

# A2

May 9, 2024

## 1 Computational Physics Blatt 04

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```
[ ]: import numpy as np
      from numpy.linalg import norm
      from functools import partial
      import matplotlib.pyplot as plt
      import matplotlib
      from scipy.optimize import curve_fit
      from scipy.stats import linregress

      matplotlib.rcParams["figure.figsize"] = [12, 8]
      %matplotlib inline

      T_MAX = 100 # Increased for nicer results
      H = 0.001 # decreased for nicer results
      SIGMA = 10
      B = 8/3
      R = 28
```

### 1.1 a)

Implementation of Runge-Kutta 5

```
[ ]: def RKDP(f, y0, t0, tmax, h, digits=None, **kwargs):
      dim = len(y0)
      coef = { # Coefficients for the Dormand-Prince method, padded with zeros
        ↪ for the 6th order
        "k2": np.array([1 / 5, 0, 0, 0, 0, 0]).reshape(-1, 1),
        "k3": np.array([3 / 40, 9 / 40, 0, 0, 0, 0]).reshape(-1, 1),
        "k4": np.array([44 / 45, -56 / 15, 32 / 9, 0, 0, 0]).reshape(-1, 1),
        "k5": np.array(
          [19372 / 6561, -25360 / 2187, 64448 / 6561, -212 / 729, 0, 0]
        ).reshape(-1, 1),
        "k6": np.array(
          [9017 / 3168, -355 / 33, 46732 / 5247, 49 / 176, -5103 / 18656, 0]
        ).reshape(-1, 1),
```

```

        -1, 1
    ), # called a6 in the literature
    "y": np.array(
        [35 / 384, 0, 500 / 1113, 125 / 192, -2187 / 6784, 11 / 84]
    ).reshape(
        -1, 1
    ), # called b in the literature
    "t": np.array(
        [0, 1 / 5, 3 / 10, 4 / 5, 8 / 9, 1]
    ), # called c in the literature
}

t = t0
y = y0
fp = partial(f, **kwargs) # set sigma, r and b
while t < tmax:
    ks = np.zeros((6, dim))
    ks[0] = h * fp(t, y)
    ks[1] = h * fp(t + h * coef["t"][1], y + np.sum(ks * coef["k2"]))
    ks[2] = h * fp(t + h * coef["t"][2], y + np.sum(ks * coef["k3"]))
    ks[3] = h * fp(t + h * coef["t"][3], y + np.sum(ks * coef["k4"]))
    ks[4] = h * fp(t + h * coef["t"][4], y + np.sum(ks * coef["k5"]))
    ks[5] = h * fp(t + h * coef["t"][5], y + np.sum(ks * coef["k6"]))
    y += np.sum(coef["y"] * ks, axis=0)
    if digits:
        y = np.round(y, digits)
    t += h
    yield t, *y

```

## 1.2 b)

Solve the Lorenz equation with  $y_0 = (2, 3, 14)$ , the parameters  $\sigma = 10$ ,  $b = \frac{3}{5}$  and  $r = 28$  in the time frame  $t_0 = 0$  to  $t_e = 50$  with a stepsize of  $h = 0.01$ . Plot the XZ-projection.

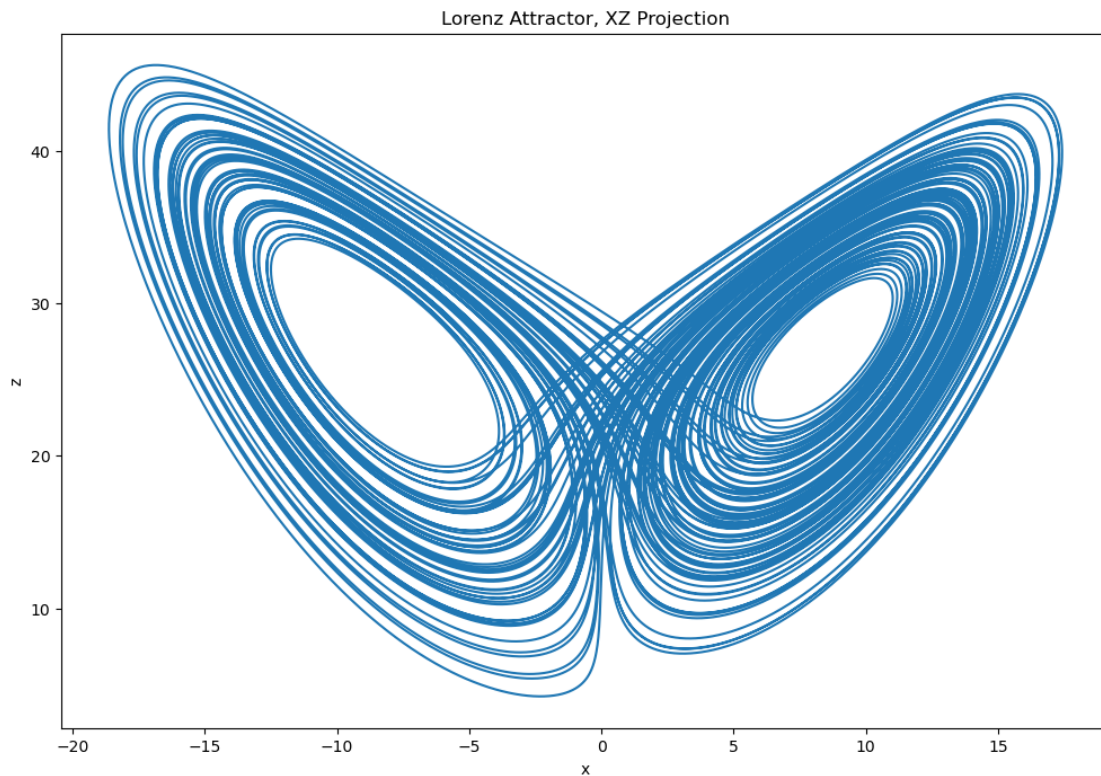
```

[ ]: def lorentz(t, y, sigma, r, b):
    x_dot = sigma * (y[1] - y[0])
    y_dot = y[0] * (r - y[2]) - y[1]
    z_dot = y[0] * y[1] - b * y[2]
    return np.array([x_dot, y_dot, z_dot])

