
Studienprojekt

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Feb 16, 2021

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GRID POINTS

Here we show what kind of grids are used for the different quadratures.

Further more, there are two options, how many points are used for the probabilistic quadratures. (one or three points)

These options are compared afterwards.

```
[1]: import matplotlib.pyplot as plt
import os
os.chdir("../")
from Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point import *
```

1.1 One point for $q = 1$

```
[2]: def grid_points(quadrature: str, q: int, a=0, b=1):
    list_of_points = []
    dim = 2

    # Because the points used for the deterministic quadratures are not changed, it_
    ↪ is faster to call them
    # at the beginning before starting the Smolyak alg.
    if "Carlo" in quadrature:
        q = q + 1

    # For every one dimensional degree of approx. a weights and points vector
    for i in range(1, (q - dim + 2)):

        # One option for Newton-Cotes and Trapezoidal quadrature, because
        if quadrature == "Newton-Cotes" or quadrature == "Trapezoidal":
            points, weights = one_dim_trapezoidal(i, a, b)

        if quadrature == "Monte Carlo (nested)" and i > 1:
            points, weights = monte_carlo_quad(i-1, a, b)
            points = np.concatenate([list_of_points[-1], points])

        if quadrature == "Monte Carlo (non-nested)" or (quadrature == "Monte Carlo_
    ↪ (nested)" and i == 1) :
            points, weights = monte_carlo_quad(i, a, b)

        if quadrature == "Quasi-Monte Carlo":
            points, weights = qmc_quad(i, a, b)
```

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```

        list_of_points.append(points)

        # Now we get the combinations of one-dimensional degrees of approx. in  $Q(q,d)$ 
        rng = list(range(q)) * dim
        rng = [x + 1 for x in rng]

        possible_combinations = list(set(i for i in itt.permutations(rng, dim) if (q -
        ↪ dim) < sum(i) < q + 1))
        number_approx = len(possible_combinations)
        meshgrid_list_x = []
        meshgrid_list_y = []
        for i in range(number_approx):

            # gridpoints and the weight vectors are put into separate lists to make it
            ↪ easier to
            # make a meshgrid
            current_tuple = possible_combinations[i]
            current_points = []

            for j in range(len(current_tuple)):
                current_points.append(list_of_points[current_tuple[j] - 1])

            # meshgrid yields a np.array with all coordinates needed for alg
            # coordinate and respective weight could be found in the same places in the
            ↪ array
            meshgrid_points = np.array(np.meshgrid(*current_points))
            meshgrid_list_x.append(meshgrid_points[0])
            meshgrid_list_y.append(meshgrid_points[1])

        return [meshgrid_list_x, meshgrid_list_y]

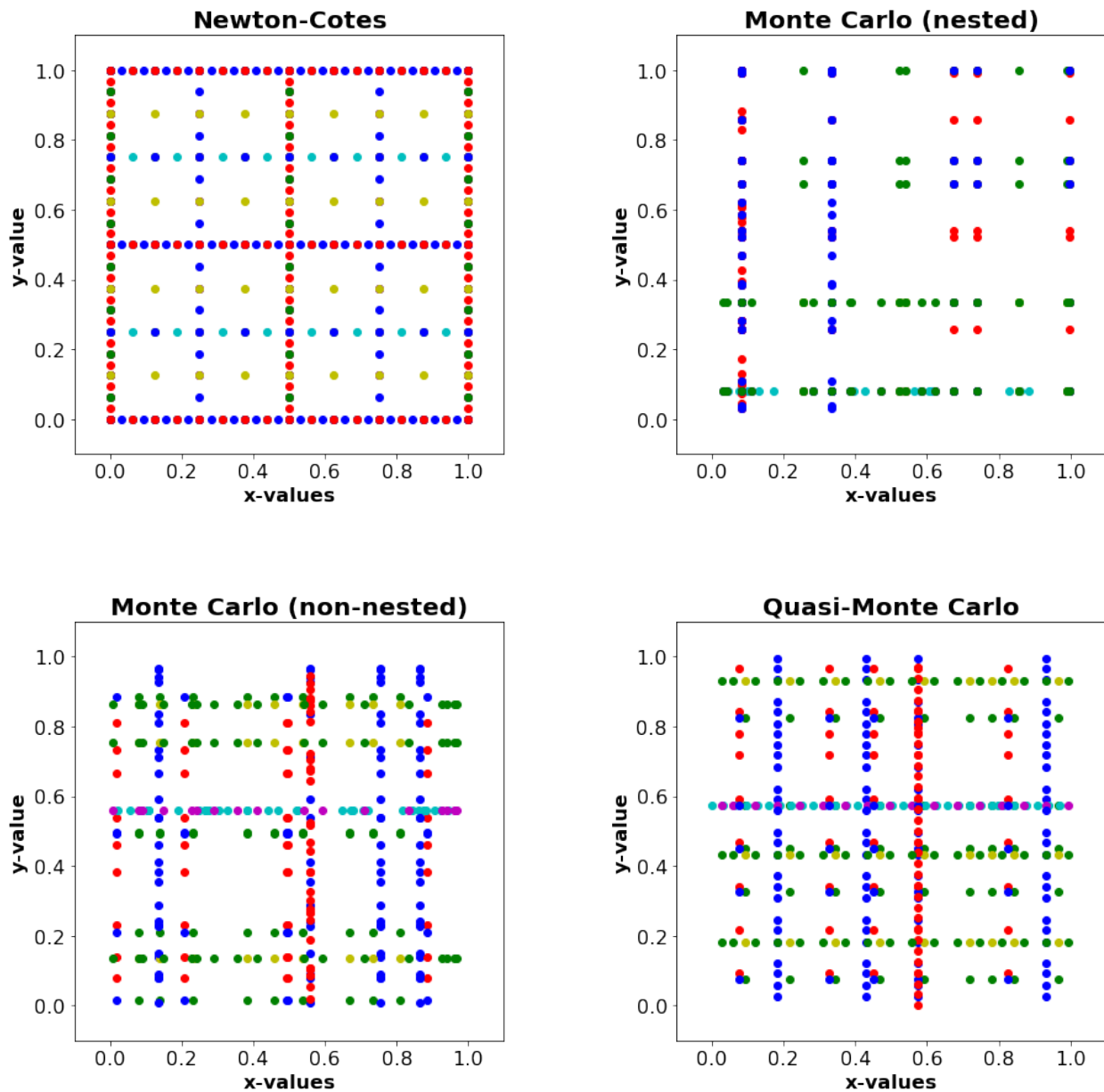
def make_plot_of_grids():
    option_list = ["Newton-Cotes",
                  "Monte Carlo (nested)",
                  "Monte Carlo (non-nested)",
                  "Quasi-Monte Carlo"]
    fig = plt.figure(figsize=(15,15))
    plt.subplots_adjust(hspace=0.4, wspace=0.4)

    for i in range(len(option_list)):
        grid_list_x, grid_list_y = grid_points(option_list[i], 6)
        plt.subplot(2,2,i+1)
        modified_scatter_plot(grid_list_x, grid_list_y, title=option_list[i], input_
        ↪ list=True)
        plt.xticks(fontsize=16)
        plt.yticks(fontsize=16)

    fig.suptitle("Gridpoints used for Smolyak-algorithm (one point)", fontsize=24,
    ↪ fontweight="bold")
    plt.show()

make_plot_of_grids()

```

Gridpoints used for Smolyak-algorithm (one point)

1.1.1 Here the grid points used for the approximation of the integral of a 2 dimensional function with the level of approximation $q = 6$ is shown, if one point is used for the one is used for probabilistic quadratures and $q = 1$. Apart from the upper left grid, the other grids change a little every time they are generated.

1.2 Three points for $q = 1$

```
[3]: from Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_three_points import *
```

```
[4]: def grid_points(quadrature: str, q: int, a=0,b=1):
    list_of_points = []
    dim = 2
    # Now for the probabilistic quadratures, only one point is used for the first_
    ↪one-dimensional
    # degree of approximation. For this we add one to q, because otherwise the number_
    ↪of points for
    # the probabilistic quadratures would be much smaller.

    # Now we apply the quadrature chosen by user
    # !! If quadratures is changed or extended in interface, please options here !!

    # Because the points used for the deterministic quadratures are not changed, it_
    ↪is faster to call them
    # at the beginning before starting the Smolyak alg.

    # For every one dimensional degree of approx. a weights and points vector
    for i in range(1, (q - dim + 2)):

        # One option for Newton-Cotes and Trapezoidal quadrature, because
        if quadrature == "Newton-Cotes" or quadrature == "Trapezoidal":
            points, weights = one_dim_trapezoidal(i, a, b)

        if quadrature == "Monte Carlo (nested)" and i > 1:
            points, weights = monte_carlo_quad(i-1, a, b)
            points = np.concatenate([list_of_points[-1], points])

        if quadrature == "Monte Carlo (non-nested)" or (quadrature == "Monte Carlo_
        ↪(nested)" and i == 1) :
            points, weights = monte_carlo_quad(i, a, b)

        if quadrature == "Quasi-Monte Carlo":
            points, weights = qmc_quad(i, a, b)

        list_of_points.append(points)

    # Now we get the combinations of one-dimensional degrees of approx. in  $Q(q,d)$ 
    rng = list(range(q)) * dim
    rng = [x + 1 for x in rng]

    possible_combinations = list(set(i for i in itt.permutations(rng, dim) if (q -
    ↪dim) < sum(i) < q + 1))
```

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```

number_approx = len(possible_combinations)
meshgrid_list_x = []
meshgrid_list_y = []
for i in range(number_approx):

    # the next two steps could be combined in one line.
    # what is done is, that first the tuple at position i is taken from the list_
    of the set I.
    # Then the regarding vectors are taken from the list with the weight and_
    points vectors and
    # are combined to a d dimensional point set

    # gridpoints and the weight vectors are put into separate lists to make it_
    easier to
    # make a meshgrid
    current_tuple = possible_combinations[i]
    current_points = []

    for j in range(len(current_tuple)):
        current_points.append(list_of_points[current_tuple[j] - 1])

    # meshgrid yields a np.array with all coordinates needed for alg
    # coordinate and respective weight could be found in the same places in the_
    array
    meshgrid_points = np.array(np.meshgrid(*current_points))
    meshgrid_list_x.append(meshgrid_points[0])
    meshgrid_list_y.append(meshgrid_points[1])

    return [meshgrid_list_x, meshgrid_list_y]

def make_plot_of_grids():
    option_list = ["Newton-Cotes",
                  "Monte Carlo (nested)",
                  "Monte Carlo (non-nested)",
                  "Quasi-Monte Carlo"]
    fig = plt.figure(figsize=(15,15))
    plt.subplots_adjust(hspace=0.4, wspace=0.4)

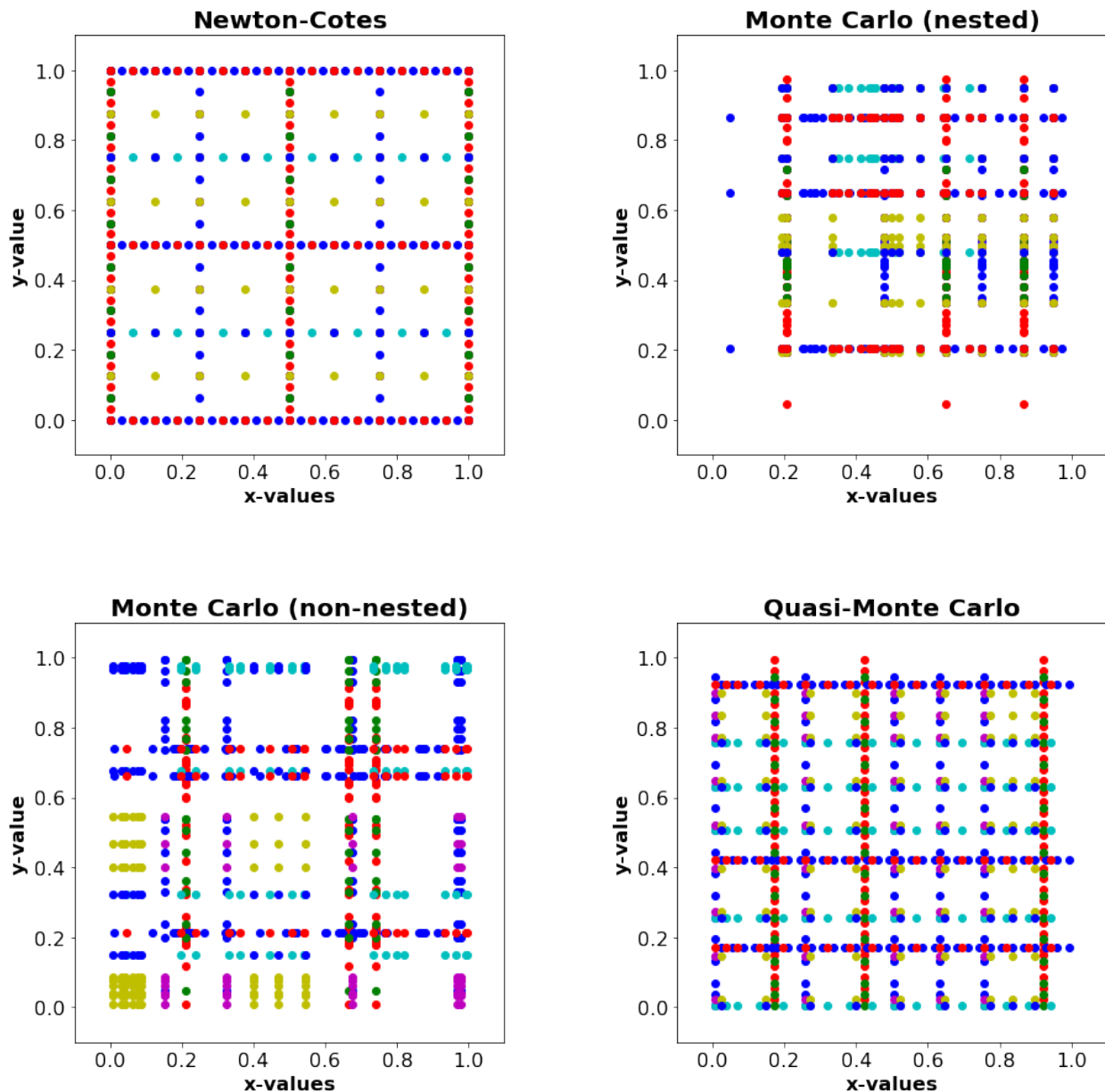
    for i in range(len(option_list)):
        grid_list_x, grid_list_y = grid_points(option_list[i], 6)
        plt.subplot(2,2,i+1)
        modified_scatter_plot(grid_list_x,grid_list_y,title=option_list[i], input_
        list=True)
        plt.xticks(fontsize=16)
        plt.yticks(fontsize=16)

    fig.suptitle("Gridpoints used for Smolyak-algorithm (one point)", fontsize=24,
    fontweight="bold")
    plt.show()

make_plot_of_grids()

```

Gridpoints used for Smolyak-algorithm (one point)



Here the grid points used for the approximation of the integral of a 2 dimensional function with the level of approximation $q = 6$ is shown, if one point is used for the one is used for probabilistic quadratures and $q = 1$. Apart from the upper left grid, the other grids change a little every time they are generated.

[]:

PROOF OF PRINCIPLE

2.1 One point for $q = 1$

We want to show that the approximations are working. For this we use an easy differential function. The result of the integral is 1. We only take low degrees of evaluation, because we only want to give a proof of principle. Quantitative evaluations comparisons are made afterwards for on a statistical basis.

```
[1]: import matplotlib.pyplot as plt
import time as time
import pandas as bearcats
import matplotlib.colors as colors

import os
os.chdir("../")
from Methodes_Studienproject.Studienprojekt_Smolyak_qmc_one_point import *
```

```
[2]: function_string = "pi**2/4 *sin(pi*x)*sin(pi*y) "
variables_string = "(x,y) "
option_list = ["Newton-Cotes",
               "Trapezoidal",
               "Monte Carlo (nested)",
               "Monte Carlo (non-nested)",
               "Quasi-Monte Carlo"]
result_list = []

for i in range(len(option_list)):
    results_quad = []
    for j in range(2,7):
        results_quad.append(controller_smolyak(function_string, variables_string,
        ↪option_list[i], j))
    result_list.append(results_quad)

for i in range(5):
    result = [result_list[i][j][0] for j in range(5) ]
    error = [result_list[i][j][1] for j in range(5) ]
    cost = [result_list[i][j][2] for j in range(5) ]
    fig = plt.figure(figsize=(15,7))
    plt.subplots_adjust(hspace=0.4, wspace=0.4)

    plt.subplot(2,2,1)
    plt.plot(list(range(2,7)),result)
    plt.hlines(1,0,10, linestyle= "dotted")
```

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```

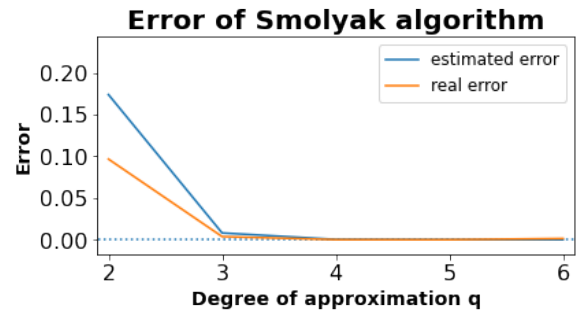
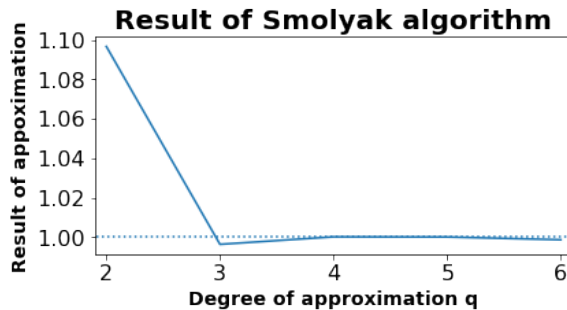
plt.xlim(1.9,6.1)
plt.xlabel("Degree of approximation q", fontsize = 14,fontweight = "bold")
plt.ylabel("Result of appoximation",fontsize = 14,fontweight = "bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.xticks(range(2,7))
plt.title("Result of Smolyak algorithm", fontsize = 20, fontweight = "bold")

plt.subplot(2,2,2)
plt.plot(list(range(2,7)),error, label= "estimated error")
plt.plot(list(range(2,7)),[abs(x-1) for x in result], label= "real error")
plt.hlines(0,0,10, linestyle= "dotted")
plt.xlim(1.9,6.1)

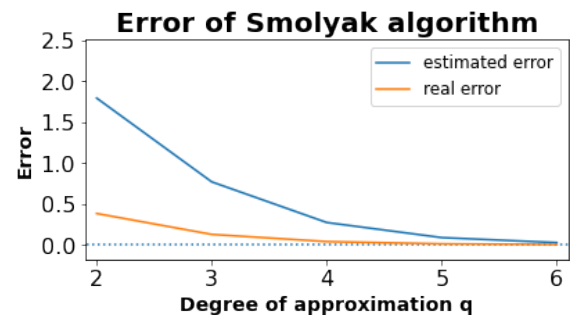
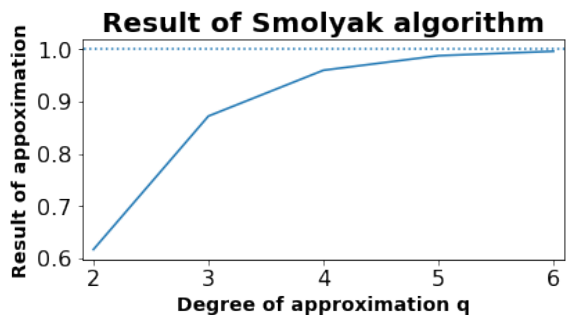
plt.xlabel("Degree of approximation q", fontsize=14,fontweight = "bold")
plt.ylabel("Error", fontsize=14,fontweight = "bold")
plt.title("Error of Smolyak algorithm",fontsize = 20, fontweight="bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.xticks(range(2,7))
plt.ylim(-max(error)*0.1,max(error)*1.4)
plt.legend(fontsize=12, loc=1)
plt.suptitle(option_list[i],fontsize=24, fontweight="bold")
plt.show()

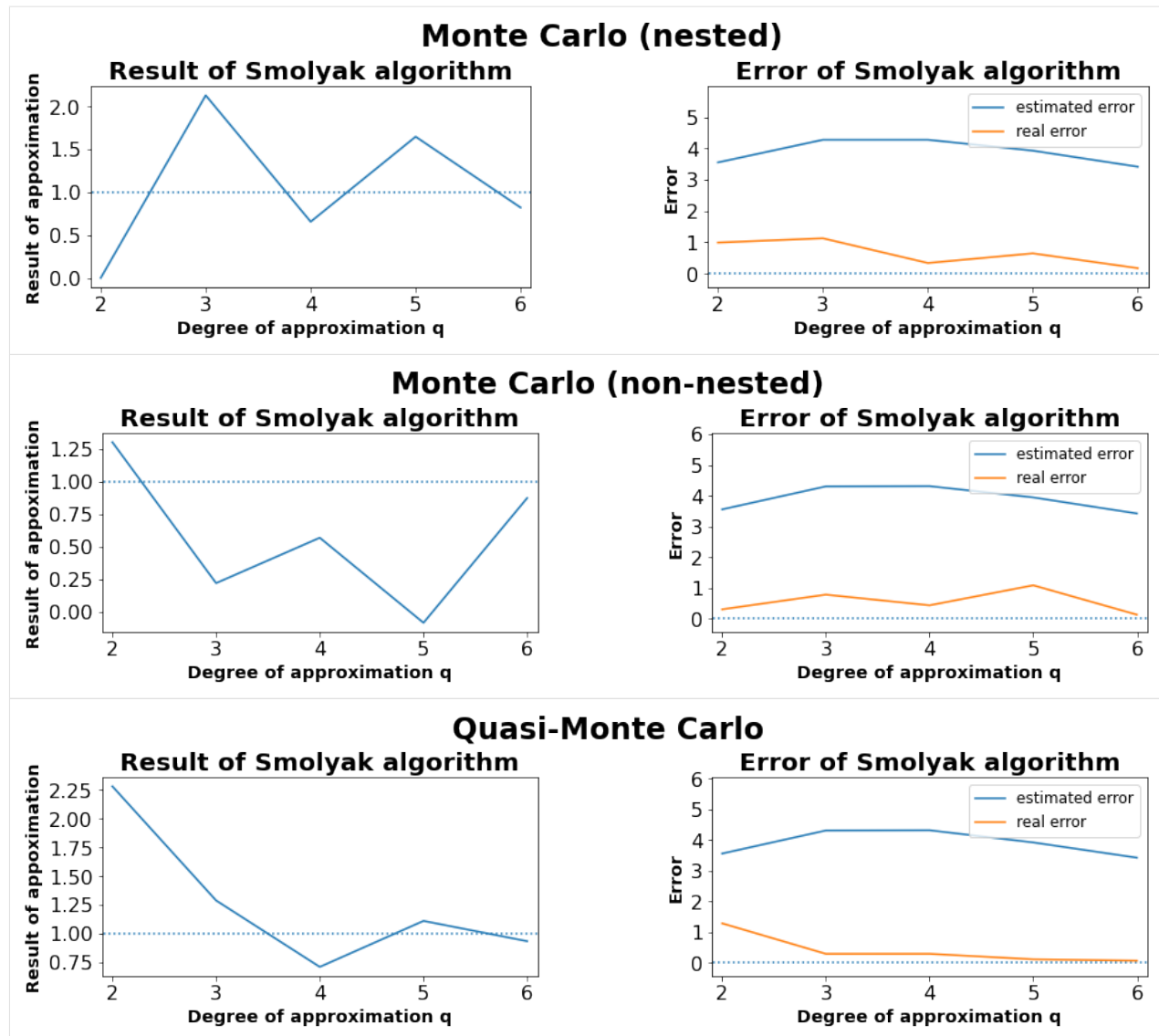
```

Newton-Cotes



Trapezoidal





The result of the approximation and the error each should converge towards the dotted line. The estimated error for the probabilistic quadratures is not really significant, because the central limit theorem could not be applied here. The probabilistic quadratures for $q_i = 1$ only uses one point. For this reason these quadratures are less exact.

2.2 Three points for $q = 1$.

