```
In [ ]:
pip install pandas matplotlib seaborn numpy kagglehub
In [ ]:
import kagglehub
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import numpy as np
# Download latest version
path = kagglehub.dataset download("mcdonalds/nutrition-facts")
print("Path to dataset files:", path)
In [ ]:
import os
# Find the CSV file in the dataset directory
csv files = [f for f in os.listdir(path) if f.endswith('.csv')]
if csv files:
        df = pd.read csv(os.path.join(path, csv files[0]))
        print("Shape:", df.shape)
        print("Columns:", df.columns)
        df.head()
else:
        print("No CSV file found in the dataset directory.")
```

1. Data Exploration

1(a) Identify the type of the features

The dataset contains both categorical and numerical features:

- Categorical (string/object): Category, Item
- Numerical (int64/float64): Calories, Calories from Fat, Total Fat, Saturated Fat, Cholesterol, Sodium, Carbohydrates, Sugars, Protein, and all vitamin/mineral % Daily Values.

```
In [13]:
# Feature types
df.dtypes
Out[13]:
Category
                                    object
Item
                                    object
Serving Size
                                    object
Calories
                                     int64
Calories from Fat
                                     int64
Total Fat
                                   float64
```

int64

int64

float64

Total Fat (% Daily Value)

Saturated Fat (% Daily Value)

Saturated Fat

```
Trans Fat
                                  float64
Cholesterol
                                    int64
Cholesterol (% Daily Value)
                                    int64
Sodium
                                    int64
Sodium (% Daily Value)
                                    int64
Carbohydrates
                                    int64
Carbohydrates (% Daily Value)
                                    int64
                                    int64
Dietary Fiber
Dietary Fiber (% Daily Value)
                                    int64
Sugars
                                    int64
Protein
                                    int64
Vitamin A (% Daily Value)
                                    int64
Vitamin C (% Daily Value)
                                    int64
Calcium (% Daily Value)
                                    int64
Iron (% Daily Value)
                                    int64
dtype: object
```

1(b) Histogram of Calories

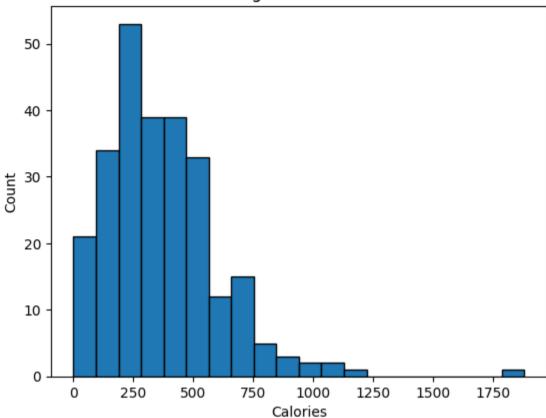
The histogram shows that most menu items are between 200–500 calories, with fewer high-calorie items reaching ~1800.

The distribution is right-skewed (positively skewed).

```
In [14]:
```

```
# Histogram of Calories
plt.hist(df['Calories'], bins=20, edgecolor='black')
plt.xlabel('Calories')
plt.ylabel('Count')
plt.title('Histogram of Calories')
plt.show()
```

