

week_8_differencing

October 16, 2023

```
[ ]: import pandas as pd # standard
import numpy as np # standard
from sklearn import tree # package to make decision tree
from sklearn.metrics import accuracy_score # for accuracy calculation
from sklearn.metrics import balanced_accuracy_score
from sklearn.metrics import roc_auc_score

import matplotlib.pyplot as plt
import seaborn as sns

import thermogram_utilities

import warnings
warnings.filterwarnings("ignore")

df = pd.read_excel("/Users/avery/OneDrive/Documents/GitHub/
↳Clinical_TLB_2023-2024/lung_cancer_tlb.xlsx")

# replace NA with control
df['CancerType'] = np.where(df['CancerType'].isna(), 'Control',
↳df['CancerType'])

# get location of cut off values
lower_column_index = df.columns.get_loc("T50")
upper_column_index = df.columns.get_loc("T85.1")
label_column_index = df.columns.get_loc("CancerType")

column_indices = np.arange(lower_column_index, upper_column_index)
column_indices = np.append(column_indices, 0)
column_indices = np.append(column_indices, 1)

column_indices = np.append(column_indices, label_column_index)

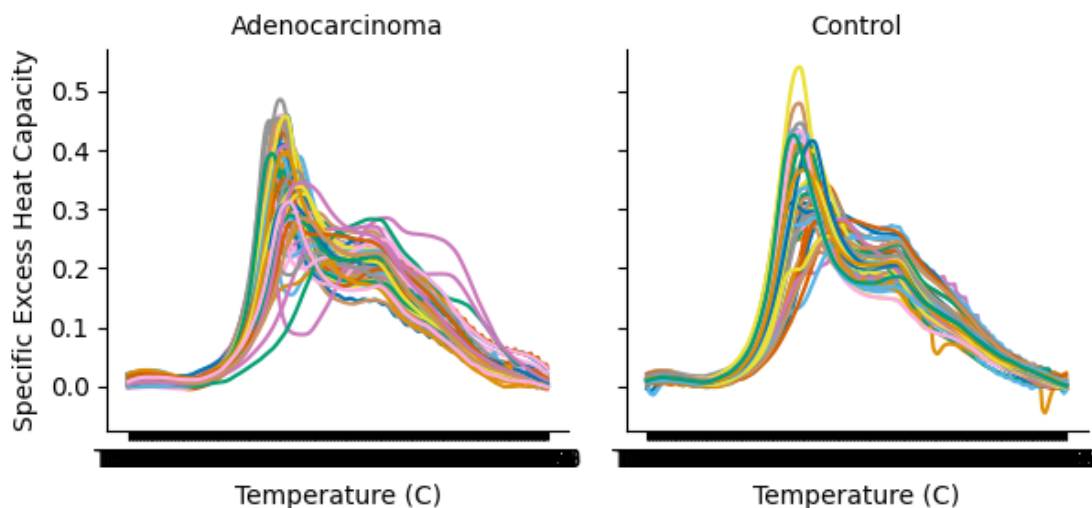
df = df.iloc[:, column_indices]
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# keep only Control and Adenocarcinoma for analysis
df_tree = df[(df['CancerType'] == 'Control') | (df['CancerType'] ==
↳ 'Adenocarcinoma')]
df_tree = df_tree.reset_index(drop=True)
```

```
[ ]: df_long = pd.melt(df_tree, id_vars=['sample_id', 'pub_id', 'CancerType'],
↳ var_name='temp', value_name='dsp' )
df_long['temperature'] = df_long['temp'].str.replace('T', '').astype(float)

g = sns.FacetGrid(df_long, col="CancerType", col_wrap= 3, hue="sample_id",
↳ palette = 'colorblind')
g.map_dataframe(sns.lineplot, x="temp", y="dsp")
g.set_axis_labels("Temperature (C)", "Specific Excess Heat Capacity")
g.set_titles(col_template="{col_name}")
```

```
[ ]: <seaborn.axisgrid.FacetGrid at 0x1c92521f9d0>
```



```
[ ]: df_long = pd.melt(df_tree, id_vars=['sample_id', 'pub_id', 'CancerType'],
↳ var_name='temp', value_name='dsp' )

median_df = thermogram_utilities.median_curve(df_long, 'CancerType', 'temp',
↳ 'dsp')

median_df['temperature'] = median_df['temperature'].str.replace('T', '').
↳ astype(float)

plt.figure(figsize=(10, 6)) # Adjust the figure size if needed
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sns.lineplot(data=median_df, x='temperature', y='median', hue='type',
             palette='colorblind')
ax = plt.gca()

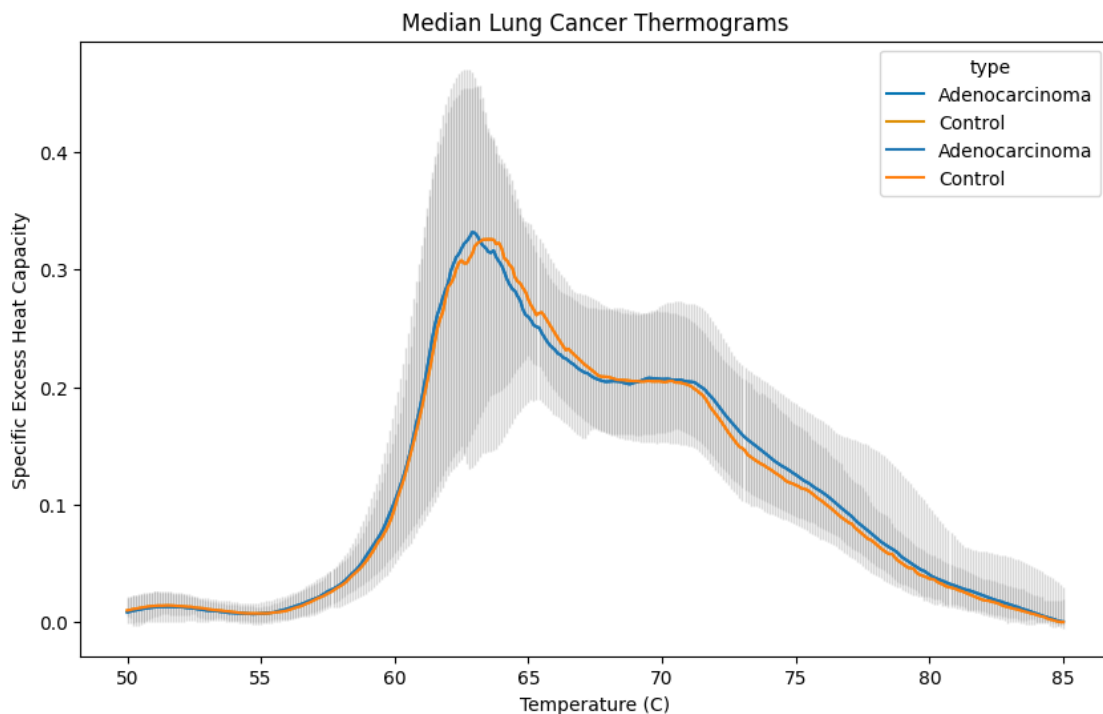
# Plot the lines with different colors for 'type'
sns.lineplot(data=median_df, x='temperature', y='median', hue='type')

# Add ribbons for each 'type' with different colors
for _, row in median_df.iterrows():
    ax.fill_between([row['temperature']], row['lower_q'], row['upper_q'],
                   alpha=0.3, color='grey')

plt.xlabel('Temperature (C)')
plt.ylabel('Specific Excess Heat Capacity')
plt.title('Median Lung Cancer Thermograms')