[ ]: y0 = np.array([2, 3, 14], dtype=float) # prevent type-casting errors
rkdp = RKDP(lorentz, y0, t0=0, tmax=T_MAX, h=H, sigma=SIGMA, r=R, b=B)
system = np.array(list(rkdp))
time = system[:, 0]
trajectory = system[:, 1:]

```

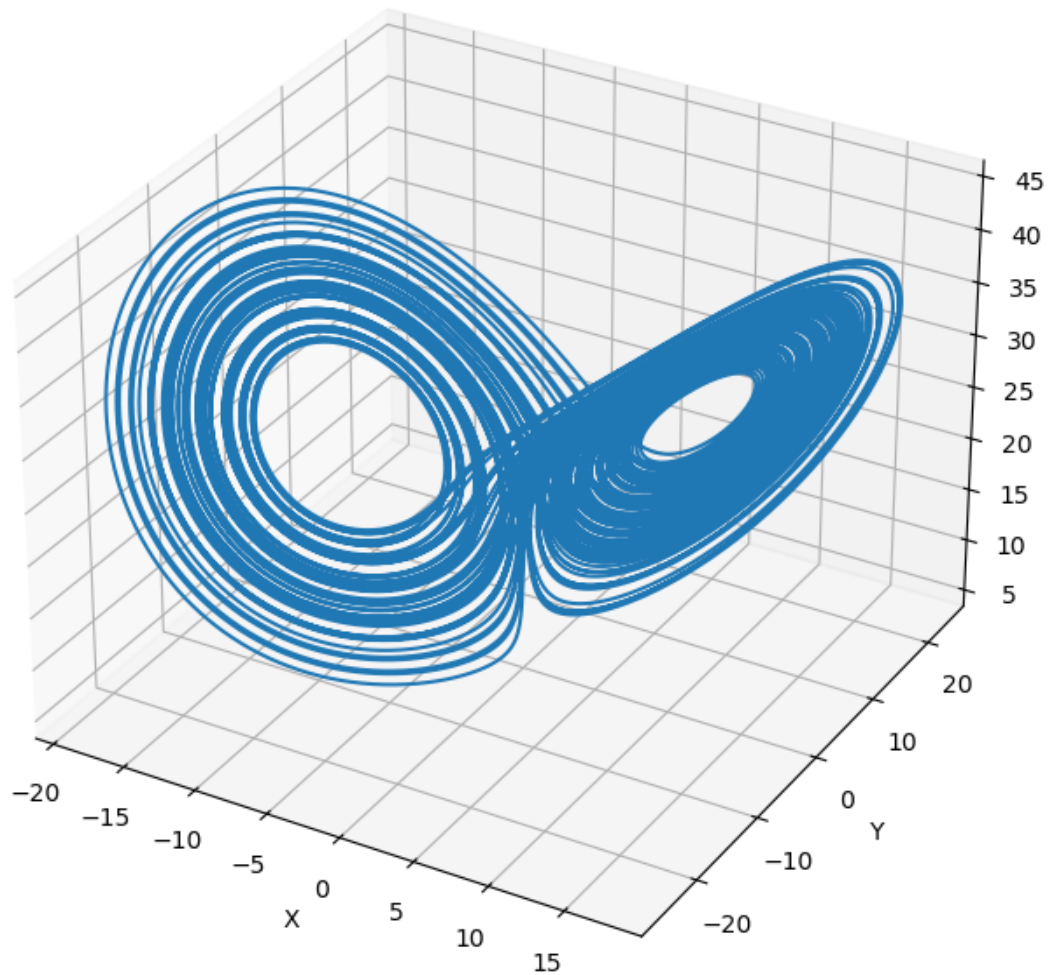
```
[ ]: plt.plot(trajectory[:, 0], trajectory[:, 2])
plt.xlabel("x")
plt.ylabel("z")
plt.title("Lorenz Attractor, XZ Projection")
None # prevent ugly output
```



```
[ ]: fig = plt.figure()
ax = fig.add_subplot(1, 1, 1, projection="3d")

ax.plot(trajectory[:, 0], trajectory[:, 1], trajectory[:, 2])
ax.set_xlabel("X")
ax.set_ylabel("Y")
ax.set_zlabel("Z") # strangely missing
plt.title("Lorenz Attractor")
None # prevent ugly output
```

## Lorenz Attractor



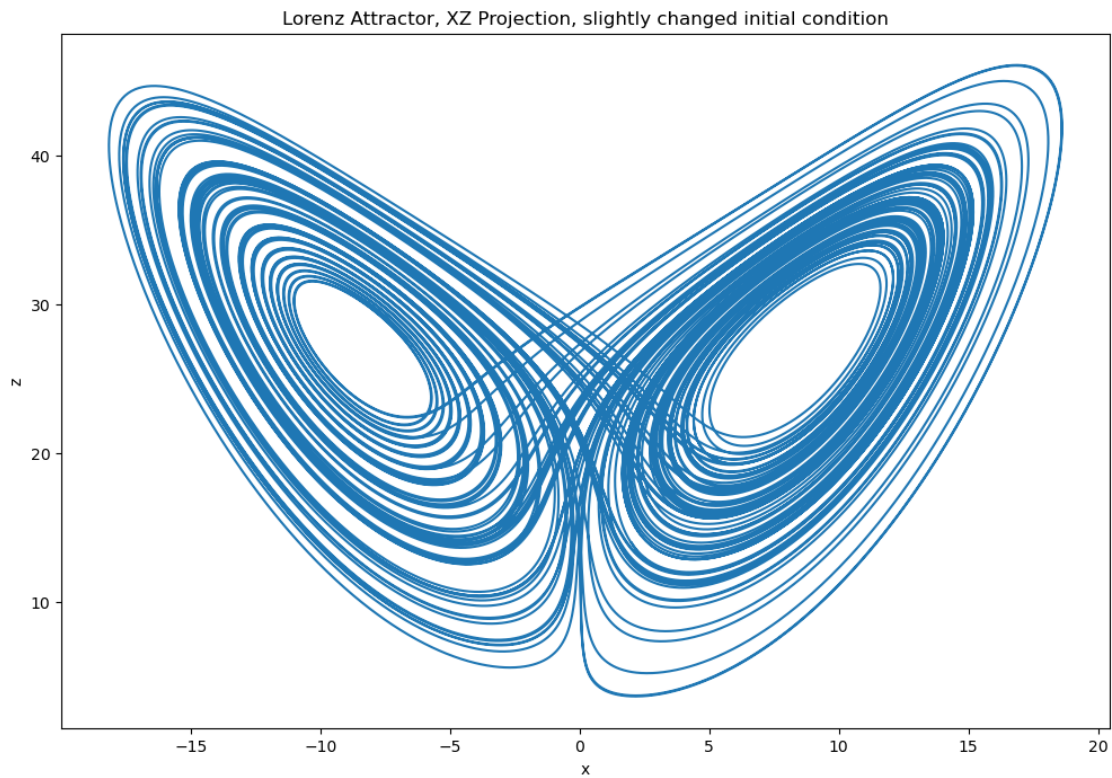
### 1.3 c)

Check the chaotic nature by using  $y_0 = (2, 3, 14 + 10^{-9})$  and compare the trajectories.