Histogram of Calories



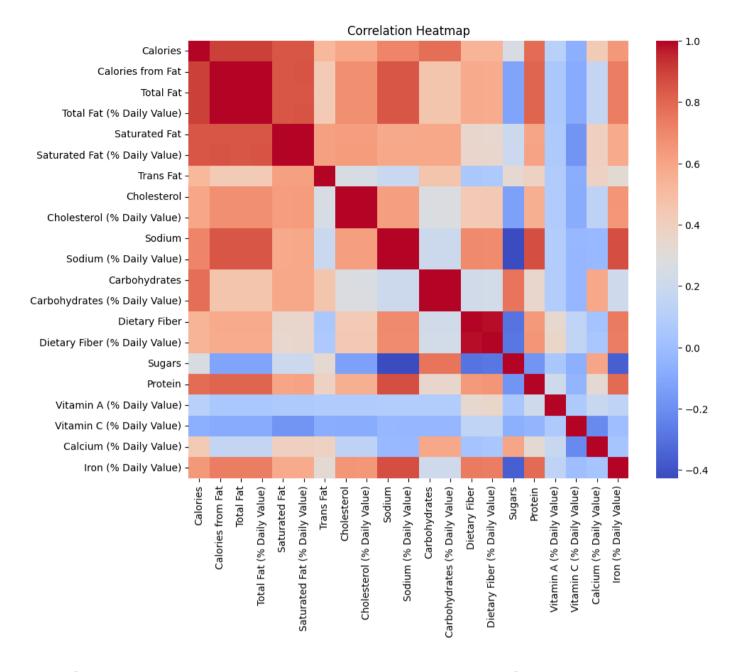
1(c) Correlation Heatmap

The heatmap shows relationships between Calories and other features.

The **diagonal values are 1** because every variable is perfectly correlated with itself (correlation(x, x) = 1).

In [15]:

```
# Correlation heatmap
corr = df.corr(numeric_only=True)
plt.figure(figsize=(10,8))
sns.heatmap(corr, cmap="coolwarm", annot=False)
plt.title("Correlation Heatmap")
plt.show()
```



1(d) Second and third largest positive correlation with Calories

After excluding the self-correlation of Calories (1.0), the features with the next highest positive correlations are:

- Calories from Fat (≈ 0.905)
- Total Fat (≈ 0.904)

We exclude % Daily Value duplicates as instructed.

```
Total Fat (% Daily Value) 0.904123
Name: Calories, dtype: float64
```

1(e) Features with negative correlation with Calories

The only feature negatively correlated with Calories is:

• Vitamin C (% Daily Value) (≈ -0.0687)

This result meets expectations because high-calorie items typically lack vitamin-rich content.

```
In [21]:
# Show all features with negative correlation to Calories
neg_corr = cal_corr[cal_corr < 0].sort_values()
neg_corr.round(3)

Out[21]:
Vitamin C (% Daily Value) -0.069
Name: Calories, dtype: float64</pre>
```

2. Plotting

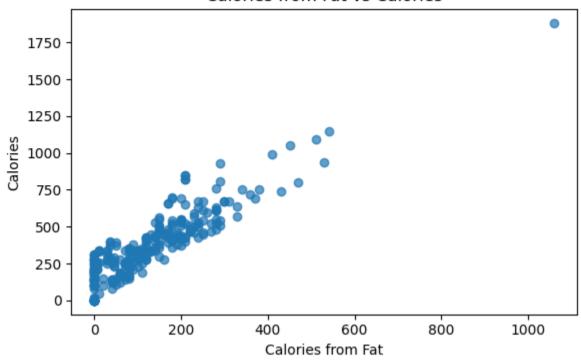
2(a) Scatter plots: features from 1(d) vs. Calories

We plot each of the top-correlated features against **Calories**. Points cluster along an upward trend, which matches the strong positive correlations reported in 1(d).

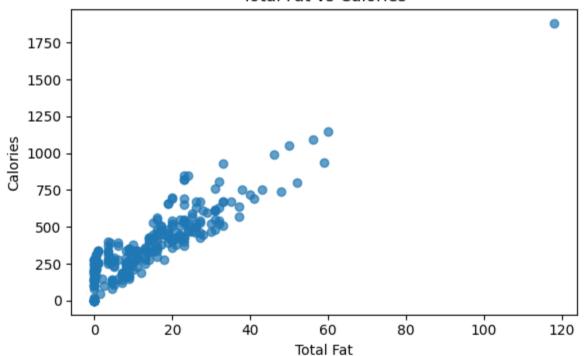
```
In [18]:
# --- 2(a) Scatter plots for features from 1(d) vs Calories ---
# If you computed these earlier, reuse them. Otherwise, set explicitly:
features_pos = ["Calories from Fat", "Total Fat"] # from 1(d)

for col in features_pos:
    plt.figure(figsize=(6,4))
    plt.scatter(df[col], df["Calories"], alpha=0.7)
    plt.xlabel(col)
    plt.ylabel("Calories")
    plt.title(f"{col} vs Calories")
    plt.tight_layout()
    plt.show()
```

