

This code is written to evaluate Equivalent Clean Air (ECA) flow rate, also called the clean air delivery rate (CADR), using two different approaches.

The first approach is called log-linear approach that is conventionally used to evaluate ECA based on first-order decay assumption. The next approach is called Integral approach which uses area under decay curve to estimate ECA.

First, we import all the needed python libraries.

```
In [1]: import numpy as np
        from scipy import optimize
        import matplotlib.pyplot as plt
        from scipy import stats
```

Next, we have to import concentration dataset alongside its measurement errors, either manually or through reading from a file.

In the following section, 8 different cases are provided as examples of concentration dataset. The examples used are as follows:

- Case #1: Ideal first order
- Case #2: Non-biological particle by Zeng et al. ([link](#)).
- Case #3: Hypothetical E. Coli by Stephens et al. [link](#).
- Case #4-6: Bacteriophage MS2 by Ratliff et al. [doi](#)
- Case #7: High res. VOC (formaldehyde) by Schumacher [doi](#)
- Case #8: Low res. VOC (formaldehyde) by Law et al. [doi](#)

## Case #1: Ideal first order

```
In [ ]: t_c = np.array([0, 15, 30, 60, 90, 120]) # Time points
        t_t = np.array([0, 15, 30, 60, 90, 120]) # Time points

        num_simulations = 10000 # Number of simulations

        c_test = np.array([ 8.00E+10, 2.29E+10, 6.57E+09,
                           5.39E+08, 4.42E+07, 3.63E+06]) # Test concentrations
        c_control = np.array([8.00E+10, 4.28E+10, 2.29E+10,
                              6.57E+09, 1.88E+09, 5.39E+08]) # Control concentrations

        c_test = np.log10(c_test)
        c_control = np.log10(c_control)

        c_bg_test=1 # If there is background
        c_bg_control=1
        c_bg_test = np.log10(c_bg_test)
        c_bg_control = np.log10(c_bg_control)

        sigma_test = np.array([0,0,0,0,0,0]) # Test standard deviations
        sigma_control = np.array([0,0,0,0,0,0]) # Control standard deviations
```

```
# Chamber
# Chamber
```

```
In [ ]: num_simulations = 10000 # Number of simulations

t_t = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10,
                11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21]) # Time points

t_c = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10,
                11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21]) # Time points

c_control = np.array([46629.5822, 45262.5924, 38944.8016, 41968.5662, 39589.2249,
                      37112.7987, 34422.0024, 33845.9813, 31813.524, 26790.9496,
                      30782.9434, 29226.4405, 27259.9242, 27281.7278, 26790.9496,
                      25085.1379, 22133.3491, 22427.7312, 21338.2308, 18838.1459,
                      19229.6865, 19123.3036]) # Test control

c_control=np.log10(c_control)

c_test = np.array([20733.1887, 17199.744, 15496.7706, 13660.3969, 11310.3118,
                  10883.8787, 9122.8056, 8170.2825, 6648.0319, 6299.9989,
                  4945.4016, 4635.3165, 3815.3086, 3194.0609, 2984.6603,
                  2455.5343, 2276.8268, 1918.0086, 1619.6262, 1389.2609,
                  1244.1919, 1231.9916]) # Control

c_test=np.log10(c_test)

sigma_control = np.array([0., 0., 0., 0., 0., 0., 0., 0., 0., 0.,
                          0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0.])

sigma_test = np.array([0., 0., 0., 0., 0., 0., 0., 0., 0., 0.,
                       0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0., 0.])

c_bg_test=113.7572667 # If there is a background
c_bg_control=607.22006143
#c_bg_test = np.log10(c_bg_test)
#c_bg_control = np.log10(c_bg_control)

V = 1296.048 # Chamber volume
#V = 36.7 # Chamber volume
```

```
In [ ]: num_simulations = 10000 # Number of simulations

t_c = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79, 80, 81, 82, 83, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 97, 98, 99])
```



```

c_bg_control=1
c_bg_test = np.log10(c_bg_test)
c_bg_control = np.log10(c_bg_control)

sigma_test = np.array([0,0,0,0,0]) # Test sta
sigma_control = np.array([0,0,0,0,0]) # Control
V = 500 # Chamber
#V = 14.2 # Chamber

