

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import warnings
import matplotlib.patches as mpatches
import matplotlib.lines as mlines
from matplotlib.patches import FancyBboxPatch, FancyArrowPatch
import matplotlib.gridspec as gridspec
warnings.filterwarnings('ignore')
```

```
!wget -q --show-progress -O food_facts.csv.gz \
"https://static.openfoodfacts.org/data/en.openfoodfacts.org.produ
print("Download complete")

food_facts.csv.gz 100%[=====>] 1.14G 32.8MB/s
Download complete
```

```
# The full dataset has 180+ columns. We only need these 8.
COLUMNS_NEEDED = [
    'product_name',
    'categories_tags',
    'ingredients_text',
    'energy_100g',
    'sugars_100g',
    'proteins_100g',
    'fat_100g',
    'fiber_100g',
]
print(f"Loaded {len(COLUMNS_NEEDED)} columns out of 180+")

Loaded 8 columns out of 180+
```

```

df_raw = pd.read_csv(
    'food_facts.csv.gz',
    sep='\t',
    nrows=500_000,
    usecols=COLUMNS_NEEDED,
    low_memory=False,
    on_bad_lines='skip'
)

print(f"Loaded {len(df_raw)} rows")
print(f"Shape: {df_raw.shape}")
df_raw.tail(5)

```

Loaded 500,000 rows  
 Shape: (500000, 8)

	product_name	categories_tags	ingredients_text	energy
499995	Orange Juice		NaN	orange juice
499996	Orange Juice Medium Pulp		NaN	NaN
499997	Black sweet tea	en:plant-based-foods-and-beverages,en:beverage...		NaN
499998	Homestyle Lemonade		NaN	NaN
499999	Large Grade A Vegetarian Fed Cage Free Eggs	en:farming-products,en:eggs,en:chicken-eggs,en...		NaN

## # Data Exploration

```

print(" DATASET OVERVIEW \n")
print(f"Total rows: {len(df_raw)}")
print(f"Total columns: {df_raw.shape[1]}")
print("\n NULL VALUE COUNT (per column) ")
print(df_raw.isnull().sum())
print("\n DATA TYPES ")
print(df_raw.dtypes)
print("\n BASIC STATISTICS (numeric columns) ")
df_raw[['sugars_100g', 'proteins_100g', 'fat_100g', 'fiber_100g']]

```

DATASET OVERVIEW

Total rows: 500,000

```
Total columns: 8
```

```
NULL VALUE COUNT (per column)
product_name          15579
categories_tags       229747
ingredients_text      231191
energy_100g           74707
fat_100g              77345
sugars_100g           93436
fiber_100g            177542
proteins_100g         76871
dtype: int64
```

```
DATA TYPES
product_name          object
categories_tags        object
ingredients_text       object
energy_100g            float64
fat_100g               float64
sugars_100g            float64
fiber_100g             float64
proteins_100g          float64
dtype: object
```

```
BASIC STATISTICS (numeric columns)
```

	sugars_100g	proteins_100g	fat_100g	fiber_100g	grid icon
<b>count</b>	406564.000000	423129.000000	422655.000000	322458.000000	
<b>mean</b>	17.542264	8.657712	14.869983	3.252585	
<b>std</b>	43.720976	23.888684	41.107267	25.655246	
<b>min</b>	-1.450000	-12.600000	0.000000	-44.700000	
<b>25%</b>	1.000000	0.770000	0.000000	0.000000	
<b>50%</b>	5.000000	5.000000	5.260000	1.400000	
<b>75%</b>	22.200000	10.800000	20.000000	3.571429	
<b>max</b>	11111.000000	7270.000000	11111.000000	11800.000000	

```
df_clean = df_raw.dropna(subset=['product_name', 'sugars_100g', 'pr  
rows_dropped = len(df_raw) - len(df_clean)  
print(f"Rows before cleaning: {len(df_raw):,}")  
print(f"Rows dropped (missing critical values): {rows_dropped:,}")  
print(f"Rows remaining: {len(df_clean):,}")
```

```
Rows before cleaning: 500,000  
Rows dropped (missing critical values): 98,913  
Rows remaining: 401,087
```

```
# Define valid ranges  
VALID_MIN = 0  
VALID_MAX = 100  
  
df_clean = df_clean[  
    (df_clean['sugars_100g']    >= VALID_MIN) & (df_clean['sugars_100g'] <= VALID_MAX)  
    (df_clean['proteins_100g'] >= VALID_MIN) & (df_clean['proteins_100g'] <= VALID_MAX)  
    (df_clean['fat_100g'].isna() | ((df_clean['fat_100g'] >= VALID_MIN) & (df_clean['fat_100g'] <= VALID_MAX))  
    (df_clean['fiber_100g'].isna() | ((df_clean['fiber_100g'] >= VALID_MIN) & (df_clean['fiber_100g'] <= VALID_MAX))  
]  
  
print(f"Rows after removing impossible values: {len(df_clean):,}")
```

```
Rows after removing impossible values: 391,528
```

```
# clean data

print(" CLEAN DATASET STATS \n")
print(df_clean[['sugars_100g', 'proteins_100g', 'fat_100g', 'fiber_100g']])

print("\n SAMPLE PRODUCTS ")
print(df_clean[['product_name', 'sugars_100g', 'proteins_100g']].sample(10))

df_clean.to_csv('food_facts_clean.csv', index=False)
print("\nClean dataset saved as food_facts_clean.csv")
print(f"Final shape: {df_clean.shape}")
```

### CLEAN DATASET STATS

	sugars_100g	proteins_100g	fat_100g	fiber_100g
count	391528.00	391528.00	390603.00	307274.00
mean	14.79	7.72	11.93	2.83
std	20.52	9.84	16.07	5.17
min	0.00	0.00	0.00	0.00
25%	1.10	0.83	0.00	0.00
50%	5.00	4.93	5.00	1.40
75%	21.25	10.53	18.75	3.57
max	100.00	100.00	100.00	100.00

### SAMPLE PRODUCTS

	product_name	sugars_100g	proteins_100g
316270	Breaded squid rings	1.000000	1.000000
130944	100% Whole Grain Rolled Oats	1.000000	1.000000
264987	Old Fashioned Grape seed oil	0.000000	0.000000
20828	Maple Syrup Sachet	66.700000	66.700000
449521	Taro Cake with Dried Shrimp	1.330000	1.330000
389545	MESQUITE BARBECUE	7.142857	7.142857
120966	Buttermilk Fudge Stripes	9.000000	9.000000
457890	Energy Drink Mix	0.000000	0.000000
457709	Cut Green Beans	0.830000	0.830000
441208	Cookie dough snack-pack	37.500000	37.500000

Clean dataset saved as food\_facts\_clean.csv  
 Final shape: (391528, 8)

```
print(" SAMPLE OF RAW categories_tags VALUES \n")

# Show 100 examples
sample_tags = df_clean['categories_tags'].dropna().sample(100, random_state=42)
for tag in sample_tags:
    print(tag)
    print("---")
```

---

```
en:snacks,en:sweet-snacks,en:confectioneries,en:candies,en:gummi-c  
---  
en:snacks  
---  
en:plant-based-foods-and-beverages,en:plant-based-foods,en:cereals  
---  
en:condiments,en:sauces,en:dips  
---  
en:desserts  
---  
en:condiments,en:sauces  
---  
en:plant-based-foods-and-beverages,en:beverages,en:plant-based-bev  
---  
en:dairies,en:fermented-foods,en:fermented-milk-products,en:desser  
---  
en:syrups  
---  
en:undefined  
---  
en:condiments,en:sauces,en:groceries  
---  
en:undefined  
---  
en:plant-based-foods-and-beverages,en:dairy-substitutes,en:milk-su  
---  
en:snacks,en:sweet-snacks,en:biscuits-and-cakes,en:cakes  
---  
en:condiments,en:sauces,en:groceries  
---  
en:plant-based-foods-and-beverages,en:plant-based-foods,en:cereals  
---  
en:snacks,en:sweet-snacks,en:biscuits-and-cakes,en:pastries  
---  
en:beverages-and-beverages-preparations,en:beverages,en:carbonated  
---  
en:undefined  
---  
en:meats-and-their-products,en:meats,en:chicken-and-its-products,e  
---  
en:dairies,en:fermented-foods,en:fermented-milk-products,en:desser  
---  
en:desserts,en:frozen-foods,en:frozen-desserts,en:ice-creams-and-s  
---  
en:snacks,en:sweet-snacks,en:biscuits-and-cakes  
---  
en:plant-based-foods-and-beverages,en:plant-based-foods,en:legumes  
---  
en:beverages,en:carbonated-drinks,en:sodas,en:sweetened-beverages  
---  
en:snacks,en:sweet-snacks,en:biscuits-and-cakes,en:pastries  
---  
en:meals,en:rice-dishes
```