```
[11]: import matplotlib.pyplot as plt
import time as time
from Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_three_points import *
```

```
[12]: function_string = "pi**2/4 *sin(pi*x)*sin(pi*y)"
variables_string = "(x,y)"
option_list = ["Newton-Cotes",
              "Trapezoidal",
```

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```

        "Monte Carlo (nested)",
        "Monte Carlo (non-nested)",
        "Quasi-Monte Carlo"]
result_list = []

for i in range(len(option_list)):
    results_quad = []
    for j in range(2,7):
        results_quad.append(controller_smolyak(function_string, variables_string,
↪option_list[i], j))
    result_list.append(results_quad)

for i in range(5):
    result = [result_list[i][j][0] for j in range(5) ]
    error   = [result_list[i][j][1] for j in range(5) ]
    cost    = [result_list[i][j][2] for j in range(5) ]
    fig = plt.figure(figsize=(15,7))
    plt.subplots_adjust(hspace=0.4, wspace=0.4)

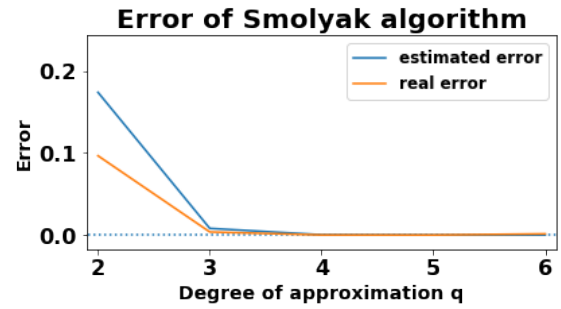
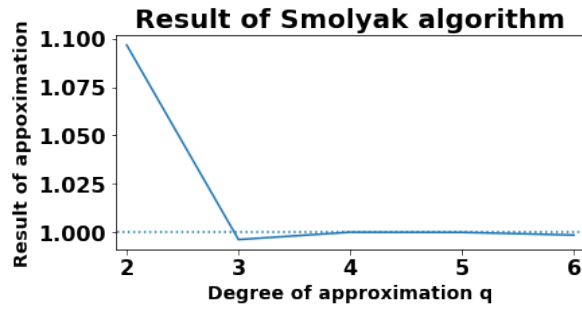
    plt.subplot(2,2,1)
    plt.plot(list(range(2,7)),result)
    plt.hlines(1,0,10, linestyle= "dotted")
    plt.xlim(1.9,6.1)
    plt.xlabel("Degree of approximation q", fontsize = 14,fontweight = "bold")
    plt.ylabel("Result of appoximation",fontsize = 14,fontweight = "bold")
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)
    plt.xticks(range(2,7))
    plt.title("Result of Smolyak algorithm", fontsize = 20, fontweight = "bold")

    plt.subplot(2,2,2)
    plt.plot(list(range(2,7)),error, label= "estimated error")
    plt.plot(list(range(2,7)),[abs(x-1) for x in result], label= "real error")
    plt.hlines(0,0,10, linestyle= "dotted")
    plt.xlim(1.9,6.1)

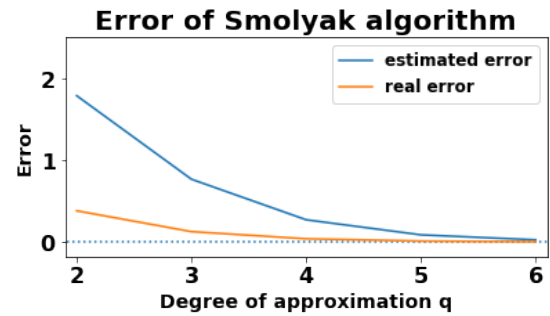
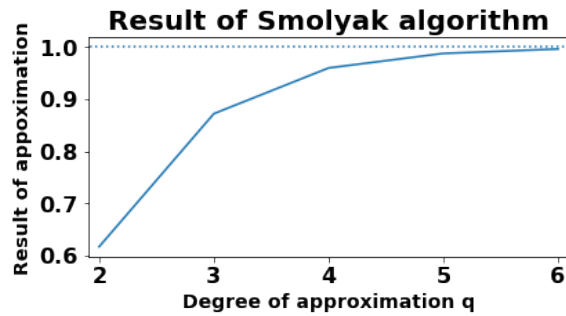
    plt.xlabel("Degree of approximation q", fontsize=14,fontweight = "bold")
    plt.ylabel("Error", fontsize=14,fontweight = "bold")
    plt.title("Error of Smolyak algorithm",fontsize = 20, fontweight="bold")
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)
    plt.xticks(range(2,7))
    plt.ylim(-max(error)*0.1,max(error)*1.4)
    plt.legend(fontsize=12, loc=1)
    plt.suptitle(option_list[i],fontsize=24, fontweight="bold")
    plt.show()

```

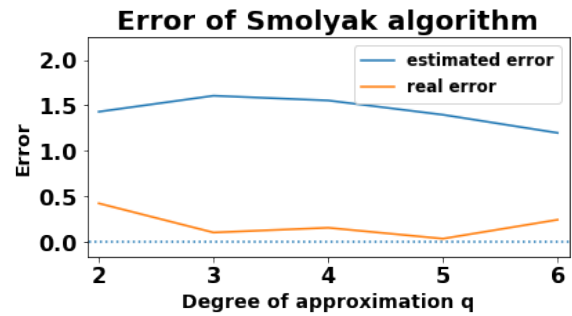
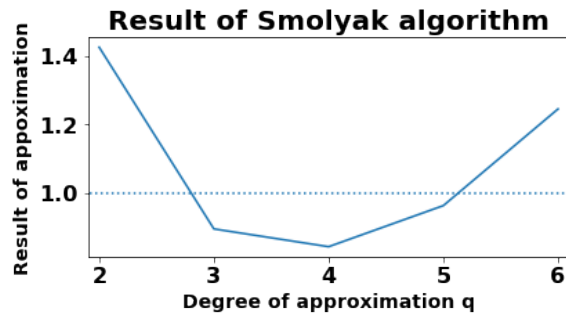
Newton-Cotes



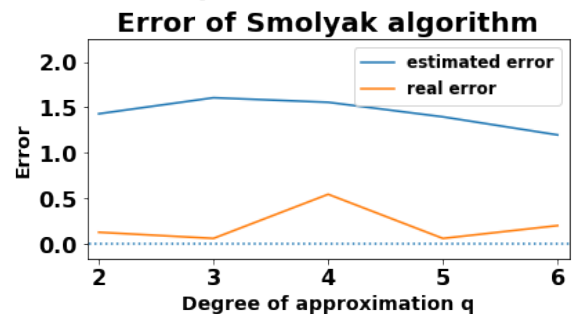
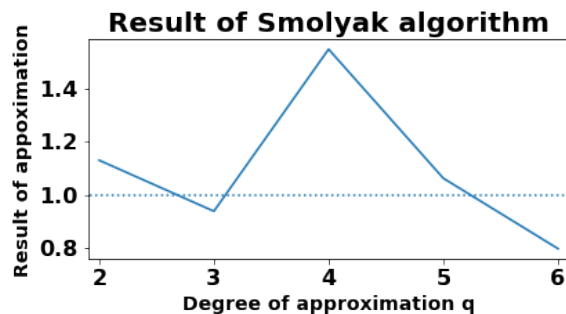
Trapezoidal

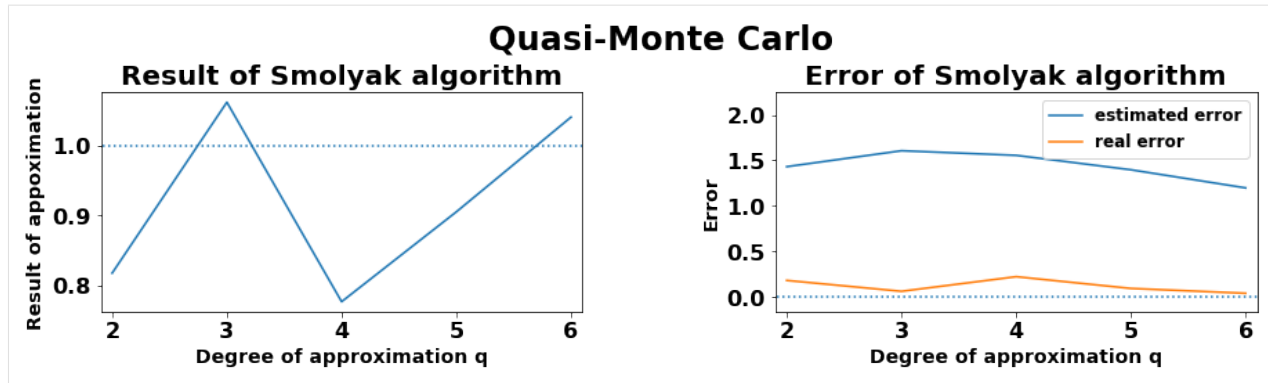


Monte Carlo (nested)



Monte Carlo (non-nested)





The result of the approximation and the error each should converge towards the dotted line. The estimated error for the probabilistic quadratures is not really significant, because the central limit theorem could not be applied here. The probabilistic quadratures for $q_i = 1$ here use three points.

2.3 Higher dimension

2.3.1 Approximation of constant

In the end, we want to show that the algorithm is working correct in different dimension. For this we first calculate the integral $f_d(\vec{x}) = 1$, to see, if the Smolyak algorithm is working well. Here the only interesting point is the difference of the result to 1. This should be sufficiently small to be explainable by the machine epsilon. After this we calculated the integral of $f_d(\vec{x}) = 2^d \cdot \prod_{i=0}^d x_i$ on the interval $[0, 1]^d$. This integral is approximated exact by the trapezoidal quadrature. Apart from that an impression of the error for the non-deterministic quadratures.

The data is loaded, because the calculation takes about 10 minutes. The calculations are repeated 10 times for $q = d + 10$.

```
[13]: color_str = [ "fuchsia", "navy", "limegreen", "red"]

plt.rc('font', size=14, weight="bold")
os.chdir("Data")
approx_simple_highdim_fct = bearcats.read_pickle("integral_over_one.pkl")
np.mean(approx_simple_highdim_fct["2", results], axis=2).shape
fig, ax = plt.subplots(2, 2, figsize=(15, 7))

for dim in range(3, 7):
    error = abs(1 - np.mean(approx_simple_highdim_fct[str(dim) + ", results"], axis=2))

    # Time plot

    plt.subplots_adjust(hspace=0.8, wspace=0.4)

    mat = ax[(dim-3)%2, int((dim-3)/2)].pcolor(error)
    fig.colorbar(mat, ax=ax[(dim-3)%2, int((dim-3)/2)])

    ax[(dim-3)%2, int((dim-3)/2)].set_ylabel("quadrature", fontsize=14, fontweight=
    ↪ "bold")
    ax[(dim-3)%2, int((dim-3)/2)].set_yticks([0.5, 1.5, 2.5, 3.5])
```

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```

ax[(dim-3)%2, int((dim-3)/2)].set_yticklabels(["N.C.", "T.", "M.C.", "Q.M.C."],
↪ fontsize=14, fontweight="bold")
ax[(dim-3)%2, int((dim-3)/2)].set_xlabel("degree of approximation q", fontsize=14, ↪
↪ fontweight="bold")
ax[(dim-3)%2, int((dim-3)/2)].set_xticks([0.5, 3.5, 6.5, 9.5])
ax[(dim-3)%2, int((dim-3)/2)].set_xticklabels([dim, dim+3, dim+6, dim+9], fontsize=14,
↪ fontweight="bold")
ax[(dim-3)%2, int((dim-3)/2)].set_title("Error in dimension " + str(dim),
↪ fontsize=14, fontweight="bold")

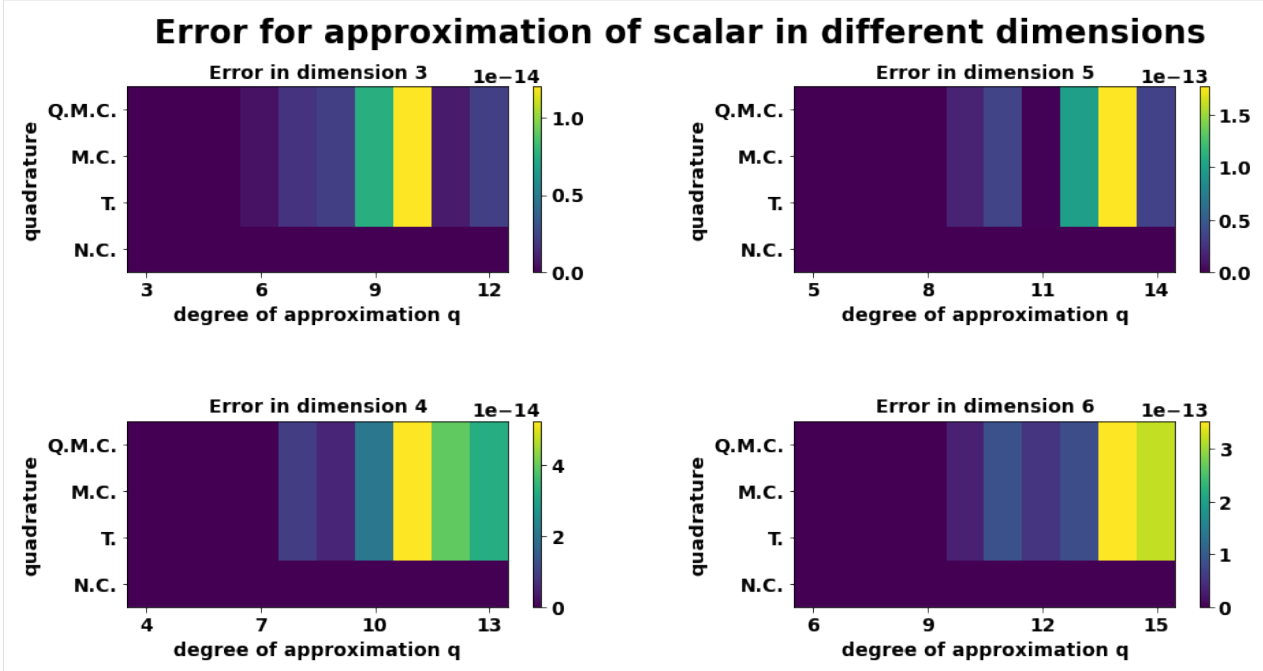
```

```

fig.suptitle("Error for approximation of scalar in different dimensions\n", ↪
↪ fontsize=24, fontweight="bold")

```

```
plt.show()
```



It turns out that the error is in the suspected range of the machine epsilon. Furthermore, it is not systematically increasing with q . This means that the different summands used in the Smolyak algorithm are summed up in a proper way. The calculation was repeated 25 times and every time the error stayed the same.

The bottom line is 0, because the Newton-Cotes algorithm could not be applied for any one-dimensional q bigger 5. Up to that point no error occurred.

2.3.2 Approximation of simple analytical function

Next we show the results of the approximation.

First we show the average of 25 calculation. As mentioned the deterministic quadratures are exact for this functions.

```

[14]: color_str = [ "fuchsia", "navy", "limegreen", "red"]

approx_simple_highdim_fct = bearcats.read_pickle("integral_over_polynomial.pkl")
fig = plt.figure(figsize=(15,15))

```

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```

for dim in range(3,7):
    approx = np.mean(approx_simple_highdim_fct[str(dim) + ", results"], axis=2)

    # Time plot

    plt.subplots_adjust(hspace=0.4, wspace=0.4)
    plt.subplot(2,2,dim-2)

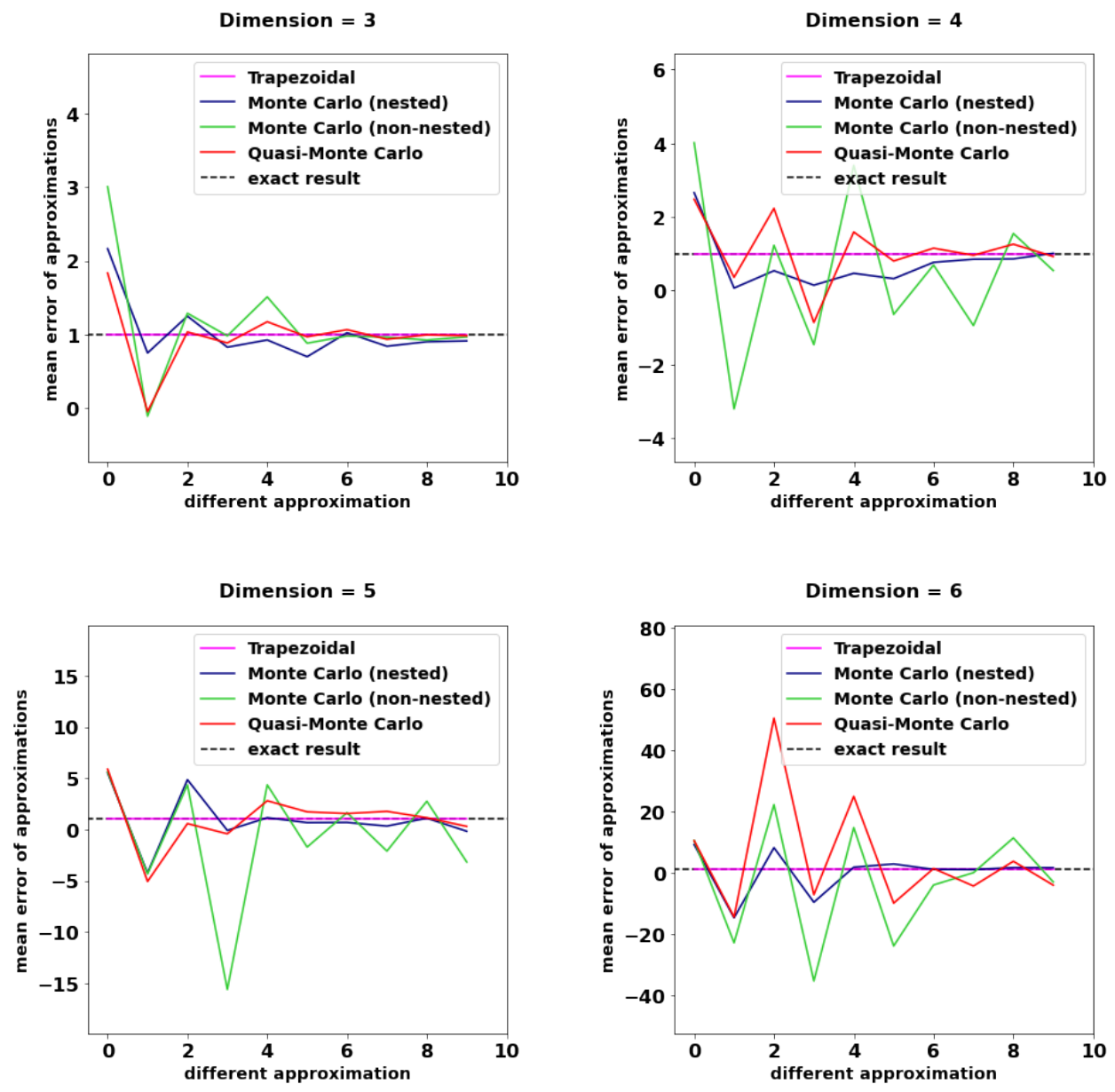
    for k_1 in range( approx.shape[0]):
        plt.plot( (approx[k_1,:]), color = color_str[k_1], label=option_list[k_1+1])

    plt.xlim(-0.5,10,5)
    plt.hlines(1, xmin=-1, xmax = 11, linestyle="--", color= "black", label="exact_
↪result")
    plt.legend(loc=1)
    plt.ylabel("mean error of approximations",fontsize = 14,fontweight = "bold")
    plt.xlabel("different approximation",fontsize = 14,fontweight = "bold")
    plt.ylim(np.max(approx) - abs(np.max(approx)-np.min(approx)) *1.2, np.max([np.
↪max(approx)*1.6,abs(np.max(approx)- abs(np.max(approx)-np.min(approx))*1.2)]))
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)
    plt.title("Dimension = " + str(dim)+"\n", fontsize = 16,fontweight = "bold")
fig.suptitle("Approximation of analytical function in different dimensions",
↪fontsize=24,fontweight="bold")

plt.show()

```

Approximation of analytical function in different dimensions



The deterministic quadratures are exact. The non deterministic quadratures tend to be relatively inaccurate especially in higher dimensions

Now the errors of the non-deterministic approximations are shown in a logarithmic scale.

```
[15]: plt.rc('font', size=14, weight="bold")
approx_simple_highdim_fct = bearcats.read_pickle("integral_over_polynomial.pkl")

fig, ax = plt.subplots(2,2,figsize=(15,7))
```

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```

for dim in range(3,7):
    error = abs(1 - np.mean(approx_simple_highdim_fct[str(dim) + ", results"][1:,:],
    ↪axis=2))