```
[ ]: y0_eps = np.array([2, 3, 14 + 1e-9], dtype=np.float64)
      rkdp2 = RKDP(lorenz, y0_eps, t0=0, tmax=T_MAX, h=H, sigma=SIGMA, r=R, b=B)
      system2 = np.array(list(rkdp2))
      time2 = system2[:, 0]
      trajectory2 = system2[:, 1:]
```

```
[ ]: plt.plot(trajectory2[:, 0], trajectory2[:, 2])
      plt.xlabel("x")
```

```
plt.ylabel("z")
plt.title("Lorenz Attractor, XZ Projection, slightly changed initial condition")
None # prevent ugly output
```

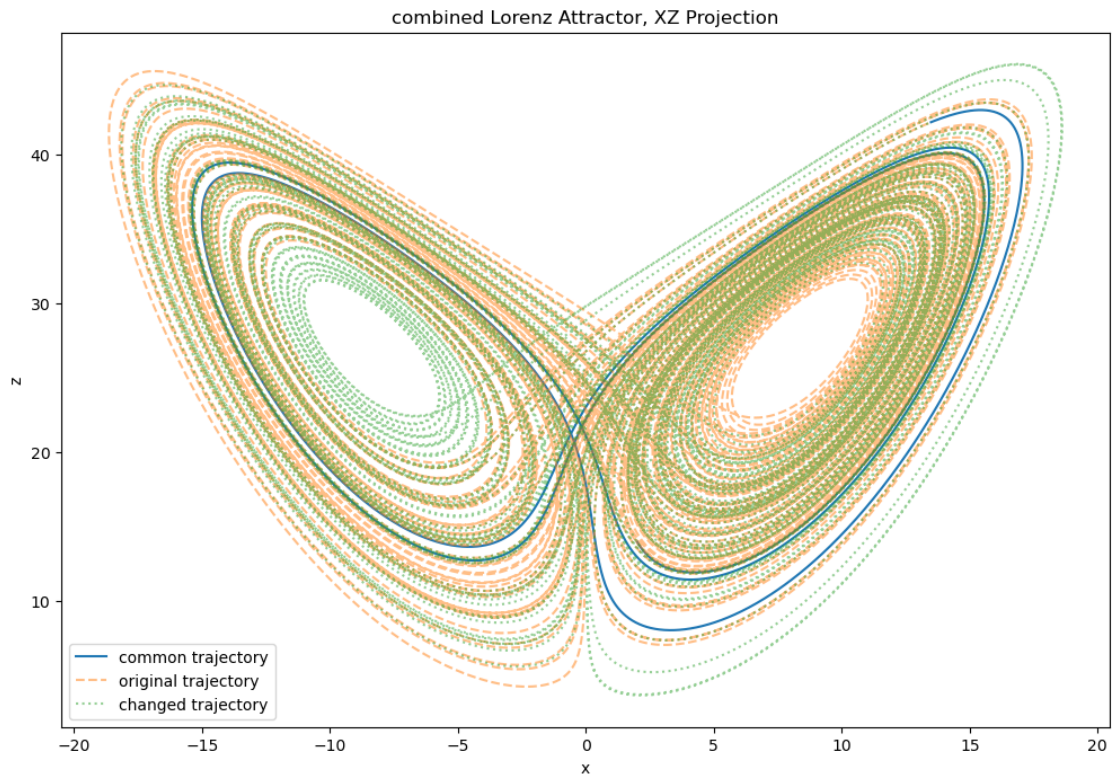


```
[ ]: traj_mask = np.all(np.isclose(trajjectory, trajjectory2, rtol=1e-7), axis=1)
div_point = np.argmin(traj_mask)
plt.plot(
    trajjectory[:div_point, 0],
    trajjectory[:div_point, 2],
    ls="solid",
    label="common trajectory",
)
plt.plot(
    trajjectory[div_point:, 0],
    trajjectory[div_point:, 2],
    ls="dashed",
    alpha=0.5,
    label="original trajectory",
)
plt.plot(
    trajjectory2[div_point:, 0],
    trajjectory2[div_point:, 2],
```

```

    ls="dotted",
    alpha=0.5,
    label="changed trajectory",
)
plt.legend()
plt.xlabel("x")
plt.ylabel("z")
plt.title("combined Lorenz Attractor, XZ Projection")
None # prevent ugly output

```



```

[ ]: plt.plot(
    trajectory[div_point:, 0],
    trajectory[div_point:, 2],
    ls="solid",
    c="tab:orange",
    label="original trajectory",
)
plt.plot(
    trajectory2[div_point:, 0],
    trajectory2[div_point:, 2],
    ls="dashed",
    c="tab:green",

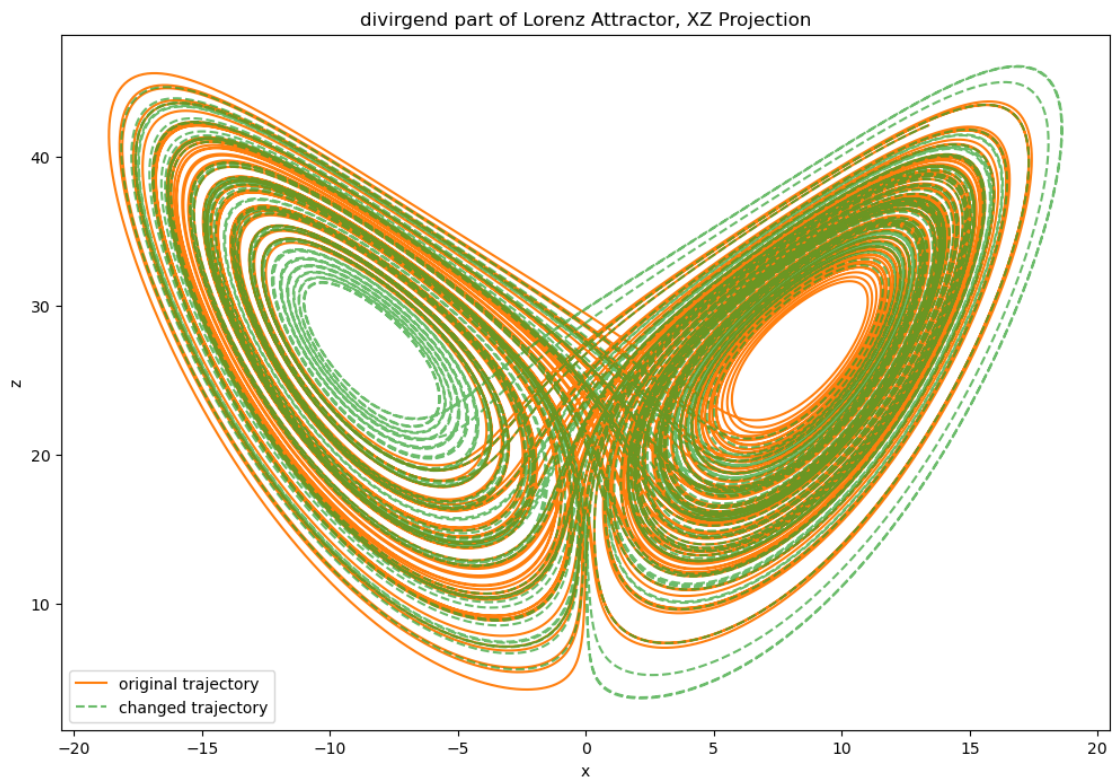
```



```

    alpha=0.7,
    label="changed trajectory",
)
plt.legend()
plt.xlabel("x")
plt.ylabel("z")
plt.title("divirgend part of Lorenz Attractor, XZ Projection")
None # prevent ugly output

```



#### 1.4 d)

Calculate the Liapunov exponent. Plot the result and create a fit to an exponential function.

