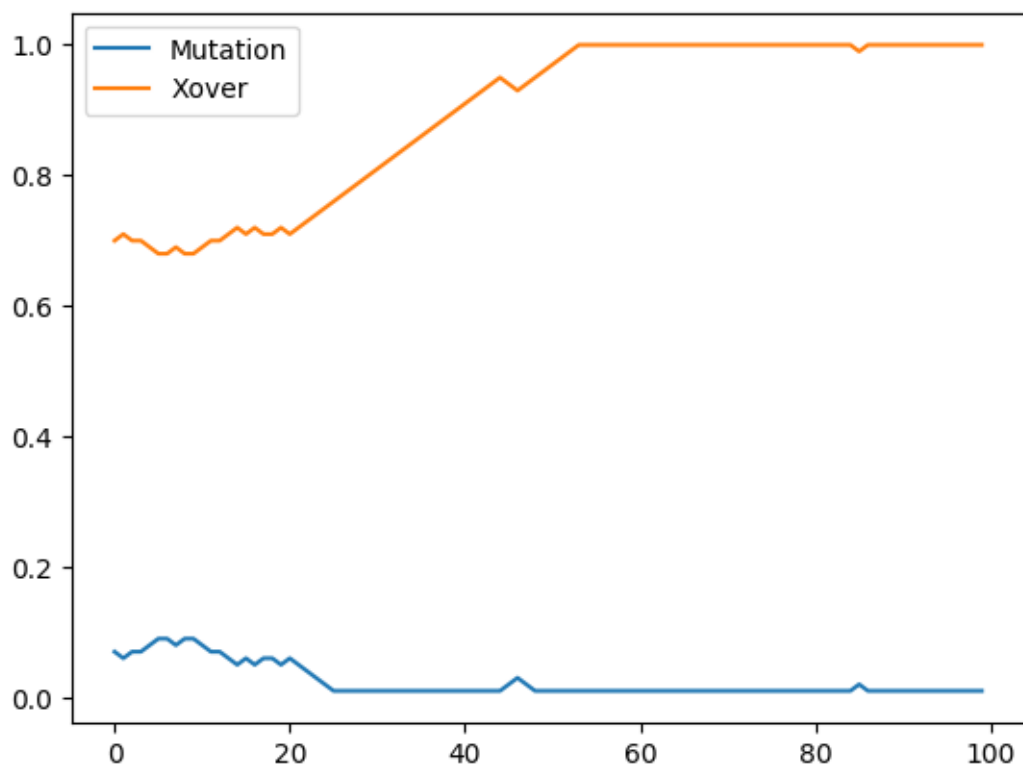
[63]: [['c', 'g', 'i', 'g', 'j', 'x'],
      ['x', 'g', 'h', 'i', 'x', 'm'],
      ['d', 'e', 'g', 'x', 'g', 'k'],
      ['d', 'g', 'x', 'h', 'x', 'x'],
      ['c', 'g', 'g', 'x', 'i', 'm'],
      ['d', 'f', 'x', 'g', 'j', 'j'],
      ['x', 'x', 'j', 'g', 'h', 'x'],
      ['x', 'd', 'i', 'i', 'x', 'j'],
      ['c', 'e', 'x', 'j', 'h', 'k'],
      ['d', 'd', 'g', 'x', 'j', 'x'],
      ['d', 'd', 'x', 'h', 'j', 'l'],
      ['c', 'e', 'i', 'j', 'h', 'l'],
      ['c', 'e', 'j', 'h', 'i', 'm'],
      ['b', 'd', 'h', 'j', 'g', 'x'],
      ['a', 'x', 'i', 'h', 'g', 'l'],
      ['b', 'd', 'i', 'x', 'i', 'k'],
      ['a', 'f', 'g', 'i', 'g', 'm'],
      ['a', 'f', 'x', 'j', 'i', 'm'],
      ['x', 'f', 'g', 'g', 'x', 'l'],

```