    # Time plot

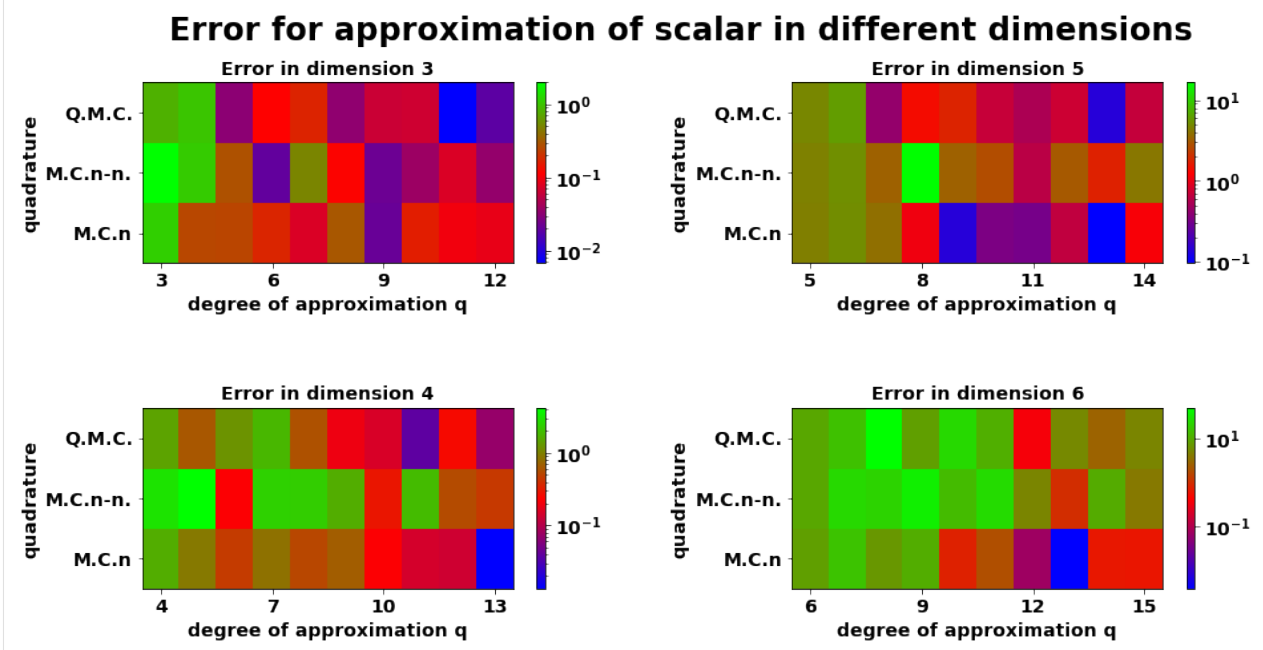
    plt.subplots_adjust(hspace=0.8, wspace=0.4)

    mat = ax[(dim-3)%2, int((dim-3)/2)].pcolor(error,norm=colors.LogNorm(vmin=error.
    ↪min(), vmax=error.max()),
        cmap='brg')
    fig.colorbar(mat, ax=ax[(dim-3)%2, int((dim-3)/2)])

    ax[(dim-3)%2, int((dim-3)/2)].set_ylabel("quadrature",fontsize=14, fontweight=
    ↪"bold")
    ax[(dim-3)%2, int((dim-3)/2)].set_yticks([0.5,1.5,2.5])
    ax[(dim-3)%2, int((dim-3)/2)].set_yticklabels(["M.C.n", "M.C.n-n.", "Q.M.C."],
    ↪fontsize=14, fontweight="bold")
    ax[(dim-3)%2, int((dim-3)/2)].set_xlabel("degree of approximation q",fontsize=14,
    ↪fontweight="bold")
    ax[(dim-3)%2, int((dim-3)/2)].set_xticks([0.5,3.5,6.5,9.5])
    ax[(dim-3)%2, int((dim-3)/2)].set_xticklabels([dim,dim+3,dim+6,dim+9],fontsize=14,
    ↪fontweight="bold")
    ax[(dim-3)%2, int((dim-3)/2)].set_title("Error in dimension "+ str(dim),
    ↪fontsize=14, fontweight="bold")

fig.suptitle("Error for approximation of scalar in different dimensions\n",
    ↪fontsize=24,fontweight="bold")
plt.show()

```



```
[1]: '''
import Methodes_Studienproject.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
↪one
import numpy as np
import time
import pandas as bearcats

# for proof of prnciple
variables = ["(x_1)"]
#function = ["1"]
functions = ["2 * x_1"]
option_list = ["Trapezoidal",
               "Monte Carlo (nested)",
               "Monte Carlo (non-nested)",
               "Quasi-Monte Carlo"]

repetitions = 25
results = {}
for dim in range(2, 7):
    print(dim)
    variables.append(variables[-1][:-1]+"", x_ "+ str(dim)+"")
    functions.append(functions[-1]+"* 2 * x_ "+str(dim))
    # functions.append(functions[-1])
    results_one_dim= np.empty((4,10,repetitions,3))
    run_times = np.empty((4))
    for k_1 in range(len(option_list)):

        start = time.time()
        for k_3 in range(dim,dim+10):
            print(k_3)
            for k_2 in range(repetitions):
                results_one_dim[k_1, k_3-dim, k_2, :] = Studieproject_one.controller_
↪smolyak(functions[-1], variables[-1], option_list[k_1], k_3)
                run_times[k_1] = (time.time()-start)/25
            new_results = { str(dim)+ " , results": results_one_dim[:, :, :, 0], str(dim)+ " , time
↪": run_times}
            results.update(new_results)
bearcats.to_pickle(results, "integral_over_polynomial.pkl")
# bearcats.to_pickle(results, "integral_over_one.pkl")
'''
print("If you want to generate data, remove quotation marks.")

If you want to generate data, remove quotation marks.
```

```
[ ]:
```


APPROXIMATE π

The way not not steady functions could be used in the project is illustrated in a well known basic example for the use of the Monte Carlo method. This is the approximation of pi. Here the it is used that the area of the (unit) circle is proportional to pi. Hence it is possible to approximate π by setting all points in the circle to 1 and all others to 0. Then the average of these value converges to π . Because we restrict our self to the interval $[0, 1]^n$, we have

$$\pi = 4 \cdot A(2, q)$$

and

$$\pi = 6 \cdot A(3, q),$$

with $A(n, q)$ being the result of the Smolyak algorithm in n dimensions and the degree of approximation q . Because the function is not derivable, it is not possible to estimate the error for the algorithm for the deterministic quadratures. We will just compare them to the value of π used by numpy.

```
[6]: import os
      os.chdir("..")

      import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
      ↪one
      import matplotlib.pyplot as plt
      import sympy as sp
      import numpy as np
      import itertools as itt
```

3.1 The functions used

```
[2]: # Functions receiving 2 or 3 matrices of the the dimension 2 or 3. Each matirce_
      ↪contains the coordinate in
      # the component x, y or z. The function returns a martice of the shape of the input_
      ↪components with 1 if
      # the 2-norm of the coordinate is smaller or equal 1 and 0 otherwise.
      def circle(X_1, X_2):
          save_mat = np.ones(X_1.shape)

          for index_vec, vectors in enumerate(X_1):
              for index_col, x_1 in enumerate(vectors):
```

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```

        if x_1**2 + X_2[index_vec, index_col]**2 > 1:
            save_mat[index_vec, index_col] = 0
    return save_mat

def ball(X_1, X_2, X_3):
    save_mat = np.ones(X_1.shape)
    for index_mat, matrix in enumerate(X_1):
        for index_vec, vectors in enumerate(matrix):
            for index_col, x_1 in enumerate(vectors):

                if x_1**2 + X_2[index_mat, index_vec, index_col]**2 + X_3[index_mat,
↪index_vec, index_col]**2 > 1:
                    save_mat[index_mat, index_vec, index_col] = 0
    return save_mat

color_str = [ "fuchsia", "navy", "limegreen", "red"]
variables_string_rewrite = "(x_1, x_2, x_3)"
variables_string_rewrite = variables_string_rewrite.replace(" ", "")
variables = variables_string_rewrite.strip('()').split(',')
variables_string_rewrite = variables_string_rewrite.strip('()').split(',')

if len(variables) == 0:
    raise Exception("There at least has to be one variable.")

for f in range(len(variables)):
    variables[f] = sp.Symbol("x_" + str(f))

option_list = ["Trapezoidal",
               "Monte Carlo (nested)",
               "Monte Carlo (non-nested)",
               "Quasi-Monte Carlo"]

# Approximation of pi with different q and degrees of approx.
lower_bound, upper_bound = [8,15]
approx_pi_circle = np.empty([len(option_list), upper_bound - lower_bound])
for index_option, option in enumerate(option_list):
    for index_q, degree_of_approx in enumerate(range(lower_bound, upper_bound)):
        approx_pi_circle[index_option, index_q] = 4 * Studieproject_one.controller_
↪smolyak(circle, variables[:2], option, degree_of_approx, function_given=True, no_
↪error=True)[0]

approx_pi_ball = np.empty([len(option_list), upper_bound - lower_bound])
for index_option, option in enumerate(option_list):
    for index_q, degree_of_approx in enumerate(range(lower_bound, upper_bound)):
        approx_pi_ball[index_option, index_q] = 6 * Studieproject_one.controller_
↪smolyak(ball, variables, option, degree_of_approx, function_given=True, no_
↪error=True)[0]

```


3.2 Illustration of the principle of approximation for 2 dimensions

```
[3]: plt.rc('font', size=14, weight="bold")
def grid_points(quadrature: str, q: int, a=0,b=1):
    list_of_points = []
    dim = 2

    # Because the points used for the deterministic quadratures are not changed, it_
    ↪is faster to call them
    # at the beginning before starting the Smolyak alg.
    if "Carlo" in quadrature:
        q = q + 1

    # For every one dimensional degree of approx. a weights and points vector
    for i in range(1, (q - dim + 2)):

        # One option for Newton-Cotes and Trapezoidal quadrature, because
        if quadrature == "Newton-Cotes" or quadrature == "Trapezoidal":
            points, weights = Studieproject_one.one_dim_trapezoidal(i, a, b)

        if quadrature == "Monte Carlo (nested)" and i > 1:
            points, weights = Studieproject_one.monte_carlo_quad(i-1, a, b)
            points = np.concatenate([list_of_points[-1], points])

        if quadrature == "Monte Carlo (non-nested)" or (quadrature == "Monte Carlo_
    ↪(nested)" and i == 1) :
            points, weights = Studieproject_one.monte_carlo_quad(i, a, b)

        if quadrature == "Quasi-Monte Carlo":
            points, weights = Studieproject_one.qmc_quad(i, a, b)

        list_of_points.append(points)

    # Now we get the combinations of one-dimensional degrees of approx. in  $Q(q,d)$ 
    rng = list(range(q)) * dim
    rng = [x + 1 for x in rng]

    possible_combinations = list(set(i for i in itt.permutations(rng, dim) if (q -
    ↪dim) < sum(i) < q + 1))
    number_approx = len(possible_combinations)
    meshgrid_list_x = []
    meshgrid_list_y = []
    for i in range(number_approx):

        # gridpoints and the weight vectors are put into separate lists to make it_
    ↪easier to
        # make a meshgrid
        current_tuple = possible_combinations[i]
        current_points = []

        for j in range(len(current_tuple)):
            current_points.append(list_of_points[current_tuple[j] - 1])

        # meshgrid yields a np.array with all coordinates needed for alg
        # coordinate and respective weight could be found in the same places in the_
    ↪array
```

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```

        meshgrid_points = np.array(np.meshgrid(*current_points))
        meshgrid_list_x.append(meshgrid_points[0])
        meshgrid_list_y.append(meshgrid_points[1])

    return [meshgrid_list_x, meshgrid_list_y]

def make_plot_of_grids():
    a = 0
    b = 1
    option_list = ["Newton-Cotes",
                  "Monte Carlo (nested)",
                  "Monte Carlo (non-nested)",
                  "Quasi-Monte Carlo"]
    fig, ax = plt.subplots(2,2,figsize=(15,15))
    plt.subplots_adjust(hspace=0.4, wspace=0.4)

    for i in range(len(option_list)):
        x, y = grid_points(option_list[i], 6)
        circle_pi = list()
        for index_x, _ in enumerate(x):

            circle_pi.append(circle(x[index_x], y[index_x]))
            ax[i%2,int(i/2)].scatter(x[index_x][circle_pi[-1]==1], y[index_x][circle_
↪pi[-1]==1], color="blue",label="1")
            ax[i%2,int(i/2)].scatter(x[index_x][circle_pi[-1]==0], y[index_x][circle_
↪pi[-1]==0], color="red",label="0")
            ax[i%2,int(i/2)].set_title(option_list[i],fontsize=14, fontweight="bold")

        # print circle
        x_circle = np.linspace(0,np.pi / 2,100)
        r_x = np.cos(x_circle)
        r_y = np.sin(x_circle)
        ax[i%2,int(i/2)].plot(r_x,r_y,color="blue")

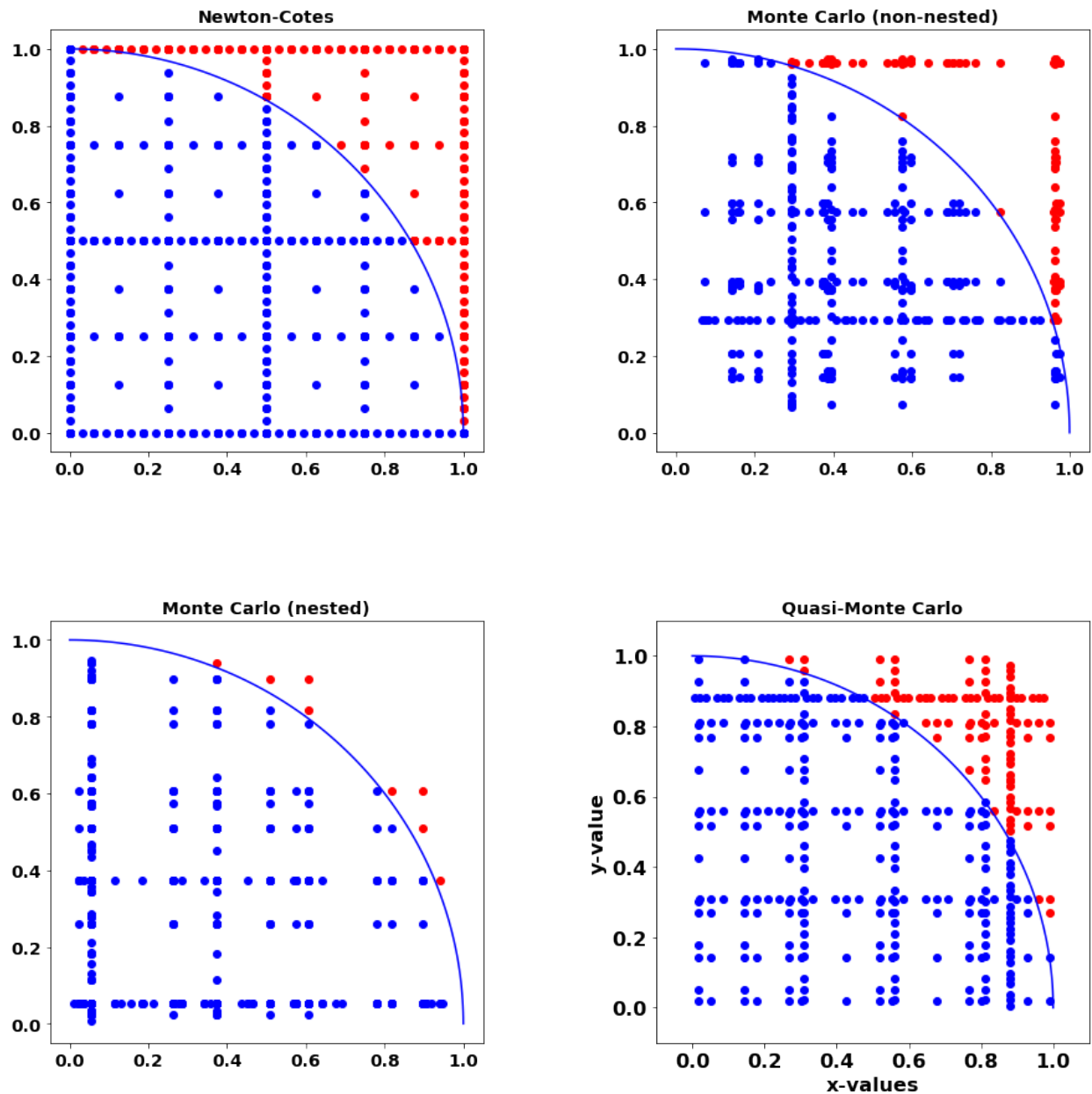
    # Some cosmetic changes in plot
    plt.xlabel("x-values", fontsize=16, fontweight="bold")
    plt.ylabel("y-value", fontsize=16, fontweight="bold")
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)
    plt.xlim(a - (b - a) / 10, b + (b - a) / 10)
    plt.ylim(a - (b - a) / 10, b + (b - a) / 10)
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)

    fig.suptitle("Illustration of principle", fontsize=24,fontweight="bold")
    plt.show()

make_plot_of_grids()

```

Illustration of principle



The blue points coincides with the points with value 1 and the red ones with the value 0.

```
[4]: fig, ax = plt.subplots(1,2,figsize=(15,5))

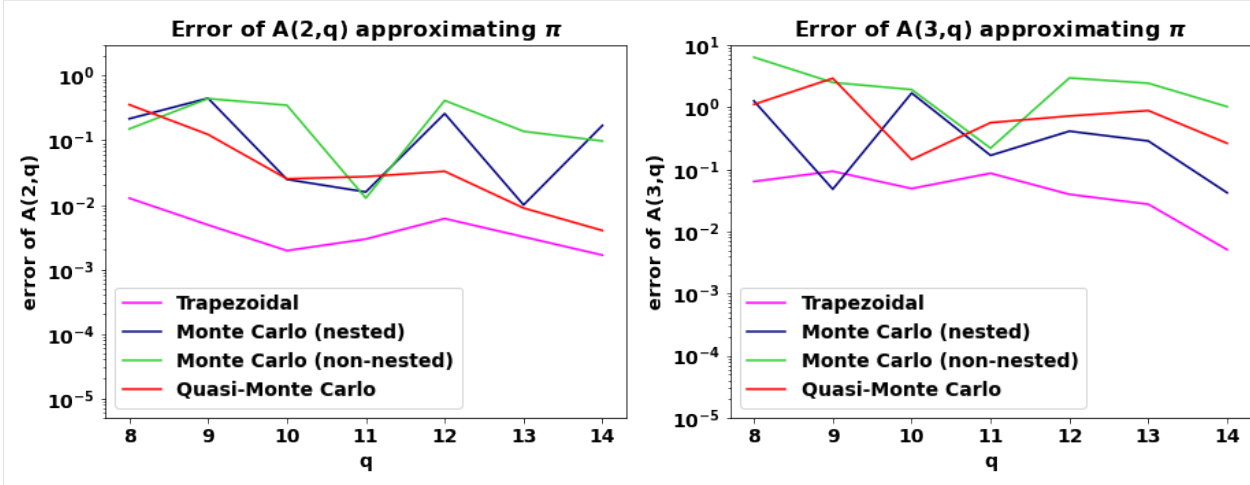
for index_option, option in enumerate(option_list):
    ax[0].plot(range(lower_bound, upper_bound), abs(np.pi-approx_pi_circle[index_
    ↪option, :]), color = color_str[index_option],
                label=option)
ax[0].set_yscale("log")
```

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```
ax[0].set_ylim(0.000005,3)
ax[0].set_ylabel("error of A(2,q)",fontsize=14, fontweight="bold")
ax[0].set_xlabel("q",fontsize=14, fontweight="bold")
ax[0].set_xticks(list(range(lower_bound, upper_bound)))
ax[0].set_xticklabels(list(range(lower_bound, upper_bound)),fontsize=14, fontweight=
    ↳ "bold")
ax[0].set_title(r"Error of A(2,q) approximating  $\pi$ ",fontsize=16, fontweight="bold")
ax[0].legend()

for index_option, option in enumerate(option_list):
    ax[1].plot(range(lower_bound, upper_bound), abs(np.pi-approx_pi_ball[index_option,
    ↳ :]), color = color_str[index_option],
        label=option)
ax[1].set_yscale("log")
ax[1].set_ylabel("error of A(3,q)",fontsize=14, fontweight="bold")
ax[1].set_xlabel("q",fontsize=14, fontweight="bold")
ax[1].set_xticks(list(range(lower_bound, upper_bound)))
ax[1].set_xticklabels(list(range(lower_bound, upper_bound)),fontsize=14, fontweight=
    ↳ "bold")
ax[1].set_title(r"Error of A(3,q) approximating  $\pi$ ",fontsize=16, fontweight="bold")
ax[1].legend()
ax[1].set_ylim(0.00001,10)
plt.show()
```



[]:

STATISTICAL EVALUATION FOR ERROR FOR PROBABILISTIC QUADRATURES

In this script we want to evaluate statistical data. For this we load data, which was calculated and saved in a .csv/ .pkt file, because generating file takes some hours.

```
[3]: import matplotlib.pyplot as plt

import os
os.chdir("../")
from Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point import *
```

4.1 First approximation of distribution

4.1.1 One point for $q = 1$

First we want to give a feeling for the distribution of the results of the Smolyak algorithm using different quadratures and the effect of higher degrees of approximation.