```

## Case #4: Ratliff et al BPI

```

In [ ]: t_c = np.array([0, 15, 30, 60, 90 ,120, 1200]) # Time poi
t_t = np.array([0, 15, 30, 60, 90 ,120, 1200]) # Time poi

c_test = np.array([8.46, 7.63, 7.22, 6.21, 5.89, 5.64, 10**0.1 ]) # Test cor
c_control = np.array([8.47, 7.68, 7.16, 7.09, 6.21, 6.10, 10**0.1 ]) # Control

sigma_test = np.array([0.000001, 0.24 ,0.51 ,0.52 ,0.53 ,0.5, 0]) # Test
sigma_control = np.array([0.000001 ,0.36, 0.39, 0.3, 0.36, 0.49, 0]) # Cor

c_bg_test=0 # If there
c_bg_control=0

V = 3000 # Chamber
#V = 85 # Chamber

```

## Case #5: Ratliff et al PCO1

```

In [ ]: t_c = np.array([0, 15, 30, 60, 90]) # Time poi
t_t = np.array([0, 15, 30, 60, 90]) # Time poi

c_test = np.array([ 8.11212724060371, 7.589987734, 6.745802232,
6.013936402, 5.394571206 ]) # Test cor

c_control = np.array([8.46745953764963, 7.684595602, 7.163725729,
7.086631702, 6.211568868]) # Control

sigma_test = np.array([ 0.108808569398572, 0.254108644, 0.169662969,
0.262771866, 0.047624259]) # Test

sigma_control = np.array([ 0.237966029977382, 0.358965447, 0.393189888,
0.301479043, 0.361953917]) # Control

c_bg_test=0 # If there
c_bg_control=0

V = 3000 # Chamber
#V = 85 # Chamber

```

## Case #6: Ratliff et al PCO2

```
In [ ]: t_c = np.array([0, 15, 30, 60, 90]) # Time po
t_t = np.array([0, 15, 30, 60, 90]) # Time po
c_test = np.array([8.032891171, 7.087704857, 6.424694684,
                  5.320557829, 4.709637767]) # Test co

c_control = np.array([8.467459538, 7.684595602, 7.163725729,
                    7.086631702, 6.211568868]) # Control

sigma_test = np.array([0.033101768, 0.029661622, 0.219839957,
                      0.117283171, 0.129724927]) # Test
sigma_control = np.array([0.23796603, 0.358965447, 0.393189888,
                        0.301479043, 0.361953917]) # Control

c_bg_test=0 # If there
c_bg_control=0

V = 3000 # Chamber
#V = 85 # Chamber
```

## Case #7: Real VOC (High Resolution) (Schumacher et. al. 2024)

```
In [46]: # full dataset

t_c = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60])

t_t = np.array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60])

c_test = np.array([833.75, 761.28, 658.93, 567.20, 499.72, 451.38, 422.54, 377.29, 321.06, 295.83, 223.47, 192.89, 168.51, 180.00, 154.03, 117.60, 114.59, 94.81, 86.12, 74.05, 65.78, 61.08, 55.48, 51.09, 47.80, 43.67, 38.95, 35.24, 32.66, 30.53, 28.45, 26.76, 27.44, 29.12, 23.70, 24.67, 21.88, 19.60, 18.40, 17.51, 19.38, 19.07, 16.17, 14.67, 13.91, 13.55, 12.91, 12.21, 11.86, 12.43, 12.45, 12.32, 11.57, 11.21, 11.55, 10.61, 10.93, 11.39, 11.73, 11.32, 11.30 ])

c_test = np.log10(c_test)
c_control = np.array([929.45, 951.57, 923.26, 906.75, 912.08, 933.73, 915.11, 904.99, 891.02, 898.66, 899.28, 881.89, 880.40, 889.34, 888.76, 896.80, 891.25, 880.13, 870.49, 862.30, 862.03, 866.09, 874.08, 864.19, 880.57, 872.92, 861.04, 856.26, 862.26, 863.67, 843.84, 844.73, 842.50, 838.72, 844.15, 843.70, 855.47, 834.44, 830.98, 823.93, 822.37, 825.25, 848.16, 812.26, 885.45, 863.72, 826.80, 833.43, 822.13, 809.45, 806.02, 800.97, 801.26, 795.55, 836.46, 827.70, 785.76, 786.31, 786.06, 851.69, 837.34]) # Co
```

```

c_control = np.log10(c_control)
sigma_test = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                        0]) # Test sta

sigma_control = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
                           0]) # Test sta

c_bg_test=9.66 # If tl
c_bg_control=250
#c_bg_test = np.log10(c_bg_test)
#c_bg_control = np.log10(c_bg_control)

V = 1034.72 # Chamber
#V = 29.3 # Chamber