```
['c', 'f', 'j', 'x', 'g', 'l'],
['b', 'e', 'h', 'i', 'h', 'j'],
['b', 'g', 'x', 'g', 'j', 'k'],
['d', 'g', 'h', 'x', 'h', 'k'],
['x', 'd', 'x', 'g', 'g', 'j'],
['d', 'f', 'i', 'i', 'h', 'm']]
```

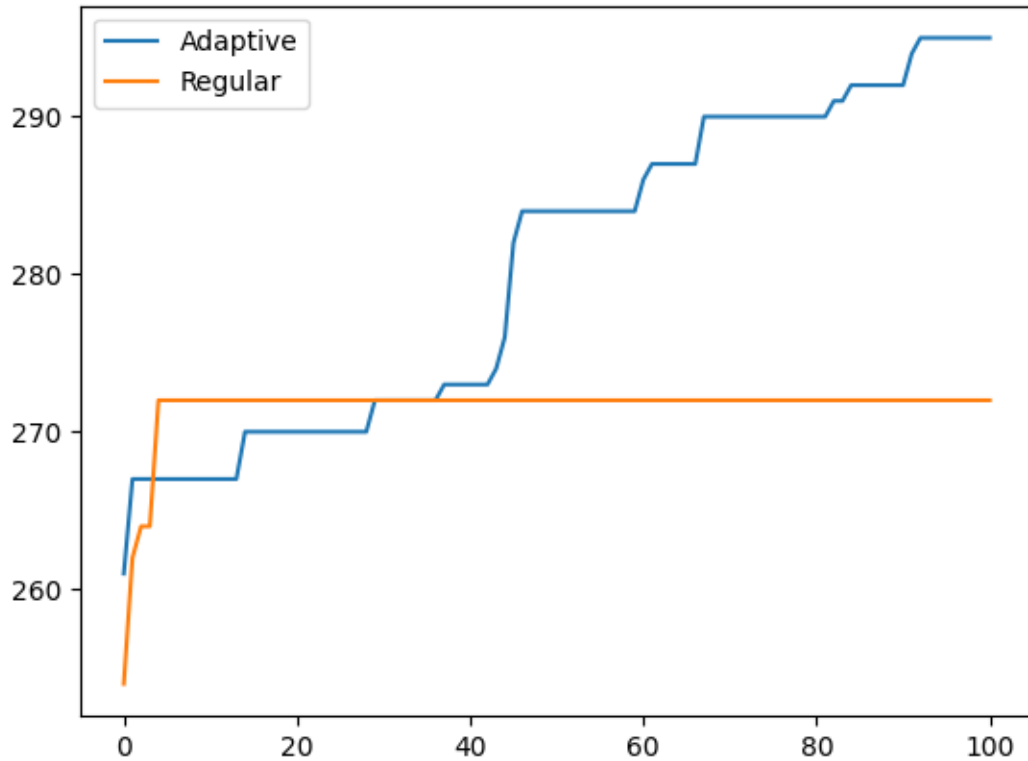
```
[64]: plt.plot(mutation_values, label="Mutation")
plt.plot(xover_values, label="Xover")
plt.legend()
```

[64]: <matplotlib.legend.Legend at 0x16876b820>



```
[65]: plt.plot(adaptive_fitness, label="Adaptive")
plt.plot(fitness_values, label="Regular")
plt.legend()
```

[65]: <matplotlib.legend.Legend at 0x168887370>



1.5 Self-Adaptive Evolutionary Algorithms

Adaptive evolutionary algorithms control some parameters, but thereby introduce new ones (e.g., when and how and how much to update the controlled parameters). What should we set these new parameters to? And should these values be constant, or also adaptive? To avoid this problem altogether we can just let our algorithm decide on its own what values to use, and turn it into a *self-adaptive* evolutionary algorithm.

We thus add the parameters we want the algorithm to adapt to the representation:

```
[66]: def get_random_solution():
    solution = L([])
    for test in range(num_tests):
        solution.append(get_random_test())

    solution.proBABILITIES = {"P_xover" : random.random(),
                              "P_mutation" : random.random()}

    return solution
```

When crossing over two individuals, for simplicity we will simply let them inherit their parents' parameter values:


```
[67]: def self_crossover(parent1, parent2):
    offspring1, offspring2 = crossover(parent1, parent2)

    # Make sure copies have the probabilities
    offspring1.probabilities = parent1.probabilities.copy()
    offspring2.probabilities = parent2.probabilities.copy()

    return offspring1, offspring2
```

Now comes the important bit: When mutating individuals, we not only alter the values (i.e., the test data in our search problem), but also the parameter values. We will add some Gaussian noise to mutate parameters by a little bit, so here's a helper function to keep values in bounds when applying the Gaussian noise:

```
[68]: def crop(value, boundary_min, boundary_max):
    if value > boundary_max:
        return boundary_max
    elif value < boundary_min:
        return boundary_min
    else:
        return value
```

The actual mutation just calls our usual mutation, and then applies the Gaussian noise – using the current mutation probability to decide whether or not to mutate each parameter:

```
[69]: def self_mutate(individual, P_mutate):

    mutated = mutate(individual, P_mutate)
    mutated.probabilities = individual.probabilities.copy()

    for p in mutated.probabilities.keys():
        if random.random() < P_mutate:
            mutated.probabilities[p] = crop(random.gauss(mutated.
↪probabilities[p], 0.05), 0.001, 1)

    return mutated
```

When given two parents, we use the average of the crossover probabilities of the parents as our crossover probability; when mutating an individual, we use its own mutation probability:

```
[70]: def self_evolution_step(population):
    new_population = []
    while len(new_population) < len(population):
        parent1 = selection(population)
        parent2 = selection(population)

        P_xover = (parent1.probabilities["P_xover"] + parent2.
↪probabilities["P_xover"])/2
```

```

        if random.random() < P_xover:
            offspring1, offspring2 = self_crossover(parent1, parent2)
        else:
            offspring1, offspring2 = parent1, parent2

        offspring1 = self_mutate(offspring1, offspring1.
↪probabilities["P_mutation"])
        offspring2 = self_mutate(offspring2, offspring2.
↪probabilities["P_mutation"])

        new_population.append(offspring1)
        new_population.append(offspring2)

    population.clear()
    population.extend(new_population)

```

The evolutionary algorithm itself is unchanged, but we add some code to keep track of the average parameters in the population so we can look at that after the evolution:

```

[71]: def self_adaptive_ga():
    population = [get_random_solution() for _ in
↪range(configuration["population_size"])]
    best_individual = max(population, key=lambda k: get_fitness(k))
    best_fitness = get_fitness(best_individual)
    print(f"Initial population, best fitness: {best_fitness}")

    iteration = 0
    max_iterations = (max_evaluations / configuration["population_size"])

    while iteration < max_iterations:
        fitness_values.append(best_fitness)
        mutation_values.append(sum([p.probabilities["P_mutation"] for p in
↪population])/len(population))
        xover_values.append(sum([p.probabilities["P_xover"] for p in
↪population])/len(population))
        iteration += 1
        self_evolution_step(population)

        current_best = max(population, key=lambda k: get_fitness(k))
        current_fitness = get_fitness(current_best)
        if current_fitness > best_fitness:
            print(f"Iteration {iteration}, best fitness: {best_fitness}")
            best_individual = copy.deepcopy(current_best)
            best_fitness = current_fitness

        fitness_values.append(best_fitness)
    return best_individual

```

Once again we compare against our initial default values.

```
[72]: max_evaluations = 10000
      fitness_values = []
      configuration['P_xover'] = 0.7
      configuration['P_mutation'] = 0.07
      configuration['population_size'] = 100
      configuration['tournament_size'] = 2

      self_values = []
      xover_values = []
      mutation_values = []

      self_adaptive_ga()
      self_values = fitness_values

      fitness_values = []
      ga()
```

Initial population, best fitness: 259

Iteration 1, best fitness: 259

Iteration 2, best fitness: 261

Iteration 5, best fitness: 263

Iteration 6, best fitness: 264

Iteration 11, best fitness: 267

Iteration 13, best fitness: 268

Iteration 14, best fitness: 269

Iteration 17, best fitness: 270

Iteration 18, best fitness: 272

Iteration 21, best fitness: 277

Iteration 22, best fitness: 279

Iteration 23, best fitness: 280

Iteration 27, best fitness: 282

Iteration 28, best fitness: 283

Iteration 34, best fitness: 285

Iteration 36, best fitness: 287

Iteration 39, best fitness: 288

Iteration 42, best fitness: 289

Iteration 47, best fitness: 290

Iteration 48, best fitness: 291

Iteration 50, best fitness: 292

Iteration 51, best fitness: 293

Iteration 55, best fitness: 294

Iteration 60, best fitness: 296

Iteration 61, best fitness: 297

Iteration 62, best fitness: 298

Iteration 67, best fitness: 299

Iteration 69, best fitness: 300