We only show the first degrees of approximation, because otherwise nothing could be seen, due to the scaling of the plot.

```
[11]: try:
        os.chdir("Data")
    except:
        pass
    results_one = load_stat_data("stat_approach_2_to_20_result_13_11_one.csv")
    results_three = load_stat_data("stat_approach_2_to_20_result_13_11_three.csv")

[12]: title_string = ["One point for q = 1 on the one dimensional level \n\n Monte Carlo_
    ↳quadrature (nested)\n", "Monte Carlo quadrature (non-nested)\n",
        "Quasi-Monte Carlo quadrature \n"]

results = results_one
for k_1 in range(len(results[0])):
    fig = plt.figure(figsize=(20,7))
    ax = plt.axes()
    boxprops = dict(linewidth=3.0, color='black')
    whiskerprops = dict(linestyle='-',linewidth=3.0, color='black')
    medianprops = dict(linestyle='-',linewidth=3.0, color='red')
    for k_2 in range(len(results[0][0])-14):
        y =results[0][k_1][k_2]
```

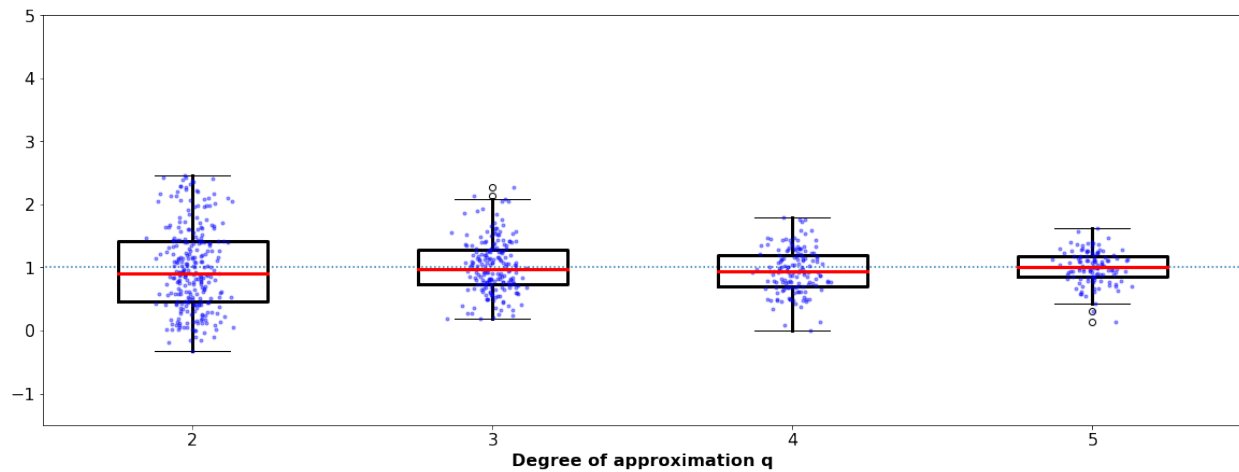
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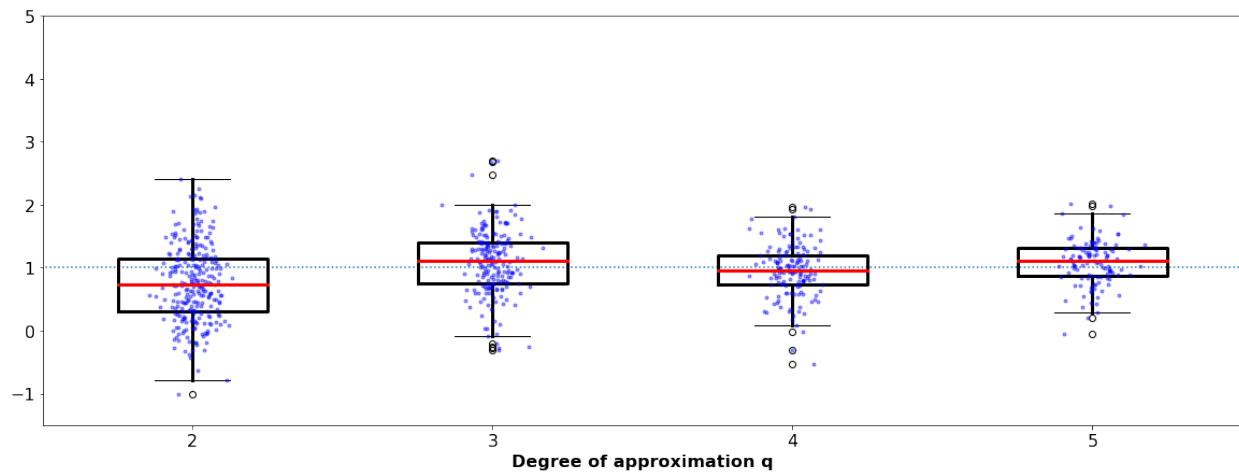
```
bp = plt.boxplot(y, positions = [k_2 + 1], widths = 0.5, boxprops=boxprops,
                whiskerprops = whiskerprops, medianprops = medianprops)
x = np.random.normal(k_2+1, 0.05, size=len(y))
plt.plot(x, y, 'b.', alpha=0.4)
plt.hlines(1,0,10, linestyle= "dotted")
plt.xlim((1.5, 5.5))
plt.ylim((-1.5,5))
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.title(title_string[k_1], fontsize = 24, fontweight = "bold")
plt.xlabel("Degree of approximation q", fontsize= 16, fontweight = "bold")
plt.show()
```

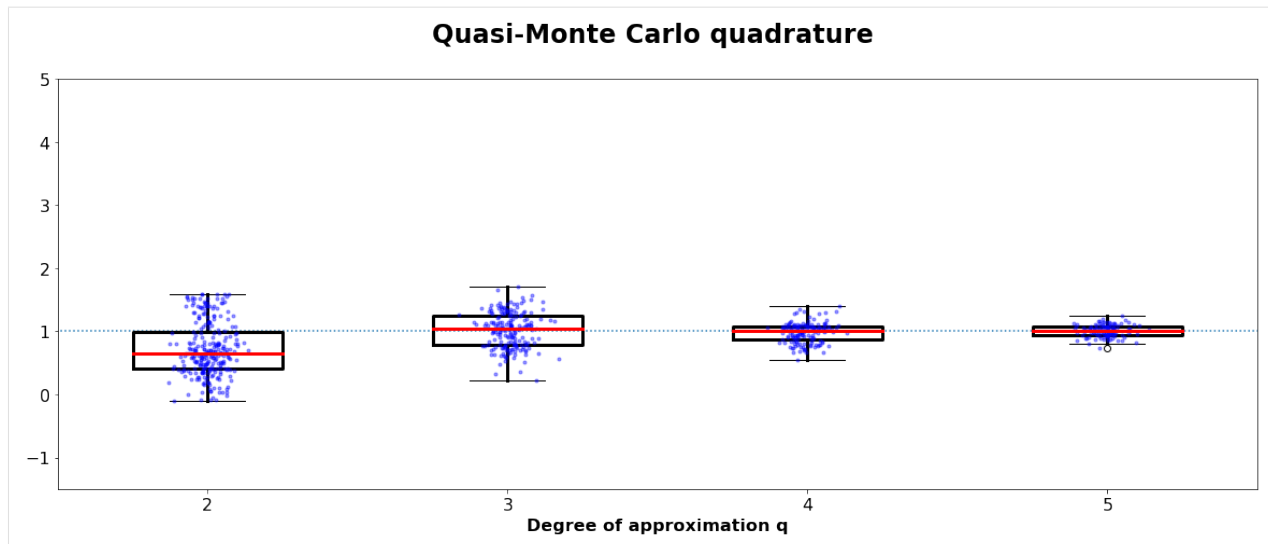
One point for $q = 1$ on the one dimensional level

Monte Carlo quadrature (nested)



Monte Carlo quadrature (non-nested)





Boxplot of at least 200 calculations of the function $\pi^2/4 \cdot \sin(\pi \cdot x) \cdot \sin(\pi \cdot y)$ using probabilistic quadratures. Here $q = 2$ implies that the integral is approximated by a point evaluation. Thus, the high variance for low q is expected.

By taking a look at the variance of the result and comparing it to the error estimation, we get a more quantitative impression about the goodness of the approximation

4.1.2 Three points

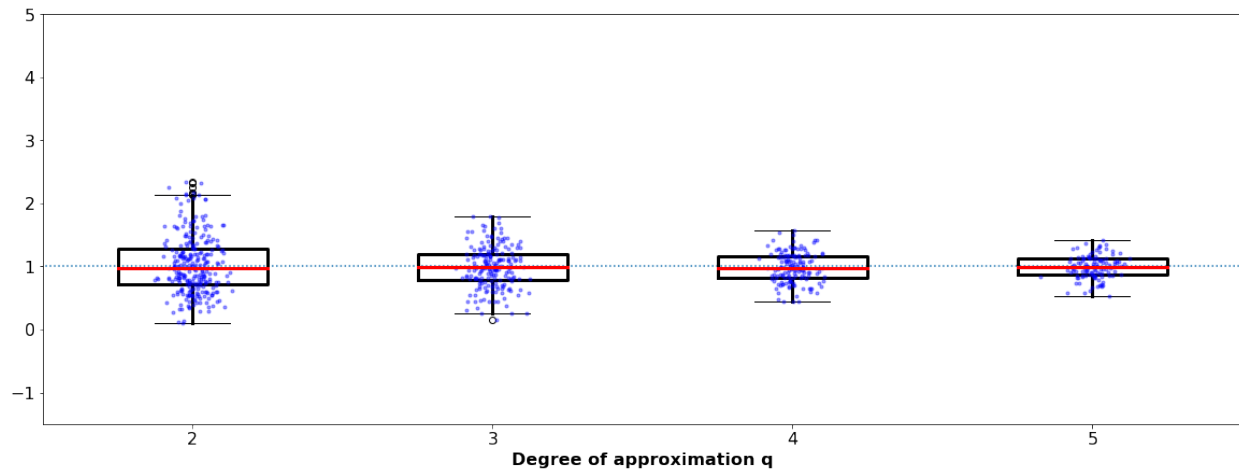
For comparison we also show the result, if for $q = 1$ on the one dimensional level we use 3 points.

```
[13]: title_string = ["Three points for q = 1 on the one dimensional level \n\n Monte Carlo ↵
↵quadrature (nested)\n", "Monte Carlo quadrature (non-nested)\n",
                    "Quasi-Monte Carlo quadrature \n"]

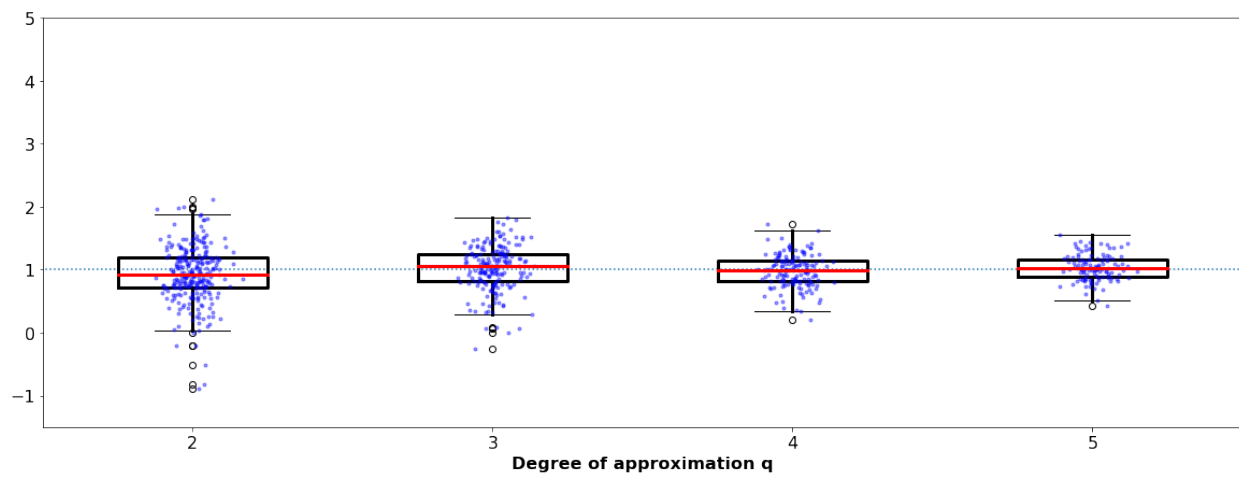
results = results_three
for k_1 in range(len(results[0])):
    fig = plt.figure(figsize=(20,7))
    ax = plt.axes()
    boxprops = dict(linewidth=3.0, color='black')
    whiskerprops = dict(linestyle='-', linewidth=3.0, color='black')
    medianprops = dict(linestyle='-', linewidth=3.0, color='red')
    for k_2 in range(len(results[0][0])-14):
        y = results[0][k_1][k_2]
        bp = plt.boxplot(y, positions = [k_2 + 1], widths = 0.5, boxprops=boxprops,
                        whiskerprops = whiskerprops, medianprops = medianprops)
        x = np.random.normal(k_2+1, 0.05, size=len(y))
        plt.plot(x, y, 'b.', alpha=0.4)
    plt.hlines(1,0,10, linestyles= "dotted")
    plt.xlim((1.5, 5.5))
    plt.ylim((-1.5,5))
    plt.xticks(fontsize=16)
    plt.yticks(fontsize=16)
    plt.title(title_string[k_1], fontsize = 24, fontweight = "bold")
    plt.xlabel("Degree of approximation q", fontsize= 16, fontweight = "bold")
    plt.show()
```

Three points for $q = 1$ on the one dimensional level

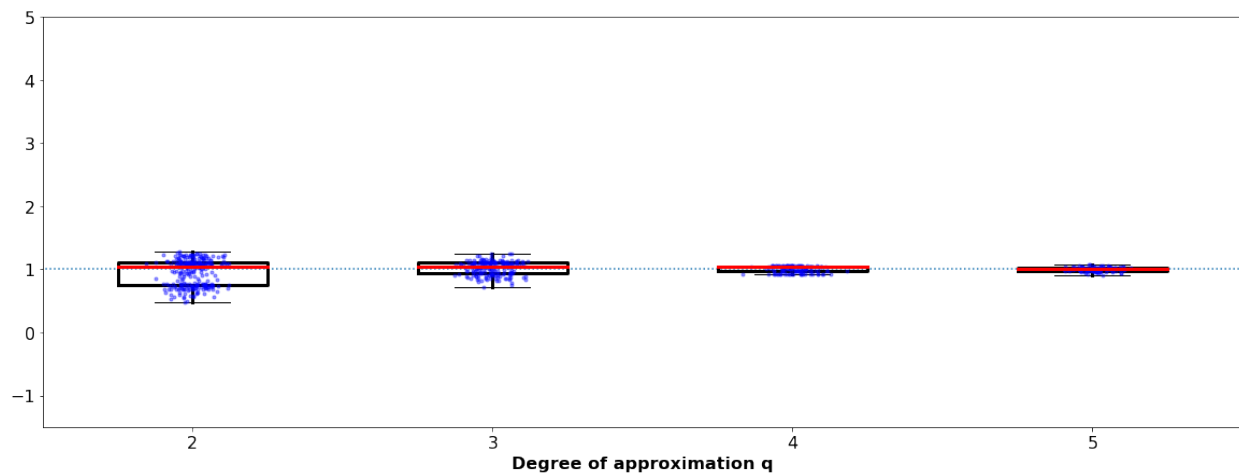
Monte Carlo quadrature (nested)



Monte Carlo quadrature (non-nested)



Quasi-Monte Carlo quadrature



The approximation in this case is much more exact. A comparison of the runtime and the smaller error will be done later on.

4.2 Statistical parameters of distribution of results

```
[14]: # Calculation of mean and variance of the result and estimated error of approximation
mean_result = np.ones((2,3,len(results[0][0])))
sigma_result = np.ones((2,3,len(results[0][0])))

mean_error = np.ones((2,3,len(results[0][0])))
sigma_error = np.ones((2,3,len(results[0][0])))

runtime = np.ones((2,3,len(results[0][0])))
runtime_sig = np.ones((2,3,len(results[0][0])))

for k_3 in range(2):
    if k_3 == 0:
        results = results_one
    else:
        results = results_three

    for k_1 in range(3):
        for k_2 in range(len(results[0][0])):
            mean_result[k_3][k_1][k_2] = np.mean(results[0][k_1][k_2])
            mean_error[k_3][k_1][k_2] = np.mean(results[1][k_1][k_2])

            # We again choose the 95 % level
            sigma_result[k_3][k_1][k_2] = 2 * np.sqrt(np.var(results[0][k_1][k_2]))
            sigma_error[k_3][k_1][k_2] = 2 * np.sqrt(np.var(results[1][k_1][k_2]))

            runtime[k_3][k_1][k_2] = np.mean(results[2][k_1][k_2])
            runtime_sig[k_3][k_1][k_2] = np.sqrt(np.var(results[2][k_1][k_2]))

# What can be found in the different dimensions of the matrices: shape: (2,3,23):
# x: x= 0 q_i uses one point, x=1 q_i uses three points;
# y: y=0 Monte Carlo (nested), y=1 Monte Carlo (non-nested), y=2 Quasi-Monte Carlo_
↳ (nested)
# z: element related to z = q-2
```

```
[15]: from sklearn import datasets, linear_model
color_code = ["r-", "g-", "b-", "r--", "g--", "b--", "r:", "g:", "b:"]
legend_label = ["Monte Carlo quadrature (nested)", "Monte Carlo quadrature (non-
↳ nested)",
                "Quasi-Monte Carlo quadrature"]

fig = plt.figure(figsize=(15,6))

for k_1 in range(6):

    # solution not beautiful. Result of afterwards implementing algorithm with one_
↳ point used for q_i = 1.
    if k_1 < 3:
        k_2 = 1
```

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```

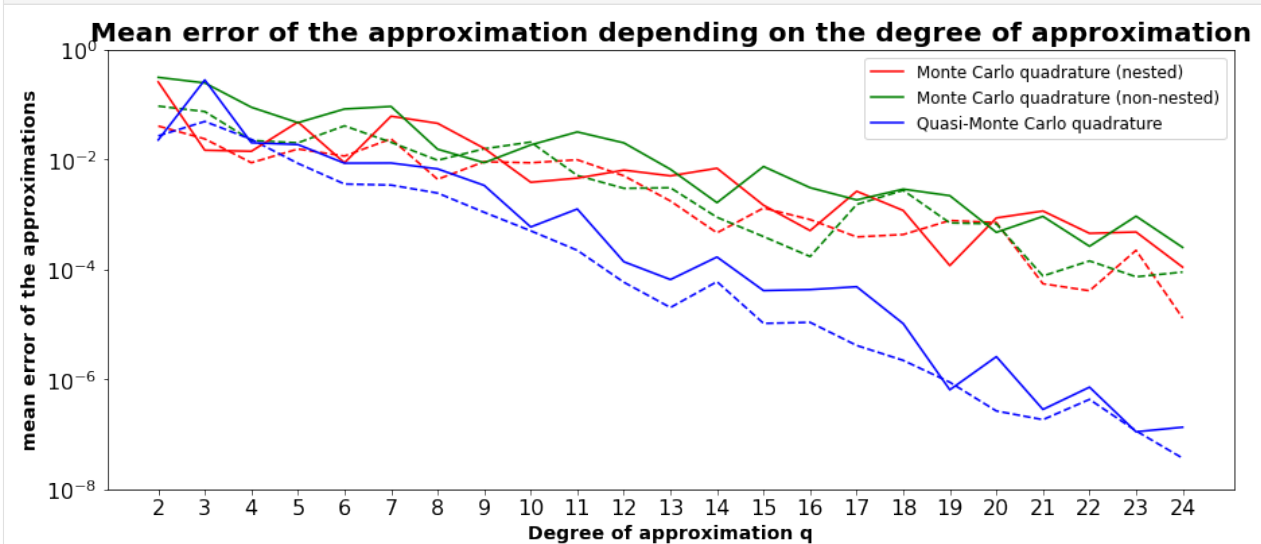
        k_1 = k_1 % 3
    else:
        k_2 = 0
        k_1 = k_1 % 3

    mean_error_real = abs(mean_result[k_2][k_1]-1)
    mean_sigma_real = sigma_result[k_2][k_1]
#     plt.subplots_adjust(hspace=0.4, wspace=0.4)

    degree_of_app = len(results[0][0])
    if k_2 == 0:
        plt.plot(list(range(2,2+degree_of_app)),mean_error_real,color_code[k_1],_
        ↪label=legend_label[k_1])
    else:
        plt.plot(list(range(2,2+degree_of_app)),mean_error_real,color_code[k_1+3*k_2])

plt.ylim(min(abs(mean_result[1][2]-1)*0.5),max(abs(mean_result[0][0]-1)*1.5))
plt.yscale("log")
plt.ylim([1e-8,1])
plt.legend(fontsize=12, loc=1)
plt.xlabel("Degree of approximation q", fontsize = 14,fontweight = "bold")
plt.ylabel("mean error of the approximations",fontsize = 14,fontweight = "bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.xticks(range(2,len(results[0][0])+2))
plt.title("Mean error of the approximation depending on the degree of approximation",
        fontsize = 20, fontweight = "bold")
plt.show()

```



This graphic shows the mean of the real error of the different approximations. The solid lines are related to the approximations using one point for $q_i = 1$, while the dotted lines were calculated for $q_i = 3$. The points for q near 24 are statistically speaking less significant because about 25 approximations were calculated.

```

[16]: fig = plt.figure(figsize=(15,6))
      for k_1 in range(6):

```

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```

# solution not beautiful. Result of afterwards implementing algorithm with one_
↪point used for q_i = 1.
    if k_1 < 3:
        k_2 = 1
        k_1 = k_1 % 3
    else:
        k_2 = 0
        k_1 = k_1 % 3
for k_1 in range(6):
    # solution not beautiful. Result of afterwards implementing algorithm with_
    ↪one point used for q_i = 1.
    if k_1 < 3:
        k_2 = 1
        k_1 = k_1 % 3
    else:
        k_2 = 0
        k_1 = k_1 % 3

mean_error_est = mean_error[k_2][k_1]

s_mean = sigma_result[k_2][k_1]
s_error = sigma_error[k_2][k_1]
mean_error_real = abs(mean_result[k_2][k_1]-1)
mean_sigma_real = sigma_result[k_2][k_1]

degree_of_app = len(results[0][0])

plt.errorbar(list(range(2,2+degree_of_app)), mean_error_est, s_error,ecolor=color_
↪code[k_1][0],
              color=(0,0,0,0),marker="_",capsize=0.5)
plt.plot(list(range(2,2+degree_of_app)),mean_error_est, color_code[k_1+6])
if k_2 == 0:
    plt.plot(list(range(2,2+degree_of_app)),np.add(mean_error_real,s_mean),color_
↪code[k_1+ k_2 * 3],label=legend_label[k_1])
else:
    plt.plot(list(range(2,2+degree_of_app)),np.add(mean_error_real,s_mean),color_
↪code[k_1+ k_2 *3])

if k_1 == 2:
    regression_data_x = [[k_1] for k_1 in list(range(2,2+degree_of_app))]
    regression_data_y = [math.log2(k_1) for k_1 in np.add(mean_error_real,s_mean)]
    regr = linear_model.LinearRegression()
    regr.fit(regression_data_x , regression_data_y)
    regr.score(regression_data_x , regression_data_y)
    exponent = regr.coef_

    regression_data_x = [[k_1] for k_1 in list(range(2,2+degree_of_app))]
    regression_data_y = [math.log2(k_1) for k_1 in np.add(mean_error_real,s_mean)]
    regr = linear_model.LinearRegression()
    regr.fit(regression_data_x , regression_data_y)
    regr.score(regression_data_x , regression_data_y)
    exponent = regr.coef_

    regression_y = [np.add(mean_error_real,s_mean)[0]*1.3*2**(exponent[0]*(x -2))_
↪ for x in list(range(0,20+degree_of_app))]
    plt.plot(list(range(0,20+degree_of_app)),regression_y, "k:")

```

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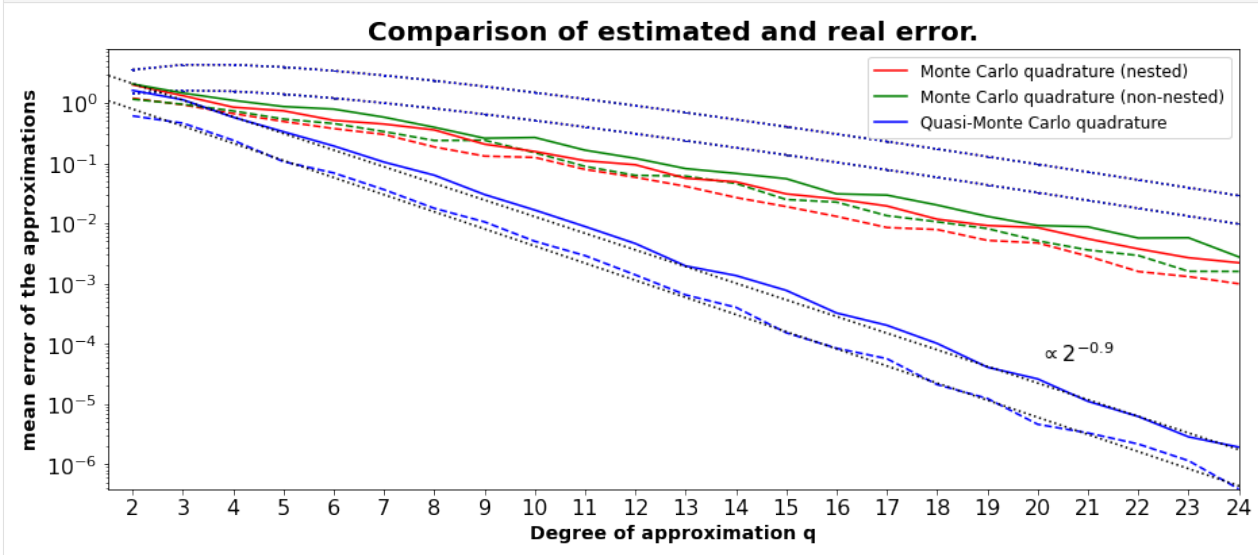
```

    if k_2 == 1:
        plt.annotate("$\propto 2^{-0.9}$", (20, regression_y[19]), xytext=(20,
        ↪ 5*10**(-5)), fontweight="bold", fontsize=16)

plt.ylim(min(abs(np.add(mean_error_real, s_mean))*0.2), max(abs(mean_error[1][2])*5))
plt.xlim(1.5, degree_of_app+0.5)
plt.yscale("log")
plt.legend(fontsize=12, loc=1)
plt.xlabel("Degree of approximation q", fontsize = 14, fontweight = "bold")
plt.ylabel("mean error of the approximations", fontsize = 14, fontweight = "bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.xticks(range(2, len(results[0][0])+2))
plt.title("Comparison of estimated and real error.",
        fontsize = 20, fontweight = "bold")

plt.show()

```



The dotted lines are the error estimations for the different quadratures. The upper one is related to the calculations using one point for $q_i = 1$. The lower lines show the error within a confidence level of 95 %". The real error bound nearly decreases proportional to $2^{**}(-1)$, if the Quasi-Monte Carlo quadrature is used. The exponent is varying a little with respect to the generated data used.

```

[18]: fig = plt.figure(figsize=(15,6))

for k_1 in range(6):

    # solution not beautiful. Result of afterwards implementing algorithm with one_
    ↪ point used for q_i = 1.
    if k_1 < 3:
        k_2 = 1
        k_1 = k_1 % 3
    else:
        k_2 = 0
        k_1 = k_1 % 3
for k_1 in range(6):
    # solution not beautiful. Result of afterwards implementing algorithm with_
    ↪ one point used for q_i = 1.