```
---  
en:dairies,en:milks  
---
```

```
# Step 1: Define what to EXCLUDE first  
EXCLUDE_PATTERNS = [  
    'beverage',  
    'juice',  
    'soda',  
    'milk',  
    'cream',  
    'ice-cream',  
    'frozen-dessert',  
    'frozen-food',  
    'mochi-ice',  
    'sorbet',  
    'soup',  
    'sauce',  
    'condiment',  
    'pasta',  
    'rice-dish',  
    'meal',  
    'bread',  
    'cereal',  
    'muesli',  
    'supplement',  
    'syrup',  
    'honey',  
    'spread',  
    'pizza',  
    'meat',  
    'chicken',  
    'vegetable',  
    'tomato',  
    'legume',  
    'canned-bean',  
]  
  
# Step 2: Define SNACK category mapping (priority ordered – first is highest priority)  
CATEGORY_MAP = [  
    # Protein / Fitness Bars – most specific, highest priority  
    ('protein-bar', 'Protein & Fitness Bars'),  
    ('energy-bar', 'Protein & Fitness Bars'),  
    ('cereal-bar', 'Protein & Fitness Bars'),  
    ('granola-bar', 'Protein & Fitness Bars'),  
    ('nutrition-bar', 'Protein & Fitness Bars'),
```

```
('sport', 'Protein & Fitness Bars'),  
  
# Chips & Savory Snacks  
('chips-and-fries', 'Chips & Savory Snacks'),  
('potato-crisp', 'Chips & Savory Snacks'),  
('crisps', 'Chips & Savory Snacks'),  
('bbq-chip', 'Chips & Savory Snacks'),  
('chip', 'Chips & Savory Snacks'),  
('pretzel', 'Chips & Savory Snacks'),  
('popcorn', 'Chips & Savory Snacks'),  
('puffed', 'Chips & Savory Snacks'),  
('salty-snack', 'Chips & Savory Snacks'),  
('salted-snack', 'Chips & Savory Snacks'),  
('savory-snack', 'Chips & Savory Snacks'),  
('crackers-appetizer', 'Chips & Savory Snacks'),  
('cracker', 'Chips & Savory Snacks'),  
  
# Cookies & Biscuits  
('biscuits-and-cakes', 'Cookies & Biscuits'),  
('biscuit', 'Cookies & Biscuits'),  
('cookie', 'Cookies & Biscuits'),  
('wafer', 'Cookies & Biscuits'),  
('brownie', 'Cookies & Biscuits'),  
('toaster-pastri', 'Cookies & Biscuits'),  
('pastri', 'Cookies & Biscuits'),  
  
# Candy & Confectionery  
('confectioner', 'Candy & Confectionery'),  
('candies', 'Candy & Confectionery'),  
('candy', 'Candy & Confectionery'),  
('gummi', 'Candy & Confectionery'),  
('gummy', 'Candy & Confectionery'),  
('jelly-bean', 'Candy & Confectionery'),  
('lollipop', 'Candy & Confectionery'),  
('caramel', 'Candy & Confectionery'),  
('chocolate', 'Candy & Confectionery'),  
('sweet-snack', 'Candy & Confectionery'),  
  
# Nuts & Seeds  
('roasted-nut', 'Nuts & Seeds'),  
('roasted-peanut', 'Nuts & Seeds'),  
('trail-mix', 'Nuts & Seeds'),  
('nut', 'Nuts & Seeds'),  
('seed', 'Nuts & Seeds'),  
('almond', 'Nuts & Seeds'),  
('peanut', 'Nuts & Seeds'),  
('cashew', 'Nuts & Seeds'),
```

```

# Dairy Snacks
('yogurt',                               'Dairy & Yogurt Snacks'),
('yoghurt',                               'Dairy & Yogurt Snacks'),
('cheese-snack',                          'Dairy & Yogurt Snacks'),
('dairy-snack',                           'Dairy & Yogurt Snacks'),


# Fruit & Vegetable Snacks (room temp only – dried/packaged)
('dried-fruit',                           'Fruit & Veg Snacks'),
('fruit-snack',                           'Fruit & Veg Snacks'),
('fruit-bar',                             'Fruit & Veg Snacks'),
('fruit-based-food',                     'Fruit & Veg Snacks'),
('fruits-based-food',                    'Fruit & Veg Snacks'),
('vegetable-chip',                        'Fruit & Veg Snacks'),
('dried-vegetable',                       'Fruit & Veg Snacks'),


# General Snacks – broad catch-all, always last
('snack',                                 'General Snacks'),
]

print(f" Exclusion patterns defined: {len(EXCLUDE_PATTERNS)}")
print(f" Category rules defined: {len(CATEGORY_MAP)}")
print(f" Covering {len(set(v for _, v in CATEGORY_MAP))} unique cat

```

Exclusion patterns defined: 30  
 Category rules defined: 56  
 Covering 8 unique categories

```

def assign_category(tags_string):
    """
    Takes a raw tags string, returns a clean category label or None

    Logic:
    1. Null/undefined tags → None (skip)
    2. Matches an exclusion pattern → None (not a snack)
    3. Matches a category keyword → return that category
    4. No match → None (uncategorized, skip)
    """

    # Guard: null values
    if pd.isna(tags_string):
        return None

    tags_lower = tags_string.lower()

    # Guard: garbage tags like en:null, en:undefined
    if tags_lower.strip() in ('en:null', 'en:undefined', 'en:null',
        return None

```

```

# Step 1: Check exclusions FIRST
# If this product matches any exclusion pattern, it's not a snack
for exclude in EXCLUDE_PATTERNS:
    if exclude in tags_lower:
        # Special case: "vegetable-chip" should NOT be excluded
        # even though it contains "vegetable"
        if exclude == 'vegetable' and 'vegetable-chip' in tags_
            continue
        return None

# Step 2: Match to a snack category
for keyword, category in CATEGORY_MAP:
    if keyword in tags_lower:
        return category

return None

# Test with real examples from your actual data
test_cases = [
    ("Mochi ice cream", "en:desserts,en:frozen-foods,en:frozen-"),
    ("Potato crisps", "en:snacks,en:salty-snacks,en:chips-and"),
    ("Peanuts", "en:legumes-and-their-products,en:nuts,"),
    ("Gummi candies", "en:snacks,en:sweet-snacks,en:confectio"),
    ("Biscuits", "en:snacks,en:sweet-snacks,en:biscuits-"),
    ("Orange juice", "en:beverages,en:fruit-juices,en:orange"),
    ("Toaster pastries", "en:snacks,en:sweet-snacks,en:biscuits-"),
    ("en:null tag", "en:null"),
    ("Soda", "en:beverages,en:carbonated-drinks,en:s"),
    ("BBQ chips", "en:bbq-chips"),
    ("Crackers", "en:snacks,en:salty-snacks,en:biscuits-"),
    ("Vegetable chips", "en:snacks,en:salty-snacks,en:vegetable")
]

print("== CLASSIFICATION TEST (with real tag patterns) ==\n")
for name, tags in test_cases:
    result = assign_category(tags)
    status = "added" if result else "(excluded)"
    print(f"{status} {name}<20> → {result}")

== CLASSIFICATION TEST (with real tag patterns) ==
(excluded) Mochi ice cream      → None
added Potato crisps           → Chips & Savory Snacks
(excluded) Peanuts             → None
added Gummi candies           → Candy & Confectionery
added Biscuits                → Cookies & Biscuits
(excluded) Orange juice        → None
added Toaster pastries         → Cookies & Biscuits

```

```
(excluded) en:null tag           → None
(excluded) Soda                 → None
added  BBQ chips               → Chips & Savory Snacks
added  Crackers                → Chips & Savory Snacks
added  Vegetable chips          → Chips & Savory Snacks
```

```
# This creates a new column called 'primary_category'
df_clean['primary_category'] = df_clean['categories_tags'].apply(as

# filter OUT rows where primary_category is None

df_snacks = df_clean[df_clean['primary_category'].notna()].copy()

print(f"\nTotal rows before snack filter: {len(df_clean)}")
print(f"Snack products identified:      {len(df_snacks)}")
print(f"Non-snack rows removed:         {len(df_clean) - len(df_snacks)}")
```