```

Iteration 73, best fitness: 301
Iteration 77, best fitness: 303
Iteration 86, best fitness: 304
Iteration 97, best fitness: 305
Initial population, best fitness: 258
Iteration 1, best fitness: 258
Iteration 3, best fitness: 259
Iteration 4, best fitness: 261
Iteration 7, best fitness: 264
Iteration 8, best fitness: 266
Iteration 11, best fitness: 267
Iteration 19, best fitness: 268
Iteration 22, best fitness: 271
Iteration 76, best fitness: 272
Iteration 88, best fitness: 273

```

```

[72]: [['x', 'f', 'x', 'x', 'h', 'x'],
      ['c', 'f', 'i', 'x', 'g', 'm'],
      ['d', 'f', 'i', 'x', 'i', 'x'],
      ['c', 'f', 'j', 'i', 'h', 'l'],
      ['c', 'f', 'x', 'j', 'x', 'j'],
      ['b', 'g', 'x', 'i', 'i', 'l'],
      ['a', 'f', 'x', 'g', 'i', 'k'],
      ['c', 'x', 'x', 'g', 'j', 'k'],
      ['a', 'd', 'g', 'x', 'x', 'l'],
      ['a', 'x', 'g', 'g', 'j', 'm'],
      ['x', 'e', 'j', 'g', 'x', 'j'],
      ['x', 'g', 'j', 'h', 'j', 'j'],
      ['x', 'x', 'x', 'h', 'h', 'm'],
      ['b', 'f', 'h', 'g', 'j', 'm'],
      ['d', 'f', 'g', 'i', 'g', 'm'],
      ['c', 'e', 'g', 'g', 'i', 'j'],
      ['b', 'd', 'g', 'h', 'x', 'k'],
      ['x', 'f', 'i', 'g', 'g', 'l'],
      ['a', 'g', 'g', 'j', 'h', 'x'],
      ['c', 'd', 'i', 'i', 'j', 'x'],
      ['b', 'x', 'h', 'x', 'g', 'x'],
      ['c', 'e', 'h', 'g', 'h', 'k'],
      ['b', 'f', 'i', 'h', 'x', 'k'],
      ['d', 'd', 'j', 'j', 'g', 'm'],
      ['b', 'g', 'i', 'j', 'g', 'j']]

```

```

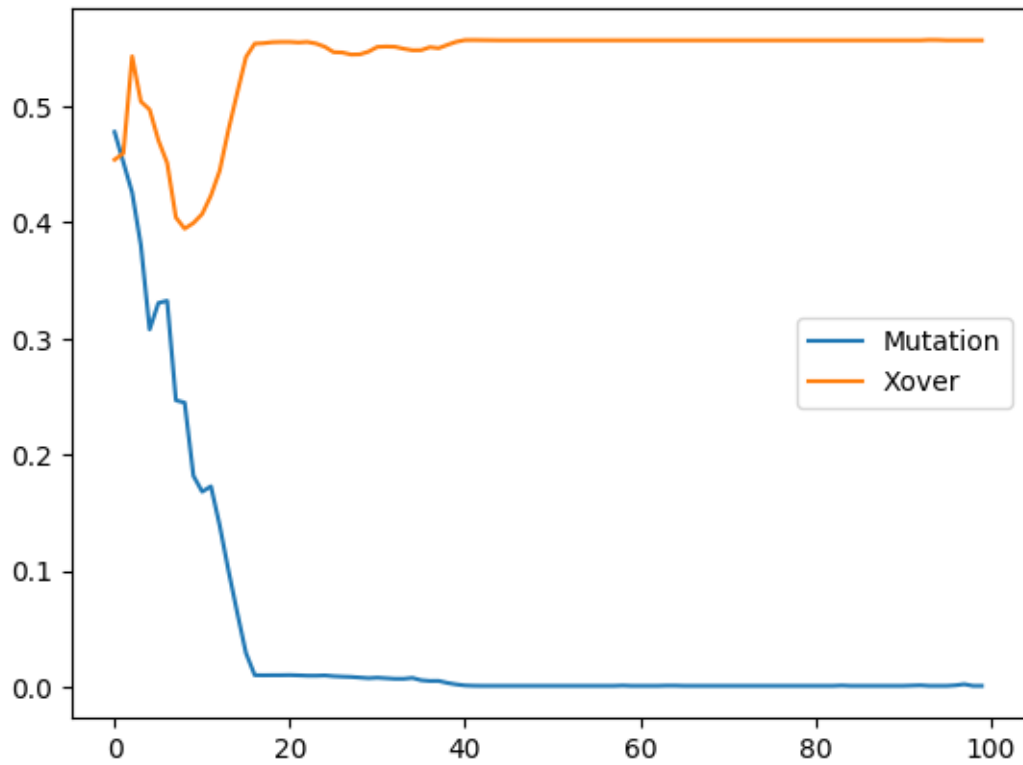
[73]: plt.plot(mutation_values, label="Mutation")
      plt.plot(xover_values, label="Xover")
      plt.legend()

```