```

## Case #8: Law et al 2024 (truncated dataset)

```

In [9]: #0-40 minutes for control and 0-25 minutes for test

t_t = np.array([0, 5, 10, 15, 20, 25]) # Time poi
t_c = np.array([0, 5, 10, 15, 20, 25, 30, 35, 40 ]) # Time poi

c_test = np.array([0.958783047, 0.502557891, 0.40034575,
                   0.332838565, 0.301776099, 0.272451193]) # Test c

c_control = np.array([0.972665911, 0.969369171, 0.959128797, 0.959302774,
                      0.95947675, 0.950973937, 0.949412555, 0.940911944,
                      0.941083718]) # Control

sigma_test = np.array([0, 0, 0, 0, 0, 0 ]) # Test sta
sigma_control = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]) # Control

c_test = np.log10(c_test)
c_control=np.log10(c_control)

c_bg_test=0.24 # If there
c_bg_control=0.65

V = 1059.44174
#V = 30

```

## Case #8: Law et al 2024 (Full dataset)

```

In [27]: t_c = np.array([0, 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60]) # Time poi

```

```

t_t = np.array([0, 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60]) # Time points

#0-60 minutes dataset (full dataset)

c_test = np.array([0.958783047, 0.502557891, 0.40034575, 0.332838565,
                   0.301776099, 0.272451193, 0.272625169, 0.251974851,
                   0.262558772, 0.262730546, 0.252494577, 0.252666351,
                   0.24242818 ]) # Test concentrations

c_test=np.log10(c_test)

c_control = np.array([0.972665911, 0.969369171, 0.959128797, 0.959302774,
                     0.95947675, 0.950973937, 0.949412555, 0.940911944,
                     0.941083718, 0.910021252, 0.910193026, 0.899954854,
                     0.901861986]) # Control concentrations

c_control=np.log10(c_control)

sigma_control = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]) # Control standard deviations
sigma_test     = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]) # Test standard deviations

c_bg_test=0.24 # If there is background
c_bg_control=0.65

V = 1059.44174 # Chamber volume
#V = 30 # Chamber volume

```

# Monte Carlo Analysis

## 1- First-order decay method:

The loss rate has been calculated based on the following formula

<!DOCTYPE html>

$$K = \frac{\sum_{i=0}^{np} t_i \ln\left(\frac{C_i - C_{bg}}{C_0 - C_{bg}}\right)}{\sum_{i=0}^{np} t_i^2}$$

Then, loss rates will be plugged into the CADR formula to obtain value for CADR

<!DOCTYPE html>

$$CADR = V(K_{Test} - K_{Control})$$

In [47]:

```

num_simulations = 10000
# Function to compute the CADR component for either test or control
def compute_component(t, c, c_bg):
    #print((c / c[0]))
    ln_c_ratio = -np.log((10**c-c_bg) / (10**c[0]-c_bg))

    numerator = np.sum((t * ln_c_ratio) )
    denominator = np.sum((t ** 2) )

    return (numerator / denominator)

# Original CADR value
CADR_original = V * (compute_component(t_t, c_test, c_bg_test) - compute_component(t_c, c_

```

```

print(f"Original CADR value = {CADR_original}", "\n")

# Monte Carlo simulations
CADR_values = np.zeros(num_simulations)

for i in range(num_simulations):
    c_test_simulated = np.random.normal((c_test), (sigma_test/3))
    c_control_simulated = np.random.normal((c_control), (sigma_control/3))
    CADR_values[i] = V * (compute_component(t_t, c_test_simulated, c_bg_test) - compute_co

# Calculate mean and standard deviation of CADR
CADR_mean = np.mean(CADR_values)
CADR_std = np.std(CADR_values)

print(f"Mean CADR value from Monte Carlo simulations = {CADR_mean}", "\n")
print(f"Standard deviation of CADR value from Monte Carlo simulations = {CADR_std}", "\n")
print(f"Error = {100*CADR_std/CADR_mean}%", "\n")

# Plotting the distribution of CADR values
plt.hist(CADR_values, bins=50, alpha=0.7, color='b', edgecolor='black')
plt.title('Distribution of CADR values from Monte Carlo simulations \n(first-order decay r
plt.xlabel('CADR ')
plt.ylabel('Frequency')

# Disable scientific notation for x-axis and y-axis
plt.ticklabel_format(style='plain', axis='x')
plt.ticklabel_format(style='plain', axis='y')

# Optionally, adjust the format of the ticks
from matplotlib.ticker import ScalarFormatter
plt.gca().xaxis.set_major_formatter(ScalarFormatter(useOffset=False))
plt.gca().yaxis.set_major_formatter(ScalarFormatter(useOffset=False))

plt.show()