```

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```

if k_1 < 3:
    k_2 = 1
    k_1 = k_1 % 3
else:
    k_2 = 0
    k_1 = k_1 % 3
runtime_mean = runtime[k_2][k_1]
runtime_sigma = runtime_sig[k_2][k_1]

degree_of_app = len(runtime_mean)

if k_2 == 0:
    plt.plot(list(range(2,2+degree_of_app)), runtime_mean, color_code[k_1],
    ↪label=legend_label[k_1])
else:
    plt.plot(list(range(2,2+degree_of_app)), runtime_mean, color_code[k_1 + 3])
    plt.errorbar(list(range(2,2+degree_of_app)), runtime_mean, runtime_sigma,
    ↪ecolor=color_code[k_1][0],
        color=(0,0,0,0), marker="_", capsize=0.5)
if k_1 == 2:

    regression_data_x = [[k_1] for k_1 in list(range(2,2+degree_of_app))]
    regression_data_y = [math.log2(k_1) for k_1 in runtime_mean]
    regr = linear_model.LinearRegression()
    regr.fit(regression_data_x , regression_data_y)
    regr.score(regression_data_x , regression_data_y)
    exponent = regr.coef_

    regression_data_x = [[k_1] for k_1 in list(range(21,2+degree_of_app))]
    regression_data_y = [math.log2(k_1) for k_1 in runtime_mean[19:degree_of_app]]
    regr = linear_model.LinearRegression()
    regr.fit(regression_data_x , regression_data_y)
    regr.score(regression_data_x , regression_data_y)
    exponent = regr.coef_
    regression_y = [2** (regr.intercept_ + exponent*x) for x in list(range(0,
    ↪20+degree_of_app))]
    plt.plot(list(range(0,20+degree_of_app)), regression_y, "k:")

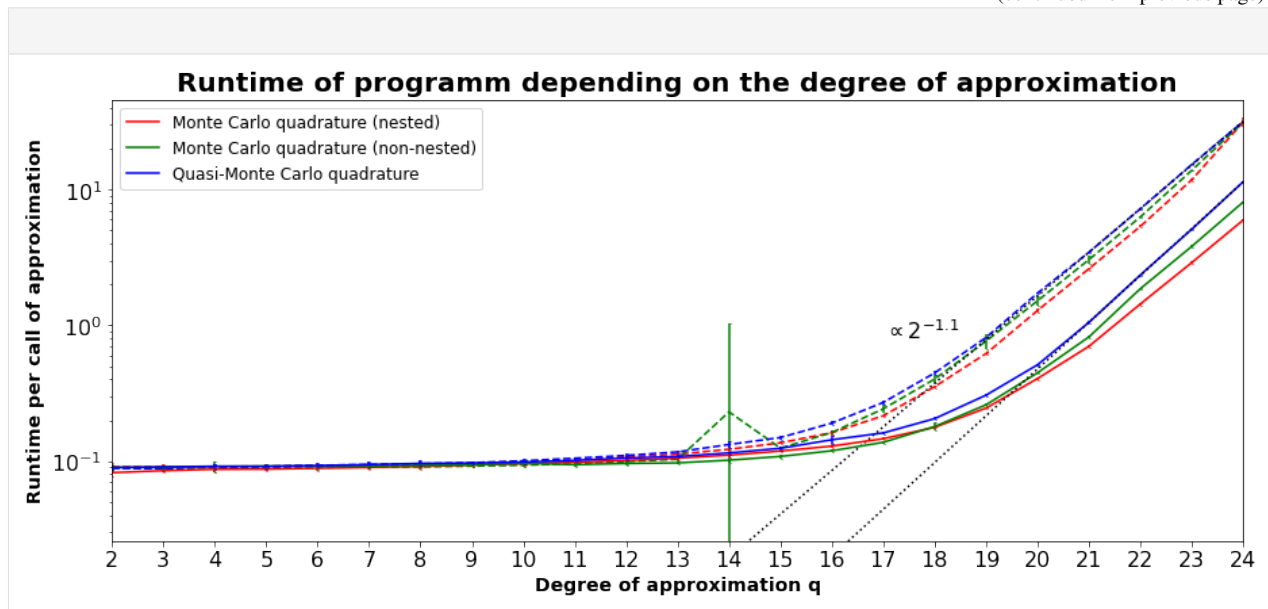
    if k_2 == 1:
        plt.annotate("$\propto 2^{-1.1}$", (20, regression_y[19]), xytext=(17,
        ↪regression_y[19]), fontweight="bold", fontsize=16)

plt.xlim(2,24)
plt.ylim(min(abs(runtime[0][2]-1)*0.5), max(abs(runtime[1][2]-1)*1.5))
plt.yscale("log")
plt.legend(fontsize=12, loc=2)
plt.xlabel("Degree of approximation q", fontsize = 14, fontweight = "bold")
plt.ylabel("Runtime per call of approximation", fontsize = 14, fontweight = "bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.xticks(range(2, len(results[0][0])+2))
plt.title("Runtime of programm depending on the degree of approximation",
    fontsize = 20, fontweight = "bold")
plt.show()

```

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It could be seen that the difference between the three quadratures is minor for this example. The variance of the runtime is relatively low. The observed runtime did not change fundamentally for changes in the implementation, such that this might be related to the algorithm itself.

4.2.1 Code used for generation of evaluated data.

```
[1]: '''
import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
↳ one

Studieproject_one.calculate_stat_data("pi^2/4 sin(pi*x)sin(pi*y)", "(x,y)", ["Monte_
↳ Carlo (nested)", "Monte Carlo (non-nested)", "Quasi-Monte Carlo"]
, "stat_approach_2_to_20_result_1_12_one.csv",
[2, 25], repetitions=2 ** 10, damping=True,
↳ damping_exp=1.2)
'''

'''
import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_three_points as
↳ Studieproject_three

Studieproject_three.calculate_stat_data("pi^2/4 sin(pi*x)sin(pi*y)", "(x,y)", ["Monte_
↳ Carlo (nested)", "Monte Carlo (non-nested)", "Quasi-Monte Carlo"]
, "stat_approach_2_to_20_result_1_12_three.csv", [2, 23], repetitions=2 ** 10,
↳ damping=True,
damping_exp=1.2)
'''
print("Approximation of analytical function in 2 dimensions. (Three points)")
Approximation of analytical function in 2 dimensions. (Three points)
```

```
[ ]:
```

EXAMPLE 15.2.6: PERIODIC FUNCTIONS (MODIFIED)

This is an easy example of a function fulfilling those conditions that make it possible to apply the results of example 15.2.6. In the example the classical Wiener measure is used. This measure so far is not implemented in this project, such that the results of this section do not hold for these examples. Nevertheless, the results are compared with the results using the Lebesgue measure. If the use of the Wiener measure is implemented, the example easily could be modified.

In example 15.2.6. in one dimension it is assumed that you have a r -times continuously differentiable function with period β . Then the norm of this space $F_1 = \tilde{C}^r([0, \beta])$ could be written as

$$\|f\|_{F_1} = |f(0)| + \max_{t \in [0, \beta]} |f^{(r)}(t)|$$

.

If we now assume $f^{(i)}(0) = f^{(i)}(\beta) = 0$ for all $i \in 1, 2, \dots, r$, we have for the d dimensional case

$$\|f\|_{F_d} = \max_{t \in [0, \beta]^d} |f^{(r, r, \dots, r)}(t)|$$

.

Using the information $N_i(f) = [f(\frac{\beta}{m_i+1}), f(\frac{2\beta}{m_i+1}), \dots, f(\frac{m_i\beta}{m_i+1})]$ we use the algorithm

$$U_i(f) = \frac{\beta}{m_i} \sum_{j_1=1}^{m_i} f\left(\frac{j_1\beta}{m_i+1}\right)$$

to approximate the integral. This coincides with the open trapezoidal quadrature. The rest of the properties of the approximation are explained afterwards.

First we take a quick look at the function

$$f_1(x, t) = \min(x, t),$$

which is in $\tilde{C}^0([0, 1])$. It is possible to calculate that the

$$\int_{[0, 1]^2} f_1(\vec{x}) d\vec{x} = \frac{1}{3}$$

holds. The real cost of the reaching of a specific error is compared with the estimated cost in the rear part of the section.

After this we take a look at some easy functions $\tilde{C}^{r-1}([0, 1])$, with $r > 0$.

Here the one dimensional function we chose is

$$f_r(x) = \left(\prod_{k=1}^{\lfloor r/2 \rfloor} \frac{2k-1}{2k} \right) \cdot \sin^r(2\pi \cdot x),$$

which especially is in $\tilde{C}^{r-1}([0,1])$, with $f(0) = f(1) = 1$. The result of the integral $\int_0^1 \prod_{k=1}^2 \sin^2(2\pi kx) dx$ could be determined analytically to be 1 for even r and 0 for uneven r and $d \in \mathbb{N}$.

```
[1]: import os
os.chdir("../")

import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
↪ one
import matplotlib.pyplot as plt
import pandas as bearcats
import scipy
import sympy as sp
import numpy as np
import itertools as itt
import math
from sklearn import datasets, linear_model
plt.rc('font', size=14, weight="bold")
```

5.1 Example on page 358

First we show the calculate the approximation of $f_1(x, t)$ and the approximation given on page 358.

```
[11]: from IPython.display import display, Math

# other properties of the example
r = 0
beta = 1
q = 2
dim = 2
m_i = 2 ** q - 1

# Specific error properties of periodic functions of this kind
C_r = np.sqrt(abs(scipy.special.bernoulli(2+2*r)[-1])/math.factorial(2*r+2))
e_one_a = C_r * beta ** ((2*r+3)/2)
e_one_b = 2 ** (-(r + 1))

# Cost and error esimation for the one dimensional error
B, C, D, F_0, F = [e_one_a, e_one_a, e_one_b, 1, 2]

# calculation of the approximation and the error and cost estimation
# of the example
option = "Trapezoidal"
deg_approx = 18
def max_fun(x, t):
    # implementation of the function given on page 358 for input like used in
    # implementation of algorithm.
    return np.min(np.array([x,t]), axis=0)

variables = []
for f in range(dim):
    variables.append(sp.Symbol("x_" + str(f)))
```

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```
result_max_fun = np.empty((deg_approx,2))
cost_and_err_estim = np.empty((20,2))
for eps_ind, epsilon in enumerate([(1/2) ** (i+3) for i in range(20)]):
    m_i = 2 ** q - 1
    cost_and_err_estim[eps_ind,:] = Studieproject_one.fast_eps_cost(dim, B, C, D, F_0,
    ↪ F, epsilon)[0::6]

for q in range(dim, dim + deg_approx):
    result_max_fun[q-2, :] = Studieproject_one.controller_smolyak(max_fun, variables,
    ↪ option, q, function_given=True, no_error=True, example_2=True)[0:3:2]

result_max_fun[:,0] = abs(result_max_fun[:,0] -1/3)
```

We here want to show that the function calculating the estimation works properly.

```
[12]: estim_cost_error = Studieproject_one.fast_eps_cost(dim, B, C, D, F_0, F, 1/8)
meaning = [r"cost(A_{\epsilon}(d))", r"\alpha", r"\alpha_0", r"\alpha_1", r"\alpha_2",
    ↪ r"\alpha_3"]

display(Math(r" $We chose $\epsilon$"+r"$ = $ \frac{1}{8}$"))
print("This leads to:")
for index, var in enumerate(meaning):
    display(Math(var+"$ = $" +str(round(estim_cost_error[index],4))))
    print("")
display(Math("$This coincides with the values displayed on the pages 368 and 369. For
    ↪ bigger values of $ \epsilon$, $ the cost-estimation is not a number. In this
    ↪ reagon the estimation will be seen to be bigger than actual values.$ "))
```

We chose $\epsilon = \frac{1}{8}$

This leads to:

$cost(A_{\epsilon}(d)) = 0.4141$

$\alpha = 1.0$

$\alpha_0 = 3.3902$

$\alpha_1 = -0.5762$

$\alpha_2 = 2.7098$

$\alpha_3 = 1.5$

This coincides with the values displayed on the pages 368 and 369. For bigger values of ϵ , the cost-estimation is not a number. In this reagon the estimation will be seen to be bigger than actual values.

```
[36]: fig = plt.figure(figsize=(15,6))
plt.plot( abs(result_max_fun[:,0]), result_max_fun[:,1], color="red", label=
    ↪ "calculations")
regression_data_x = [[k_1] for k_1 in np.log(abs(result_max_fun[-5:,0]))]
```

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```

regression_data_y = [[k_2] for k_2 in np.log(abs(result_max_fun[-5:,1]))]
regr = linear_model.LinearRegression()
regr.fit(regression_data_x , regression_data_y)
regr.score(regression_data_x , regression_data_y)
exponent = regr.coef_

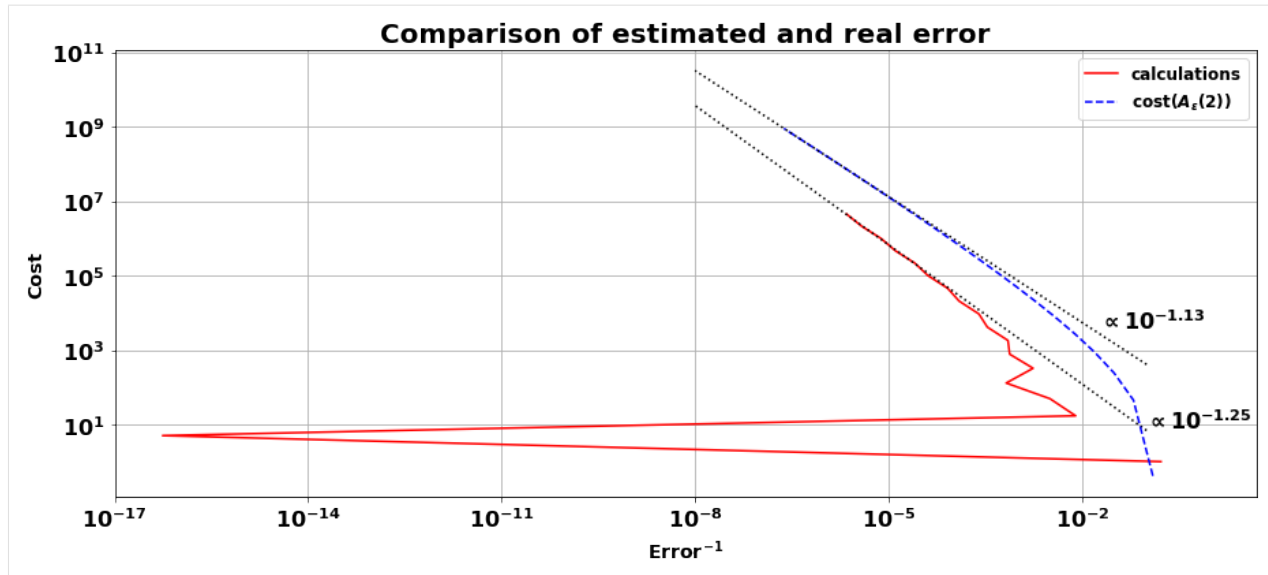
regression_y = [np.e**(regr.intercept_ + exponent[0][0] * np.log(x)) for x in np.
↳ linspace(10**(-1),10**(-8),30)]
plt.plot(np.linspace(10**(-1),10**(-8),30),regression_y, "k:")
plt.annotate("$\propto 10^{-1.25}$", (np.linspace(10**(-1),10**(-8),30)[2],regression_
↳ y[1]),xytext=(np.linspace(10**(-1),10**(-8),30)[2],regression_y[1]*1.1), fontweight=
↳ "bold",fontsize=16)

plt.plot( [(1/2) ** (i+3) for i in range(20)], cost_and_err_estim[:,0], color="blue",
↳ linestyle="--",label="$A_{\epsilon}(2)$")
regression_data_x = [[k_1] for k_1 in np.log10([(2) ** (-i-3) for i in range(15,20)])]
regression_data_y = [[k_2] for k_2 in np.log10(cost_and_err_estim[-5:,0])]
regr = linear_model.LinearRegression()
regr.fit(regression_data_x , regression_data_y)
regr.score(regression_data_x , regression_data_y)
exponent = regr.coef_

regression_y = [10**(regr.intercept_ + exponent[0][0] * np.log10(x)) for x in np.
↳ linspace(10**(-1),10**(-8),30)]
plt.plot(np.linspace(10**(-1),10**(-8),30),regression_y, "k:")
plt.annotate("$\propto 10^{-1.13}$", (np.linspace(10**(-1),10**(-8),30)[25],
↳ regression_y[25]),xytext=(np.linspace(10**(-1),10**(-8),30)[25]*1.3,regression_
↳ y[25]), fontweight="bold",fontsize=16)

plt.xlim(10**(-17), 5)
plt.yscale("log")
plt.xscale("log")
plt.grid(axis="both")
plt.legend(fontsize=12)
plt.ylabel("Cost", fontsize = 14,fontweight = "bold")
plt.xlabel(r"Error$^{-1}$",fontsize = 14,fontweight = "bold")
plt.yticks(fontsize=16,fontweight="bold")
plt.xticks(fontsize=16,fontweight="bold")
plt.title("Comparison of estimated and real error",
          fontsize = 20, fontweight = "bold")
plt.show()

```



The error scales in a surprisingly nice way, such that the error bound holds for the function $f_1(\vec{x})$ for almost every point. We once again remind the reader that the error bound could not be applied to the calculation without changing the measure used in this project.

5.2 Example 2: Functions in $f \in \tilde{C}^r([0, 1])$ with $r > 0$ and higher dimensions

[6]: # We from now on use l instead of r to make clear that the so far the results of \tilde{C}^r
 ↳ section 15.2.6 could not be applied.