```
Total rows before snack filter: 391,528
Snack products identified:      54,169
Non-snack rows removed:         337,359
```

```
category_counts = df_snacks['primary_category'].value_counts()

print(" CATEGORY DISTRIBUTION \n")
print(category_counts.to_string())
print(f"\nTotal snack products: {len(df_snacks)}")

# Visualize it
import matplotlib.pyplot as plt

fig, ax = plt.subplots(figsize=(10, 5))
category_counts.plot(kind='barh', ax=ax, color='steelblue', edgecolor='black')
ax.set_xlabel('Number of Products')
ax.set_title('Products per Category', fontsize=14, fontweight='bold')
ax.invert_yaxis()
plt.tight_layout()
plt.savefig('category_distribution.png', dpi=150, bbox_inches='tight')
plt.show()
print("Chart saved")
```

## CATEGORY DISTRIBUTION

primary_category	
General Snacks	16835
Candy & Confectionery	15717
Cookies & Biscuits	13810
Chips & Savory Snacks	7620
Nuts & Seeds	105

Fruit & Veg Snacks	38
Protein & Fitness Bars	24
Dairy & Yogurt Snacks	20

Total snack products: 54,169

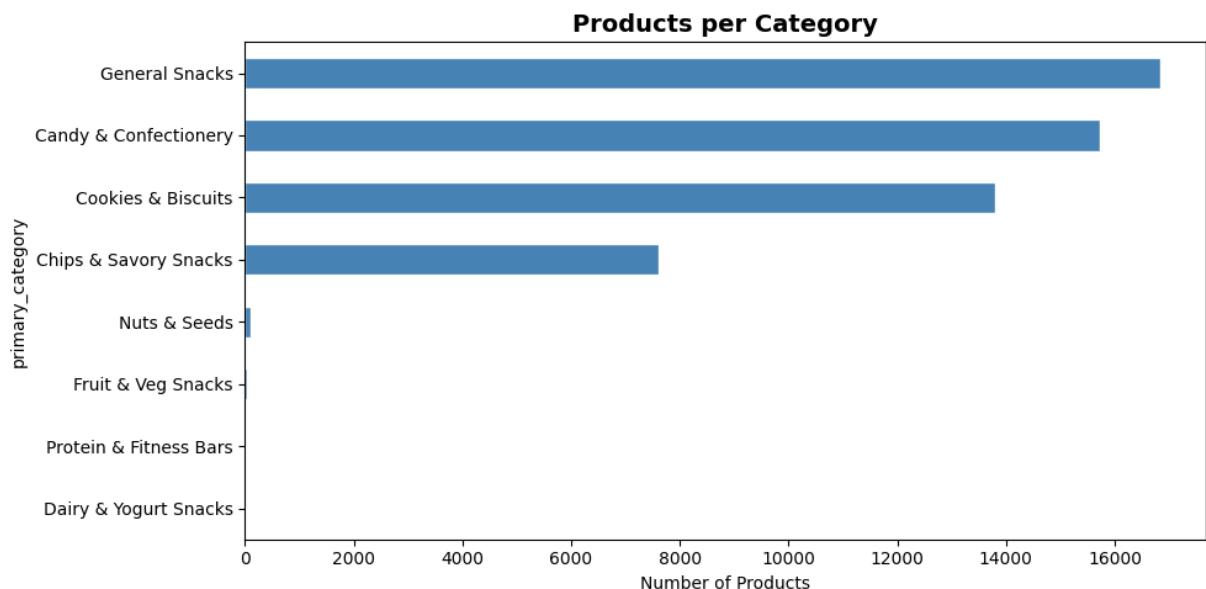


Chart saved

```
summary = df_snacks.groupby('primary_category').agg(  
    product_count = ('product_name', 'count'),  
    avg_sugar = ('sugars_100g', 'mean'),  
    avg_protein = ('proteins_100g', 'mean'),  
    avg_fat = ('fat_100g', 'mean'),  
    median_sugar = ('sugars_100g', 'median'),  
    median_protein = ('proteins_100g', 'median'),  
).round(2).sort_values('avg_sugar', ascending=False)  
  
print(" NUTRITIONAL PROFILE BY CATEGORY \n")  
print(summary.to_string())  
  
# Save snack dataset  
df_snacks.to_csv('food_facts_snacks.csv', index=False)  
print("\n Snack dataset saved as food_facts_snacks.csv")
```

#### NUTRITIONAL PROFILE BY CATEGORY

primary_category	product_count	avg_sugar	avg_protein	avg_f
Candy & Confectionery	15717	54.00	3.37	12.
Fruit & Veg Snacks	38	51.72	3.62	3.
Cookies & Biscuits	13810	29.06	5.28	16.
Nuts & Seeds	105	19.08	10.36	27.
General Snacks	16835	17.54	10.70	26.
Dairy & Yogurt Snacks	20	15.63	8.83	10.
Chips & Savory Snacks	7620	8.23	5.15	14.
Protein & Fitness Bars	24	4.24	0.47	0.

Snack dataset saved as food\_facts\_snacks.csv

## ▼ Data Visualization

```
plt.rcParams.update({
    'figure.facecolor':      'white',
    'axes.facecolor':        '#F8F9FA',
    'axes.grid':             True,
    'grid.color':            '#E0E0E0',
    'grid.linewidth':         0.7,
    'grid.alpha':             0.8,
    'axes.spines.top':       False,
    'axes.spines.right':     False,
    'axes.spines.left':      True,
    'axes.spines.bottom':    True,
    'axes.labelsize':         12,
    'axes.titlesize':        14,
    'axes.titleweight':      'bold',
    'xtick.labelsize':       10,
    'ytick.labelsize':       10,
    'legend.fontsize':        10,
    'legend.framealpha':     0.9,
    'font.family':            'sans-serif',
})

CATEGORY_COLORS = {
    'Candy & Confectionery': '#E63946',
    'Cookies & Biscuits':     '#F4A261',
    'Chips & Savory Snacks': '#E9C46A',
    'General Snacks':          '#A8DADC',
    'Fruit & Veg Snacks':     '#52B788',
    'Nuts & Seeds':            '#2D6A4F',
    'Dairy & Yogurt Snacks': '#457B9D',
    'Protein & Fitness Bars': '#1D3557',
}
```

```
try:  
    df_snacks  
    print(f"df_snacks in memory: {len(df_snacks):,} rows")  
except NameError:  
    df_snacks = pd.read_csv('food_facts_snacks.csv')  
    print(f"Reloaded: {len(df_snacks):,} rows")  
  
# Cap at 99th percentile for visual clarity – all data still in df_  
# We only cap for PLOTTING, not for stats  
p99_sugar = df_snacks['sugars_100g'].quantile(0.99)  
p99_protein = df_snacks['proteins_100g'].quantile(0.99)  
  
df_plot = df_snacks[  
    (df_snacks['sugars_100g'] <= p99_sugar) &  
    (df_snacks['proteins_100g'] <= p99_protein)  
].copy()  
  
print(f"Rows available for plotting: {len(df_plot):,}")  
print(f"(99th pct: sugar={p99_sugar:.1f}g, protein={p99_protein:.1f}g)")  
  
df_snacks in memory: 54,169 rows  
Rows available for plotting: 53,112  
(99th pct: sugar=91.7g, protein=33.3g)
```

```
fig, ax = plt.subplots(figsize=(14, 9))  
  
PLOT_ORDER = [  
    'General Snacks',  
    'Candy & Confectionery',  
    'Cookies & Biscuits',  
    'Fruit & Veg Snacks',  
    'Dairy & Yogurt Snacks',  
    'Chips & Savory Snacks',  
    'Nuts & Seeds',  
    'Protein & Fitness Bars',  
]  
  
for category in PLOT_ORDER:  
    subset = df_plot[df_plot['primary_category'] == category]  
    ax.scatter(  
        subset['sugars_100g'],  
        subset['proteins_100g'],  
        c=CATEGORY_COLORS[category],  
        label=f"{category} (n={len(subset):,})",  
        alpha=0.25,  
        s=8,  
        linewidths=0,
```

```
        rasterized=True,
    )

# Blue Ocean Zone rectangle
blue_ocean_rect = FancyBboxPatch(
    (0, 15),
    10, p99_protein - 15,
    boxstyle="round,pad=0.3",
    facecolor='#1D3557',
    edgecolor='#1D3557',
    alpha=0.12,
    linewidth=2,
    linestyle='--',
    zorder=5
)
ax.add_patch(blue_ocean_rect)