```

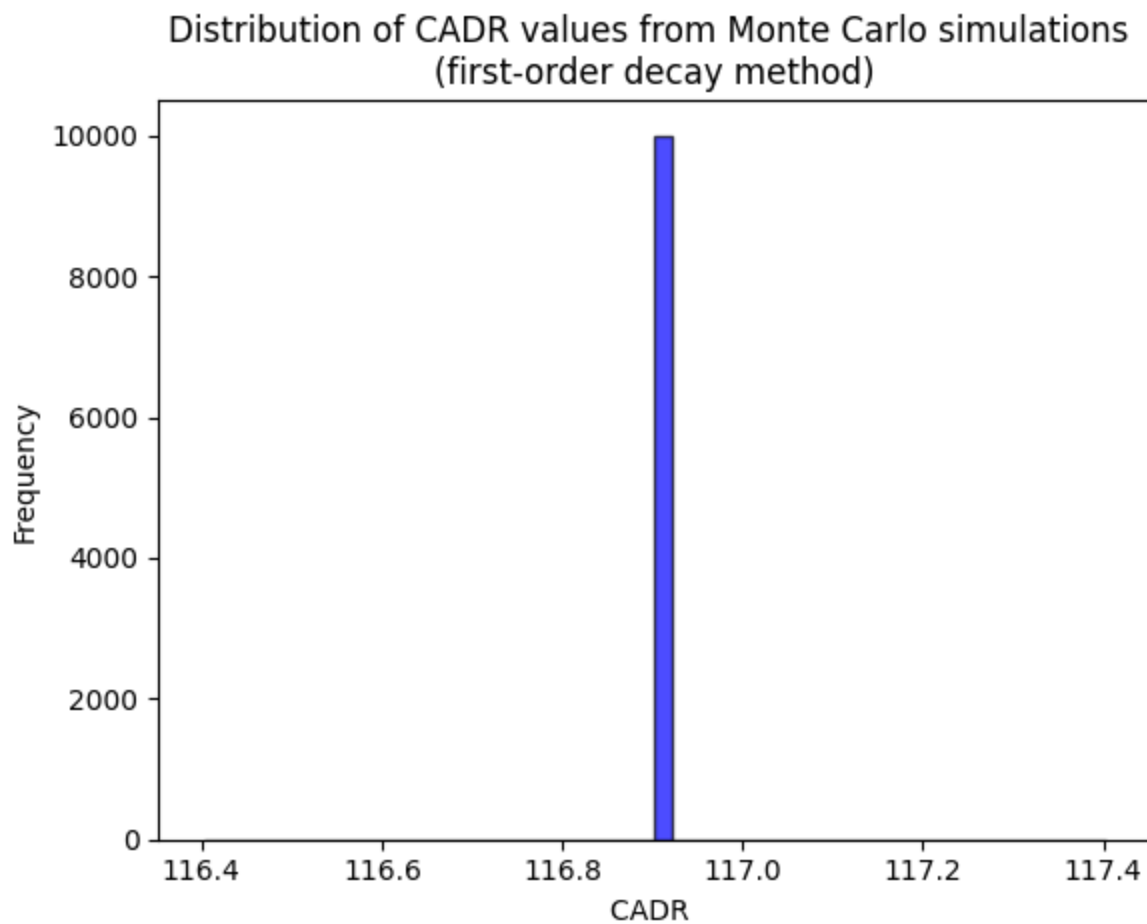
Original CADR value = 116.90209908956523

Mean CADR value from Monte Carlo simulations = 116.90209908956517

Standard deviation of CADR value from Monte Carlo simulations = 5.684341886080802e-14

Error = 4.862480597312211e-14%





The CADR base on the integral method can be calculated as follows:

<!DOCTYPE html>

$$\text{CADR}_{\text{Integral}} = \frac{VC_{\text{test}}(0)}{\int_0^{\infty} (C_{\text{test}}(t) - C_{bg, \text{control}}) dt} - \frac{VC_{\text{control}}(0)}{\int_0^{\infty} (C_{\text{control}}(t) - C_{bg, \text{control}}) dt}$$

Where the integral in the denominator can be calculated using either trapezoidal or logarithmic trapezoidal method. here, we use logarithmic trapezoidal method which looks like the following:

<!DOCTYPE html>

$$\int_0^{\infty} C(t) dt = \sum_{n=1}^{\infty} \left( \frac{t_i - t_{i-1}}{\ln \left( \frac{C(t_{i-1}) - C_{bg}}{C(t_i) - C_{bg}} \right)} (C(t_{i-1}) - C(t_i)) \right)$$