```
beta = 1
```

```
try:
    os.chdir("Data")
except:
    pass
```

```
color_str = ["navy", "crimson", "forestgreen", "cornflowerblue", "darkorange"]
```

```
# Structure of data: [r, dim, q, :]
result_per_func = bearcats.read_pickle("approx_per_function.pkl")
result_per_func = result_per_func['results']#[:-1, :, :, :]
```

```
fig, ax = plt.subplots(result_per_func.shape[0], figsize=(15, 25))
```

```
for l in range(result_per_func.shape[0]):
    result_per_func_l = result_per_func[l, :, :, :]
    l_real = 0
```

```
''' These calculations are not made because the estimations are not met.
    In further implementation it might be possible to use them
```

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```

# Specific error properties of periodic functions of this kind
C_r = np.sqrt(abs(scipy.special.bernoulli(2+2*r_real)[-1])/math.factorial(2*r_
↪real+2))
e_one_a = C_r * beta ** ((2*r_real+3)/2)
e_one_b = 2 ** (-(r_real + 1))

# Cost and error esimation for the one dimensional error
B, C, D, F_0, F = [e_one_a, e_one_a, e_one_b, 1, 2]
'''

for dim in range(result_per_func_l.shape[0]):
    dim_real = dim+2
    # We want to take a look at the error, not the value of the approximation.
    if l%2==0:
        error = abs(result_per_func_l[dim,:,0]-1)
    else:
        error = abs(result_per_func_l[dim,:,0])+np.max(abs(result_per_func_l[:, :,
↪0]))/1000

    cost = result_per_func_l[dim,:,2]

    '''
    cost_estim = []
    epsilons = [(1/2)**(20+l+k_1) for k_1 in range(error.shape[0])]
    for eps in epsilons:
        cost_estim.append(Studieproject_one.fast_eps_cost(dim_real, B, C, D, F_0,
↪F, eps)[0])
    '''

    plt.subplots_adjust(hspace=0.4, wspace=0.4)
    ax[l].plot(cost, error, label="dimension d = "+ str(dim+2))

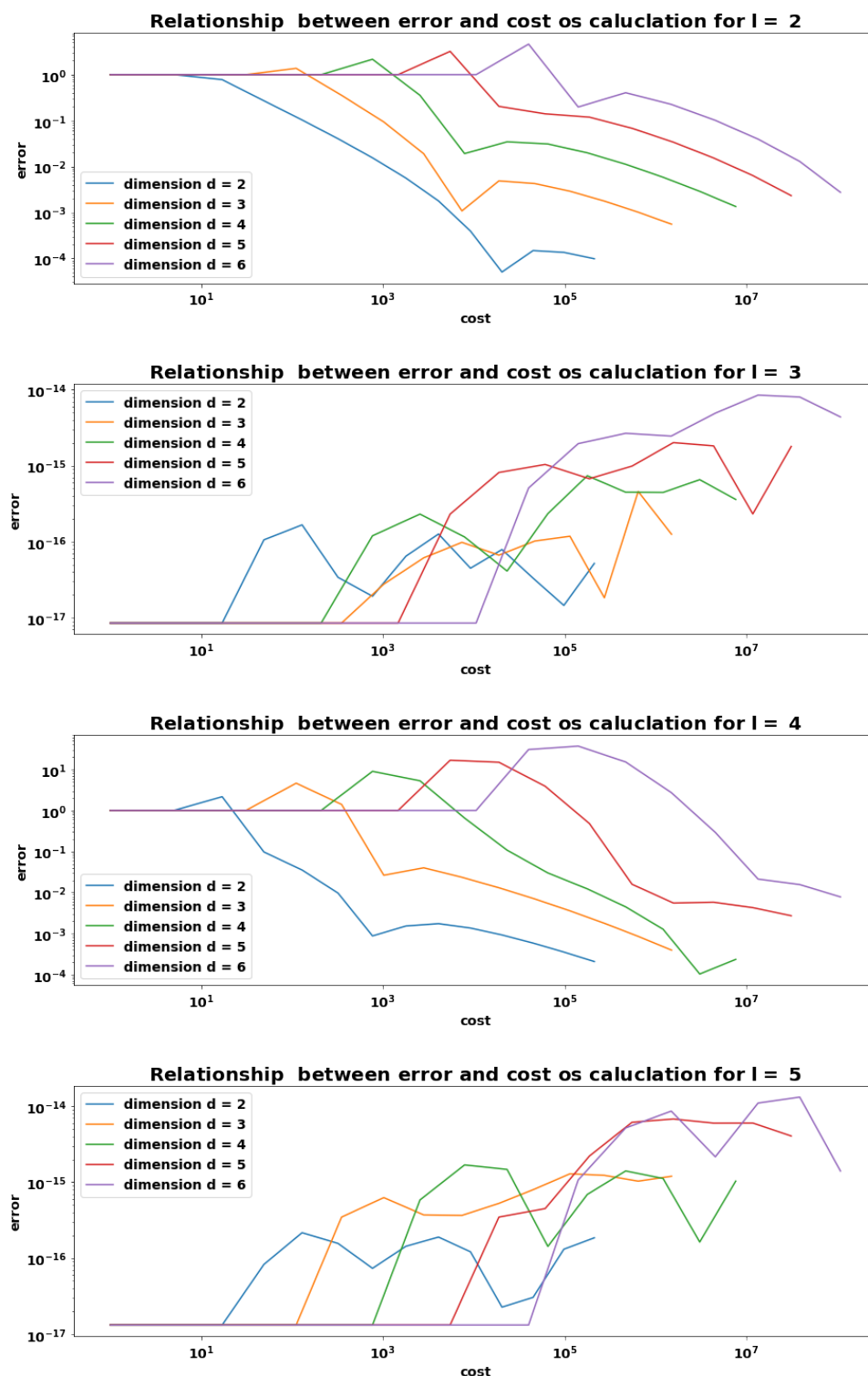
    ax[l].set_xscale("log")
    ax[l].set_yscale("log")
    ax[l].legend()
    ax[l].set_title(r"Relationship between error and cost os caluclation for $\mathbf{f}$
↪{l} = $ " + str(l+2), fontsize=20, fontweight="bold")
    ax[l].set_xlabel("cost", fontsize=14, fontweight="bold")
    ax[l].set_ylabel("error", fontsize=14, fontweight="bold")

fig.suptitle("Error for approximation of scalar in different dimensions", fontsize=24,
↪fontweight="bold")

plt.show()

```

Error for approximation of scalar in different dimensions



For l odd the results were shifted, because for small q the error is 0. Because of the symmetry of the function and the points used for the calculation the only source of an inexact approximation is the machine error.

For l even it could easily be seen that the results of section 15.2.6 could not be applied, because the approximation would need to have the tendency to get more precise with increasing r . This could not be observed.

```
[2]: '''
option = "Trapezoidal"
deg_approx = 15
max_dim = 6
r_max = 5

results_example_2= np.empty((r_max-1 , max_dim-1, deg_approx, 3))
for r in range(2, r_max+1):
    print(" ")
    norm_factor = 1
    variables = ["(x_1)"]
    for k in range(1,int(np.ceil(r/2))+1):
        norm_factor = norm_factor * (2*k/(2*k-1))

    functions = [str(norm_factor) + " sin(x_"+str(1)+"*2*pi)** "+str(r)]

    for dim in range(2,max_dim+1):

        variables.append(variables[-1][:-1]+", x_"+ str(dim)+")")
        functions.append(functions[-1]+ " * " + str(norm_factor) + "* sin(x_
→"+str(dim)+"*2*pi)** " + str(r))

        for q in range(dim,dim+deg_approx):
            print(str(r) + " " +str(dim)+ " " + str(q))
            results_example_2[r-2,dim-2, q - dim, :] = Studieproject_one.controller_
→smolyak(functions[-1], variables[-1], option, q, example_2=True)

results_example_2 = { "results": results_example_2}

bearcats.to_pickle(results_example_2,"approx_per_function.pkl")

print("If you want to generate data, remove quotation marks.")
'''
print("Remove quotes to generate data.")

Remove quotes to generate data.
```

```
[ ]:
```

EXAMPLE 15.3.8 PERTUBATED COULOMB POTENTIAL (PCP)

We want to implement the example 15.3.8 in the book Tractability of Multivariate Problems Vol 2 for 2 particles The function itself is

$$f(x_1, x_2, \dots, x_l) = \sum_{1 \leq i < j \leq l} \left(\frac{1}{\sqrt{\|x_i - x_j\|_2^2 + \alpha}} \right)$$

Where x_i , $i = 1, 2, \dots, l$ are three dimensional vectors. We first choose $l = 2$. Using the results of the example we calculate the function by integrating

$$\tilde{f}_\alpha(z_1, z_2, \dots, z_{3l}) = \left(\prod_{j=1}^{6-1} \left(j + \frac{1}{2} \right) \right) \cdot \frac{(z_1 - z_4)^4 (z_2 - z_5)^4 (z_3 - z_6)^4}{((z_1 - z_4)^2 + (z_2 - z_5)^2 + (z_3 - z_6)^2 + \alpha)^{13}}$$

over the interval $[0, 1]^6$. For higher dimension the algorithm gets extremely slow.

```
[3]: import matplotlib.pyplot as plt
import matplotlib.colors as colors
import time as time
import numpy as np
import pandas as bearcats
import scipy.integrate
import sympy

import os
os.chdir("../")
import Methodes_Studienprojekt.Smolyak_qmc_one_point as Studieproject_
↪ one
import Methodes_Studienprojekt.Smolyak_qmc_three_points as _
↪ Studienprojekt_three
```

6.1 Results of approximations

We calculated the integral for $\alpha = 10^{-i}$ for $i = 0, \dots, 5$ using different quadratures up to a degree of approximation of up to 23, depending on the runtime of the algorithm. Furthermore it was possible to calculate the integral for $i = 0$ and $i = 1$ in an acceptable amount of time and with a sufficient accuracy. On this basis we also show a lower bound of the integral for the smaller α which was estimated on page 393 of the same book. First we show the results of the approximation.

```
[26]: try:
        os.chdir("Data")
    except:
        pass
    # Settings needed for plotting of results.
    option_list = ["Newton-Cotes",
                  "Trapezoidal",
                  "Monte Carlo (non-nested)",
                  "Quasi-Monte Carlo"]
    color_str = [ "fuchsia", "navy", "limegreen", "red"]
    alpha = [1, 0.1, 0.01, 0.001, 0.0001, 0.00001]

    # Load the result the integration of the function calculated by scipy methode nquad
    result_scipy_calc_basic = bearcats.read_pickle("integral_Coulomb_alpha_
    ↪"+str(alpha[0])+".pkl")

    for k_1 in range(len(alpha)):
        # Load single approximation of the PCP different alphas and a wider range of
        ↪degree of approximation q.
        data = bearcats.read_pickle("test_higher_dim_approx_alpha_"+ str(alpha[k_1]) +".
        ↪pkl")

        # Integral could only be calculated relative exactly by the nquad for the first
        ↪2 alphas.
        if k_1 < 2:
            result_scipy_calc = bearcats.read_pickle("integral_Coulomb_alpha_
            ↪"+str(alpha[k_1])+".pkl")
            # For alpha < 1 the PCP was approximated using the trapezoidal quadrature and q =
            ↪19. Higher approximations
            # would easily have taken about 5 hours.
            if k_1 > 0:
                single_approx_trap = bearcats.read_pickle("test_higher_dim_approx_one_alpha_"
                ↪+ str(alpha[k_1]) + "_only_trap.pkl").Trapezoidal[0]

            fig = plt.figure(figsize=(15,12))
            plt.subplots_adjust(hspace=0.4, wspace=0.4)

            # First figure showing the absolut value of the approx in a logarithmical scale
            plt.subplot(2,1,1)

            for k_2 in range(len(option_list)):

                approx = data[option_list[k_2]][0][:17,0]
                error = data[option_list[k_2]][0][:17,1]

                end_of_approx = np.where(approx==0)[0][0]

                if end_of_approx == 0:
                    end_of_approx = np.where(approx==0)[0][-1]+1

                if k_2 < 2:
                    plt.plot(list(range(6,6+end_of_approx)), abs(approx[:end_of_approx]),
                    ↪color = color_str[k_2], label=option_list[k_2])

                else:
```

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```

plt.plot(list(range(6+end_of_approx,6+17)), abs(approx[end_of_approx:]),
color = color_str[k_2], label=option_list[k_2])

if k_1 > 0:
    plt.plot(6+13, abs(single_approx_trap[0]), color="navy", marker="o")

if k_1 < 2:
    plt.hlines(result_scipy_calc[0], xmin = 5.5, xmax= 6.5+17, linestyle="--",
color= "black", label="exact result")

if k_1 > 0:
    plt.hlines(result_scipy_calc_basic[0]*6*(0.1**(-2))**int(np.log10(alpha[0]/
alpha[k_1])), xmin = 5.5, xmax= 6.5+17, linestyle=":", color= "orange", label=
"lower bound of result")

plt.xlim(5.5,6.5+17)
plt.ylim(np.min(abs(data[option_list[1]][0][np.where(data[option_list[1]][0][:,0]!
=0)][:,0]))
/100,np.max(data[option_list[2]][0][:,0])*10)
plt.legend(loc=4, prop={"size":14, "weight":"bold"})
plt.ylabel("result of approximation",fontsize = 14,fontweight = "bold")
plt.xlabel("degree of approximaition q",fontsize = 14,fontweight = "bold")
plt.yscale("log")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.title("Absolut value of approximation" , fontsize=18,fontweight="bold")

# Second subplot showing the absolut values.
plt.subplot(2,1,2)

for k_2 in range(1, len(option_list)):
    approx = data[option_list[k_2]][0][:17,0]
    error = data[option_list[k_2]][0][:17,1]
    end_of_approx = np.where(approx==0)[0][0]

    if end_of_approx ==0:
        end_of_approx = np.where(approx==0)[0][-1]+1

    if k_2 < 2:
        plt.plot(list(range(6,6+end_of_approx)), approx[:end_of_approx], color =
color_str[k_2], label=option_list[k_2])

    else:
        plt.plot(list(range(6+end_of_approx,6+17)), approx[end_of_approx:], color_
= color_str[k_2], label=option_list[k_2])

    if k_1 > 0:
        plt.plot(6+13, single_approx_trap[0], color="navy", marker="o")

plt.xlim(5.5,6.5+17)
plt.ylim(np.min([np.max(data[option_list[2]][0][np.where(data[option_list[2]][0][:
:17,0]!=0),0))- abs(
np.min(data[option_list[2]][0][np.where(data[option_list[2]][0][:17,0]!=0),
0])-np.max(data[option_list[2]][0][
np.where(data[option_list[2]][0][:17,0]!=0),0]))*1.3,0]),np.max([np.
max(data[option_list[2]][0][:17,0])*1.1,np.max(data[option_list[3]][0][:17,0])*1.
1]))

```

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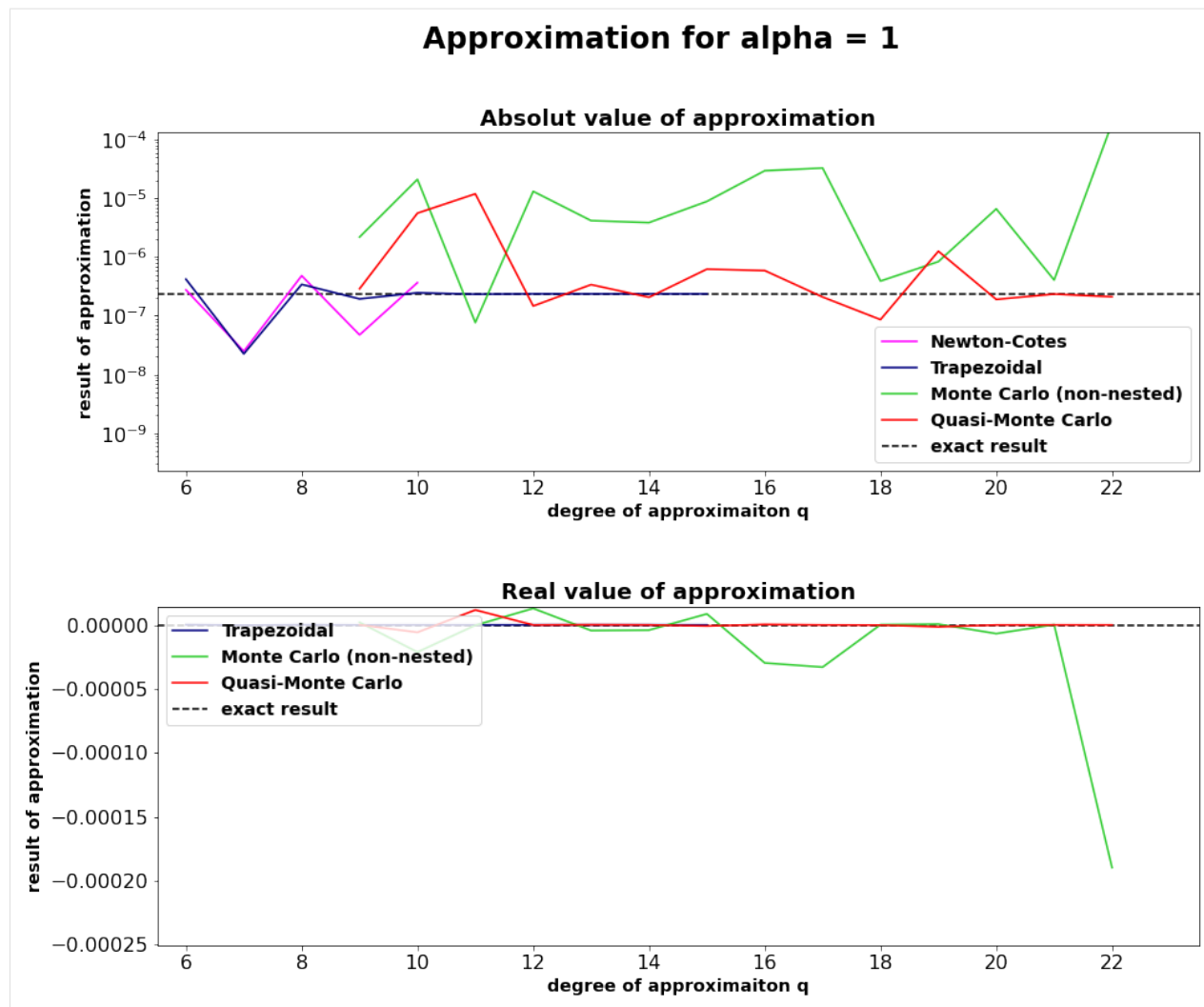
```
if k_1 < 2:
    plt.hlines(result_scipy_calc[0], xmin = 5.5, xmax= 6.5+17, linestyle="--",
    ↪color= "black", label="exact result")

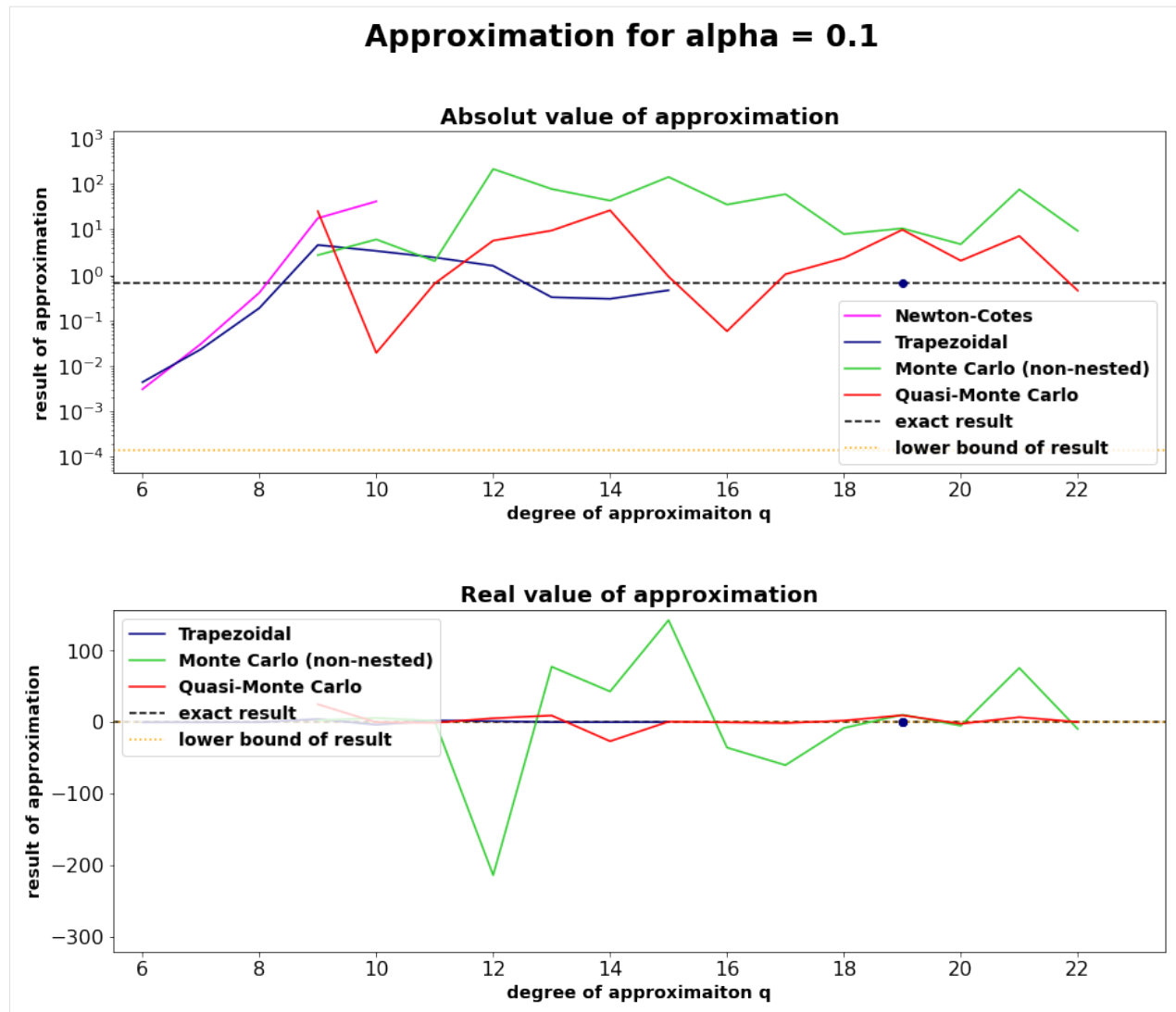
if k_1 > 0:
    plt.hlines(result_scipy_calc_basic[0]*6*(0.1**(-2))**int(np.log10(alpha[0]/
    ↪alpha[k_1])), xmin = 5.5, xmax= 6.5+17, linestyle=":", color= "orange", label=
    ↪"lower bound of result")

plt.legend(loc=2, prop={"size":14, "weight":"bold"})
plt.ylabel("result of approximation",fontsize = 14,fontweight = "bold")
plt.xlabel("degree of approximaiton q",fontsize = 14,fontweight = "bold")
plt.xticks(fontsize=16)
plt.yticks(fontsize=16)
plt.ticklabel_format(style="sci", axis="both")
plt.title("Real value of approximation", fontsize=18,fontweight="bold")

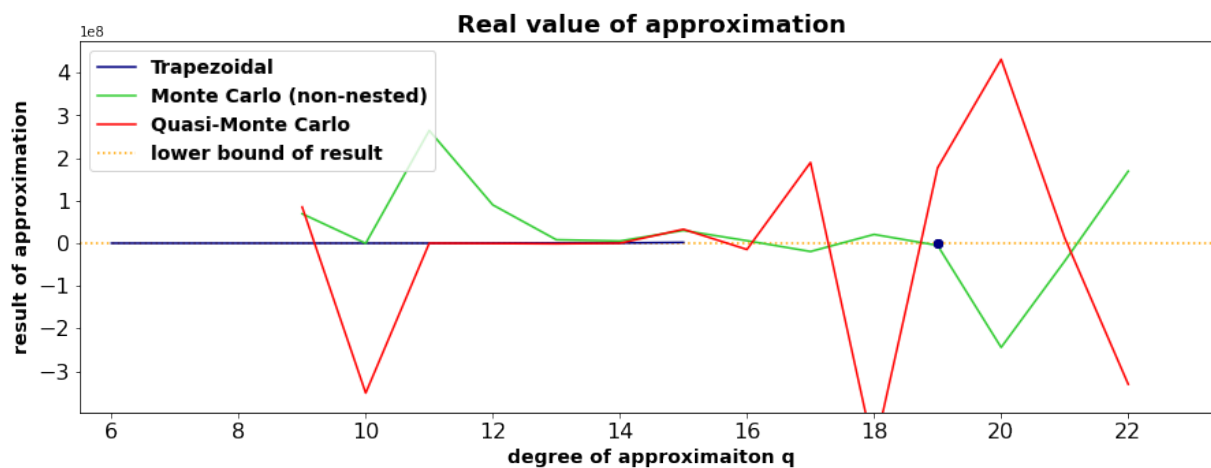
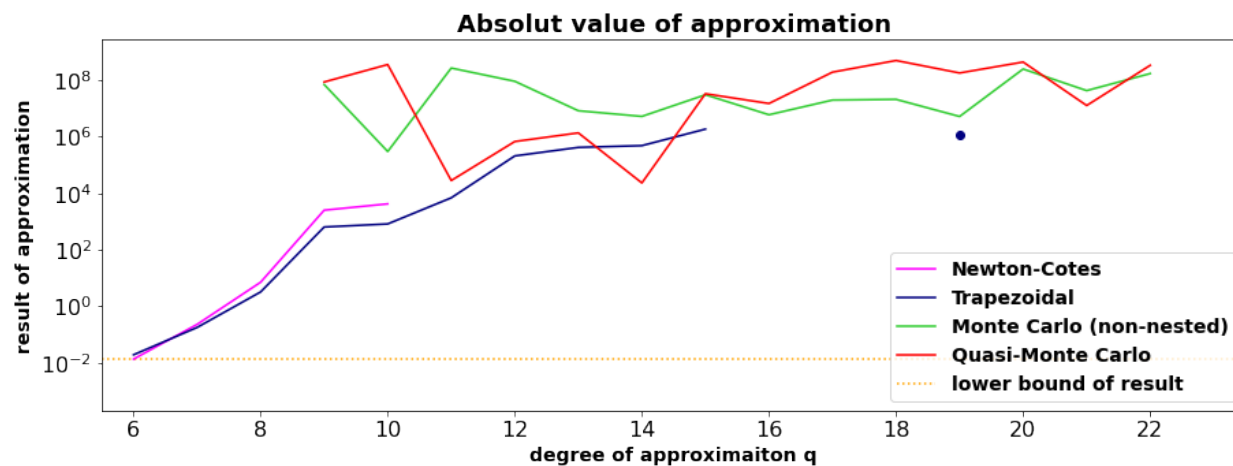
fig.suptitle("Approximation for alpha = " + str(alpha[k_1]), fontsize=24,
    ↪fontweight="bold")