```
[ ]: delta_y = np.linalg.norm(trajectory2 - trajectory, axis=1)
```

```

def liapunov(t, lambda_, a):
    return np.exp(lambda_ * t) * a

```

```
[ ]:
```

```

# automation for the search of the plateau start by fitting a function to the
# data. This only works if the data has the increas-plateau form. If not
# return NaN. As the min and max value are already good estimations for b and
# c, this fit should converge rather quickly.

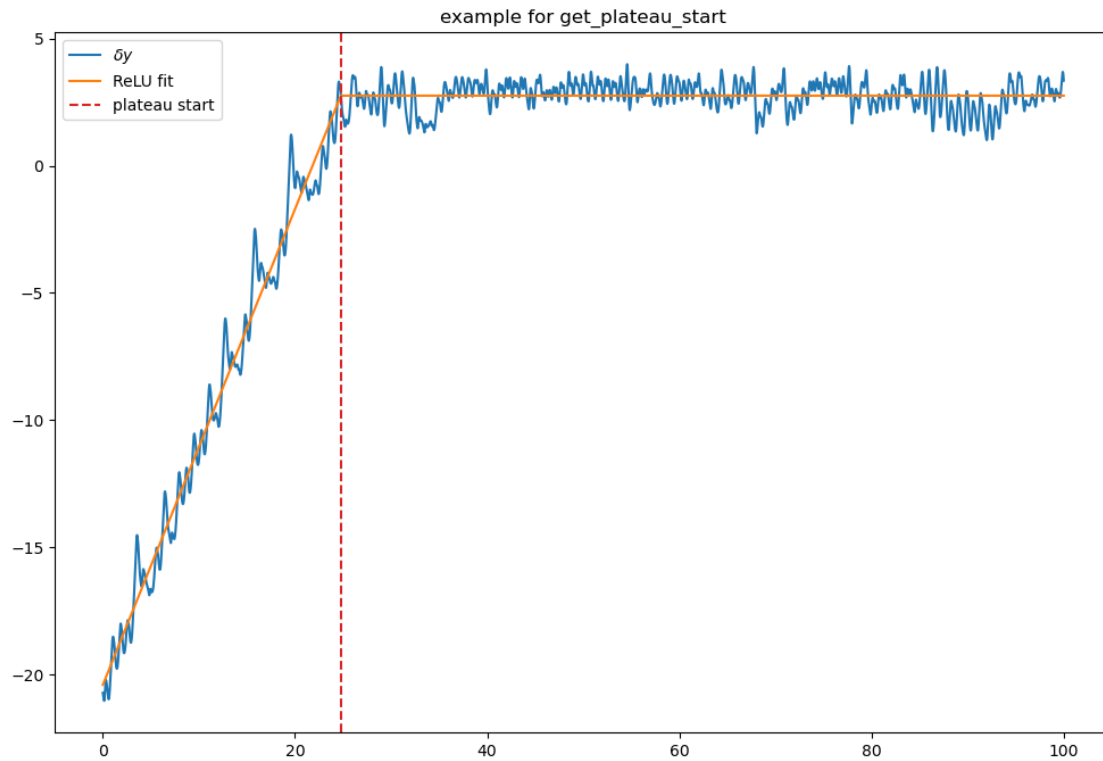
def relu(x, a, b, c):
    return np.minimum(a * x + b, c)

def get_plateau_start(
    delta_y, time
):
    # This was way more work than doing part f) by hand
    """Fitts a relu function to the data and returns the time at which the
    function plateaus."""
    logY = np.log(delta_y)
    try:
        params, cov = curve_fit(
            relu,
            time[delta_y > 0],
            logY[delta_y > 0],
            p0=[1, np.min(logY), np.max(logY)],
        )
        t_cutoff = (params[2] - params[1]) / params[0] # x = (c-b)/a
    except RuntimeError: # if the fit fails, return nan
        t_cutoff = np.nan
    return t_cutoff

# example plot of what we are doing
logY = np.log(delta_y)
params, cov = curve_fit(relu, time, logY, p0=[1, np.min(logY), np.max(logY)])
plt.plot(time, np.log(delta_y), label="$\delta y$")
plt.plot(time, relu(time, *params), label="ReLU fit")
plt.axvline(
    get_plateau_start(delta_y, time), c="tab:red", ls="--", label="plateau
    start"
)
plt.legend()
plt.title("example for get_plateau_start")
None # prevent ugly output

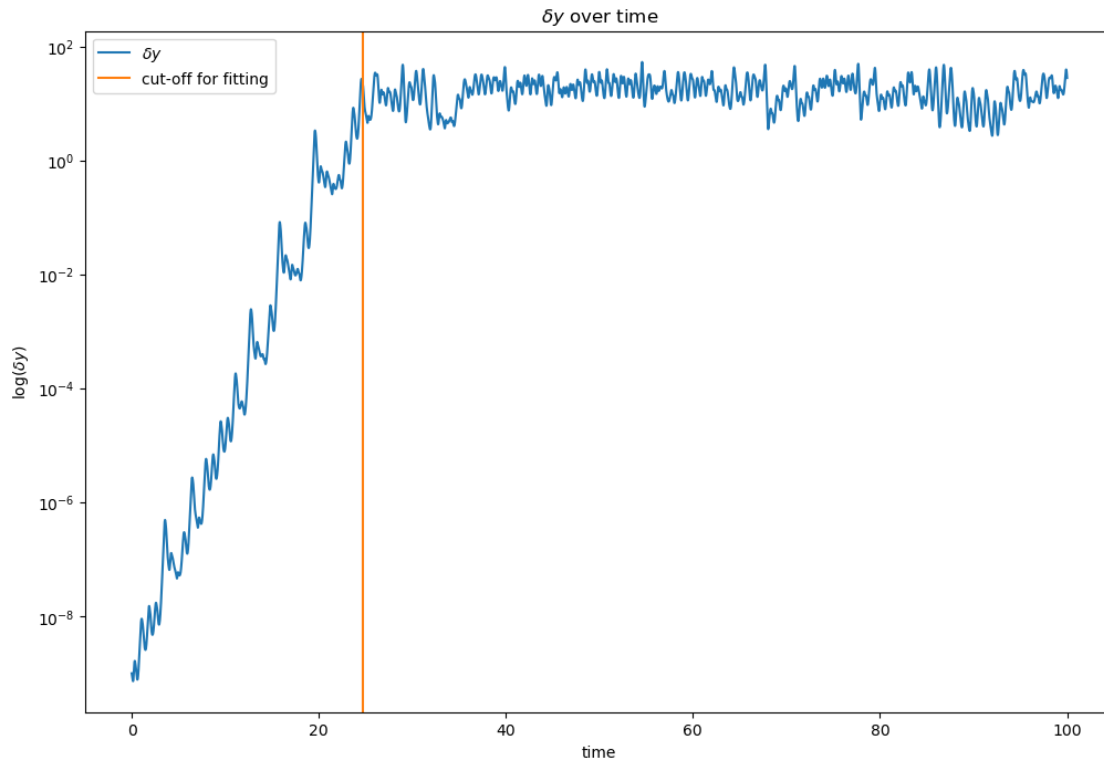
```





```
[ ]: delta_y = np.linalg.norm(trajecory2 - trajectory, axis=1)
t_cutoff = get_plateau_start(delta_y, time)

