

Praktische Optimierung Blatt 05

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```
In [ ]: import numpy as np
        from scipy.stats import multivariate_normal
        import matplotlib.pyplot as plt
        from multiprocessing import Pool
        from tqdm import tqdm
        from functools import partial
```

Aufgabe 1

```
In [ ]: def f(x): return 1.5 * x[0]**2 + x[1]**2 + 21 * np.sin(x[0]) * np.cos(x[1])
```

```
In [ ]: np.random.seed(1)
```

(a) Threshold Accepting

```
In [ ]: def ThresholdAccepting(func : callable, X0 : np.ndarray, T0, gamma, iters
        d = X0.shape[0]
        X = X0
        T = T0
        dist = multivariate_normal(np.zeros_like(X0), np.identity(d))
        for i in range(iters):
            Z = np.array(dist.rvs(1))
            Y = X + Z
            if func(Y) < func(X) + T:
                X = Y
            T = gamma * T
        return {'x' : X, 'fun' : func(X)}
```

```
In [ ]: xs_ta = []
        fns_ta = []
        for i in range(20):
            result = ThresholdAccepting(f, np.array([5,5]), 1, 0.5)
            xs_ta.append(result['x'])
            fns_ta.append(result['fun'])
```

(b) Simulated Annealing

```
In [ ]: def SimulatedAnnealing(func : callable, X0 : np.ndarray, T0, gamma, iters
        d = X0.shape[0]
        X = X0
        T = T0
        dist = multivariate_normal(np.zeros_like(X0), np.identity(d))
        for i in range(iters):
            Z = np.array(dist.rvs(1))
            Y = X + Z
            fun_y = func(Y)
            fun_x = func(X)
            if fun_y < fun_x:
                X = Y
            elif fun_y > fun_x:
                U = np.random.uniform(1)
                if U < np.exp(-(fun_y - fun_x) / T):
                    X = Y
            T = gamma * T
        return {'x' : X, 'fun' : func(X)}
```

```
In [ ]: xs_sa = []
        funs_sa = []
        for i in range(20):
            result = SimulatedAnnealing(f, np.array([5,5]), 1, 0.5)
            xs_sa.append(result['x'])
            funs_sa.append(result['fun'])
```

Ergebnisse

```
In [ ]: funs_ta = np.array(funs_ta)
        funs_sa = np.array(funs_sa)

        funs_ta.sort()
        funs_sa.sort()

        mean_sa = funs_sa.mean()
        median_sa = (funs_sa[9] + funs_sa[10]) / 2
        std_sa = funs_sa.std()