# Blue Ocean label inside the box
ax.text(
    5, p99_protein * 0.88, \
    ' BLUE OCEAN ZONE\n' \
    'High Protein (>15g) + Low Sugar (<10g)\n' \
    'Under-served: few products exist here',
    fontsize=10,
    fontweight='bold',
    color='#1D3557',
    ha='center',
    va='top',
    zorder=6,
    bbox=dict(
        boxstyle='round,pad=0.4',
        facecolor='white',
        edgecolor='#1D3557',
        alpha=0.92,
        linewidth=1.5
    )
)

# Quadrant divider lines
ax.axhline(y=15, color="#555555", linewidth=1.2, linestyle=':', alpha=0.92)
ax.axvline(x=10, color="#555555", linewidth=1.2, linestyle=':', alpha=0.92)

# Quadrant labels in the three "non-blue-ocean" corners
ax.text(50, p99_protein * 0.92,
    ' High Sugar + High Protein\n(Uncommon – specialty product',
    fontsize=8.5, color="#555", ha='center', style='italic',
    bbox=dict(boxstyle='round', facecolor='#FFF9C4', alpha=0.7,
```

```
ax.text(50, 2,
        ' SUGAR TRAP\n'
        'High Sugar + Low Protein\n'
        '29,527 products – over-served',
        fontsize=9, color='#7f1d1d', ha='center', fontweight='bold',
        bbox=dict(boxstyle='round', facecolor='#FEE2E2', alpha=0.8)

ax.text(3, 2,
        ' Low Sugar + Low Protein\n(Savory/neutral snacks)',
        fontsize=8.5, color="#555", ha='center', style='italic',
        bbox=dict(boxstyle='round', facecolor='#FFF9C4', alpha=0.7,

# Arrow pointing to the opportunity
ax.annotate(
    'Market Opportunity\npoints here →',
    xy=(5, 22),
    xytext=(18, 26),
    fontsize=9,
    color='#1D3557',
    fontweight='bold',
    arrowprops=dict(
        arrowstyle='->', color='#1D3557', lw=1.8
    )
)

# Titles and labels
ax.set_title(
    'Sugar vs. Protein: The Snack Market Landscape\n'
    'Each dot = 1 product | 53,112 snack products from Open Food
    fontsize=15,
    fontweight='bold',
    pad=18
)
ax.set_xlabel('Sugar (g per 100g) ← Lower is healthier', fontsize=14)
ax.set_ylabel('Protein (g per 100g) ↑ Higher is better', fontsize=14)

# Axis limits with breathing room
ax.set_xlim(-1, p99_sugar + 2)
ax.set_ylim(-0.5, p99_protein + 3)

# Legend – sorted, with product counts
handles, labels = ax.get_legend_handles_labels()
ax.legend(
    handles, labels,
    title='Category (n = product count)',
    title_fontsize=10,
    loc='upper right',
    framealpha=0.95,
```

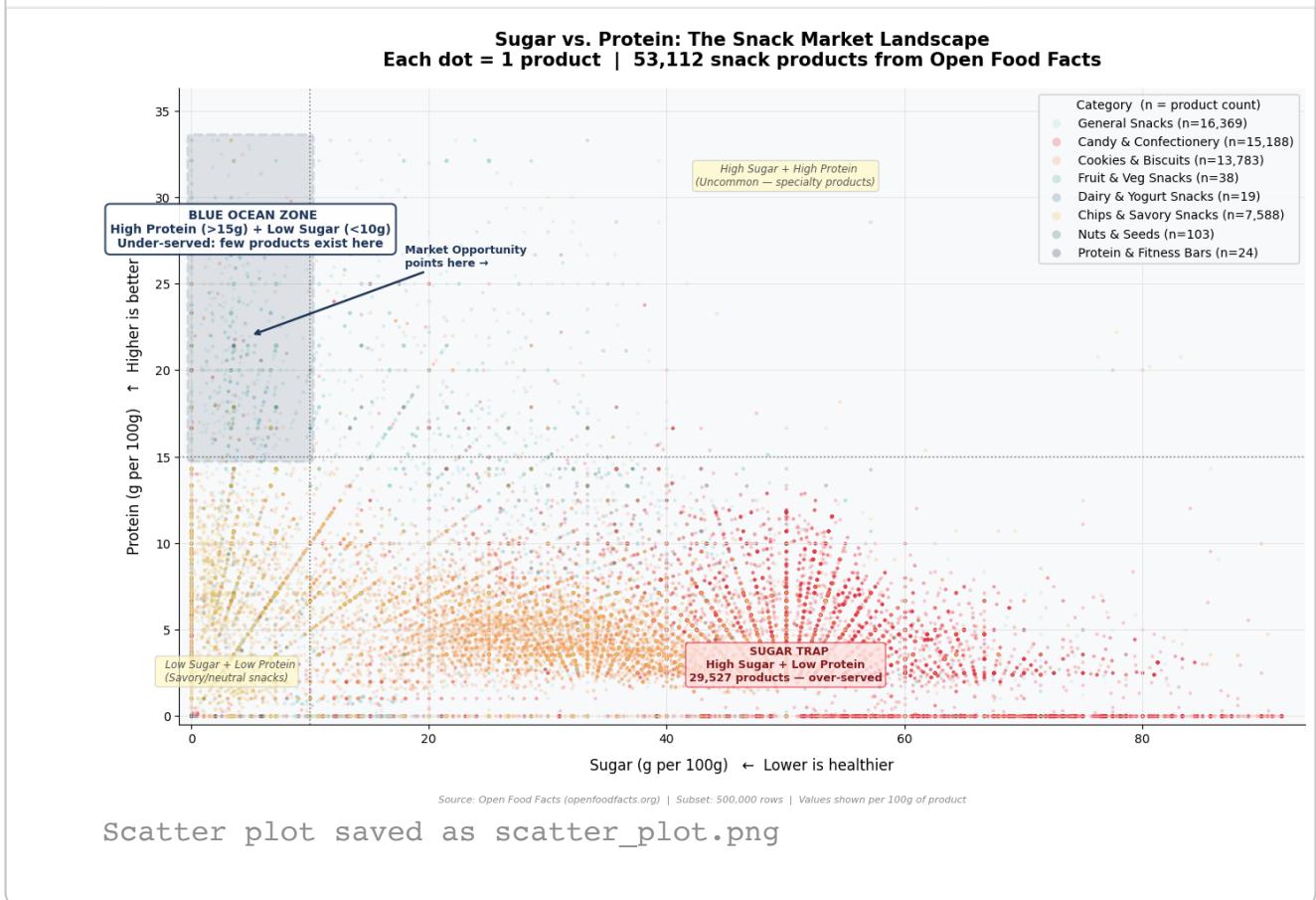
```

        edgecolor='#cccccc',
        markerscale=2.5
    )

# Footer annotation
fig.text(
    0.5, 0.01,
    'Source: Open Food Facts (openfoodfacts.org) | Subset: 500,000',
    'Values shown per 100g of product',
    ha='center', fontsize=8, color="#888888", style='italic'
)

plt.tight_layout(rect=[0, 0.03, 1, 1])
plt.savefig('scatter_plot.png', dpi=180, bbox_inches='tight')
plt.show()
print("Scatter plot saved as scatter_plot.png")

```



```

summary = df_snacks.groupby('primary_category').agg(
    product_count = ('product_name', 'count'),
    avg_sugar     = ('sugars_100g', 'mean'),
    avg_protein   = ('proteins_100g', 'mean'),
    median_sugar  = ('sugars_100g', 'median'),
    median_protein = ('proteins_100g', 'median'),
).round(2).reset_index()

```

```
# Sort by avg_sugar descending – worst at top, best at bottom
summary = summary.sort_values('avg_sugar', ascending=True)

fig, ax = plt.subplots(figsize=(13, 7))

x = np.arange(len(summary))
bar_width = 0.38

bars_sugar = ax.bar(
    x - bar_width/2,
    summary['avg_sugar'],
    width=bar_width,
    color='#E63946',
    label='Avg Sugar (g per 100g)',
    edgecolor='white',
    linewidth=0.5,
    zorder=3
)

bars_protein = ax.bar(
    x + bar_width/2,
    summary['avg_protein'],
    width=bar_width,
    color='#1D3557',
    label='Avg Protein (g per 100g)',
    edgecolor='white',
    linewidth=0.5,
    zorder=3
)