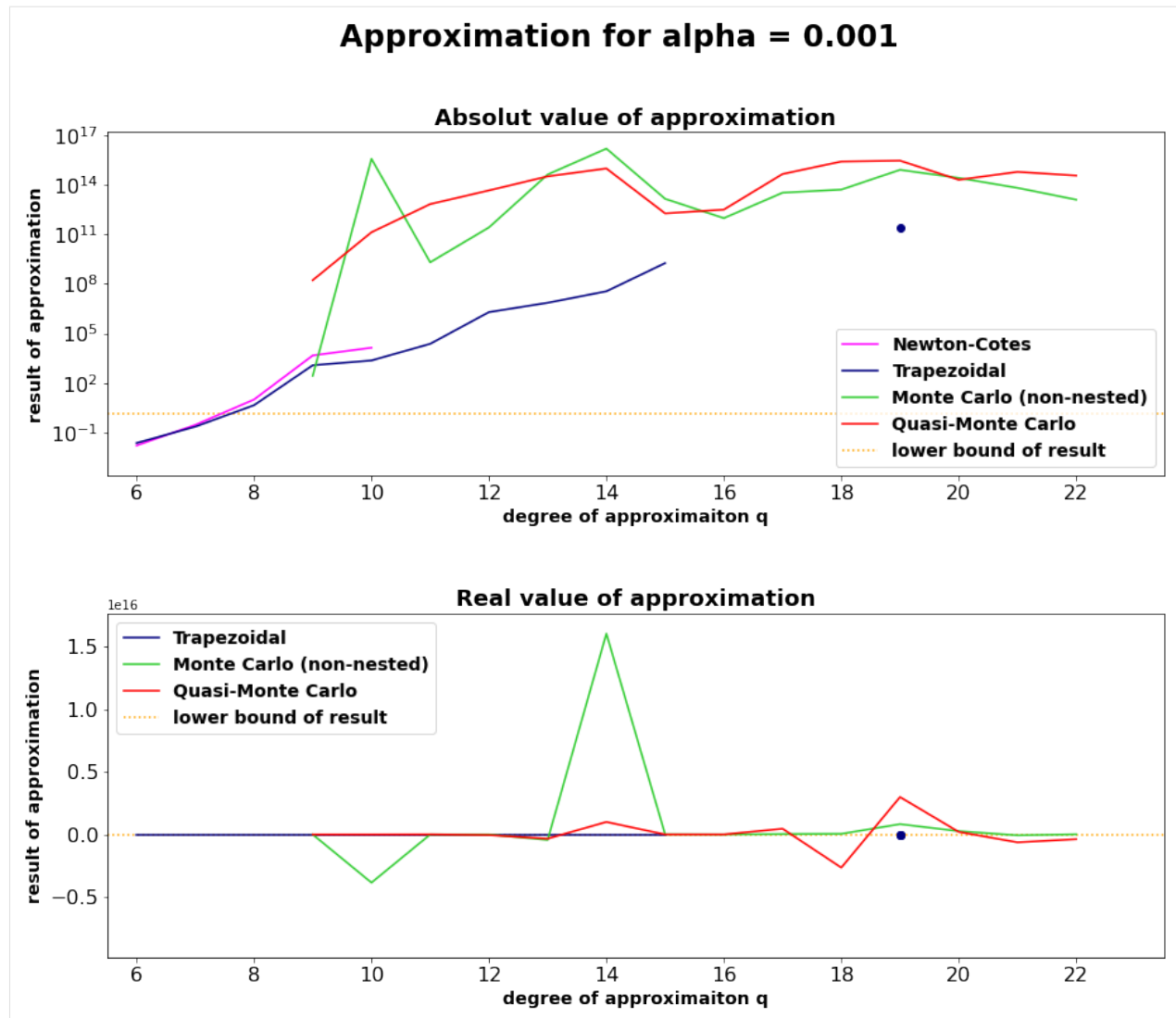
plt.show()
```



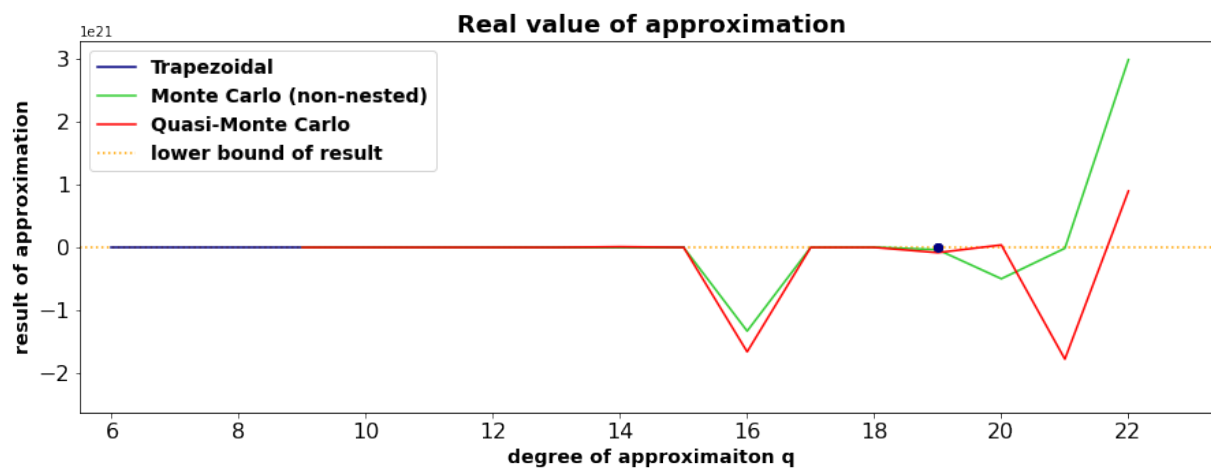
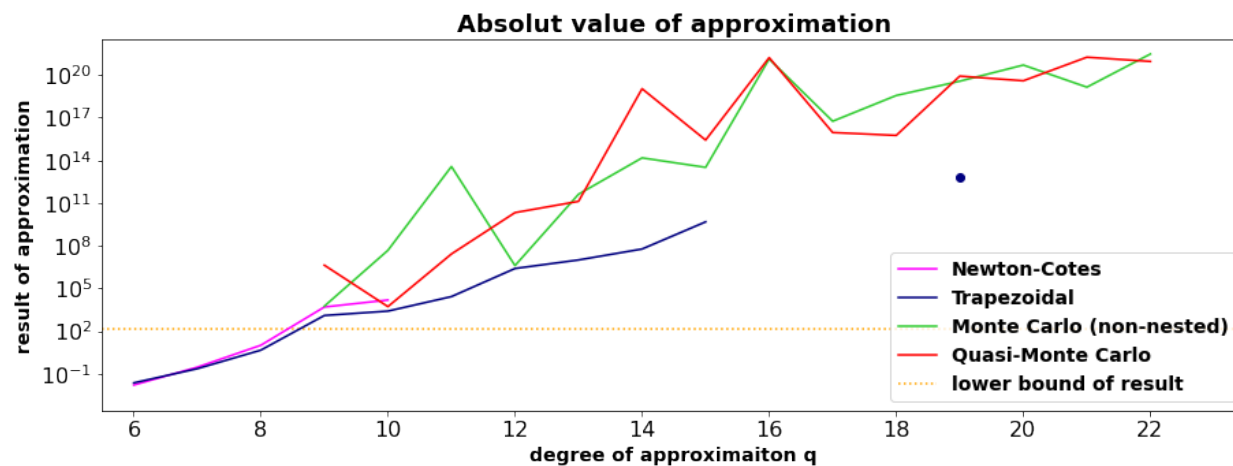


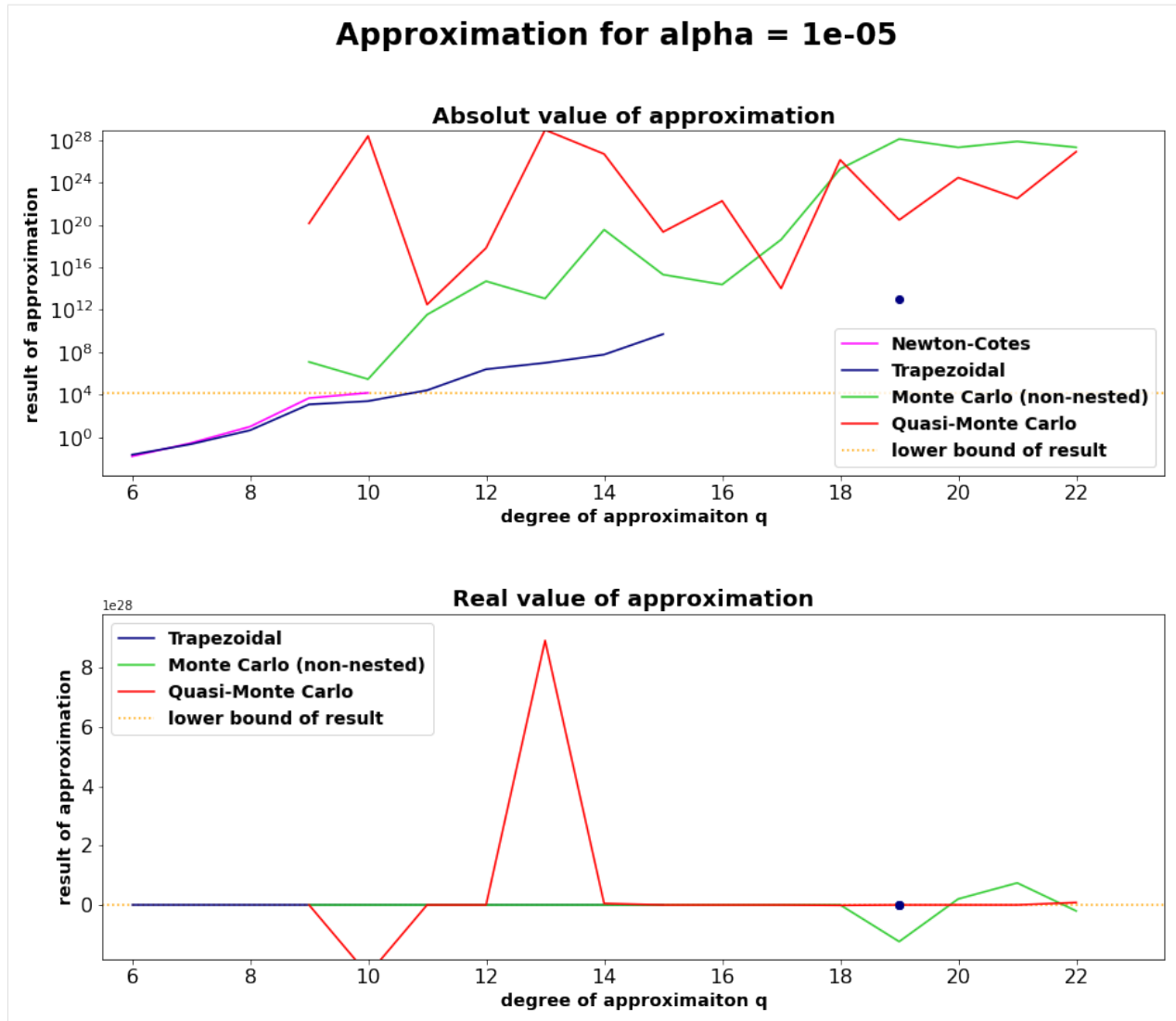
Approximation for alpha = 0.01





Approximation for $\alpha = 0.0001$





Often the deterministic quadratures the Smolyak algorithm seems approximate the integral for bigger α in a proper way. Never the less the error bounds are far away from being sufficiently small. In most cases these are bigger than the absolute value of the results. For $\alpha = 0.01$ even the approximation using the trapezoidal algorithm for $q = 19$ is negative.

Especially the result for the non-deterministic quadratures are varying by several orders of magnitude. Aaprt from this the error estimation for the non-deterministic quadratures are also varying by several magnitudes depending on whether the estimation of the variance includes points near the maximum of the function. In this case it is not possible to say, applying these has advantages compared to especially the trapezoidal quadrature.

It should be noted, that reducing the number of points used by the one dimensional quadratures for $q = 1$ for this function implies that for $q < 4$ the result of the approximation is 0.

For the sake of completeness, we show the error estimation of the different algorithms below.

6.2 Error estimation

```
[27]: plt.rc('font', size=14, weight="bold")
approx_simple_highdim_fct = bearcats.read_pickle("integral_over_one.pkl")

alpha = [1, 0.1, 0.01, 0.001, 0.0001, 0.00001]
fig, ax = plt.subplots(3,2,figsize=(18,10))
plt.subplots_adjust(hspace=0.4, wspace=0.4)

for k_1 in range(len(alpha)):
    data = bearcats.read_pickle("test_higher_dim_approx_alpha_"+ str(alpha[k_1]) + ".
    ↪pkl")
    if k_1 < 2:
        result_scipy_calc = bearcats.read_pickle("integral_Coulomb_alpha_
    ↪"+str(alpha[k_1])+".pkl")

    error = list()
    for k_2 in range(len(option_list)):
        error.append(data[option_list[k_2]][0][:17,1])
    error = np.vstack(error)

    if k_1 > 0:
        single_approx_trap = bearcats.read_pickle("test_higher_dim_approx_one_alpha_"
    ↪+ str(alpha[k_1]) + "_only_trap.pkl").Trapezoidal[0]
        error[1,13] = single_approx_trap[1]

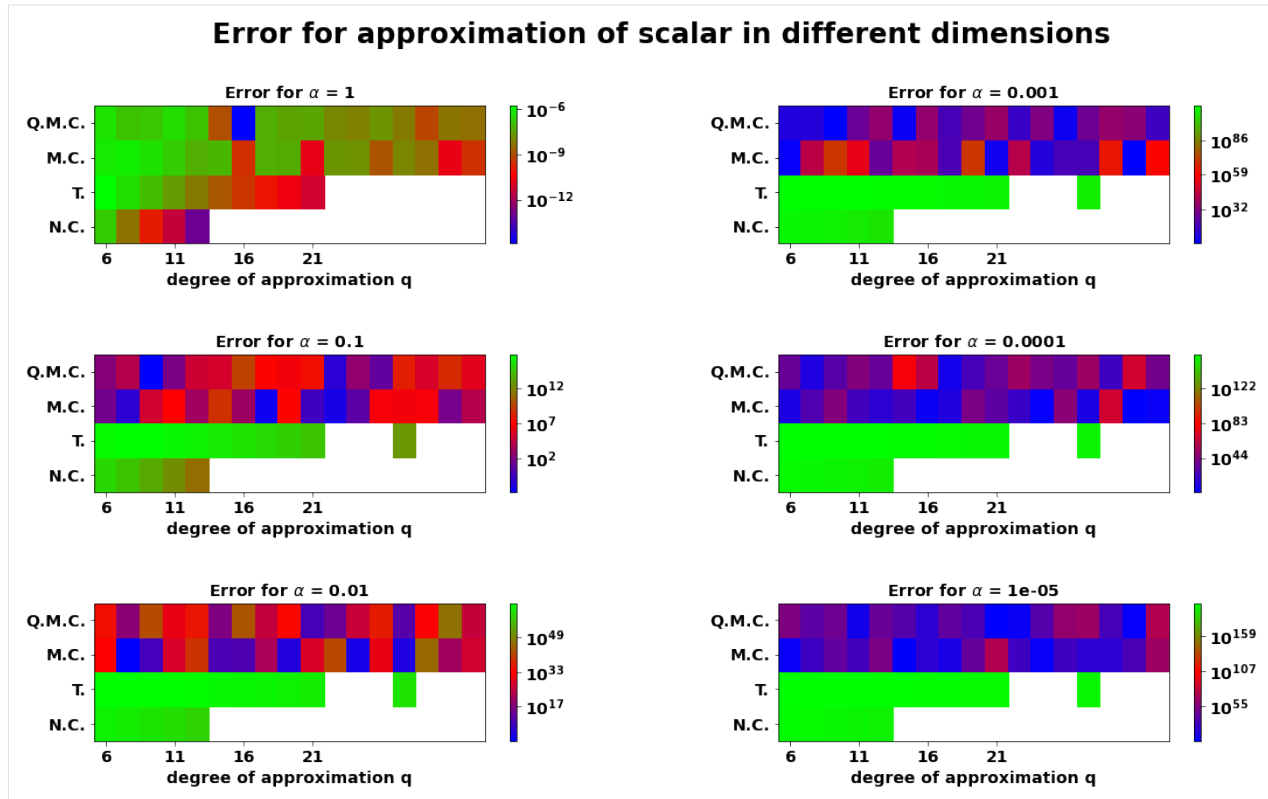
    plt.subplots_adjust(hspace=0.8, wspace=0.4)

    mat = ax[(k_1)%3, int((k_1)/3)].pcolor(error,norm=colors.LogNorm(vmin=error[error!
    ↪=0].min(), vmax=error.max()),
        cmap='brg')
    fig.colorbar(mat, ax=ax[(k_1)%3, int((k_1)/3)])

    ax[(k_1)%3, int((k_1)/3)].set_yticks([0.5,1.5,2.5,3.5])
    ax[(k_1)%3, int((k_1)/3)].set_yticklabels(["N.C.", "T.", "M.C.", "Q.M.C."],
    ↪fontsize=14, fontweight="bold")
    ax[(k_1)%3, int((k_1)/3)].set_xlabel("degree of approximation q",fontsize=14,
    ↪fontweight="bold")
    ax[(k_1)%3, int((k_1)/3)].set_xticks([0.5,3.5,6.5,9.5])
    ax[(k_1)%3, int((k_1)/3)].set_xticklabels([6,6+5,6+10,6+15],fontsize=14,
    ↪fontweight="bold")
    ax[(k_1)%3, int((k_1)/3)].set_title(r"Error for $\alpha$ = "+ str(alpha[k_1]),
    ↪fontsize=14, fontweight="bold")

fig.suptitle("Error for approximation of scalar in different dimensions\n",
    ↪fontsize=24,fontweight="bold")

plt.show()
```



6.3 Algorithm used for calculation of data

Below you see the algorithm, with which the integral was approximated.

```
[1]: '''
import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
    ↳ one
print("remove quotes, if you want to generate data.")
alpha = [1, 0.1, 0.01, 0.001, 0.0001, 0.00001]
dim = 6
option_list = ["Newton-Cotes",
               "Trapezoidal",
               "Monte Carlo (non-nested)",
               "Quasi-Monte Carlo"]

degree_of_approx = [5, 10, 17, 17, 17]
approx_result = np.zeros((max(degree_of_approx), len(option_list), 2))

for k_3 in range( len(alpha)):
    print("")
    print("alpha = "+ str(alpha[k_3]))
    print("")
    prefactor = np.sum(np.array(list(range(1,6)))+0.5)
    function_string = str(prefactor) + " * ((z_1 - z_4) ** 4 * (z_2 - z_5) ** 4 * (z_
    ↳ 3 - z_6)**4)/(((z_1 - z_4) ** 2 + (z_2 - z_5) ** 2 + (z_3 - z_6) ** 2 + " +_
    ↳ str(alpha[k_3])+ ") ** (13))"
    variables_string = "(z_1 , z_2 , z_3, z_4 , z_5, z_6)"
```

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```

    for k_1 in range(len(option_list)):
        print(option_list[k_1])
        for k_2 in range(degree_of_approx[k_1]):
            approx_result[k_2,k_1,:] = Studieproject_one.controller_smolyak(function_
↪string, variables_string, option_list[k_1], k_2 + 6)[0:2]
            print(approx_result[k_2,k_1,:])
            data = {"Newton-Cotes": [approx_result[:,0,:]], "Trapezoidal": [approx_result[:,1,:
↪]], "Monte Carlo (non-nested)": [approx_result[:,2,:]], "Quasi-Monte Carlo": [approx_
↪result[:,3,:]]}
            data = bearcats.DataFrame(data=data)
            bearcats.to_pickle(data, "test_higher_dim_approx_alpha_"+ str(alpha[k_3]) + ".pkl")
'''
'''
import scipy
import numpy as np
import pandas as bearcats
alpha = [1, 0.1, 0.01, 0.001, 0.0001, 0.00001]
dim = 6

approx_result = np.zeros((max(degree_of_approx), len(option_list), 2))

for k_3 in range(2, len(alpha)):

    prefactor = np.sum(np.array(list(range(1, 6))))+0.5)
    function_string = str(prefactor) + " * ((z_1 - z_4) ** 4 * (z_2 - z_5) ** 4 * (z_
↪3 - z_6)**4)/(((z_1 - z_4) ** 2 + (z_2 - z_5) ** 2 + (z_3 - z_6) ** 2 + " +
↪str(alpha[k_3]) + ") ** (13))"
    variables_string = "(z_1 , z_2 , z_3, z_4 , z_5, z_6)"
    f = Studieproject_one.rewrite_function(function_string, variables_string)[0]
    borders = [[0,1],[0,1],[0,1],[0,1],[0,1],[0,1]]
    integral = scipy.integrate.nquad(f, borders, opts = {"epsrel": 1/alpha[k_3]**2,
↪"epsabs": 3e4/(alpha[k_3])**2, "limit": 1 })
    print(integral)
    bearcats.to_pickle(integral, "integral_Coulomb_alpha_"+str(alpha[k_3])+".pkl")
'''
print("Remove quotes to approximate function either using the Smolyak algorithm or
↪the scipy.integral package in python.")

Remove quotes to approximate function either using the Smolyak algorithm or the scipy.
↪integral package in python.