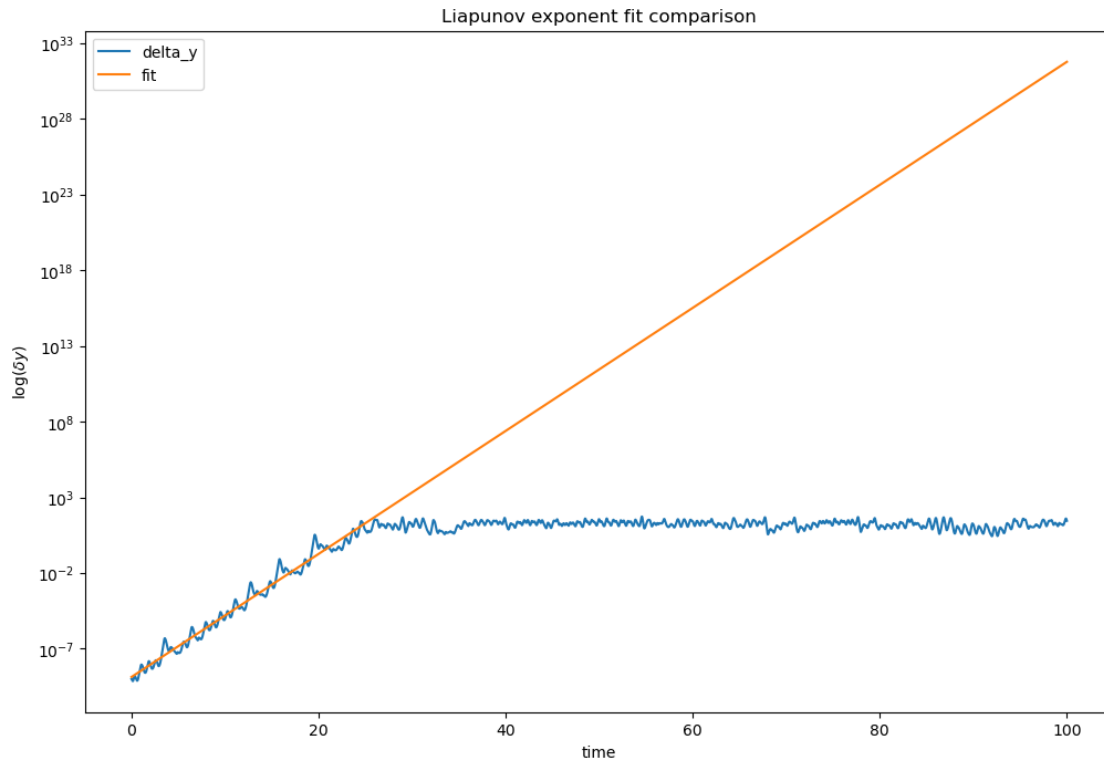
plt.plot(time, delta_y, label="$\delta y$")
plt.yscale("log")
plt.ylabel(r"log($\delta y$)")
plt.axvline(x=t_cutoff, c="tab:orange", label="cut-off for fitting")
plt.xlabel("time")
plt.title("$\delta y$ over time")
plt.legend()
None # prevent ugly output
```



```
[ ]: linFit = linregress(
    time[time < t_cutoff], np.log(delta_y[time < t_cutoff])
) # if a linear regression is possible, it performs better than an ordinary
  ↳ minimizer
print(f"Lyapunov exponent: {linFit.slope}")
print(f"a value: {np.exp(linFit.intercept)}")
```

Lyapunov exponent: 0.9357312391505839  
a value: 1.3669578951590576e-09

```
[ ]: plt.plot(time, delta_y, label="delta_y")
plt.plot(time, liapunov(time, linFit.slope, np.exp(linFit.intercept)),
  ↳ label="fit")
plt.yscale("log")
plt.ylabel(r"log($\delta y$)")
plt.xlabel("time")
plt.legend()
plt.title("Liapunov exponent fit comparison")
None # prevent ugly output
```



### 1.5 e)

Create errors by rounding the values to 10 digits. Compare them to the trajectory from b).

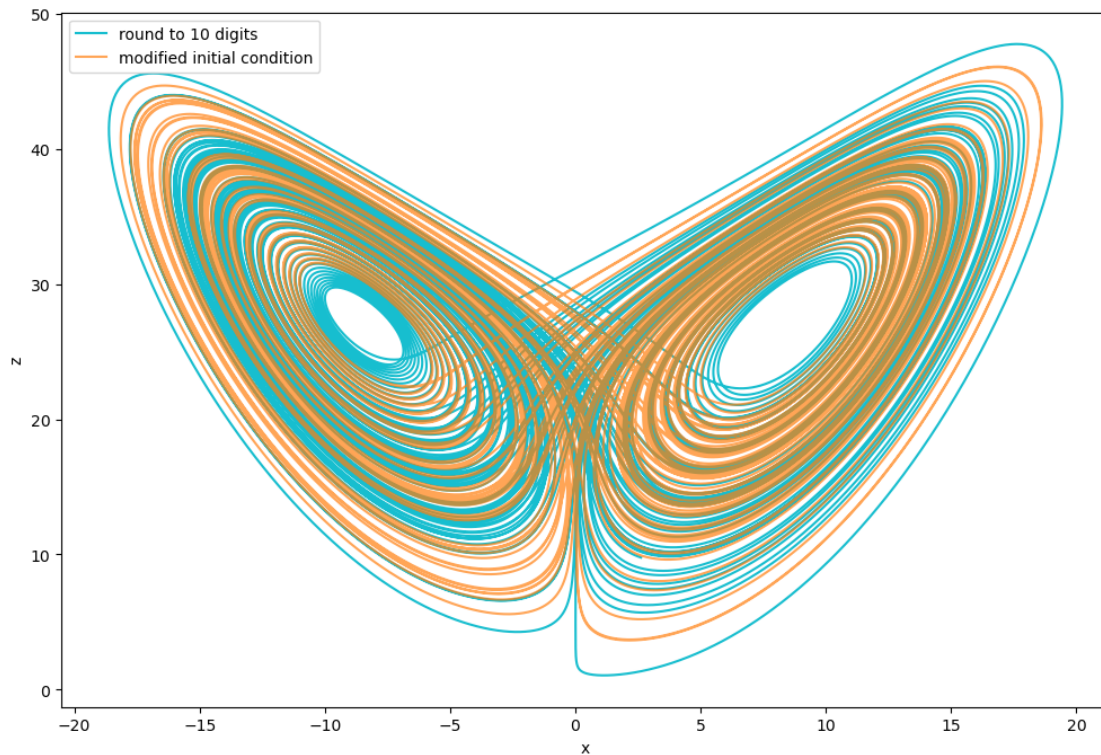
```
[ ]: rkdp3 = RKDP(
    lorentz,
    y0,
    t0=0,
    tmax=T_MAX,
    h=H,
    digits=10,
    sigma=SIGMA,
    r=R,
    b=B,
)
system3 = np.array(list(rkdp3))
time3 = system3[:, 0]
trajectory3 = system3[:, 1:]