```

```
[ ]: Text(0.5, 1.0, 'Median Lung Cancer Thermograms')
```



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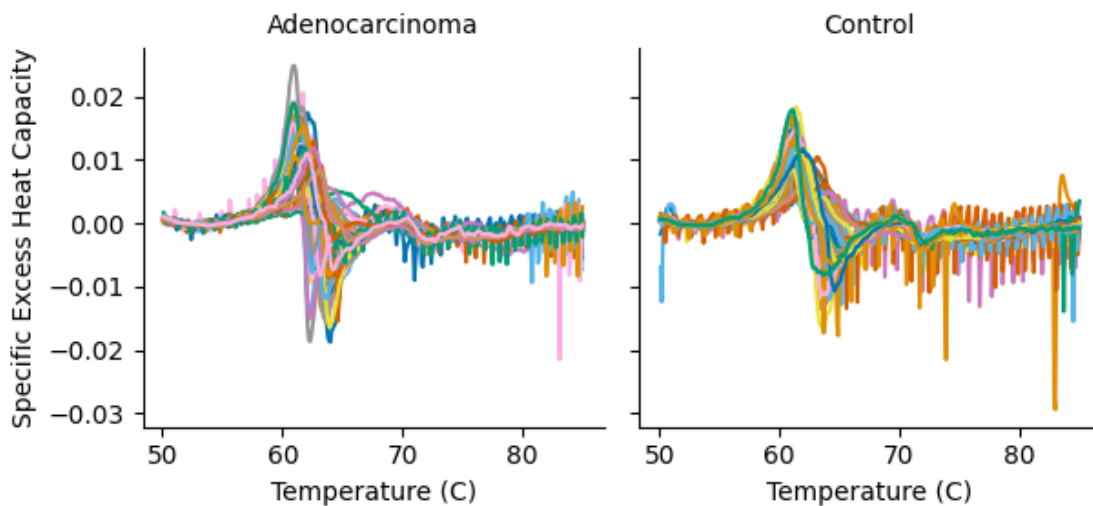
[ ]: df_tree_dif1 = df_tree.select_dtypes(include=['number']).diff(axis = 1)
non_numeric_columns = df_tree.select_dtypes(exclude=['number'])
df_tree_dif1 = pd.concat( [df_tree_dif1, non_numeric_columns], axis=1)
df_tree_dif1 = df_tree_dif1.iloc[:, 1:]

```