In [48]:

```
import numpy as np
import matplotlib.pyplot as plt

num_simulations = 10
# Function to compute the integral
def compute_integral(t, c, c_bg):
    N = len(t)
    integral_sum = 0.0

    for i in range(1, N):
        t_diff = t[i] - t[i-1]
        ln_c_ratio = np.log((10**c[i-1]-c_bg) / (10**c[i]-c_bg))
        #print(ln_c_ratio)
```

```

        c_diff = 10**c[i-1] - 10**c[i]
        integral_sum += (t_diff / ln_c_ratio) * c_diff
    #print(integral_sum)
    return integral_sum

# Function to compute the CADR component for either test or control
def compute_component(t, c, c_bg):

    integral_c = compute_integral(t, c, c_bg)
    #print(integral_c)
    return (10**c[0]) / integral_c

# Original CADR value
CADR_original = V* (compute_component(t_t, c_test, c_bg_test) - compute_component(t_c, c_c

print("test",compute_component(t_t, c_test, c_bg_test))
print("control",compute_component(t_c, c_control, c_bg_control))
print(f"Original CADR value = {CADR_original}\n")

# Monte Carlo simulations
CADR_values = np.zeros(num_simulations)
'''
for i in range(num_simulations):
    c_test_simulated = np.random.normal(c_test, sigma_test)
    plt.plot(t_t, c_test_simulated)
    c_control_simulated = np.random.normal(c_control, sigma_control)
    plt.plot(t_c, c_control_simulated)

    CADR_values[i] = V* (compute_component(t_t, c_test_simulated) - compute_component(t_c,
#print(CADR_values)
plt.show()
'''

#fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(14, 6))

for i in range(num_simulations):
    c_test_simulated = np.random.normal(c_test, sigma_test/3)
    c_control_simulated = np.random.normal(c_control, sigma_control/3)

    #ax1.plot(t_t, c_test_simulated, alpha=0.1)
    #ax2.plot(t_c, c_control_simulated, alpha=0.1)

    CADR_values[i] = V * (compute_component(t_t, c_test_simulated, c_bg_test) - compute_co
# Customize plots
'''
ax1.set_title('Test Simulations')
ax1.set_xlabel('Time')
ax1.set_ylabel('Concentration')

ax2.set_title('Control Simulations')
ax2.set_xlabel('Time')
ax2.set_ylabel('Concentration')

plt.tight_layout()
plt.show()
'''

# Print CADR values if needed
# print(CADR_values)

# Calculate mean and standard deviation of CADR
CADR_mean = np.mean(CADR_values)
CADR_std = np.std(CADR_values)

print(f"Mean CADR value from Monte Carlo simulations = {CADR_mean}\n")

```

```

print(f"Standard deviation of CADR value from Monte Carlo simulations = {CADR_std}\n")
print(f"Error = {CADR_std / CADR_mean * 100}%\n")

# Plotting the distribution of CADR values
plt.hist(CADR_values, bins=50, alpha=0.7, color='b', edgecolor='black')
plt.title('Distribution of CADR values from Monte Carlo simulations \n(Integral method)')
plt.xlabel('CADR ')
plt.ylabel('Frequency')

# Disable scientific notation for x-axis and y-axis
plt.ticklabel_format(style='plain', axis='x')
plt.ticklabel_format(style='plain', axis='y')

# Optionally, adjust the format of the ticks
from matplotlib.ticker import ScalarFormatter
plt.gca().xaxis.set_major_formatter(ScalarFormatter(useOffset=False))
plt.gca().yaxis.set_major_formatter(ScalarFormatter(useOffset=False))

plt.show()