# Value labels on top of every bar
for bar in bars_sugar:
    ax.text(
        bar.get_x() + bar.get_width() / 2,
        bar.get_height() + 0.4,
        f'{bar.get_height():.1f}g',
        ha='center', va='bottom', fontsize=8.5, color='#C0392B', fc
    )

for bar in bars_protein:
    ax.text(
        bar.get_x() + bar.get_width() / 2,
        bar.get_height() + 0.4,
        f'{bar.get_height():.1f}g',
        ha='center', va='bottom', fontsize=8.5, color='#1D3557', fc
    )

# Product count labels below each category on X axis
```

```
ax.set_xticks(x)
ax.set_xticklabels([
    f'{row["primary_category"]}\n(n={row["product_count"]},)}'
    for _, row in summary.iterrows()
], fontsize=9.5, rotation=20, ha='right')

# Highlight the best opportunity bar pair
# Chips & Savory Snacks: low sugar, high product count
best_idx = summary[summary['primary_category'] == 'Chips & Savory S
if len(best_idx) > 0:
    best_pos = list(summary.index).index(best_idx[0])
    ax.axvspan(
        best_pos - 0.5, best_pos + 0.5,
        alpha=0.10, color='#2D6A4F', zorder=0
    )
    ax.text(
        best_pos, summary['avg_sugar'].max() * 0.75,
        '⭐ Best opportunity\n(large market, low sugar)',
        ha='center', fontsize=8.5, color='#2D6A4F', fontweight='bol
        bbox=dict(boxstyle='round', facecolor='#D8F3DC', edgecolor=
    )

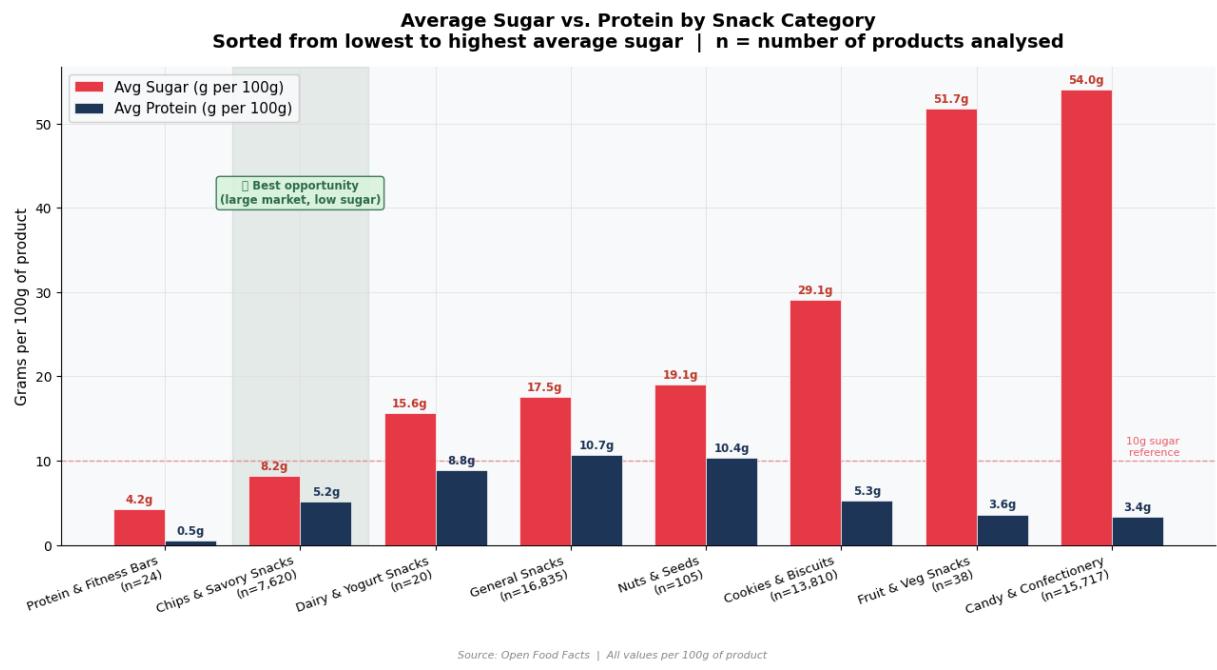
ax.set_title(
    'Average Sugar vs. Protein by Snack Category\n'
    'Sorted from lowest to highest average sugar | n = number of
    fontsize=14, fontweight='bold', pad=15
)
ax.set_ylabel('Grams per 100g of product', fontsize=11)
ax.set_xlabel('')
ax.legend(fontsize=11, loc='upper left')

# Reference line at 10g sugar – the "decent" threshold
ax.axhline(y=10, color='#E63946', linewidth=1, linestyle='--', alph
ax.text(
    len(summary) - 0.5, 10.4,
    '10g sugar\nreference',
    fontsize=8, color='#E63946', ha='right', va='bottom', alpha=0.8
)

fig.text(
    0.5, 0.01,
    'Source: Open Food Facts | All values per 100g of product',
    ha='center', fontsize=8, color="#888", style='italic'
)

plt.tight_layout(rect=[0, 0.03, 1, 1])
plt.savefig('bar_chart.png', dpi=180, bbox_inches='tight')
plt.show()
```

```
print("Bar chart saved as bar_chart.png")
```



Bar chart saved as bar\_chart.png

```
# Calculate the opportunity score
summary['protein_sugar_ratio'] = summary['avg_protein'] / (summary['avg_sugar'] * np.log1p(summary['product_count']))
summary['opportunity_score'] = (
    summary['protein_sugar_ratio'] * np.log1p(summary['product_count'])
).round(2)

score_min = summary['opportunity_score'].min()
score_max = summary['opportunity_score'].max()
summary['score_norm'] = (
    (summary['opportunity_score'] - score_min) / (score_max - score_min)
).round(1)

summary_sorted = summary.sort_values('score_norm', ascending=True)

fig, ax = plt.subplots(figsize=(12, 6))

# Color each bar: green ≥ 60, yellow 30–60, red < 30
bar_colors = [
    '#2D6A4F' if s >= 60 else '#E9C46A' if s >= 30 else '#E63946'
    for s in summary_sorted['score_norm']
]

bars = ax.barh(
    summary_sorted['primary_category'],
    summary_sorted['score_norm'],
    color=bar_colors,
```

```
        edgecolor='white',
        linewidth=0.8,
        height=0.6,
        zorder=3
    )

# Rich labels on each bar
for i, (_, row) in enumerate(summary_sorted.iterrows()):
    # Score value at end of bar
    ax.text(
        row['score_norm'] + 1.2,
        i,
        f'{row["score_norm"]:.0f}',
        va='center', fontsize=11, fontweight='bold',
        color='#333'
    )
    # Inline stats inside the bar
    ax.text(
        1.5, i,
        f" Sugar: {row['avg_sugar']:.1f}g | "
        f"Protein: {row['avg_protein']:.1f}g | "
        f"Products: {row['product_count']:,}",
        va='center', fontsize=8.5, color='white', fontweight='bold',
        alpha=0.95
    )

# Threshold line and label
ax.axvline(x=60, color='#2D6A4F', linewidth=1.5, linestyle='--', alpha=0.95)
ax.text(61, -0.6, 'High\nOpportunity\nThreshold',
        fontsize=8, color='#2D6A4F', va='bottom')