```
[ ]: df_long = pd.melt(df_tree_dif1, id_vars=['sample_id', 'pub_id', 'CancerType'],
    ↪var_name='temp', value_name='dsp' )
df_long['temp'] = df_long['temp'].str.replace('T', '').astype(float)

g = sns.FacetGrid(df_long, col="CancerType", col_wrap= 3, hue="sample_id",
    ↪palette = 'colorblind')
g.map_dataframe(sns.lineplot, x="temp", y="dsp")
g.set_axis_labels("Temperature (C)", "Specific Excess Heat Capacity")
g.set_titles(col_template="{col_name}")
```

```
[ ]: <seaborn.axisgrid.FacetGrid at 0x1c95c063130>
```



```
[ ]: df_tree_dif1 = df_tree.select_dtypes(include=['number']).diff(axis = 1)
df_tree_dif2 = df_tree_dif1.diff(axis = 1)
non_numeric_columns = df_tree.select_dtypes(exclude=['number'])
df_tree_dif2 = pd.concat( [df_tree_dif2, non_numeric_columns], axis=1)
df_tree_dif2 = df_tree_dif2.iloc[:, 2:]

#df_tree_dif2
```

```
[ ]:
      T50.2    T50.3    T50.4    T50.5    T50.6    T50.7    T50.8  \
0   -2.200000e-04 -0.000002  0.00025  0.00032 -0.00035 -0.00017 -0.00031
1   -1.400000e-04  0.00023  0.00005 -0.00052  0.00034 -0.00021 -0.00002
2   -2.600000e-04  0.00006 -0.00021  0.00003  0.00006 -0.00019  0.00007
3    1.700000e-04  0.00028 -0.00038 -0.00007 -0.00017 -0.00047  0.00024
4   -1.000000e-04 -0.00061 -0.00040  0.00054  0.00023 -0.00039  0.00016
..      ...      ...      ...      ...      ...      ...      ...
118  1.600000e-04 -0.00016 -0.00021 -0.00001 -0.00017 -0.00009 -0.00006
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119  1.000000e-05  0.00010  0.00001 -0.00003 -0.00007 -0.00002 -0.00006
120 -1.000000e-05  0.00002 -0.00010 -0.00010 -0.00006 -0.00003  0.00002
121  2.900000e-04 -0.00045 -0.00034 -0.00020 -0.00008 -0.00011 -0.00001
122  1.734723e-18 -0.00005 -0.00007 -0.00004 -0.00004 -0.00004 -0.00001

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      T50.9      T51      T51.1  ...      T84.4      T84.5  \
0   0.00012 -1.400000e-04 -0.00032  ...  0.000080  2.100000e-04
1   -0.00020  1.400000e-04 -0.00020  ... -0.000070  6.000000e-05
2   -0.00040 -8.000000e-05 -0.00014  ... -0.000062  1.090000e-04
3   -0.00018  1.100000e-04  0.00019  ... -0.000740  3.800000e-04
4   -0.00026 -4.100000e-04  0.00010  ... -0.000670 -5.400000e-04
..   ...      ...      ...  ...  ...      ...
118 -0.00006 -4.000000e-05 -0.00009  ...  0.000140  4.336809e-19
119 -0.00001 -6.000000e-05 -0.00008  ...  0.000080  7.000000e-05
120 -0.00008 -1.000000e-04 -0.00009  ...  0.000150  1.500000e-04
121 -0.00011 -1.734723e-18  0.00006  ... -0.000080 -1.300000e-04
122 -0.00009 -1.200000e-04 -0.00004  ...  0.000160  1.600000e-04

```

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      T84.6      T84.7      T84.8      T84.9      T85  sample_id  \
0  -6.100000e-05  0.000269  0.000072  6.280000e-04  0.000193    C67801
1   1.300000e-04  0.000220  0.000259  6.000000e-05  0.000573    C67901
2  -4.220000e-04  0.000920  0.000580 -4.336809e-19  0.000050    C55701
3  -2.600000e-04  0.000180  0.000480 -3.000000e-04  0.000460    C67201
4   1.381000e-03  0.001220  0.000022  2.713000e-04  0.000096    L15701
..   ...      ...      ...      ...      ...
118  1.700000e-04  0.000050  0.000330  1.550000e-04  0.000556        44
119  5.000000e-05  0.000200  0.000089  1.050000e-04  0.000501        45
120  1.700000e-04  0.000000 -0.000150 -5.000000e-05  0.000090        46
121 -8.673617e-19  0.000190  0.000080  5.000000e-05  0.000150        47
122  3.000000e-05 -0.000160  0.000020  2.000000e-05 -0.000090        48

