

notebook

January 19, 2026

```
[64]: import pandas as pd
import matplotlib.pyplot as plt
import matplotlib.colors as mcolors
```

```
[65]: df = pd.read_csv("data/Canadian_Health_Survey_Sample_cleaned.csv")
```

0.0.1 Analyzing dataset

```
[66]: df.head()
```

```
[66]:
```

	Province	Gender	Age	Income	BMI	PhysicalActivity \
0	Ontario	Female	59	80572.0	25.7	3.8
1	New Brunswick	Male	38	125739.0	24.2	3.4
2	Saskatchewan	Male	30	75947.0	28.3	3.9
3	Saskatchewan	Female	79	113966.0	31.2	1.6
4	Newfoundland and Labrador	Female	24	101828.0	27.8	4.5

	Smoking	SelfRatedHealth	StressLevel	BMI_Category	IncomeBracket \
0	Yes	Good	6	Overweight	80-110k
1	No	Good	9	Normal	>110k
2	No	Excellent	10	Overweight	50-80k
3	No	Fair	6	Obese	>110k
4	Yes	Good	5	Overweight	80-110k

	SelfRatedHealth_Num
0	3
1	3
2	5
3	2
4	3

```
[67]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 825 entries, 0 to 824
Data columns (total 12 columns):
#   Column                                Non-Null Count  Dtype
---  -

```

```

0    Province      825 non-null    object
1    Gender        825 non-null    object
2    Age           825 non-null    int64
3    Income        825 non-null    float64
4    BMI           825 non-null    float64
5    PhysicalActivity 825 non-null    float64
6    Smoking       825 non-null    object
7    SelfRatedHealth 825 non-null    object
8    StressLevel   825 non-null    int64
9    BMI_Category  825 non-null    object
10   IncomeBracket 825 non-null    object
11   SelfRatedHealth_Num 825 non-null    int64

```

dtypes: float64(3), int64(3), object(6)

memory usage: 77.5+ KB

```
[68]: df.describe(include='all')
```

```

[68]:      Province Gender      Age      Income      BMI \
count      825      825  825.000000    825.000000  825.000000
unique        10        2        NaN          NaN          NaN
top      Quebec   Male        NaN          NaN          NaN
freq         97      426        NaN          NaN          NaN
mean         NaN      NaN  49.408485   77545.210909   27.472039
std          NaN      NaN  18.187646   22145.238035    4.953750
min          NaN      NaN  18.000000    5000.000000   13.500000
25%          NaN      NaN  33.000000   64115.000000   24.200000
50%          NaN      NaN  49.000000   78130.000000   27.488861
75%          NaN      NaN  66.000000   90100.000000   30.700000
max          NaN      NaN  80.000000  250000.000000   60.000000

      PhysicalActivity Smoking SelfRatedHealth StressLevel BMI_Category \
count      825.000000      825          825    825.000000          825
unique          NaN        2          5          NaN            4
top          NaN       No      Good          NaN      Overweight
freq          NaN      640      254          NaN            333
mean         3.503246      NaN          NaN    5.294545          NaN
std         1.899122      NaN          NaN    2.121156          NaN
min         0.000000      NaN          NaN    1.000000          NaN
25%         2.100000      NaN          NaN    4.000000          NaN
50%         3.500000      NaN          NaN    5.000000          NaN
75%         4.700000      NaN          NaN    7.000000          NaN
max         10.400000      NaN          NaN   10.000000          NaN

      IncomeBracket SelfRatedHealth_Num
count          825      825.000000
unique          4          NaN
top       50-80k          NaN

```

freq	394	NaN
mean	NaN	3.208485
std	NaN	1.147042
min	NaN	1.000000
25%	NaN	2.000000
50%	NaN	3.000000
75%	NaN	4.000000
max	NaN	5.000000

0.0.2 Cleaning & Standardizing Categories

- Normalizing gender values and smoking values

```
[69]: # normalizing gender values to Male or Female
df['Gender'] = df['Gender'].str.lower()
df['Gender'] = df['Gender'].replace({
    'male': 'Male',
    'm': 'Male',
    'female': 'Female',
    'f': 'Female'
})

# normalizing smoking to Yes or No
df['Smoking'] = df['Smoking'].str.lower()
df['Smoking'] = df['Smoking'].replace({
    'yes': 'Yes',
    'y': 'Yes',
    'no': 'No',
    'n': 'No'
})
```

0.0.3 Data Cleaning

- Filled missing BMI and Income values using averages (mean/median) to preserve dataset size without distorting quantitative distributions.
- Dropped rows with missing StressLevel because it is ordinal, and imputing an ordered scale would distort the meaning of the categories.
- Filled missing PhysicalActivity values using the mean, since it is a discrete numeric variable and will not affect categorical encodings in later visualizations.

```
[70]: df['BMI'] = df['BMI'].fillna(df['BMI'].mean())
df['Income'] = df['Income'].fillna(df['Income'].median())
df = df.dropna(subset=['StressLevel'])
df['PhysicalActivity'] = df['PhysicalActivity'].fillna(df['PhysicalActivity'].
    ↪mean())
```