# Legend
legend_patches = [
    mpatches.Patch(color='#2D6A4F', label='High Opportunity (score > 60)'),
    mpatches.Patch(color='#E9C46A', label='Medium Opportunity (score between 40 and 60)'),
    mpatches.Patch(color='#E63946', label='Low Opportunity (score < 40)')
]
ax.legend(handles=legend_patches, loc='lower right', fontsize=9, frameon=False)

ax.set_title(
    'Market Opportunity Score by Snack Category\n'
    'Formula: (Avg Protein ÷ Avg Sugar) × log(Product Count) '
    '- rewards healthy nutrition in large, established markets',
    fontsize=13, fontweight='bold', pad=15
)
ax.set_xlabel('Opportunity Score (0 = avoid → 100 = highest priority)')
ax.set_xlim(0, 115)
ax.set_ylabel('')
```

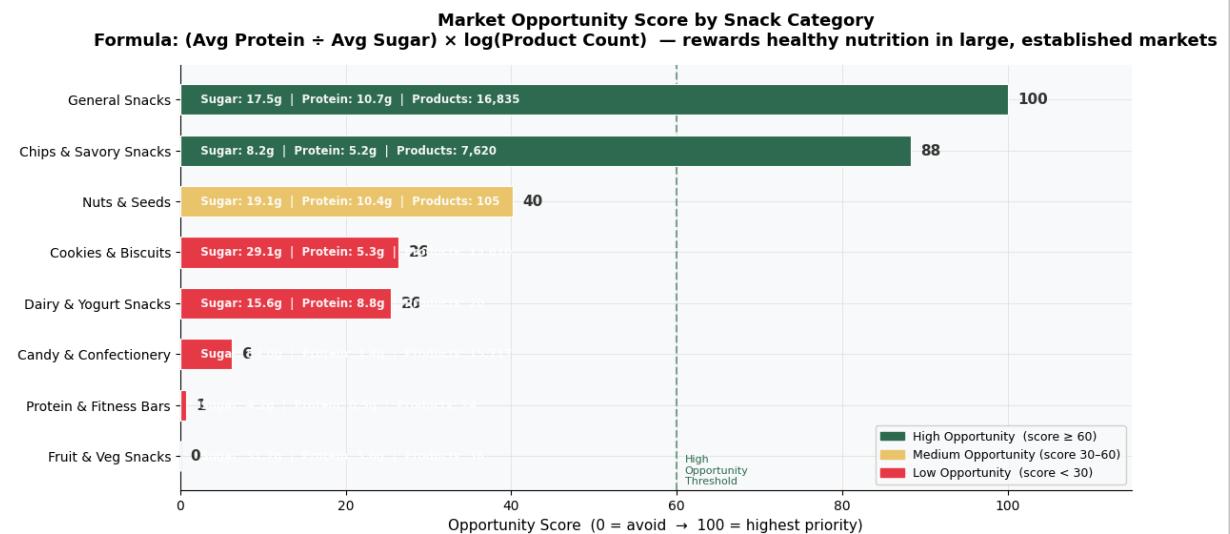
```

fig.text(
    0.5, 0.01,
    'Source: Open Food Facts | Score normalised 0–100 | '
    'Higher score = large market with poor current nutritional profile'
    ha='center', fontsize=8, color='#888', style='italic'
)

plt.tight_layout(rect=[0, 0.04, 1, 1])
plt.savefig('opportunity_score.png', dpi=180, bbox_inches='tight')
plt.show()
print(" Opportunity score chart saved as opportunity_score.png")

print("\n==== FINAL OPPORTUNITY SCORES ===")
print(summary_sorted[['primary_category', 'product_count',
                     'avg_sugar', 'avg_protein', 'score_norm']]
      .sort_values('score_norm', ascending=False)
      .to_string(index=False))

```



Opportunity score chart saved as opportunity\_score.png

```

==== FINAL OPPORTUNITY SCORES ====
   primary_category  product_count  avg_sugar  avg_protein  score
   General Snacks           16835     17.54       10.70
   Chips & Savory Snacks        7620      8.23        5.15
   Nuts & Seeds                  105     19.08       10.36
   Cookies & Biscuits            13810     29.06        5.28
   Dairy & Yogurt Snacks             20     15.63        8.83
   Candy & Confectionery            15717     54.00        3.37
   Protein & Fitness Bars                 24      4.24        0.47
   Fruit & Veg Snacks                   38     51.72        3.62

```

```

# Target category: Chips & Savory Snacks
# General Snacks scored #1 but it's a catch-all bucket – vague, not
# "Build a General Snack" tells R&D nothing.
# Chips & Savory Snacks is a SPECIFIC, REAL shelf category (score=8
# with 7,620 products proving consumer demand, and avg protein of c
# – a clear, measurable gap to fill.

TARGET_CATEGORY = 'Chips & Savory Snacks'

target = df_snacks[df_snacks['primary_category'] == TARGET_CATEGORY]

# Products in the 75th–90th percentile for protein = "best in class"
p75_protein = target['proteins_100g'].quantile(0.75)
p90_protein = target['proteins_100g'].quantile(0.90)
p25_sugar   = target['sugars_100g'].quantile(0.25)
median_sugar = target['sugars_100g'].median()

blue_ocean_products = target[
    (target['proteins_100g'] >= 15) &
    (target['sugars_100g']    <= 10)
]

blue_ocean_pct = len(blue_ocean_products) / len(target) * 100

# Total snack market context
total_snacks = len(df_snacks)
total_high_protein_low_sugar = len(df_snacks[
    (df_snacks['proteins_100g'] >= 15) &
    (df_snacks['sugars_100g']    <= 10)
])
blue_ocean_market_pct = total_high_protein_low_sugar / total_snacks

print(" STORY 4: The recommendation")

print(f"""
TARGET CATEGORY: {TARGET_CATEGORY}
Product count: {len(target)} products (established market)
Avg sugar: {target['sugars_100g'].mean():.1f}g per 100g
Avg protein: {target['proteins_100g'].mean():.1f}g per 100g

BENCHMARK – Top 25% protein in this category: >{p75_protein:.1f}g
BENCHMARK – Top 10% protein in this category: >{p90_protein:.1f}g
LOW SUGAR target (bottom 25%): <{p25_sugar:.1f}g

BLUE OCEAN GAP in {TARGET_CATEGORY}:
Products with ≥15g protein AND ≤10g sugar: {len(blue_ocean_products)}
That's only {blue_ocean_pct:.1f}% of the category – nearly empty
""")

```

### ACROSS ALL SNACK CATEGORIES:

Products in the Blue Ocean zone: {total\_high\_protein\_low\_sugar} c

That's only {blue\_ocean\_market\_pct:.1f}% of the entire snack market")

```
print("  Recommendation (Story 4 Acceptance Criteria)")
print(f""")
```

"Based on the data, the biggest market opportunity is in CHIPS & SAVORY SNACKS, specifically targeting products with {p75\_protein:.0f}g+ of protein and less than {p25\_sugar:.0f}g

This category contains {len(target):,} products – proving strong consumer demand – yet only {blue\_ocean\_pct:.1f}% of those products sit in the High Protein + Low Sugar quadrant.

Across all {total\_snacks:,} snack products analysed, just {blue\_ocean\_market\_pct:.1f}% meet the Blue Ocean criteria ( $\geq 15\text{g protein}$  AND  $\leq 10\text{g sugar}$ ), confirming the market gap is real and wide."")

### STORY 4: The recommendation

TARGET CATEGORY: Chips & Savory Snacks

Product count: 7,620 products (established market)

Avg sugar: 8.2g per 100g

Avg protein: 5.2g per 100g

BENCHMARK – Top 25% protein in this category: >7.1g

BENCHMARK – Top 10% protein in this category: >10.0g

LOW SUGAR target (bottom 25%): <0.0g

BLUE OCEAN GAP in Chips & Savory Snacks:

Products with  $\geq 15\text{g protein}$  AND  $\leq 10\text{g sugar}$ : 111 of 7,620

That's only 1.5% of the category – nearly empty

### ACROSS ALL SNACK CATEGORIES:

Products in the Blue Ocean zone: 3176 of 54,169

That's only 5.9% of the entire snack market

Recommendation (Story 4 Acceptance Criteria)

"Based on the data, the biggest market opportunity is in CHIPS & SAVORY SNACKS, specifically targeting products with 7g+ of protein and less than 0g of sugar.

This category contains 7,620 products – proving strong consumer demand – yet only 1.5% of those products sit in the High Protein + Low Sugar quadrant.

Across all 54,169 snack products analysed, just 5.9% meet the Blue Ocean criteria ( $\geq 15\text{g protein}$ ,

$\leq 10\text{g sugar}$ ), confirming the market gap is real and wide."

```
import re
from collections import Counter

# For products in the Blue Ocean zone (high protein, low sugar), we
# Step 1: Isolate the Blue Ocean products across ALL categories
blue_ocean_all = df_snacks[
    (df_snacks['proteins_100g'] >= 15) &
    (df_snacks['sugars_100g'] <= 10) &
    (df_snacks['ingredients_text'].notna()) # Must have ingredients
].copy()

print(f"Blue Ocean products with ingredient data: {len(blue_ocean_all)}")
print(f"\nSample ingredients text:")
print(blue_ocean_all['ingredients_text'].iloc[0][:300])
print("...")
```

Blue Ocean products with ingredient data: 3072

Sample ingredients text:

Nuts (Peanuts, Almonds), Prebiotic Blend (Tapioca Fiber, Vegetable Fiber, ...)