```

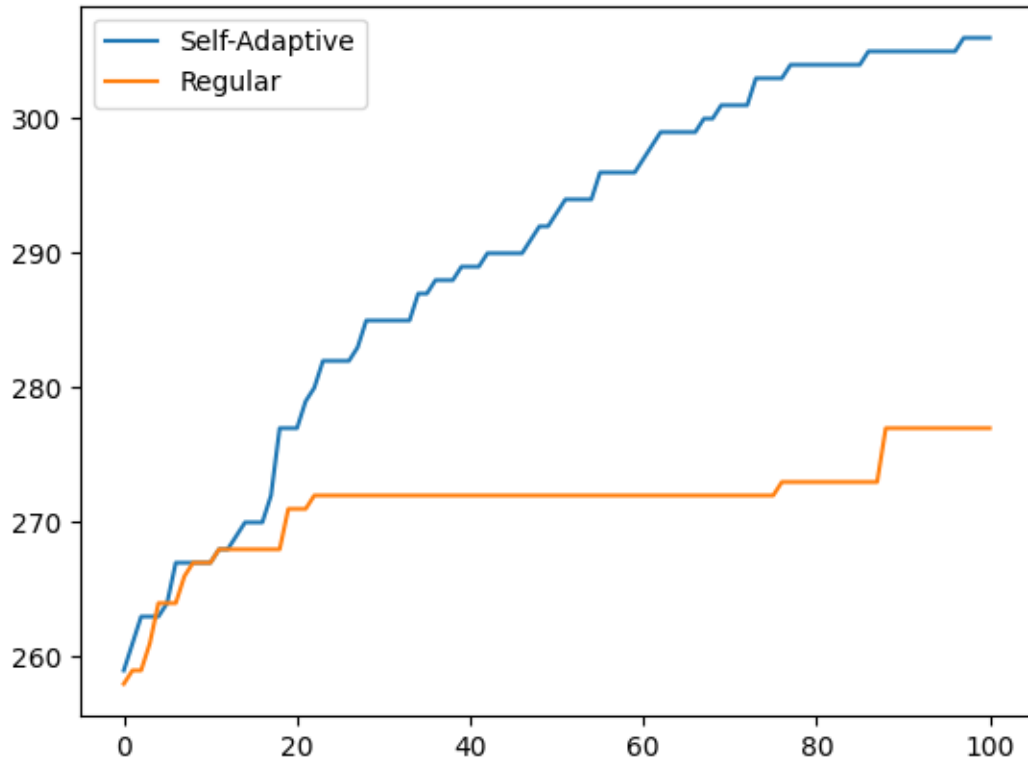
[73]: <matplotlib.legend.Legend at 0x168885ba0>

```



```
[74]: plt.plot(self_values, label="Self-Adaptive")  
      plt.plot(fitness_values, label="Regular")  
      plt.legend()
```

```
[74]: <matplotlib.legend.Legend at 0x168884190>
```



1.6 Generalisation

Throughout this notebook, when we compared algorithms we only looked at individual runs; it's difficult to generalise from an individual run. Furthermore, we always applied the search to the exact same problem. In order to better compare the different configurations, we need to consider *different* problems.