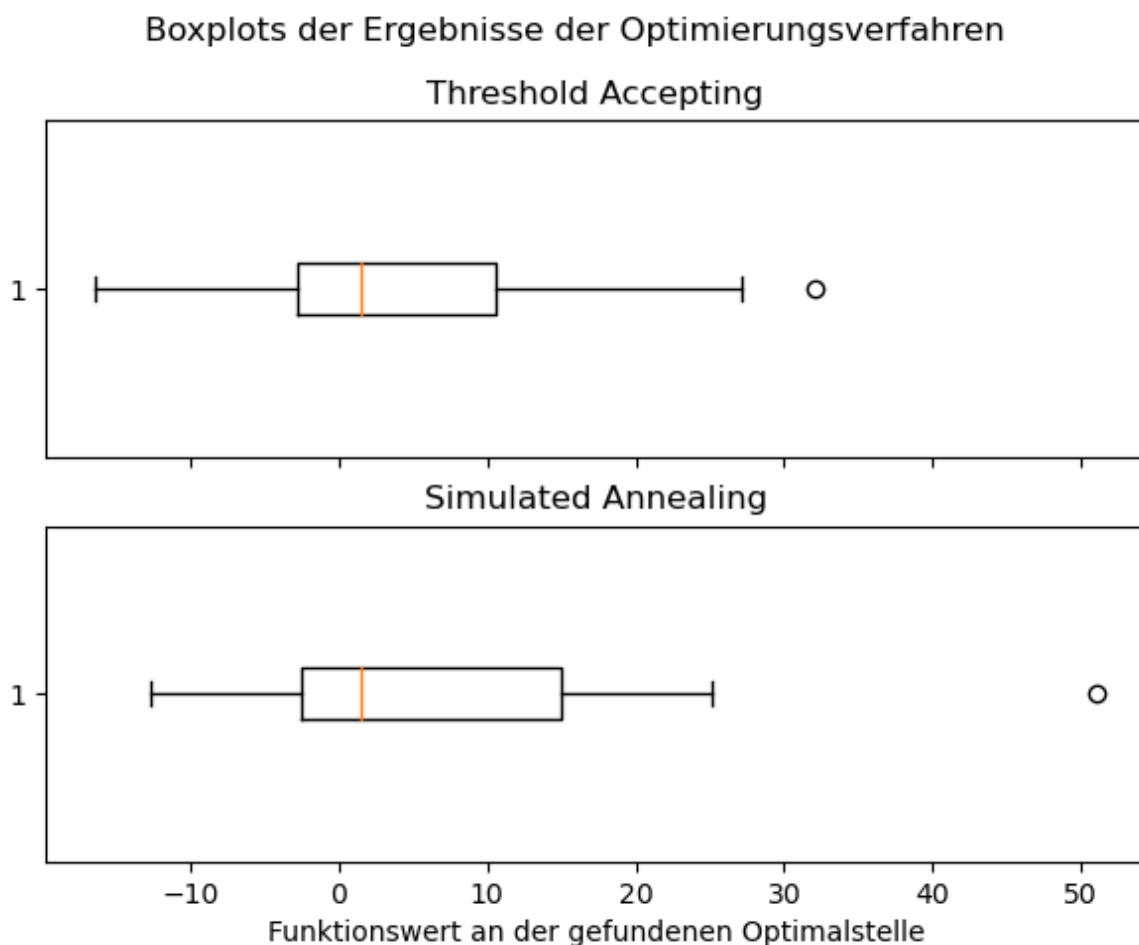
        mean_ta = funs_ta.mean()
        median_ta = (funs_ta[9] + funs_ta[10]) / 2
        std_ta = funs_ta.std()
```

```
In [ ]: print(f'Funktion >-----< Mean >-----< Median >-----< Std >-')
        print(f'-----')
        print(f'Threshold Accept      {mean_ta : .2f}\t      {median_ta : .2f}\t')
        print(f'-----')
        print(f'Simulated Annealing    {mean_sa : .2f}\t      {median_sa : .2f}\t')
```

```
Funktion >-----< Mean >-----< Median >-----< Std >-
-----
Threshold Accept      5.18          1.50          11.62
-----
Simulated Annealing   6.31          1.51          14.01
```

```
In [ ]: fig, ax = plt.subplots(2, sharex=True)
fig.suptitle('Boxplots der Ergebnisse der Optimierungsverfahren')
fig.tight_layout(pad=1.5)
plt.xlabel('Funktionswert an der gefundenen Optimalstelle')
ax[0].set_title('Threshold Accepting')
ax[0].boxplot(funs_ta, vert=False)
ax[1].set_title('Simulated Annealing')
ax[1].boxplot(funs_sa, vert=False)
```