```
# Step 2: Define protein source keywords to search for
# These are the most common protein-contributing ingredients in snacks
# We search for these as substrings – "whey protein isolate" matches
```

```
PROTEIN_SOURCES = {
    # Animal proteins
    'whey': 'Whey Protein',
    'casein': 'Casein',
    'egg white': 'Egg White',
    'egg': 'Egg',
    'milk protein': 'Milk Protein',
    'chicken': 'Chicken',
    'tuna': 'Tuna',
    'salmon': 'Salmon',
    'beef': 'Beef',

    # Plant proteins
    'pea protein': 'Pea Protein',
    'soy protein': 'Soy Protein',
    'soy': 'Soy / Soybean',
    'peanut': 'Peanut',
    'almond': 'Almond',
```

```

'sunflower seed':      'Sunflower Seed',
'pumpkin seed':       'Pumpkin Seed',
'hemp':                'Hemp Protein',
'quinoa':              'Quinoa',
'chickpea':            'Chickpea',
'lentil':               'Lentil',
'rice protein':        'Rice Protein',

# Common high-protein bases
'oat':                 'Oats',
'nut':                 'Nuts (general)',
'seed':                'Seeds (general)',

}

# Step 3: Count occurrences of each protein source
protein_counts = Counter()

for ingredients in blue_ocean_all['ingredients_text'].dropna():
    ingredients_lower = ingredients.lower()
    for keyword, label in PROTEIN_SOURCES.items():
        if keyword in ingredients_lower:
            protein_counts[label] += 1
            break # Count each product once per keyword match – av

# Step 4: Convert to DataFrame for display and charting
protein_df = pd.DataFrame(
    protein_counts.most_common(15),
    columns=['Protein Source', 'Product Count']
)
protein_df['Pct of Blue Ocean Products'] = (
    protein_df['Product Count'] / len(blue_ocean_all) * 100
).round(1)

print(" TOP PROTEIN SOURCES IN BLUE OCEAN PRODUCTS")
print(protein_df.to_string(index=False))

```

TOP PROTEIN SOURCES IN BLUE OCEAN PRODUCTS			
Protein Source	Product Count	Pct of Blue Ocean Products	
Peanut	799	26.0	
Soy / Soybean	669	21.8	
Almond	397	12.9	
Beef	135	4.4	
Nuts (general)	125	4.1	
Whey Protein	111	3.6	
Soy Protein	62	2.0	
Sunflower Seed	48	1.6	
Pumpkin Seed	47	1.5	
Seeds (general)	29	0.9	
Chicken	26	0.8	

Egg	22	0.7
Oats	22	0.7
Chickpea	13	0.4
Lentil	11	0.4

```
# Step 5: Visualise it
fig, ax = plt.subplots(figsize=(11, 6))

# Color bars: animal proteins blue, plant proteins green
animal_proteins = {
    'Whey Protein', 'Casein', 'Egg White', 'Egg',
    'Milk Protein', 'Chicken', 'Tuna', 'Salmon', 'Beef'
}

bar_colors = [
    '#1D3557' if src in animal_proteins else '#2D6A4F'
    for src in protein_df['Protein Source']
]

bars = ax.barh(
    protein_df['Protein Source'],
    protein_df['Product Count'],
    color=bar_colors,
    edgecolor='white',
    linewidth=0.6,
    height=0.65
)

# Value labels
for i, (_, row) in enumerate(protein_df.iterrows()):
    ax.text(
        row['Product Count'] + 0.3,
        i,
        f'{row["Product Count"]}\n({row["Pct of Blue Ocean Products"]})',
        va='center', fontsize=9.5, color="#333"
    )

# Legend for color coding
legend_patches = [
    mpatches.Patch(color='#1D3557', label='Animal-based protein'),
    mpatches.Patch(color='#2D6A4F', label='Plant-based protein'),
]
ax.legend(handles=legend_patches, loc='lower right', fontsize=10)

ax.set_title(
    'Top Protein Sources in Blue Ocean Snack Products\n'
    f'Based on {len(blue_ocean_all)} products with ≥15g protein & ≤ '
    f'size=13, fontweight="bold", pad=15
```

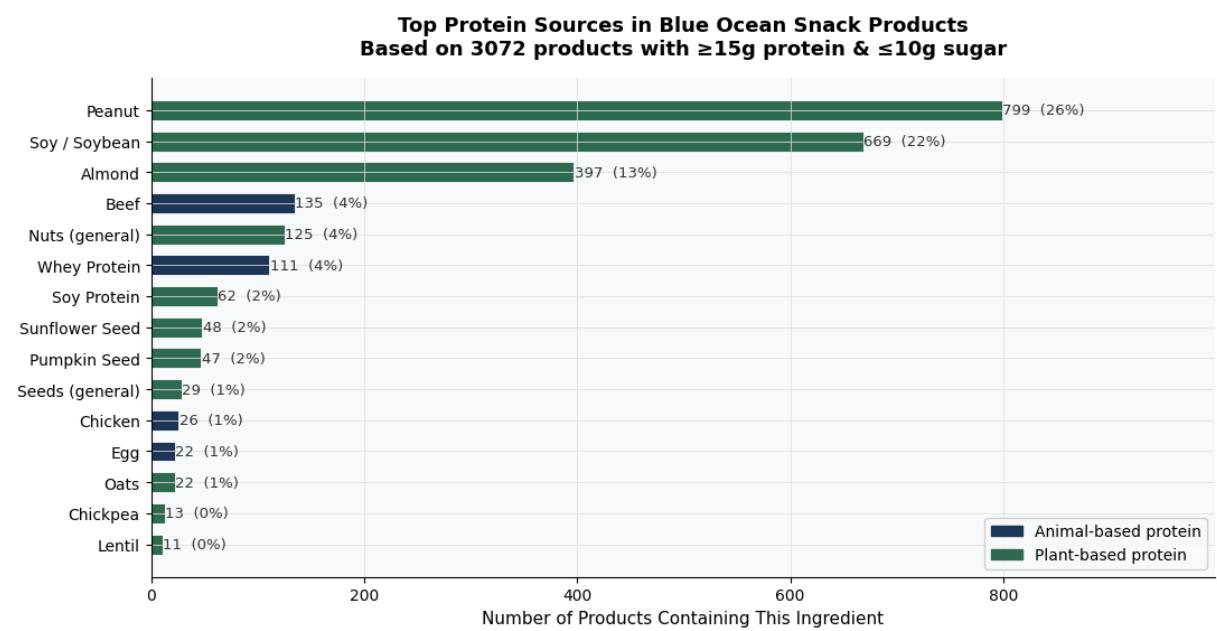
```

)
ax.set_xlabel('Number of Products Containing This Ingredient', font
ax.invert_yaxis() # Highest count at top
ax.set_xlim(0, protein_df['Product Count'].max() * 1.25)

fig.text(
    0.5, 0.01,
    'Source: Open Food Facts ingredients_text column | '
    '% = share of all Blue Ocean products containing this ingredient'
    ha='center', fontsize=8, color='#888', style='italic'
)

plt.tight_layout(rect=[0, 0.04, 1, 1])
plt.savefig('protein_sources.png', dpi=180, bbox_inches='tight')
plt.show()
print(" Protein sources chart saved as protein_sources.png")

```



Protein sources chart saved as protein\_sources.png

```

# =====
# CANDIDATE'S CHOICE: "The Reformulation Gap Analysis"
#
# WHY I ADDED THIS:
# The scatter plot shows WHERE the market is. The opportunity score
#
# This chart quantifies the DISTANCE each category needs to travel
# to reach the Blue Ocean zone ( $\geq 15\text{g}$  protein,  $\leq 10\text{g}$  sugar).
# A category close to the target = easy reformulation win.
# A category far from the target = requires new product architectur

```

```
# This transforms the analysis from "here's the gap" to "here's now"
```

```
TARGET_PROTEIN = 15.0    # Blue Ocean protein floor
TARGET_SUGAR   = 10.0    # Blue Ocean sugar ceiling

# Calculate gap per category
summary_gap = df_snacks.groupby('primary_category').agg(
    avg_sugar      = ('sugars_100g', 'mean'),
    avg_protein    = ('proteins_100g', 'mean'),
    product_count  = ('product_name', 'count'),
).round(2).reset_index()

# Gap = how far the AVERAGE product in this category is from the target
# Positive gap = needs to increase (protein) or decrease (sugar)
# Negative gap = already meets or beats the target
summary_gap['protein_gap'] = (TARGET_PROTEIN - summary_gap['avg_protein'])
summary_gap['sugar_gap']   = (summary_gap['avg_sugar'] - TARGET_SUGAR)