```

test 0.1272083774282268

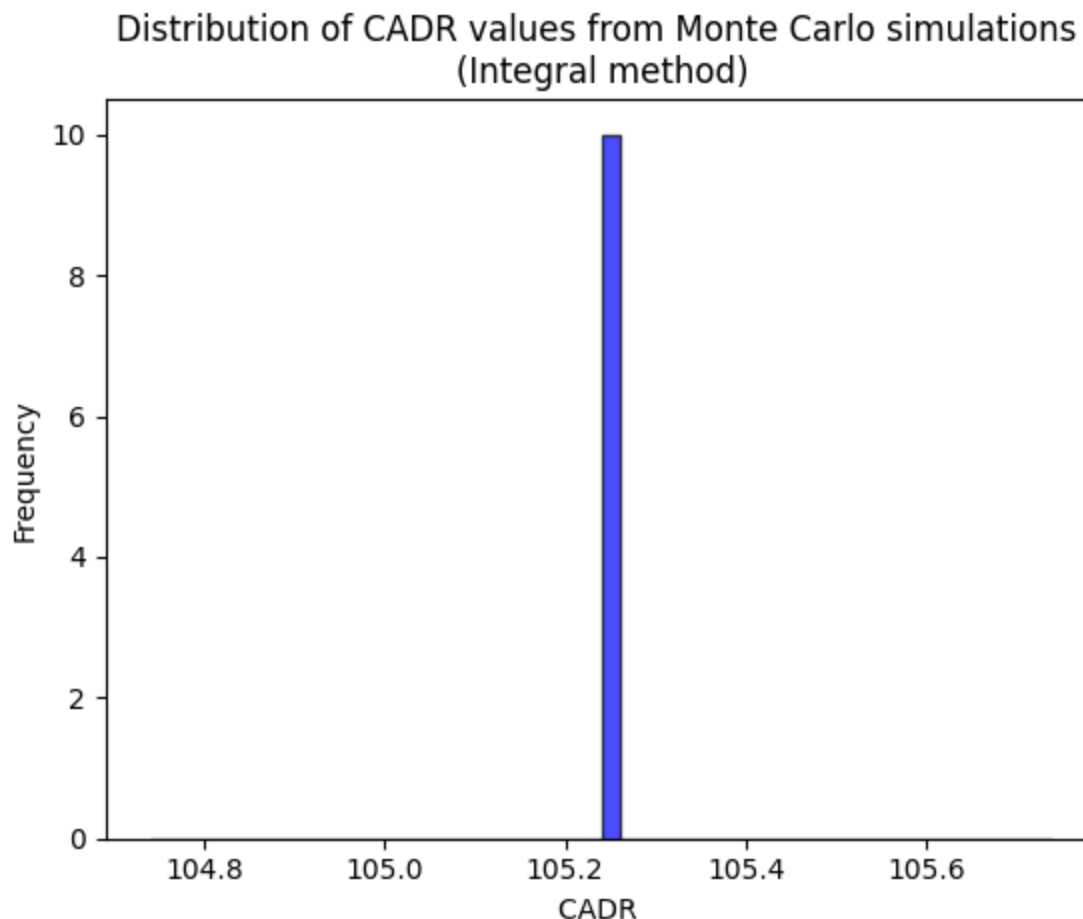
control 0.025499043930522245

Original CADR value = 105.24068155674486

Mean CADR value from Monte Carlo simulations = 105.24068155674486

Standard deviation of CADR value from Monte Carlo simulations = 0.0

Error = 0.0%



In [11]:

```

import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import t
from scipy import stats

```

```

num_simulations = 10000

# Define the compute_k1 function using the first part of the new formula
def compute_k1(t, c, c_bg):
    ln_c_ratio = -np.log((10**c-c_bg) / (10**c[0]-c_bg))
    numerator = np.sum(t * ln_c_ratio)
    denominator = np.sum(t ** 2)
    return numerator / denominator

# Define the compute_k2 function using the second part of the new formula
def compute_k2(t, c, k, c_bg, alpha=0.05):
    ln_c_ratio = -np.log((10**c-c_bg) / (10**c[0]-c_bg))

    # Calculate the mean of the time points
    t_mean = np.mean(t)

    # Calculate the second term
    residual_sum = np.sum(((ln_c_ratio - k * t)) ** 2)
    #print(k)
    critical_value = stats.t.ppf(1- alpha / 2, len(t) - 1)
    #second_term = critical_value * np.sqrt(residual_sum / ((len(t) - 1) * np.sum((t) ** 2)))
    second_term = np.sqrt(residual_sum / ((len(t) - 1) * np.sum((t) ** 2)))

    return second_term

# Monte Carlo simulations for the first formula

k1_values_test = np.zeros(num_simulations)
k1_values_control = np.zeros(num_simulations)

for i in range(num_simulations):
    c_test_simulated = np.random.normal(c_test, sigma_test / 3)
    c_control_simulated = np.random.normal(c_control, sigma_control / 3)
    k1_values_test[i] = compute_k1(t_t, c_test_simulated, c_bg_test)
    k1_values_control[i] = compute_k1(t_c, c_control_simulated, c_bg_control)

# Monte Carlo simulations for the second formula
k2_values_test = np.zeros(num_simulations)
k2_values_control = np.zeros(num_simulations)

for i in range(num_simulations):
    c_test_simulated = np.random.normal(c_test, sigma_test / 3)
    c_control_simulated = np.random.normal(c_control, sigma_control / 3)
    k_test = compute_k1(t_t, c_test_simulated, c_bg_test)
    k_control = compute_k1(t_c, c_control_simulated, c_bg_control)
    k2_values_test[i] = compute_k2(t_t, c_test_simulated, k_test, c_bg_test)
    k2_values_control[i] = compute_k2(t_c, c_control_simulated, k_control, c_bg_control)

# Calculate mean and standard deviation for both formulas
k1_mean_test = np.mean(k1_values_test)
k1_std_test = np.std(k1_values_test)
k1_mean_control = np.mean(k1_values_control)
k1_std_control = np.std(k1_values_control)

k2_mean_test = np.mean(k2_values_test)
k2_std_test = np.std(k2_values_test)
k2_mean_control = np.mean(k2_values_control)
k2_std_control = np.std(k2_values_control)

print(f"Mean k1 value for test from Monte Carlo simulations = {k1_mean_test}\n")
print(f"Standard deviation of k1 value for test from Monte Carlo simulations = {k1_std_test}\n")
print(f"Mean k1 value for control from Monte Carlo simulations = {k1_mean_control}\n")
print(f"Standard deviation of k1 value for control from Monte Carlo simulations = {k1_std_control}\n")