[ ]: plt.plot(trajectory3[:, 0], trajectory3[:, 2], c="tab:cyan", label="round to 10_
    ↪digits")
plt.plot(
    trajectory2[:, 0],
```

```

trajectory2[:, 2],
c="tab:orange",
label="modified initial condition",
alpha=0.7,
)
plt.legend()
plt.xlabel("x")
plt.ylabel("z")
None # prevent ugly output

```

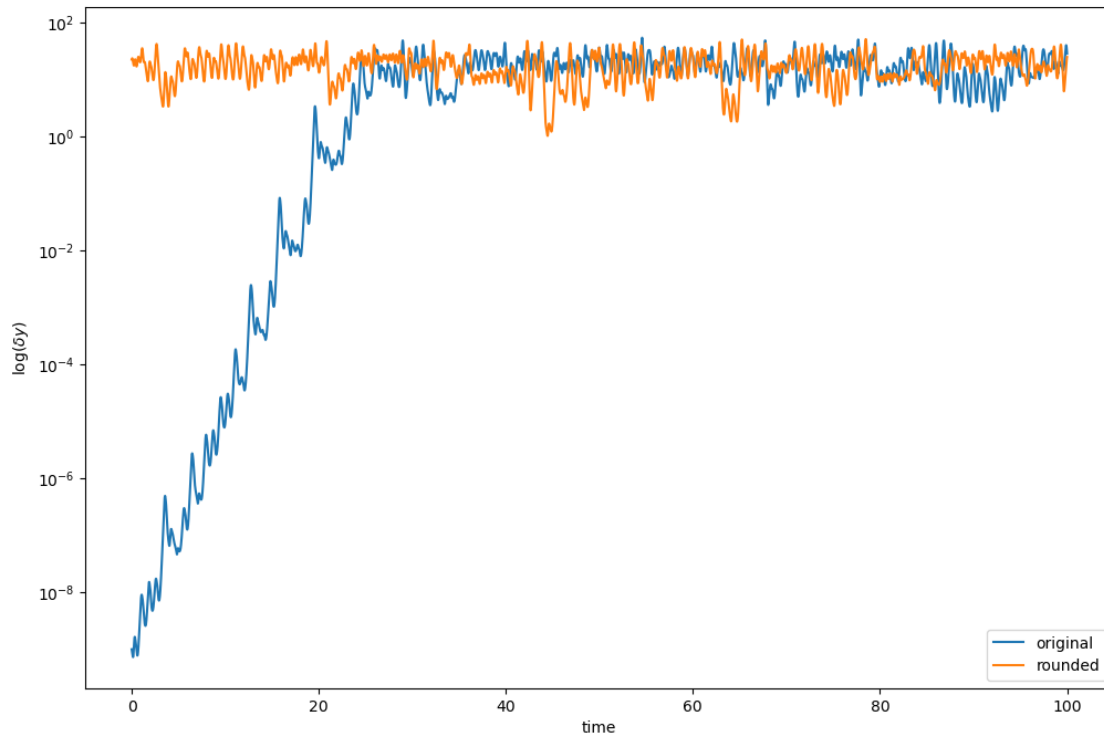


```

[ ]: delta_y3 = np.linalg.norm(trajectory3 - trajectory, axis=1)

plt.plot(time, delta_y, label="original")
plt.plot(time, delta_y3, label="rounded")
plt.yscale("log")
plt.ylabel(r"log($\delta y$)")
plt.xlabel("time")
plt.legend()
None # prevent ugly output

```



When introducing errors due to rounding,  $\delta y$  stays in the same (big) order of magnitude. The “natural” divergence has an exponential increasing error instead.

## 1.6 f)

Lower  $r$  to find a  $r_{crit}$  where the system is no longer chaotic.

```
[ ]: def liapFromLorePara(r, sigma, b, t0, tmax, h, digits=None):
    """calculates the liapunov exponent of the lorentz system for given
    ↪ parameters."""
    y0 = np.array([2, 3, 14], dtype=float) # prevent type-casting errors
    rkdp0 = RKDP(lorentz, y0, t0, tmax, h, digits=digits, sigma=sigma, r=r, b=b)
    system0 = np.array(list(rkdp0))
    time0 = system0[:, 0]
    trajectory0 = system0[:, 1:]

    y1 = np.array([2, 3, 14 + 1e-9], dtype=float) # prevent type-casting errors
    rkdp1 = RKDP(lorentz, y1, t0, tmax, h, digits=digits, sigma=sigma, r=r, b=b)
    system1 = np.array(list(rkdp1))
    time1 = system1[:, 0]
    trajectory1 = system1[:, 1:]

    delta_y = np.linalg.norm(trajectory0 - trajectory1, axis=1)
    t_cutoff = get_plateau_start(delta_y, time0)
```

```

    if np.isnan(t_cutoff):
        return np.NaN
    if len(time0[time0 < t_cutoff]) < 2 or len(time0[time0 < t_cutoff]) == len(
        time0
    ): # check if plateau start is in the time frame
        return 0
    linFit = linregress(time0[time0 < t_cutoff], np.log(delta_y[time0 <
↪t_cutoff]))
    return linFit.slope

```

```

[ ]: from multiprocessing import Pool, cpu_count

rs = np.linspace(1, 30, 59)

liapF = partial(liapFromLorePara, sigma=SIGMA, b=B, t0=0, tmax=T_MAX, h=H)

with Pool(cpu_count() - 2) as p: # use all but 2 cores
    liap = np.array(p.map(liapF, rs))

```

```

/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:14: OptimizeWarning: Covariance of the
parameters could not be estimated
    params, cov = curve_fit(
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)

```

```

/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:12: RuntimeWarning: divide by zero
encountered in log
    logY = np.log(delta_y)
/tmp/ipykernel_16137/1903625063.py:14: OptimizeWarning: Covariance of the
parameters could not be estimated
    params, cov = curve_fit(
/tmp/ipykernel_16137/1903625063.py:14: OptimizeWarning: Covariance of the
parameters could not be estimated
    params, cov = curve_fit(
/tmp/ipykernel_16137/1903625063.py:14: OptimizeWarning: Covariance of the
parameters could not be estimated
    params, cov = curve_fit(

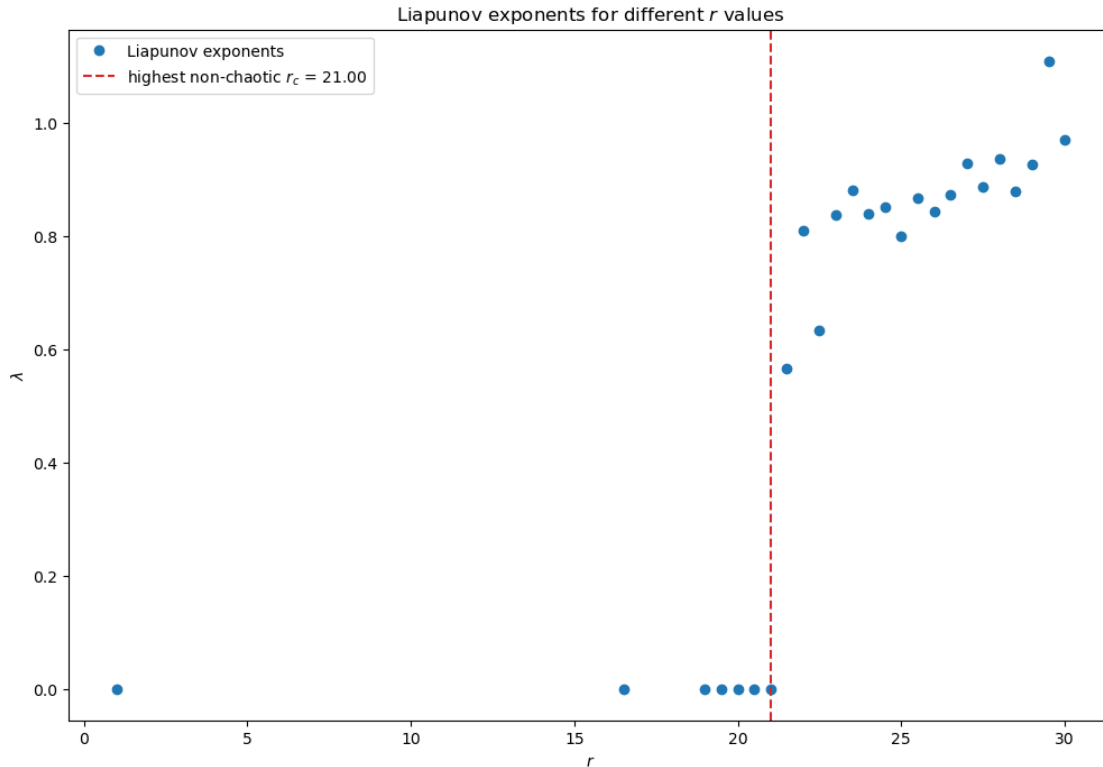
```

```

[ ]: r_crit = rs[liap == 0][-1]
plt.plot(rs, liap, marker="o", ls="none", label="Liapunov exponents")
plt.axvline(
    r_crit, c="tab:red", ls="--", label=f"highest non-chaotic $r_c$ = {r_crit:.
↪2f}"
)
plt.xlabel("$r$")
plt.ylabel("$\lambda$")
plt.legend()
plt.title("Liapunov exponents for different $r$ values")
None # prevent ugly output

```





### 1.7 g)

Implement the dynamic step-size using a safety factor of  $S = 0.95$ . Compare the compute time to the fixed approach.

```
[ ]: def RKF(f, y0, t0, tmax, h0, atol, rtol, hmax, S=0.95, alpha=0.2, **kwargs):
    dim = len(y0)
    coef = { # Coefficients for the Dormand-Prince method, padded with zeros
    ↪ for the 6th order
        "k2": np.array([1 / 5, 0, 0, 0, 0, 0]).reshape(-1, 1),
        "k3": np.array([3 / 40, 9 / 40, 0, 0, 0, 0]).reshape(-1, 1),
        "k4": np.array([44 / 45, -56 / 15, 32 / 9, 0, 0, 0]).reshape(-1, 1),
        "k5": np.array(
            [19372 / 6561, -25360 / 2187, 64448 / 6561, -212 / 729, 0, 0]
        ).reshape(-1, 1),
        "k6": np.array(
            [9017 / 3168, -355 / 33, 46732 / 5247, 49 / 176, -5103 / 18656, 0]
        ).reshape(
            -1, 1
        ), # called a6 in the literature
        "y5": np.array(
            [35 / 384, 0, 500 / 1113, 125 / 192, -2187 / 6784, 11 / 84]
```

```

        ).reshape(
            -1, 1
        ), # called b in the literature
        "y4": np.array(
            [5179 / 57600, 0, 7571 / 16695, 393 / 640, -92097 / 339200, 187 / 2100]
        ).reshape(
            -1, 1
        ), # called b* in the literature
        "t": np.array(
            [0, 1 / 5, 3 / 10, 4 / 5, 8 / 9, 1]
        ), # called c in the literature
    }

    t = t0
    y = y0
    h = h0
    fp = partial(f, **kwargs)
    while t < tmax:
        ks = np.zeros((6, dim))
        ks[0] = h * fp(t, y)
        ks[1] = h * fp(t + h * coef["t"][1], y + np.sum(ks * coef["k2"]))
        ks[2] = h * fp(t + h * coef["t"][2], y + np.sum(ks * coef["k3"]))
        ks[3] = h * fp(t + h * coef["t"][3], y + np.sum(ks * coef["k4"]))
        ks[4] = h * fp(t + h * coef["t"][4], y + np.sum(ks * coef["k5"]))
        ks[5] = h * fp(t + h * coef["t"][5], y + np.sum(ks * coef["k6"]))
        y5 = np.sum(coef["y5"] * ks, axis=0)
        y4 = np.sum(coef["y4"] * ks, axis=0) + y5 / 40
        scale = atol + rtol * np.max([norm(y), norm(y5)], axis=0)
        err = np.sqrt(np.mean(((y5 - y4) / scale) ** 2))

        if err < 1:
            t += h
            h = np.min(
                [S * h * (1 / err) ** alpha, hmax]
            ) # here we check for the maximum step size
            y += y5
            yield t, *y
        else:
            h = S * h * (1 / err) ** alpha

```

```

[ ]: rkf = RKF(
    lorentz,
    y0,
    t0=0,
    tmax=10,
    h0=0.01,

```

```

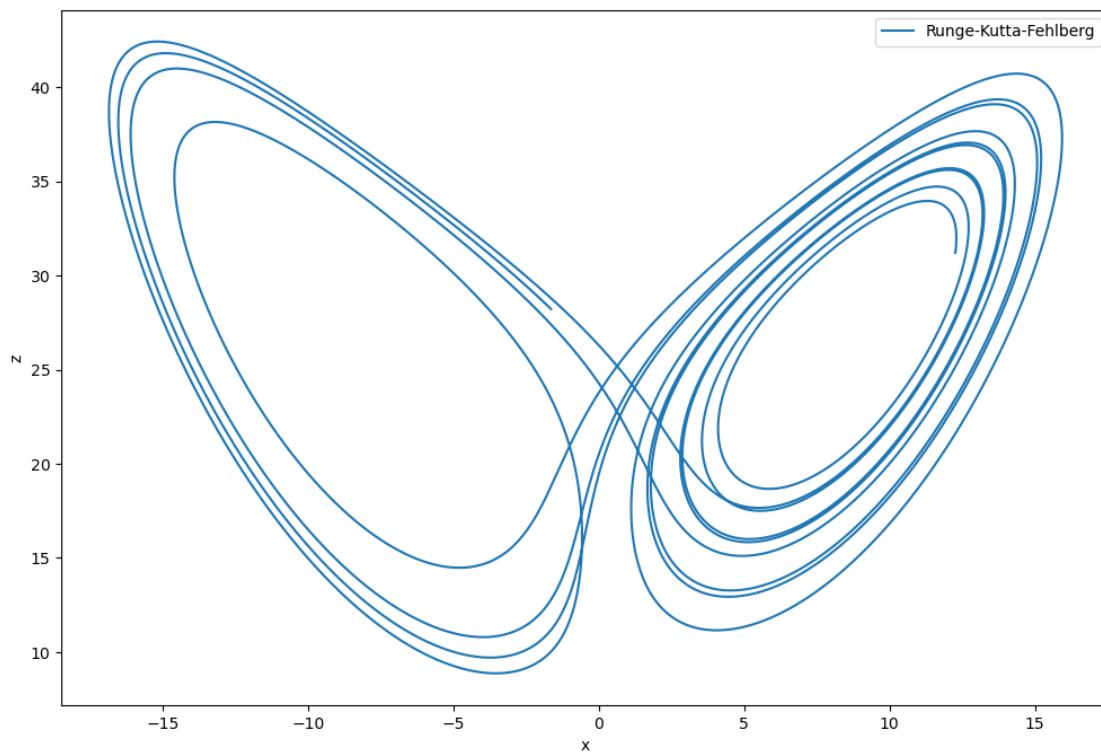
    hmax=2,
    atol=1e-8, # 1e-12 takes way too long
    rtol=1e-8, # took around 10 minutes
    sigma=SIGMA,
    r=R,
    b=B,
)
system_rkf = np.array(list(rkf))
time_rkf = system_rkf[:, 0]
trajectory_rkf = system_rkf[:, 1:]

```

```

[ ]: plt.plot(trajectory_rkf[:, 0], trajectory_rkf[:, 2],
             ↪label="Runge-Kutta-Fehlberg")
plt.legend()
plt.xlabel("x")
plt.ylabel("z")
None # prevent ugly output

```



```

[ ]: hs = np.logspace(-5, -2, 5)

def hToDeltaY(h):

```

```

rkdp_ = RKDP(lorentz, y0, t0=0, tmax=10, h=h, sigma=SIGMA, r=R, b=B)
system_ = np.array(list(rkdp_))
time_temp = system_[:, 0]
trajectory_ = system_[:, 1:]
return (norm(trajectory_[-1] - trajectory_rkf[-1]), h)

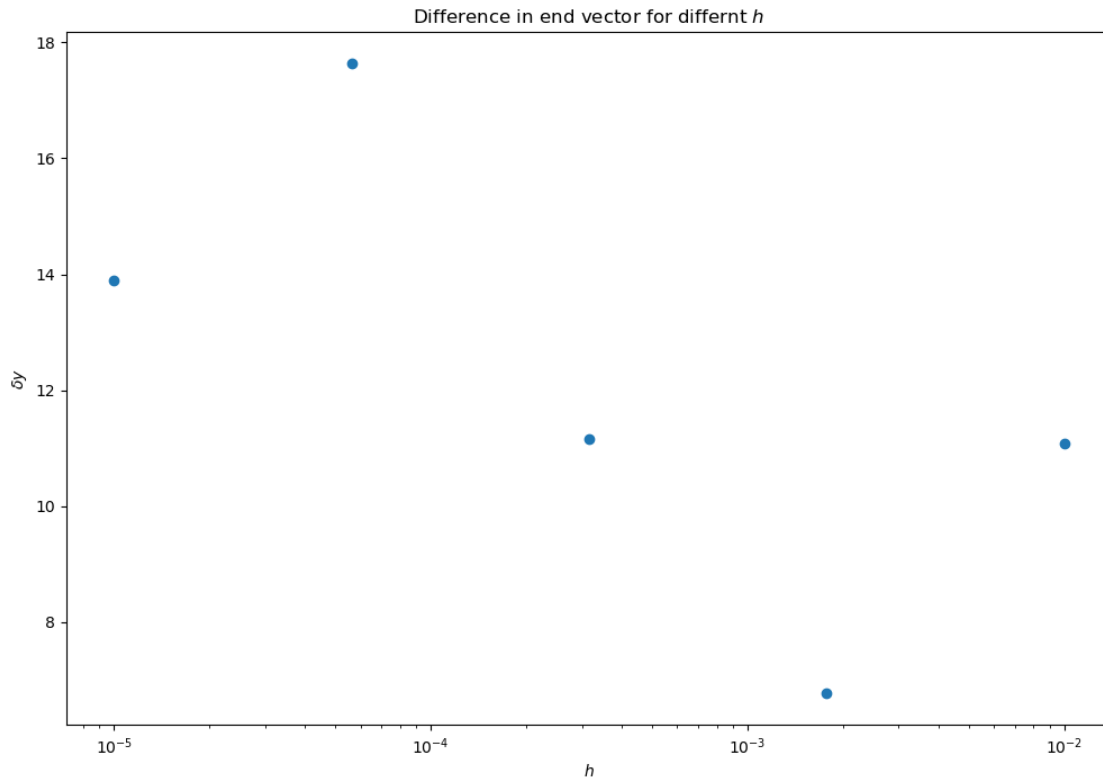
with Pool(cpu_count() - 2) as p: # use all but 2 cores
    pool_result = np.array(p.map(hToDeltaY, hs))

```

```

[ ]: plt.scatter(pool_result[:, 1], pool_result[:, 0])
plt.xscale("log")
plt.xlabel("$h$")
plt.ylabel("$\delta y$")
plt.title("Difference in end vector for differnt $h$")
None # prevent ugly output

```



We are not even close to  $\delta y(t_e) < 10^{-6}$ . But reducing  $h$  any more leads to unreasonable computation time. We might have an error in this implementation.

## 2 A1

### 2.1 a)

ODE integration methods with a dynamik stepsize have the advantage, that generally they run as fast as possible. Due to the overhead it is possible that a given equation computes slower than a fixed stepsize, but this should rarely be the case.

### 2.2 b)

The stepsize estimator is based on the different error estimator of the runge kutta methods. In the rk4 method we have  $err = O(h^{-5}) - O(h^{-4})$ . So for a given error we can calculate the optimal  $h$