0.1 Enforcing Correct Data Types

```
[71]: df['Age'] = df['Age'].astype(int)
df['Income'] = df['Income'].astype(float)
df['BMI'] = df['BMI'].astype(float)
df['StressLevel'] = df['StressLevel'].astype(int)
```

0.1.1 Enhancing the Dataset with Derived Features

```
[72]: # create new column
df['BMI_Category'] = None

# create column values based on constraints
df.loc[df['BMI'] < 18.5, 'BMI_Category'] = 'Underweight'
df.loc[(df['BMI'] >= 18.5) & (df['BMI'] <= 24.9), 'BMI_Category'] = 'Normal'
df.loc[(df['BMI'] >= 25) & (df['BMI'] <= 29.9), 'BMI_Category'] = 'Overweight'
df.loc[df['BMI'] >= 30, 'BMI_Category'] = 'Obese'

# create new column
df['IncomeBracket'] = None

# create column values based on constraints
df.loc[df['Income'] < 50000, 'IncomeBracket'] = '<50k'
df.loc[(df['Income'] >= 50000) & (df['Income'] < 80000), 'IncomeBracket'] = '50-80k'
df.loc[(df['Income'] >= 80000) & (df['Income'] < 110000), 'IncomeBracket'] = '80-110k'
df.loc[df['Income'] >= 110000, 'IncomeBracket'] = '>110k'
```

0.1.2 Bar Chart - Average Life/Health Proxy by Province Analysis

- This bar chart is created to compare the average self-rated health score across provinces.

```
[73]: # create new column
df['SelfRatedHealth_Num'] = None

# create column values based on constraints
df.loc[df['SelfRatedHealth'] == 'Poor', 'SelfRatedHealth_Num'] = 1
df.loc[df['SelfRatedHealth'] == 'Fair', 'SelfRatedHealth_Num'] = 2
df.loc[df['SelfRatedHealth'] == 'Good', 'SelfRatedHealth_Num'] = 3
df.loc[df['SelfRatedHealth'] == 'Very Good', 'SelfRatedHealth_Num'] = 4
df.loc[df['SelfRatedHealth'] == 'Excellent', 'SelfRatedHealth_Num'] = 5

# group each province
groups = df.groupby('Province')

# get the average score for each province
avg_score = groups['SelfRatedHealth_Num'].mean()
```

```

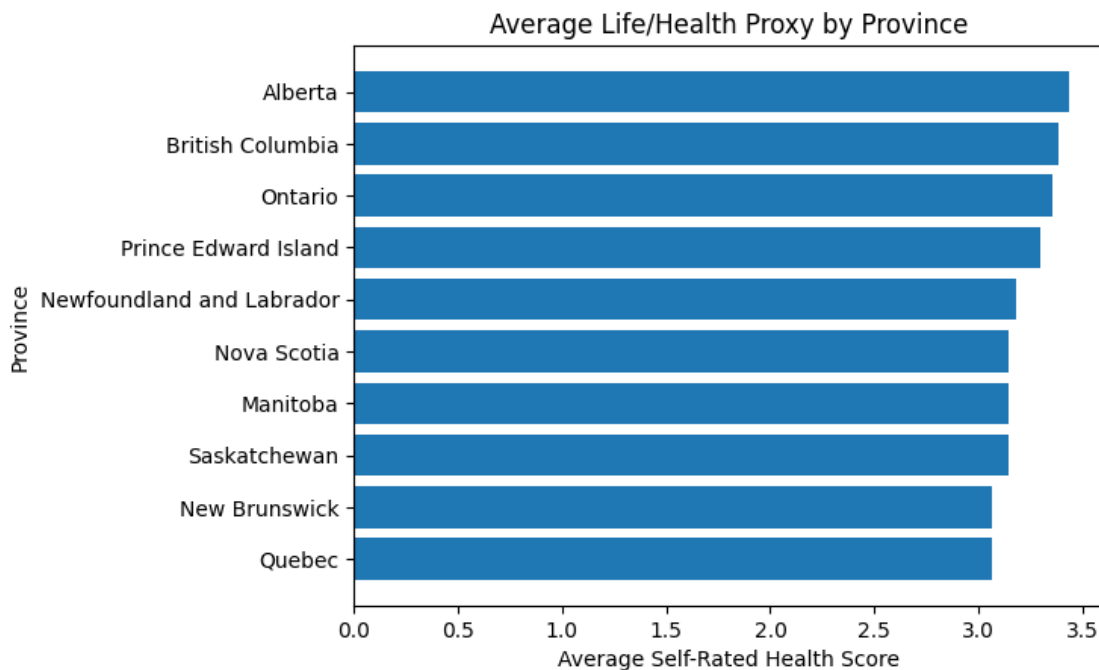
avg_score = avg_score.sort_values() # better visualization

# plot
plt.barh(avg_score.index, avg_score.values)

plt.xlabel('Average Self-Rated Health Score')
plt.ylabel('Province')
plt.title('Average Life/Health Proxy by Province')

plt.show()