```
[75]: from IPython.utils import io

def run_times(algorithm, configuration, repetitions):
    global fitness_values
    result = []
    for i in range(repetitions):
        fitness_values = []
        with io.capture_output() as captured:
            algorithm()
        result.append(fitness_values[-1])
    return result
```

```
[76]: parameters = { 0 : ["a", "b", "c", "d"],
                      1 : ["d", "e", "f", "g", "x"],
                      2 : ["g", "h", "i"]}
```

```

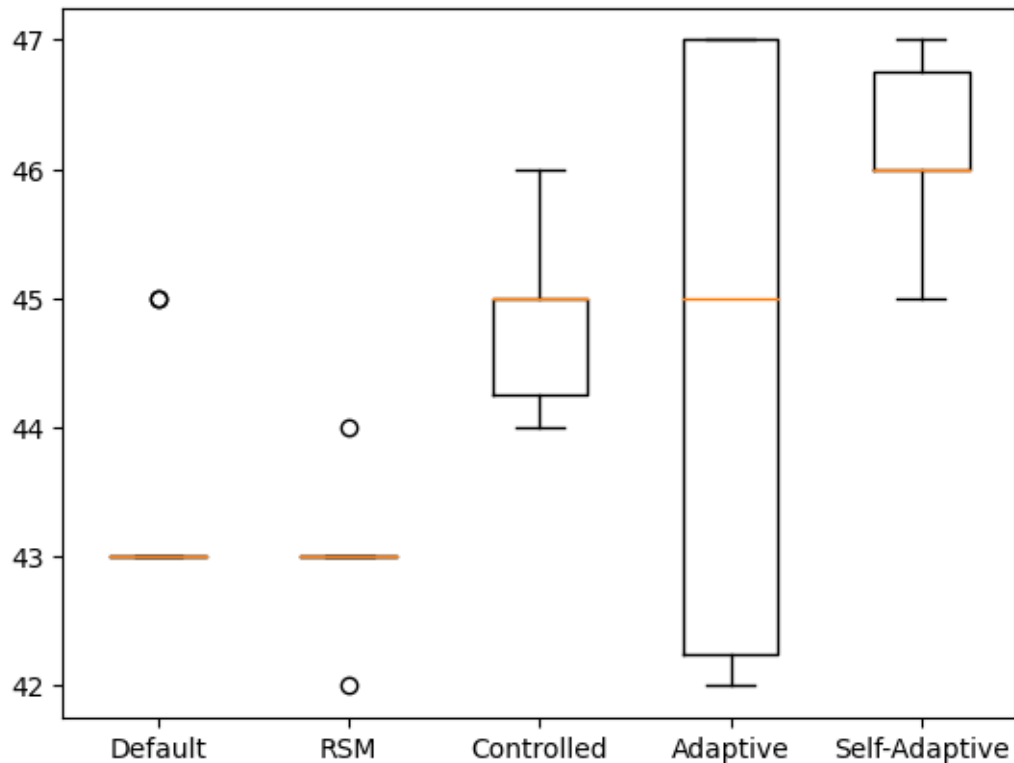
num_tests = 20
configuration['P_xover'] = 0.7
configuration['P_mutation'] = 0.07
configuration['population_size'] = 100
configuration['tournament_size'] = 2
results = {
    "Default"      : run_times(ga, configuration, 10),
    "RSM"          : run_times(ga, rsm_config, 10),
    "Controlled"   : run_times(controlled_ga, configuration, 10),
    "Adaptive"     : run_times(adaptive_ga, configuration, 10),
    "Self-Adaptive": run_times(self_adaptive_ga, configuration, 10)
}
fig, ax = plt.subplots()
ax.boxplot(results.values())
ax.set_xticklabels(results.keys())

```

```

[76]: [Text(1, 0, 'Default'),
      Text(2, 0, 'RSM'),
      Text(3, 0, 'Controlled'),
      Text(4, 0, 'Adaptive'),
      Text(5, 0, 'Self-Adaptive')]