```
Out[ ]: {'whiskers': [<matplotlib.lines.Line2D at 0x7ff128ce3730>,
<matplotlib.lines.Line2D at 0x7ff128ce38b0>],
'caps': [<matplotlib.lines.Line2D at 0x7ff128ce3b50>,
<matplotlib.lines.Line2D at 0x7ff128ce3df0>],
'boxes': [<matplotlib.lines.Line2D at 0x7ff128ce3490>],
'medians': [<matplotlib.lines.Line2D at 0x7ff128c6f0d0>],
'fliers': [<matplotlib.lines.Line2D at 0x7ff128c6f370>],
'means': []}
```



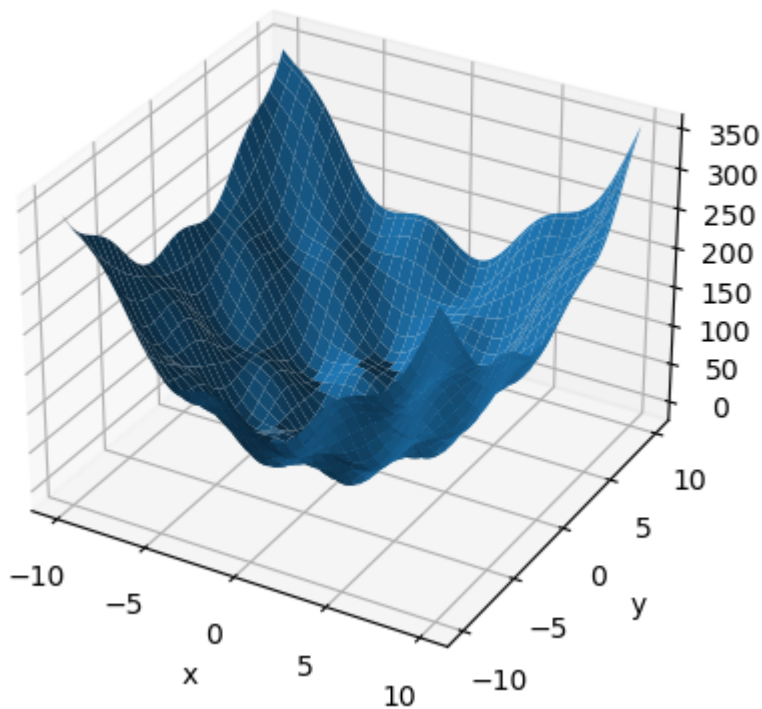
```
In [ ]: ax = plt.figure().add_subplot(projection='3d')

x = np.linspace(-10, 10, 101)
y = np.linspace(-10, 10, 101)
X, Y = np.meshgrid(x, y)
Z = 1.5 * X**2 + Y**2 + 21 * np.sin(X) * np.cos(Y) + 0.5 * (np.abs(X))**2

plt.title('3D-Plot der Funktion f')
plt.xlabel('x')
plt.ylabel('y')
ax.set_axisbelow(True)
ax.plot_surface(X, Y, Z)
```

```
Out[ ]: <mpl_toolkits.mplot3d.art3d.Poly3DCollection at 0x7ff128c24580>
```

3D-Plot der Funktion f



Interpretation

Die Threshold Accepting funktion scheint hier etwas bessere Ergebnisse zu liefern. Die Werte liegen im Mittel etwas tiefer und weisen eine geringere Streuung auf. Außerdem ist im Boxplot von Simulated Annealing zu sehen, dass hier 3 Outlierwerte vorliegen während bei TA nur ein Wert deutlich größer ist.

Aufgabe 2

Hilfsfunktion für übersichtlichen Code

```
In [ ]: def checkValid(solution : np.ndarray):  
    s1 = solution[0] + solution[1] + solution[2]  
    s2 = solution[3] + solution[4] + solution[5]  
    s3 = solution[6] + solution[7] + solution[8]  
  
    if not s1 == s2 == s3:  
        return False  
  
    s4 = solution[0] + solution[3] + solution[6]  
    s5 = solution[1] + solution[4] + solution[7]  
    s6 = solution[2] + solution[5] + solution[8]  
  
    if not s4 == s5 == s6:  
        return False  
  
    s7 = solution[0] + solution[4] + solution[8]  
    s8 = solution[6] + solution[4] + solution[2]  
  
    return s7 == s8
```

Rekombinationsfunktionen

```

In [ ]: def order_based_crossover(p1 : np.ndarray, p2 : np.ndarray):
    l = p1.shape[0]
    child = np.zeros_like(p1)
    ind = np.random.choice(np.arange(l), 2, replace=False)
    if ind[1] < ind[0]:
        tmp = ind[1]
        ind[1] = ind[0]
        ind[0] = tmp

    child[ind[0]:ind[1]+1] = p1[ind[0]:ind[1]+1]

    current_index = ind[1] + 1

    for i in range(ind[1]+1, ind[1]+1+l):
        if p2[i % l] not in child:
            child[current_index % l] = p2[i % l]
            current_index += 1

    return child

def partially_mapped_crossover(p1 : np.ndarray, p2 : np.ndarray):
    l = p1.shape[0]
    child = p2.copy()
    ind = np.random.choice(np.arange(l), 2, replace=False)
    if ind[1] < ind[0]:
        tmp = ind[1]
        ind[1] = ind[0]
        ind[0] = tmp

    for i in range(ind[0], ind[1]+1):
        j = np.where(child == p1[i])[0][0]
        tmp = child[i]
        child[i] = child[j]
        child[j] = tmp

    return child

def no_recombination(p1 : np.ndarray, p2 : np.ndarray):
    if np.random.choice(np.arange(2), 1)[0] == 0:
        return p1.copy()
    return p2.copy()

```

Mutationsfunktionen