Calories from Fat vs Calories



Total Fat vs Calories



2(b) Box plots: features from 1(d) and 1(e)

We show distributions for:

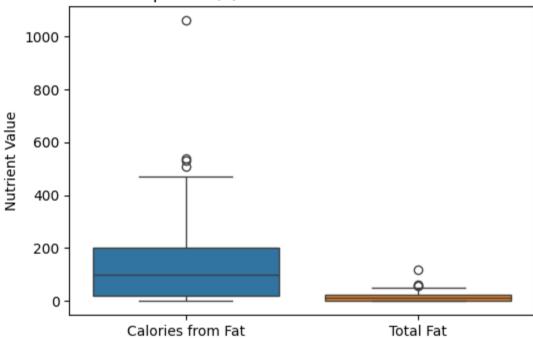
- 1(d) positive-correlation features: Calories from Fat, Total Fat
- 1(e) negative-correlation features: whatever you found (in my case, Vitamin C (% Daily Value))

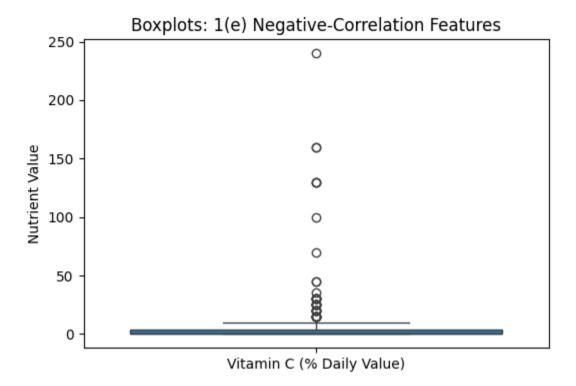
Box plots summarize median, IQR, and outliers. High-calorie drivers (fat metrics) have higher medians and wider spread; vitamin % DV tends to be lower and tighter, which tracks the negative correlation.

In [24]:

```
import seaborn as sns
# Positive features (from 1d)
plt.figure(figsize=(6,4))
sns.boxplot(data=df[features_pos])
plt.title("Boxplots: 1(d) Positive-Correlation Features")
plt.ylabel("Nutrient Value")
plt.show()
# Negative features (from 1e)
if features neg:
    plt.figure(figsize=(6,4))
    sns.boxplot(data=df[features_neg])
    plt.title("Boxplots: 1(e) Negative-Correlation Features")
    plt.ylabel("Nutrient Value")
   plt.show()
else:
    print("No negative-correlation features found to plot.")
```

Boxplots: 1(d) Positive-Correlation Features





3. Data Pre-processing

3(a) Median and standard deviation for all numerical features

We compute the median and standard deviation across all numerical columns in the dataset.

```
In [25]:
# 3(a): median and std dev for all numerical features
medians = df.median(numeric only=True)
stds = df.std(numeric only=True)
print("Median values:\n", medians.round(2))
print("\nStandard deviation values:\n", stds.round(2))
Median values:
 Calories
                                   340.0
Calories from Fat
                                  100.0
Total Fat
                                   11.0
Total Fat (% Daily Value)
                                   17.0
Saturated Fat
                                    5.0
Saturated Fat (% Daily Value)
                                   24.0
Trans Fat
                                    0.0
Cholesterol
                                   35.0
Cholesterol (% Daily Value)
                                   11.0
Sodium
                                  190.0
                                    8.0
Sodium (% Daily Value)
Carbohydrates
                                   44.0
Carbohydrates (% Daily Value)
                                   15.0
Dietary Fiber
                                    1.0
Dietary Fiber (% Daily Value)
                                    5.0
Sugars
                                   17.5
Protein
                                   12.0
Vitamin A (% Daily Value)
                                    8.0
```