```

[]:

VIEW

This notebook provides the possibility to use the methodes implemented by entering a function and a degree of approximation. Regarding the quadrature the option

- Newton-Cotes,
- Trapezoidal,
- Monte-Carlo (nested and not nested) and
- Quasi-Monte Carlo

exist. An other way is calling the View function.

```
[1]: # The packages needed here are Tkinter, because the View is implemented using it
# and of course the packages of the projekt
import tkinter as tk

import os
os.chdir("../")
import Methodes_Studienprojekt.Studienprojekt_Smolyak_qmc_one_point as Studieproject_
↳ one
```

```
[76]: from tkinter import messagebox as mbox
# View/ Main

# In this script we make an interface with tkinter
# The interface has: - An entry line for the function
#                   - An entry line for the variables
#                   - number of evaluations
#                   - Perhaps an entry line to verify dimension
#                   - button to start calculation
#                   - A separate popup for result showing of
#                   - Drop down menu for choosing of quadrature
#                   - Several popup for warnings
# import tkinter as tk

# own files imported
# import My_Controller as cont
# import controller_smolyak as cont

# create main viewer
root = tk.Tk()
root.title("Application for calculation of integral with Smolak algorithm")
root.geometry("800x300")
```

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```

# Label for description of functionality of viewer
description = tk.Label(root, text="To use the application you need first need to_
↳enter a function in the "
                        "entry.\n"
                        "After this the tuple of variables and in the end_
↳the quadrature you want to be "
                        "used.\n"
                        "The calculation starts after clicking start.",_
↳anchor="w", justify="left")
description.grid(row=0, column=0, sticky="w", columnspan=3)

#
# Column for entry of function
func_label = tk.Label(root, text="Please enter function: ", justify="center", anchor=
↳"w", pady=10, padx=10)
func_label.grid(row=1, column=0, sticky="w")

func_entry = tk.Entry(root, width=50, borderwidth=5)
func_entry.grid(row=1, column=1, sticky="e", padx=20, pady=20, columnspan=2)
func_entry.insert(0, "2 x y")

# Column for entry of variables
var_label = tk.Label(root, text="Please enter variables: ", justify="center", anchor=
↳"w", pady=10, padx=10)
var_label.grid(row=2, column=0, sticky="w")

var_entry = tk.Entry(root, width=50, borderwidth=5)
var_entry.grid(row=2, column=1, sticky="e", padx=20, pady=20, columnspan=2)
var_entry.insert(0, "(x , y)")

# Drop down menu for quadrature

# list of quadratures available
option_list = ["Newton-Cotes",
               "Trapezoidal",
               "Monte Carlo (nested)",
               "Monte Carlo (non-nested)",
               "Quasi-Monte Carlo"]
option = tk.StringVar()
option.set(option_list[0])
which_quad = tk.OptionMenu(root, option, *option_list)
which_quad.grid(row=3, column=2)

degree_label = tk.Label(root, text="Please choose degree of approximation: ", justify=
↳"center", anchor="w",
                        pady=10, padx=10)
degree_label.grid(row=3, column=0, sticky="w")

degree_of_approx = tk.Entry(root, width=10, borderwidth=5)
degree_of_approx.grid(row=3, column=1)
degree_of_approx.insert(0, 2)

# !! Here we also could think about a button making it possible to define the_
↳epsilon-error!!

```

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```

# Start button
def start():

    # read out entries
    function_string = func_entry.get()
    variables_string = var_entry.get()
    quadrature = option.get()
    degree = int(degree_of_approx.get())

    # We want to open an error box, if the degree of approximation is not big enough.
    ↪ Hence, we need to
    # know the dimension of the function. Here we get this by counting the commas.
    if degree < (variables_string.count(",")+1):
        mbox.showerror("Input error", "The degree of the approximation needs to be at
        ↪ least as high as the "
                                "dimension of m the function. \n\nIn this case
        ↪ this would be q = "
                                + str(variables_string.count(",") + 1) + ".")

    else:

        # ask controller for benchmarks of approximation
        result_approx, error, cost = Studieproject.controller_smolyak(function_string,
                                                                    variables_
        ↪ string, quadrature, degree)

        # Create new window where results are shown
        result_window = tk.Toplevel(root)
        result_window.title("Result")
        result_window.geometry("800x200")

        # Label for result of approx.
        result_label = tk.Label(result_window, text="The result of the approximation
        ↪ is: " + str(result_approx),
                                justify="left", anchor="e", pady=10, padx=10)
        result_label.grid(row=0, column=0, columnspan=2)

        # Label for error
        error_label = tk.Label(result_window, text="Estimated error: " + str(error),
        ↪ justify="center", anchor="w",
                                pady=10, padx=10)
        error_label.grid(row=1, column=0)

        # Label for cost estimation
        cost_label = tk.Label(result_window, text="Estimated number of evaluations: "
        ↪ + str(cost), justify="center",
                                anchor="w", pady=10, padx=10)
        cost_label.grid(row=1, column=2)

        # Quit button of result window
        quit_button_result = tk.Button(result_window, text="Quit", command=root.
        ↪ destroy, width=10, borderwidth=5,
                                padx=5, pady=5)
        quit_button_result.grid(row=2, column=2)

```

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```
# Start button
start_button = tk.Button(root, text="Start", command=start, width=10, borderwidth=5,
↳ padx=5, pady=5)

start_button.grid(row=4, column=0)

# Quit button
quit_button = tk.Button(root, text="Quit", command=root.destroy, width=10,
↳ borderwidth=5, padx=5, pady=5)
quit_button.grid(row=4, column=2)

root.mainloop()
```

7.1 Overview over methods defined for project

Furthermore, an overview should be given over the methods defined in the project. These methods are written in alphabetic order.

```
[80]: def write_functions():
    functions= [function for function in dir(Studieproject_one) if
        ((not function.startswith('__') and function.find("_") > 0 and
↳ function != "parse_expr") or function == 'smolyak')]
    for k_1 in range(len(functions)):
        print('\033[1m'+functions[k_1])
        print("")
    print('\033[0m')
write_functions()
```

```
calculate_stat_data
controller_smolyak
cost_smolyak
epsilon_cost
error_smolyak
find_all_sensible_combinations
find_next_tuples_index
find_right_position
implicit_multiplication
load_stat_data
modified_scatter_plot
monte_carlo_quad
newton_cotes
next_step
one_dim_newton_cotes
one_dim_trapezoidal
qmc_quad
rewrite_function
slice_find
smolyak
sort_tuples
standard_transformations
sum_tuple_shape
```


[]:

DOCUMENTATION

`Studienprojekt_Smolyak_qmc_one_point.View()`
View of project

Returns None

`Studienprojekt_Smolyak_qmc_one_point.calculate_stat_data` (*function_string_stat_calc:*
str, var_string: str, option_list_quad: str,
filename: str, degree_of_app_range:
list, repetitions=1024,
damping=True, damping_exp=1.5)

This method calculates and saves the result, error and runtime Smolyak approximation for the quadratures # given in option_list several times.

Parameters

- **function_string_stat_calc** (*string*) – function for which the integral should be approximated.
- **var_string** (*string*) – variables of function for which the integral should be approximated.
- **option_list_quad** (*list*) – list with the quadratures that should be used.
- **filename** (*string*) – name of the file in which you want to save the data
- **degree_of_app_range** (*list*) – list with degrees of approximation, for which data should be generated repetitions: Times the results are generated.
- **repetitions** (*int*) – Times the results are generated.
- **damping** (*boolean, optional*) – boolean saying, whether the number of repetition should decrease according to the degree of approximation. Depending on the range of degrees of approximation and the computer recommendable, default is True.
- **damping_exp** (*float*) – damping exponent of the degree of approximation, if damping is true.

Raises

- **Exception** – If one of the quadrature is not defined.
- **Exception** – If the filename is not valid
- **Exception** – If degree_of_app_range does not have the right length.

Returns None

```
Studienprojekt_Smolyak_qmc_one_point.controller_smolyak (function_string_control:
                                                         str,                vari-
                                                         ables_string_control:
                                                         str,  quadrature:    str,
                                                         q:    int, no_error=False,
                                                         function_given=False,
                                                         example_2=False,    a=0,
                                                         b=1)
```

Controller of project

Parameters

- **function_string_control** (*str*) – string in which the function is defined
- **variables_string_control** (*str*) – string for safe use, to be sure that right variables are used in function
- **quadrature** (*str*) – variable with 4 options to choose quadrature that should be used for approx
- **q** (*int*) – degree of approximation
- **no_error** (*boolean, optional*) – Decides, if error should be calculated. (Sometimes the time can be reduced relevantly.) Default is set to False.
- **function_given** (*boolean, optional*) – Needs to be set to true, if a callable function already is given. In this case the input function_string_control needs to be a callable function and variable_string_control need to be a list of sympy.variables, default set to False.
- **example_2** (*boolean, optional*) – Defined for example 2. In this case the quadrature is open, default set to False.
- **a** (*float, optional*) – lower limit of interval, default set to 0.
- **b** (*float, optional*) – upper limit of interval, default set to 1.

Raises Exception – If the type of quadrature chosen is not defined.

Returns result of approximation, upper bound of error of approximation and upper bound of cost.

Return type [float, float, float]

```
Studienprojekt_Smolyak_qmc_one_point.cost_smolyak (q:    int, dim:    int, nested=True,
                                                         is_rand=False,  example_2=False,
                                                         give_param=False)
```

Methode giving back an estimation of the cardinality of the information needed for the algorithm. Due to the definition of the sequences of the one dimensional algorithms used, the cost could be calculated relatively quickly

Parameters

- **q** (*int*) – Degree of approximation
- **dim** (*int*) – Dimension
- **nested** (*boolean, optional*) – Decides if information is nested, default True.
- **is_rand** (*boolean, optional*) – Decides if quadrature is probabilistic, default is False.
- **example_2** (*boolean, optional*) – Mainly defined for example 2. Decides, if the quadrature is open, default is False.
- **give_param** (*boolean, optional*) – Decides, if the cost estimation or the parameters are returned, default is False.

Returns estimated cost of approximation.

Return type float

```
Studienprojekt_Smolyak_qmc_one_point.epsilon_cost(function_string: str, variable_string: str, quadrature: str, epsilon: <module 'numbers' from '/usr/local/Cellar/python@3.9/3.9.0_5/Frameworks/Python.framework/Versions/3.9/Resources/Python.framework/Versions/3.9/Headers/numbers.h', nested=False, open=False)
```

Methode calculating the epsilon error :param function_string: string of function :type function_string: str :param variable_string: string of variables :type variable_string: str :param quadrature: string of one-dimensional quadrature to use :type quadrature: str :param epsilon: upper bound of error :type epsilon: float :param nested: Decides if the equation for nested or not nested information is used, default is :type nested: boolean, optional :param False.: :param open: Decides if a open or closed quadrature is used, default is :type open: boolean, optional :param False.:

Raises Exception – Exception: If the quadrature chosen is not implemented.

Returns The estimated error and cost estimation.

Return type [float, float]

```
Studienprojekt_Smolyak_qmc_one_point.error_smolyak(f, quadrature: str, variables, q: int, function_string_error: str, a=0, b=1, give_parameters=False)
```

Parameters

- **f** (*function*) – callable sympy function
- **quadrature** (*str*) – quadrature used
- **variables** (*sympy.variables*) – sympy variables of function
- **q** (*int*) – degree of approximation
- **function_string_error** (*str*) – string of function. (Needed, because analytically deriving strings is easier than
- **functions**) (*deriving*) –
- **a** (*float, optional*) – lower limit of interval, default to 0.
- **b** (*float, optional*) – upper limit of interval, default to 1.
- **give_parameters** (*boolean, optional*) – Boolean defined to get the parameters of the error estimation instead of
- **error estimation** (*the*) –
- **set to False.** (*default*) –

Raises

- **Exception** – If the quadrature chosen is not implemented
- **Exception** – If deterministic quadratures are used and the function can not be derived sufficiently many times.

Returns error of estimation

Return type float

Studienprojekt_Smolyak_qmc_one_point.**fast_eps_cost** (*dim: int, B: float, C: float, D: float, F_0: float, F: float, epsilon: float, q=0, is_nested=True*)

Function for the fast calc. of eps cost, if all parameters are given :param dim: dimension :type dim: int :param B: Estim. of norm of operator :type B: float :param C: Estim. of linear part of approx error :type C: float :param D: Estim. of exp. part of approx error :type D: float :param F_0: Estim. of linear part of costs :type F_0: float :param F: Estim. of exp. part of cost :type F: float :param epsilon: epsilon error chosen :type epsilon: float :param q: degree of approximation :type q: int, optional :param is_nested: Decides if the equation for nested or not nested information is used, default is :type is_nested: boolean, optional :param True.:

Returns The epsilon cost

Studienprojekt_Smolyak_qmc_one_point.**find_all_sensible_combinations** (*dim: int, q: int*)

Methode finding all tuples of combinations of one dim approximations necessary to calc. Smol. alg.. :param dim: dimension of vectors :type dim: int :param q: maximal cumulative sum :type q: int

Returns List of all combination of degrees of approximation needed for later calculations.

Return type list

Studienprojekt_Smolyak_qmc_one_point.**find_next_tuples_index** (*current_tuple: tuple, tuple_list: list*)

Method to find the function evaluation matrix with the most reusable function values

Parameters

- **current_tuple** (*tuple*) – Tuple saving the degree of approximation currently used for the approximation
- **tuple_list** (*list*) – list of tuples for which the evaluations are stored.

Returns The index of the evaluations with the most reusable evaluations.

Return type integer

Studienprojekt_Smolyak_qmc_one_point.**find_right_position** (*current_vector: numpy.array, pointer_current_position: int, max_value: int*)

Decides, where to add one. :param current vector: Current combination of degrees of approximation. :type current vector: np.array :param pointer_current_position: index of dimension, in which needs to be changed next. :type pointer_current_position: int :param max_value: upper bound of degrees of approximation needed in set. :type max_value: int

Returns highest position that needs to be changed in next step

Return type pointer_current_position (int)

Studienprojekt_Smolyak_qmc_one_point.**load_stat_data** (*filename: str*)

Method loading the formerly saved data to the memory. !! This method only should be used for the calculate_stat_data, because the data structure is relatively specific. :param filename: string of name of file that should be loaded.

Raises Exception – If the file name has the wrong format.

Returns Contains the results of the approximation.

Return type List

Studienprojekt_Smolyak_qmc_one_point.modified_scatter_plot(*x*, *y*, *title*='Gridpoints used for Smolyak-algorithm', *input_list*=False, *a*=0, *b*=1)

Methode quickly applying some basic modifications to a scatter plot of the gridpoints used for the Smolyak algorithm :param *x*: list of *x*-values of points :type *x*: list :param *y*: list of *y*-values of points :type *y*: list :param *title*: title of plot :type *title*: str :param *input_list*: Boolean saying, whether input is list. If True, the points in different

lists are plotted in different colours, default is False.

Parameters

- **a**(*float*, *optional*) – lower border of interval
- **b**(*float*, *optional*) – upper border of interval

Returns None

Studienprojekt_Smolyak_qmc_one_point.monte_carlo_quad(*q*: int, *a*=0, *b*=1)

Monte Carlo quadrature

This function computes the nodes and weights for the 1D Monte carlo quadrature.

Parameters

- **q**(*int*) – degree of approximation. Number of nodes, $m = 2^{(q-1)+1}$.
- **a**(*int*, *optional*) – Lower boundary of integral. Defaults to 0.
- **b**(*int*, *optional*) – Upper boundary of integral. Defaults to 1.

Raises

- **Exception** – *a* and *b* need to be real numbers.
- **Exception** – *b* as the upper boundary has to be bigger than *a*.
- **Exception** – *q* needs to be an integer.

Returns nodes and weights for the 1D Monte carlo quadrature of degree *q*.

Return type [array, array]

Studienprojekt_Smolyak_qmc_one_point.next_step(*current_vector*: numpy.array, *pointer_current_position*: int, *max_value*: int)

Methode returning, were and what to change in the vectors determining, what one dimensional degree of approximation should be applied. (Internal function)

Parameters

- **vector**(*current*) – full actual vector. (Could be reduced in most cases)
- **pointer_current_position**(*int*) – position currently worked at
- **max_value**(*int*) – necessary to decide when to skip to the next position

Returns The position at which the value needs to be incremented, the next value and a pointer showing, in which dimension the next action needs to be performed.

Return type [int, int, int]

`Studienprojekt_Smolyak_qmc_one_point.one_dim_newton_cotes(q: int, a=0, b=1)`

Newton-Cotes quadrature

This function computes the nodes and weights for the 1D Newton-Cotes quadrature. This is a program helping me getting into programming. On the internet better implementation of this very basic and not very precise quadrature. Apart from this, the quadrature will be used for the testing of the viewer at the start. !! Effectively we have the open Newton-cotes formula of the degree $(2^q - 1)!!$

Parameters

- **q**(*int*) – Degree of quadrature.
- **a**(*int*, *optional*) – Lower boundary of integral. Defaults to 0.
- **b**(*int*, *optional*) – Upper boundary of integral. Defaults to 1.

Raises

- **Exception** – a and b need to be real numbers.
- **Exception** – b as the upper boundary has to be bigger than a.
- **Exception** – q needs to be an integer.
- **Exception** – Please choose $q < 6$ for the Newton-Cotes approximation to avoid numerical errors.

Returns nodes and weights for the 1D Newton-Cotes quadrature of degree q.

Return type [array, array]

`Studienprojekt_Smolyak_qmc_one_point.one_dim_trapezoidal(q: int, a=0, b=1, example_2=False)`

Trapezoidal quadrature

This function computes the nodes and the weights for the 1D trapezoidal quadrature.

Parameters

- **q**(*int*) – Degree of quadrature. The number of points used for the quadrature then is 2^q
- **a**(*int*, *optional*) – Lower boundary of integral. Defaults to 0.
- **b**(*int*, *optional*) – Upper boundary of integral. Defaults to 1.
- **example_2**(*bool*, *optional*) – Boolean only defined for the example with the periodic, derivable functions. If this variable is true, the borders of the interval are not used.

Raises

- **Exception** – a and b need to be real numbers.
- **Exception** – b as the upper boundary has to be bigger than a.
- **Exception** – q needs to be an integer.

Returns nodes and the weights for the 1D trapezoidal quadrature of degree q.

Return type [array, array]

`Studienprojekt_Smolyak_qmc_one_point.qmc_quad(q: int, a=0, b=1)`

Quasi-Monte Carlo quadrature

This function computes the nodes and weights for the 1D Quasi Monte carlo quadrature using lattice rules.

Parameters

- **q**(*int*) – degree of approximation. Number of nodes, $m = 2^q$.

- **a**(*int, optional*) – Lower boundary of integral. Defaults to 0.
- **b**(*int, optional*) – Upper boundary of integral. Defaults to 1.

Raises

- **Exception** – a and b need to be real numbers.
- **Exception** – b as the upper boundary has to be bigger than a.
- **Exception** – q needs to be an integer.

Returns nodes and weights for the 1D QMC quadrature of degree q.

Return type [array, array]

Studienprojekt_Smolyak_qmc_one_point.**rewrite_function**(*function_string_rewrite: str, variables_string_rewrite: str, a=0, b=1*) → list

Parameters

- **function_string_rewrite**(*str*) – function to be approximated
- **variables_string_rewrite**(*str*) – variables used in function
- **a**(*float*) – lower border of interval
- **b**(*float*) – upper border of interval

Raises

- **Exception** – If no variable defined.
- **Exception** – If the set of variables used in function and defined in variable string do not coincide.

Returns

function and variable used for approximation. String rewritten with standardized variables.

Return type [sympy.function, sympy.variables, string]

Studienprojekt_Smolyak_qmc_one_point.**slice_find**(*current_shape: tuple, nearest_shape: tuple, open_quad=False*)

Method giving returning slices that make it possible to find old evaluations can be used for the calculation of the current summand and the points for that further evaluation are needed. :param current_shape: Tuple storing the number of evaluations used for the current approximation. :type current_shape: tuple :param nearest_shape: Tuple storing the number of evaluations used for a former evaluation :type nearest_shape: tuple :param open_quad: Only defined for example 2. If this is true, the borders of the interval are not

used, default to False.

Raises Exception – A unforeseen combination of values occurred.

Returns

Three d dimensional lists of slices are returned. In these the slice at the i-th position coincide with the points used in this dimension.

list1: Positions at which a new evaluation is needed. list2: Positions at which old evaluations can be used. list3: Positions of the weights multiplied with the old evaluations.

Return type [list1, list2, list3]

Studienprojekt_Smolyak_qmc_one_point.**smolyak** (*f, dim: int, weights: list, points: list, q: int, is_nested=False, example_2=False*) → list

The Smolyak algorithm

Method to compute an approximation of the integral of a function using the Smolyak algorithm.

Parameters

- **f** (*function*) – Callable sympy function.
- **dim** (*int*) – Dimension of the function f.
- **weights** (*list*) – Numpy array with saved 1D weights corresponding to the points.
- **points** (*list*) – Points on which the function is evaluated for the 1D integral approx.
- **q** (*int*) – Degree of approximation.
- **is_nested** (*bool, optional*) – Boolean defined for the case that the information used is nested. Defaults to False.
- **example_2** (*bool, optional*) – Boolean only defined for the example with the periodic, derivable functions. If this variable is true, the borders of the interval are not used.

Raises

- **Exception** – The list of points and weights given for the Smolyak have incorrect lengths.
- **Exception** – The dimensions of the 1D gridpoints and weights do not match.
- **Exception** – Something went wrong with the usage of the nested information.
- **Exception** – Up to now the input vectors should be one dimensional but are not.

Returns result of approximation

Return type float

Studienprojekt_Smolyak_qmc_one_point.**sort_tuples** (*list_of_tuples: list, q: int*)

If we want to use the advantages of nested information, for this it is easier to first need to sort the list of tuples in a way, that we first evaluate the function in a systematical way. This sorting is done here. Starting with the first dimension we sort the list in a rising order.

This means: input = [(2,1,4),(1,2,2),(1,1,1),(2,1,2)] -> [(1,1,1),(1,2,2),(2,1,2),(2,1,4)]

Parameters

- **list_of_tuples** (*list*) – List of tuples that needs to be sorted in a specific order.
- **q** (*int*) – Degree of approximation

Returns Sorted list of tuples.

Return type list

Studienprojekt_Smolyak_qmc_one_point.**sum_tuple_shape** (*old_shape: tuple, new_shape: tuple*)

Function summing up the dimension of the old information used in the way necessary here. This means that if the dimension is the same, this value is saved. In other case, the values are summed up. Example: (33,3) + (32,3) = (65,3).

Parameters

- **old_shape** (*tuple*) – Dimension of the used information.
- **new_shape** (*tuple*) – Dimension of the used information.

Raises **Exception** – The input data needs to have the same lengths.

Returns Combined shape of information.

Return type tuple

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