```
In [ ]: def two_swap(p):
        child = p.copy()
        ind = np.random.choice(np.arange(p.shape[0]), 2, replace=False)
        tmp = child[ind[0]]
        child[ind[0]] = child[ind[1]]
        child[ind[1]] = tmp
        return child

def one_translocation(p):
    child = p.copy()
    ind = np.random.choice(np.arange(p.shape[0]), 2, replace=False)
    if ind[1] < ind[0]:
        tmp = ind[0]
        ind[0] = ind[1]
        ind[1] = tmp
    tmp = child[ind[0]]
    for i in range(ind[0], ind[1]):
        child[i] = child[i+1]
    child[ind[1]] = tmp
    return child

def global_mutation(p : np.ndarray, k):
    child = p.copy()
    for i in range(k):
        child = two_swap(child)
    return child

def global_mutation_2(p : np.ndarray):
    return global_mutation(p, k=2)

def global_mutation_4(p : np.ndarray):
    return global_mutation(p, k=4)
```

Zielfunktionen

```

In [ ]: def first(individual : np.ndarray):
        s1 = individual[0] + individual[1] + individual[2]
        s2 = individual[3] + individual[4] + individual[5]
        s3 = individual[6] + individual[7] + individual[8]
        s4 = individual[0] + individual[3] + individual[6]
        s5 = individual[1] + individual[4] + individual[7]
        s6 = individual[2] + individual[5] + individual[8]
        s7 = individual[0] + individual[4] + individual[8]
        s8 = individual[6] + individual[4] + individual[2]
        S = np.array([s1, s2, s3, s4, s5, s6, s7, s8])

        return S.max() - S.min()

def second(individual : np.ndarray):
    s1 = individual[0] + individual[1] + individual[2]
    s2 = individual[3] + individual[4] + individual[5]
    s3 = individual[6] + individual[7] + individual[8]
    s4 = individual[0] + individual[3] + individual[6]
    s5 = individual[1] + individual[4] + individual[7]
    s6 = individual[2] + individual[5] + individual[8]
    s7 = individual[0] + individual[4] + individual[8]
    s8 = individual[6] + individual[4] + individual[2]
    S = np.array([s1, s2, s3, s4, s5, s6, s7, s8])

    sum = 0
    for i in range(7):
        for j in range(i+1, 8):
            sum += np.abs(S[i] - S[j])
    return sum

def third(individual : np.ndarray):
    s1 = individual[0] + individual[1] + individual[2]
    s2 = individual[3] + individual[4] + individual[5]
    s3 = individual[6] + individual[7] + individual[8]
    s4 = individual[0] + individual[3] + individual[6]
    s5 = individual[1] + individual[4] + individual[7]
    s6 = individual[2] + individual[5] + individual[8]
    s7 = individual[0] + individual[4] + individual[8]
    s8 = individual[6] + individual[4] + individual[2]
    S = np.array([s1, s2, s3, s4, s5, s6, s7, s8])

    sum = 0
    for i in range(1, 8):
        sum += np.abs(S[i] - S[0])
    return sum

```

EA


```

In [ ]: def magic_square_ea(fitness : callable, mu : int, lam : int, recombine :
        alphabet = np.arange(9) + 1
        parents = np.zeros((mu, 9))

        # Haltekriterium
        foundValidSolution = False

        # initiale Eltern generierung
        for i in range(mu):
            parents[i] = np.random.permutation(alphabet)

        fit_vals_parents = np.apply_along_axis(fitness, axis=1, arr=parents)
        count = parents.shape[0]