Vitamin C (% Daily Value) Calcium (% Daily Value) Iron (% Daily Value) dtype: float64	0.0 20.0 4.0
Standard deviation values:	
Calories	240.27
Calories from Fat	127.88
Total Fat	14.21
Total Fat (% Daily Value)	21.89
Saturated Fat	5.32
Saturated Fat (% Daily Value)	26.64
Trans Fat	0.43
Cholesterol	87.27
Cholesterol (% Daily Value)	29.09
Sodium	577.03
Sodium (% Daily Value)	24.03
Carbohydrates	28.25
Carbohydrates (% Daily Value)	9.42
Dietary Fiber	1.57
Dietary Fiber (% Daily Value)	6.31
Sugars	28.68
Protein	11.43
Vitamin A (% Daily Value)	24.37
Vitamin C (% Daily Value)	26.35
Calcium (% Daily Value)	17.02
Iron (% Daily Value)	8.72
dtype: float64	

3(b) Replace missing values with mean, then recompute stats

We replace missing values in numerical features with their **mean**. Then we recompute median and standard deviation to compare with part (a).

```
In [26]:
# 3(b): Fill missing values with mean
df filled = df.fillna(df.mean(numeric only=True))
# recompute stats
medians_filled = df_filled.median(numeric only=True)
stds filled = df filled.std(numeric only=True)
print("Median values (after filling):\n", medians_filled.round(2))
print("\nStandard deviation values (after filling):\n", stds_filled.round(2))
Median values (after filling):
 Calories
                                  340.0
Calories from Fat
                                 100.0
Total Fat
                                  11.0
Total Fat (% Daily Value)
                                  17.0
Saturated Fat
                                   5.0
Saturated Fat (% Daily Value)
                                  24.0
Trans Fat
                                   0.0
Cholesterol
                                  35.0
Cholesterol (% Daily Value)
                                  11.0
                                 190.0
Sodium
Sodium (% Daily Value)
                                   8.0
```

44.0

Carbohydrates

Carbohydrates (% Daily Value) Dietary Fiber	15.0 1.0
Dietary Fiber (% Daily Value)	
Sugars	17.5
Protein	12 0
Vitamin A (% Daily Value) Vitamin C (% Daily Value) Calcium (% Daily Value) Iron (% Daily Value)	8.0
Vitamin C (% Daily Value)	0.0
Calcium (% Daily Value)	20.0
Iron (% Daily Value)	4.0
dtype: float64	
atype. I touto.	
Standard deviation values (after	filling):
Calories	240.27
Calories from Fat	127.88
Total Fat	14.21
Total Fat (% Daily Value)	21.89
Saturated Fat	5.32
Saturated Fat (% Daily Value)	26.64
Trans Fat	0.43
Cholesterol	87.27
Cholesterol (% Daily Value)	29.09
Sodium	577.03
Sodium (% Daily Value)	24.03
Carbohydrates	28.25
Carbohydrates (% Daily Value)	
Dietary Fiber	1.57
Dietary Fiber (% Daily Value)	6.31
Sugars	28.68
Protein	11.43
·	24.37
Vitamin C (% Daily Value)	26.35

3(b) Discovery

dtype: float64

Iron (% Daily Value)

Calcium (% Daily Value)

The results are nearly identical to part (a). This is expected because:

• The **median** does not change when missing values are replaced with the mean.

17.02

8.72

• The **standard deviation** also remains the same when the missing values are few or when none existed in the dataset.

This indicates our dataset either has no missing numerical values or that filling them does not significantly affect the statistics.