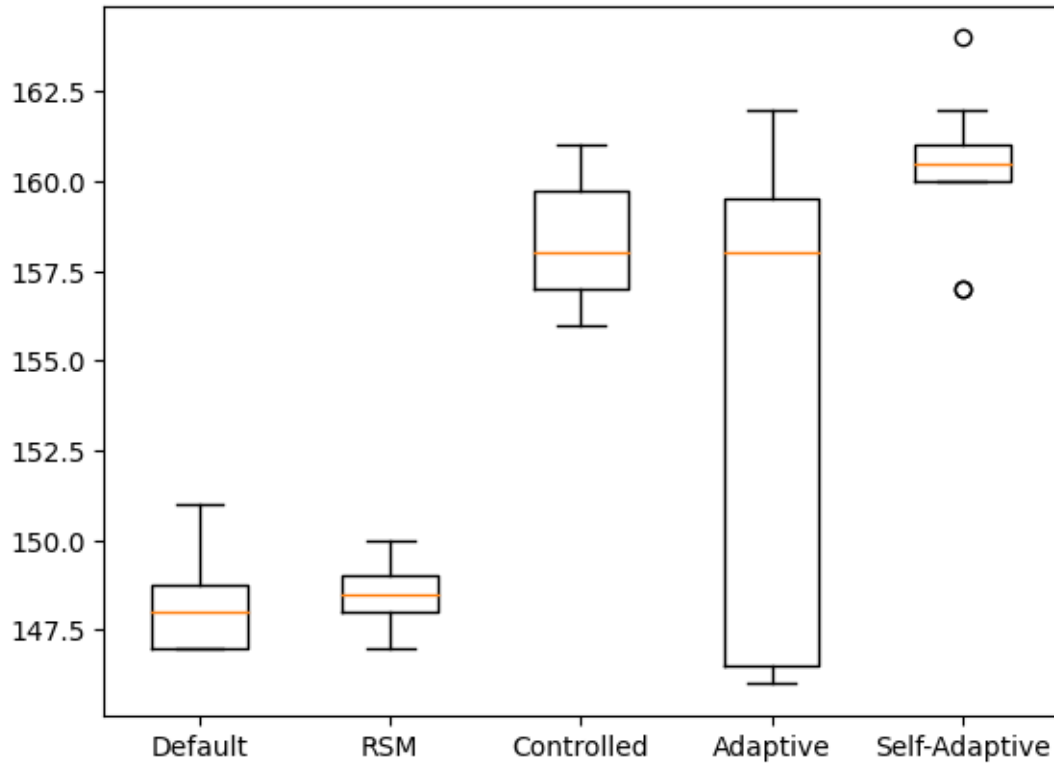
```



```
[77]: parameters = { 0 : ["a", "b", "c", "d", "e", "f", "g"],
                    1 : ["d", "e", "f", "g", "x"],
                    2 : ["d", "e", "f", "g"],
                    3 : ["d", "e", "f"],
                    4 : ["g", "h"]}

num_tests = 30
configuration['P_xover'] = 0.7
configuration['P_mutation'] = 0.07
configuration['population_size'] = 100
configuration['tournament_size'] = 2
results = {
    "Default" : run_times(ga, configuration, 10),
    "RSM" : run_times(ga, rsm_config, 10),
    "Controlled" : run_times(controlled_ga, configuration, 10),
    "Adaptive" : run_times(adaptive_ga, configuration, 10),
    "Self-Adaptive" : run_times(self_adaptive_ga, configuration, 10)
}
fig, ax = plt.subplots()
ax.boxplot(results.values())
ax.set_xticklabels(results.keys())
```