```

```

      pub_id      CancerType
0         L14  Adenocarcinoma
1         L15  Adenocarcinoma
2          L4  Adenocarcinoma
3         L13  Adenocarcinoma
4         L83  Adenocarcinoma
..   ...      ...
118     C39      Control
119     C40      Control
120     C41      Control
121     C42      Control
122     C43      Control

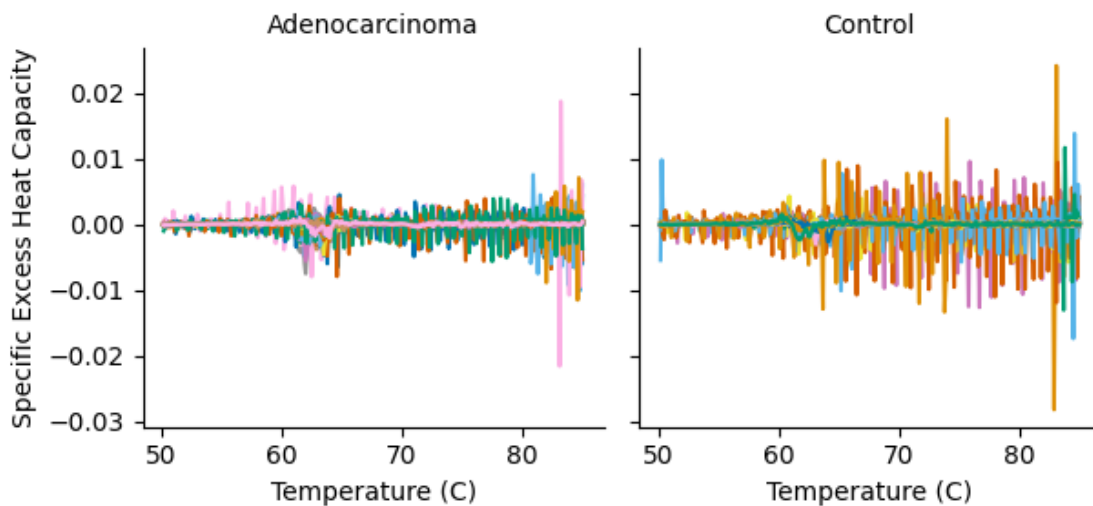
```

[123 rows x 352 columns]

```
[ ]: df_long = pd.melt(df_tree_dif2, id_vars=['sample_id', 'pub_id', 'CancerType'],
    ↪var_name='temp', value_name='dsp' )
df_long['temp'] = df_long['temp'].str.replace('T', '').astype(float)

g = sns.FacetGrid(df_long, col="CancerType", col_wrap= 3, hue="sample_id",
    ↪palette = 'colorblind')
g.map_dataframe(sns.lineplot, x="temp", y="dsp")
g.set_axis_labels("Temperature (C)", "Specific Excess Heat Capacity")
g.set_titles(col_template="{col_name}")
```

```
[ ]: <seaborn.axisgrid.FacetGrid at 0x1c95ef2bbe0>
```



```
[ ]: df_tree_dif1 = df_tree.select_dtypes(include=['number']).diff(axis = 1)
df_tree_dif2 = df_tree_dif1.diff(axis = 1)
df_tree_dif3 = df_tree_dif2.diff(axis = 1)
non_numeric_columns = df_tree.select_dtypes(exclude=['number'])
df_tree_dif3 = pd.concat( [df_tree_dif3, non_numeric_columns], axis=1)
df_tree_dif3 = df_tree_dif3.iloc[:, 3:]

#df_tree_dif3
```

```
[ ]:      T50.3    T50.4      T50.5      T50.6      T50.7    T50.8  \
0    0.00020  0.00027  7.000000e-05 -6.700000e-04  1.800000e-04 -0.00014
1    0.00037 -0.00018 -5.700000e-04  8.600000e-04 -5.500000e-04  0.00019
2    0.00032 -0.00027  2.400000e-04  3.000000e-05 -2.500000e-04  0.00026
3    0.00011 -0.00066  3.100000e-04 -1.000000e-04 -3.000000e-04  0.00071
4   -0.00051  0.00021  9.400000e-04 -3.100000e-04 -6.200000e-04  0.00055
..      ...      ...      ...      ...      ...      ...
```

```

118 -0.00032 -0.00005 2.000000e-04 -1.600000e-04 8.000000e-05 0.00003
119 0.00009 -0.00009 -4.000000e-05 -4.000000e-05 5.000000e-05 -0.00004
120 0.00003 -0.00012 -1.734723e-18 4.000000e-05 3.000000e-05 0.00005
121 -0.00074 0.00011 1.400000e-04 1.200000e-04 -3.000000e-05 0.00010
122 -0.00005 -0.00002 3.000000e-05 -1.734723e-18 -1.734723e-18 0.00003