# Clamp negatives to 0 - "already meets target" = 0 gap, not negative
summary_gap['protein_gap_display'] = summary_gap['protein_gap'].clip(lower=0)
summary_gap['sugar_gap_display']   = summary_gap['sugar_gap'].clip(lower=0)

# Reformulation difficulty score:
# Combine both gaps into one number (equal weight)
# Higher = harder to reformulate
summary_gap['difficulty'] = (
    summary_gap['protein_gap_display'] + summary_gap['sugar_gap_display']
).round(1)

summary_gap = summary_gap.sort_values('difficulty', ascending=True).

print("== REFORMULATION GAP TABLE ==\n")
print(summary_gap[[
    'primary_category', 'avg_sugar', 'avg_protein',
    'protein_gap', 'sugar_gap', 'difficulty'
]].to_string(index=False))
```

== REFORMULATION GAP TABLE ==

primary_category	avg_sugar	avg_protein	protein_gap	sugar_gap
Chips & Savory Snacks	8.23	5.15	9.8	-1
Dairy & Yogurt Snacks	15.63	8.83	6.2	5
General Snacks	17.54	10.70	4.3	7
Nuts & Seeds	19.08	10.36	4.6	9
Protein & Fitness Bars	4.24	0.47	14.5	-5
Cookies & Biscuits	29.06	5.28	9.7	19
Fruit & Veg Snacks	51.72	3.62	11.4	41

Candy & Confectionery	54.00	3.37	11.6	44
-----------------------	-------	------	------	----

```
# Build the chart
fig, axes = plt.subplots(1, 2, figsize=(16, 7))
fig.suptitle(
    "The Reformulation Gap Analysis\n"
    "How far is each category from the Blue Ocean target "
    "(≥15g protein + ≤10g sugar)?",
    fontsize=14, fontweight='bold', y=1.01
)

categories = summary_gap['primary_category']
x = np.arange(len(categories))
bar_h = 0.4

# LEFT PANEL: Sugar gap (how much sugar to cut)
ax1 = axes[0]

colors_sugar = [
    '#E63946' if g > 20 else '#F4A261' if g > 5 else '#52B788'
    for g in summary_gap['sugar_gap_display']
]

bars1 = ax1.bars(
    x, summary_gap['sugar_gap_display'],
    height=bar_h, color=colors_sugar,
    edgecolor='white', linewidth=0.6
)

for i, (gap, avg) in enumerate(zip(
    summary_gap['sugar_gap_display'],
    summary_gap['avg_sugar']
)):
    if gap == 0:
        ax1.text(0.3, i, f" Avg {avg:.1f}g – already at target",
                 va='center', fontsize=8.5, color='#2D6A4F', fontwei
    else:
        ax1.text(gap + 0.3, i,
                 f"Cut {gap:.0f}g (avg now: {avg:.1f}g)",
                 va='center', fontsize=8.5, color='#333')

ax1.set_yticks(x)
ax1.set_yticklabels(categories, fontsize=9.5)
ax1.set_xlabel('Sugar reduction needed (g per 100g)', fontsize=10)
ax1.set_title('Sugar to Cut\nunto reach ≤10g target',
              fontsize=11, fontweight='bold', pad=10)
ax1.axvline(x=0, color='#333', linewidth=1)
```

```

ax1.set_xlim(0, summary_gap['sugar_gap_display'].max() * 1.35)

# Reference: Chips & Savory as the benchmark
best_sugar_row = summary_gap[
    summary_gap['primary_category'] == 'Chips & Savory Snacks'
].iloc[0]
ax1.axvline(
    x=best_sugar_row['sugar_gap_display'],
    color='#1D3557', linewidth=1.5, linestyle='--', alpha=0.6
)
ax1.text(
    best_sugar_row['sugar_gap_display'] + 0.3,
    len(categories) - 0.7,
    'Chips benchmark',
    fontsize=8, color='#1D3557'
)

# — RIGHT PANEL: Protein gap (how much protein to add) —
ax2 = axes[1]

colors_protein = [
    '#E63946' if g > 10 else '#F4A261' if g > 5 else '#52B788'
    for g in summary_gap['protein_gap_display']
]

bars2 = ax2.barih(
    x, summary_gap['protein_gap_display'],
    height=bar_h, color=colors_protein,
    edgecolor='white', linewidth=0.6
)

for i, (gap, avg) in enumerate(zip(
    summary_gap['protein_gap_display'],
    summary_gap['avg_protein']
)):
    if gap == 0:
        ax2.text(0.3, i, f"Avg {avg:.1f}g – already at target",
                 va='center', fontsize=8.5, color='#2D6A4F', fontweight='bold')
    else:
        ax2.text(gap + 0.2, i,
                 f"Add {gap:.0f}g (avg now: {avg:.1f}g)",
                 va='center', fontsize=8.5, color="#333")

ax2.set_yticks(x)
ax2.set_yticklabels(categories, fontsize=9.5)
ax2.set_xlabel('Protein increase needed (g per 100g)', fontsize=10)
ax2.set_title(' Protein to Add\nto reach ≥15g target',
              fontsize=11, fontweight='bold', pad=10)
ax2.axvline(x=0, color='black', linewidth=1)

```

```

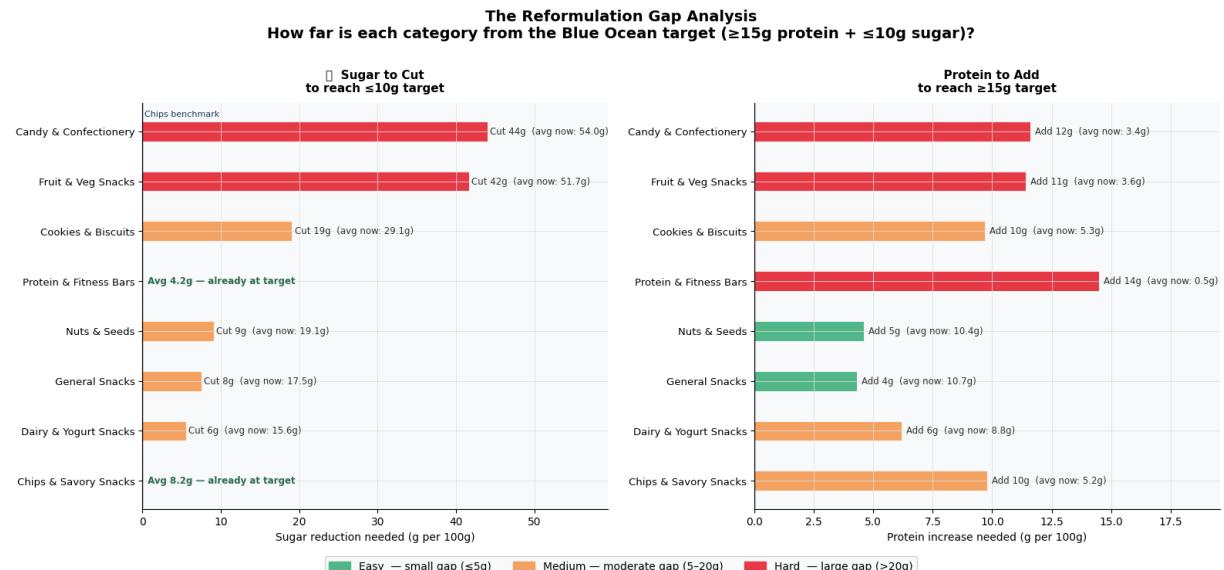
ax2.set_xlim(0, summary_gap['protein_gap_display'].max() * 1.35)

# Shared legend for color coding
legend_patches_easy = [
    mpatches.Patch(color='#52B788', label='Easy – small gap ( $\leq 5g$ )'),
    mpatches.Patch(color='#F4A261', label='Medium – moderate gap (5– $>20g$ )'),
    mpatches.Patch(color='#E63946', label='Hard – large gap ( $>20g$ )')
]
fig.legend(
    handles=legend_patches_easy,
    loc='lower center',
    ncol=3,
    fontsize=10,
    framealpha=0.9,
    bbox_to_anchor=(0.5, -0.05)
)

fig.text(
    0.5, -0.08,
    'Source: Open Food Facts | Gaps calculated from category average already meets this target',
    ha='center', fontsize=8, color="#888", style='italic'
)

plt.tight_layout()
plt.savefig('reformulation_gap.png', dpi=180, bbox_inches='tight')
plt.show()
print("Reformulation gap chart saved as reformulation_gap.png")

```



Reformulation gap chart saved as reformulation\_gap.png