        # Hauptschleife
        while not foundValidSolution:

            children = np.zeros((lam, 9))
            # Rekombination der Eltern mit anschließender Mutation des Kindes
            for i in range(lam):
                # Auswahl von 2 unterschiedlichen Eltern
                indices = np.random.choice(np.arange(mu), 2, replace=False)
                p1 = parents[indices[0]]
                p2 = parents[indices[1]]
                # Mutation und Rekombination
                children[i] = mutate(recombine(p1, p2))

            population = np.concatenate([parents, children])
            fit_vals_children = np.apply_along_axis(fitness, axis=1, arr=children)

            fit_vals = np.concatenate([fit_vals_parents, fit_vals_children])
            count += children.shape[0]
            # wähle die mu besten Individuen
            ind = np.argpartition(fit_vals, mu)[:mu]
            parents = population[ind]
            fit_vals_parents = fit_vals[ind]

            # Überprüfe Haltekriterium
            for individual in parents:
                if checkValid(individual):
                    return {'Quadrat' : individual, 'Anzahl' : count, 'mu' :
                            'lam' : lam, 'recombine' : recombine.__name__, 'm
                            , 'fit' : fitness.__name__}

```

Testen der Parameterkombinationen

```

In [ ]: mus = [5, 10, 20]
        lambdas = [1, 5, 10]
        #mutations = [two_swap, one_translocation, global_mutation_2, global_muta
        mutations = [global_mutation_4]
        recombinations = [order_based_crossover, partially_mapped_crossover, no_r
        fits = [first, second, third]

```

```

In [ ]: fit, mu, lam, rec, mut = np.meshgrid(fits, mus, lambdas, recombinations,

```

```
In [ ]: mu = mu.flatten()
        lam = lam.flatten()
        mut = mut.flatten()
        rec = rec.flatten()
        fit = fit.flatten()
```

gloable Mutation mit $k = 4$

```
In [ ]: results_g4 = []
        with Pool(81) as p:
            results_g4 = p.starmap(magic_square_ea, zip(fit, mu, lam, rec, mut))
```

```
In [ ]: vals_first = []
        vals_second = []
        vals_third = []

        results_first = []
        results_second = []
        results_third = []

        for res in results_g4:
            if res['fit'] == first.__name__:
                vals_first.append(res['Anzahl'])
                results_first.append(res)
            elif res['fit'] == second.__name__:
                vals_second.append(res['Anzahl'])
                results_second.append(res)
            else:
                vals_third.append(res['Anzahl'])
                results_third.append(res)
```

```
In [ ]: vals_first = np.array(vals_first)
        vals_second = np.array(vals_second)
        vals_third = np.array(vals_third)

        results_first = np.array(results_first)
        results_second = np.array(results_second)
        results_third = np.array(results_third)
```

```
In [ ]: max_5_first = np.argpartition(vals_first, -5)[-5:]
        max_5_second = np.argpartition(vals_second, -5)[-5:]
        max_5_third = np.argpartition(vals_third, -5)[-5:]

        min_5_first = np.argpartition(vals_first, 5)[:5]
        min_5_second = np.argpartition(vals_second, 5)[:5]
        min_5_third = np.argpartition(vals_third, 5)[:5]
```

```
In [ ]: best_5_first = results_first[min_5_first]
        best_5_second = results_second[min_5_second]
        best_5_third = results_third[min_5_third]

        worst_5_first = results_first[max_5_first]
        worst_5_second = results_second[max_5_second]
        worst_5_third = results_third[max_5_third]
```

```
In [ ]: def print_result(res : dict):
        print(f'Mu: {res["mu"]}\tlambda: {res["lam"]}\tcombine: {res["recombi
```

```
In [ ]: print('Zielfunktion 1:\nTop-5:\n')
        for res in best_5_first:
            print_result(res)
        print('\nBottom-5:\n')
        for res in worst_5_first:
            print_result(res)
```

Zielfunktion 1:

Top-5:

```
Mu: 5   lambda: 5       combine: partially_mapped_crossover   mutate: gl
obal_mutation_4 #Auswertungen: 570
Mu: 10  lambda: 10      combine: no_recombination           mutate: global_mut
ation_4 #Auswertungen: 1480
Mu: 10  lambda: 1       combine: partially_mapped_crossover   mutate: gl
obal_mutation_4 #Auswertungen: 1532
Mu: 10  lambda: 1       combine: no_recombination           mutate: global_mut
ation_4 #Auswertungen: 2286
Mu: 10  lambda: 1       combine: order_based_crossover       mutate: global_mut
ation_4 #Auswertungen: 2591
```

Bottom-5:

```
Mu: 5   lambda: 1       combine: order_based_crossover       mutate: global_mut
ation_4 #Auswertungen: 16903
Mu: 5   lambda: 5       combine: order_based_crossover       mutate: global_mut
ation_4 #Auswertungen: 22510
Mu: 20  lambda: 5       combine: order_based_crossover       mutate: global_mut
ation_4 #Auswertungen: 21220
Mu: 5   lambda: 10      combine: no_recombination           mutate: global_mut
ation_4 #Auswertungen: 21725
Mu: 10  lambda: 5       combine: order_based_crossover       mutate: global_mut
ation_4 #Auswertungen: 22830
```

```
In [ ]: print('Zielfunktion 2:\nTop-5:\n')
        for res in best_5_second:
            print_result(res)
        print('\nBottom-5:\n')
        for res in worst_5_second:
            print_result(res)
```

Zielfunktion 2:

Top-5:

```
Mu: 20  lambda: 5      combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 1010
Mu: 10  lambda: 1      combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 1776
Mu: 5   lambda: 1      combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 1175
Mu: 10  lambda: 10     combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 1950
Mu: 20  lambda: 5      combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 2170
```

Bottom-5:

```
Mu: 20  lambda: 10     combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 12000
Mu: 10  lambda: 1      combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 17480
Mu: 20  lambda: 10     combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 21120
Mu: 5   lambda: 10     combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 12285
Mu: 5   lambda: 10     combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 18695
```

```
In [ ]: print('Zielfunktion 3:\nTop-5:\n')
        for res in best_5_third:
            print_result(res)
        print('\nBottom-5:\n')
        for res in worst_5_third:
            print_result(res)
```

Zielfunktion 3:

Top-5:

```
Mu: 5   lambda: 5      combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 3030
Mu: 10  lambda: 10     combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 2410
Mu: 20  lambda: 1      combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 3216
Mu: 10  lambda: 5      combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 5055
Mu: 5   lambda: 1      combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 5486
```

Bottom-5:

```
Mu: 20  lambda: 1      combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 39025
Mu: 10  lambda: 1      combine: no_recombination      mutate: global_mutation_4 #Auswertungen: 40474
Mu: 5   lambda: 10     combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 50215
Mu: 20  lambda: 10     combine: partially_mapped_crossover  mutate: global_mutation_4 #Auswertungen: 54220
Mu: 10  lambda: 10     combine: order_based_crossover  mutate: global_mutation_4 #Auswertungen: 68550
```

Ergebnisse

Ich konnte leider nicht alle Parameterkombinationen testen, da der EA für manche Kombinationen sehr lange in lokalen Minima stecken bleibt (ich habe die Ausführung nach ~3h abgebrochen). Von den Ergebnissen die ich beobachten konnte scheint es mir so, als wären Parameter optimal, welche eine höhere Anzahl an Nachkommen haben und die für eine größere Abweichung von den Eltern sorgen. Dadurch bleibt der Algorithmus nicht so sehr in lokalen Minima stecken. Hat man pro Iteration nur 1 Nachkommen, welcher sich nur an einer Stelle von einem der Elternteile unterscheidet ist es sehr schwierig aus lokalen Minima wieder heraus zu kommen. Daher vermute ich, dass die Mutationsfunktion **global_k4** die beste ist.

Für **global_k4** scheint die Zielfunktion 1 die beste zu sein, gefolgt von 2 und dann 3.

Die Parameterkombination:

$\mu = 5$, $\lambda = 5$, Rekombination = partially-mapped crossover, Zielfunktion = 1

war die beste bzgl. der Mutationsfunktion **global_k4**.