```

```

      T50.9      T51      T51.1      T51.2 ...      T84.4      T84.5 \
0  4.300000e-04 -0.00026 -0.00018 0.00003 ... 0.000910 1.300000e-04
1 -1.800000e-04 0.00034 -0.00034 0.00057 ... 0.000440 1.300000e-04
2 -4.700000e-04 0.00032 -0.00006 0.00075 ... -0.000177 1.710000e-04
3 -4.200000e-04 0.00029 0.00008 -0.00033 ... -0.001170 1.120000e-03
4 -4.200000e-04 -0.00015 0.00051 0.00020 ... -0.000220 1.300000e-04
..      ...      ...      ...      ...      ...      ...
118 1.734723e-18 0.00002 -0.00005 0.00002 ... 0.000290 -1.400000e-04
119 5.000000e-05 -0.00005 -0.00002 0.00004 ... 0.000100 -1.000000e-05
120 -1.000000e-04 -0.00002 0.00001 0.00002 ... 0.000190 8.673617e-19
121 -1.000000e-04 0.00011 0.00006 -0.00004 ... -0.000030 -5.000000e-05
122 -8.000000e-05 -0.00003 0.00008 -0.00008 ... 0.000080 0.000000e+00

```

```

      T84.6      T84.7      T84.8      T84.9      T85 sample_id pub_id \
0 -0.000271 0.000330 -0.000197 5.560000e-04 -0.000435 C67801 L14
1 0.000070 0.000090 0.000039 -1.990000e-04 0.000513 C67901 L15
2 -0.000531 0.001342 -0.000340 -5.800000e-04 0.000050 C55701 L4
3 -0.000640 0.000440 0.000300 -7.800000e-04 0.000760 C67201 L13
4 0.001921 -0.000161 -0.001199 2.497000e-04 -0.000176 L15701 L83
..      ...      ...      ...      ...      ...
118 0.000170 -0.000120 0.000280 -1.750000e-04 0.000401 44 C39
119 -0.000020 0.000150 -0.000111 1.600000e-05 0.000396 45 C40
120 0.000020 -0.000170 -0.000150 1.000000e-04 0.000140 46 C41
121 0.000130 0.000190 -0.000110 -3.000000e-05 0.000100 47 C42
122 -0.000130 -0.000190 0.000180 -4.336809e-19 -0.000110 48 C43

```

```

      CancerType
0  Adenocarcinoma
1  Adenocarcinoma
2  Adenocarcinoma
3  Adenocarcinoma
4  Adenocarcinoma
..      ...
118      Control
119      Control
120      Control
121      Control
122      Control

```

[123 rows x 351 columns]

```
[ ]: df_long = pd.melt(df_tree_dif3, id_vars=['sample_id', 'pub_id', 'CancerType'],
    ↪var_name='temp', value_name='dsp' )
df_long['temp'] = df_long['temp'].str.replace('T', '').astype(float)

g = sns.FacetGrid(df_long, col="CancerType", col_wrap= 3, hue="sample_id",
    ↪palette = 'colorblind')
g.map_dataframe(sns.lineplot, x="temp", y="dsp")
g.set_axis_labels("Temperature (C)", "Specific Excess Heat Capacity")
g.set_titles(col_template="{col_name}")
```

```
[ ]: <seaborn.axisgrid.FacetGrid at 0x1c95f0e06d0>
```