```



0.1.3 Scatter Plot - Income vs BMI Analysis

- This scatter plot is created to examine the relationship between Income and BMI while also showing how StressLevel and Gender influence the pattern.

```

[74]: # create two new dataframes only containing the gender column where one is only
      ↪ males and the other is only females
male_df = df[df['Gender'] == 'Male']
female_df = df[df['Gender'] == 'Female']

# for males
plt.scatter(
    male_df['Income'],
    male_df['BMI'],

```

```

    c = male_df['StressLevel'], # colormap with stress level
    cmap = 'viridis',
    marker = 'o', # circle for males
    label = 'Male' # for legend
)

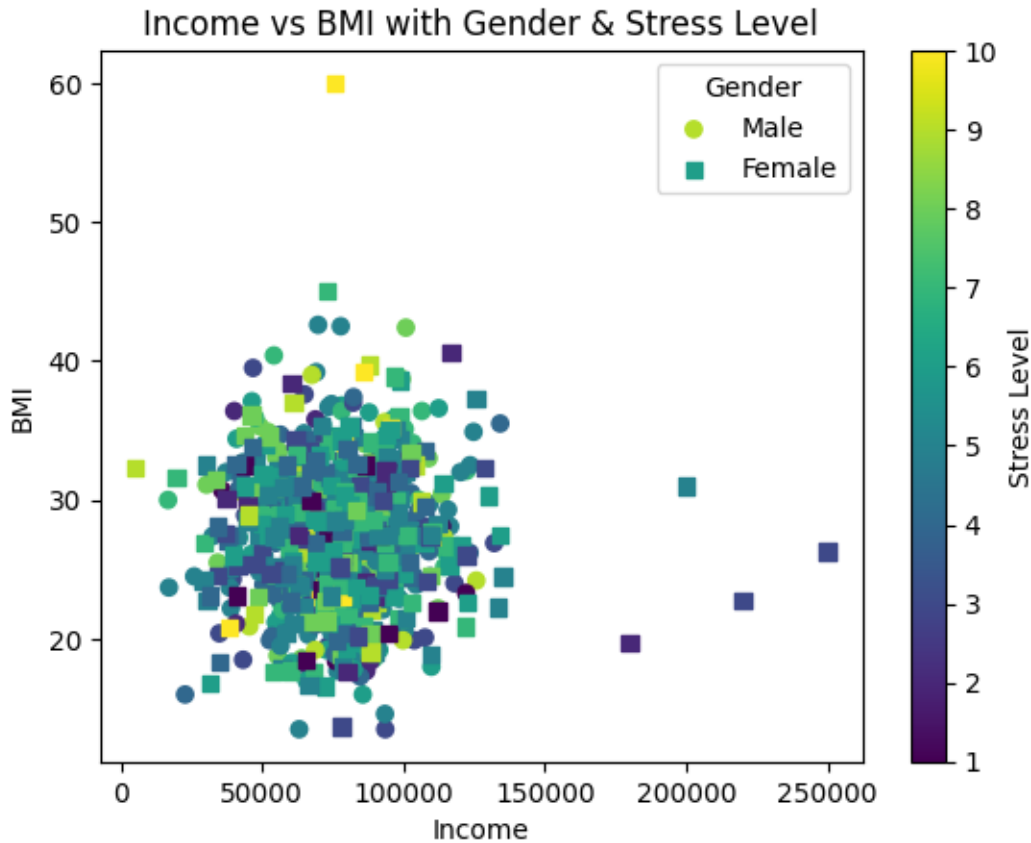
# for females
plt.scatter(
    female_df['Income'],
    female_df['BMI'],
    c = female_df['StressLevel'], # colormap with stress level
    cmap = 'viridis',
    marker = 's', # squares for females
    label = 'Female' # for legend
)

# Colorbar for StressLevel
colorbar = plt.colorbar()
colorbar.set_label('Stress Level')

plt.xlabel('Income')
plt.ylabel('BMI')
plt.title('Income vs BMI with Gender & Stress Level')

plt.legend(title='Gender')
plt.show()

```



0.1.4 Diverging Bar Chart - Life Proxy Difference by Province Analysis

- This diverging bar chart is used to show how each province's average self-rated health score differs from the overall national average.
- The values can be both above and below zero, a diverging visualization is the correct choice as it highlights positive vs. negative deviation around a meaningful midpoint.

```
[75]: groups = df.groupby('Province')

# get the average score for each province
province_avg = groups['SelfRatedHealth_Num'].mean()

# get the overall average
ovr_avg = df['SelfRatedHealth_Num'].mean()

life_proxy = province_avg - ovr_avg
life_proxy = life_proxy.sort_values()
life_proxy = life_proxy.astype(float)

# color map
```

```

cmap = plt.cm.coolwarm # choose coolwarm color map

low = life_proxy.min()
high = life_proxy.max()
center = 0

color_map = mcolors.TwoSlopeNorm(vmin = low, vcenter = center, vmax = high)
colors = cmap(color_map(life_proxy.values)) # mapping each value to a color

plt.barh(life_proxy.index, life_proxy.values, color = colors)

plt.xlabel('Δ Life Proxy')
plt.ylabel('Province')
plt.title('Δ Life Proxy by Province')

plt.show()

```