```
[77]: [Text(1, 0, 'Default'),
      Text(2, 0, 'RSM'),
      Text(3, 0, 'Controlled'),
      Text(4, 0, 'Adaptive'),
      Text(5, 0, 'Self-Adaptive')]
```

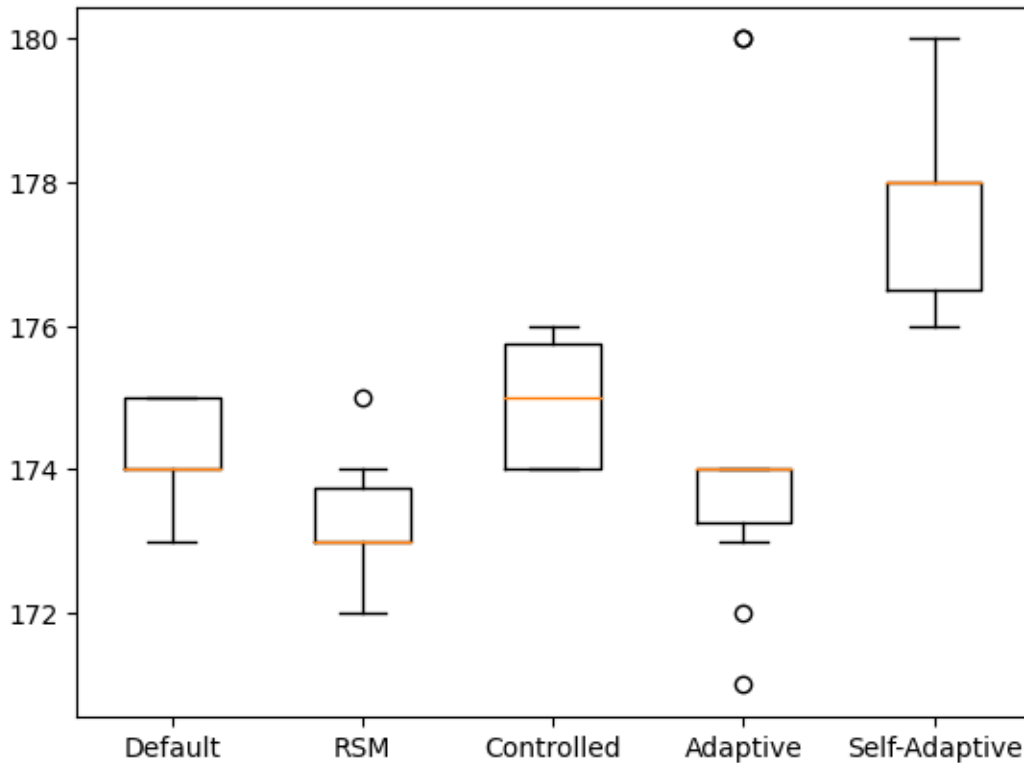



```
[78]: parameters = { 0 : ["0", "1"],
                    1 : ["0", "1"],
                    2 : ["0", "1"],
                    3 : ["0", "1"],
                    4 : ["0", "1"],
                    5 : ["0", "1"],
                    6 : ["0", "1"],
                    7 : ["0", "1"],
                    8 : ["0", "1"],
                    9 : ["0", "1"]}

num_tests = 6 # 10 is required
configuration['P_xover'] = 0.7
configuration['P_mutation'] = 0.07
configuration['population_size'] = 100
configuration['tournament_size'] = 2
results = {
    "Default" : run_times(ga, configuration, 10),
    "RSM" : run_times(ga, rsm_config, 10),
    "Controlled" : run_times(controlled_ga, configuration, 10),
    "Adaptive" : run_times(adaptive_ga, configuration, 10),
    "Self-Adaptive" : run_times(self_adaptive_ga, configuration, 10)
}
```

```
fig, ax = plt.subplots()
ax.boxplot(results.values())
ax.set_xticklabels(results.keys())
```

```
[78]: [Text(1, 0, 'Default'),
      Text(2, 0, 'RSM'),
      Text(3, 0, 'Controlled'),
      Text(4, 0, 'Adaptive'),
      Text(5, 0, 'Self-Adaptive')]
```



1.7 Outlook: Hyper-Heuristics

In this chapter we considered how to tune and adjust the quantitative parameter values for a given search algorithm. However, sometimes different qualitative parameters (e.g., different types of mutation operators) may be beneficial at different times throughout the search, and sometimes even entirely different algorithms may perform better at different points during the search. Indeed there might not be a single optimal heuristic for a given problem type. The idea of *hyper heuristics* is to apply different heuristics to different parts or phases of the solution process. A hyper-heuristic is a search method for selecting or generating heuristics to solve computational search problems, and so it can, essentially, produce entirely new and adaptive meta-heuristic search algorithms.

An example for a hyper-heuristic application in software engineering can be found in the following article:

- Jia, Y., Cohen, M. B., Harman, M., & Petke, J. Learning combinatorial interaction test generation strategies using hyperheuristic search. In Proceedings of the 37th International Conference on Software Engineering-Volume 1 (pp. 540-550). IEEE Press, 2015

In this article, a selective hyper-heuristic is applied to combinatorial interaction testing, i.e. the example problem we used in this notebook. Simulated annealing is the central meta-heuristic, but six different neighbourhood exploration operators are defined, and reinforcement learning is applied in order to select which of them should be applied at each iteration of the simulated annealing algorithm.