```

```

print(f"Mean k2 value for test from Monte Carlo simulations = {k2_mean_test}\n")
print(f"Standard deviation of k2 value for test from Monte Carlo simulations = {k2_std_test}\n")
print(f"Mean k2 value for control from Monte Carlo simulations = {k2_mean_control}\n")
print(f"Standard deviation of k2 value for control from Monte Carlo simulations = {k2_std_control}\n")

# Plotting the distribution of k1 values
plt.hist(k1_values_test, bins=50, alpha=0.7, color='b', edgecolor='black', label='k1 Test')
plt.hist(k1_values_control, bins=50, alpha=0.7, color='r', edgecolor='black', label='k1 Control')
plt.title('Distribution of k1 values from Monte Carlo simulations')
plt.xlabel('k1')
plt.ylabel('Frequency')
plt.legend()
plt.show()

# Plotting the distribution of k2 values
plt.hist(k2_values_test, bins=50, alpha=0.7, color='b', edgecolor='black', label='k2 Test')
plt.hist(k2_values_control, bins=50, alpha=0.7, color='r', edgecolor='black', label='k2 Control')
plt.title('Distribution of k2 values from Monte Carlo simulations')
plt.xlabel('k2')
plt.ylabel('Frequency')
plt.legend()
plt.show()

```

Mean k1 value for test from Monte Carlo simulations = 0.12891970374259107

Standard deviation of k1 value for test from Monte Carlo simulations = 2.7755575615628914e-17

Mean k1 value for control from Monte Carlo simulations = 0.0026821435337412926

Standard deviation of k1 value for control from Monte Carlo simulations = 4.336808689942018e-19

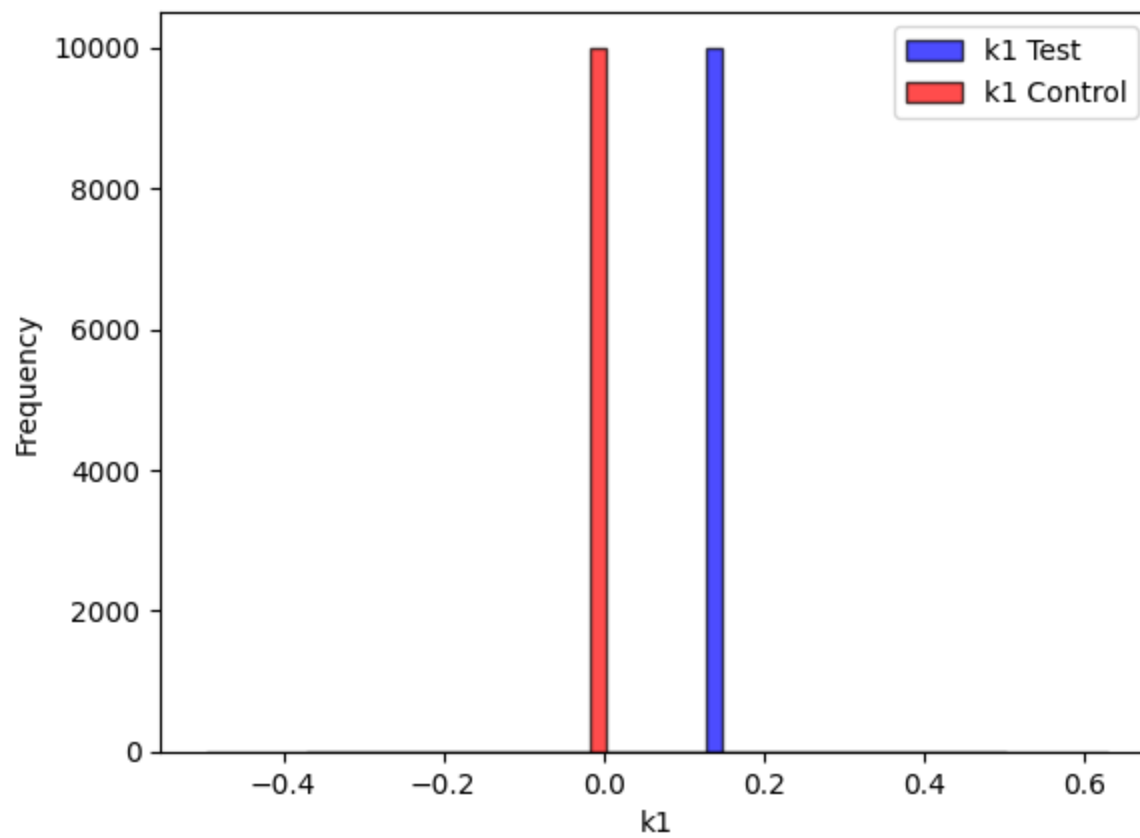
Mean k2 value for test from Monte Carlo simulations = 0.005654409953692421

Standard deviation of k2 value for test from Monte Carlo simulations = 8.673617379884035e-19

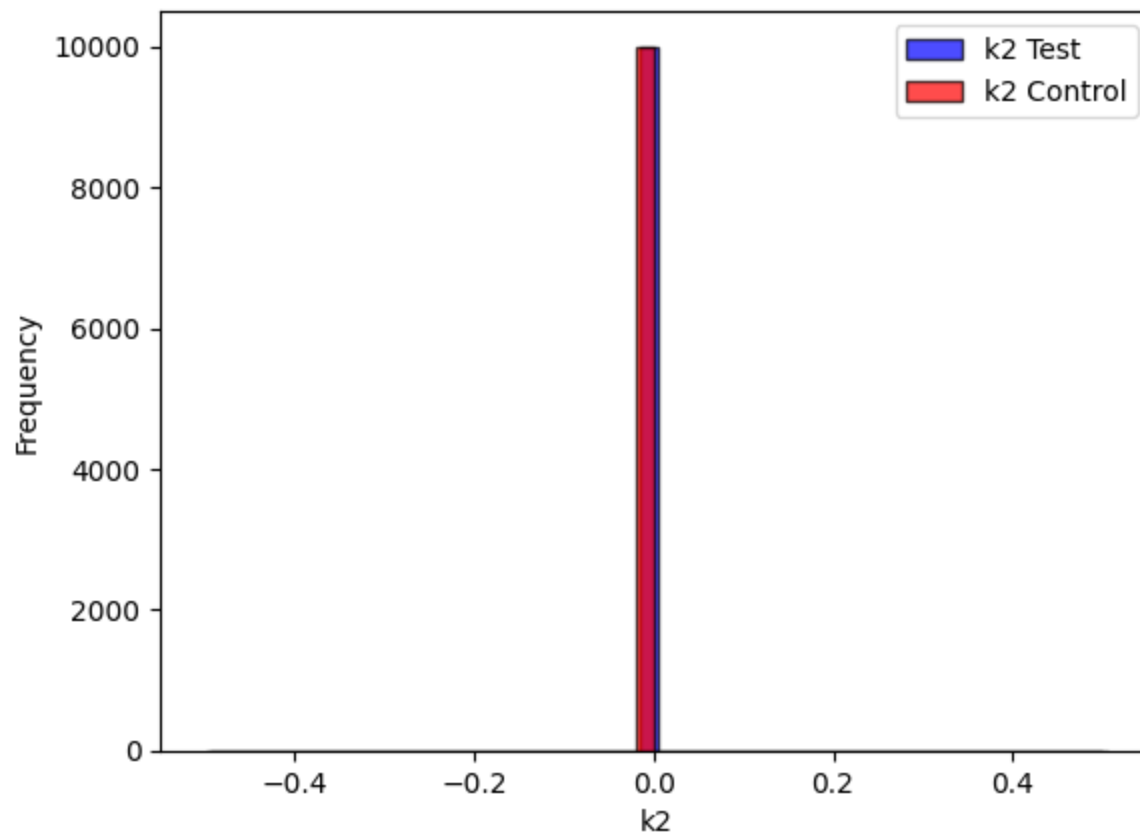
Mean k2 value for control from Monte Carlo simulations = 0.0001176333295346026

Standard deviation of k2 value for control from Monte Carlo simulations = 4.0657581468206416e-20

Distribution of k1 values from Monte Carlo simulations



Distribution of k2 values from Monte Carlo simulations



In [ ]: